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Volume 2

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### Volume 2

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<b>Version</b>	<b>Revision History</b>	<b>Year</b>	<b>Part Number</b>
1.0	Original Issue	1984	STLB-USM-PERFCT-1.0
1.1	Fixed bugs and added significant changes to functionality.	1989	STLB-USM-PERFCT-EN8901-1.1
2.0	Added routines to enhance functionality	1991	STLB-USM-PERFCT-EN9109- 2.0
3.0	No changes were made / reprint only	1994	Vol. 1 - 5114A, Vol. 2- 5115A

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# Chapter 8: Time Series Analysis and Forecasting

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## Usage Notes

The name of a time series routine is a combination of three sets of one or two letters. The first set specifies the type of model. The second set identifies the particular method. The final set specifies the general procedure. The table below summarizes the naming convention of the time series analysis and forecasting routines.

The names and meanings of arguments are consistent within a set of routines pertaining to a particular topic. For example, `XCNTR` corresponds to the constant used to center the time series `x` in all of the spectral analysis routines. Note that `IPRINT` always represents the printing option, the values and possible choices of output necessarily depend on the given routine. An option argument always begins with the letter “I,” and a leading dimension argument always begins with “LD.”

The routines in this chapter assume the time series does not contain any missing observations. If missing values are present, they should be set to `NaN` (see the routine `AMACH`, page 1334), and the routine will return an appropriate error message. To enable fitting of the model, the missing values must be replaced by appropriate estimates.

Naming Conventions in Chapter 8	
Meaning	Abbreviation
Nonseasonal ARMA	NS*
Transfer Function	TF*
Multichannel	M*
Periodogram	P*
Cross Periodogram	CP*
Spectral Density	S*
Cross-Spectral Density	CS*
Preliminary	*P*
Method of Moments	*MM*
Least-Squares	*LS*
Box-Jenkins	*BJ*
Spectral Window	*SW*
Weights	*WE*
Estimation	*E
Forecast	*F
Fast Fourier Transform	*FFT
Periodogram	*P
Data	*D

The “\*” represents one or more letters.

## General Methodology

A major component of the model identification step concerns determining if a given time series is stationary. The sample correlation functions computed by routines *ACF* (page 637), *PACF* (page 641), *CCF* (page 644), and *MCCF* (page 649) may be used to diagnose the presence of nonstationarity in the data, as well as to indicate the type of transformation require to induce stationarity. The family of power transformations provided by routine *BCTR* (page 629) coupled with the ability to difference the transformed data using routine *DIFF* (page 633) affords a convenient method of transforming a wide class of nonstationary time series to stationarity.

The “raw” data, transformed data, and sample correlation functions also provide insight into the nature of the underlying model. Typically, this information is displayed in graphical form via time series plots, plots of the lagged data, and various correlation function plots. The routines in Chapter 16, “Line Printer Graphics,” provide the necessary tools to produce the visual displays of this quantitative information.

The observed time series may also be compared with time series generated from various theoretical models to help identify possible candidates for model fitting. The routine *RNARM* (page 1232) in Chapter 18, “Random Number Generation” may be used to generate a time series according to a specified autoregressive moving average model.



## Time Domain Methodology

Once the data are transformed to stationarity, a tentative model in the time domain is often proposed and parameter estimation, diagnostic checking and forecasting are performed.

### Autoregressive Moving Average Model

A parsimonious, yet comprehensive, class of stationary time series models consists of the nonseasonal autoregressive moving average (ARMA) processes defined by

$$\phi(B)(W_t - \mu) = \theta(B)A_t \quad t \in ZZ$$

where

$$ZZ = \{ \dots, -2, -1, 0, 1, 2, \dots \}$$

denotes the set of integers,  $B$  is the backward shift operator defined by  $B^k W_t = W_{t-k}$ ,  $\mu$  is the mean of  $W_t$ ,

$$\phi(B) = 1 - \phi_1 B - \phi_2 B^2 - \dots - \phi_p B^p \quad p \geq 0$$

$$\theta(B) = 1 - \theta_1 B - \theta_2 B^2 - \dots - \theta_q B^q \quad q \geq 0$$

The model is of order  $(p, q)$  and is referred to as an ARMA( $p, q$ ) model.

An equivalent version of the ARMA( $p, q$ ) model is given by

$$\phi(B)W_t = \theta_0 + \theta(B)A_t \quad t \in ZZ$$

where  $\theta_0$  is an overall constant defined by

$$\theta_0 = \mu \left( 1 - \sum_{i=1}^p \phi_i \right)$$

See Box and Jenkins (1976, pages 92–93) for a discussion of the meaning and usefulness of the overall constant.

If the “raw” data  $\{Z_t\}$  are homogeneous nonstationary, then differencing induces stationarity and the model is called autoregressive *integrated* moving average (ARIMA). Parameter estimation is performed on the stationary time series

$$W_t = \nabla^d Z_t$$

where

$$\nabla^d = (1 - B)^d$$

is the backward difference operator with period 1 and order  $d, d > 0$ .

Typically, routine NSPE (page 664) is first applied to the transformed data to provide preliminary parameter estimates. These estimates are used as initial

values in an estimation procedure. In particular, routine `NSLSE` (page 669) may be used to compute conditional or unconditional least-squares estimates of the parameters, depending on the choice of the backcasting length. Parameter estimates from either `NSPE` or `NSLSE` may be input to routine `NSBJF` (page 680) to produce forecasts with associated probability limits. The routines for preliminary parameter estimation, least squares parameter estimation, and forecasting follow the approach of Box and Jenkins (1976, programs 2–4, pages 498–509).

### Transfer Function Model

Define  $\{x_t\}$  and  $\{y_t\}$  by

$$x_t = \begin{cases} X_t - \hat{\mu}_X & d = 0 \\ \nabla^d X_t & d > 0 \end{cases}$$

and

$$y_t = \begin{cases} Y_t - \hat{\mu}_Y & d = 0 \\ \nabla^d Y_t & d > 0 \end{cases}$$

where  $\{X_t\}$  and  $\{Y_t\}$  for  $t = (-d + 1), \dots, n$  represent the undifferenced input and undifferenced output series with

$$\hat{\mu}_X \text{ and } \hat{\mu}_Y$$

estimates of their respective means. The differenced input and differenced output series may be obtained using the routine `DIFF` (page 633) following any preliminary transformation of the data.

The transfer function model is defined by

$$Y_t = \delta^{-1}(B)\omega(B)X_{t-b}$$

or equivalently,

$$y_t = \delta^{-1}(B)\omega(B)x_{t-b} + n_t$$

where  $n_t = \nabla^d N_t$  for  $d \geq 0$ , and the left-hand side and right-hand side transfer function polynomial operators are, respectively,

$$\delta(B) = 1 - \delta_1 B - \delta_2 B^2 - \dots - \delta_r B^r$$

$$\omega(B) = \omega_0 - \omega_1 B - \omega_2 B^2 - \dots - \omega_s B^s$$

with  $r \geq 0$ ,  $s \geq 0$ , and  $b \geq 0$ . The noise process  $\{N_t\}$  and the input process  $\{X_t\}$  are assumed to be independent, with the noise process given by the ARIMA model

$$\phi(B)n_t = \theta(B)A_t$$

where

$$\phi(B) = 1 - \phi_1 B - \phi_2 B^2 - \dots - \phi_p B^p$$

$$\theta(B) = 1 - \theta_1 B - \theta_2 B^2 - \dots - \theta_q B^q$$

with  $p \geq 0$  and  $q \geq 0$ .

The impulse response weights  $\{v_k\}$  of the transfer function

$$v(B) = \delta^{-1}(B)\omega(B) = v_0 + v_1 B + v_2 B^2 + \dots$$

and the differenced noise series  $\{n_t\}$  are estimated using routine `IRNSE` (page 685). Preliminary estimates of the transfer function parameters and noise model parameters are computed by routine `TFPE` (page 689).

### Multichannel Time Series

A multichannel time series  $X$  is simply a multivariate time series whose channels correspond to interrelated univariate time series. In this setting, the model-building process is a logical extension of the procedures used to identify, estimate, and forecast univariate time series. In particular, the multichannel cross-correlation function computed by routine `MCCF` (page 649) may help identify a tentative model. A particular regression model may be fit using routine `MLSE` (page 694), with the Wiener filter estimated using routine `MWFE` (page 700). The Wiener forecast function for a single channel may be obtained by routine `SPWF` (page 677). The state space approach to fitting many time domain models is available through routine `KALMN` (page 705).

### Frequency Domain Methodology

An alternative method of time series analysis with much less emphasis on the form of the model may be performed in the frequency domain.

#### Spectral Analysis

Let  $\{X(t)\}$  denote a continuous-parameter stationary process with mean

$$\mu = E[X(t)]$$

and autocovariance function

$$\sigma(k) = \text{cov}\{X(t), X(t+k)\} = E\{[X(t) - \mu][X(t+k) - \mu]\} \quad k \in \mathbf{R}$$

Similarly, let  $\{X_t\}$  denote a discrete-parameter stationary process with mean

$$\mu = E[X_t]$$

and autocovariance function

$$\sigma(k) = \text{cov}\{X_t, X_{t+k}\} = E\{[X_t - \mu][X_{t+k} - \mu]\} \quad k \in \mathbf{ZZ}$$

Note that  $\sigma(0) = \sigma^2$  is the variance of the process.

The routines for the spectral analysis of time series are concerned with the estimation of the spectral density of a stationary process given a finite realization  $\{X_t\}$  for  $t = 1, \dots, n$  where  $n = \text{NOBS}$ . This realization consists of values sampled at equally spaced time intervals in the continuous-parameter case or of values observed consecutively in the discrete-parameter case. Hence, we need only develop methodology concerned with the spectral analysis of discrete-parameter stationary processes and later account for the time sampling in the continuous-parameter model.

The nonnormalized spectral density  $h(\omega)$  and the autocovariance function  $\sigma(k)$  of the stationary process form a Fourier transform pair. The relationship in the continuous-parameter case is given by

$$h(\omega) = \frac{1}{2\pi} \int_{-\infty}^{\infty} \sigma(k) e^{-i\omega k} dk$$

$$\sigma(k) = \int_{-\pi}^{\pi} h(\omega) e^{i\omega k} d\omega$$

Similarly, the normalized spectral density  $f(\omega)$  and the autocorrelation function  $\rho(k) = \sigma(k)/\sigma(0)$  of the stationary process form a Fourier transform pair. The relationship in the continuous-parameter case is given by

$$f(\omega) = \frac{h(\omega)}{\sigma(0)}$$

$$= \frac{1}{2\pi} \int_{-\infty}^{\infty} \rho(k) e^{-i\omega k} dk$$

$$\rho(k) = \int_{-\pi}^{\pi} f(\omega) e^{i\omega k} d\omega$$

The discrete-parameter analogs of the above equations involve summation over  $k$  instead of integration over  $dk$ . Also, the normalized spectral density  $f(\omega)$  satisfies

$$\int_{-\pi}^{\pi} f(\omega) d\omega = 1$$

**Discrete Fourier Transform.** The discrete Fourier transform of the sequence  $\{Z_t\}$  for  $t = 1, \dots, N$  is defined by

$$\zeta(\omega_p) = \sum_{t=1}^N Z_t e^{-i\omega_p t}$$

over the discrete set of frequencies

$$\omega_p = \frac{2\pi p}{N} \quad p = 0, \pm 1, \dots, \pm \lfloor N/2 \rfloor$$

where the function  $\lfloor r \rfloor$  determines the greatest integer less than or equal to  $r$ . An alternative representation of  $\zeta(\omega_p)$  in terms of cosine and sine transforms is

$$\zeta(\omega_p) = \alpha(\omega_p) - i\beta(\omega_p)$$

where

$$\alpha(\omega_p) = \sum_{t=1}^N Z_t \cos(\omega_p t)$$

$$\beta(\omega_p) = \sum_{t=1}^N Z_t \sin(\omega_p t)$$

The fast Fourier transform algorithm implemented in the IMSL MATH/LIBRARY routine `FFTCF` is used to compute the discrete Fourier transform. All of the frequency domain routines that output a periodogram utilize the fast Fourier transform algorithm.

**Centering and Padding.** Consider the centered and padded realization

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  defined by

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu} & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases} \quad (1)$$

where  $N = (n + n_0)$  and

$$\hat{\mu} = \text{XCNTR}$$

is

$$\hat{\mu} = \begin{cases} \mu & \mu \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t & \mu \text{ unknown} \end{cases} \quad (2)$$

Centering the data simplifies the formulas for estimation of the periodogram and spectral density. The addition of  $n_0 = \text{NPAD}$  zeros to the end of the data is called *padding*. This procedure increases the effective length of the data from  $n$  to  $N$  in an effort to

- increase the computational efficiency of the Fourier transformation of the series by providing a more suitable series length  $N$  (Priestley 1981, page 577).
- obtain the periodogram ordinates required to give the exact expression of the sample autocovariances in terms of the inverse Fourier transformation of the periodogram (Priestley 1981, page 579).
- produce periodogram ordinates over a more refined range of frequencies  $\omega_p$ .

Any desired filtering, prewhitening, or data tapering should be performed prior to estimating the spectral density. The resulting estimate may be adjusted accordingly.

**Periodogram.** The periodogram of the sample sequence  $\{X_t\}$ ,  $t = 1, \dots, n$  computed with the centered and padded sequence

$$\{\tilde{X}_t\}, \quad t = 1, \dots, N$$

is defined by

$$I_{n,N,\tilde{X}}(\omega_p) = K \left| \sum_{t=1}^N \tilde{X}_t e^{-i\omega_p t} \right|^2 = K \left| \zeta_{\tilde{X}}(\omega_p) \right|^2$$

where  $K$  is the scale factor

$$K = \begin{cases} \frac{2}{n} & \text{for the usual periodogram} \\ \frac{1}{2\pi n} & \text{for the modified periodogram} \end{cases}$$

The scale factor of the usual periodogram relates the ordinates to the sum of squares of

$$X_t - \hat{\mu}$$

(Fuller 1976, pages 276–277). If the first ordinate (corresponding to  $p = 0$ ) is replaced by one-half of its value, then if  $N$  is odd, the sum of the  $\lfloor N/2 \rfloor + 1$  ordinates corresponding to  $p = 0, 1, \dots, \lfloor N/2 \rfloor$  is

$$\frac{N}{n} \sum_{t=1}^n (X_t - \hat{\mu})^2$$

The modified periodogram is an asymptotically unbiased estimate of the nonnormalized spectral density function at each frequency  $\omega_p$  (Priestley 1981, page 417). The argument `IPVER` is used to specify the version of the periodogram.

**Spectral Density.** The relationship between the sample autocovariance function and estimate of the nonnormalized spectral density function is similar to the theoretical situation previously discussed.

Define the sample autocovariance function of the  $X_t$  process by

$$\hat{\sigma}(k) = \frac{1}{n} \sum_{t=1}^{n-|k|} \{[X_t - \hat{\mu}][X_{t+|k|} - \hat{\mu}]\} \quad k = 0, \pm 1, \dots, \pm(n-1)$$

where

$$\hat{\mu}$$

is given by Equation 2. Note that

$$\hat{\sigma}(0) = \hat{\sigma}^2$$

is the sample variance. The nonnormalized spectral density may be estimated directly from the sample autocovariances by

$$\hat{h}(\omega) = \frac{1}{2\pi} \sum_{k=-(n-1)}^{(n-1)} \lambda_n(k) \hat{\sigma}(k) e^{-i\omega k}$$

The sequence of weights  $\{\lambda_n(k)\}$  called the *lag window* decreases at a rate appropriate for consistent estimation of  $h(\omega)$ .

An algebraically equivalent method of estimating  $h(\omega)$  consists of locally smoothing the modified periodogram in a neighborhood of  $\omega$ . Let

$$I_{n,N,\tilde{X}}$$

denote the modified periodogram of the centered and padded realization

$$\{\tilde{X}_t\}$$

defined in Equation 1. Then, an estimate of the nonnormalized spectral density is given by

$$\hat{h}(\omega) = \frac{2\pi}{N} \sum_{p=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} I_{n,N,\tilde{X}}(\omega_p) W_n(\omega - \omega_p) \quad (3)$$

where

$$W_n(\theta) = \frac{1}{2\pi} \sum_{k=-(n-1)}^{(n-1)} \lambda_n(k) e^{-i\theta k}$$

The *spectral window*  $W_n(\theta)$  is the discrete Fourier transform of the lag window  $\lambda_n(k)$ . We note that for  $N = 2n - 1$ , the modified periodogram and autocovariances,

$$I_{n,2n-1,\tilde{X}}(\omega_p) \text{ and } \hat{\sigma}(k)$$

form the discrete Fourier transform pair

$$I_{n,N,\tilde{X}}(\omega_p) = \frac{1}{2\pi} \sum_{k=-(n-1)}^{n-1} \hat{\sigma}(k) e^{-i\omega_p k}, \quad p = 0, \pm 1, \dots, \pm(n-1)$$

$$\hat{\sigma}(k) = \frac{2\pi}{N} \sum_{p=-(n-1)}^{n-1} I_{n,N,\tilde{X}}(\omega_p) e^{i\omega_p k}, \quad k = 0, \pm 1, \dots, \pm(n-1)$$

This relationship is exact and recovers the  $(n - 1)$  sample autocovariances only when  $n_0 = (n - 1)$  zeros are padded, since then  $\lfloor N/2 \rfloor = (n - 1)$ .

Another method of estimating  $h(\omega)$  is given by

$$\hat{h}(\omega) = \sum_j w_j I_{n,N,\tilde{X}}(\omega_{p,j}) \quad (4)$$

where

$$\omega_{p,j} = \frac{2\pi\{p(\omega) + j\}}{N}$$

and  $p(\omega)$  is the integer such that  $\omega_{p,0}$  is closest to  $\omega$ . The sequence of  $m$  weights  $\{w_j\}$  for  $j = -[m/2], \dots, (m - [m/2] - 1)$  is fixed in the sense that they do not depend on the frequency,  $\omega$ , and satisfy  $\sum_j w_j = 1$ . Priestley (1981, page 581) notes that if we write

$$w_j = \frac{2\pi}{N} W_n(\omega - \omega_{p,j})$$

then Equation 4 and Equation 3 are quite similar except that the weights  $\{w_j\}$  depend on  $\omega$ . In fact, if  $p(\omega) = 0$  and  $m = N$ , these equations are equivalent.

Given estimates

$$\hat{h}(\omega) \text{ and } \hat{\sigma}(0)$$

the estimate of the normalized spectral density is given by

$$\hat{f}(\omega) = \frac{\hat{h}(\omega)}{\hat{\sigma}(0)}$$

This follows directly from the definition of  $f(\omega)$ .

**Spectral Window.** The following spectral windows  $W_n(\theta)$  are available in routines containing the argument `ISWVER`.

#### Modified Bartlett

$$W_n(\theta) = \frac{1}{2\pi M} \left\{ \frac{\sin(M\theta/2)}{\sin(\theta/2)} \right\}^2 = F_M(\theta)$$

where  $F_M(\theta)$  corresponds to the Fejér kernel of order  $M$ .

#### Daniell

$$W_n(\theta) = \begin{cases} M/2\pi & -\pi/M \leq \theta \leq \pi/M \\ 0 & \text{otherwise} \end{cases}$$

#### Tukey

$$W_n(\theta) = aD_M\left(\theta - \frac{\pi}{M}\right) + (1 - 2a)D_M(\theta) + aD_M\left(\theta + \frac{\pi}{M}\right)$$



for  $0 < a \leq 0.25$ , where  $D_M(\theta)$  represents the Dirichlet kernel. The Tukey-Hanning window is obtained when  $a = 0.23$ , and the Tukey-Hamming window is obtained when  $a = 0.25$ .

### Parzen

$$W_n(\theta) = \frac{6\pi}{M} [F_{M/2}(\theta)]^2 \left\{ 1 - \frac{2}{3} \sin^2(\theta/2) \right\}$$

where  $M$  is even. If  $M$  is odd, then  $M + 1$  is used instead of  $M$  in the above formula.

### Bartlett-Priestley

$$W_n(\theta) = \begin{cases} \frac{3M}{4\pi} \left\{ 1 - \left( \frac{M\theta}{\pi} \right)^2 \right\} & |\theta| \leq \pi/M \\ 0 & |\theta| > \pi/M \end{cases}$$

The *window parameter*  $M$  is inversely proportional to the *bandwidth* of the spectral window. Priestley (1981, pages 520–522) discusses a number of definitions of bandwidth and concludes that the particular definition adopted is of little significance. The choice of spectral window bandwidth, and hence, the choice of  $M$ , is a more important problem. One practical choice for  $M$  is the last lag at which the estimated autocorrelation function

$$\hat{\rho}(k)$$

is significantly different from zero, i.e.,

$$\hat{\rho}(k) \approx 0 \text{ for } k > M$$

The estimated autocorrelations and their associated estimated standard errors can be computed using routine `ACF` (page 637). See Priestley (1981, pages 528–556) for alternative strategies of determining the window parameter  $M$ .

Since the spectral window is the Fourier transform of the lag window, we estimate the spectral density function by application of a particular spectral window to the periodogram. Note that  $M$  is directly related to the rate of decay of the lag window.

**Time Interval.** Consider the continuous-parameter stationary process  $\{X(t)\}$  and let  $\{X_t\}$  denote a realization of this process sampled at equal time intervals  $\Delta t = \text{TINT}$ . Although the spectral density of  $X(t)$  extends over the frequency range  $(-\pi, \pi)$ , the spectral density of  $X_t$  is unique over the restricted frequency range  $(-\pi/\Delta t, \pi/\Delta t)$ . This problem of aliasing or *spectrum folding* is inherent to spectral analysis, see Blackman and Tukey (1958) and Priestley (1981) for further discussion.

In practice, the  $\{X_t\}$  realization is treated as a discrete parameter process with spectral density

$$h_X^\dagger(\omega)$$

defined over the frequency range  $(-\pi, \pi)$ . This corresponds to setting  $\Delta t = 1$ . The transformation of the spectral density to the restricted frequency range  $(-\pi/\Delta t, \pi/\Delta t)$  is given by

$$h_X(\omega) = \Delta t h_X^\dagger(\omega \Delta t) \quad |\omega| \leq \pi / \Delta t$$

Priestley (1981, pages 507–508) considers a method of choosing  $\Delta t$ . A similar transformation is performed for the estimated spectral density.

**Frequency Scale.** The argument `IFSCAL` is used to specify the scale of the frequencies at which to estimate the spectral density. The `NF` frequencies are contained in the argument `F`.

**Approximate Confidence Intervals for Spectral Ordinates.** An approximate  $(1 - \alpha)100\%$  confidence interval for the value of the nonnormalized spectral density function  $h(\omega)$  at a particular frequency  $\omega$  is given by the formula (Priestley 1981, page 468)

$$\left( \frac{DF \times \hat{h}(\omega)}{\chi_{DF,1-\alpha/2}^2}, \frac{DF \times \hat{h}(\omega)}{\chi_{DF,\alpha/2}^2} \right)$$

Routine `CHIIN` (page 1132) using argument `P = 1 - \alpha/2` and `P = \alpha/2` can be used to compute the percentage point

$$\chi_{DF,P}^2$$

Also, routine `CHIIN` should be used with degrees of freedom (`DF`), which depend upon the version of the spectral window (`ISWVER`), as given in the following table (Priestley 1981, page 467).

<code>ISWVER</code>	Window	<code>DF</code>
1	Modified Bartlett	$3n/M$
2	Daniell	$2n/M$
3	Tukey-Hamming	$2.5164n/M$
4	Tukey-Hanning	$2.2/3n/M$
5	Parzen	$3.708614n/M$
6	Bartlett-Priestley	$1.4n/M$

If one of the windows above is not specified and the user provides relative weights, such as with routine `SWED` (page 741), the weights are normalized to sum to one in the actual computations. Given all  $m$  ( $m$  odd) normalized weights  $w_j$ , then for  $2\pi \lfloor m/2 \rfloor / n < \omega < \pi(1 - 2 \lfloor m/2 \rfloor / n)$  the degrees of freedom for a confidence interval on  $h(\omega)$  are given by Fuller (1976, page 296)

$$DF = \frac{2}{\sum_{j=-\lfloor m/2 \rfloor}^{\lfloor m/2 \rfloor} w_j^2}$$

Frequently, confidence intervals on the  $\ln h(\omega)$  are suggested because this produces fixed width intervals. The interval is

$$\left( \ln \hat{h}(\omega) + \ln \left[ \frac{DF}{\chi_{DF, 1-\alpha/2}^2} \right], \ln \hat{h}(\omega) + \ln \left[ \frac{DF}{\chi_{DF, \alpha/2}^2} \right] \right)$$

### Cross-Spectral Analysis

The routines for cross-spectral analysis are concerned with the estimation of the crossspectral density of two jointly stationary processes given finite realizations  $\{X_t\}$  and  $\{Y_t\}$  for  $t = 1, \dots, n$ . These realizations consist of values sampled at equally spaced time intervals in the continuous-parameter case or of values observed consecutively in the discreteparameter case. Again, we develop methodology concerned with the cross-spectral analysis of discrete-parameter stationary processes and later account for the time sampling in the continuous-parameter model.

Let  $\mu_X$  and  $\sigma_{XX}(k)$  denote the mean and autocovariance function of the  $X_t$  process; similarly, define  $\mu_Y$  and  $\sigma_{YY}(k)$ , with respect to the  $Y_t$  process. Define the cross-covariance function between  $X_t$  and  $Y_t$  by

$$\sigma_{XY}(k) = \text{cov}\{[X_t - \mu_X][Y_{t+k} - \mu_Y]\} \quad k \in \mathbb{Z}$$

Then, the nonnormalized cross-spectral density  $h_{XY}(\omega)$  and the cross-covariance function  $\sigma_{XY}(k)$  form a Fourier transform pair. The relationship in the continuous-parameter case is given by

$$h_{XY}(\omega) = \frac{1}{2\pi} \int_{-\infty}^{\infty} \sigma_{XY}(k) e^{-i\omega k} dk$$

$$\sigma_{XY}(k) = \int_{-\pi}^{\pi} h_{XY}(\omega) e^{i\omega k} d\omega$$

Similarly, the normalized cross-spectral density  $f_{XY}(\omega)$  and the cross-correlation function  $\rho_{XY}(k) = \sigma_{XY}(k)/[\sigma_{XX}(0)\sigma_{YY}(0)]$  form a Fourier transform pair. The relationship in the continuous-parameter case is given by

$$f_{XY}(\omega) = \frac{h_{XY}(\omega)}{\sigma_{XX}(0)\sigma_{YY}(0)} = \frac{1}{2\pi} \int_{-\infty}^{\infty} \rho_{XY}(k) e^{-i\omega k} dk$$

$$\rho_{XY}(k) = \int_{-\pi}^{\pi} f_{XY}(\omega) e^{i\omega k} d\omega$$

The discrete-parameter analogs of the above equations involve summation over  $k$  instead of integration over  $dk$ .

The cross-spectral density function is often written in terms of real and imaginary components, since in general, the function is complex-valued. In particular,

$$h_{XY}(\omega) = c_{XY}(\omega) - iq_{XY}(\omega)$$

where the *cospectrum* and *quadrature spectrum* of the  $X_t$  and  $Y_t$  process are respectively defined by

$$c_{XY}(\omega) = \frac{1}{2\pi} \int_{-\infty}^{\infty} \frac{1}{2} [\sigma_{XY}(k) + \sigma_{XY}(-k)] \cos k\omega \, dk$$

$$q_{XY}(\omega) = \frac{1}{2\pi} \int_{-\infty}^{\infty} \frac{1}{2} [\sigma_{XY}(k) - \sigma_{XY}(-k)] \sin k\omega \, dk$$

The *polar form* of  $h_{XY}(\omega)$  is defined by

$$h_{XY}(\omega) = \alpha_{XY}(\omega) e^{i\phi_{XY}(\omega)}$$

where the *cross-amplitude spectrum* is

$$\alpha_{XY}(\omega) = |h_{XY}(\omega)| = [c_{XY}^2(\omega) + q_{XY}^2(\omega)]^{1/2}$$

and the *phase spectrum* is

$$\phi_{XY}(\omega) = \tan^{-1}[-q_{XY}(\omega) / c_{XY}(\omega)]$$

The *coherency spectrum* is defined by

$$w_{XY}(\omega) = \frac{h_{XY}(\omega)}{[h_{XX}(\omega)h_{YY}(\omega)]^{1/2}}$$

For a given frequency  $\omega$ , the *coherency*  $|w_{XY}(\omega)|$  lies between zero and one, inclusive, and reflects the linear relationship between the random coefficients. See Priestley (1981, pages 654–661) for additional information concerning the interpretation of the components of the cross-spectral density.

**Centering and Padding.** The centered and padded realizations

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

are defined as in Equation 1 with centering constants

$$\hat{\mu}_X = \text{XCNTR} \text{ and } \hat{\mu}_Y = \text{YCNTR}$$

Any desired filtering, prewhitening, or data tapering should be performed prior to estimating the crossspectral density. The resulting estimate may be adjusted accordingly.

**Cross Periodogram.** The cross periodogram of the sample sequences  $\{X_t\}$  and  $\{Y_t\}$ ,  $t = 1, \dots, n$  computed with the padded sequences

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

$t = 1, \dots, N$  is defined by

$$I_{n,N,\tilde{X}\tilde{Y}}(\omega_p) = \left( \sum_{t=1}^N \tilde{X}_t e^{-i\omega_p t} \right) \left( \sum_{t=1}^N \tilde{Y}_t e^{i\omega_p t} \right) = K \zeta_{\tilde{X}}(\omega_p) \zeta_{\tilde{Y}}^*(\omega_p)$$

where  $K$  is the scale factor

$$K = \begin{cases} \frac{2}{n} & \text{for the usual cross periodogram} \\ \frac{1}{2\pi n} & \text{for the modified cross periodogram} \end{cases}$$

The scale factor option is maintained for compatibility with the spectral routines. The argument `IPVER` is used to specify the version of the periodogram used to compute the cross periodogram.

**Cross-Spectral Density Estimation.** The relationship between the sample cross-covariance function and estimate of the nonnormalized cross-spectral density function is similar to the theoretical situation previously discussed.

Define the sample cross-covariance function between the  $X_t$  and  $Y_t$  process by

$$\tilde{\sigma}_{XY}(k) = \begin{cases} \frac{1}{n} \sum_{t=1}^{n-k} \{ [X_t - \hat{\mu}_X][Y_{t+k} - \hat{\mu}_Y] \} & k = 0, 1, \dots, (n-1) \\ \frac{1}{n} \sum_{t=1-k}^n \{ [X_t - \hat{\mu}_X][Y_{t+k} - \hat{\mu}_Y] \} & k = -1, -2, \dots, -(n-1) \end{cases}$$

The nonnormalized cross-spectral density may be estimated directly from the sample cross-covariances by

$$\tilde{h}_{XY}(\omega) = \frac{1}{2\pi} \sum_{k=-(n-1)}^{(n-1)} \lambda_n(k) \hat{\sigma}_{XY}(k) e^{-i\omega k}$$

The sequence of weights  $\{\lambda_n(k)\}$  called the *lag window* decreases at a rate appropriate for consistent estimation of  $h_{XY}(\omega)$ .

An algebraically equivalent method of estimating  $h_{XY}(\omega)$ , consists of locally smoothing the modified cross periodogram in a neighborhood of  $\omega$ . Let

$$I_{n,N,\tilde{X}\tilde{Y}}$$

denote the modified cross periodogram of the centered and padded realizations

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

Then, an estimate of the nonnormalized cross-spectral density is given by

$$\hat{h}_{XY}(\omega) = \frac{2\pi}{N} \sum_{p=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} I_{n,N,\tilde{X}\tilde{Y}}(\omega_p) W_n(\omega - \omega_p) \quad (5)$$

where  $W_n(\theta)$  is the spectral window.

Another method of estimating  $h_{XY}(\omega)$  is given by

$$\hat{h}_{XY}(\omega) = \sum_j w_j I_{n,N,\tilde{X}\tilde{Y}}(\omega_{p,j}) \quad (6)$$

where  $\omega_{p,j}$ ,  $p(\omega)$ , and the weights  $\{w_j\}$  are as defined in the univariate setting.

Given estimates

$$\hat{h}_{XY}(\omega), \hat{\sigma}_{XX}(0), \text{ and } \hat{\sigma}_{YY}(0)$$

the estimate of the normalized cross-spectral density is given by

$$\hat{f}_{XY}(\omega) = \frac{\hat{h}_{XY}(\omega)}{\hat{\sigma}_{XX}(0)\hat{\sigma}_{YY}(0)}$$

This follows directly from the definition of  $f_{XY}(\omega)$ .

## BCTR/DBCTR (Single/Double precision)

Perform a forward or an inverse Box-Cox (power) transformation.

### Usage

CALL BCTR (NOBS, Z, IPRINT, IDIR, POWER, SHIFT, X)

### Arguments

**NOBS** — Number of observations in Z. (Input)

NOBS must be greater than or equal to one.

**Z** — Vector of length NOBS containing the data. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.

1 Print Z and the transformed data, X.

**IDIR** — Direction of transformation option. (Input)

#### IDIR Action

0 Forward transformation.

1 Inverse transformation.

**POWER** — Exponent parameter in the power transformation. (Input)

**SHIFT** — Shift parameter in the power transformation. (Input)

SHIFT must satisfy the relation  $\min(z(i)) + \text{SHIFT} > 0$ .

**X** — Vector of length NOBS containing the transformed data. (Output)

If Z is not needed, then X and Z can occupy the same storage locations. In this case, IPRINT = 1 will print two identical vectors.

### Comments

1. Informational errors

Type	Code	
4	1	For the specified forward transformation, the minimum element of X will underflow.
4	2	For the specified forward transformation, the maximum element of X will overflow.
4	3	For the specified inverse transformation, the maximum element of X will overflow.
4	4	For the specified inverse transformation, the minimum element of X will underflow.

2. The forward transformation is performed prior to fitting a model. Differencing of the data is done after the data are transformed.

3. The inverse transformation is performed on results such as forecasts and their corresponding probability limits.

### Algorithm

Routine BCTR performs a forward or inverse Box-Cox transformation of the  $n = \text{NOBS}$  observations  $\{Z_t\}$  for  $t = 1, 2, \dots, n$ .

The forward transformation is useful in the analysis of linear models or models with nonnormal errors or nonconstant variance (Draper and Smith 1981, page 222). In the time series setting, application of the appropriate transformation and subsequent differencing of a series may enable model identification and parameter estimation in the class of homogeneous stationary autoregressive-moving average models. The inverse transformation may later be applied to certain results of the analysis, such as forecasts and probability limits of forecasts, in order to express the results in the scale of the original data. A brief note concerning the choice of transformations in ARIMA models is given in Box and Jenkins (1976, page 328). The class of power transformations discussed by Box and Cox (1964) is defined by

$$X_t = \begin{cases} \frac{(Z_t + \xi)^\lambda - 1}{\lambda} & \lambda \neq 0 \\ \ln(Z_t + \xi) & \lambda = 0 \end{cases}$$

where  $Z_t + \xi > 0$  for all  $t$ . Since

$$\lim_{\lambda \rightarrow 0} \frac{(Z_t + \xi)^\lambda - 1}{\lambda} = \ln(Z_t + \xi)$$

the family of power transformations is continuous.

Let  $\lambda = \text{POWER}$  and  $\xi = \text{SHIFT}$ ; then, the computational formula utilized by routine BCTR is given by

$$X_t = \begin{cases} (Z_t + \xi)^\lambda & \lambda \neq 0 \\ \ln(Z_t + \xi) & \lambda = 0 \end{cases}$$

where  $Z_t + \xi > 0$  for all  $t$ . The computational and Box-Cox formulas differ only in the scale and the origin of the transformed data. Consequently, the general analysis of the data is unaffected (Draper and Smith 1981, page 225).

The inverse transformation is computed by

$$X_t = \begin{cases} Z_t^{1/\lambda} - \xi & \lambda \neq 0 \\ \exp(Z_t) - \xi & \lambda = 0 \end{cases}$$

where  $\{Z_t\}$  now represents the result computed by BCTR for a forward transformation of the original data using parameters  $\lambda$  and  $\xi$ .

### Example 1

Consider the Airline Data (Box and Jenkins 1976, page 531) consisting of the monthly total number of international airline passengers from January 1949 through December 1960. Routine BCTR is used to compute a forward Box-Cox transformation of the first 12 observations. In the transformation SHIFT and POWER are each set to zero, which corresponds to taking natural logarithms of the data.

```

INTEGER IPRINT, NOBS
PARAMETER (IPRINT=1, NOBS=12)
C
INTEGER      IDIR, NCOL, NROW
REAL         POWER, SHIFT, X(NOBS), Z(144)
EXTERNAL    BCTR, GDATA
C
CALL GDATA (4, 0, NROW, NCOL, Z, 144, 1)
C
IDIR = 0
C
POWER = 0.0
SHIFT = 0.0
C
CALL BCTR (NOBS, Z, IPRINT, IDIR, POWER, SHIFT, X)
C
END

```

### Output

Output from BCTR

I	Z	X
1	112.00	4.7185



2	118.00	4.7707
3	132.00	4.8828
4	129.00	4.8598
5	121.00	4.7958
6	135.00	4.9053
7	148.00	4.9972
8	148.00	4.9972
9	136.00	4.9127
10	119.00	4.7791
11	104.00	4.6444
12	118.00	4.7707

## Example 2

The estimated standard errors of forecasts (lead times 1 through 12 at origin July 1957) using the transformed Airline Data (Box and Jenkins 1976, page 311) may be converted back to their original scale using routine BCTR. The backward Box-Cox transformation with SHIFT and POWER each set to zero corresponds to using the exponential function.

```

INTEGER      NOBS
PARAMETER    (NOBS=12)
C
INTEGER      IDIR, IPRINT
REAL         POWER, SD(NOBS), SHIFT, X(NOBS)
EXTERNAL     BCTR, SSCAL
C
                                Standard errors of forecasts
DATA SD/3.7, 4.3, 4.8, 5.3, 5.8, 6.2, 6.6, 6.9, 7.2, 7.6, 8.0,
&          8.2/
C
CALL SSCAL (NOBS, 1.0E-02, SD, 1)
C
                                Backward direction
IDIR = 1
C
                                Transformation parameters
POWER = 0.0
SHIFT = 0.0
C
                                Transform standard errors from
C
                                log scale to original scale
IPRINT = 1
CALL BCTR (NOBS, SD, IPRINT, IDIR, POWER, SHIFT, X)
C
END

```

## Output

Output from BCTR

I	Z	X
1	0.037000	1.0377
2	0.043000	1.0439
3	0.048000	1.0492
4	0.053000	1.0544
5	0.058000	1.0597
6	0.062000	1.0640
7	0.066000	1.0682
8	0.069000	1.0714
9	0.072000	1.0747
10	0.076000	1.0790

11 0.080000 1.0833  
12 0.082000 1.0855

---

## DIFF/DDIFF (Single/Double precision)

Difference a time series.

### Usage

```
CALL DIFF (NOBSZ, Z, NDIFF, IPER, IORD, IPRINT, IMISS,  
          NLOST, NOBSX, X)
```

### Arguments

**NOBSZ** — Number of observations in the time series *Z*. (Input)  
NOBSZ must be greater than or equal to one.

**Z** — Vector of length NOBSZ containing the time series. (Input)

**NDIFF** — Number of differences to perform. (Input)  
NDIFF must be greater than or equal to one.

**IPER** — Vector of length NDIFF containing the periods at which *Z* is to be differenced. (Input)  
The elements of IPER must be greater than or equal to one.

**IORD** — Vector of length NDIFF containing the order of each difference given in IPER. (Input)  
The elements of IORD must be greater than or equal to zero.

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.  
1 Print the number of observations lost because of differencing *Z*, the number of observations in the differenced series *X*, and the differenced series *X*.

**IMISS** — Missing value option. (Input)

#### IMISS Action

0 Include missing values in *X*.  
1 Exclude missing values from *X*.

**NLOST** — Number of observations lost because of differencing the time series *Z*. (Output)  
$$NLOST = IPER(1) * IORD(1) + \dots + IPER(NDIFF) * IORD(NDIFF).$$

**NOBSX** — Number of observations in the differenced series *X*. (Output)  
$$NOBSX = NOBSZ - IMISS * NLOST.$$

**X** — Vector of length NOBSX containing the differenced series. (Output)

## Comments

- Automatic workspace usage is

DIFF NOBSZ units, or  
DDIFF 2 \* NOBSZ units.

Workspace may be explicitly provided, if desired, by use of D2FF/DD2FF. The reference is

```
CALL D2FF (NOBSZ, Z, NDIFF, IPER, IORD, IPRINT,
           IMISS, NLOST, NOBSX, X, XWK)
```

The additional argument is

**XWK** — Work vector of length equal to NOBSZ.

- A value is considered to be missing if it is not itself in the data set or if it is the result of an operation involving missing value(s). In differencing, missing values occur at the beginning of the differenced series since  $x(i) = z(i) - z(i - k)$  is not defined for  $k$  greater than or equal to  $i$ .

## Algorithm

Routine DIFF performs  $m = \text{NDIFF}$  successive backward differences of period  $s_i = \text{IPER}(i)$  and order  $d_i = \text{IORD}(i)$  for  $i = 1, \dots, m$  on the  $n = \text{NOBSZ}$  observations  $\{Z_t\}$  for  $t = 1, 2, \dots, n$ .

Consider the backward shift operator  $B$  given by

$$B^k Z_t = Z_{t-k}, \text{ for all } k$$

Then, the *backward difference operator* with period  $s$  is defined by

$$\nabla_s Z_t = (1 - B^s) Z_t = Z_t - Z_{t-s}, s \geq 0$$

Note that  $B^s Z_t$  and  $\nabla_s Z_t$  are defined only for  $t = (s + 1), \dots, n$ . Repeated differencing with period  $s$  is simply

$$\nabla_s^d Z_t = (1 - B^s)^d Z_t = \sum_{j=0}^d \frac{d!}{j!(d-j)!} (-1)^j B^{sj} Z_t$$

where  $d \geq 0$  is the order of differencing. Note that

$$\nabla_s^d Z_t$$

is defined only for  $t = (sd + 1), \dots, n$ .

The general difference formula used in routine DIFF is given by

$$X_t = \begin{cases} \text{NaN} & t = 1, \dots, n_L \\ \nabla_{s_1}^{d_1} \nabla_{s_2}^{d_2} \dots \nabla_{s_m}^{d_m} Z_t & t = n_L + 1, \dots, n \end{cases}$$

where  $n_L = \text{NLOST}$  represents the number of observations “lost” because of differencing and NaN (not a number) represents the missing value code. See the routine `AMACH` (page 1334) in the “Machine-Dependent Constants” section of the Reference Material. Note that  $n_L = \sum_j s_j d_j$ .

A homogeneous stationary time series may be arrived at by appropriately differencing a homogeneous nonstationary time series (Box and Jenkins 1976, page 85). Preliminary application of an appropriate transformation followed by differencing of a series may enable model identification and parameter estimation in the class of homogeneous stationary autoregressive-moving average models.

### Example

Consider the Airline Data (Box and Jenkins 1976, page 531) consisting of the monthly total number of international airline passengers from January 1949 through December 1960. Routine `DIFF` is used to compute

$$X_t = \nabla_1 \nabla_{12} X_t = (Z_t - Z_{t-12}) - (Z_{t-1} - Z_{t-13})$$

For the first invocation of `DIFF` with `IMISS = 0`,  $X_1, X_2, \dots, X_{13}$  are set to the missing value code (NaN) and the equation is applied for  $t = 14, 15, \dots, 24$ . For the second invocation of `DIFF` with `IMISS = 1`, the missing values are excluded from the output array containing the differenced series.

```

C      INTEGER      IPRINT, NDIFF, NOBSZ
C      PARAMETER    (IPRINT=1, NDIFF=2, NOBSZ=24)

C      INTEGER      IMISS, IORD(NDIFF), IPER(NDIFF), NCOL, NLOST, NOBSX,
C      &            NROW
C      REAL         X(NOBSZ), Z(144)
C      EXTERNAL     DIFF, GDATA

C                                     Periods of differencing
C      DATA IPER/1, 12/

C                                     Orders of differencing
C      DATA IORD/1, 1/

C                                     Airline Data
C      CALL GDATA (4, 0, NROW, NCOL, Z, 144, 1)
C                                     Nonseasonal and seasonal difference
C                                     first 24 observations in Z
C
C                                     Include missing values in result X
C      IMISS = 0
C      CALL DIFF (NOBSZ, Z, NDIFF, IPER, IORD, IPRINT, IMISS, NLOST,
C      &          NOBSX, X)
C                                     Exclude missing values in result X
C      IMISS = 1
C      CALL DIFF (NOBSZ, Z, NDIFF, IPER, IORD, IPRINT, IMISS, NLOST,
C      &          NOBSX, X)

C      END

```

### Output

Output from `DIFF/D2FF`

NLOST = 13  
NOBSX = 24

I	Z(I)	X(I)
1	112.00	NaN
2	118.00	NaN
3	132.00	NaN
4	129.00	NaN
5	121.00	NaN
6	135.00	NaN
7	148.00	NaN
8	148.00	NaN
9	136.00	NaN
10	119.00	NaN
11	104.00	NaN
12	118.00	NaN
13	115.00	NaN
14	126.00	5.000
15	141.00	1.000
16	135.00	-3.000
17	125.00	-2.000
18	149.00	10.000
19	170.00	8.000
20	170.00	0.000
21	158.00	0.000
22	133.00	-8.000
23	114.00	-4.000
24	140.00	12.000

Output from DIFF/D2FF

NLOST = 13  
NOBSX = 11

I	Z(I)	X(I)
1	112.00	5.000
2	118.00	1.000
3	132.00	-3.000
4	129.00	-2.000
5	121.00	10.000
6	135.00	8.000
7	148.00	0.000
8	148.00	0.000
9	136.00	-8.000
10	119.00	-4.000
11	104.00	12.000
12	118.00	
13	115.00	
14	126.00	
15	141.00	
16	135.00	
17	125.00	
18	149.00	
19	170.00	
20	170.00	
21	158.00	
22	133.00	
23	114.00	
24	140.00	

---

## ACF/DACF (Single/Double precision)

Compute the sample autocorrelation function of a stationary time series.

### Usage

```
CALL ACF (NOBS, X, IPRINT, ISEOPT, IMEAN, XMEAN, MAXLAG,  
         ACV, AC, SEAC)
```

### Arguments

**NOBS** — Number of observations in the time series  $x$ . (Input)

NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

- 0 No printing is performed.
- 1 Print the mean and variance.
- 2 Print the mean, variance, and autocovariances.
- 3 Print the mean, variance, autocovariances, autocorrelations, and standard errors of autocorrelations.

**ISEOPT** — Option for computing standard errors of autocorrelations. (Input)

#### ISEOPT Action

- 0 No standard errors of autocorrelations are computed.
- 1 Compute standard errors of autocorrelations using Bartlett's formula.
- 2 Compute standard errors of autocorrelations using Moran's formula.

**IMEAN** — Option for computing the mean. (Input)

#### IMEAN Action

- 0 XMEAN is user specified.
- 1 XMEAN is set to the arithmetic mean of  $x$ .

**XMEAN** — Estimate of the mean of time series  $x$ . (Input, if IMEAN = 0; output, if IMEAN = 1)

**MAXLAG** — Maximum lag of autocovariances, autocorrelations, and standard errors of autocorrelations to be computed. (Input)

MAXLAG must be greater than or equal to one and less than NOBS.

**ACV** — Vector of length MAXLAG + 1 containing the variance and autocovariances of the time series  $x$ . (Output)

ACV(0) contains the variance of the series  $x$ . ACV( $k$ ) contains the autocovariance of lag  $k$  where  $k = 1, \dots, \text{MAXLAG}$ .

**AC** — Vector of length MAXLAG + 1 containing the autocorrelations of the time series  $x$ . (Output)

AC(0) = 1. AC( $k$ ) contains the autocorrelation of lag  $k$  where  $k = 1, \dots, \text{MAXLAG}$ .

**SEAC** — Vector of length `MAXLAG` containing the standard errors of the autocorrelations of the time series `x`. (Output)

The standard error of  $AC(k)$  is  $SEAC(k)$  where  $k = 1, \dots, \text{MAXLAG}$ . If `ISEOPT = 0`, then **SEAC** may be dimensioned of length 1.

### Algorithm

Routine `ACF` estimates the autocorrelation function of a stationary time series given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  for  $t = 1, 2, \dots, n$ .

Let

$$\hat{\mu} = \text{XMEAN}$$

be the estimate of the mean  $\mu$  of the time series  $\{X_t\}$  where

$$\hat{\mu} = \begin{cases} \mu, & \mu \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t & \mu \text{ unknown} \end{cases}$$

The autocovariance function  $\sigma(k)$  is estimated by

$$\hat{\sigma}(k) = \frac{1}{n} \sum_{t=1}^{n-k} (X_t - \hat{\mu})(X_{t+k} - \hat{\mu}), \quad k = 0, 1, \dots, K$$

where  $K = \text{MAXLAG}$ . Note that

$$\hat{\sigma}(0)$$

is an estimate of the sample variance. The autocorrelation function  $\rho(k)$  is estimated by

$$\hat{\rho}(k) = \frac{\hat{\sigma}(k)}{\hat{\sigma}(0)}, \quad k = 0, 1, \dots, K$$

Note that

$$\hat{\rho}(0) \equiv 1$$

by definition.

The standard errors of the sample autocorrelations may be optionally computed according to argument `ISEOPT`. One method (Bartlett 1946) is based on a general asymptotic expression for the variance of the sample autocorrelation coefficient of a stationary time series with independent, identically distributed normal errors. The theoretical formula is

$$\text{var}\{\hat{\rho}(k)\} = \frac{1}{n} \sum_{i=-\infty}^{\infty} \left[ \rho^2(i) + \rho(i-k)\rho(i+k) - 4\rho(i)\rho(k)\rho(i-k) + 2\rho^2(i)\rho^2(k) \right]$$

where

$$\hat{\rho}(k)$$

assumes  $\mu$  is unknown. For computational purposes, the autocorrelations  $\rho(k)$  are replaced by their estimates

$$\hat{\rho}(k)$$

for  $|k| \leq K$ , and the limits of summation are bounded because of the assumption that  $\rho(k) = 0$  for all  $k$  such that  $|k| > K$ .

A second method (Moran 1947) utilizes an exact formula for the variance of the sample autocorrelation coefficient of a random process with independent, identically distributed normal errors. The theoretical formula is

$$\text{var}\{\hat{\rho}(k)\} = \frac{n-k}{n(n+2)}$$

where  $\mu$  is assumed to be equal to zero. Note that this formula does not depend on the autocorrelation function.

### Example

Consider the Wolfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Routine ACF computes the estimated autocovariances, estimated autocorrelations, and estimated standard errors of the autocorrelations.

```

INTEGER      IPRINT, MAXLAG, NOBS
PARAMETER   ( IPRINT=3, MAXLAG=20, NOBS=100)
C
INTEGER      IMEAN, ISEOPT, NCOL, NROW
REAL         AC(0:MAXLAG), ACV(0:MAXLAG), RDATA(176,2),
&           SEAC(MAXLAG), X(NOBS), XMEAN
EXTERNAL     ACF, GDATA
C
EQUIVALENCE (X(1), RDATA(22,2))
C                                     Wolfer Sunspot Data for
C                                     years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C                                     Compute standard errors
ISEOPT = 1
C                                     Center on arithmetic mean
IMEAN = 1
C                                     Compute sample ACF
CALL ACF (NOBS, X, IPRINT, ISEOPT, IMEAN, XMEAN, MAXLAG, ACV,
&        AC, SEAC)
C
END

```

### Output

Output from ACF/A2F

```

Mean      =      46.976
Variance  =     1382.9

```



Lag	ACV	AC	SEAC
0	1382.9	1.00000	
1	1115.0	0.80629	0.03478
2	592.0	0.42809	0.09624
3	95.3	0.06891	0.15678
4	-236.0	-0.17062	0.20577
5	-370.0	-0.26756	0.23096
6	-294.3	-0.21278	0.22899
7	-60.4	-0.04371	0.20862
8	227.6	0.16460	0.17848
9	458.4	0.33146	0.14573
10	567.8	0.41061	0.13441
11	546.1	0.39491	0.15068
12	398.9	0.28848	0.17435
13	197.8	0.14300	0.19062
14	26.9	0.01945	0.19549
15	-77.3	-0.05588	0.19589
16	-143.7	-0.10394	0.19629
17	-202.0	-0.14610	0.19602
18	-245.4	-0.17743	0.19872
19	-230.8	-0.16691	0.20536
20	-142.9	-0.10332	0.20939

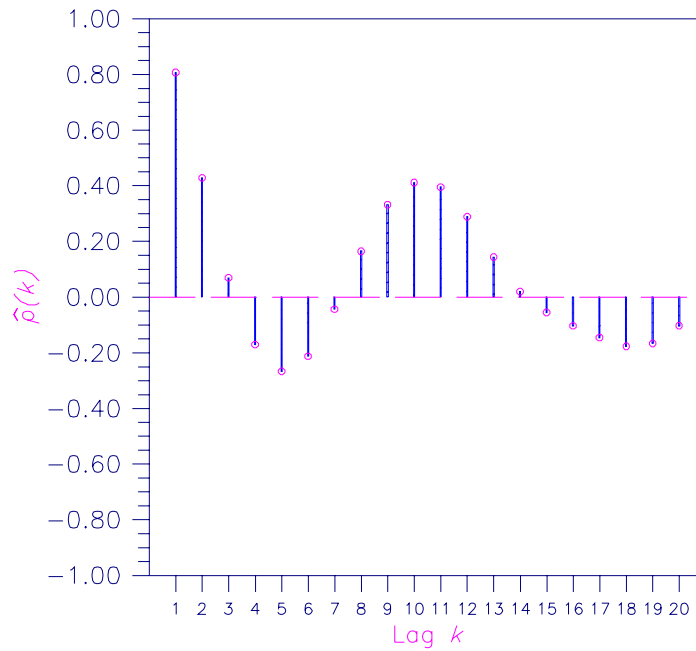


Figure 8-1 Sample Autocorrelation Function

---

## PACF/DPACF (Single/Double precision)

Compute the sample partial autocorrelation function of a stationary time series.

### Usage

CALL PACF (MAXLAG, AC, PAC)

### Arguments

**MAXLAG** — Maximum lag of partial autocorrelations to be computed. (Input)

**AC** — Vector of length MAXLAG+ 1 containing the autocorrelations of the time series  $X$ . (Input)

$AC(0) = 1$ .  $AC(k)$  contains the autocorrelation of lag  $k$  where  $k = 1, \dots, \text{MAXLAG}$ .

**PAC** — Vector of length MAXLAG containing the partial autocorrelations of the time series  $X$ . (Output)

The partial autocorrelation of lag  $k$  corresponds to  $PAC(k)$  where  $k = 1, \dots, \text{MAXLAG}$ .

### Comments

Automatic workspace storage is

PACF 2 \* MAXLAG units, or  
DPACF 4 \* MAXLAG units.

Workspace may be explicitly provided, if desired, by use of P2CF/DP2CF. The reference is

CALL P2CF (MAXLAG, AC, PAC, WK)

The additional argument is

**WK** — Work vector of length 2 \* MAXLAG.

### Algorithm

Routine PACF estimates the partial autocorrelations of a stationary time series given the  $K = \text{MAXLAG}$  sample autocorrelations

$$\hat{\rho}(k)$$

for  $k = 0, 1, \dots, K$ . Consider the AR( $k$ ) process defined by

$$X_t = \phi_{k1} X_{t-1} + \phi_{k2} X_{t-2} + \dots + \phi_{kk} X_{t-k} + A_t$$

where  $\phi_{kj}$  denotes the  $j$ -th coefficient in the process. The set of estimates

$$\{\hat{\phi}_{kk}\}$$

for  $k = 1, \dots, K$  is the sample partial autocorrelation function. The autoregressive parameters

$$\{\hat{\phi}_{kj}\}$$

for  $j = 1, \dots, k$  are approximated by Yule-Walker estimates for successive AR( $k$ ) models where  $k = 1, \dots, K$ . Based on the sample Yule-Walker equations

$$\hat{\rho}(j) = \hat{\phi}_{k1}\hat{\rho}(j-1) + \hat{\phi}_{k2}\hat{\rho}(j-2) + \dots + \hat{\phi}_{kk}\hat{\rho}(j-k), \quad j = 1, 2, \dots, k$$

a recursive relationship for  $k = 1, \dots, K$  was developed by Durbin (1960). The equations are given by

$$\hat{\phi}_{kk} = \begin{cases} \hat{\rho}(1) & k = 1 \\ \frac{\hat{\rho}(k) - \sum_{j=1}^{k-1} \hat{\phi}_{k-1,j} \hat{\rho}(k-j)}{1 - \sum_{j=1}^{k-1} \hat{\phi}_{k-1,j} \hat{\rho}(j)} & k = 2, \dots, K \end{cases}$$

and

$$\hat{\phi}_{kj} = \begin{cases} \hat{\phi}_{k-1,j} - \hat{\phi}_{kk} \hat{\phi}_{k-1,k-j} & j = 1, 2, \dots, k-1 \\ \hat{\phi}_{kk} & j = k \end{cases}$$

This procedure is sensitive to rounding error and should not be used if the parameters are near the nonstationarity boundary. A possible alternative would be to estimate  $\{\phi_{kk}\}$  for successive AR( $k$ ) models using least squares (IMSL routine NSLSE, page 669) or maximum likelihood. Based on the hypothesis that the true process is AR( $p$ ), Box and Jenkins (1976, page 65) note

$$\text{var}\{\hat{\phi}_{kk}\} \simeq \frac{1}{n} \quad k \geq p + 1$$

See Box and Jenkins (1976, pages 82–84) for more information concerning the partial autocorrelation function.

### Example

Consider the Wolfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Routine PACF to used to compute the estimated partial autocorrelations.

```

C      INTEGER      IMEAN, IPRINT, ISEOPT, MAXLAG, NOBS
PARAMETER      (IMEAN=1, IPRINT=0, ISEOPT=0, MAXLAG=20, NOBS=100)

C      INTEGER      NCOL, NROW
REAL           AC(0:MAXLAG), ACV(0:MAXLAG), PAC(MAXLAG),
&             RDATA(176,2), SEAC(1), X(NOBS), XMEAN
CHARACTER     CLABEL(2)*4, RLABEL(1)*6
EXTERNAL      ACF, GDATA, PACF, WRRRL

C      EQUIVALENCE (X(1), RDATA(22,2))

C      DATA RLABEL/'NUMBER'/, CLABEL/'Lag ', 'PACF'/
C                                 Wolfer Sunspot Data for
C                                 years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)

```

```

C          Compute sample ACF
  CALL ACF (NOBS, X, IPRINT, ISEOPT, IMEAN, XMEAN, MAXLAG, ACV,
&          AC, SEAC)
C          Compute sample PACF
  CALL PACF (MAXLAG, AC, PAC)
C          Print results
  CALL WRRRL (' ', 20, 1, PAC, 20, 0, '(F8.3)', RLABEL, CLABEL)
C
  END

```

### Output

Lag	PACF
1	0.806
2	-0.635
3	0.078
4	-0.059
5	-0.001
6	0.172
7	0.109
8	0.110
9	0.079
10	0.079
11	0.069
12	-0.038
13	0.081
14	0.033
15	-0.035
16	-0.131
17	-0.155
18	-0.119
19	-0.016
20	-0.004

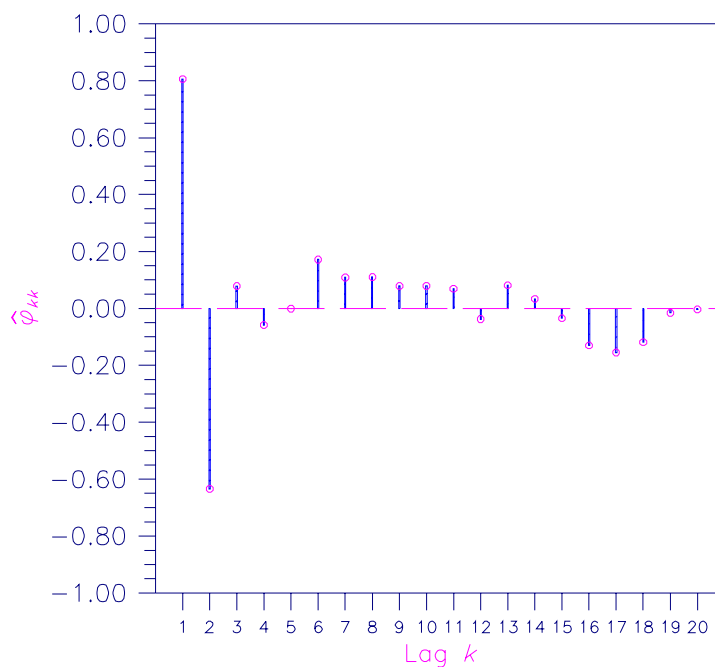


Figure 8-2 Sample Partial Autocorrelation Function

---

## CCF/DCCF (Single/Double precision)

Compute the sample cross-correlation function of two stationary time series.

### Usage

```
CALL CCF (NOBS, X, Y, MAXLAG, IPRINT, ISEOPT, IMEAN, XMEAN,
          YMEAN, XVAR, YVAR, CCV, CC, SECC)
```

### Arguments

**NOBS** — Number of observations in each time series. (Input)  
 NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the first time series. (Input)

**Y** — Vector of length NOBS containing the second time series. (Input)

**MAXLAG** — Maximum lag of cross-covariances and cross-correlations to be computed. (Input)  
 MAXLAG must be greater than or equal to one and less than NOBS.

**IPRINT** — Printing option. (Input)

### IPRINT Action

- 0 No printing is performed.
- 1 Print the means and variances.
- 2 Print the means, variances, and cross-covariances.
- 3 Print the means, variances, cross-covariances, cross-correlations, and standard errors of cross-correlations.

**ISEOPT** — Option for computing standard errors of cross correlations. (Input)

**ISEOPT Action**

- 0 No standard errors of cross-correlations are computed.
- 1 Compute standard errors of cross-correlations using Bartlett's formula.
- 2 Compute standard errors of cross-correlations using Bartlett's formula with the assumption of no cross-correlation.

**IMEAN** — Option for computing the mean. (Input)

**IMEAN Action**

- 0 XMEAN and YMEAN are user specified.
- 1 XMEAN and YMEAN are set to the arithmetic means of X and Y.

**XMEAN** — Estimate of the mean of time series X. (Input, if IMEAN = 0; output, if IMEAN = 1)

**YMEAN** — Estimate of the mean of time series Y. (Input, if IMEAN = 0; output, if IMEAN = 1)

**XVAR** — Variance of the time series X. (Output)

**YVAR** — Variance of the time series Y. (Output)

**CCV** — Vector of length  $2 * \text{MAXLAG} + 1$  containing the cross-covariances between the time series X and Y. (Output)

The cross-covariance between X and Y at lag  $k$  corresponds to  $\text{CCV}(k)$  where  $k = -\text{MAXLAG}, \dots, -1, 0, 1, \dots, \text{MAXLAG}$ .

**CC** — Vector of length  $2 * \text{MAXLAG} + 1$  containing the cross-correlations between the time series X and Y. (Output)

The cross-correlation between X and Y at lag  $k$  corresponds to  $\text{CC}(k)$  where  $k = -\text{MAXLAG}, \dots, -1, 0, 1, \dots, \text{MAXLAG}$ .

**SECC** — Vector of length  $2 * \text{MAXLAG} + 1$  containing the standard errors of the crosscorrelations between the time series X and Y. (Output)

The standard error of  $\text{CC}(k)$  is  $\text{SECC}(k)$  where  $k = -\text{MAXLAG}, \dots, -1, 0, 1, \dots, \text{MAXLAG}$ .

**Comments**

- 1. Automatic workspace usage is
  - CCF 2 \* (MAXLAG + 1) units, or
  - DCCF 4 \* (MAXLAG + 1) units.

Workspace may be explicitly provided, if desired, by use of C2F/DC2F. The reference is

```
CALL C2F (NOBS, X, Y, MAXLAG, IPRINT, ISEOPT, IMEAN,
          XMEAN, YMEAN, XVAR, YVAR, CCV, CC, SECC,
          ACX, ACY)
```

The additional arguments are as follows:

**ACX** — Work vector of length equal to MAXLAG + 1.

**ACY** — Work vector of length equal to MAXLAG + 1.

2. If ISEOPT = 0, then no workspace is needed and SECC, ACX, and ACY can be dimensioned with length 1.
3. Autocovariances, autocorrelations, and standard errors of autocorrelations may be obtained by setting the first and second time series equal.

### Algorithm

Routine CCF estimates the cross-correlation function of two jointly stationary time series given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  and  $\{Y_t\}$  for  $t = 1, 2, \dots, n$ .

Let

$$\hat{\mu}_X = \text{XMEAN}$$

be the estimate of the mean  $\mu_X$  of the time series  $\{X_t\}$  where

$$\hat{\mu}_X = \begin{cases} \mu_X & \mu_X \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t & \mu_X \text{ unknown} \end{cases}$$

The autocovariance function of  $\{X_t\}$ ,  $\sigma_X(k)$ , is estimated by

$$\hat{\sigma}_X(k) = \frac{1}{n} \sum_{t=1}^{n-k} (X_t - \hat{\mu}_X)(X_{t+k} - \hat{\mu}_X), \quad k = 0, 1, \dots, K$$

where  $K = \text{MAXLAG}$ . Note that

$$\hat{\sigma}_X(0)$$

is equivalent to the sample variance  $\text{XVAR}$ . The autocorrelation function  $\rho_X(k)$  is estimated by

$$\hat{\rho}_X(k) = \frac{\hat{\sigma}_X(k)}{\hat{\sigma}_X(0)} \quad k = 0, 1, \dots, K$$

Note that

$$\hat{\rho}_X(0) \equiv 1$$

by definition. Let

$$\hat{\mu}_Y = \text{YMEAN}, \hat{\sigma}_Y(k), \text{ and } \hat{\rho}_Y(k)$$

be similarly defined.

The cross-covariance function  $\sigma_{XY}(k)$  is estimated by

$$\hat{\sigma}_{XY}(k) = \begin{cases} \frac{1}{n} \sum_{t=1}^{n-k} (X_t - \hat{\mu}_X)(Y_{t+k} - \hat{\mu}_Y) & k = 0, 1, \dots, K \\ \frac{1}{n} \sum_{t=1-k}^n (X_t - \hat{\mu}_X)(Y_{t+k} - \hat{\mu}_Y) & k = -1, -2, \dots, -K \end{cases}$$

The cross-correlation function  $\rho_{XY}(k)$  is estimated by

$$\hat{\rho}_{XY}(k) = \frac{\hat{\sigma}_{XY}(k)}{[\hat{\sigma}_X(0)\hat{\sigma}_Y(0)]^{1/2}} \quad k = 0, \pm 1, \dots, \pm K$$

The standard errors of the sample cross-correlations may be optionally computed according to argument `ISEOPT`. One method is based on a general asymptotic expression for the variance of the sample cross-correlation coefficient of two jointly stationary time series with independent, identically distributed normal errors given by Bartlett (1978, page 352). The theoretical formula is

$$\begin{aligned} \text{var}\{\hat{\rho}_{XY}(k)\} &= \frac{1}{n-k} \sum_{i=-\infty}^{\infty} [\rho_X(i)\rho_Y(i) + \rho_{XY}(i-k)\rho_{XY}(i+k) \\ &\quad - 2\rho_{XY}(k)\{\rho_X(i)\rho_{XY}(i+k) + \rho_{XY}(-i)\rho_Y(i+k)\} \\ &\quad + \rho_{XY}^2(k)\left\{\rho_X(i) + \frac{1}{2}\rho_X^2(i) + \frac{1}{2}\rho_Y^2(i)\right\}] \end{aligned}$$

For computational purposes, the autocorrelations  $\rho_X(k)$  and  $\rho_Y(k)$  and the cross-correlations  $\rho_{XY}(k)$  are replaced by their corresponding estimates for  $|k| \leq K$ , and the limits of summation are equal to zero for all  $k$  such that  $|k| > K$ .

A second method evaluates Bartlett's formula under the additional assumption that the two series have no cross-correlation. The theoretical formula is

$$\text{var}\{\hat{\rho}_{XY}(k)\} = \frac{1}{n-k} \sum_{i=-\infty}^{\infty} \rho_X(i)\rho_Y(i) \quad k \geq 0$$

For additional special cases of Bartlett's formula, see Box and Jenkins (1976, page 377).

An important property of the cross-covariance coefficient is  $\sigma_{XY}(k) = \sigma_{YX}(-k)$  for  $k \geq 0$ . This result is used in the computation of the standard error of the sample crosscorrelation for lag  $k < 0$ . In general, the cross-covariance function is not symmetric about zero so both positive and negative lags are of interest.



## Example

Consider the Gas Furnace Data (Box and Jenkins 1976, pages 532–533) where  $X$  is the input gas rate in cubic feet/minute and  $Y$  is the percent  $\text{CO}_2$  in the outlet gas. Routine CCF is used to compute the cross-covariances and cross-correlations between time series  $X$  and  $Y$  with lags from  $-\text{MAXLAG} = -10$  through lag  $\text{MAXLAG} = 10$ . In addition, the estimated standard errors of the estimated cross-correlations are computed. In the first invocation with  $\text{ISEOPT} = 1$ , the standard errors are based on the assumption that autocorrelations and cross-correlations for lags greater than  $\text{MAXLAG}$  or less than  $-\text{MAXLAG}$  are zero. In the second invocation with  $\text{ISEOPT} = 2$ , the standard errors are based on the additional assumption that all cross-correlations for  $X$  and  $Y$  are zero.

```
C
  INTEGER      IPRINT, MAXLAG, NOBS
  PARAMETER   (IPRINT=3, MAXLAG=10, NOBS=296)
C
  INTEGER      IMEAN, ISEOPT, NCOL, NROW
  REAL        CC(-MAXLAG:MAXLAG), CCV(-MAXLAG:MAXLAG),
&            RDATA(296,2), SECC(-MAXLAG:MAXLAG), X(NOBS), XMEAN,
&            XVAR, Y(NOBS), YMEAN, YVAR
  EXTERNAL    CCF, GDATA
C
  EQUIVALENCE (X(1), RDATA(1,1)), (Y(1), RDATA(1,2))
C
  CALL GDATA (7, 0, NROW, NCOL, RDATA, 296, 2)
C
  IMEAN = 1
C
  ISEOPT = 1
C
  CALL CCF (NOBS, X, Y, MAXLAG, IPRINT, ISEOPT, IMEAN, XMEAN,
&          YMEAN, XVAR, YVAR, CCV, CC, SECC)
C
  ISEOPT = 2
C
  CALL CCF (NOBS, X, Y, MAXLAG, IPRINT, ISEOPT, IMEAN, XMEAN,
&          YMEAN, XVAR, YVAR, CCV, CC, SECC)
C
  END
```

## Output

Output from CCF/C2F

```
Mean of series X      = -0.056834
Variance of series X =  1.1469

Mean of series Y      =  53.509
Variance of series Y =  10.219
```

Lag	CCV	CC	SECC
-10	-0.40450	-0.11815	0.158148
-9	-0.50849	-0.14853	0.155750
-8	-0.61437	-0.17946	0.152735

-7	-0.70548	-0.20607	0.149087
-6	-0.77617	-0.22672	0.145055
-5	-0.83147	-0.24287	0.141300
-4	-0.89132	-0.26035	0.138421
-3	-0.98060	-0.28643	0.136074
-2	-1.12477	-0.32854	0.132159
-1	-1.34704	-0.39347	0.123531
0	-1.65853	-0.48445	0.107879
1	-2.04865	-0.59841	0.087341
2	-2.48217	-0.72503	0.064141
3	-2.88541	-0.84282	0.046946
4	-3.16536	-0.92459	0.044097
5	-3.25344	-0.95032	0.048234
6	-3.13113	-0.91459	0.049155
7	-2.83919	-0.82932	0.047562
8	-2.45302	-0.71652	0.053478
9	-2.05269	-0.59958	0.071566
10	-1.69466	-0.49500	0.093933

Output from CCF/C2F

Mean of series X = -0.056834  
Variance of series X = 1.1469

Mean of series Y = 53.509  
Variance of series Y = 10.219

Lag	CCV	CC	SECC
-10	-0.40450	-0.11815	0.16275
-9	-0.50849	-0.14853	0.16247
-8	-0.61437	-0.17946	0.16219
-7	-0.70548	-0.20607	0.16191
-6	-0.77617	-0.22672	0.16163
-5	-0.83147	-0.24287	0.16135
-4	-0.89132	-0.26035	0.16107
-3	-0.98060	-0.28643	0.16080
-2	-1.12477	-0.32854	0.16052
-1	-1.34704	-0.39347	0.16025
0	-1.65853	-0.48445	0.15998
1	-2.04865	-0.59841	0.16025
2	-2.48217	-0.72503	0.16052
3	-2.88541	-0.84282	0.16080
4	-3.16536	-0.92459	0.16107
5	-3.25344	-0.95032	0.16135
6	-3.13113	-0.91459	0.16163
7	-2.83919	-0.82932	0.16191
8	-2.45302	-0.71652	0.16219
9	-2.05269	-0.59958	0.16247
10	-1.69466	-0.49500	0.16275

---

## MCCF/DMCCF (Single/Double precision)

Compute the multichannel cross-correlation function of two mutually stationary multichannel time series.

## Usage

```
CALL MCCF (NOBSX, NCHANX, X, LDX, NOBSY, NCHANY, Y, LDY,  
           MAXLAG, IPRINT, IMEAN, XMEAN, YMEAN, XVAR, YVAR,  
           CCV, LDCCV, MDCCV, CC, LDCC, MDCC)
```

## Arguments

**NOBSX** — Number of observations in each channel of the first time series  $X$ . (Input)

NOBSX must be greater than or equal to two.

**NCHANX** — Number of channels in the first time series  $X$ . (Input)

NCHANX must be greater than or equal to one.

**X** — NOBSX by NCHANX matrix containing the first time series. (Input)

Each row of  $X$  corresponds to an observation of a multivariate time series and each column of  $X$  corresponds to a univariate time series.

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement of the calling program. (Input)

LDX must be greater than or equal to NOBSX.

**NOBSY** — Number of observations in each channel of the second time series  $Y$ . (Input)

NOBSY must be greater than or equal to two.

**NCHANY** — Number of channels in the second time series  $Y$ . (Input)

NCHANY must be greater than or equal to one.

**Y** — NOBSY by NCHANY matrix containing the second time series. (Input)

Each row of  $Y$  corresponds to an observation of a multivariate time series and each column of  $Y$  corresponds to a univariate time series.

**LDY** — Leading dimension of  $Y$  exactly as specified in the dimension statement of the calling program. (Input)

LDY must be greater than or equal to NOBSY.

**MAXLAG** — Maximum lag of cross-covariances and cross-correlations to be computed. (Input)

MAXLAG must be greater than or equal to one and less than the minimum of NOBSX and NOBSY.

**IPRINT** — Printing option. (Input)

### IPRINT Action

- 0 No printing is performed.
- 1 Print the means and variances.
- 2 Print the means, variances, and cross-covariances.
- 3 Print the means, variances, cross-covariances, and cross-correlations.

**IMEAN** — Option for computing the means. (Input)

### IMEAN Action

- 0 XMEAN and YMEAN are user-specified.

1        *XMEAN* and *YMEAN* are set to the arithmetic means of their respective channels.

*XMEAN* — Vector of length *NCHANX* containing the means of the channels of *X*.  
(Input, if *IMEAN* = 0; output, if *IMEAN* = 1)

*YMEAN* — Vector of length *NCHANY* containing the means of the channels of *Y*.  
(Input, if *IMEAN* = 0; output, if *IMEAN* = 1)

*XVAR* — Vector of length *NCHANX* containing the variances of the channels of *X*.  
(Output)

*YVAR* — Vector of length *NCHANY* containing the variances of the channels of *Y*.  
(Output)

*CCV* — Array of size *NCHANX* by *NCHANY* by  $2 * \text{MAXLAG} + 1$  containing the cross-covariances between the channels of *X* and *Y*. (Output)

The cross-covariance between channel *i* of the *X* series and channel *j* of the *Y* series at lag *k* corresponds to  $CCV(i, j, k)$  where  $i = 1, \dots, NCHANX$ ,  $j = 1, \dots, NCHANY$ , and  $k = -\text{MAXLAG}, \dots, -1, 0, 1, \dots, \text{MAXLAG}$ .

*LDCCV* — Leading dimension of *CCV* exactly as specified in the dimension statement in the calling program. (Input)  
*LDCCV* must be greater than or equal to *NCHANX*.

*MDCCV* — Middle dimension of *CCV* exactly as specified in the dimension statement in the calling program. (Input)  
*MDCCV* must be greater than or equal to *NCHANY*.

*CC* — Array of size *NCHANX* by *NCHANY* by  $2 * \text{MAXLAG} + 1$  containing the cross-correlations between the channels of *X* and *Y*. (Output)

The cross-correlation between channel *i* of the *X* series and channel *j* of the *Y* series at lag *k* corresponds to  $CC(i, j, k)$  where  $i = 1, \dots, NCHANX$ ,  $j = 1, \dots, NCHANY$ , and  $k = -\text{MAXLAG}, \dots, -1, 0, 1, \dots, \text{MAXLAG}$ .

*LDCC* — Leading dimension of *CC* exactly as specified in the dimension statement in the calling program. (Input)  
*LDCC* must be greater than or equal to *NCHANX*.

*MDCC* — Middle dimension of *CC* exactly as specified in the dimension statement in the calling program. (Input)  
*MDCC* must be greater than or equal to *NCHANY*.

## Comments

1. For a given lag *k*, the multichannel cross-covariance coefficient is defined as the array of dimension *NCHANX* by *NCHANY* whose components are the single-channel cross-covariance coefficients  $CCV(i, j, k)$ . A similar definition holds for the multichannel cross-correlation coefficient.
2. Multichannel autocovariances and autocorrelations may be obtained by setting the first and second time series equal.

### Algorithm

Routine `MCCF` estimates the multichannel cross-correlation function of two mutually stationary multichannel time series. Define the multichannel time series  $X$  by

$$X = (X_1, X_2, \dots, X_p)$$

where

$$X_j = (X_{1j}, X_{2j}, \dots, X_{nj})^T, \quad j = 1, 2, \dots, p$$

with  $n = \text{NOBSX}$  and  $p = \text{NCHANX}$ . Similarly, define the multichannel time series  $Y$  by

$$Y = (Y_1, Y_2, \dots, Y_q)$$

where

$$Y_j = (Y_{1j}, Y_{2j}, \dots, Y_{mj})^T, \quad j = 1, 2, \dots, q$$

with  $m = \text{NOBSY}$  and  $q = \text{NCHANY}$ . The columns of  $X$  and  $Y$  correspond to individual channels of multichannel time series and may be examined from a univariate perspective. The rows of  $X$  and  $Y$  correspond to observations of  $p$ -variate and  $q$ -variate time series, respectively, and may be examined from a multivariate perspective. Note that an alternative characterization of a multivariate time series  $X$  considers the columns to be observations of the multivariate time series while the rows contain univariate time series. For example, see Priestley (1981, page 692) and Fuller (1976, page 14).

Let

$$\hat{\mu}_X = \text{XMEAN}$$

be the row vector containing the means of the channels of  $X$ . In particular,

$$\hat{\mu}_X = (\hat{\mu}_{X_1}, \hat{\mu}_{X_2}, \dots, \hat{\mu}_{X_p})$$

where for  $j = 1, 2, \dots, p$

$$\hat{\mu}_{X_j} = \begin{cases} \mu_{X_j} & \mu_{X_j} \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_{tj} & \mu_{X_j} \text{ unknown} \end{cases}$$

Let

$$\hat{\mu}_Y = \text{YMEAN}$$

be similarly defined. The cross-covariance of lag  $k$  between channel  $i$  of  $X$  and channel  $j$  of  $Y$  is estimated by

$$\hat{\sigma}_{X_i Y_j}(k) = \begin{cases} \frac{1}{N} \sum_t (X_{ti} - \hat{\mu}_{X_i})(Y_{t+k,j} - \hat{\mu}_{Y_j}) & k = 0, 1, \dots, K \\ \frac{1}{N} \sum_t (X_{ti} - \hat{\mu}_{X_i})(Y_{t+k,j} - \hat{\mu}_{Y_j}) & k = -1, -2, \dots, -K \end{cases}$$

where  $i = 1, \dots, p, j = 1, \dots, q$ , and  $K = \text{MAXLAG}$ . The summation on  $t$  extends over all possible cross-products with  $N$  equal to the number of cross-products in the sum.

Let

$$\hat{\sigma}_X(0) = \text{XVAR}$$

be the row vector consisting of the estimated variances of the channels of  $X$ . In particular,

$$\hat{\sigma}_X(0) = (\hat{\sigma}_{X_1}(0), \hat{\sigma}_{X_2}(0), \dots, \hat{\sigma}_{X_p}(0))$$

where

$$\hat{\sigma}_{X_j}(0) = \frac{1}{n} \sum_{t=1}^n (X_{tj} - \hat{\mu}_{X_j})^2 \quad j = 1, 2, \dots, p$$

Let

$$\hat{\sigma}_Y(0) = \text{YVAR}$$

be similarly defined. The cross-correlation of lag  $k$  between channel  $i$  of  $X$  and channel  $j$  of  $Y$  is estimated by

$$\hat{\rho}_{X_i Y_j}(k) = \frac{\hat{\sigma}_{X_i Y_j}(k)}{[\hat{\sigma}_{X_i}(0) \hat{\sigma}_{Y_j}(0)]^{1/2}} \quad k = 0, \pm 1, \dots, \pm K$$

### Example

Consider the Wolfer Sunspot Data ( $Y$ ) (Box and Jenkins 1976, page 530) along with data on northern light activity ( $X_1$ ) and earthquake activity ( $X_2$ ) (Robinson 1967, page 204) to be a three-channel time series. Routine `MCCF` is used to compute the cross-covariances and cross-correlations between  $X_1$  and  $Y$  and between  $X_2$  and  $Y$  with lags from  $-\text{MAXLAG} = -10$  through lag  $\text{MAXLAG} = 10$ :

```

C
  INTEGER      IPRINT, LDCC, LDCCV, LDX, LDY, MAXLAG, MDCC, MDCCV,
&             NCHANX, NCHANY, NOBSX, NOBSY
  PARAMETER   (IPRINT=3, MAXLAG=10, NCHANX=2, NCHANY=1, NOBSX=100,
&             NOBSY=100, LDCC=NCHANX, LDCCV=NCHANX, LDX=NOBSX,
&             LDY=NOBSY, MDCC=NCHANY, MDCCV=NCHANY)

  INTEGER      IMEAN, NCOL, NROW
  REAL         CC(LDCC, MDCC, -MAXLAG:MAXLAG), CCV(LDCCV, MDCCV, -
&             MAXLAG:MAXLAG), RDATA(100, 4), X(LDX, NCHANX),
&             XMEAN(NCHANX), XVAR(NCHANX), Y(LDY, NCHANY),
```

```

&          YMEAN(NCHANY), YVAR(NCHANY)
EXTERNAL  GDATA, MCCF
C
EQUIVALENCE (X(1,1), RDATA(1,3)), (X(1,2), RDATA(1,4))
EQUIVALENCE (Y(1,1), RDATA(1,2))
C
CALL GDATA (8, 0, NROW, NCOL, RDATA, 100, 4)
C          Option to estimate channel means
C          IMEAN = 1
C          Compute multichannel CCVF and CCF
CALL MCCF (NOBSX, NCHANX, X, LDX, NOBSY, NCHANY, Y, LDY, MAXLAG,
&          IPRINT, IMEAN, XMEAN, YMEAN, XVAR, YVAR, CCV, LDCCV,
&          MDCCV, CC, LDCC, MDCC)
C
END

```

### Output

Channel means of X from MCCF

	1	2
	63.43	97.97

Channel variances of X

	1	2
	2643.7	1978.4

Channel means of Y from MCCF

	46.94
--	-------

Channel variances of Y

	1383.8
--	--------

Multichannel cross-covariance between X and Y from MCCF

Lag K = -10

1	-20.51
2	70.71

Lag K = -9

1	65.02
2	38.14

Lag K = -8

1	216.6
2	135.6

Lag K = -7

1	246.8
2	100.4

Lag K = -6

1	142.1
2	45.0

Lag K = -5

1	50.70
2	-11.81

Lag K = -4

1	72.68
2	32.69
Lag K = -3	
1	217.9
2	-40.1
Lag K = -2	
1	355.8
2	-152.6
Lag K = -1	
1	579.7
2	-213.0
Lag K = 0	
1	821.6
2	-104.8
Lag K = 1	
1	810.1
2	55.2
Lag K = 2	
1	628.4
2	84.8
Lag K = 3	
1	438.3
2	76.0
Lag K = 4	
1	238.8
2	200.4
Lag K = 5	
1	143.6
2	283.0
Lag K = 6	
1	253.0
2	234.4
Lag K = 7	
1	479.5
2	223.0
Lag K = 8	
1	724.9
2	124.5
Lag K = 9	
1	925.0
2	-79.5
Lag K = 10	
1	922.8
2	-279.3



Multichannel cross-correlation between X and Y from MCCF

Lag K = -10  
1 -0.01072  
2 0.04274

Lag K = -9  
1 0.03400  
2 0.02305

Lag K = -8  
1 0.1133  
2 0.0819

Lag K = -7  
1 0.1290  
2 0.0607

Lag K = -6  
1 0.07431  
2 0.02718

Lag K = -5  
1 0.02651  
2 -0.00714

Lag K = -4  
1 0.03800  
2 0.01976

Lag K = -3  
1 0.1139  
2 -0.0242

Lag K = -2  
1 0.1860  
2 -0.0923

Lag K = -1  
1 0.3031  
2 -0.1287

Lag K = 0  
1 0.4296  
2 -0.0633

Lag K = 1  
1 0.4236  
2 0.0333

Lag K = 2  
1 0.3285  
2 0.0512

Lag K = 3  
1 0.2291  
2 0.0459

Lag K = 4

```

1  0.1248
2  0.1211

Lag K =      5
1  0.0751
2  0.1710

Lag K =      6
1  0.1323
2  0.1417

Lag K =      7
1  0.2507
2  0.1348

Lag K =      8
1  0.3790
2  0.0752

Lag K =      9
1  0.4836
2 -0.0481

Lag K =     10
1  0.4825
2 -0.1688

```

---

## ARMME/DARMME (Single/Double precision)

Compute method of moments estimates of the autoregressive parameters of an ARMA model.

### Usage

CALL ARMME (MAXLAG, ACV, IPRINT, NPMA, NPAR, PAR)

### Arguments

**MAXLAG** — Maximum lag of the sample autocovariances of the time series  $W$ . (Input)

MAXLAG must be greater than or equal to NPAR + NPMA.

**ACV** — Vector of length MAXLAG + 1 containing the sample autocovariances of  $W$ . (Input)

The  $k$ -th sample autocovariance of  $W$  is denoted by  $ACV(k)$ ,  $k = 0, 1, \dots, MAXLAG$ .

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.

1 Print the estimates of the autoregressive parameters.

**NPMA** — Number of moving average parameters. (Input)

NPMA must be greater than or equal to zero.

**NPAR** — Number of autoregressive parameters. (Input)  
 NPAR must be greater than or equal to one.

**PAR** — Vector of length NPAR containing the estimates of the autoregressive parameters. (Output)

### Comments

- Automatic workspace usage is

ARMME  $2 * \text{NPAR} * (\text{NPAR} + 1)$  units, or  
 DARMME  $\text{NPAR} * (4 * \text{NPAR} + 3)$  units.

Workspace may be explicitly provided, if desired, by use of  
 A2MME/DA2MME. The reference is

```
CALL A2MME (MAXLAG, ACV, IPRINT, NPMA, NPAR, PAR, A,
            FAC, IPVT, WK)
```

The additional arguments are as follows:

**A** — Work vector of length equal to  $\text{NPAR}^2$ .

**FAC** — Work vector of length equal to  $\text{NPAR}^2$ .

**IPVT** — Work vector of length equal to NPAR.

**WK** — Work vector of length equal to NPAR.

- Informational error

Type	Code	
4	1	The problem is ill-conditioned. Transformation of the data or increased precision in the calculations may be appropriate.

- The sample autocovariance function may be obtained using the routine ACF (page 637).
- The first element of ACV must be the sample variance of the time series.

### Algorithm

Routine ARMME determines the autoregressive parameters of an ARMA process using the extended Yule-Walker equations given the  $K = \text{MAXLAG}$  autocovariances  $\sigma(k)$  for  $k = 1, \dots, K$ .

Suppose the time series  $\{W_t\}$  is generated by an ARMA( $p, q$ ) model

$$W_t = \theta_0 + \phi_1 W_{t-1} + \dots + \phi_p W_{t-p} + A_t - \theta_1 A_{t-1} - \dots - \theta_q A_{t-q}, \quad t \in \{0, \pm 1, \pm 2, \dots\}$$

where  $p = \text{NPAR}$  and  $q = \text{NPMA}$ . Since  $W_t$  depends only on the innovations  $A_t$  that have occurred up through time  $t$ , the  $p$  autoregressive parameters are related to the autocovariances of lags  $k = q + 1, \dots, q + p$  by the set of equations

$$\sigma(q + 1) = \phi_1 \sigma(q) + \phi_2 \sigma(q - 1) + \dots + \phi_p \sigma(q - p + 1)$$

$$\sigma(q+2) = \phi_1 \sigma(q+1) + \phi_2 \sigma(q) + \dots + \phi_p \sigma(q-p+2)$$

⋮

$$\sigma(q+p) = \phi_1 \sigma(q+p-1) + \phi_2 \sigma(q+p-2) + \dots + \phi_p \sigma(q)$$

This general system of linear equations is called the extended Yule-Walker equations. For  $q = 0$ , the system is referred to as the Yule-Walker equations. The equivalent matrix version is given by

$$\Sigma \phi = \sigma$$

where

$$\begin{aligned} \phi &= (\phi_1, \dots, \phi_p)^T \\ \Sigma_{ij} &= \sigma(|q+i-j|) \quad i, j = 1, \dots, p \\ \sigma_i &= \sigma(q+i) \quad i = 1, \dots, p \end{aligned}$$

The overall constant  $\theta_0$  is defined by

$$\theta_0 = \begin{cases} \mu & p = 0 \\ \mu(1 - \sum_{i=1}^p \phi_i) & p > 0 \end{cases}$$

where  $\mu$  is the mean of  $W_t$ .

In practice, the autocovariance function is estimated by the sample autocovariances

$$\hat{\sigma}(k)$$

for  $k = 1, \dots, K$ . The solution of the extended Yule-Walker equations using these sample moments yields the *method of moments* estimates of the autoregressive parameters. The overall constant may then be estimated given an estimate of  $\mu$ . Note that the extended Yule-Walker equations may be analogously defined in terms of autocorrelations instead of autocovariances. See Box and Jenkins (1976, pages 189–191) for some comments concerning the initial estimation of autoregressive parameters using the Yule-Walker equations.

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Routine `ARMME` is invoked first to compute the method of moments estimates for the autoregressive parameters of an ARMA(2, 0) model given the sample autocovariances computed from routine `ACF` (page 637). Then, `ARMME` is invoked a second time to compute estimated autoregressive parameters for an ARMA(2, 1) model.

INTEGER    IMEAN, IPRINT, ISEOPT, MAXLAG, NOBS

```

PARAMETER (IMEAN=1, IPRINT=1, ISEOPT=0, MAXLAG=4, NOBS=100)
C
INTEGER    NCOL, NOUT, NPAR, NPMA, NROW
REAL      AC(0:MAXLAG), ACV(0:MAXLAG), PAR(2), RDATA(176,2),
&         SEAC(1), W(100), WMEAN
EXTERNAL  ACF, ARMME, GDATA, UMACH
C
EQUIVALENCE (W(1), RDATA(22,2))
C
CALL UMACH (2, NOUT)
C
C           Wolfer Sunspot Data for
C           years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C
C           Compute sample ACV
CALL ACF (NOBS, W, 0, ISEOPT, IMEAN, WMEAN, MAXLAG, ACV, AC,
&        SEAC)
C
C           Compute estimates of autoregressive
C           parameters for ARMA(2,0) model
C           (Box and Jenkins, page 83)
WRITE (NOUT,*) 'ARMA(2,0) Model'
NPMA = 2
NPMA = 0
CALL ARMME (MAXLAG, ACV, IPRINT, NPMA, NPAR, PAR)
C
C           Compute estimates of autoregressive
C           parameters for ARMA(2,1) model
WRITE (NOUT,*) ' '
WRITE (NOUT,*) 'ARMA(2,1) Model'
NPMA = 1
CALL ARMME (MAXLAG, ACV, IPRINT, NPMA, NPAR, PAR)
C
END

```

### Output

ARMA(2,0) Model

```

Output PAR
  1      2
1.318  -0.635

```

ARMA(2,1) Model

```

Output PAR
  1      2
1.244  -0.575

```

---

## MAMME/DMAMME (Single/Double precision)

Compute method of moments estimates of the moving average parameters of an ARMA model.

### Usage

```

CALL MAMME (MAXLAG, ACV, IPRINT, NPAR, PAR, RELERR, MAXIT,
           NPMA, PMA)

```

## Arguments

**MAXLAG** — Maximum lag of the sample autocovariances of the time series  $W$ . (Input)

MAXLAG must be greater than or equal to  $\text{NPAR} + \text{NPMA}$ .

**ACV** — Vector of length  $\text{MAXLAG} + 1$  containing the sample autocovariances of  $W$ . (Input)

The  $k$ -th sample autocovariance of  $W$  is denoted by  $\text{ACV}(k)$ ,  $k = 0, 1, \dots, \text{MAXLAG}$ .

**IPRINT** — Printing option. (Input)

### IPRINT Action

0 No printing is performed.

1 Print the estimates of the moving average parameters.

**NPAR** — Number of autoregressive parameters. (Input)

NPAR must be greater than or equal to zero.

**PAR** — Vector of length NPAR containing the estimates of the autoregressive parameters. (Input)

**RELERR** — Stopping criterion for use in the nonlinear equation solver. (Input)

If  $\text{RELERR} = 0.0$ , then the default value  $\text{RELERR} = 100.0 * \text{AMACH}(4)$  is used. See the documentation for routine *AMACH* (page 1334).

**MAXIT** — The maximum number of iterations allowed in the nonlinear equation solver. (Input)

If  $\text{MAXIT} = 0$ , then the default value  $\text{MAXIT} = 200$  is used.

**NPMA** — Number of moving average parameters. (Input)

NPMA must be greater than or equal to one.

**PMA** — Vector of length NPMA containing the estimates of the moving average parameters. (Output)

## Comments

1. Automatic workspace usage is

MAMME  $\text{NPAR} + 1 + (\text{NPMA} + 1) * (12 + 1.5 * \text{NPMA})$  units, or

DMAMME  $2 * (\text{NPAR} + 1 + (\text{NPMA} + 1) * (12 + 1.5 * \text{NPMA}))$  units.

Workspace may be explicitly provided, if desired, by use of

M2MME/DM2MME. The reference is

```
CALL M2MME (MAXLAG, ACV, IPRINT, NPAR, PAR, RELERR,
            MAXIT, NPMA, PMA, PARWK, ACVMOD, TAUINI,
            TAU, FVEC, FJAC, R, QTF, WKNLN)
```

The additional arguments are as follows:

**PARWK** — Work vector of length equal to  $\text{NPAR} + 1$ .

**ACVMOD** — Work vector of length equal to NPMA + 1.

**TAUINI** — Work vector of length equal to NPMA + 1.

**TAU** — Work vector of length equal to NPMA + 1.

**FVEC** — Work vector of length equal to NPMA + 1.

**FJAC** — Work vector of length equal to (NPMA + 1)<sup>2</sup>.

**R** — Work vector of length equal to (NPMA + 1) \* (NPMA + 2)/2.

**QTF** — Work vector of length equal to NPMA + 1.

**WKNLN** — Work vector of length equal to 5 \* (NPMA + 1).

2. Informational error

Type Code

4 1 The nonlinear equation solver did not converge to RELERR within MAXIT iterations.

3. The sample autocovariance function may be computed using the routine ACF (page 637).

4. The autoregressive parameter estimates may be computed using the routine ARMME (page 657).

**Algorithm**

Routine MAMME estimates the moving average parameters of an ARMA process based on a system of nonlinear equations given  $K = \text{MAXLAG}$  autocovariances  $\sigma(k)$  for  $k = 1, \dots, K$  and  $p = \text{NPAR}$  autoregressive parameters  $\phi_i$  for  $i = 1, \dots, p$ .

Suppose the time series  $\{W_t\}$  is generated by an ARMA( $p, q$ ) model

$$\phi(B)W_t = \theta_0 + \theta(B)A_t, \quad t \in 2 \{0, \pm 1, \pm 2, \dots\}$$

where  $p = \text{NPAR}$  and  $q = \text{NPMA}$  Let

$$W_t' = \phi(B)W_t$$

then the autocovariances of the *derived* moving average process  $W_t = \theta(B)A_t$  are given by

$$\sigma'(k) = \begin{cases} \sigma(k) & p = 0 \\ \sum_{i=0}^p \sum_{j=0}^p \phi_i \phi_j \sigma(|k+i-j|) & p \geq 1, \phi_0 \equiv -1 \end{cases}$$

where  $\sigma(k)$  denotes the autocovariance function of the original  $W_t$  process. The iterative procedure for determining the moving average parameters is based on the relation

$$\sigma'(k) = \begin{cases} (1 + \theta_1^2 + \dots + \theta_q^2) \sigma_A^2 & k = 0 \\ (-\theta_k + \theta_1 \theta_{k+1} + \dots + \theta_{q-k} \theta_q) \sigma_A^2 & k \geq 1 \end{cases}$$

Let  $\tau = (\tau_0, \tau_1, \dots, \tau_q)^T$  and  $f = (f_0, f_1, \dots, f_q)^T$  where

$$\tau_j = \begin{cases} \sigma_A & j = 0 \\ -\theta_j / \tau_0 & j = 1, \dots, q \end{cases}$$

and

$$f_j = \sum_{i=0}^{q-j} \tau_i \tau_{i+j} - \sigma'(j) \quad j = 0, 1, \dots, q$$

Then, the value of  $\tau$  at the  $(i + 1)$ -th iteration is determined by

$$\tau^{i+1} = \tau^i - (T^i)^{-1} f^i$$

The estimation procedure begins with the initial value

$$\tau^0 = (\sqrt{\sigma'(0)}, 0, \dots, 0)^T$$

and terminates at iteration  $i$  when either  $\|f^i\|$  is less than RELERR or  $i$  equals MAXIT. The moving average parameters are determined from the final estimate of  $\tau$  by setting  $\theta_j = -\tau_j / \tau_0$  for  $j = 1, \dots, q$ . The random shock variance is determined according to

$$\sigma_A^2 = \begin{cases} \sigma(0) - \sum_{i=1}^p \phi_i \sigma(i) & q = 0 \\ \tau_0^2 & q \geq 1 \end{cases}$$

In practice, both the autocovariances and the autoregressive parameters are estimated. The solution of the system of nonlinear equations using these sample moments yields the method of moments estimates of the moving average parameters and the random shock variance. Note that autocorrelations  $\rho(k)$  may be used instead of autocovariances  $\sigma(k)$  to compute  $\sigma'(k)$  for  $k = 1, \dots, K$ . See Box and Jenkins (1976, pages 203–204) for additional motivation concerning the initial estimation of moving average parameters using a Newton-Raphson algorithm.

### Example

Consider the Wölfer Sunspot Data (Box and Jenkins 1976, page 530) consisting of the number of sunspots observed each year from 1770 through 1869. Routine MAMME is invoked to compute the method of moments estimates for the moving average parameter of an ARMA(2,1) model given the sample autocovariances computed from routine ACF (page 637) and given the estimated autoregressive parameters computed from routine ARMME (page 657).

```

C
  INTEGER      IMEAN, IPRINT, ISEOPT, LDX, MAXLAG, NDX, NOBS,
&             NOPRIN, NPAR, NPMA
  PARAMETER   (IMEAN=1, IPRINT=1, ISEOPT=0, LDX=176, MAXLAG=4,
&             NDX=2, NOBS=100, NOPRIN=0, NPAR=2, NPMA=1)
  INTEGER      MAXIT, NCOL, NROW

```



```

REAL      AC(0:MAXLAG), ACV(0:MAXLAG), PAR(2), PMA(1),
&         RDATA(LDX,NDX), RELERR, SEAC(1), W(100), WMEAN
EXTERNAL  ACF, ARMME, GDATA, MAMME
C
EQUIVALENCE (W(1), RDATA(22,2))
C
C           Wolfer Sunspot Data for
C           years 1770 through 1869
CALL GDATA (2, NOPRIN, NROW, NCOL, RDATA, LDX, NDX)
C
C           Compute sample ACV
CALL ACF (NOBS, W, NOPRIN, ISEOPT, IMEAN, WMEAN, MAXLAG, ACV,
&         AC, SEAC)
C
C           Compute estimates of autoregressive
C           parameters for ARMA(2,1) model
CALL ARMME (MAXLAG, ACV, NOPRIN, NPMA, NPAR, PAR)
C
C           Convergence parameters
MAXIT = 0
RELERR = 0.0
C
C           Compute estimate of moving average
C           parameter for ARMA(2,1) model
CALL MAMME (MAXLAG, ACV, IPRINT, NPAR, PAR, RELERR, MAXIT, NPMA,
&         PMA)
C
END

```

### Output

Output PMA from MAMME/M2MME  
-0.1241

---

## NSPE/DNSPE (Single/Double precision)

Compute preliminary estimates of the autoregressive and moving average parameters of an ARMA model.

### Usage

```
CALL NSPE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, NPMA,
          RELERR, MAXIT, CONST, PAR, PMA, AVAR)
```

### Arguments

**NOBS** — Number of observations in the stationary time series *W*. (Input)  
NOBS must be greater than NPAR + NPMA + 1.

**W** — Vector of length NOBS containing the stationary time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.

1 Print the mean of the time series, the estimate of the overall constant, the estimates of the autoregressive parameters, the estimates of the moving average parameters, and the estimate of the random shock variance.

**IMEAN** — Option for centering the time series *x*. (Input)

**IMEAN Action**

- 0 WMEAN is user specified.  
 1 WMEAN is set to the arithmetic mean of x.

**WMEAN** — Constant used to center the time series x. (Input, if *IMEAN* = 0; output, if *IMEAN* = 1)

**NPAR** — Number of autoregressive parameters. (Input)  
 NPAR must be greater than or equal to zero.

**NPMA** — Number of moving average parameters. (Input)  
 NPMA must be greater than or equal to zero.

**RELERR** — Stopping criterion for use in the nonlinear equation solver. (Input)  
 If *RELERR* = 0.0, then the default value *RELERR* = 100.0 \* *AMACH*(4) is used. See the documentation for routine *AMACH* (page 1334).

**MAXIT** — The maximum number of iterations allowed in the nonlinear equation solver. (Input)  
 If *MAXIT* = 0, then the default value *MAXIT* = 200 is used.

**CONST** — Estimate of the overall constant. (Output)

**PAR** — Vector of length *NPAR* containing the autoregressive parameter estimates. (Output)

**PMA** — Vector of length *NPMA* containing the moving average parameter estimates. (Output)

**AVAR** — Estimate of the random shock variance. (Output)

**Comments**

- Automatic workspace usage is

NSPE 14 + NPMA \* (14.5 + 1.5 \* NPMA) + 2 \* NPAR \* (NPAR + 2)  
 units, or

DNspe 28 + NPMA \* (14 + 3 \* NPMA) + NPAR \* (4 \* NPAR + 7) units.

Workspace may be explicitly provided, if desired, by use of *N2PE/DN2PE*. The reference is

```
CALL N2PE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR,
           NPMA, RELERR, MAXIT, CONST, PAR, PMA,
           AVAR, ACV, PARWK, ACVMOD, TAUINI, TAU,
           FVEC, FJAC, R, QTF, WKNLN, A, FAC, IPVT,
           WKARMM)
```

The additional arguments are as follows:

**ACV** — Work vector of length equal to *NPAR* + *NPMA* + 1.

**PARWK** — Work vector of length equal to *NPAR* + 1.

**ACVMOD** — Work vector of length equal to *NPMA* + 1.

**TAUINI** — Work vector of length equal to NPMA + 1.  
**TAU** — Work vector of length equal to NPMA + 1.  
**FVEC** — Work vector of length equal to NPMA + 1.  
**FJAC** — Work vector of length equal to (NPMA + 1)<sup>2</sup>.  
**R** — Work vector of length equal to (NPMA + 1) \* (NPMA + 2)/2.  
**QTF** — Work vector of length equal to NPMA + 1.  
**WKNLN** — Work vector of length equal to 5 \* (NPMA + 1).  
**A** — Work vector of length equal to NPAR<sup>2</sup>.  
**FAC** — Work vector of length equal to NPAR<sup>2</sup>.  
**IPVT** — Work vector of length equal to NPAR.  
**WKARMM** — Work vector of length equal to NPAR.

2. Informational error
 

Type	Code	
4	1	The nonlinear equation solver did not converge to RELEERR within MAXIT iterations.
3. The value of WMEAN is used in the computation of the sample autocovariances of  $w$  in the process of obtaining the preliminary autoregressive parameter estimates. Also, WMEAN is used to obtain the value of CONST.

### Algorithm

Routine NSPE computes preliminary estimates of the parameters of an ARMA process given a sample of  $n = \text{NOBS}$  observations  $\{W_t\}$  for  $t = 1, 2, \dots, n$ .

Suppose the time series  $\{W_t\}$  is generated by an ARMA( $p, q$ ) model of the form

$$\phi(B)W_t = \theta_0 + \theta(B)A_t \quad t \in \{0, \pm 1, \pm 2, \dots\}$$

where  $B$  is the backward shift operator,

$$\phi(B) = 1 - \phi_1(B) - \phi_2(B)^2 - \dots - \phi_p(B)^p$$

$$\theta(B) = 1 - \theta_1(B) - \theta_2(B)^2 - \dots - \theta_q(B)^q$$

$p = \text{NPAR}$  and  $q = \text{NPMA}$ . Let

$$\hat{\mu} = \text{WMEAN}$$

be the estimate of the mean of the time series  $\{W_t\}$  where

$$\hat{\mu} = \begin{cases} \mu & \mu \text{ known} \\ \frac{1}{n} \sum_{t=1}^n W_t & \mu \text{ unknown} \end{cases}$$

The autocovariance function  $\sigma(k)$  is estimated by

$$\hat{\sigma}(k) = \frac{1}{n} \sum_{t=1}^{n-k} (W_t - \hat{\mu})(W_{t+k} - \hat{\mu}) \quad k = 0, 1, \dots, K$$

where  $K = p + q$ . Note that

$$\hat{\sigma}(0)$$

is an estimate of the sample variance.

Given the sample autocovariances, the routine `ARMME` (page 657) is used to compute the method of moments estimates of the autoregressive parameters using the extended Yule-Walker equations

$$\hat{\Sigma} \hat{\phi} = \hat{\sigma}$$

where

$$\begin{aligned} \hat{\phi} &= (\hat{\phi}_1, \dots, \hat{\phi}_p)^T \\ \hat{\Sigma}_{ij} &= \hat{\sigma}(|q + i - j|) \quad i, j = 1, \dots, p \\ \hat{\sigma}_i &= \hat{\sigma}(q + i) \quad i = 1, \dots, p \end{aligned}$$

The overall constant  $\theta_0$  is estimated by

$$\hat{\theta}_0 = \begin{cases} \hat{\mu} & p = 0 \\ \hat{\mu}(1 - \sum_{i=1}^p \hat{\phi}_i) & p > 0 \end{cases}$$

The moving average parameters are estimated using the routine `MAMME` (page 660). Let

$$W'_t = \phi(B)W_t$$

then the autocovariances of the *derived* moving average process

$$W'_t = \theta(B)A_t$$

are estimated by

$$\hat{\sigma}'(k) = \begin{cases} \hat{\sigma}(k) & p = 0 \\ \sum_{i=0}^p \sum_{j=0}^p \hat{\phi}_i \hat{\phi}_j \hat{\sigma}(|k + i - j|) & p \geq 1, \hat{\phi}_0 \equiv -1 \end{cases}$$

The iterative procedure for determining the moving average parameters is based on the relation

$$\sigma'(k) = \begin{cases} (1 + \theta_1^2 + \dots + \theta_q^2)\sigma_A^2 & k = 0 \\ (-\theta_k + \theta_1\theta_{k+1} + \dots + \theta_{q-k}\theta_q)\sigma_A^2 & k \geq 1 \end{cases}$$

where  $\sigma(k)$  denotes the autocovariance function of the original  $W_t$  process.

Let  $\tau = (\tau_0, \tau_1, \dots, \tau_q)^T$  and  $f = (f_0, f_1, \dots, f_q)^T$  where

$$\tau_j = \begin{cases} \sigma_A & j = 0 \\ -\theta_j / \tau_0 & j = 1, \dots, q \end{cases}$$

and

$$f_j = \sum_{i=0}^{q-j} \tau_i \tau_{i+j} - \hat{\sigma}'(j) \quad j = 0, 1, \dots, q$$

Then, the value of  $\tau$  at the  $(i + 1)$ -th iteration is determined by

$$\tau^{i+1} = \tau^i - (T^i)^{-1} f^i$$

The estimation procedure begins with the initial value

$$\tau^0 = (\sqrt{\hat{\sigma}'(0)}, 0, \dots, 0)^T$$

and terminates at iteration  $i$  when either  $\|f^i\|$  is less than `RELEERR` or  $i$  equals `MAXIT`. The moving average parameter estimates are obtained from the final estimate of  $\tau$  by setting

$$\hat{\theta}_j = -\tau_j / \tau_0 \quad \text{for } j = 1, \dots, q$$

The random shock variance is estimated by

$$\hat{\sigma}_A^2 = \begin{cases} \hat{\sigma}(0) - \sum_{i=1}^p \hat{\phi}_i \hat{\sigma}(i) & q = 0 \\ \tau_0^2 & q \geq 0 \end{cases}$$

See Box and Jenkins (1976, pages 498–500) for a description of a similar routine.

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Routine `NSPE` (page 664) is used to compute preliminary estimates

$\hat{\theta}_0$  (output in `CONST`)

$\hat{\phi}_1, \phi_2$ , (output in `PAR`)

$\hat{\theta}_1$  (output in `PMA`)

$\hat{\sigma}_A^2$  (output in AVAR)

for the following ARMA (2, 1) model

$$w_t = \theta_0 + \phi_1 w_{t-1} + \phi_2 w_{t-2} - \theta_1 A_{t-1} + A_t$$

where the errors  $A_t$  are independently distributed each normal with mean zero and variance

$\sigma_A^2$

```
INTEGER      IPRINT, LDX, NDX, NOBS, NOPRIN, NPAR, NPMA
PARAMETER    (IPRINT=1, LDX=176, NDX=2, NOBS=100, NOPRIN=0, NPAR=2,
&            NPMA=1)
C
INTEGER      IMEAN, MAXIT, NCOL, NROW
REAL         AVAR, CONST, PAR(NPAR), PMA(NPMA), RDATA(LDX,NDX),
&            RELERR, W(NOBS), WMEAN
EXTERNAL     GDATA, NSPE
C
EQUIVALENCE (W(1), RDATA(22,2))
C                                     Wolfer Sunspot Data for
C                                     years 1770 through 1869
CALL GDATA (2, NOPRIN, NROW, NCOL, RDATA, LDX, NDX)
C                                     Convergence parameters
MAXIT = 0
RELERR = 0.0
C                                     Compute preliminary parameter
C                                     estimates for ARMA(2,1) model
IMEAN = 1
CALL NSPE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, NPMA, RELERR,
&         MAXIT, CONST, PAR, PMA, AVAR)
C
END
```

### Output

Results from NSPE/N2PE

```
WMEAN =      46.9760
CONST =      15.5440
AVAR =       287.242
```

```
PAR
  1      2
1.244  -0.575
```

```
PMA
-0.1241
```

---

## NSLSE/DNSLSE (Single/Double precision)

Compute least-squares estimates of parameters for a nonseasonal ARMA model.

## Usage

```
CALL NSLSE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, PAR,  
           LAGAR, NPMA, PMA, LAGMA, MAXBC, TOLBC, TOLSS,  
           CONST, COV, LDCOV, NA, A, AVAR)
```

## Arguments

**NOBS** — Number of observations in the stationary time series *w*. (Input)  
NOBS must be greater than IARDEG + IMADEG where IARDEG = max(LAGAR(*i*))  
and IMADEG = max(LAGMA(*j*)).

**W** — Vector of length NOBS containing the stationary time series. (Input)

**IPRINT** — Printing option. (Input)

### IPRINT Action

- 0 No printing is performed.
- 1 Print the least-squares estimates of the parameters, their associated standard errors, and the residual sum of squares at the final iteration.
- 2 Print the least-squares estimates of the parameters and the residual sum of squares at each iteration and at the final iteration. Print the standard errors of the parameters at the final iteration.

**IMEAN** — Option for centering the time series *w*. (Input)

### IMEAN Action

- 0 *w* is not centered.
- 1 *w* is centered about WMEAN. Centering the time series *w* about WMEAN is equivalent to inclusion of the overall constant in the model.

**WMEAN** — Estimate of the mean of the time series *w*. (Input/Output, if IMEAN = 1; not used if IMEAN = 0)

For IMEAN = 1, on input, WMEAN contains the preliminary estimate, on output, WMEAN contains the final estimate.

**NPAR** — Number of autoregressive parameters. (Input)

NPAR must be greater than or equal to zero.

**PAR** — Vector of length NPAR containing the autoregressive parameters.(Input/Output)

On input, PAR contains the preliminary estimate. On output, PAR contains the final estimate.

**LAGAR** — Vector of length NPAR containing the order of the autoregressive parameters. (Input)

The elements of LAGAR must be greater than or equal to one.

**NPMA** — Number of moving average parameters. (Input)

NPMA must be greater than or equal to zero.

**PMA** — Vector of length NPMA containing the moving average parameters.(Input/Output)

On input, PMA contains the preliminary estimate. On output, PMA contains the final estimate.

**LAGMA** — Vector of length NPMA containing the order of the moving average parameters. (Input)

The elements of LAGMA must be greater than or equal to one.

**MAXBC** — Maximum length of backcasting. (Input)

MAXBC must be greater than or equal to zero.

**TOLBC** — Tolerance level used to determine convergence of the backcast algorithm. (Input)

Backcasting terminates when the absolute value of a backcast is less than TOLBC. Typically, TOLBC is set to a fraction of WSTDEV where WSTDEV is an estimate of the standard deviation of the time series. If TOLBC = 0.0, then TOLBC = 0.01 \* WSTDEV is used.

**TOLSS** — Tolerance level used to determine convergence of the nonlinear least-squares algorithm. (Input)

TOLSS represents the minimum relative decrease in sum of squares between two iterations required to determine convergence. Hence, TOLSS must be greater than or equal to zero and less than one where TOLSS = 0.0 specifies the default value is to be used. The default value is

$\max\{10^{-10}, \text{EPS}^{2/3}\}$  for single precision and

$\max\{10^{-20}, \text{EPS}^{2/3}\}$  for double precision

where EPS = AMACH(4) for single precision and EPS = DMACH(4) for double precision. See the documentation for routine AMACH (page 1334).

**CONST** — Estimate of the overall constant. (Output)

For IMEAN = 0, CONST is set to zero. For IMEAN = 1, CONST = WMEAN \* (1 - PAR(1) - PAR(2) - ... - PAR(NPAR)).

**COV** — NP by NP variance-covariance matrix of the estimates of the parameters where NP = IMEAN + NPAR + NPMA. (Output)

The ordering of variables in COV is WMEAN (if defined), PAR, and PMA. NP must 1 or more.

**LDCOV** — Leading dimension of COV exactly as specified in the dimension statement in the calling program. (Input)

**NA** — Number of residuals computed (including backcasts). (Output)

If NB values of the time series are backcast, then NA = NOBS - IARDEG + NB.

**A** — Vector of length NOBS - IARDEG + MAXBC containing the residuals (including backcasts) at the final parameter estimate point in the first NA locations. (Output)

**AVAR** — Estimate of the random shock variance. (Output)

$\text{AVAR} = (\text{A}(1)^2 + \dots + \text{A}(\text{NA})^2) / (\text{NOBS} - \text{IMEAN} - \text{NPAR} - \text{NPMA})$ .



## Comments

1. Automatic workspace usage is

NSLSE  $(14 + M) * NP + 4M + NOBS + 2 * MAXBC + IMADEG - 1$  units, or

DNLSLSE  $2 * ((13+M) * NP + 4M + NOBS + 2 * MAXBC + IMADEG - 1) + NP$   
units where  $NP = NPAR + NPMA + IMEAN$  and  $M = NOBS -$   
 $IARDEG + MAXBC$ .

Workspace may be explicitly provided, if desired, by use of  
N2LSE/DN2LSE. The reference is

```
CALL N2LSE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR,  
           PAR, LAGAR, NPMA, PMA, LAGMA, MAXBC,  
           TOLBC, TOLSS, CONST, COV, LDCOV, NA, A,  
           AVAR, XGUESS, XSCALE, FSCALE, X, FVEC,  
           FJAC, LDFJAC, RWKUNL, IWKUNL, WKNSRE,  
           AI, FCST)
```

The additional arguments are as follows:

**XGUESS** — Work vector of length NP.

**XSCALE** — Work vector of length NP.

**FSCALE** — Work vector of length M.

**X** — Work vector of length NP.

**FVEC** — Work vector of length M.

**FJAC** — Work vector of length  $M * NP$ .

**LDFJAC** — Integer scalar equal to M.

**RWKUNL** — Work vector of length  $10 * NP + 2 * M - 1$ .

**IWKUNL** — Work vector of length NP.

**WKNSRE** — Work vector of length  $NOBS + MAXBC$ .

**AI** — Work vector of length IMADEG.

**FCST** — Work vector of length MAXBC.

2. Informational error

Type	Code	
3	1	Least-squares estimation of the parameters has failed to converge. Increase MAXBC and/or TOLBC and/or TOLSS. The estimates of the parameters at the last iteration may be used as new starting values.

### Algorithm

Routine `NLSSE` computes least-squares estimates of parameters for a nonseasonal ARMA model given a sample of  $n = \text{NOBS}$  observations  $\{W_t\}$  for  $t = 1, 2, \dots, n$ .

Suppose the time series  $\{W_t\}$  is generated by a nonseasonal ARMA model of the form

$$\phi(B)(W_t - \mu) = \theta(B)A_t \quad t \in \{0, \pm 1, \pm 2, \dots\}$$

where  $B$  is the backward shift operator,  $\mu$  is the mean of  $W_t$ ,

$$\phi(B) = 1 - \phi_1 B^{l_\phi(1)} - \phi_2 B^{l_\phi(2)} - \dots - \phi_p B^{l_\phi(p)} \quad p \geq 0$$

$$\theta(B) = 1 - \theta_1 B^{l_\theta(1)} - \theta_2 B^{l_\theta(2)} - \dots - \theta_q B^{l_\theta(q)} \quad q \geq 0$$

with  $p = \text{NPAR}$  and  $q = \text{NPMA}$ . Without loss of generality, we assume

$$1 \leq l_\phi(1) \leq l_\phi(2) \leq \dots \leq l_\phi(p)$$

$$1 \leq l_\theta(1) \leq l_\theta(2) \leq \dots \leq l_\theta(q)$$

so that the nonseasonal ARMA model is of order  $(p', q')$  where  $p' = l_\phi(p)$  and  $q' = l_\theta(q)$ . Note that the usual hierarchical model assumes

$$l_\phi(i) = i \quad 1 \leq i \leq p$$

$$l_\theta(j) = j \quad 1 \leq j \leq q$$

Consider the sum of squares function

$$S_T(\mu, \phi, \theta) = \sum_{-T+1}^n$$

where

$$[A_t] = E[A_t | \mu, \phi, \theta, W]$$

and  $T$  is the *backward origin*. The random shocks  $\{A_t\}$  are assumed to be independent and identically distributed

$$N(0, \sigma_A^2)$$

random variables. Hence, the *log-likelihood function* is given by

$$l(\mu, \phi, \theta, \sigma_A) = f(\mu, \phi, \theta) - n \ln \sigma_A - \frac{S_T(\mu, \phi, \theta)}{2\sigma_A^2}$$

where  $f(\mu, \phi, \theta)$  is a function of  $\mu$ ,  $\phi$ , and  $\theta$ .

For  $T = 0$ , the log-likelihood function is *conditional* on the past values of both  $W_t$  and  $A_t$  required to initialize the model. The method of selecting these initial values usually introduces transient bias into the model (Box and Jenkins 1976, pages 210–211). For  $T = \infty$ , this dependency vanishes, and the estimation

problem concerns maximization of the *unconditional* log-likelihood function. Box and Jenkins (1976, page 213) argue that

$$S_{\infty}(\mu, \phi, \theta) / 2\sigma_A^2$$

dominates

$$l(\mu, \phi, \theta, \sigma_A^2)$$

The parameter estimates that minimize the sum of squares function are called *least-squares estimates*. For large  $n$ , the unconditional least-squares estimates are approximately equal to the maximum likelihood estimates.

In practice, a finite value of  $T$  will enable sufficient approximation of the unconditional sum of squares function. The values of  $[A_t]$  needed to compute the unconditional sum of squares are computed iteratively with initial values of  $W_t$  obtained by back-forecasting. The residuals (including backcasts), estimate of random shock variance, and covariance matrix of the final parameter estimates are also computed. Note that application of an appropriate transformation using routine `BCTR` (page 629) followed by differencing using routine `DIFF` (page 633) allows for fitting of nonseasonal ARIMA models. The algorithm for nonseasonal ARIMA models is developed in Chapter 7 of Box and Jenkins (1976). The extension to multiplicative seasonal ARIMA models is given in Box and Jenkins (1976, pages 500–504).

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Routine `NSPE` (page 664) is first invoked to compute preliminary estimates for an ARMA(2, 1) model. Then, `NSLSE` is invoked with the preliminary estimates as input in order to compute the least-squares estimates

$$\hat{\theta}_0 \text{ (output in CONST)}$$

$$\hat{\phi}_1, \hat{\phi}_2, \text{ (output in PAR)}$$

$$\hat{\theta}_1 \text{ (output in PMA)}$$

$$\hat{\sigma}_A^2 \text{ (output in AVAR)}$$

for the ARMA(2, 1) model

$$w_t = \theta_0 + \phi_1 w_{t-1} + \phi_2 w_{t-2} - \theta_1 A_{t-1} + A_t$$

where the errors  $A_t$  are independently distributed each normal with mean zero and variance

$$\sigma_A^2$$

Note at the end of the output a warning error appears. Most of the time this error message can be ignored. There are three general reasons this error can occur.

1. Convergence was declared using the criterion based on TOLSS, but the gradient of the residual sum of squares function was nonzero. This occurred in this example. Either the message can be ignored or TOLSS can be reduced to allow more iterations and a slightly more accurate solution.
2. Convergence is declared based on the fact that a very small step was taken, but the gradient of the residual sum of squares function was nonzero. The message can usually be ignored. However, sometimes the algorithm is making very slow progress and is not near a minimum.
3. Convergence is not declared after 100 iterations.

Examination of the history of iterations using IPRINT = 2 and trying a smaller value for TOLSS can help you determine what caused the error message.

```

INTEGER      IARDEG, IMEAN, LDCOV, LDX, MAXBC, MDX, NOBS, NP,
&            NPAR, NPMA
PARAMETER    (IARDEG=2, IMEAN=1, LDX=176, MAXBC=10, MDX=2,
&            NOBS=100, NPAR=2, NPMA=1, NP=NPAR+NPMA+IMEAN,
&            LDCOV=NP)
C
INTEGER      IPRINT, LAGAR(NPAR), LAGMA(NPMA), MAXIT, NA, NCOL,
&            NROW
REAL         A(NOBS-IARDEG+MAXBC), AVAR, CONST, COV(LDCOV,NP),
&            PAR(NPAR), PMA(NPMA), RELERR, TOLBC, TOLSS, W(NOBS),
&            WMEAN, X(LDX,MDX)
EXTERNAL     GDATA, NSLSE, NSPE
C
EQUIVALENCE (W(1), X(22,2))
C
DATA LAGAR/1, 2/, LAGMA/1/
C
C                                Wolfer Sunspot Data for
C                                years 1770 through 1869
IPRINT = 0
CALL GDATA (2, IPRINT, NROW, NCOL, X, LDX, MDX)
C
C                                Convergence parameters
MAXIT = 0
RELERR = 0.0
C
C                                Compute preliminary parameter
C                                estimates for ARMA(2,1) model
IPRINT = 1
CALL NSPE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, NPMA, RELERR,
&         MAXIT, CONST, PAR, PMA, AVAR)
C
TOLBC = 0.0
TOLSS = 0.125
IPRINT = 2
C
CALL NSLSE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, PAR, LAGAR,
&         NPMA, PMA, LAGMA, MAXBC, TOLBC, TOLSS, CONST, COV,
&         LDCOV, NA, A, AVAR)
C
END

```

## Output

Results from NSPE/N2PE

WMEAN = 46.9760  
CONST = 15.5440  
AVAR = 287.242

PAR  
1 2  
1.244 -0.575

PMA  
-0.1241

-----  
Iteration 1

WMEAN = 52.638233185

PAR  
1 2  
1.264 -0.606

PMA  
-0.1731

Residual SS (including backcasts) = 23908.66210937500  
Number of residuals = 108  
Number of backcasts = 10

-----  
Iteration 2

WMEAN = 54.756504059

PAR  
1 2  
1.360 -0.688

PMA  
-0.1411

Residual SS (including backcasts) = 23520.71484375000  
Number of residuals = 108  
Number of backcasts = 10

-----  
Final Results, Iteration 3

Parameter	Estimate	Std. Error	t-ratio
WMEAN	53.9187279	5.5178852	9.7716293
		PAR	
1	1.3925704	0.0960639	14.4962845
2	-0.7329484	0.0866115	-8.4624796
		PMA	
1	-0.1375125	0.1223797	-1.1236545
CONST =	18.3527489		

AVAR = 243.4830170

Residual SS (including backcasts) = 23374.3691406  
Number of residuals = 108

Residual SS (excluding backcasts) = 20931.7519531  
Number of residuals = 98

\*\*\* WARNING ERROR 1 from NSLSE. Least squares estimation of the parameters  
\*\*\* has failed to converge. Increase MAXBC and/or TOLBC and/or  
\*\*\* TOLSS. The estimates of the parameters at the last iteration  
\*\*\* may be used as new starting values.

---

## SPWF/DSPWF (Single/Double precision)

Compute the Wiener forecast operator for a stationary stochastic process.

### Usage

CALL SPWF (NOBS, W, IWMEAN, WMEAN, WNADJ, EPS, MLFOP, LFOP,  
FOP)

### Arguments

**NOBS** — Number of observations in the stationary time series  $w$ . (Input)  
NOBS must be greater than or equal to two.

**W** — Vector of length NOBS containing the stationary time series. (Input)

**IWMEAN** — Option for estimation of the mean of  $w$ . (Input)

#### **IWMEAN Action**

0 WMEAN is user specified.

1 WMEAN is set equal to the arithmetic mean of  $w$ .

**WMEAN** — Estimate of the mean of the time series  $w$ . (Input, if IWMEAN = 0;  
output, if IWMEAN = 1)

WMEAN is used to center the time series  $w$  prior to estimation of the forecast  
operator.

**WNADJ** — White noise adjustment factor. (Input)

WNADJ must be greater than or equal to zero.

**EPS** — Bound on the normalized mean square error. (Input)

EPS must be in the range (0, 1) inclusive.

**MLFOP** — Maximum length of the forecast operator. (Input)

MLFOP must be greater than or equal to one and less than NOBS.

**LFOP** — Length of the estimated forecast operator. (Output)

**FOP** — Vector of length LFOP containing the estimated forecast operator  
coefficients. (Output)

## Comments

1. Automatic workspace storage is

SPWF NOBS + 2 \* MLFOP + 1 units, or  
DSPWF 2 \* (NOBS + 2 \* MLFOP + 1) units.

Workspace may be explicitly provided, if desired, by use of S2WF/DS2WF. The reference is

```
CALL S2WF (NOBS, W, IWMEAN, WMEAN, WNADJ, EPS,  
          MLFOP, LFOP, FOP, CW, WK)
```

The additional arguments are as follows:

**CW** — Vector of length NOBS containing the centered time series  $w$ .  
(Output)

**WK** — Vector of length 2 \* MLFOP + 1. (Output)

2. Informational error

Type	Code	
3	5	No operator could be found of length less than or equal to MLFOP that produced a normalized mean square error less than EPS.

- 3 The length of the forecast operator is determined by the arguments EPS and MLFOP. Iteration to a longer forecast operator stops when either the normalized mean square error is less than EPS, or the operator reaches the maximum allowable length, MLFOP.

- 4 The white noise adjustment factor, WNADJ, is used to modify the estimate of the variance of the time series  $w$  used in the computation of the autocorrelation function of  $w$ . In the absence of white noise, WNADJ should be set to zero.

## Algorithm

Routine SPWF performs least-squares estimation of parameters for successive autoregressive models of a stationary stochastic process given a sample of  $n = \text{NOBS}$  observations  $\{W_t\}$  for  $t = 1, \dots, n$ .

Let

$$\hat{\mu} = \text{WMEAN}$$

be the estimate of the mean  $\mu$  of the stochastic process  $\{W_t\}$  where

$$\hat{\mu} = \begin{cases} \mu & \mu \text{ known} \\ \frac{1}{n} \sum_{t=1}^n W_t & \mu \text{ unknown} \end{cases}$$

Consider the autoregressive model of order  $k$  defined by

$$\phi_k(B)\tilde{W}_t = A_t \quad k \geq 0$$

where

$$\tilde{W}_t = W_t - \hat{\mu}$$

and

$$\phi_k(B) = 1 - \phi_{1k}B - \phi_{2k}B^2 - \dots - \phi_{kk}B^k \quad k \geq 1$$

Successive AR( $k$ ) models are fit to the centered data using Durbin's algorithm (1960) based on the sample autocovariances

$$\hat{\sigma}(k) = \frac{1}{n} \sum_{t=1}^{n-k} (W_t - \hat{\mu})(W_{t+k} - \hat{\mu}) \quad k \geq 0$$

Note that the variance

$$\hat{\sigma}^*(0)$$

used in the fitting algorithm is adjusted by the amount  $\delta = \text{WNADJ}$  according to

$$\hat{\sigma}^*(0) = (1 + \delta)\hat{\sigma}(0)$$

See Robinson (1967, page 96).

Iteration to the next higher order model terminates when either the expected mean square error of the model is less than `EPS` or when  $k = \text{MLFOP}$ . The forecast operator  $\phi = (\phi_1, \phi_2, \dots, \phi_{k^*})^T$  for  $k^* = \text{LFOP}$  is contained in `FOP`. See also Craddock (1969).

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Application of routine `SPWF` to these data produces the following results:

```

INTEGER      MLFOP, NOBS
PARAMETER    (MLFOP=1, NOBS=100)
C           INTEGER      I, IMEAN, LFOP, NCOL, NOUT, NROW

REAL         EPS, FOP(MLFOP), RDATA(176,2), W(NOBS), WMEAN, WNADJ
EXTERNAL     GDATA, SPWF, UMACH
C
EQUIVALENCE (W(1), RDATA(22,2))
C                                     Wolfer Sunspot Data for
C                                     years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C                                     Center on arithmetic mean
IMEAN = 0
WMEAN = 46.976
C                                     White noise adjustment
WNADJ = 0.0
C                                     Bound on normalized MSE

```



```

      EPS = 0.1
C          Determine autoregressive model
      CALL SPWF (NOBS, W, IMEAN, WMEAN, WNADJ, EPS, MLFOP, LFOP, FOP)
C          Print results
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99997) LFOP
99997  FORMAT (/, 1X, 'Forecast operator length, LFOP = ', I2)
      WRITE (NOUT,99998)
99998  FORMAT (/, 1X, ' I          FOP(I)')
      DO 10 I=1, LFOP
          WRITE (NOUT,99999) I, FOP
99999  FORMAT (1X, I2, 2X, F12.4)
      10 CONTINUE
C
      END

```

### Output

```

*** WARNING  ERROR 5 from SPWF.  No operator could be found of length less
***          than or equal to 1 which produced a normalized mean square
***          error less than 1.000000E-01.

```

Forecast operator length, LFOP = 1

```

I          FOP(I)
1          0.8063

```

---

## NSBJF/DNSBJF (Single/Double precision)

Compute Box-Jenkins forecasts and their associated probability limits for a nonseasonal ARMA model.

### Usage

```

CALL NSBJF (NOBS, W, IPRINT, NPAR, PAR, LAGAR, NPMA, PMA,
           LAGMA, ICONST, CONST, AVAR, ALPHA, MXBKOR,
           MXLEAD, FCST, LDFCST)

```

### Arguments

**NOBS** — Number of observations in the time series *W*. (Input)  
NOBS must be greater than  $ICONST + \max(LAGAR(i)) + \max(LAGMA(j))$ .

**W** — Vector of length NOBS containing the time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.  
1 Print the forecasts for lead times  $l = 1, \dots, MXLEAD$  at each origin  $t = (NOBS - MXBKOR), \dots, NOBS$ , the  $100(1 - ALPHA)\%$  probability limit deviations, and the psi weights.

**NPAR** — Number of autoregressive parameters. (Input)  
NPAR must be greater than or equal to zero.

**PAR** — Vector of length *NPAR* containing the autoregressive parameters. (Input)

**LAGAR** — Vector of length *NPAR* containing the order of the autoregressive parameters. (Input)

The elements of *LAGAR* must be greater than zero.

**NPMA** — Number of moving average parameters. (Input)

*NPMA* must be greater than or equal to zero.

**PMA** — Vector of length *NPMA* containing the moving average parameters. (Input)

**LAGMA** — Vector of length *NPMA* containing the order of the moving average parameters. (Input)

The elements of *LAGMA* must be greater than zero.

**ICONST** — Option for including the overall constant in the model. (Input)

**ICONST Action**

0 No overall constant is included.

1 The overall constant is included.

**CONST** — Estimate of the overall constant. (Input)

**AVAR** — Estimate of the random shock variance. (Input)

*AVAR* must be greater than 0.

**ALPHA** — Value in the exclusive interval (0, 1) used to specify the 100(1 - *ALPHA*)% probability limits of the forecasts. (Input)

Typical choices for *ALPHA* are 0.10, 0.05, and 0.01.

**MXBKOR** — Maximum backward origin. (Input)

*MXBKOR* must be greater than or equal to zero and less than or equal to

*NOBS* - max(*MAXAR*, *MAXMA*) where *MAXAR* = max(*LAGAR*(*i*)) and

*MAXMA* = max(*LAGMA*(*j*)). Forecasts at origins *NOBS* - *MXBKOR* through *NOBS* are generated.

**MXLEAD** — Maximum lead time for forecasts. (Input)

*MXLEAD* must be greater than zero.

**FCST** — *MXLEAD* by (*MXBKOR* + 3) matrix defined below. (Output)

Column	Content
<i>j</i>	Forecasts for lead times $l = 1, \dots, \text{MXLEAD}$ at origins $\text{NOBS} - \text{MXBKOR} - 1 + j, j = 1, \dots, \text{MXBKOR} + 1$ .
<i>MXBKOR</i> + 2	Deviations from each forecast that give the 100(1 - <i>ALPHA</i> )% probability limits.
<i>MXBKOR</i> + 3	Psi weights of the infinite order moving average form of the model.

**LDFCST** — Leading dimension of *FCST* exactly as specified in the dimension statement in the calling program. (Input)

*LDFCST* must be greater than or equal to *MXLEAD*.

## Comments

1. Automatic workspace usage is

NSBJF  $5 + \text{IARDEG} + \text{IMADEG} + 3 * \text{MXLEAD}$  units, or

DNSBJF  $7 + 2 * \text{IARDEG} + 2 * \text{IMADEG} + 5 * \text{MXLEAD}$  units, where  
 $\text{IARDEG} = \max(\text{LAGAR}(i))$  and  $\text{IMADEG} = \max(\text{LAGMA}(j))$ .

Workspace may be explicitly provided, if desired, by use of  
N2BJF/DN2BJF. The reference is

```
CALL N2BJF (NOBS, W, IPRINT, NPAR, PAR, LAGAR, NPMA,  
           PMA, LAGMA, ICONST, CONST, AVAR, ALPHA,  
           MXBKOR, MXLEAD, FCST, LDFCST, PARH,  
           PMAH, PSIH, PSI, LAGPSI)
```

The additional arguments are as follows:

**PARH** — Work vector of length equal to  $\text{IARDEG} + 1$ .

**PMAH** — Work vector of length equal to  $\text{IMADEG} + 1$ .

**PSIH** — Work vector of length equal to  $\text{MXLEAD} + 1$ .

**PSI** — Work vector of length equal to  $\text{MXLEAD} + 1$ .

**LAGPSI** — Work vector of length equal to  $\text{MXLEAD} + 1$ .

2. If the  $w$  series has been transformed using a Box-Cox transformation with parameters **POWER** and **SHIFT**, the forecasts and probability limits for the original series may be obtained by application of routine **BCTR** (page 629) with the same parameters and argument **IDIR** set equal to one.

## Algorithm

Routine **NSBJF** computes Box-Jenkins forecasts and their associated probability limits for a nonseasonal ARMA model given a sample of  $n = \text{NOBS}$  observations  $\{W_t\}$  for  $t = 1, 2, \dots, n$ .

Suppose the time series  $\{W_t\}$  is generated by a nonseasonal ARMA model of the form

$$\phi(B)W_t = \theta_0 + \theta(B)A_t \quad t \in \{0, \pm 1, \pm 2, \dots\}$$

where  $B$  is the backward shift operator,  $\theta_0 = \text{CONST}$ ,

$$\phi(B) = 1 - \phi_1 B^{l_\phi(1)} - \phi_2 B^{l_\phi(2)} - \dots - \phi_p B^{l_\phi(p)}$$

$$\theta(B) = 1 - \theta_1 B^{l_\theta(1)} - \theta_2 B^{l_\theta(2)} - \dots - \theta_q B^{l_\theta(q)}$$

$p = \text{NPAR}$  and  $q = \text{NPMA}$ . Without loss of generality, we assume

$$1 \leq l_\phi(1) \leq l_\phi(2) \leq \dots \leq l_\phi(p)$$

$$1 \leq l_\theta(1) \leq l_\theta(2) \leq \dots \leq l_\theta(q)$$

so that the nonseasonal ARMA model is of order  $(p', q')$  where  $p' = l_\phi(p)$  and  $q' = l_\theta(q)$ . Note that the usual hierarchical model assumes

$$l_\phi(i) = i \quad 1 \leq i \leq p$$

$$l_\theta(j) = j \quad 1 \leq j \leq q$$

The Box-Jenkins forecast at origin  $t$  for lead time  $l$  of  $W_{t+l}$  is defined in terms of the difference equation

$$\begin{aligned} \hat{W}_t(l) = & \theta_0 + \phi_1[W_{t+l-l_\phi(1)}] + \dots + \phi_p[W_{t+l-l_\phi(p)}] \\ & + [A_{t+l}] - \theta_1[A_{t+l-l_\theta(1)}] - \dots - \theta_q[A_{t+l-l_\theta(q)}] \end{aligned}$$

where

$$[W_{t+k}] = \begin{cases} W_{t+k} & k = 0, -1, -2, \dots \\ \hat{W}_t(k) & k = 1, 2, \dots \end{cases}$$

$$[A_{t+k}] = \begin{cases} W_{t+k} - \hat{W}_{t+k-1}(1) & k = 0, -1, -2, \dots \\ 0 & k = 1, 2, \dots \end{cases}$$

The  $100(1 - \alpha)\%$  probability limits for  $W_{t+l}$  are given by

$$\hat{W}_t(l) \pm z_{\alpha/2} \left\{ 1 + \sum_{j=1}^{l-1} \psi_j^2 \right\}^{1/2} \sigma_A$$

where  $z_{(1-\alpha/2)}$  is the  $100(1 - \alpha/2)$  percentile of the standard normal distribution,

$$\sigma_A^2 = \text{AVAR}$$

and  $\{\psi_j\}$  are the parameters of the random shock form of the difference equation. Note that the forecasts are computed for lead times  $l = 1, 2, \dots, L$  at origins  $t = (n - b), (n - b + 1), \dots, n$  where  $L = \text{MXLEAD}$  and  $b = \text{MXBKOR}$ .

The Box-Jenkins forecasts minimize the mean square error

$$E[W_{t+l} - \hat{W}_t(l)]^2$$

Also, the forecasts may be easily updated according to the equation

$$\hat{W}_{t+1}(l) = \hat{W}_t(l+1) + \psi_l A_{t+1} \quad (7)$$

This approach and others are given in Chapter 5 of Box and Jenkins (1976).

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Routine NSBJF is used to compute forecasts and 95% probability limits for the forecasts for an ARMA(2, 1) model fit using routine NSPE (page 664). With  $MXBKOR = 3$ , columns one through four of FCST give forecasts given the data through 1866, 1867, 1868, and 1869, respectively. Column 5 gives the deviations from the forecast for computing probability limits, and column six gives the psi weights, which can be used to update forecasts once more data is available. For example, the forecast for the 102-nd observation (year 1871) given the data through the 100-th observation (year 1869) is 77.21, and 95% probability limits are given by  $77.21 \pm 56.30$ . After observation 101 ( $W_{101}$  for year 1870) is available, the forecast can be updated by using equation 7 with the psi weight ( $\psi_1 = 1.37$ ) and the one-step-ahead forecast error for observation 101 ( $W_{101} - 83.72$ ) to give

$$77.21 + 1.37 (W_{101} - 83.72)$$

Since this updated forecast is one step ahead, the 95% probability limits are now given by the forecast  $\pm 33.22$ .

```

INTEGER      LDFCST, MXBKOR, MXLEAD, NOBS, NPAR, NPMA
PARAMETER    (MXBKOR=3, MXLEAD=12, NOBS=100, NPAR=2, NPMA=1,
&            LDFCST=MXLEAD)
C
INTEGER      ICONST, IMEAN, IPRINT, LAGAR(NPAR), LAGMA(NPMA),
&            MAXIT, NCOL, NROW
REAL         ALPHA, AVAR, CONST, FCST(LDFCST,MXBKOR+3), PAR(NPAR),
&            PMA(NPMA), RDATA(176,2), RELERR, W(NOBS), WMEAN
CHARACTER    CLABEL(MXBKOR+4)*40, RLABEL(1)*6
EXTERNAL     GDATA, NSBJF, NSPE, WRRRL
C
EQUIVALENCE (W(1), RDATA(22,2))
C
DATA LAGAR(1), LAGAR(2)/1, 2/
DATA LAGMA(1)/1/
DATA RLABEL/'NUMBER'/, CLABEL/'%/Lead%/Time',
&          '%/Forecast%/From 1866', '%/Forecast%/From 1867',
&          '%/Forecast%/From 1868', '%/Forecast%/From 1869',
&          ' Deviation %/ for 95% %/Prob. Limits', '%/%/Psi'/
C
C                               Wolfer Sunspot Data for
C                               years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C                               Compute preliminary parameter
C                               estimates for ARMA(2,1) model
C
IMEAN  = 1
MAXIT  = 0
RELERR = 0.0
IPRINT = 0
CALL NSPE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, NPMA, RELERR,
&          MAXIT, CONST, PAR, PMA, AVAR)
C
C                               Include constant in forecast model

```

```

        ICONST = 1
C
C                               Specify 95 percent probability
C                               limits for forecasts
        ALPHA = 0.05
C
C                               Compute forecasts
        IPRINT = 0
        CALL NSBJF (NOBS, W, IPRINT, NPAR, PAR, LAGAR, NPMA, PMA, LAGMA,
&                  ICONST, CONST, AVAR, ALPHA, MXBKOR, MXLEAD, FCST,
&                  LDFCST)
C
C                               Print results
        CALL WRRRL ('FCST', MXLEAD, MXBKOR+3, FCST, LDFCST, 0,
&                  '(5F9.2,F6.3)', RLABEL, CLABEL)
C
        END

```

### Output

Lead Time	FCST				Deviation for 95%		Psi
	Forecast From 1866	Forecast From 1867	Forecast From 1868	Forecast From 1869	Prob. Limits		
1	18.28	16.62	55.19	83.72	33.22	1.368	
2	28.92	32.02	62.76	77.21	56.30	1.127	
3	41.01	45.83	61.89	63.46	67.62	0.616	
4	49.94	54.15	56.46	50.10	70.64	0.118	
5	54.09	56.56	50.19	41.38	70.75	-0.208	
6	54.13	54.78	45.53	38.22	71.09	-0.326	
7	51.78	51.17	43.32	39.30	71.91	-0.286	
8	48.84	47.71	43.26	42.46	72.53	-0.169	
9	46.53	45.47	44.46	45.77	72.75	-0.045	
10	45.35	44.69	45.98	48.08	72.77	0.041	
11	45.21	44.99	47.18	49.04	72.78	0.077	
12	45.71	45.82	47.81	48.91	72.82	0.072	

---

## IRNSE/DIRNSE (Single/Double precision)

Compute estimates of the impulse response weights and noise series of a univariate transfer function model.

### Usage

```
CALL IRNSE (NOBS, X, Y, IPRINT, NPAR, PAR, NPMA, PMA,
           MWTIR, MWTSN, WTIR, SNOISE, XPW, YPW)
```

### Arguments

**NOBS** — Number of observations in each time series. (Input)

NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the input time series. (Input)

**Y** — Vector of length NOBS containing the output time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.

1            Print the estimates of the impulse response weights and the noise series.

**NPAR** — Number of prewhitening autoregressive parameters. (Input)  
NPAR must be greater than or equal to zero.

**PAR** — Vector of length NPARG containing the prewhitening autoregressive parameters. (Input)

**NPMA** — Number of prewhitening moving average parameters. (Input)  
NPMA must be greater than or equal to zero.

**PMA** — Vector of length NPMA containing the prewhitening moving average parameters. (Input)

**MWTIR** — Maximum index of the impulse response weights. (Input)  
MWTIR must be greater than or equal to zero and less than or equal to NOBS - 1.

**MWTSN** — Maximum index of the impulse response weights used to compute the noise series. (Input)  
MWTSN must be greater than or equal to zero and less than or equal to MWTIR.

**WTIR** — Vector of length MWTIR + 1 containing the impulse response weight estimates. (Output)  
The impulse response weight estimate of index  $k$  is given by  $WTIR(k)$  for  $k = 0, 1, \dots, MWTIR$ .

**SNOISE** — Vector of length NOBS - MWTSN containing the noise series based on the impulse response weight estimates. (Output)

**XPW** — Vector of length NOBS - NPARG containing the prewhitened input time series  $X$ . (Output)

**YPW** — Vector of length NOBS - NPARG containing the prewhitened output time series  $Y$ . (Output)

### Comments

1.            Automatic workspace usage is

IRNSE 3 + 4 \* MWTIR units, or  
DIRNSE 6 + 8 \* MWTIR units.

Workspace may be explicitly provided, if desired, by use of  
I2NSE/DI2NSE. The reference is

```
CALL I2NSE (NOBS, X, Y, IPRINT, NPARG, PAR, NPMA,  
           PMA, MWTIR, MWTSN, WTIR, SNOISE, XPW,  
           YPW, ACPWX, ACPWY, CCPW)
```

The additional arguments are as follows:

**ACPWX** — Vector of length MWTIR + 1 containing the estimated autocorrelation function of  $PWX$ . (Output)

**ACPWY** — Vector of length MWTIR + 1 containing the estimated autocorrelation function of  $PWY$ . (Output)

**CCPW** — Vector of length  $2 * \text{MWTIR} + 1$  containing the estimated cross-correlation function of **PWX** and **PWY**. (Output)

2. The input series **X** and output series **Y** are assumed to be the result of transforming and differencing the raw input and output series. The routines **BCTR** (page 629) and **DIFF** (page 633) may be used, respectively, to perform a Box-Cox transformation and difference the raw input and output series.
3. Note that the prewhitened input and output are computed at time  $t = \text{NPAR} + 1$  through  $t = \text{NOBS}$ . Also, the noise series is computed at time  $t = \text{MWTSN} + 1$  through  $t = \text{NOBS}$ .

### Algorithm

Routine **IRNSE** estimates the impulse response weights and noise series of a transfer function model given a sample of  $n = \text{NOBS}$  observations of the input  $\{x_t\}$  and output  $\{y_t\}$  for  $t = 1, 2, \dots, n$ . Define  $\{x_t\}$  and  $\{y_t\}$ , respectively, by

$$x_t = \begin{cases} X_t - \hat{\mu}_X & d = 0 \\ \nabla^d X_t & d > 0 \end{cases}$$

and

$$y_t = \begin{cases} Y_t - \hat{\mu}_Y & d = 0 \\ \nabla^d Y_t & d > 0 \end{cases}$$

where  $\{X_t\}$  and  $\{Y_t\}$  for  $t = (-d + 1), \dots, n$  represent the undifferenced input and output series with

$$\hat{\mu}_X \text{ and } \hat{\mu}_Y$$

estimates of their respective means. The differenced input and output series may be obtained using the routine **DIFF** (page 633) following any preliminary transformation of the data.

The transfer function model is defined by

$$Y_t = v(B)X_t + N_t$$

or, equivalently,

$$y_t = v(B)x_t + n_t$$

with transfer function

$$v(B) = v_0 + v_1 B + v_2 B^2 + \dots$$

and differenced noise series  $n_t = \nabla^d N_t$ .

The prewhitened input and output series are computed for  $t = (p + 1), \dots, n$  according to



$$\alpha_t = \phi(B)x_t + \theta_1(B)\alpha_t$$

$$\beta_t = \phi(B)y_t + \theta_1(B)\beta_t$$

where

$$\phi(B) = 1 - \phi_1 B - \phi_2 B^2 - \dots - \phi_p B^p$$

$$\theta(B) = \theta_1 B + \theta_2 B^2 + \dots + \theta_q B^q$$

The parameters of the prewhitening transformation may be estimated roughly using the routine NSPE (page 664) or more precisely using the routine NSLSE (page 669). The correlation relationship between  $\{\alpha_t\}$ ,  $\{\beta_t\}$ , and  $\{n_t\}$  may be further examined using the routines ACF (page 637), PACF (page 641), and CCF (page 644).

The *impulse response weights*  $\{v_k\}$  are estimated by

$$\hat{v}_k = \frac{\hat{\sigma}_\beta}{\hat{\sigma}_\alpha} \hat{\rho}_{\alpha\beta}(k) \quad k = 0, 1, \dots, K$$

where  $K = \text{MWTIR}$ ,

$$\hat{\sigma}_\alpha \text{ and } \hat{\sigma}_\beta$$

denote the standard deviation of  $\{\alpha_t\}$  and  $\{\beta_t\}$ ;

$$\hat{\rho}_{\alpha\beta}(k) \text{ for } k = 0, 1, \dots, K$$

represents the cross-correlation function between  $\{\alpha_t\}$  and  $\{\beta_t\}$ . The differenced noise series  $\{n_t\}$  for  $t = (K' + 1), \dots, n$  is reconstructed using the model

$$n_t = y_t - \hat{v}_0 x_t - \hat{v}_1 x_{t-1} - \dots - \hat{v}_{K'} x_{t-K'}$$

where  $K' = \text{MWTSN}$ .

### Example

Consider the Gas Furnace Data (Box and Jenkins 1976, pages 532–533) where  $X$  is the input gas rate in cubic feet/minute and  $Y$  is the percent  $\text{CO}_2$  in the outlet gas. Application of routine IRNSE to these data produces the following results:

```

C      INTEGER      IPRINT, LDX, MWTIR, MWTSN, NDX, NOBS, NOPRIN, NPAR,
&      NPMA
C      PARAMETER    ( IPRINT=0, LDX=296, MWTIR=10, NDX=2, NOBS=296,
&      NOPRIN=0, NPAR=3, NPMA=0, MWTSN=MWTIR)
C
C      INTEGER      NCOL, NROW
C      REAL         PAR(NPAR), PMA(1), RDATA(296,2), SNOISE(NOBS-MWTSN),
&      WTIR(MWTIR+1), X(NOBS), XPW(NOBS-NPAR), Y(NOBS),
&      YPW(NOBS-NPAR)
C      EXTERNAL     GDATA, IRNSE, WRRRN
C
C      EQUIVALENCE (X(1), RDATA(1,1)), (Y(1), RDATA(1,2))
C                                     Gas Furnace Data

```

```

CALL GDATA (7, NOPRIN, NROW, NCOL, RDATA, LDX, NDX)
C                                     Specify AR parameters for
C                                     prewhitening transformation
PAR(1) = 1.97
PAR(2) = -1.37
PAR(3) = 0.34
C                                     Compute estimates of impulse
C                                     response weights and noise series
CALL IRNSE (NOBS, X, Y, IPRINT, NPAR, PAR, NPMA, PMA, MWTIR,
&          MWTSN, WTIR, SNOISE, XPW, YPW)
C                                     Print results
CALL WRRRN ('WTIR', 1, 11, WTIR, 1, 0)
CALL WRRRN ('SNOISE', 1, 20, SNOISE, 1, 0)
C
END

```

### Output

```

                                     WTIR
      1      2      3      4      5      6      7      8
-0.0355  0.0716 -0.0764 -0.5655 -0.6549 -0.8936 -0.5358 -0.3482

      9     10     11
-0.0782  0.0277 -0.1364

                                     SNOISE
      1      2      3      4      5      6      7      8      9     10
53.21  53.49  53.72  54.05  53.98  53.95  53.69  53.02  52.56  52.33

     11     12     13     14     15     16     17     18     19     20
52.47  52.69  52.57  52.63  52.81  53.14  53.21  53.20  53.05  52.88

```

---

## TFPE/DTFPE (Single/Double precision)

Compute preliminary estimates of parameters for a univariate transfer function model.

### Usage

```

CALL TFPE (IPRINT, NPLHS, NPRHS, NPNAR, NPNMA, NDELAY,
          MWTIR, WTIR, NSNOIS, SNOISE, RELERR, MAXIT,
          PLHS, PRHS, PNAR, PNMA, AVAR)

```

### Arguments

**IPRINT** — Printing option. (Input)

#### IPRINT Action

- 0 No printing is performed.
- 1 Print estimates of transfer function parameters, estimates of noise model parameters, and an of the random shock variance.

**NPLHS** — Number of left-hand side transfer function parameters. (Input)  
NPLHS must be greater than or equal to zero.

***NPRHS*** — Number of right-hand side transfer function parameters (excluding the index 0 parameter). (Input)

*NPRHS* must be greater than or equal to zero.

***NPNAR*** — Number of noise autoregressive parameters. (Input)

*NPNAR* must be greater than or equal to zero.

***NPNMA*** — Number of noise moving average parameters. (Input)

*NPNMA* must be greater than or equal to zero.

***NDELAY*** — Time delay parameter. (Input)

*NDELAY* must be greater than or equal to zero.

***MWTIR*** — Maximum index of the impulse response weights. (Input)

*MWTIR* must be greater than or equal to *NPLHS* + *NPRHS* + *NDELAY*.

***WTIR*** — Vector of length *MWTIR* + 1 containing the impulse response weight estimates. (Input)

The impulse response weight estimate of index *k* is given by *WTIR*(*k*) for *k* = 0, 1, ..., *MWTIR*.

***NSNOIS*** — Number of elements in the noise series. (Input)

*NSNOIS* must be greater than or equal to *NPNAR* + *NPNMA* + 1.

***SNOISE*** — Vector of length *NSNOIS* containing the noise series. (Input)

***RELERR*** — Stopping criterion for use in the nonlinear equation solver. (Input)

If *RELERR* = 0.0, then the default value *RELERR* = 100.0 \* *AMACH*(4) is used. See the documentation for routine *AMACH* (page 1334).

***MAXIT*** — The maximum number of iterations allowed in the nonlinear equation solver. (Input)

If *MAXIT* = 0, then the default value *MAXIT* = 200 is used.

***PLHS*** — Vector of length *NPLHS* containing the estimates of the left-hand side transfer function parameters. (Output)

The LHS weight estimates are *PLHS*(*k*), *k* = 1, ..., *NPLHS*.

***PRHS*** — Vector of length *NPRHS* + 1 containing the estimates of the right-hand side transfer function parameters. (Output)

The RHS weight estimates are *PRHS*(*k*), *k* = 0, ..., *NPRHS*.

***PNAR*** — Vector of length *NPNAR* containing the estimates of the noise autoregressive parameters. (Output)

***PNMA*** — Vector of length *NPNMA* containing the estimates of the noise moving average parameters. (Output)

***AVAR*** — Estimate of the random shock variance. (Output)

## Comments

1. Automatic workspace usage is

TFPE  $2 * \max(\text{NPLHS}, \text{NPNAR}) * (\max(\text{NPLHS}, \text{NPNAR}) + 1) + 2 * \text{NPNAR} + 1 + (\text{NPNMA} + 1) * (13 + 1.5 * \text{NPNMA}) + \text{NPLHS units,}$   
or

DTFPE  $2 * (2 * \max(\text{NPLHS}, \text{NPNAR}) * (\max(\text{NPLHS}, \text{NPNAR}) + .75) + 2 * \text{NPNAR} + 1 + (\text{NPNMA} + 1) * (13 + 1.5 * \text{NPNMA}) + \text{NPLHS})$   
units.

Workspace may be explicitly provided, if desired, by use of T2PE/DT2PE. The reference is

```
CALL T2PE ( IPRINT, NPLHS, NPRHS, NPNAR, NPNMA,
            NDELAY, MWTIR, WTIR, NSNOIS, SNOISE,
            RELERR, MAXIT, PLHS, PRHS, PNAR, PNMA,
            AVAR, A, FAC, IPVT, WK, ACV, PARWK,
            ACVMOD, TAUINI, TAU, FVEC, FJAC, R, QTF,
            WKNLN, H)
```

The additional arguments are as follows:

*A* — Work vector of length  $(\max(\text{NPLHS}, \text{NPNAR}))^2$ .

*FAC* — Work vector of length  $(\max(\text{NPLHS}, \text{NPNAR}))^2$ .

*IPVT* — Work vector of length  $\max(\text{NPLHS}, \text{NPNAR})$ .

*WK* — Work vector of length  $\max(\text{NPLHS}, \text{NPNAR})$ .

*ACV* — Work vector of length  $\text{NPNAR} + \text{NPNMA} + 1$ .

*PARWK* — Work vector of length  $\text{NPNAR} + 1$ .

*ACVMOD* — Work vector of length  $\text{NPNMA} + 1$ .

*TAUINI* — Work vector of length  $\text{NPNMA} + 1$ .

*TAU* — Work vector of length  $\text{NPNMA} + 1$ .

*FVEC* — Work vector of length  $\text{NPNMA} + 1$ .

*FJAC* — Work vector of length  $(\text{NPNMA} + 1)^2$ .

*R* — Work vector of length  $(\text{NPNMA} + 1) * (\text{NPNMA} + 2)/2$ .

*QTF* — Work vector of length  $\text{NPNMA} + 1$ .

*WKNLN* — Work vector of length  $5 * (\text{NPNMA} + 1)$ .

*H* — Work vector of length  $\text{NPLHS}$ .

2. Informational error

Type	Code	
4	1	The nonlinear equation solver did not converge to RELERR within MAXIT iterations.

3. The impulse response weight estimates and the noise series may be computed using routine IRNSE (page 685).

### Algorithm

Routine `TFPE` computes preliminary estimates of the parameters of a transfer function model given a sample of  $n = \text{NOBS}$  observations of the differenced input  $\{x_t\}$  and differenced output  $\{y_t\}$  for  $t = 1, 2, \dots, n$ .

Define  $\{x_t\}$  and  $\{y_t\}$ , respectively, by

$$x_t = \begin{cases} X_t - \hat{\mu}_X & d = 0 \\ \nabla^d X_t & d > 0 \end{cases}$$

and

$$y_t = \begin{cases} Y_t - \hat{\mu}_Y & d = 0 \\ \nabla^d Y_t & d > 0 \end{cases}$$

where  $\{X_t\}$  and  $\{Y_t\}$  for  $t = (-d + 1), \dots, n$  represent the undifferenced input and output series with

$$\hat{\mu}_X \text{ and } \hat{\mu}_Y$$

estimates of their respective means. The differenced input and output series may be obtained using the routine `DIFF` (page 633) following any preliminary transformation of the data.

The transfer function model is defined by

$$Y_t = \delta^{-1}(B)\omega(B)X_{t-b} + N_t$$

or, equivalently,

$$y_t = \delta^{-1}(B)\omega(B)x_{t-b} + n_t$$

where  $n_t = \nabla^d N_t$  and the left-hand side and right-hand side transfer function polynomial operators are

$$\begin{aligned} \delta(B) &= 1 - \delta_1 B - \delta_2 B^2 - \dots - \delta_r B^r \\ \omega(B) &= \omega_0 - \omega_1 B - \omega_2 B^2 - \dots - \omega_s B^s \end{aligned}$$

with  $r = \text{NPLHS}$ ,  $s = \text{NPRHS}$ , and  $b = \text{NDELAY}$ . The noise process  $\{N_t\}$  and the input process  $\{X_t\}$  are assumed to be independent with the noise process given by the ARIMA model

$$\phi(B)n_t = \theta(B)A_t$$

where

$$\begin{aligned} \phi(B) &= 1 - \phi_1 B - \phi_2 B^2 - \dots - \phi_p B^p \\ \theta(B) &= 1 - \theta_1 B - \theta_2 B^2 - \dots - \theta_q B^q \end{aligned}$$

with  $p = \text{NPNAR}$  and  $q = \text{NPNMA}$ .

The impulse response weights and the transfer function parameters are related by

$$v_k = \begin{cases} 0 & k = 0, 1, \dots, b-1 \\ \sum_{j=1}^r \delta_j v_{k-j} + \omega_0 & k = b \\ \sum_{j=1}^r \delta_j v_{k-j} - \omega_{k-b} & k = b+1, b+2, \dots, b+s \\ \sum_{j=1}^r \delta_j v_{k-j} & k = b+s+1, b+s+2, \dots \end{cases}$$

See Abraham and Ledolter (1983, page 341). The  $r$  left-hand side transfer function parameters are estimated using the difference equation given as the last case above. The resulting estimates

$$\hat{\delta}_1, \dots, \hat{\delta}_r$$

are then substituted into the middle two cases to determine the  $s+1$  estimates

$$\hat{\omega}_0, \hat{\omega}_1, \dots, \hat{\omega}_s$$

The noise series parameters are estimated using the routine NSPE (page 664). The impulse response weights  $\{v_k\}$  and differenced noise series  $\{n_t\}$  may be computed using the routine IRNSE (page 685). See Box and Jenkins (1976, pages 511–513).

### Example

Consider the Gas Furnace Data (Box and Jenkins 1976, pages 532–533) where  $X$  is the input gas rate in cubic feet/minute and  $Y$  is the percent  $\text{CO}_2$  in the outlet gas. The data is retrieved by routine GDATA (page 1302). Routine IRNSE (page 685) computes the impulse response weights. Application of routine TFPE to these weights produces the following results:

```

INTEGER      MWTIR, MWTSN, NDELAY, NOBS, NPAR, NPLHS, NPMA, NPNAR,
&            NPNMA, NPRHS, NSNOIS
PARAMETER    (MWTIR=10, NDELAY=3, NOBS=100, NPAR=3, NPLHS=2,
&            NPMA=0, NPNAR=2, NPNMA=0, NPRHS=2, MWTSN=MWTIR,
&            NSNOIS=NOBS-MWTSN)
C
INTEGER      IPRINT, MAXIT, NCOL, NROW
REAL         AVAR, PAR(NPAR), PLHS(NPLHS), PMA(1), PNAR(NPNAR),
&            PNMA(1), PRHS(NPRHS+1), RDATA(296,2), RELERR,
&            SNOISE(NOBS-MWTSN), WTIR(MWTIR+1), X(NOBS),
&            XPW(NOBS-NPAR), Y(NOBS), YPW(NOBS-NPAR)
EXTERNAL     GDATA, IRNSE, TFPE, WROPT
C
EQUIVALENCE (X(1), RDATA(1,1)), (Y(1), RDATA(1,2))
C
C                                     Gas Furnace Data
CALL GDATA (7, 0, NROW, NCOL, RDATA, 296, 2)
C
C                                     Specify AR parameters for
C                                     prewhitening transformation
PAR(1) = 1.97
PAR(2) = -1.37
PAR(3) = 0.34
C
C                                     Compute estimates of impulse

```

```

C                                     response weights and noise series
      IPRINT = 0
      CALL IRNSE (NOBS, X, Y, IPRINT, NPAR, PAR, NPMA, PMA, MWTIR,
&               MWTSN, WTIR, SNOISE, XPW, YPW)
C                                     Convergence parameters
      RELERR = 0.0
      MAXIT = 0
C                                     Compute preliminary estimates of
C                                     transfer function parameters
      CALL WROPT (-6, 1, 1)
      IPRINT = 1
      CALL TFPE (IPRINT, NPLHS, NPRHS, NPNAR, NPNMA, NDELAY, MWTIR,
&               WTIR, NSNOIS, SNOISE, RELERR, MAXIT, PLHS, PRHS,
&               PNAR, PNMA, AVAR)
C
      END

```

### Output

```

PLHS from TFPE/T2PE
      1          2
0.120342      0.326149

      PRHS from TFPE/T2PE
      1          2          3
-0.623240      0.318698      0.362488

PNAR from TFPE/T2PE
      1          2
1.64679      -0.70916

PNMA is not written since NPNMA = 0

AVAR from TFPE/T2PE =      2.85408E-02

```

---

## MLSE/DMLSE (Single/Double precision)

Compute least-squares estimates of a linear regression model for a multichannel time series with a specified base channel.

### Usage

```
CALL MLSE (NOBSX, NCHANX, X, LDX, IMEAN, XMEAN, NDIFF,
          NDPREG, LAG, CONST, NPREG, PREG)
```

### Arguments

**NOBSX** — Number of observations in each channel of the time series *x*. (Input)  
NOBSX must be less than or equal LDX and greater than  $\max(\text{NDPREG}(i) + \text{LAG}(i))$   
for  $i = 1, 2, \dots, \text{NCHANX}$ .

**NCHANX** — Number of channels in the time series *x*. (Input)  
NCHANX must be greater than or equal to one.

**X** — NOBSX by NCHANX matrix containing the time series. (Input)  
 Each row of X corresponds to an observation of a multivariate time series, and each column of X corresponds to a univariate time series. The base time series or output channel is contained in the first column.

**LDX** — Leading dimension of X exactly as specified in the dimension statement of the calling program. (Input)

**IMEAN** — Option for computation of the means of the channels of X. (Input)

**IMEAN Action**

- 0 XMEAN is user specified.
- 1 XMEAN is set to the vector of arithmetic means of the channels of X.

**XMEAN** — Vector of length NCHANX containing the means of the channels of X. (Input, if IMEAN = 0; output, if IMEAN = 1)

**NDIFF** — Vector of length NCHANX containing the order of differencing for each channel of X. (Input)

The elements of NDIFF must be greater than or equal to zero.

**NDPREG** — Vector of length NCHANX containing the number of regression parameters in the differenced form of the model for each channel of X. (Input)

The elements of NDPREG must be greater than or equal to zero.

**LAG** — Vector of length NCHANX containing the amount of time that each channel of X is to lag the base series. (Input)

The elements of LAG must be greater than or equal to zero.

**CONST** — Estimate of the overall constant. (Output)

**NPREG** — Number of regression parameters in the undifferenced model. (Output)

$$NPREG = IADD + (NDPREG(1) + NDIFF(1)) + \dots + (NDPREG(NCHANX) + NDIFF(NCHANX))$$

where

$$IADD = NDIFF(1), \text{ if } NDPREG(1) = 0$$

$$IADD = \max(0, \min(LAG(1) - 1, NDIFF(1))), \text{ if } NDPREG(1) > 0.$$

**PREG** — Vector of length NPREG containing the regression parameters in the undifferenced model. (Output)

The parameter estimates are concatenated as follows.

Channel 1: REG(*i*), *i* = 1, 2, ..., IADD + NDPREG(1) + NDIFF(1)

Channel *j*: PREG(*i*), *i* = I(*j*) + 1, I(*j*) + 2, ..., I(*j*) + NDPREG(*j*) + NDIFF(*j*)

where

$$I(j) = IADD + NDPREG(1) + NDIFF(1) + \dots + NDPREG(j - 1) + NDIFF(j - 1)$$

for *j* = 2, 3, ..., NCHANX.



## Comments

1. Automatic workspace storage is

MLSE NOBSX \* NCHANX + 2 \* NSUM<sup>2</sup> + NSUM + max(IADD, NCHANX + NSUM) units, or

DMLSE 2 \* (NOBSX \* NCHANX + 2 \* NSUM<sup>2</sup> + max(IADD, NCHANX + NSUM)) + NSUM units.

Workspace may be explicitly provided, if desired, by use of M2SE/DM2SE. The reference is

```
CALL M2SE (NOBSX, NCHANX, X, LDX, IMEAN, XMEAN,  
          NDIFF, NDPREG, LAG, CONST, NPREG, PREG,  
          XWK, IWK)
```

The additional arguments are as follows:

**XWK** — Work vector of length NOBSX \* NCHANX + 2 \* NSUM<sup>2</sup> + max(IADD, NCHANX + NSUM), where NSUM = NDPREG(1) + ... + NDPREG(NCHANX).

**IWK** — Work vector of length NSUM.

2. Prior to parameter estimation, the channels of X are centered and/or differenced according to XMEAN and NDIFF, respectively.
3. The undifferenced predictive form of the model is

$$x(t, 1) = \text{CONST} + \text{PREG}(1) * x(t-1, 1) + \dots + \text{PREG}(\text{IADD}) * x(t - \text{IADD}, 1) + \text{PREG}(\text{IADD} + 1) * x(t - \text{LAG}(1), 1) + \dots + \text{PREG}(\text{IADD} + \text{NDPREG}(1) + \text{NDIFF}(1)) * x(t - \text{LAG}(1) + 1 - \text{NDPREG}(1) - \text{NDIFF}(1), 1) + \dots + \text{PREG}(I(j) + 1) * x(t - \text{LAG}(j), j) + \dots + \text{PREG}(I(j) + \text{NDPREG}(j) + \text{NDIFF}(j)) * x(t - \text{LAG}(j) + 1 - \text{NDPREG}(j) - \text{NDIFF}(j), j) + \dots$$

where

$$I(j) = \text{IADD} + \text{NDPREG}(1) + \text{NDIFF}(1) + \dots + \text{NDPREG}(j-1) + \text{NDIFF}(j-1)$$

for  $j = 2, 3, \dots, \text{NCHANX}$ .

## Algorithm

Routine MLSE performs least-squares estimation of a linear regression model for a multichannel time series with a specified base channel.

Define the multichannel time series X by

$$X = (X_1, X_2, \dots, X_m)$$

where

$$X_j = (X_{1j}, X_{2j}, \dots, X_{nj})^T \quad j = 1, 2, \dots, m$$

with  $n = \text{NOBSX}$  and  $m = \text{NCHANX}$ . The columns of  $X$  correspond to individual channels of a multichannel time series and may be examined from a univariate perspective. The rows of  $X$  correspond to observations of an  $m$ -variate time series and may be examined from a multivariate perspective. Note that an alternative characterization of the multivariate time series  $X$  considers the columns of  $X$  to be observations of an  $m$ -variate time series with the rows of  $X$  containing univariate time series. For example, see Priestley (1981, page 692) and Fuller (1976, page 14).

The model is formed by regressing the base series  $X_1$  on its previous values and on the remaining channels  $X_2, \dots, X_m$ . The differenced form of the model is given by

$$X_{t1} = \theta_0 + \phi_1(B)\nabla^{d_1} B^{l_1} X_{t1} + \phi_2(B)\nabla^{d_2} B^{l_2} X_{t2} \\ + \dots + \phi_m(B)\nabla^{d_m} B^{l_m} X_{tm}$$

where  $\theta_0 = \text{CONST}$  is the overall constant,  $d_k = \text{NDIFF}(k)$  is the order of differencing  $X_k$ ,  $l_k = \text{LAG}(k)$  is the amount  $X_k$  lags  $X_1$ ,

$$\phi_k(B) = \phi_{1k} + \phi_{2k}B + \dots + \phi_{p_k,k}B^{p_k-1}$$

and  $p_k = \text{NDPREG}(k)$  for  $k = 1, 2, \dots, m$ .

The undifferenced form of the model is given by

$$X_{t1} = \theta_0 + \phi_1(B)X_{t-l_1,1} + \phi_2(B)X_{t-l_2,2} + \dots + \phi_m(B)X_{t-l_m,m}$$

where the undifferenced parameters  $\varphi_k = \text{PREG}(k)$  are defined by

$$\varphi_k(B) = \phi_k(B)\nabla^{d_k} \\ = \varphi_1 + \varphi_2B + \dots + \varphi_{p_k+d_k}B^{p_k+d_k-1}$$

for  $k = 1, 2, \dots, m$ . Note that if  $l_1 \geq d_1 \geq 0$ , the base series terms  $X_{t-j,1}$  at lags  $j = 1, \dots, (l_1 - 1)$  are omitted from the right-hand side of the above model when  $d_1 \geq 1$ . In the actual computations, these terms are included.

### Example 1

Consider the Wölfer Sunspot Data (Box and Jenkins 1976, page 530) along with data on northern light activity and earthquake activity (Robinson 1967, page 204) to be a three-channel time series. Routine MLSE is applied to these data to examine the regressive relationship between the channels.

```

C
INTEGER      LDX, NCHANX, NOBSX
PARAMETER   (NCHANX=3, NOBSX=100, LDX=NOBSX)

INTEGER      I, IMEAN, LAG(NCHANX), NCOL, NDIFF(NCHANX),
&            NDPREG(NCHANX), NOUT, NPREG, NROW
REAL        CONST, PREG(20), RDATA(100,4), X(LDX,NCHANX),
&            XMEAN(NCHANX)

```

```

EXTERNAL  GDATA, MLSE, UMACH
C
EQUIVALENCE (X(1,1), RDATA(1,2)), (X(1,2), RDATA(1,3)),
&           (X(1,3), RDATA(1,4))
C
DATA NDIFF(1), NDIFF(2), NDIFF(3)/1, 1, 0/
DATA LAG(1), LAG(2), LAG(3)/1, 2, 1/
DATA NDPREG(1), NDPREG(2), NDPREG(3)/2, 1, 3/
C
CALL GDATA (8, 0, NROW, NCOL, RDATA, 100, 4)
C                               Option to estimate channel means
IMEAN = 1
C                               Compute regression parameters
CALL MLSE (NOBSX, NCHANX, X, LDX, IMEAN, XMEAN, NDIFF, NDPREG,
&         LAG, CONST, NPREG, PREG)
C
C                               Print results
CALL UMACH (2, NOUT)
WRITE (NOUT,99993)
99993 FORMAT (//, 1X, '      Results of MLSE/M2SE')
WRITE (NOUT,99994)
99994 FORMAT (1X, ' I  NDIFF(I)  LAG(I)  NDPREG(I)      XMEAN(I)')
DO 10 I=1, NCHANX
    WRITE (NOUT,99995) I, NDIFF(I), LAG(I), NDPREG(I), XMEAN(I)
99995   FORMAT (1X, 4(I3,6X), F12.4)
    10 CONTINUE
WRITE (NOUT,99996) CONST
99996   FORMAT (1X, 'Overall constant, CONST = ', F12.4)
WRITE (NOUT,99997) NPREG
99997   FORMAT (//, 1X, 'Total number of parameters, NPREG = ', I2)
WRITE (NOUT,99998)
99998   FORMAT (//, 1X, ' I          PREG(I)')
DO 20 I=1, NPREG
    WRITE (NOUT,99999) I, PREG(I)
99999   FORMAT (1X, I2, 5X, F12.4)
    20 CONTINUE
C
END

```

### Output

```

Results of MLSE/M2SE
I  NDIFF(I)  LAG(I)  NDPREG(I)      XMEAN(I)
1      1      1      2          46.9400
2      1      2      1          63.4300
3      0      1      3          97.9700
Overall constant, CONST =      -7.2698

Total number of parameters, NPREG = 8

I          PREG(I)
1          -0.1481
2          -1.3444
3           0.4925
4          -0.0302
5           0.0302
6          -0.0210
7           0.0187
8           0.0765

```

## Example 2

Consider the Gas Furnace Data (Box and Jenkins 1976, pages 532–533) where  $X_1$  is the percent  $\text{CO}_2$  in the outlet gas and  $X_2$  is the input gas rate in cubic feet/minute. Application of routine MLSE to these data produces the following results:

```
      INTEGER      LDX, NCHANX, NOBSX
      PARAMETER    (NCHANX=2, NOBSX=296, LDX=NOBSX)
C
      INTEGER      I, IMEAN, LAG(NCHANX), NCOL, NDIFF(NCHANX),
&                NDPREG(NCHANX), NOUT, NPREG, NROW
      REAL         CONST, PREG(20), RDATA(296,2), X(LDX,NCHANX),
&                XMEAN(NCHANX)
      EXTERNAL     GDATA, MLSE, SCOPY, UMACH
C
      DATA NDIFF(1), NDIFF(2)/0, 0/
      DATA LAG(1), LAG(2)/1, 3/
      DATA NDPREG(1), NDPREG(2)/2, 3/
C
      CALL GDATA (7, 0, NROW, NCOL, RDATA, 296, 2)
C
      CALL SCOPY (NOBSX, RDATA(1,2), 1, X(1,1), 1)
      CALL SCOPY (NOBSX, RDATA(1,1), 1, X(1,2), 1)
C
      IMEAN = 1
C
      CALL MLSE (NOBSX, NCHANX, X, LDX, IMEAN, XMEAN, NDIFF, NDPREG,
&             LAG, CONST, NPREG, PREG)
C
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99993)
99993 FORMAT (1X, 'Results of MLSE/M2SE on Gas Furnace Data')
      WRITE (NOUT,99994)
99994 FORMAT (1X, ' I      NDIFF(I)  LAG(I)  NDPREG(I)      XMEAN(I)')
      DO 10 I=1, NCHANX
          WRITE (NOUT,99995) I, NDIFF(I), LAG(I), NDPREG(I), XMEAN(I)
99995   FORMAT (1X, 4(I3,6X), F12.4)
      10 CONTINUE
      WRITE (NOUT,99996) CONST
99996   FORMAT (1X, 'Overall constant, CONST = ', F12.4)
      WRITE (NOUT,99997) NPREG
99997   FORMAT (1X, 'Total number of parameters, NPREG = ', I2)
      WRITE (NOUT,99998)
99998   FORMAT (1X, ' I      PREG(I)')
      DO 20 I=1, NPREG
          WRITE (NOUT,99999) I, PREG(I)
99999   FORMAT (1X, I2, 5X, F12.4)
      20 CONTINUE
C
      END
```

## Output

```
Results of MLSE/M2SE on Gas Furnace Data
I      NDIFF(I)  LAG(I)  NDPREG(I)      XMEAN(I)
```

1	0	1	2	53.5091
2	0	3	3	-0.0568

Overall constant, CONST = 2.6562  
Total number of parameters, NPREG = 5

I	PREG(I)
1	1.6063
2	-0.6561
3	-0.4837
4	-0.1653
5	0.5052

---

## MWFE/DMWFE (Single/Double precision)

Compute least-squares estimates of the multichannel Wiener filter coefficients for two mutually stationary multichannel time series.

### Usage

```
CALL MWFE (NCHX, MLFIL, CXX, LDCXX, MDCXX, NCHZ, CZX,
           LDCZX, MDCZX, EPS, TRACE, LFIL, FIL, LDFIL,
           MDFIL ENMS)
```

### Arguments

**NCHX** — Number of input channels. (Input)  
NCHX must be greater than or equal to one.

**MLFIL** — Maximum length of the Wiener filter. (Input)  
MLFIL must be greater than or equal to one.

**CXX** — Array of size NCHX by NCHX by MLFIL containing the autocovariances of the input time series X. (Input)

**LDCXX** — Leading dimension of CXX exactly as specified in the dimension statement of the calling program. (Input)  
LDCXX must be greater than or equal to NCHX.

**MDCXX** — Middle dimension of CXX exactly as specified in the dimension statement of the calling program. (Input)  
MDCXX must be greater than or equal to NCHX.

**NCHZ** — Number of channels in desired output time series. (Input)  
NCHZ must be greater than or equal to one.

**CZX** — Array of size NCHZ by NCHX by MLFIL containing the cross-covariances between the desired output time series Z and the input time series X. (Input)

**LDCZX** — Leading dimension of CZX exactly as specified in the dimension statement of the calling program. (Input)  
LDCZX must be greater than or equal to NCHZ.

**MDCZX** — Middle dimension of CZX exactly as specified in the dimension statement of the calling program. (Input)  
MDCZX must be greater than or equal to NCHX.

**EPS** — Lower bound for the normalized mean square error. (Input)

**TRACE** — Trace of the autocovariance matrix of the desired output time series  $z$  at time lag zero. (Input)

**LFIL** — Length of the Wiener filter. (Output)

**FIL** — Array of size  $NCHZ$  by  $NCHX$  by  $MLFIL$  containing the multichannel Wiener filter coefficients. (Output)

**LDFIL** — Leading dimension of **FIL** exactly as specified in the dimension statement of the calling program. (Input)

$LDFIL$  must be greater than or equal to  $NCHZ$ .

**MDFIL** — Middle dimension of **FIL** exactly as specified in the dimension statement of the calling program. (Input)

$MDFIL$  must be greater than or equal to  $NCHX$ .

**ENMS** — Vector of length  $MLFIL$  containing the normalized mean square error corresponding to each filter length. (Output)

### Comments

1. Automatic workspace usage is

$MFFE$   $NCHX * NCHX * (2 * MLFIL + 12) + NCHZ + NCHX$  units,  
 $DMWFE$   $2 * (NCHX * NCHX * (2 * MLFIL + 12) + NCHZ) + NCHX$  units

Workspace may be explicitly provided, if desired, by use of  $M2FE/DM2FE$ . The reference is

```
CALL M2FE (NCHX, MLFIL, CXX, LDCXX, MDCXX, NCHZ,  
          CZX, LDCZX, MDCZX, EPS, TRACE, LFIL, FIL,  
          LDFIL, MDFIL, ENMS, IWK, WK)
```

The additional arguments are as follows:

**IWK** — Work vector of length  $NCHX$ .

**WK** — Work vector of length  $NCHX * NCHX * (2 * MLFIL + 12) + NCHZ$ .

2. The length of the filter is determined by the arguments **EPS** and **MLFIL**. Iteration to a longer filter stops when either the normalized mean square error **ENMS** is less than **EPS**, or the filter reaches the maximum allowable length, **MLFIL**.
3. The routine **MCCF** (page 649) may be used to obtain the input arguments **CXX**, **CZX**, and **TRACE**. For **TRACE**, routine **MCCF** may be used to obtain the autocovariances of the desired output series  $z$ . In particular,  $TRACE = ZVAR(1) + \dots + ZVAR(NCHZ)$ .
4. For a given lag  $k$ , the multichannel cross-covariance coefficient between  $z$  and  $x$  is defined as the array of size  $NCHZ$  by  $NCHX$  whose elements are the single-channel crosscovariance coefficients  $CZX(i, j, k)$ .

### Algorithm

Routine `MFFE` computes least-squares estimates of the multichannel Wiener filter coefficients for two mutually stationary multichannel time series.

Define the multichannel time series  $X$  by

$$X = (X_1, X_2, \dots, X_p)$$

where

$$X_j = (X_{1j}, X_{2j}, \dots, X_{nj})^T \quad j = 1, 2, \dots, p$$

with  $p = \text{NCHX}$ . Similarly, define the multichannel time series  $Z$  by

$$Z = (Z_1, Z_2, \dots, Z_q)$$

where

$$Z_j = (Z_{1j}, Z_{2j}, \dots, Z_{mj})^T \quad j = 1, 2, \dots, q$$

with  $q = \text{NCHZ}$ . The columns of  $X$  and  $Z$  correspond to individual channels of multichannel time series and may be examined from a univariate perspective. The rows of  $X$  and  $Z$  correspond to observations of  $p$ -variate and  $q$ -variate time series and may be examined from a multivariate perspective. Note that an alternative characterization of a multivariate time series  $X$  considers the columns of  $X$  to be observations of a  $p$ -variate time series with the rows of  $X$  containing univariate time series. For example, see Priestley (1981, page 692) and Fuller (1976, page 14).

Let

$$\hat{\mu}_X$$

be the row vector containing the means of the channels of  $X$ . In particular,

$$\hat{\mu}_X = (\hat{\mu}_{X_1}, \hat{\mu}_{X_2}, \dots, \hat{\mu}_{X_p})$$

where for  $j = 1, 2, \dots, p$

$$\hat{\mu}_{X_j} = \begin{cases} \mu_{X_j} & \mu_{X_j} \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_{tj} & \mu_{X_j} \text{ unknown} \end{cases}$$

Let

$$\hat{\mu}_Z$$

be similarly defined. In what follows, assume the channels of both  $X$  and  $Z$  have been centered about their respective means

$$\hat{\mu}_X \text{ and } \hat{\mu}_Z$$

Suppose the desired output is the multichannel time series  $Z$  defined by the model

$$Z_{t\bullet} = X_{t\bullet}\phi_0 + X_{(t-1)\bullet}\phi_1 + \dots + X_{(t-K)\bullet}\phi_K$$

where

$$X_{t\bullet} = (X_{t1}, X_{t2}, \dots, X_{tp})$$

$$Z_{t\bullet} = (Z_{t1}, Z_{t2}, \dots, Z_{tp})$$

and  $\phi_k$  is the array of dimension  $p \times q$  containing the Wiener filter coefficients

$$\phi_k = \begin{bmatrix} \phi_{11k} & \phi_{12k} & \cdots & \phi_{1qk} \\ \phi_{21k} & \phi_{22k} & \cdots & \phi_{2qk} \\ \vdots & \vdots & & \vdots \\ \phi_{p1k} & \phi_{p2k} & \cdots & \phi_{pqk} \end{bmatrix}$$

for  $k = 1, \dots, K$ . The array  $\phi_k$  is the  $(k + 1)$ -st level of the 3-dimensional array `FIL`.

The filter coefficients are computed by solving a set of normal equations. The algorithm utilizes the block Toeplitz (or Töplitz) matrix structure of these equations and is given by Robinson (1967, pages 238–246). In particular, the required input consists of the multichannel autocovariance matrices  $\Sigma_X$ ,  $\Sigma_Z$ , and the multichannel cross-covariance matrix  $\Sigma_{ZX}$ . The routine `MCCF` (page 649) may be used to estimate these covariance matrices.

Note that successively longer filters are estimated until either the normalized mean square error is less than `EPS` or the filter length  $K = \text{LFIL}$  equals `MLFIL`. The normalized mean square error is defined by

$$Q_k = 1 - \frac{\text{tr} \sum_{k=0}^K \Sigma_{ZX}(k) \phi_k}{\text{tr} \Sigma_Z(0)}$$

where  $\text{tr} \Sigma_Z(0) = \text{TRACE}$  is the trace of the multichannel autocorrelation coefficient of the desired output at lag zero. The values of  $Q_k$  for the successive filters of length  $k = 1, 2, \dots, K$  are contained in `ENMS`.

### Example

Consider the Wölfer Sunspot Data (Box and Jenkins 1976, page 530) along with data on northern light activity and earthquake activity (Robinson 1967, page 204) to be a three-channel time series. Routine `MWFE` applied to these data determines the following Wiener filter:

```

C                               SPECIFICATIONS FOR PARAMETERS
      INTEGER      IMEAN, IPRINT, LDCXX, LDCZX, LDFIL, LDX, LDZ, MAXLAG,
&                MDCXX, MDCZX, MDFIL, MLFIL, NCHANX, NCHANZ, NOBSX,
&                NOBSZ
      PARAMETER    ( IMEAN=1, IPRINT=0, MLFIL=3, NCHANX=3, NCHANZ=3,
&                NOBSX=99, NOBSZ=99, LDCXX=NCHANX, LDCZX=NCHANZ,
&                LDFIL=NCHANX, LDX=NOBSX, LDZ=NOBSZ, MAXLAG=MLFIL-1,
&                MDCXX=NCHANX, MDCZX=NCHANX, MDFIL=NCHANZ )
C
      INTEGER      I, J, K, LFIL, NCOL, NOUT, NROW

```



```

REAL      CVXX(LDCXX,MDCXX,-MAXLAG:MAXLAG), CVXX1(3,3,3),
&         CVZX(LDCZX,MDCZX,-MAXLAG:MAXLAG), CVZX1(3,3,3),
&         CXX(LDCXX,MDCXX,-MAXLAG:MAXLAG),
&         CZX(LDCZX,MDCZX,-MAXLAG:MAXLAG), ENMS(MLFIL), EPS,
&         FIL(LDFIL,MDFIL,MLFIL), R(0:2), RDATA(100,4), SSUM,
&         TRACE, X(LDX,NCHANX), XMEAN(NCHANX), XVAR(NCHANX),
&         YMEAN, YVAR, Z(LDZ,NCHANZ), ZMEAN(NCHANZ),
&         ZVAR(NCHANZ)
EXTERNAL  GDATA, MCCF, MWFE, SCOPY, SSUM, UMACH

C
EQUIVALENCE (CVXX(1,1,0), CVXX1(1,1,1)), (CVZX(1,1,0), CVZX1(1,1,
&         1))
C                                     Wolfer sunspot numbers
C                                     Northern lights activity
C                                     Earthquake activity
CALL GDATA (8, 0, NROW, NCOL, RDATA, 100, 4)
C
CALL SCOPY (NOBSX, RDATA(1,2), 1, X(1,1), 1)
CALL SCOPY (NOBSX, RDATA(1,3), 1, X(1,2), 1)
CALL SCOPY (NOBSX, RDATA(1,4), 1, X(1,3), 1)
C
CALL SCOPY (NOBSZ, RDATA(2,2), 1, Z(1,1), 1)
CALL SCOPY (NOBSZ, RDATA(2,3), 1, Z(1,2), 1)
CALL SCOPY (NOBSZ, RDATA(2,4), 1, Z(1,3), 1)
C                                     Compute multichannel ACF of Z
CALL MCCF (NOBSZ, NCHANZ, Z, LDZ, NOBSZ, NCHANZ, Z, LDZ, MAXLAG,
&         IPRINT, IMEAN, XMEAN, YMEAN, XVAR, YVAR, CVXX, LDCXX,
&         MDCXX, CXX, LDCXX, MDCXX)
C                                     Compute TRACE
TRACE = SSUM(NCHANZ,XVAR,1)
C                                     Compute multichannel ACF of X
CALL MCCF (NOBSX, NCHANX, X, LDX, NOBSX, NCHANX, X, LDX, MAXLAG,
&         IPRINT, IMEAN, XMEAN, ZMEAN, XVAR, ZVAR, CVXX, LDCXX,
&         LDCXX, CXX, LDCXX, LDCXX)
C                                     Compute multichannel CCF of Z and X
CALL MCCF (NOBSZ, NCHANZ, Z, LDZ, NOBSX, NCHANX, X, LDX, MAXLAG,
&         IPRINT, IMEAN, XMEAN, ZMEAN, XVAR, ZVAR, CVZX, LDCZX,
&         MDCZX, CZX, LDCZX, MDCZX)
C                                     Bound normalized MSE to be positive
EPS = 0.0
C                                     Reverse the LAG direction and scale
C                                     to agree with Robinson (1967)
R(0) = 99.D0
R(1) = 98.D0
R(2) = 97.D0
TRACE = TRACE*R(0)
DO 10 K=0, MAXLAG
  DO 10 J=1, NCHANX
    DO 10 I=1, NCHANX
      CVXX(I,J,K) = CVXX(I,J,-K)*R(K)
      CVZX(I,J,K) = CVZX(I,J,-K)*R(K)
10 CONTINUE
C                                     Compute multichannel Wiener filter
CALL MWFE (NCHANX, MLFIL, CVXX1, LDCXX, MDCXX, NCHANZ, CVZX1,
&         LDCZX, MDCZX, EPS, TRACE, LFIL, FIL, LDFIL, MDFIL,
&         ENMS)
C                                     Print results
CALL UMACH (2, NOUT)
WRITE (NOUT,99994) LFIL

```

```

99994 FORMAT (1X, 'Number of filter coefficients, LFIL = ', I3)
      DO 30 K=1, LFIL
          WRITE (NOUT,99995) K
99995  FORMAT (//, 1X, 'Wiener filter coefficient of index K = ', I3)
          DO 20 I=1, NCHANX
              WRITE (NOUT,99996) (FIL(I,J,K),J=1,NCHANZ)
99996  FORMAT (1X, 3F12.4)
          20 CONTINUE
          30 CONTINUE
          WRITE (NOUT,99997)
99997  FORMAT (//, 1X, 'Normalized mean square error')
          WRITE (NOUT,99998)
99998  FORMAT (1X, ' K           ENMS(K)')
          DO 40 K=1, LFIL
              WRITE (NOUT,99999) K, ENMS(K)
99999  FORMAT (1X, I2, 5X, F12.4)
          40 CONTINUE
C
      END

```

### Output

```

Number of filter coefficients, LFIL = 3

Wiener filter coefficient of index K = 1
 1.3834      0.0348      0.0158
 0.0599      0.8266      0.0629
-0.1710     -0.0332     -0.1205

Wiener filter coefficient of index K = 2
-0.7719     -0.0183     -0.0318
-0.0040     -0.2328      0.0484
-0.2170      0.1912     -0.0667

Wiener filter coefficient of index K = 3
 0.0516      0.0563     -0.0138
-0.0568      0.1084     -0.1731
 0.0007      0.2177     -0.0152

Normalized mean square error
 K           ENMS(K)
 1           0.6042
 2           0.5389
 3           0.5174

```

---

## KALMN/DKALMN (Single/Double precision)

Perform Kalman filtering and evaluate the likelihood function for the state-space model.

### Usage

```

CALL KALMN (NY, Y, NB, Z, LDZ, R, LDR, IT, T, LDT, IQ, Q,
           LDQ, TOL, B, COVB, LDCOV, N, SS, ALNDET, V,
           COVV, LDCOVV)

```

## Arguments

***NY*** — Number of observations for current update. (Input)

If  $NY = 0$ , no update is performed.

***Y*** — Vector of length  $NY$  containing the observations. (Input)

***NB*** — Number of elements in the state vector. (Input)

***Z*** —  $NY$  by  $NB$  matrix relating the observations to the state vector in the observation equation. (Input)

***LDZ*** — Leading dimension of  $Z$  exactly as specified in the dimension statement in the calling program. (Input)

***R*** —  $NY$  by  $NY$  matrix such that  $R * \sigma^2$  is the variance-covariance matrix of errors in the observation equation. (Input)

$\sigma^2$  is a positive unknown scalar. Only elements in the upper triangle of  $R$  are referenced.

***LDR*** — Leading dimension of  $R$  exactly as specified in the dimension statement in the calling program. (Input)

***IT*** — Transition matrix option. (Input)

**IT      Action**

0       $T$  is the transition matrix in the state equation.

1      The identity is the transition matrix in the state equation.

***T*** —  $NB$  by  $NB$  transition matrix in the state equation. (Input, if  $IT = 0$ )

If  $IT = 1$ , then  $T$  is not referenced and can be a vector of length one.

***LDT*** — Leading dimension of  $T$  exactly as specified in the dimension statement in the calling program. (Input)

***IQ*** — State equation error option. (Input)

**IQ      Action**

0      There is an error term in the state equation.

1      There is no error term in the state equation.

***Q*** —  $NB$  by  $NB$  matrix such that  $Q * \sigma^2$  is the variance-covariance matrix of the error vector in the state equation. (Input, if  $IQ = 0$ )

$\sigma^2$  is a positive unknown scalar. If  $IQ = 1$ , then  $Q$  is not referenced and can be a vector of length one. If  $IQ = 0$ , only the elements in the upper triangle of  $Q$  are referenced.

***LDQ*** — Leading dimension of  $Q$  exactly as specified in the dimension statement in the calling program. (Input)

***TOL*** — Tolerance used in determining linear dependence. (Input)

For **KALMN**,  $TOL = 100.0 * AMACH(4)$  is a common choice. For **DKALMN**,

$TOL = 100.0 * DMACH(4)$  is a common choice. See the documentation for routine **AMACH/DMACH** (Reference Material).

**B** — Estimated state vector of length NB. (Input/Output)

The input is the estimated state vector at time  $k$  given the observations thru time  $k - 1$ . The output is the estimated state vector at time  $k + 1$  given the observations thru time  $k$ . On the first call to KALMN, the input B must be the prior mean of the state vector at time 1.

**COVB** — NB by NB matrix such that  $COVB * \sigma^2$  is the mean squared error matrix for B. (Input/Output)

Before the first call to KALMN,  $COVB * \sigma^2$  must equal the variance-covariance matrix of the state vector.

**LDCOV** — Leading dimension of COVB exactly as specified in the dimension statement in the calling program. (Input)

**N** — Rank of the variance-covariance matrix for all the observations. (Input/Output)

N must be initialized to zero before the first call to KALMN. In the usual case when the variance-covariance matrix is nonsingular, N equals the sum of the NY's from the invocations to KALMN.

**SS** — Generalized sum of squares. (Input/Output)

SS must be initialized to zero before the first call to KALMN. The estimate of  $\sigma^2$  is given by  $SS/N$ .

**ALNDET** — Natural log of the product of the nonzero eigenvalues of  $P$  where  $P * \sigma^2$  is the variance-covariance matrix of the observations. (Input/Output)

Although ALNDET is computed, KALMN avoids the explicit computation of  $P$ . ALNDET must be initialized to zero before the first call to KALMN. In the usual case when  $P$  is nonsingular, ALNDET is the natural log of the determinant of  $P$ .

**V** — Vector of length NY containing the one-step-ahead prediction error. (Output)

If Y is not needed, then V and Y can occupy the same storage locations.

**COVV** — NY by NY matrix such that  $COVV * \sigma^2$  is the variance-covariance matrix of v. (Output)

If R is not needed, then COVV and R can occupy the same storage locations.

**LDCOV** — Leading dimension of COVV exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Automatic workspace usage is

KALMN  $NY^2 + NB^2 + NB * NY + \max(NB, NY)$  units, or

DKALMN  $2 * (NY^2 + NB^2 + NB * NY + \max(NB, NY))$  units.

Workspace may be explicitly provided, if desired, by use of K2LMN/DK2LMN. The reference is

```
CALL K2LMN (NY, Y, NB, Z, LDZ, R, LDR, IT, T, LDT,
           IQ, Q, LDQ, TOL, B, COVB, LDCOV, N, SS,
           ALNDET, V, COVV, LDCOVV, COVVCH, WK1,
           WK2)
```

The additional arguments are as follows.

**COVVCH** — Work vector of length  $NY * NY$  containing the Cholesky factor of the **COVV** matrix. If **R** and **COVV** are not needed, **COVVCH**, **R**, and **COVV** can occupy the same storage locations and **LDR** must equal **LDCOVV**.

**WK1** — Work vector of length  $NB * NB$ .

**WK2** — Work vector of length  $NB * NY + \max(NB, NY)$ .

## 2. Informational errors

Type	Code	
4	1	$R + Z * COVB * Z^T$ is not nonnegative definite within the tolerance defined by <b>TOL</b> . Either <b>TOL</b> is too small, or <b>R</b> or <b>COVB</b> is not nonnegative definite.
4	2	The system of equations $COVVCH^T * x = V$ is inconsistent. The variance-covariance matrix of the observations is inconsistent with the observations input in <b>Y</b> .
4	3	The system of equations $COVVCH^T * x = Z * COVB$ is inconsistent. The Cholesky factorization to compute <b>COVVCH</b> may be based on too large a value for <b>TOL</b> . The input of a smaller value for <b>TOL</b> may be appropriate.

3. If **R**, **Q**, and **T** are known functions of unknown parameters, **KALMN** can be used in conjunction with routine **UMINF** (IMSL MATH/LIBRARY) to perform maximum likelihood estimation of these unknown parameters. **UMINF** should be used to minimize the function

$$N * \text{ALOG}(SS/N) + \text{ALNDET};$$

4. In order to maintain acceptable numerical accuracy, the double precision routine, **DKALMN**, is usually required.

## Algorithm

Routine **KALMN** is based on a recursive algorithm given by Kalman (1960), which has come to be known as the Kalman filter. The underlying model is known as the state-space model. The model is specified stage by stage where the stages generally correspond to time points at which the observations become available. The routine **KALMN** avoids many of the computations and storage requirements that would be necessary if one were to process all the data at the end of each stage in order to estimate the state vector. This is accomplished by using previous computations and retaining in storage only those items essential for processing of future observations.

The notation used here follows that of Sallas and Harville (1981). Let  $y_k$  (input in  $\mathcal{Y}$ ) be the  $n_k \times 1$  vector of observations that become available at time  $k$ . The subscript  $k$  is used here rather than  $t$ , which is more customary in time series, to emphasize that the model is expressed in stages  $k = 1, 2, \dots$  and that these stages need not correspond to equally spaced time points. In fact, they need not correspond to time points of any kind. The *observation equation* for the state-space model is

$$y_k = Z_k b_k + e_k \quad k = 1, 2, \dots$$

Here,  $Z_k$  (input in  $\mathcal{Z}$ ) is an  $n_k \times q$  known matrix and  $b_k$  is the  $q \times 1$  state vector. The state vector  $b_k$  is allowed to change with time in accordance with the *state equation*

$$b_{k+1} = T_{k+1} b_k + w_{k+1} \quad k = 1, 2, \dots$$

starting with  $b_1 = \mu_1 + w_1$ .

The change in the state vector from time  $k$  to  $k + 1$  is explained in part by the *transition matrix*  $T_{k+1}$  (input in  $\mathcal{T}$ ), which is assumed known. It is assumed that the  $q$ -dimensional  $w_k$ 's ( $k = 1, 2, \dots$ ) are independently distributed multivariate normal with mean vector 0 and variance-covariance matrix  $\sigma^2 Q_k$ , that the  $n_k$ -dimensional  $e_k$ 's ( $k = 1, 2, \dots$ ) are independently distributed multivariate normal with mean vector 0 and variance-covariance matrix  $\sigma^2 R_k$ , and that the  $w_k$ 's and  $e_k$ 's are independent of each other. Here,  $\mu_1$  is the mean of  $b_1$  and is assumed known,  $\sigma^2$  is an unknown positive scalar.  $Q_{k+1}$  (input in  $\mathcal{Q}$ ) and  $R_k$  (input in  $\mathcal{R}$ ) are assumed known.

Denote the estimator of the realization of the state vector  $b_k$  given the observations  $y_1, y_2, \dots, y_j$  by

$$\hat{\beta}_{k|j}$$

By definition, the mean squared error matrix for

$$\hat{\beta}_{k|j}$$

is

$$\sigma^2 C_{k|j} = E(\hat{\beta}_{k|j} - b_k)(\hat{\beta}_{k|j} - b_k)^T$$

At the time of the  $k$ -th invocation, we have

$$\hat{\beta}_{k|k-1}$$

and  $C_{k|k-1}$ , which were computed from the  $(k-1)$ -st invocation, input in  $\mathcal{B}$  and  $\mathcal{COVB}$ , respectively. During the  $k$ -th invocation, routine  $\mathcal{KALMN}$  computes the filtered estimate

$$\hat{\beta}_{k|k}$$

along with  $C_{k|k}$ . These quantities are given by the *update equations*:

$$\begin{aligned}\hat{\beta}_{k|k} &= \hat{\beta}_{k|k-1} + C_{k|k-1} Z_k^T H_k^{-1} v_k \\ C_{k|k} &= C_{k|k-1} - C_{k|k-1} Z_k^T H_k^{-1} Z_k C_{k|k-1}\end{aligned}$$

where

$$v_k = y_k - Z_k \hat{\beta}_{k|k-1}$$

and where

$$H_k = R_k + Z_k C_{k|k-1} Z_k^T$$

Here,  $v_k$  (stored in  $v$ ) is the one-step-ahead prediction error, and  $\sigma^2 H_k$  is the variance-covariance matrix for  $v_k$ .  $H_k$  is stored in  $COVV$ . The “start-up values” needed on the first invocation of  $KALMN$  are

$$\hat{\beta}_{1|0} = \mu_1$$

and  $C_{1|0} = Q_1$  input via  $B$  and  $COVB$ , respectively. Computations for the  $k$ -th invocation are completed by  $KALMN$  computing the one-step-ahead estimate

$$\hat{\beta}_{k+1|k}$$

along with  $C_{k+1|k}$  given by the *prediction equations*:

$$\begin{aligned}\hat{\beta}_{k+1|k} &= T_{k+1} \hat{\beta}_{k|k} \\ C_{k+1|k} &= T_{k+1} C_{k|k} T_{k+1}^T + Q_{k+1}\end{aligned}$$

If both the filtered estimates and one-step-ahead estimates are needed by the user at each time point,  $KALMN$  can be invoked twice for each time point—first with  $IT = 1$  and  $IQ = 1$  to produce

$$\hat{\beta}_{k|k}$$

and  $C_{k|k}$ , and second with  $NY = 0$  to produce

$$\hat{\beta}_{k+1|k}$$

and  $C_{k+1|k}$  (With  $IT = 1$  and  $IQ = 1$ , the prediction equations are skipped. With  $NY = 0$ , the update equations are skipped.)

Often, one desires the estimate of the state vector more than one-step-ahead, i.e., an estimate of

$$\hat{\beta}_{k|j}$$

is needed where  $k > j + 1$ . At time  $j$ , KALMN is invoked to compute

$$\hat{\beta}_{j+1|j}$$

Subsequent invocations of KALMN with NY = 0 can compute

$$\hat{\beta}_{j+2|j}, \hat{\beta}_{j+3|j}, \dots, \hat{\beta}_{k|j}$$

Computations for

$$\hat{\beta}_{k|j}$$

and  $C_{kj}$  assume the variance-covariance matrices of the errors in the observation equation and state equation are known up to an unknown positive scalar multiplier,  $\sigma^2$ . The maximum likelihood estimate of  $\sigma^2$  based on the observations  $y_1, y_2, \dots, y_m$ , is given by

$$\hat{\sigma}^2 = SS / N$$

where

$$N = \sum_{k=1}^m n_k \text{ and } SS = \sum_{k=1}^m v_k^T H_k^{-1} v_k$$

If  $\sigma^2$  is known, the  $R_k$ 's and  $Q_k$ 's can be input as the variance-covariance matrices exactly. The earlier discussion is then simplified by letting  $\sigma^2 = 1$ .

In practice, the matrices  $T_k$ ,  $Q_k$ , and  $R_k$  are generally not completely known. They may be known functions of an unknown parameter vector  $\theta$ . In this case, KALMN can be used in conjunction with an optimization program (see routine UMINF, (IMSL MATH/LIBRARY) to obtain a maximum likelihood estimate of  $\theta$ . The natural logarithm of the likelihood function for  $y_1, y_2, \dots, y_m$  differs by no more than an additive constant from

$$L(\theta, \sigma^2; y_1, y_2, \dots, y_m) = -\frac{1}{2} N \ln \sigma^2 - \frac{1}{2} \sum_{k=1}^m \ln[\det(H_k)] - \frac{1}{2} \sigma^{-2} \sum_{k=1}^m v_k^T H_k^{-1} v_k$$

(Harvey 1981, page 14, equation 2.21). Here,

$$\sum_{k=1}^m \ln[\det(H_k)]$$

(stored in ALNDET) is the natural logarithm of the determinant of  $V$  where  $\sigma^2 V$  is the variance-covariance matrix of the observations.

Minimization of  $-2L(\theta, \sigma^2; y_1, y_2, \dots, y_m)$  over all  $\theta$  and  $\sigma^2$  produces maximum likelihood estimates. Equivalently, minimization of  $-2L_c(\theta; y_1, y_2, \dots, y_m)$  where



$$L_c(\theta; y_1, y_2, \dots, y_m) = -\frac{1}{2} N \ln\left(\frac{SS}{N}\right) - \frac{1}{2} \sum_{k=1}^m \ln[\det(H_k)]$$

produces maximum likelihood estimates

$$\hat{\theta} \text{ and } \hat{\sigma}^2 = SS / N$$

The minimization of  $-2L_c(\theta; y_1, y_2, \dots, y_m)$  instead of  $-2L(\theta, \sigma^2; y_1, y_2, \dots, y_m)$ , reduces the dimension of the minimization problem by one. The two optimization problems are equivalent since

$$\hat{\sigma}^2(\theta) = SS(\theta) / N$$

minimizes  $-2L(\theta, \sigma^2; y_1, y_2, \dots, y_m)$  for all  $\theta$ , consequently,

$$\hat{\sigma}^2(\theta)$$

can be substituted for  $\sigma^2$  in  $L(\theta, \sigma^2; y_1, y_2, \dots, y_m)$  to give a function that differs by no more than an additive constant from  $L_c(\theta; y_1, y_2, \dots, y_m)$ .

The earlier discussion assumed  $H_k$  to be nonsingular. If  $H_k$  is singular, a modification for singular distributions described by Rao (1973, pages 527–528) is used. The necessary changes in the preceding discussion are as follows:

1. Replace

$$H_k^{-1}$$

by a generalized inverse.

2. Replace  $\det(H_k)$  by the product of the nonzero eigenvalues of  $H_k$ .
3. Replace  $N$  by

$$\sum_{k=1}^m \text{rank}(H_k)$$

Maximum likelihood estimation of parameters in the Kalman filter is discussed by Sallas and Harville (1988) and Harvey (1981, pages 111–113).

### Example 1

Routine `KALMN` is used to compute the filtered estimates and one-step-ahead estimates for a scalar problem discussed by Harvey (1981, pages 116–117). The observation equation and state equation are given by

$$\begin{aligned} y_k &= b_k + e_k \\ b_{k+1} &= b_k + w_{k+1} \quad k = 1, 2, 3, 4 \end{aligned}$$

where the  $e_k$ 's are identically and independently distributed normal with mean 0 and variance  $\sigma^2$ , the  $w_k$ 's are identically and independently distributed normal with mean 0 and variance  $4\sigma^2$ , and  $b_1$  is distributed normal with mean 4 and

variance  $16\sigma^2$ . Two invocations of KALMN are needed for each time point in order to compute the filtered estimate and the one-step-ahead estimate. The first invocation uses  $IQ = 1$  and  $IT = 1$  so that the prediction equations are skipped in the computations. The second invocation uses  $NY = 0$  so that the update equations are skipped in the computations.

This example also computes the one-step-ahead prediction errors. Harvey (1981, page 117) contains a misprint for the value  $v_4$  that he gives as 1.197. The correct value of  $v_4 = 1:003$  is computed by KALMN.

```

C      INTEGER      LD COVB, LD COVV, LDQ, LDR, LDT, LDZ, NB, NOBS, NY
PARAMETER      (NB=1, NOBS=4, NY=1, LD COVB=NB, LD COVV=NY, LDQ=NB,
&              LDR=NY, LDT=NB, LDZ=NY)
C
C      INTEGER      I, IQ, IT, N, NOUT
REAL           ALNDET, AMACH, B(NB), COVB(LD COVB,NB),
&              COVV(LD COVV,NY), Q(LDQ,NB), R(LDR,NY), SS, T(LDT,NB),
&              TOL, V(NY), Y(NY), YDATA(NOBS), Z(LDZ,NB)
EXTERNAL      AMACH, KALMN, UMACH
C
C      DATA YDATA/4.4, 4.0, 3.5, 4.6/, Z/1.0/, R/1.0/, Q/4.0/, T/1.0/
C
C      TOL = 100.*AMACH(4)
CALL UMACH (2, NOUT)
C
C              Initial estimates for state vector
C              and variance-covariance matrix.
C              Initialize SS and ALNDET.
C
B(1)           = 4.0
COVB(1,1)     = 16.0
N              = 0
SS            = 0.0
ALNDET        = 0.0
WRITE (NOUT,99998)
C
C      DO 10 I=1, NOBS
C              Update
C              Y(1) = YDATA(I)
C              IQ = 1
C              IT = 1
CALL KALMN (NY, Y, NB, Z, LDZ, R, LDR, IT, T, LDT, IQ, Q,
&          LDQ, TOL, B, COVB, LD COVB, N, SS, ALNDET, V,
&          COVV, LD COVV)
WRITE (NOUT,99999) I, I, B(1), COVB(1,1), N, SS, ALNDET,
&          V(1), COVV(1,1)
C              Prediction
C              IQ = 0
C              IT = 0
CALL KALMN (0, Y, NB, Z, LDZ, R, LDR, IT, T, LDT, IQ, Q, LDQ,
&          TOL, B, COVB, LD COVB, N, SS, ALNDET, V, COVV,
&          LD COVV)
WRITE (NOUT,99999) I + 1, I, B(1), COVB(1,1), N, SS, ALNDET,
&          V(1), COVV(1,1)
10 CONTINUE
99998 FORMAT (' k/j', ' B', ' COVB', ' N', ' SS',
&          ' ALNDET', ' V', ' COVV')
99999 FORMAT (I2, '/', I1, 2F8.3, I2, 4F8.3)
END

```

### Output

k/j	B	COVB	N	SS	ALNDET	V	COVV
1/1	4.376	0.941	1	0.009	2.833	0.400	17.000
2/1	4.376	4.941	1	0.009	2.833	0.400	17.000
2/2	4.063	0.832	2	0.033	4.615	-0.376	5.941
3/2	4.063	4.832	2	0.033	4.615	-0.376	5.941
3/3	3.597	0.829	3	0.088	6.378	-0.563	5.832
4/3	3.597	4.829	3	0.088	6.378	-0.563	5.832
4/4	4.428	0.828	4	0.260	8.141	1.003	5.829
5/4	4.428	4.828	4	0.260	8.141	1.003	5.829

### Example 2

Routine `KALMN` is used with routine `UMINF` (IMSL MATH/LIBRARY) to find a maximum likelihood estimate of the parameter  $\theta$  in a MA(1) time series represented by  $y_k = \varepsilon_k - \theta\varepsilon_{k-1}$ . Routine `RNARM` (page 1232) is used to generate 200 random observations from an MA(1) time series with  $\theta = 0.5$  and  $\sigma^2 = 1$ . The MA(1) time series is cast as a state-space model of the following form (see Harvey 1981, pages 103–104, 112):

$$y_k = \begin{pmatrix} 1 & 0 \end{pmatrix} b_k$$

$$b_k = \begin{pmatrix} 0 & 1 \\ 0 & 0 \end{pmatrix} b_{k-1} + w_k$$

where the two-dimensional  $w_k$ 's are independently distributed bivariate normal with mean 0 and variance  $\sigma^2 Q_k$  and

$$Q_1 = \begin{pmatrix} 1 + \theta^2 & -\theta \\ -\theta & \theta^2 \end{pmatrix}$$

$$Q_k = \begin{pmatrix} 1 & -\theta \\ -\theta & \theta^2 \end{pmatrix} \quad k = 2, 3, \dots, 200$$

The warning error that is printed as part of the output is not serious and indicates that `UMINF` is generally used for multi-parameter minimization.

```

INTEGER      NOBS, NTHETA
PARAMETER    (NOBS=200, NTHETA=1)

C
  INTEGER     IADIST, IPARAM(7), ISEED, LAGAR(1), LAGMA(1), NOUT,
&            NPAR, NPMA
  REAL        A(NOBS+1), AVAR, CONST, FSCALE, FVALUE, PAR(1),
&            PMA(1), RPARAM(7), THETA(NTHETA), WI(1), XGUESS(1),
&            XSCALE(1), YDATA(NOBS), SNRM2
  COMMON      /MA1/ YDATA
  EXTERNAL    FCN, RNARM, RNSET, UMACH, UMINF, SNRM2

C
  ISEED = 123457
  CALL RNSET (ISEED)
  PMA(1)  = 0.5
  LAGMA(1) = 1
  CONST   = 0.0

```

```

NPAR      = 0
NPMA      = 1
IADIST    = 0
AVAR      = 1.0
CALL RNARM (NOBS, CONST, NPAR, PAR, LAGAR, NPMA, PMA, LAGMA,
&          IADIST, AVAR, A, WI, YDATA)
C          Use UMINF to find maximum likelihood
C          estimate of the MA parameter THETA.
XGUESS(1) = 0.0
XSCALE(1) = 1.0
FSCALE    = 1.0
IPARAM(1) = 0
CALL UMINF (FCN, NTHETA, XGUESS, XSCALE, FSCALE, IPARAM, RPARAM,
&          THETA, FVALUE)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) ' '
WRITE (NOUT,*) '* * * Final Estimate for THETA * * *'
WRITE (NOUT,*) 'Maximum likelihood estimate, THETA = ', THETA(1)
END
C          Use KALMN to evaluate the likelihood.
SUBROUTINE FCN (NTHETA, THETA, FUNC)
INTEGER    NTHETA
REAL      THETA(NTHETA), FUNC
C
INTEGER    LDCOVB, LDCOVV, LDQ, LDR, LDT, LDZ, NB, NOBS, NY
PARAMETER (NB=2, NOBS=200, NY=1, LDCOVB=NB, LDCOVV=NY, LDQ=NB,
&          LDR=NY, LDT=NB, LDZ=NY)
C
INTEGER    I, IQ, IT, N
REAL      ABS, ALNDET, ALOG, AMACH, B(NB), COVB(LDCOVB,NB),
&          COVV(LDCOVV,NY), Q(LDQ,NB), R(LDR,NY), SS, T(LDT,NB),
&          TOL, V(NY), Y(NY), YDATA(NOBS), Z(LDZ,NB)
COMMON    /MA1/ YDATA
INTRINSIC ABS, ALOG
EXTERNAL  AMACH, KALMN
C
DATA T/0.0, 0.0, 1.0, 0.0/, Z/1.0, 0.0/
C
IF (ABS(THETA(1)) .GT. 1.0) THEN
C          Estimate out of parameter space.
C          Set function to a large number.
      FUNC = 1.E10
      RETURN
END IF
TOL      = 100.*AMACH(4)
IQ       = 0
Q(1,1)  = 1.0
Q(1,2)  = -THETA(1)
Q(2,1)  = -THETA(1)
Q(2,2)  = THETA(1)**2
IT      = 0
C          No error in the
C          observation equation.
R(1,1)  = 0.0
C          Initial estimates for state vector
C          and variance-covariance matrix.
C          Initialize SS and ALNDET.
B(1)    = 0.0
B(2)    = 0.0

```

```

COVB(1,1) = 1.0 + THETA(1)**2
COVB(1,2) = -THETA(1)
COVB(2,1) = -THETA(1)
COVB(2,2) = THETA(1)**2
N          = 0
SS         = 0.0
ALNDET     = 0.0
C
DO 10 I=1, NOBS
  Y(1) = YDATA(I)
  CALL KALMN (NY, Y, NB, Z, LDZ, R, LDR, IT, T, LDT, IQ, Q,
&           LDQ, TOL, B, COVB, LDCOV, N, SS, ALNDET, V,
&           COV, LDCOVV)
10 CONTINUE
FUNC = N*ALOG(SS/N) + ALNDET
RETURN
END

```

### Output

```

*** WARNING  ERROR 1 from U5INF.  This routine may be inefficient for a
***          problem of size N = 1.
Here is a traceback of subprogram calls in reverse order:
Routine name          Error type  Error code
-----
U5INF                 6          1    (Called internally)
U3INF                 0          0    (Called internally)
U2INF                 0          0    (Called internally)
UMINF                 0          0
USER                  0          0
* * * Final Estimate for THETA * * *
Maximum likelihood estimate, THETA = 0.452842

```

---

## LOFCF/DLOFCF (Single/Double precision)

Perform lack-of-fit test for a univariate time series or transfer function given the appropriate correlation function.

### Usage

```
CALL LOFCF (NOBS, LAGMIN, LAGMAX, CF, NPFREE, Q, PVALUE)
```

### Arguments

**NOBS** — Number of observations in the stationary time series. (Input)  
NOBS must be greater than or equal to two.

**LAGMIN** — Minimum lag of the correlation function. (Input)  
LAGMIN corresponds to the lower bound of summation in the lack of fit test statistic. Generally, LAGMIN is set to one if CF is an autocorrelation function and is set to zero if CF is a cross correlation function.

**LAGMAX** — Maximum lag of the correlation function. (Input)  
LAGMAX corresponds to the upper bound of summation in the lack of fit test statistic. LAGMAX must be greater than or equal to LAGMIN and less than NOBS.

**CF** — Vector of length  $\text{LAGMAX} + 1$  containing the correlation function. (Input)  
 The correlation coefficient for lag  $k$  is given by  $\text{CF}(k + 1)$ ,  $k = \text{LAGMIN}, \text{LAGMIN} + 1, \dots, \text{LAGMAX}$ .

**NPFREE** — Number of free parameters in the formulation of the time series model. (Input)  
 NPFREE must be greater than or equal to zero and less than LAGMAX.

**Q** — Lack of fit test statistic. (Output)

**PVALUE** —  $p$ -value of the test statistic  $Q$ . (Output)  
 Under the null hypothesis,  $Q$  has an approximate chi-squared distribution with  $\text{LAGMAX} - \text{LAGMIN} + 1 - \text{NPFREE}$  degrees of freedom.

### Comments

Routine LOFCF may be used to diagnose lack of fit in both ARMA and transfer function models. Typical arguments for these situations are

Model	LAGMIN	LAGMAX	NPFREE
ARMA( $p, q$ )	1	$\sqrt{\text{NOBS}}$	$p + q$
Transfer function	0	$\sqrt{\text{NOBS}}$	$r + s$

See the “Algorithm” section for further information.

### Algorithm

Routine LOFCF performs a portmanteau lack of fit test for a time series or transfer function containing  $n$  observations given the appropriate sample correlation function

$$\hat{\rho}(k)$$

for  $k = L, L + 1, \dots, K$  where  $L = \text{LAGMIN}$  and  $K = \text{LAGMAX}$ .

The basic form of the test statistic  $Q$  is

$$Q = n(n + 2) \sum_{k=L}^K (n - k)^{-1} \hat{\rho}(k)$$

with  $L = 1$  if

$$\hat{\rho}(k)$$

is an autocorrelation function and  $L = 0$  if

$$\hat{\rho}(k)$$

is a cross-correlation function. Given that the model is adequate,  $Q$  has a chi-squared distribution with  $K - L + 1 - m$  degrees of freedom where  $m = \text{NPFREE}$  is the number of parameters estimated in the model. If the mean of the time

series is estimated, Woodfield (1990) recommends not including this in the count of the parameters estimated in the model. Thus, for an ARMA( $p, q$ ) model set  $NPFREE = p + q$  regardless of whether the mean is estimated or not. The original derivation for time series models is due to Box and Pierce (1970) with the above modified version discussed by Ljung and Box (1978). The extension of the test to transfer function models is discussed by Box and Jenkins (1976, pages 394–395).

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. An ARMA(2,1) with nonzero mean is fitted using routine NSLSE (page 669). The autocorrelations of the residuals are estimated using routine ACF (page 637). A portmanteau lack of fit test is computed using 10 lags with LOFCF.

The warning message from NSLSE in the output can be ignored. (See the example for routine NSLSE for a full explanation of the warning message.)

```

INTEGER      IARDEG, IMEAN, IPRINT, ISEOPT, LAGMAX, LAGMIN, LDCOV,
&            LDX, MAXBC, MDX, NOBS, NP, NPAR, NPFREE, NPMA
PARAMETER    (IARDEG=2, IMEAN=1, IPRINT=0, ISEOPT=0, LAGMAX=10,
&            LAGMIN=1, LDX=176, MAXBC=10, MDX=2, NOBS=100, NPAR=2,
&            NPFREE=4, NPMA=1, NP=NPAR+NPMA+IMEAN, LDCOV=NP)
C
INTEGER      LAGAR(NPAR), LAGMA(NPMA), MAXIT, NA, NCOL, NOUT, NROW
REAL         A(NOBS-IARDEG+MAXBC), ACV(LAGMAX+1), AVAR,
&            CF(LAGMAX+1), CONST, COV(LDCOV, NP), PAR(NPAR),
&            PMA(NPMA), PVALUE, Q, RELERR, SEAC(LAGMAX), TOLBC,
&            TOLSS, W(NOBS), WMEAN, X(LDX, MDX)
EXTERNAL     ACF, GDATA, LOFCF, NSLSE, NSPE, UMACH
C
EQUIVALENCE (W(1), X(22,2))
C
DATA LAGAR/1, 2/, LAGMA/1/
C
CALL UMACH (2, NOUT)
C
C                               Wolfer Sunspot Data for
C                               years 1770 through 1869
CALL GDATA (2, IPRINT, NROW, NCOL, X, LDX, MDX)
C
C                               Convergence parameters
MAXIT = 0
RELERR = 0.0
C
C                               Compute preliminary parameter
C                               estimates for ARMA(2,1) model
CALL NSPE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, NPMA, RELERR,
&         MAXIT, CONST, PAR, PMA, AVAR)
C
C                               Compute least squares estimates
C                               for model
TOLBC = 0.0
TOLSS = 0.125
C
CALL NSLSE (NOBS, W, IPRINT, IMEAN, WMEAN, NPAR, PAR, LAGAR,
&         NPMA, PMA, LAGMA, MAXBC, TOLBC, TOLSS, CONST, COV,

```

```

&          LDCOV, NA, A, AVAR)
C          Compute autocorrelations of the
C          residuals
CALL ACF (NOBS-IARDEG+MAXBC, A, IPRINT, ISEOPT, IMEAN, WMEAN,
&          LAGMAX, ACV, CF, SEAC)
C
CALL LOFCF (NOBS, LAGMIN, LAGMAX, CF, NPFREE, Q, PVALUE)
C
WRITE (NOUT,99998) Q
WRITE (NOUT,99999) LAGMAX - LAGMIN + 1 - NPFREE, PVALUE
C
99998 FORMAT (/4X, 'Lack of Fit statistic (Q) = ', F12.3)
99999 FORMAT (/4X, 'Degrees of freedom (LAGMAX-LAGMIN+1-NPFREE) = ',
&          I8/4X, 'P-value (PVALUE) = ', F12.4)
END

```

### Output

```

***WARNING  ERROR 1 from NSLSE.  Least squares estimation of the parameters
***          has failed to converge.  Increase MAXBC and/or TOLBC and/or
***          TOLSS.  The estimates of the parameters at the last iteration
***          may be used as new starting values.

```

Lack of Fit statistic (Q) = 14.572

Degrees of freedom (LAGMAX-LAGMIN+1-NPFREE) = 6

P-value (PVALUE) = 0.9761

---

## DIRIC/DDIRIC (Single/Double precision)

Compute the Dirichlet kernel.

### Usage

DIRIC(M, RANGLE, EPS)

### Arguments

*M* — Spectral window parameter. (Input)

*RANGLE* — Angle in radians. (Input)

*EPS* — Positive bound on  $|RANGLE|$  that determines when an approximation to the Dirichlet kernel is appropriate. (Input)

*EPS* must be between 0 and  $\pi$  inclusive. The approximation is used when  $|RANGLE|$  is less than *EPS*.

*DIRIC* — Function value. (Output)

### Comments

1. The Dirichlet kernel is equivalent to the truncated periodogram spectral window. The spectral window parameter denotes the truncation point in the weighted sum of sample autocovariances used to estimate the spectral density.



2. The Dirichlet kernel produces negative values for certain values of RANGLE. Thus, spectral windows that use the Dirichlet kernel may also take on negative values.
3. The Dirichlet kernel is defined between  $-\pi$  and  $\pi$ , inclusive, and is zero otherwise.

### Algorithm

Routine DIRIC evaluates the Dirichlet kernel,  $D_M(\theta)$ , for a given parameter  $M$ , angle  $\theta = \text{RANGLE}$ , and tolerance  $\varepsilon = \text{EPS}$ . The computational form of the function is given by

$$D_M(\theta) = \begin{cases} \frac{(2M+1)}{2\pi} \left( \frac{\sin\left[\left(M + \frac{1}{2}\right)\theta\right]}{\left(M + \frac{1}{2}\right)\theta} \right) & |\theta| < \varepsilon \\ \frac{1}{2\pi} \left( \frac{\sin\left[\left(M + \frac{1}{2}\right)\theta\right]}{\sin(\theta/2)} \right) & \varepsilon \leq |\theta| \leq \pi \\ 0 & |\theta| > \pi \end{cases}$$

The first case is an approximation to  $D_M(\theta)$  for small  $\theta$ , and the second case is the usual theoretical definition.

In spectral analysis, the Dirichlet kernel corresponds to the truncated periodogram spectral window, and  $M$  is called the spectral window parameter. Since the Dirichlet kernel may be negative for certain values of  $\theta$ , the truncated periodogram estimate of the spectral density may also be negative. This is an undesirable property since the true spectral density is a nonnegative function. See Priestley (1981, pages 437–438) and Anderson (1971, pages 508–511) for further discussion.

### Example

In this example, DIRIC is used to compute the Dirichlet kernel at  $\theta = \pm k\pi/(2M+1)$  for  $k = 0, 1, \dots, (2M+1)$  where  $M = 5$  and  $\varepsilon = 0.01$ .

```

C                                     SPECIFICATIONS FOR LOCAL VARIABLES
INTEGER      K, M, NOUT
REAL         DIRIC, EPS, PI, REAL, THETA, WT
INTRINSIC   REAL
EXTERNAL    DIRIC, UMACH

C
M           = 5
EPS        = .01
PI         = 3.14159

C
CALL UMACH (2, NOUT)

C
WRITE (NOUT,99998)

```

```

99998 FORMAT ( ' K      THETA      WEIGHT ' )
      DO 10 K=0, 2*M + 1
          THETA = (PI*REAL(K))/REAL(2*M+1)
          WT    = DIRIC(M,THETA,EPS)
          WRITE (NOUT,99999) K, THETA, WT
99999  FORMAT (1X, I2, 2(3X,F8.5))
      10 CONTINUE
C
      RETURN
      END

```

### Output

K	THETA	WEIGHT
0	0.00000	1.75070
1	0.28560	1.11833
2	0.57120	0.00000
3	0.85680	-0.38312
4	1.14240	0.00000
5	1.42800	0.24304
6	1.71359	0.00000
7	1.99919	-0.18919
8	2.28479	0.00000
9	2.57039	0.16587
10	2.85599	0.00000
11	3.14159	-0.15915

---

## FEJER/DFEJER (Single/Double precision)

Compute the Fejér kernel.

### Usage

FEJER(M, RANGLE, EPS)

### Arguments

*M* — Spectral window parameter. (Input)

*RANGLE* — Angle in radians. (Input)

*EPS* — Positive bound on  $|RANGLE|$  that determines when an approximation to the Fejér kernel is appropriate. (Input)

*EPS* must be between 0 and  $\pi$  inclusive. The approximation is used when  $|RANGLE|$  is less than *EPS*.

*FEJER* — Function value. (Output)

### Comments

1. The Fejér kernel is equivalent to the modified Bartlett spectral window. The spectral window parameter denotes the truncation point in the weighted sum of sample autocovariances used to estimate the spectral density.

2. The Fejér kernel produces nonnegative values for all values of RANGLE. Thus, spectral windows based on the Fejér kernel are always nonnegative.
3. The Fejér kernel is defined between  $-\pi$  and  $\pi$ , inclusive, and is zero otherwise.

### Algorithm

Routine FEJER evaluates the Fejér kernel,  $F_M(\theta)$ , for a given parameter  $M$ , angle  $\theta = \text{RANGLE}$ , and tolerance  $\varepsilon = \text{EPS}$ . The computational form of the function is given by

$$F_M(\theta) = \begin{cases} \frac{M}{2\pi} \left( \frac{\sin[M\theta/2]}{M\theta/2} \right)^2 & |\theta| < \varepsilon \\ \frac{1}{2\pi M} \left( \frac{\sin[M\theta/2]}{\sin(\theta/2)} \right)^2 & \varepsilon \leq |\theta| \leq \pi \\ 0 & |\theta| > \pi \end{cases}$$

The first case is an approximation to  $F_M(\theta)$  for small  $\theta$ , and the second case is the usual theoretical definition.

In spectral analysis, the Fejér kernel corresponds to the modified Bartlett spectral window, and  $M$  is called the spectral window parameter. Since the Fejér kernel is nonnegative for all values of  $\theta$ , the modified Bartlett estimate of the spectral density is also nonnegative. This is a desirable property since the true spectral density is a nonnegative function. See Priestley (1981, pages 439–440) and Anderson (1971, pages 508–511) for further discussion.

### Example

In this example, FEJER is used to compute the Fejér kernel at  $\theta = \pm k\pi/M$  for  $k = 0, 1, \dots, M$  where  $M = 11$  and  $\varepsilon = 0.01$ .

```

C
      INTEGER      K, M, NOUT
      REAL         EPS, FEJER, PI, REAL, THETA, WT
      INTRINSIC    REAL
      EXTERNAL     FEJER, UMACH
C
      M      = 11
      EPS    = .01
      PI     = 3.14159265
      CALL UMACH (2, NOUT)
C
      WRITE (NOUT,99998)
99998 FORMAT (' K      THETA      WEIGHT ')
      DO 10 K=0, M
          THETA = (PI*REAL(K))/REAL(M)
          WT    = FEJER(M, THETA, EPS)

```

```

          WRITE (NOUT,99999) K, THETA, WT
99999    FORMAT (1X, I2, 2(3X,F8.5))
      10 CONTINUE
C
      RETURN
      END

```

### Output

K	THETA	WEIGHT
0	0.00000	1.75070
1	0.28560	0.71438
2	0.57120	0.00000
3	0.85680	0.08384
4	1.14240	0.00000
5	1.42800	0.03374
6	1.71360	0.00000
7	1.99920	0.02044
8	2.28479	0.00000
9	2.57039	0.01572
10	2.85599	0.00000
11	3.14159	0.00000

---

## PFFT/DPFFT (Single/Double precision)

Compute the periodogram of a stationary time series using a fast Fourier transform.

### Usage

```
CALL PFFT (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, IPVER, PM,
          LDPM)
```

### Arguments

**NOBS** — Number of observations in the stationary time series  $x$ . (Input)  
 NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the stationary time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.

1 Print the periodogram, and the cosine and sine transformations of the centered and padded time series.

**XCNTR** — Constant used to center the time series  $x$ . (Input)

**NPAD** — Number of zeroes used to pad the centered time series. (Input)  
 NPAD must be greater than or equal to zero. The length of the centered and padded time series is  $N = \text{NOBS} + \text{NPAD}$ .

**IFSCAL** — Option for frequency scale. (Input)

**IFSCAL Action**

- 0 Frequency in radians per unit time
- 1 Frequency in cycles per unit time

**IPVER** — Option for version of the periodogram. (Input)

**IPVER Action**

- 0 Compute usual periodogram.
- 1 Compute modified periodogram.

Refer to the algorithm section for further details.

**PM** —  $(\lfloor N/2 \rfloor + 1)$  by 5 matrix that contains a summarization of the periodogram analysis. (Output)

For  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ , the  $(k + 1)$ -st element of the  $j$ -th column of **PM** is defined as

**Col. Description**

- 1 Frequency,  $\omega_k$  where  $\omega_k = 2\pi k/N$  for **IFSCAL** = 0 and  $\omega_k = k/N$  for **IFSCAL** = 1.
- 2 Period,  $p_k$  where  $p_k = 2\pi/\omega_k$  for **IFSCAL** = 0 and  $p_k = 1/\omega_k$  for **IFSCAL** = 1. If  $\omega_k = 0$ ,  $p_k$  is set to missing.
- 3 Periodogram ordinate,  $I(\omega_k)$ .
- 4 Cosine transformation coefficient,  $A(\omega_k)$ .
- 5 Sine transformation coefficient,  $B(\omega_k)$ .

**LDPM** — Leading dimension of **PM** exactly as specified in the dimension statement of the calling program. (Input)

**LDPM** must be greater than or equal to  $\lfloor N/2 \rfloor + 1$ .

**Comments**

1. Automatic workspace storage is

**PFFT**  $10N + 15$  units, or  
**DPFFT**  $20N + 30$  units.

Workspace may be explicitly provided, if desired, by use of **P2FT/DP2FT**. The reference is

CALL **P2FT** (**NOBS**, **X**, **IPRINT**, **XCNTR**, **NPAD**, **IFSCAL**,  
**IPVER**, **PM**, **LDPM**, **CX**, **COEF**, **WFFTC**, **CPY**)

The additional arguments are as follows:

**CX** — Complex work vector of length  $N$ .

**COEF** — Complex work vector of length  $N$ .

**WFFTC** — Work vector of length  $4N + 15$ .

**CPY** — Work vector of length  $2N$ .

2. The centered and padded time series is defined by
 
$$\begin{aligned} \text{CX}(j) &= X(j) - \text{XCNTR} && \text{for } j = 1, \dots, \text{NOBS} \\ \text{CX}(j) &= 0 && \text{for } j = \text{NOBS} + 1, \dots, N \end{aligned}$$
 where  $N = \text{NOBS} + \text{NPAD}$ .
3. The periodogram  $I(\omega)$  is an even function of the frequency  $\omega$ . The relation  $I(-\omega) = I(\omega)$  for  $\omega > 0.0$  recovers the periodogram for negative frequencies.
4. Since  $\cos(\omega)$  is an even function of  $\omega$  and  $\sin(\omega)$  is an odd function of  $\omega$ , the cosine and sine transformations, respectively, satisfy  $A(-\omega) = A(\omega)$  and  $B(-\omega) = -B(\omega)$  for  $\omega > 0.0$ . Similarly, the complex Fourier coefficients, stored in `COEF`, satisfy  $\text{COEF}(-\omega) = \text{conj}(\text{COEF}(\omega))$ .
5. Computation of the  $2 * \text{NOBS} - 1$  autocovariances of  $X$  using the inverse Fourier transform of the periodogram requires  $\text{NPAD} = \text{NOBS} - 1$ .

### Algorithm

Routine `PFFT` computes the periodogram of a stationary time series given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  for  $t = 1, 2, \dots, n$ .

Let

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  represent the centered and padded data where  $N = \text{NOBS} + \text{NPAD}$ ,

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu}_X & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_X = \text{XCNTR}$$

is determined by

$$\hat{\mu}_X = \begin{cases} \mu_X & \mu_X \text{ known} \\ \bar{X} = \frac{1}{n} \sum_{t=1}^n X_t & \mu_X \text{ unknown} \end{cases}$$

The discrete Fourier transform of

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  is defined by

$$\zeta_{\tilde{X}}(\omega_k) = \sum_{t=1}^N \tilde{X}_t e^{-i\omega_k t}$$

over the discrete set of frequencies

$$\omega_k = \frac{2\pi k}{N} \quad k = 0, \pm 1, \dots, \pm \lfloor N/2 \rfloor$$

An alternative representation of

$$\zeta_{\tilde{X}}(\omega_k)$$

in terms of cosine and sine transforms is

$$\zeta_{\tilde{X}}(\omega_k) = \alpha_{\tilde{X}}(\omega_k) - i\beta_{\tilde{X}}(\omega_k)$$

where

$$\alpha_{\tilde{X}}(\omega_k) = \sum_{t=1}^n \tilde{X}_t \cos(\omega_k t)$$

and

$$\beta_{\tilde{X}}(\omega_k) = \sum_{t=1}^n \tilde{X}_t \sin(\omega_k t)$$

The fast Fourier transform algorithm is used to compute the discrete Fourier transform. The periodogram of the sample sequence  $\{X_t\}$ ,  $t = 1, \dots, n$  computed with the centered and padded sequence

$$\{\tilde{X}_t\}$$

$t = 1, \dots, N$  is defined by

$$I_{n,N,\tilde{X}}(\omega_k) = K \left| \sum_{t=1}^N \tilde{X}_t e^{-i\omega_k t} \right|^2 = K |\zeta_{\tilde{X}}(\omega_k)|^2$$

where  $K$  is the scale factor

$$K = \begin{cases} \frac{2}{n} & \text{for the usual periodogram} \\ \frac{1}{2\pi n} & \text{for the modified periodogram} \end{cases}$$

The scale factor of the usual periodogram relates the ordinates to the sum of squares of

$$X_t - \hat{\mu}_X$$

(Fuller 1976, pages 276–277). If the first ordinate (corresponding to  $k = 0$ ) is replaced by one-half of its value, then if  $N$  is odd, the sum of the  $\lfloor N/2 \rfloor + 1$  ordinates corresponding to  $k = 0, 1, \dots, \lfloor N/2 \rfloor$  is

$$\frac{N}{n} \sum_{t=1}^n (X_t - \hat{\mu}_X)^2$$

For  $N$  even, if the first ordinate (corresponding to  $k = 0$ ) and the last ordinate (corresponding to  $k = N/2$ ) are each replaced by one-half of their values, then the same relationship holds. The modified periodogram is an asymptotically unbiased estimate of the nonnormalized spectral density function at each frequency  $\omega_k$  (Priestley 1981, page 417). The argument `IPVER` is used to specify the version of the periodogram.

The alternative representation of the discrete Fourier transform implies

$$I_{n,N,\tilde{X}}(\omega_k) = A_{\tilde{X}}^2(\omega_k) + B_{\tilde{X}}^2(\omega_k)$$

where

$$A_{\tilde{X}}(\omega_k) = K^{1/2} \alpha_{\tilde{X}}(\omega_k)$$

and

$$B_{\tilde{X}}(\omega_k) = K^{1/2} \beta_{\tilde{X}}(\omega_k)$$

represent the (scaled) cosine and sine transforms, respectively. Since the periodogram is an even function of the frequency, it is sufficient to estimate the periodogram at the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N} \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

Use of the centered data

$$\{\tilde{X}_t\}$$

(without padding) instead of the original data  $\{X_t\}$  for  $t = 1, \dots, n$  does not affect the asymptotic sampling properties of the periodogram. In fact,

$$I_{n,n,\tilde{X}}(\omega_k) \equiv I_{n,n,X}(\omega_k) \quad \omega_k \neq 0$$

For  $\omega_k = 0$ , both

$$I_{n,n,\tilde{X}}(0) = 0$$

and

$$I_{n,n,X}(0) = K \left( \sum_{t=1}^n X_t \right)^2 = Kn^2 \bar{X}^2$$

reflect the mean of the data. See Priestley (1981, page 417) for further discussion.

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set



for this example consists of the number of sunspots observed from 1770 through 1869. Application of routine PFFT to these data produces the following results.

```

INTEGER      IPRINT, LDPM, NOBS
PARAMETER   (IPRINT=0, LDPM=100, NOBS=100)
C
INTEGER      IFSCAL, IPVER, NCOL, NPAD, NROW
REAL         PM(LDPM,5), RDATA(176,2), REAL, SSUM, X(NOBS), XCNTR
CHARACTER    CLABEL(6)*9, FMT*20, RLABEL(1)*6
INTRINSIC    REAL
EXTERNAL     GDATA, PFFT, SSUM, WRRRL
C
EQUIVALENCE (X(1), RDATA(22,2))
C
DATA RLABEL/'NONE'/, CLABEL/' ', 'Frequency', 'Period',
&          'I(w(k))', 'A(w(k))', 'B(w(k))'/
C
C          Wolfer Sunspot Data for
C          years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C
C          Center on arithmetic mean
XCNTR = SSUM(NOBS,X,1)/REAL(NOBS)
C
C          Pad standard amount
NPAD = NOBS - 1
C
C          Frequency in radians per unit time
IFSCAL = 0
C
C          Modified periodogram version
IPVER = 1
C
C          Compute the periodogram
CALL PFFT (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, IPVER, PM, LDPM)
C
C          Print results
FMT = '(F9.4, F6.2, 3F10.2) '
CALL WRRRL (' ', 20, 5, PM, LDPM, 0, FMT, RLABEL, CLABEL)
C
END

```

### Output

Frequency	Period	I(w(k))	A(w(k))	B(w(k))
0.0000	NaN	0.00	0.00	0.00
0.0316	199.00	183.97	3.72	-13.04
0.0631	99.50	1363.37	35.45	-10.32
0.0947	66.33	2427.09	29.31	39.60
0.1263	49.75	1346.64	-21.74	29.56
0.1579	39.80	139.74	-11.69	-1.79
0.1894	33.17	44.67	-4.65	4.80
0.2210	28.43	123.47	-11.11	-0.33
0.2526	24.88	176.04	-4.79	-12.37
0.2842	22.11	143.06	9.92	-6.69
0.3157	19.90	44.17	6.43	1.68
0.3473	18.09	38.95	5.40	3.13
0.3789	16.58	63.20	7.14	3.49
0.4105	15.31	537.64	0.89	23.17
0.4420	14.21	944.68	-30.73	-0.75
0.4736	13.27	162.02	-0.95	-12.69
0.5052	12.44	908.09	-24.51	-17.53
0.5368	11.71	3197.84	34.84	-44.54
0.5683	11.06	1253.82	19.69	29.43
0.5999	10.47	850.45	-8.75	-27.82

---

## SSWD/DSSWD (Single/Double precision)

Estimate the nonnormalized spectral density of a stationary time series using a spectral window given the time series data.

### Usage

```
CALL SSWD (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, NF, F,  
          TINT, ISWVER, NM, M, PM, LDPM, SM, LDSM)
```

### Arguments

**NOBS** — Number of observations in the stationary time series  $x$ . (Input)  
NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the stationary time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

- 0 No printing is performed.
- 1 Print the periodogram, cosine transform and sine transform of the centered and padded time series, and the spectral density estimate based on a specified version of a spectral window for a given set of spectral window parameters.

**XCNTR** — Constant used to center the time series  $x$ . (Input)

**NPAD** — Number of zeroes used to pad the centered time series. (Input)  
NPAD must be greater than or equal to zero.

**IFSCAL** — Option for frequency scale. (Input)

#### IFSCAL Action

- 0 Frequency in radians per unit time.
- 1 Frequency in cycles per unit time.

**NF** — Number of frequencies at which to evaluate the spectral density estimate. (Input)

**F** — Vector of length NF containing the frequencies at which to evaluate the spectral density estimate. (Input)

The units of  $F$  correspond to the scale specified by **IFSCAL**. The elements of  $F$  must be in the range  $(-\pi/TINT, \pi/TINT)$ , inclusive for **IFSCAL** = 0 and  $(-1/(2 * TINT), 1/(2 * TINT))$  inclusive for **IFSCAL** = 1.

**TINT** — Time interval at which the series is sampled. (Input)

For a discrete parameter process, usually **TINT** = 1. For a continuous parameter process, **TINT** > 0. **TINT** is used to adjust the spectral density estimate.

**ISWVER** — Option for version of the spectral window. (Input)

#### ISWVER Action

- 1 Modified Bartlett

- 2 Daniell
- 3 Tukey-Hamming
- 4 Tukey-Hanning
- 5 Parzen
- 6 Bartlett-Priestley

Refer to the “Algorithm” section for further details.

**NM** — Number of spectral window parameters  $M$  used to compute the spectral density estimate for a given spectral window version. (Input)  
**NM** must be greater than or equal to one.

**M** — Vector of length **NM** containing the values of the spectral window parameters  $M$ . (Input)

For the Parzen spectral window (**ISWVER** = 5), all values of the spectral window parameters  $M$  must be even.

**PM** —  $(\lfloor N/2 \rfloor + 1)$  by 5 matrix that contains a summarization of the periodogram analysis. (Output)

For  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ , the  $(k + 1)$ -st element of the  $j$ -th column of **PM** is defined as

**Col. Description**

- 1 Frequency,  $\omega_k$  where  $\omega_k = 2\pi k/N$  for **IFSCAL** = 0 or  $\omega_k = k/N$  for **IFSCAL** = 1.
- 2 Period,  $p_k$  where  $p_k = 2\pi/\omega_k$  for **IFSCAL** = 0 and  $p_k = 1/\omega_k$  for **IFSCAL** = 1. If  $\omega_k = 0$ ,  $p_k$  is set to missing.
- 3 Periodogram ordinate,  $I(\omega_k)$ .
- 4 Cosine transformation coefficient,  $A(\omega_k)$ .
- 5 Sine transformation coefficient,  $B(\omega_k)$ .

Note  $N = \text{NOBS} + \text{NPAD}$ .

**LDPM** — Leading dimension of **PM** exactly as specified in the dimension statement of the calling program. (Input)

**LDPM** must be greater than or equal to  $\lfloor N/2 \rfloor + 1$ .

**SM** —  $\text{NF}$  by **NM** + 2 matrix containing a summarization of the spectral analysis. (Output)

The  $k$ -th element of the  $j$ -th column of **SM** is defined as

**Col. Description**

- 1 Frequency,  $F(k)$ .
- 2 Period,  $p_k$  where  $p_k = 2\pi/F(k)$  for **IFSCAL** = 0 and  $p_k = 1/F(k)$  for **IFSCAL** = 1. If  $F(k) = 0$ ,  $p_k$  is set to missing.
- 3 Spectral density estimate at  $F(k)$  using the spectral window parameter  $M(1)$ .
- NM** + 2 Spectral density estimate at  $F(k)$  using the spectral window parameter  $M(\text{NM})$ .

where  $k = 1, \dots, \text{NF}$ .

**LDSM** — Leading dimension of *SM* exactly as specified in the dimension statement of the calling program. (Input)  
**LDSM** must be greater than or equal to **NF**.

### Comments

- Automatic workspace storage is

**SSWD**  $10N + 15$  units, or  
**DSSWD**  $20N + 30$  units.

Workspace may be explicitly provided, if desired, by use of **S2WD/DS2WD**. The reference is

CALL **S2WD** (**NOBS**, **X**, **IPRINT**, **XCNTR**, **NPAD**, **IFSCAL**, **NF**,  
**F**, **TINT**, **ISWVER**, **NM**, **M**, **PM**, **LDPM**, **SM**,  
**LDSM**, **CX**, **COEF**, **WFFTC**, **CPY**)

The additional arguments are as follows:

**CX** — Complex vector of length  $N$  containing the centered and padded time series  $x$ . (Output)

**COEF** — Complex vector of length  $N$  containing the Fourier coefficients of the finite Fourier transform of **CX**. (Output)

Note that **COEF**( $k$ ) is the appropriately scaled Fourier coefficient at frequency  $\omega_k$ ,  $k = 0, 1, \dots, N - 1$ .

**WFFTC** — Vector of length  $4N + 15$ .

**CPY** — Vector of length  $2N$ .

- The normalized spectral density estimate is obtained by dividing the nonnormalized spectral density estimate in matrix **SM** by an estimate of the variance of  $x$ .

### Algorithm

Routine **SSWD** estimates the nonnormalized spectral density function of a stationary time series using a spectral window given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  for  $t = 1, 2, \dots, n$ .

Let

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  represent the centered and padded data where  $N = \text{NOBS} + \text{NPAD}$ ,

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu}_X & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_X = \text{XCNTR}$$

is determined by

$$\hat{\mu}_X = \begin{cases} \mu_X & \mu_X \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t & \mu_X \text{ unknown} \end{cases}$$

The modified periodogram of

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  is estimated by

$$I_{n,N,\tilde{X}}(\omega_k) = A_{\tilde{X}}^2(\omega_k) + B_{\tilde{X}}^2(\omega_k)$$

where

$$A_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \cos(\omega_k t)$$

and

$$B_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \sin(\omega_k t)$$

represent the

$$\tilde{X}_t$$

cosine and sine transforms, respectively, and  $K$  is the scale factor equal to  $1/(2\pi n)$ . Since the periodogram is an even function of the frequency, it is sufficient to estimate the periodogram over the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N} \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

The routine `PFFT` (page 723) is used to compute the modified periodogram of

$$\tilde{X}_t$$

The estimate of the nonnormalized spectral density  $h_X(\omega)$  is computed according to

$$\hat{h}_X(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} I_{n,N,\tilde{X}}(\omega_k) W_n(\omega - \omega_k)$$

where the spectral window  $W_n(\theta)$  is specified by argument `ISWVER`. The following spectral windows  $W_n(\theta)$  are available.

### Modified Bartlett

$$W_n(\theta) = \frac{1}{2\pi M} \left\{ \frac{\sin(M\theta/2)}{\sin(\theta/2)} \right\}^2 = F_M(\theta)$$

where  $F_M(\theta)$  corresponds to the Fejér kernel of order  $M$ .

### Daniell

$$W_n(\theta) = \begin{cases} M/2\pi & -\pi/M \leq \theta \leq \pi/M \\ 0 & \text{otherwise} \end{cases}$$

### Tukey

$$W_n(\theta) = aD_M\left(\theta - \frac{\pi}{M}\right) + (1-2a)D_M(\theta) + aD_M\left(\theta + \frac{\pi}{M}\right) \quad 0 < a \leq 0.25$$

where  $D_M(\theta)$  represents the Dirichlet kernel. The Tukey-Hamming window is obtained when  $a = 0.23$  and the Tukey-Hanning window is obtained when  $a = 0.25$ .

### Parzen

$$W_n(\theta) = \frac{6\pi}{M} [F_{M/2}(\theta)]^2 \left\{ 1 - \frac{2}{3} \sin^2(\theta/2) \right\}$$

where  $M$  is even. If  $M$  is odd, then  $M + 1$  is used instead of  $M$  in the above formula.

### Bartlett-Priestley

$$W_n(\theta) = \begin{cases} \frac{3M}{4\pi} \left\{ 1 - \left( \frac{M\theta}{\pi} \right)^2 \right\} & |\theta| \leq \pi/M \\ 0 & |\theta| > \pi/M \end{cases}$$

The argument  $NM$  specifies the number of window parameters  $M$  and corresponds to the number of spectral density estimates to be computed for a given spectral window. The nonnormalized spectral density is estimated over the set of frequencies

$$\omega = f_i \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale specified by the argument `IFSCAL` but are transformed to the scale of radians per unit time for computational purposes.

The above formula for

$$\hat{h}_X(\omega)$$

assumes the data  $\{X_t\}$  correspond to a realization of a discrete-parameter stationary process observed consecutively in time. In this case, the observations are equally spaced in time with interval  $\Delta t = \text{TINT}$  equivalent to one. However, if the data correspond to a realization of a continuous-parameter stationary process recorded at equal time intervals, then the estimate of the nonnormalized spectral density must be adjusted for the effect of aliasing. In general, the estimate of  $h_X(\omega)$  is given by

$$\hat{h}_X(\omega) = \Delta t \hat{h}_X(\omega) \quad |\omega| \leq \pi / \Delta t$$

Note that the frequency  $\omega$  of the desired spectral estimate is assumed to be input in a form already adjusted for the time interval  $\Delta t$ . Approximate confidence intervals for  $h(\omega)$  can be computed using formulas given in the introduction.

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Application of routine SSWD to these data produces the following results:

```

INTEGER      IPRINT, LDPM, LDSM, NF, NM, NOBS
REAL         PI
PARAMETER    (IPRINT=0, NF=20, NM=3, NOBS=100, PI=3.141592654,
&            LDPM=NOBS, LDSM=NF)
C
INTEGER      I, IFSCAL, ISWVER, M(NM), NCOL, NPAD, NROW
REAL         F(NF), PM(LDPM,5), RDATA(176,2), FLOAT, SM(LDSM,5),
&            SSUM, TINT, X(NOBS), XCNTR
CHARACTER    CLABEL(6)*9, FMT*20, RLABEL(1)*6, TITLE*60
INTRINSIC    REAL
EXTERNAL     GDATA, SSUM, SSWD, WRRRL
C
EQUIVALENCE (X(1), RDATA(22,2))
C
DATA RLABEL/'NONE'/, CLABEL/' ', 'Frequency', 'Period',
&         'M = 10', 'M = 20', 'M = 30'/
C                                     Wolfer Sunspot Data for
C                                     years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C                                     Center on arithmetic mean
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
C                                     Pad standard amount
NPAD = NOBS - 1
C                                     Frequency in radians per unit time
IFSCAL = 0
C                                     Determine frequencies at which
C                                     to evaluate spectral density
DO 10 I=1, NF
    F(I) = PI*FLOAT(I)/FLOAT(NF)
10 CONTINUE
C                                     Time interval for discrete data
TINT = 1.0

```

```

C                               Spectral window parameters
M(1) = 10
M(2) = 20
M(3) = 30

C                               Compute spectral density using
C                               the Parzen window
ISWVER = 5
CALL SSWD (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, NF, F, TINT,
&         ISWVER, NM, M, PM, LDPM, SM, LDSM)
C                               Print results
TITLE = 'Spectral Density Using the Parzen Window'
FMT    = '(F9.4, F6.2, 3F10.2)'
CALL WRRRL (TITLE, NF, 5, SM, LDSM, 0, FMT, RLABEL, CLABEL)
C                               Compute spectral density using
C                               the Bartlett-Priestley window
ISWVER = 6
CALL SSWD (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, NF, F, TINT,
&         ISWVER, NM, M, PM, LDPM, SM, LDSM)
C                               Print results
TITLE = '%/Spectral Density Using the Bartlett-Priestley '//
&       'Window'
CALL WRRRL (TITLE, NF, 5, SM, LDSM, 0, FMT, RLABEL, CLABEL)
C
END

```

### Output

```

Spectral Density Using the Parzen Window
Frequency  Period    M = 10    M = 20    M = 30
0.1571    40.00    659.64    617.42    619.73
0.3142    20.00    666.95    554.70    339.61
0.4712    13.33    653.73    770.64    860.49
0.6283    10.00    598.77    857.80    1046.13
0.7854     8.00    497.47    582.85    550.77
0.9425     6.67    367.72    266.33    186.98
1.0996     5.71    240.65    121.46    104.79
1.2566     5.00    142.41     76.17     76.74
1.4137     4.44     81.28     54.20     47.19
1.5708     4.00     49.13     40.16     41.39
1.7279     3.64     32.57     27.58     26.46
1.8850     3.33     22.44     16.52     14.40
2.0420     3.08     15.53     10.93     9.87
2.1991     2.86     11.19     8.30      8.32
2.3562     2.67     8.66      6.18      5.86
2.5133     2.50     6.93      4.75      4.22
2.6704     2.35     5.51      4.62      4.35
2.8274     2.22     4.47      4.91      5.24
2.9845     2.11     3.61      4.23      4.75
3.1416     2.00     2.62      2.44      2.27

```

```

Spectral Density Using the Bartlett-Priestley Window
Frequency  Period    M = 10    M = 20    M = 30
0.1571    40.00    604.34    712.73    757.61
0.3142    20.00    564.28    176.81    107.08
0.4712    13.33    767.63    927.14    981.10
0.6283    10.00    900.32    1190.30   1172.23
0.7854     8.00    607.45    494.85    571.65
0.9425     6.67    237.16    127.65     87.36
1.0996     5.71    103.34    113.93    135.34

```



1.2566	5.00	75.74	74.88	57.57
1.4137	4.44	52.64	44.98	38.59
1.5708	4.00	38.50	44.56	50.59
1.7279	3.64	27.35	25.28	21.76
1.8850	3.33	15.68	13.84	13.10
2.0420	3.08	10.33	9.79	7.41
2.1991	2.86	7.95	8.31	8.67
2.3562	2.67	6.04	5.86	7.08
2.5133	2.50	4.56	3.67	2.90
2.6704	2.35	4.44	4.38	4.06
2.8274	2.22	4.99	5.62	5.40
2.9845	2.11	4.31	5.07	5.08
3.1416	2.00	2.43	2.23	2.44

---

## SSWP/DSSWP (Single/Double precision)

Estimate the nonnormalized spectral density of a stationary time series using a spectral window given the periodogram.

### Usage

CALL SSWP (N, PX, NF, F, ISWVER, M, SX)

### Arguments

*N* — Number of observations in the centered and padded time series *x*. (Input)  
*N* must be greater than or equal to two.

*PX* — Vector of length  $\lfloor N/2 \rfloor + 1$  containing the (modified) periodogram of *x*. (Input)

The periodogram ordinate evaluated at (angular) frequency  $\omega_k = 2\pi k/N$  is given by  $PX(k + 1)$ ,  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ .

*NF* — Number of (angular) frequencies. (Input)  
*NF* must be greater than or equal to one.

*F* — Vector of length *NF* containing the (angular) frequencies at which the spectral density is estimated. (Input)  
The elements of *F* must be in the range  $(-\pi, \pi)$  inclusive.

*ISWVER* — Option for version of the spectral window. (Input)

#### *ISWVER* Action

- |   |                    |
|---|--------------------|
| 1 | Modified Bartlett  |
| 2 | Daniell            |
| 3 | Tukey-Hamming      |
| 4 | Tukey-Hanning      |
| 5 | Parzen             |
| 6 | Bartlett-Priestley |

Refer to the “Algorithm” section for further details.

**M** — Spectral window parameter. (Input)  
**M** must be greater than or equal to one and less than **N**.

**SX** — Vector of length **NF** containing the estimate of the spectral density of the time series **X**. (Output)

### Comments

1. The periodogram of **X** may be computed using the routine **PFFT** (page 723). Estimation of the spectral density of **X** using the modified periodogram preserves the scale of the spectral density up to adjustment for the time sampling interval.
2. The time sampling interval, **TINT**, is assumed to be equal to one. This assumption is appropriate for discrete parameter processes. The adjustment for continuous parameter processes (**TINT** > 0.0) involves multiplication of the frequency vector **F** by  $1/\text{TINT}$  and multiplication of the spectral density estimate by **TINT**.
3. To convert the frequency scale from radians per unit time to cycles per unit time, multiply **F** by  $1/(2\pi)$ .

### Algorithm

Routine **SSWP** estimates the nonnormalized spectral density function of a stationary time series using a spectral window given the modified periodogram of the appropriately centered and padded data

$$\{\tilde{X}_t\} \quad \text{for } t = 1, \dots, N$$

The routine **PFFT** (page 723) may be used to obtain the modified periodogram

$$I_{N,\tilde{X}}(\omega_k)$$

over the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

The symmetry of the periodogram is used to recover the ordinates at negative frequencies.

The estimate of the nonnormalized spectral density  $h_X(\omega)$  is computed according to

$$\hat{h}_X(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} I_{n,N,\tilde{X}}(\omega_k) W_n(\omega - \omega_k)$$

where the spectral window  $W_n(\theta)$  is specified by argument **ISWVER**. The following spectral windows  $W_n(\theta)$  are available.

### Modified Bartlett

$$W_n(\theta) = \frac{1}{2\pi M} \left\{ \frac{\sin(M\theta/2)}{\sin(\theta/2)} \right\}^2 = F_M(\theta)$$

where  $F_M(\theta)$  corresponds to the Fejer kernel of order  $M$ .

### Daniell

$$W_n(\theta) = \begin{cases} M/2\pi & -\pi/M \leq \theta \leq \pi/M \\ 0 & \text{otherwise} \end{cases}$$

### Tukey

$$W_n(\theta) = aD_M\left(\theta - \frac{\pi}{M}\right) + (1-2a)D_M(\theta) + aD_M\left(\theta + \frac{\pi}{M}\right), \quad 0 < a \leq 0.25$$

where  $D_M(\theta)$  represents the Dirichlet kernel. The Tukey-Hamming window is obtained when  $a = 0.23$ , and the Tukey-Hanning window is obtained when  $a = 0.25$ .

### Parzen

$$W_n(\theta) = \frac{6\pi}{M} [F_{M/2}(\theta)]^2 \left\{ 1 - \frac{2}{3} \sin^2(\theta/2) \right\}$$

where  $M$  is even. If  $M$  is odd, then  $M + 1$  is used instead of  $M$  in the above formula.

### Bartlett-Priestley

$$W_n(\theta) = \begin{cases} \frac{3M}{4\pi} \left\{ 1 - \left( \frac{M\theta}{\pi} \right)^2 \right\} & |\theta| \leq \pi/M \\ 0 & |\theta| > \pi/M \end{cases}$$

Only one window parameter  $M$  may be specified so that only one estimate of  $h_X(\omega)$  is computed. The nonnormalized spectral density is estimated over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale of radians per unit time. The time sampling interval  $\Delta t$  is assumed to be equal to one.

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set

for this example consists of the number of sunspots observed from 1770 through 1869. Application of routine SSWP to these data produces the following results:

```

INTEGER      IPRINT, LDPM, LDSM, NF, NM, NOBS
REAL         PI
PARAMETER    (IPRINT=0, NF=20, NM=3, NOBS=100, PI=3.141592654,
&           LDPM=NOBS, LDSM=NF)
C
INTEGER      I, IFSCAL, IPVER, ISWVER, J, M(NM), N, NCOL, NPAD,
&           NROW
REAL         F(NF), PM(LDPM,5), PX(LDPM), RDATA(176,2), FLOAT,
&           SM(NF,5), SSUM, SX(NF), X(NOBS), XCNTR
CHARACTER    CLABEL(6)*9, FMT*20, RLABEL(1)*6, TITLE*60
INTRINSIC    FLOAT
EXTERNAL     GDATA, PFFT, SCOPY, SSUM, SSWP, WRRRL
C
EQUIVALENCE (PX(1), PM(1,3)), (F(1), SM(1,1))
EQUIVALENCE (X(1), RDATA(22,2))
C
DATA RLABEL/'NONE'/, CLABEL/' ', 'Frequency', 'Period',
&         'M = 10', 'M = 20', 'M = 30'/
C                                     Wolfer Sunspot Data for
C                                     years 1770 through 1869
CALL GDATA (2, 0, NROW, NCOL, RDATA, 176, 2)
C                                     Center on arithmetic mean
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
C                                     Pad standard amount
NPAD = NOBS - 1
C                                     Frequency in radians per unit time
IFSCAL = 0
C                                     Modified periodogram version
IPVER = 1
C                                     Compute periodogram
CALL PFFT (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, IPVER, PM, LDPM)
C                                     Number of observations used to
C                                     compute the periodogram
N = NOBS + NPAD
C                                     Determine frequency and period
C                                     at which to evaluate the spectral
C                                     density
DO 10 I=1, NF
    SM(I,1) = PI*FLOAT(I)/FLOAT(NF)
    SM(I,2) = 2.0*FLOAT(NF)/FLOAT(I)
10 CONTINUE
C                                     Spectral window parameters
M(1) = 10
M(2) = 20
M(3) = 30
C                                     Compute spectral density using
C                                     the Parzen window
ISWVER = 5
DO 20 J=1, NM
    CALL SSWP (N, PX, NF, F, ISWVER, M(J), SX)
C                                     Copy into SM
    CALL SCOPY (NF, SX, 1, SM(1,2+J), 1)
20 CONTINUE
C                                     Print results
TITLE = 'Spectral Density Using the Parzen Window'
FMT    = '(F9.4, F6.2, 3F10.2)'

```

```

C      CALL WRRRL (TITLE, NF, 5, SM, LDSM, 0, FMT, RLABEL, CLABEL)
C      Compute spectral density using
C      the Bartlett-Priestley window
      ISWVER = 6
      DO 30 J=1, NM
        CALL SSWP (N, PX, NF, F, ISWVER, M(J), SX)
C      Copy into SM
        CALL SCOPY (NF, SX, 1, SM(1,2+J), 1)
30 CONTINUE
C      Print results
      TITLE = '%/Spectral Density Using the Bartlett-Priestley '//
& 'Window'
      CALL WRRRL (TITLE, NF, 5, SM, LDSM, 0, FMT, RLABEL, CLABEL)
C
      END

```

### Output

```

Spectral Density Using the Parzen Window
Frequency  Period    M = 10    M = 20    M = 30
0.1571    40.00    659.64    617.42    619.73
0.3142    20.00    666.95    554.70    339.61
0.4712    13.33    653.73    770.64    860.49
0.6283    10.00    598.77    857.80    1046.13
0.7854     8.00    497.47    582.85    550.77
0.9425     6.67    367.72    266.33    186.98
1.0996     5.71    240.65    121.46    104.79
1.2566     5.00    142.41     76.17     76.74
1.4137     4.44     81.28     54.20     47.19
1.5708     4.00     49.13     40.16     41.39
1.7279     3.64     32.57     27.58     26.46
1.8850     3.33     22.44     16.52     14.40
2.0420     3.08     15.53     10.93     9.87
2.1991     2.86     11.19      8.30     8.32
2.3562     2.67      8.66      6.18     5.86
2.5133     2.50      6.93      4.75     4.22
2.6704     2.35      5.51      4.62     4.35
2.8274     2.22      4.47      4.91     5.24
2.9845     2.11      3.61      4.23     4.75
3.1416     2.00      2.62      2.44     2.27

```

```

Spectral Density Using the Bartlett-Priestley Window
Frequency  Period    M = 10    M = 20    M = 30
0.1571    40.00    604.34    712.73    757.61
0.3142    20.00    564.28    176.81    107.08
0.4712    13.33    767.63    927.14    981.10
0.6283    10.00    900.32    1190.30   1172.23
0.7854     8.00    607.45    494.85    571.65
0.9425     6.67    237.16    127.65     87.36
1.0996     5.71    103.34    113.93    135.34
1.2566     5.00     75.74     74.88     57.57
1.4137     4.44     52.64     44.98     38.59
1.5708     4.00     38.50     44.56     50.59
1.7279     3.64     27.35     25.28     21.76
1.8850     3.33     15.68     13.84     13.10
2.0420     3.08     10.33      9.79      7.41
2.1991     2.86      7.95      8.31      8.67
2.3562     2.67      6.04      5.86      7.08
2.5133     2.50      4.56      3.67      2.90

```

2.6704	2.35	4.44	4.38	4.06
2.8274	2.22	4.99	5.62	5.40
2.9845	2.11	4.31	5.07	5.08
3.1416	2.00	2.43	2.23	2.44

---

## SWED/DSWED (Single/Double precision)

Estimation of the nonnormalized spectral density of a stationary time series based on specified periodogram weights given the time series data.

### Usage

```
CALL SWED (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, NF, F,
           TINT, NWT, WT, PM, LDPM, SM, LDSM)
```

### Arguments

**NOBS** — Number of observations in the stationary time series *x*. (Input)  
NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the stationary time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

- 0 No printing is performed.
- 1 Print the periodogram, cosine and sine transforms of the centered and padded time series, and the spectral density estimate based on a specified weight sequence.

**XCNTR** — Constant used to center the time series *x*. (Input)

**NPAD** — Number of zeroes used to pad the centered time series. (Input)  
NPAD must be greater than or equal to zero. The length of the centered and padded time series is  $N = \text{NOBS} + \text{NPAD}$ .

**IFSCAL** — Option for frequency scale. (Input)

#### IFSCAL Action

- 0 Frequency in radians per unit time.
- 1 Frequency in cycles per unit time.

**NF** — Number of frequencies at which to evaluate the spectral density estimate. (Input)

NF must be greater than zero.

**F** — Vector of length NF containing the frequencies at which to evaluate the spectral density estimate. (Input)

The units of *F* correspond to the scale specified by *IFSCAL*. The elements of *F* must be in the range  $(-\pi/\text{TINT}, \pi/\text{TINT})$ , inclusive, for *IFSCAL* = 0 and  $(-1/(2 * \text{TINT}), 1/(2 * \text{TINT}))$ , inclusive, for *IFSCAL* = 1.

**TINT** — Time interval at which the series is sampled. (Input)  
 For a discrete parameter process, usually  $TINT = 1.0$ . For a continuous parameter process,  $TINT > 0.0$ .  $TINT$  is used to adjust the spectral density estimate.

**NWT** — Number of weights. (Input)  
 $NWT$  must be greater than or equal to one.

**WT** — Vector of length  $NWT$  containing the weights used to smooth the periodogram. (Input)  
 The actual weights are the values in  $WT$  normalized to sum to 1 with the current periodogram ordinate taking the middle weight for  $NWT$  odd or the weight to the right of the middle for  $NWT$  even.

**PM** —  $(\lfloor N/2 \rfloor + 1)$  by 5 matrix that contains a summarization of the periodogram analysis. (Output)  
 For  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ , the  $(k + 1)$ -st element of the  $j$ -th column of  $PM$  is defined as

Col.	Description
1	Frequency, $\omega_k$ where $\omega_k = 2\pi k/N$ for $IFSCAL = 0$ or $\omega_k = k/N$ for $IFSCAL = 1$ .
2	Period, $p_k$ where $p_k = 2\pi/\omega_k$ for $IFSCAL = 0$ and $p_k = 1/\omega_k$ for $IFSCAL = 1$ . If $\omega_k = 0$ , $p_k$ is set to the missing value or NaN (not a number).
3	Periodogram ordinate, $I(\omega_k)$ .
4	Cosine transformation coefficient, $A(\omega_k)$ .
5	Sine transformation coefficient, $B(\omega_k)$ .

**LDPM** — Leading dimension of  $PM$  exactly as specified in the dimension statement in the calling program. (Input)  
 $LDPM$  must be greater than or equal to  $\lfloor N/2 \rfloor + 1$ .

**SM** —  $NF$  by 3 matrix containing a summarization of the spectral analysis. (Output)  
 The  $k$ -th element of the  $j$ -th column of  $SM$  is defined as

Col.	Description
1	Frequency, $F(k)$ .
2	Period, $p_k$ where $p_k = 2\pi/F(k)$ for $IFSCAL = 0$ and $p_k = 1/F(k)$ for $IFSCAL = 1$ . If $F(k) = 0$ , $p_k$ is set to missing.
3	Spectral density estimate at $F(k)$ using the specified weights $WT$ , where $k = 1, \dots, NF$ .

**LDSM** — Leading dimension of  $SM$  exactly as specified in the dimension statement in the calling program. (Input)  
 $LDSM$  must be greater than or equal to  $NF$ .

## Comments

1. Automatic workspace storage is

SWED  $10N + 15$  or  
DSWED  $20N + 30$  units.

Workspace may be explicitly provided, if desired, by use of S2ED/DS2ED. The reference is

```
CALL S2ED (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL,  
          NF, F, TINT, NWT, WT, PM, LDPM, SM, LDSM,  
          CX, COEF, WFFTC, CPY)
```

The additional arguments are as follows:

**CX** — Complex vector of length  $N$  containing the centered and padded time series  $x$ . (Output)

**COEF** — Complex vector of length  $N$  containing the Fourier coefficients of the finite Fourier transform of **CX**. (Output)

Note that  $\text{COEF}(k + 1)$  is the appropriately scaled Fourier coefficient at frequency  $\omega_k$ ,  $k = 0, 1, \dots, N - 1$ .

**WFFTC** — Work vector of length  $4N + 15$ .

**CPY** — Work vector of length  $2N$ .

2. The centered and padded time series is defined by

$$\begin{aligned} \text{CX}(j) &= x(j) - \text{XCNTR} && \text{for } j = 1, \dots, \text{NOBS} \\ \text{CX}(j) &= 0 && \text{for } j = \text{NOBS} + 1, \dots, N \end{aligned}$$

where  $N = \text{NOBS} + \text{NPAD}$ .

3. The normalized spectral density estimate is obtained by dividing the nonnormalized spectral density estimate in matrix **SM** by an estimate of the variance of  $x$ .

## Algorithm

Routine **SWED** estimates the nonnormalized spectral density function of a stationary time series using a fixed sequence of weights, given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$ , for  $t = 1, 2, \dots, n$ .

Let

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

represent the centered and padded data where  $N = \text{NOBS} + \text{NPAD}$ ,

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu}_X, & t = 1, \dots, n \\ 0, & t = (n + 1), \dots, N \end{cases}$$

and

$$\hat{\mu}_X = \text{XCNTR}$$



is determined by

$$\hat{\mu}_X = \begin{cases} \mu_X, & \mu_X \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t, & \mu_X \text{ unknown} \end{cases}$$

The modified periodogram of

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

is estimated by

$$I_{n,N,\tilde{X}}(\omega_k) = A_{\tilde{X}}^2(\omega_k) + B_{\tilde{X}}^2(\omega_k)$$

where

$$A_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \cos(\omega_k t)$$

and

$$B_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \sin(\omega_k t)$$

represent the

$$\tilde{X}_t$$

cosine and sine transforms, respectively, and  $K$  is the scale factor equal to  $1/(2\pi n)$ . Since the periodogram is an even function of the frequency, it is sufficient to estimate the periodogram at the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

(Here,  $\lfloor a \rfloor$  means the greatest integer less than or equal to  $a$ .) The routine `PFFFT` (page 723) is used to compute the modified periodogram of

$$\{\tilde{X}_t\}$$

Consider the sequence of  $m = \text{NWT}$  weights

$$\{w_j\} \text{ for } j = -\lfloor m/2 \rfloor, \dots, (m - \lfloor m/2 \rfloor - 1)$$

where

$$\sum_j w_j = 1$$

These weights are fixed in the sense that they do not depend on the frequency  $\omega$  at which to estimate the nonnormalized spectral density  $h_X(\omega)$ . The estimate of the nonnormalized spectral density is computed according to

$$\hat{h}_X(\omega) = \sum_j \omega_j I_{n,N,\tilde{X}}(\omega_{k,j})$$

where

$$\omega_{k,j} = \frac{2\pi\{k(\omega) + j\}}{N}$$

and  $k(\omega)$  is the integer such that  $\omega_{k,0}$  is closest to  $\omega$ . The weights specified by argument `WT` may be relative since they are normalized to sum to one in the actual computation of

$$\hat{h}_X(\omega)$$

Usually,  $m$  is odd with the weights symmetric about the middle weight  $w_0$ . If  $m$  is even, the weight to the right of the middle is considered  $w_0$ . Note that periodogram ordinate

$$I_{n,N,\tilde{X}}(0)$$

is replaced by

$$I_{n,N,\tilde{X}}(\omega_1)$$

and the sum reflects at each end. The nonnormalized spectral density is estimated over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale specified by the argument `IFSCAL` but are transformed to the scale of radians per unit time for computational purposes.

The above formula for

$$\hat{h}_X(\omega)$$

assumes the data  $\{X_t\}$  correspond to a realization of a discrete-parameter stationary process observed consecutively in time. In this case, the observations are equally spaced in time with interval  $\Delta t = \text{TINT}$  equivalent to one. However, if the data correspond to a realization of a continuous-parameter stationary process recorded at equal time intervals, then the estimate of the nonnormalized spectral density must be adjusted for the effect of aliasing. In general, the estimate of  $h_X(\omega)$  is given by

$$\hat{h}_X(\omega) = \Delta t \hat{h}_X(\omega), \quad |\omega| \leq \pi / \Delta t$$

Note that the frequency  $\omega$  of the desired spectral estimate is assumed to be input in a form already adjusted for the time interval  $\Delta t$ .

Approximate confidence intervals for  $h(\omega)$  can be computed using formulas given in the introduction.

## Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Application of routine SWED to these data produces the following results:

```
INTEGER      LDPM, LDRDAT, LDSM, NDRDAT, NF, NOBS, NPAD, NWT
PARAMETER    (LDRDAT=176, NDRDAT=2, NF=20, NOBS=100, NWT=7,
&            LDSM=NF, NPAD=NOBS-1, LDPM=(NOBS+NPAD)/2+1)
C
INTEGER      I, IFSCAL, IPRINT, NROW, NVAR
REAL         ASIN, F(NF), PI, PM(LDPM,5), RDATA(LDRDAT,NDRDAT),
&            REAL, SM(LDSM,3), SSUM, TINT, WT(NWT), X(NOBS), XCNTR
CHARACTER    CLABEL(4)*20, FMT*20, RLABEL(1)*4, TITLE*28
INTRINSIC    ASIN, FLOAT
EXTERNAL     GDATA, SSUM, SWED, WRRRL
C
EQUIVALENCE (X(1), RDATA(22,2))
C
DATA WT/1.0, 2.0, 3.0, 4.0, 3.0, 2.0, 1.0/
DATA IPRINT/0/, IFSCAL/0/, TINT/1.0/
DATA FMT/'(F9.4, F6.2, F9.4)'/
DATA RLABEL/'NONE'/
DATA CLABEL/' ', '%/Frequency', '%/Period', 'Spectral%/Estimates'
& /
DATA TITLE/'Results of Spectral Analysis'/
C                               Initializations
PI = 2.0*ASIN(1.0)
DO 10 I=1, NF
    F(I) = PI*FLOAT(I)/FLOAT(NF)
10 CONTINUE
C                               Wolfer Sunspot Data for years
C                               1770 through 1869
CALL GDATA (2, 0, NROW, NVAR, RDATA, LDRDAT, NDRDAT)
C                               Center on arithmetic mean
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
C                               Spectral density
CALL SWED (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, NF, F, TINT,
&          NWT, WT, PM, LDPM, SM, LDSM)
C                               Print Results
CALL WRRRL (TITLE, NF, 3, SM, LDSM, 0, FMT, RLABEL, CLABEL)
C
END
```

## Output

```
Results of Spectral Analysis
Spectral
Frequency  Period  Estimates
0.1571    40.00  710.8386
0.3142    20.00  116.3940
0.4712    13.33  937.1508
0.6283    10.00  1209.8268
0.7854     8.00  538.9236
0.9425     6.67  84.9561
1.0996     5.71  128.0791
1.2566     5.00  55.0304
```

1.4137	4.44	40.2022
1.5708	4.00	46.4240
1.7279	3.64	21.0053
1.8850	3.33	12.1449
2.0420	3.08	8.8654
2.1991	2.86	7.2589
2.3562	2.67	6.8078
2.5133	2.50	3.3873
2.6704	2.35	3.9504
2.8274	2.22	5.7418
2.9845	2.11	4.4652
3.1416	2.00	4.1216

---

## SWEP/DSWEP (Single/Double precision)

Estimation of the nonnormalized spectral density of a stationary time series based on specified periodogram weights given the periodogram.

### Usage

CALL SWEP (N, PX, NF, F, NWT, WT, SX)

### Arguments

**N** — Number of observations in the appropriately centered and padded time series  $x$ . (Input)

**N** must be greater than or equal to two.

**PX** — Vector of length  $\lfloor N/2 \rfloor + 1$  containing the (modified) periodogram of  $x$ . (Input)

The periodogram ordinate evaluated at (angular) frequency  $\omega_k = 2\pi k/N$  is given by  $PX(k + 1)$ ,  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ .

**NF** — Number of (angular) frequencies. (Input)

**NF** must be greater than or equal to one.

**F** — Vector of length **NF** containing the (angular) frequencies at which the spectral density is estimated. (Input)

The elements of **F** must be in the range  $(-\pi, \pi)$  inclusive.

**NWT** — Number of weights. (Input)

**NWT** must be greater than or equal to one.

**WT** — Vector of length **NWT** containing the weights used to smooth the periodogram. (Input)

The actual weights are the values in **WT** normalized to sum to 1 with the current periodogram ordinate taking the middle weight for **NWT** odd or the weight to the right of the middle for **NWT** even.

**SX** — Vector of length **NF** containing the estimate of the spectral density of the time series  $x$ . (Output)

### Comments

1. The periodogram of  $x$  may be computed using the routine `PFFT` (page 723). Estimation of the spectral density of  $x$  using the modified periodogram preserves the scale of the spectral density up to adjustment for the time sampling interval.
2. The time sampling interval, `TINT`, is assumed to be equal to one. This assumption is appropriate for discrete parameter processes. The adjustment for continuous parameter processes (`TINT` > 0) involves multiplication of the frequency vector  $F$  by  $1/TINT$  and multiplication of the spectral density estimate by `TINT`.
3. To convert the frequency scale from radians per unit time to cycles per unit time, multiply  $F$  by  $1/(2\pi)$ .

### Algorithm

Routine `SWEP` estimates the nonnormalized spectral density function of a stationary time series using a fixed sequence of weights given the modified periodogram of the appropriately centered and padded data

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

The routine `PFFT` (page 723) may be used to obtain the modified periodogram

$$I_{n,N,\tilde{X}}(\omega_k)$$

over the discrete set of nonnegative frequencies

$$w_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

(Here,  $\lfloor a \rfloor$  means the greatest integer less than or equal to  $a$ .) The symmetry of the periodogram is used to recover the ordinates at negative frequencies.

Consider the sequence of  $m = NWT$  weights  $\{w_j\}$  for  $j = -\lfloor m/2 \rfloor, \dots, (m - \lfloor m/2 \rfloor - 1)$  where  $\sum_j w_j = 1$ . These weights are fixed in the sense that they do not depend on the frequency  $\omega$  at which to estimate the nonnormalized spectral density  $h_X(\omega)$ . The estimate of the nonnormalized spectral density is computed according to

$$\hat{h}_X(\omega) = \sum_j \omega_j I_{n,N,\tilde{X}}(\omega_{k,j})$$

where

$$\omega_{k,j} = \frac{2\pi\{k(\omega) + j\}}{N}$$

and  $k(\omega)$  is the integer such that  $\omega_{k,0}$  is closest to  $\omega$ . The weights specified by argument `WT` may be relative since they are normalized to sum to one in the actual computation of

$$\hat{h}_X(\omega)$$

Usually,  $m$  is odd with the weights symmetric about the middle weight  $w_0$ . If  $m$  is even, the weight to the right of the middle is considered  $w_0$ . Note that periodogram ordinate

$$I_{n,N,\tilde{X}}(0)$$

is replaced by

$$I_{n,N,\tilde{X}}(\omega_1)$$

and the sum reflects at each end.

The nonnormalized spectral density estimate is computed over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale of radians per unit time. The time sampling interval  $\Delta t$  is assumed to be equal to one.

Approximate confidence intervals for  $h(\omega)$  can be computed using formulas given in the introduction.

### Example

Consider the Wölfer Sunspot Data (Anderson 1971, page 660) consisting of the number of sunspots observed each year from 1749 through 1924. The data set for this example consists of the number of sunspots observed from 1770 through 1869. Application of routine SWEF to these data produces the following results:

```

INTEGER      LDPM, LDRDAT, N, NDRDAT, NF, NOBS, NPAD, NWT
PARAMETER    (LDRDAT=176, NDRDAT=2, NF=20, NOBS=100, NWT=7,
&            NPAD=NOBS-1, LDPM=(NOBS+NPAD)/2+1, N=NOBS+NPAD)
C
INTEGER      I, IFSCAL, IPRINT, IPVER, NROW, NVAR
REAL         ASIN, F(NF), PI, PM(LDPM,5), RDATA(LDRDAT,NDRDAT),
&            FLOAT, SM(NF,2), SSUM, SX(NF), TINT, WT(NWT), X(NOBS),
&            XCNTR
CHARACTER    CLABEL(3)*30, FMT*20, RLABEL(1)*4, TITLE*28
INTRINSIC    ASIN, FLOAT
EXTERNAL     GDATA, PFFT, SCOPY, SSUM, SWEF, WRRRL
C
EQUIVALENCE (X(1), RDATA(22,2))
C
DATA WT/1., 2., 3., 4., 3., 2., 1./
DATA IPRINT/0/, TINT/1.0/, IPVER/1/, IFSCAL/0/
DATA FMT/'(F9.4)'/
DATA CLABEL/'      ', '%/Frequency', 'Spectral%/Estimates'/
DATA RLABEL/'NONE'/
DATA TITLE/'Results of Spectral Analysis'/
C
                                Initialization
PI = 2.0*ASIN(1.0)
DO 10 I=1, NF
    F(I) = PI*FLOAT(I)/FLOAT(NF)

```

```

10 CONTINUE
C                               Wolfer Sunspot Data for years
C                               1770 through 1869
  CALL GDATA (2, 0, NROW, NVAR, RDATA, LDRDAT, NDRDAT)
C                               Compute mean
  XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
C                               Compute modified periodogram
  CALL PFFT (NOBS, X, IPRINT, XCNTR, NPAD, IFSCAL, IPVER, PM, LDPM)
C
C                               Compute spectral density
  CALL SWEP (N, PM(1,3), NF, F, NWT, WT, SX)
C
C                               Print results
C
C                               Copy the frequencies to the output
C                               matrix
  CALL SCOPY (NF, F, 1, SM(1,1), 1)
C                               Copy the spectral estimates to the
C                               output matrix
  CALL SCOPY (NF, SX, 1, SM(1,2), 1)
C                               Call printing routine
  CALL WRRRL (TITLE, NF, 2, SM, NF, 0, FMT, RLABEL, CLABEL)
C
  END

```

### Output

Results of Spectral Analysis

Frequency	Spectral Estimates
0.1571	710.8386
0.3142	116.3940
0.4712	937.1508
0.6283	1209.8268
0.7854	538.9236
0.9425	84.9561
1.0996	128.0791
1.2566	55.0304
1.4137	40.2022
1.5708	46.4240
1.7279	21.0053
1.8850	12.1449
2.0420	8.8654
2.1991	7.2589
2.3562	6.8078
2.5133	3.3873
2.6704	3.9504
2.8274	5.7418
2.9845	4.4652
3.1416	4.1216

---

## CPFFT/DCPFFT (Single/Double precision)

Compute the cross periodogram of two stationary time series using a fast Fourier transform.

## Usage

CALL CPFFT (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL,  
IPVER, CPM, LDCPM)

## Arguments

**NOBS** — Number of observations in each stationary time series  $X$  and  $Y$ . (Input)  
NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the first stationary time series. (Input)

**Y** — Vector of length NOBS containing the second stationary time series. (Input)

**IPRINT** — Printing option. (Input)

### IPRINT Action

- 0 No printing is performed.
- 1 Print the periodogram, cosine and sine series, and the real and imaginary components of the cross periodogram.

**XCNTR** — Constant used to center the time series  $X$ . (Input)

**YCNTR** — Constant used to center the time series  $Y$ . (Input)

**NPAD** — Number of zeroes used to pad each centered time series. (Input)  
NPAD must be greater than or equal to zero. The length of each centered and padded time series is  $N = \text{NOBS} + \text{NPAD}$ .

**IFSCAL** — Option for frequency scale. (Input)

### IFSCAL Action

- 0 Frequency in radians per unit time
- 1 Frequency in cycles per unit time

**IPVER** — Option for version of the periodogram. (Input)

### IPVER Action

- 0 Compute usual periodogram.
- 1 Compute modified periodogram.

Refer to the algorithm section for further details.

**CPM** —  $(\lfloor N/2 \rfloor + 1)$  by 10 matrix containing a summarization of the results of the cross periodogram analysis. (Output)

For  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ , the  $(k + 1)$ -st element of the  $j$ -th column of CPM is defined as

### Col. Description

- 1 Frequency,  $\omega_k$  where  $\omega_k = 2\pi k/N$  for  $\text{IFSCAL} = 0$  or  $\omega_k = k/N$  for  $\text{IFSCAL} = 1$ .
- 2 Period,  $p_k$  where  $p_k = 2\pi/\omega_k$  for  $\text{IFSCAL} = 0$  and  $p_k = 1/\omega_k$  for  $\text{IFSCAL} = 1$ . If  $\omega_k = 0$ ,  $p_k$  is set to missing.



- 3 X periodogram ordinate,  $I_X(\omega_k)$
- 4 X cosine transformation coefficient,  $A_X(\omega_k)$
- 5 X sine transformation coefficient,  $B_X(\omega_k)$
- 6 Y periodogram ordinate,  $I_Y(\omega_k)$
- 7 Y cosine transformation coefficient,  $A_Y(\omega_k)$
- 8 Y sine transformation coefficient,  $B_Y(\omega_k)$
- 9 Real part of the XY cross periodogram ordinate  $I_{XY}(\omega_k)$ .
- 10 Imaginary part of the XY cross periodogram ordinate  $I_{XY}(\omega_k)$ .

**LDCPM** — Leading dimension of CPM exactly as specified in the dimension statement of the calling program. (Input)  
 LDCPM must be greater than or equal to  $\lfloor N/2 \rfloor + 1$ .

### Comments

1. Automatic workspace storage is

CPFFT  $10N + 15$  units, or  
 DCPFFT  $20N + 30$  units.

Workspace may be explicitly provided, if desired, by use of C2FFT/DC2FFT. The reference is

```
CALL C2FFT (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD,
           IFSCAL, IPVER, CPM, LDCPM, CX, COEF,
           WFFTC, CPY)
```

The additional arguments are as follows:

**CX** — Complex work vector of length  $N$ .

**COEF** — Complex work vector of length  $N$ .

**WFFTC** — Work vector of length  $4N + 15$ .

**CPY** — Work vector of length  $2N$ .

2. The centered and padded time series are defined by

$$\begin{aligned} CX(j) &= X(j) - XCNTR && \text{for } j = 1, \dots, \text{NOBS} \\ CX(j) &= 0 && \text{for } j = \text{NOBS} + 1, \dots, N \end{aligned}$$

and

$$\begin{aligned} CY(j) &= Y(j) - YCNTR && \text{for } j = 1, \dots, \text{NOBS} \\ CY(j) &= 0 && \text{for } j = \text{NOBS} + 1, \dots, N \end{aligned}$$

where  $N = \text{NOBS} + \text{NPAD}$ .

3. The cross periodogram  $I_{XY}(\omega)$  is complex valued in general. The relation  $I_{XY}(-\omega) = \text{conj}(I_{XY}(\omega))$  for  $\omega > 0.0$  recovers the cross periodogram for negative frequencies since  $\text{real}(I_{XY}(-\omega)) = \text{real}(I_{XY}(\omega))$  and  $\text{imag}(I_{XY}(-\omega)) = -\text{imag}(I_{XY}(\omega))$ . The periodogram  $I(\omega)$  is an even function of the frequency  $\omega$ . The relation  $I(-\omega) = I(\omega)$  for  $\omega > 0.0$  recovers the periodogram for negative frequencies.

4. Since  $\cos(\omega)$  is an even function of  $\omega$  and  $\sin(\omega)$  is an odd function of  $\omega$ , the cosine and sine transformations, respectively, satisfy  $A(-\omega) = A(\omega)$  and  $B(-\omega) = -B(\omega)$  for  $\omega > 0.0$ . Similarly, the complex Fourier coefficients, stored in `COEF`, satisfy  $\text{COEF}(-\omega) = \text{conj}(\text{COEF}(\omega))$ .
5. Computation of the  $2 * \text{NOBS} - 1$  cross-covariances of  $X$  and  $Y$  using the inverse Fourier transform of the cross periodogram requires  $\text{NPAD} = \text{NOBS} - 1$ .

### Algorithm

Routine `CPFFT` computes the cross periodogram of two jointly stationary time series given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  and  $\{Y_t\}$  for  $t = 1, 2, \dots, n$ .

Let

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

represent the centered and padded data where  $N = \text{NOBS} + \text{NPAD}$ ,

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu}_X & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_X = \text{XCNTR}$$

is determined by

$$\hat{\mu}_X = \begin{cases} \mu_X & \mu_X \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t & \mu_X \text{ unknown} \end{cases}$$

Similarly, let

$$\{\tilde{Y}_t\} \text{ for } t = 1, \dots, N$$

represent the centered and padded data where

$$\tilde{Y}_t = \begin{cases} Y_t - \hat{\mu}_Y & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_Y = \text{YCNTR}$$

is determined by

$$\hat{\mu}_Y = \begin{cases} \mu_Y & \mu_Y \text{ known} \\ \frac{1}{n} \sum_{t=1}^n Y_t & \mu_Y \text{ unknown} \end{cases}$$

The periodogram of the sample sequence  $\{X_t\}$ ,  $t = 1, \dots, n$  computed with the padded sequence

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

is defined by

$$I_{n,N,\tilde{X}}(\omega_k) = A_{\tilde{X}}^2(\omega_k) + B_{\tilde{X}}^2(\omega_k)$$

where

$$A_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \cos(\omega_k t)$$

and

$$B_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \sin(\omega_k t)$$

represent the

$$\tilde{X}_t$$

cosine and sine transforms, respectively, and  $K$  is the scale factor

$$K = \begin{cases} \frac{2}{n} & \text{for the usual periodogram,} \\ \frac{1}{2\pi n} & \text{for the modified periodogram} \end{cases}$$

The periodogram of the sample sequence  $\{Y_t\}$ ,  $t = 1, \dots, n$  computed with the padded sequence

$$\{\tilde{Y}_t\} \text{ for } t = 1, \dots, N$$

is defined by

$$I_{n,N,\tilde{Y}}(\omega_k) = A_{\tilde{Y}}^2(\omega_k) + B_{\tilde{Y}}^2(\omega_k)$$

where

$$A_{\tilde{Y}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{Y}_t \cos(\omega_k t)$$

and

$$B_{\tilde{Y}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{Y}_t \sin(\omega_k t)$$

represent the

$$\{\tilde{Y}_t\}$$

cosine and sine transforms, respectively. Since the periodogram is an even function of the frequency, it is sufficient to estimate the periodogram at the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N} \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

(Here,  $\lfloor a \rfloor$  means the greatest integer less than or equal to  $a$ ). The routine `PPFFT` (page 723) is used to compute the periodograms of both

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

according to the version specified by the argument `IPVER`. The computational formula for the cross periodogram is given by

$$I_{n,N,\tilde{X}\tilde{Y}}(\omega_k) = \Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} + i\Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\}$$

where

$$\Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} = A_{\tilde{X}}(\omega_k)A_{\tilde{Y}}(\omega_k) + B_{\tilde{X}}(\omega_k)B_{\tilde{Y}}(\omega_k)$$

and

$$\Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} = A_{\tilde{X}}(\omega_k)B_{\tilde{Y}}(\omega_k) - B_{\tilde{X}}(\omega_k)A_{\tilde{Y}}(\omega_k)$$

The real part of the (modified) cross periodogram represents the 'raw' sample cospectrum and the negative of the imaginary part of the (modified) cross periodogram represents the 'raw' sample quadrature spectrum (Priestley 1981, page 695). The relationship between the cross periodogram and its complex conjugate is given by

$$I_{n,N,\tilde{X}\tilde{Y}}(-\omega_k) \equiv I_{n,N,\tilde{X}\tilde{Y}}^*(\omega_k), \quad 0 \leq \omega_k \leq \pi$$

and may be used to recover the cross periodogram at negative frequencies.

### Example

Consider the Robinson Multichannel Time Series Data (Robinson 1967, page 204) where  $X$  is the Wölfer sunspot number and  $Y$  is the northern light activity for the time period from 1770 through 1869. Application of routine `CPFFT` to these data produces the following results. Note that `CPFFT` sets `CPM` (1, 2) to the missing value code via routine `AMACH` (page 1334). The printing of `CPM` (1, 2) depends on the computer.

```

C
INTEGER      IPRINT, LDCPM, LDRDAT, NDRDAT, NOBS, NPAD
PARAMETER   (IPRINT=0, LDRDAT=100, NDRDAT=4, NOBS=100,
&           NPAD=NOBS-1, LDCPM=(NOBS+NPAD)/2+1)

INTEGER      IFSCAL, IPVER, NRCOL, NRROW
REAL         CPM(LDCPM,10), FLOAT, RDATA(LDRDAT,NDRDAT), SSUM,
&           X(NOBS), XCNTR, Y(NOBS), YCNTR
CHARACTER    CLABEL1(6)*9, CLABEL2(6)*9, FMT*7, RLABEL(1)*6,
&           TITLE*41

```

```

INTRINSIC  FLOAT
EXTERNAL  CPFFT, GDATA, SSUM, WRRRL
C
EQUIVALENCE (X(1), RDATA(1,2)), (Y(1), RDATA(1,3))
C
DATA TITLE/'Results of the Cross Periodogram Analysis'/
DATA FMT/'(F10.3)'/
DATA CLABEL1/'k+1', 'w(k)', 'p(k)', 'IX(w(k))', 'AX(w(k))',
& 'BX(w(k))'/
DATA CLABEL2/'k+1', 'IY(w(k))', 'AY(w(k))', 'BY(w(k))',
& 'Real IXY', 'Imag. IXY'/
DATA RLABEL/'NUMBER'/
C
C                               Robinson Data
CALL GDATA (8, 0, NRROW, NRCOL, RDATA, LDRDAT, NDRDAT)
C                               Center on arithmetic means
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
YCNTR = SSUM(NOBS,Y,1)/FLOAT(NOBS)
C                               Frequency in radians per unit time
IFSCAL = 0
C                               Modified periodogram version
IPVER = 1
C                               Compute the cross periodogram
CALL CPFFT (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL,
& IPVER, CPM, LDCPM)
C
C                               Print results (First 10 rows)
CALL WRRRL (TITLE, 10, 5, CPM, LDCPM, 0, FMT, RLABEL, CLABEL1)
CALL WRRRL ('%/', 10, 5, CPM(1,6), LDCPM, 0, FMT, RLABEL,
& CLABEL2)
C
END

```

### Output

```

Results of the Cross Periodogram Analysis
k+1      w(k)      p(k)      IX(w(k))  AX(w(k))  BX(w(k))
1         0.000      NaN        0.000     0.000     0.000
2         0.032    199.000    184.159     3.742    -13.044
3         0.063    99.500    1364.408    35.457    -10.354
4         0.095    66.333    2433.933    29.411    39.610
5         0.126    49.750    1351.002    -21.749    29.631
6         0.158    39.800    140.421    -11.716    -1.773
7         0.189    33.167    44.117     -4.671     4.722
8         0.221    28.429    121.186    -11.003    -0.343
9         0.253    24.875    176.275     -4.782    -12.386
10        0.284    22.111    144.867    10.038    -6.642

k+1      IY(w(k))  AY(w(k))  BY(w(k))  Real IXY  Imag. IXY
1         0.000     0.000     0.000     0.000     0.000
2        1689.212   -37.480   -16.866     79.776   -552.014
3        4113.003    41.232   -49.122    1970.577  -1314.779
4        3255.785    44.214    36.068    2729.031   -690.474
5        1757.663    -8.162    41.122    1396.006   -652.513
6        1002.050   -30.107    9.778     335.410   -167.954
7         62.360    -6.825    3.972     50.636    13.678
8        1481.396   -38.096    5.487     417.288   -73.451
9        1274.161   -17.176   -31.291    469.704   -63.095
10        488.479   -12.442   -18.267    -3.570   -265.992

```

---

## CSSWD/DCSSWD (Single/Double precision)

Estimate the nonnormalized cross-spectral density of two stationary time series using a spectral window given the time series data.

### Usage

```
CALL CSSWD (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL,  
           NF, F, TINT, ISWVER, NM, M, CPM, LDCEM, CSM,  
           LDCSM)
```

### Arguments

**NOBS** — Number of observations in each stationary time series  $X$  and  $Y$ . (Input)  
NOBS must be greater than or equal to two.

**X** — Vector of length NOBS containing the first stationary time series. (Input)

**Y** — Vector of length NOBS containing the second stationary time series. (Input)

**IPRINT** — Printing option. (Input)

#### IPRINT Action

0 No printing is performed.

1 Print the cross periodogram and cross-spectral density estimate based on a specified version of a spectral window for a given set of spectral window parameters.

**XCNTR** — Constant used to center the time series  $X$ . (Input)

**YCNTR** — Constant used to center the time series  $Y$ . (Input)

**NPAD** — Number of zeroes used to pad each centered time series. (Input)  
NPAD must be greater than or equal to zero. The length of each centered and padded time series is  $N = \text{NOBS} + \text{NPAD}$ .

**IFSCAL** — Option for frequency scale. (Input)

#### IFSCAL Action

0 Frequency in radians per unit time.

1 Frequency in cycles per unit time.

**NF** — Number of frequencies at which to evaluate the cross-spectral density estimate. (Input)

**F** — Vector of length NF containing the frequencies at which to evaluate the cross-spectral density estimate. (Input)

The units of  $F$  correspond to the scale specified by **IFSCAL**. The elements of  $F$  must be in the range  $(-\pi/\text{TINT}, \pi/\text{TINT})$ , inclusive, for **IFSCAL** = 0 and  $(-1/(2 * \text{TINT}), 1/(2 * \text{TINT}))$ , inclusive, for **IFSCAL** = 1.

**TINT** — Time interval at which the series are sampled. (Input)  
 For a discrete parameter process, usually  $TINT = 1$ . For a continuous parameter process,  $TINT > 0$ .  $TINT$  is used to adjust the cross-spectral density estimate.

**ISWVER** — Option for version of the spectral window. (Input)

**ISWVER Action**

- 1 Modified Bartlett
- 2 Daniell
- 3 Tukey-Hamming
- 4 Tukey-Hanning
- 5 Parzen
- 6 Bartlett-Priestley

Refer to the “Algorithm” section for further details.

**NM** — Number of spectral window parameters  $M$  used to compute the cross-spectral density estimate for a given spectral window version. (Input)  
 $NM$  must be greater than or equal to one.

**M** — Vector of length  $NM$  containing the values of the spectral window parameter  $M$ . (Input)

For the Parzen spectral window ( $ISWVER = 5$ ), all values of the spectral window parameters  $M$  must be even.

**CPM** —  $(\lfloor N/2 \rfloor + 1)$  by 10 matrix containing a summarization of the cross periodogram analysis. (Output)

For  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ , the  $(k + 1)$ -st element of the  $j$ -th column of  $CPM$  is defined as

**Col. Description**

- 1 Frequency,  $\omega_k$  where  $\omega_k = 2\pi k/N$  for  $IFSCAL = 0$  or  $\omega_k = k/N$  for  $IFSCAL = 1$ .
- 2 Period,  $p_k$  where  $p_k = 2\pi/\omega_k$  for  $IFSCAL = 0$  and  $p_k = 1/\omega_k$  for  $IFSCAL = 1$ . If  $\omega_k = 0$ ,  $p_k$  is set to missing.
- 3 X periodogram ordinate,  $I_X(\omega_k)$
- 4 X cosine transformation coefficient,  $A_X(\omega_k)$
- 5 X sine transformation coefficient,  $B_X(\omega_k)$
- 6 Y periodogram ordinate,  $I_Y(\omega_k)$
- 7 Y cosine transformation coefficient,  $A_Y(\omega_k)$
- 8 Y sine transformation coefficient,  $B_Y(\omega_k)$
- 9 Real part of the XY cross periodogram ordinate  $I_{XY}(\omega_k)$ .
- 10 Imaginary part of the XY cross periodogram ordinate  $I_{XY}(\omega_k)$ .

Note  $N = NOBS + NPAD$ .

**LDCPM** — Leading dimension of  $CPM$  exactly as specified in the dimension statement of the calling program. (Input)

$LDCPM$  must be greater than or equal to  $\lfloor N/2 \rfloor + 1$ .

**CSM** —  $NF$  by  $(NM * 7 + 2)$  matrix containing a summarization of the cross-spectral analysis. (Output)

The  $k$ -th element of the  $j$ -th column of **CSM** is defined as

Col.	Description
1	Frequency, $F(k)$ .
2	Period, $p_k$ where $p_k = 2\pi/F(k)$ for $IFSCAL = 0$ and $p_k = 1/F(k)$ for $IFSCAL = 1$ . If $F(k) = 0$ , $p_k$ is set to missing.
3	X spectral density estimate at $F(k)$ using the spectral window parameter $M(1)$ .
4	Y spectral density estimate at $F(k)$ using the spectral window parameter $M(1)$ .
5	Cospectrum estimate at $F(k)$ using the spectral window parameter $M(1)$ .
6	Quadrature spectrum estimate at $F(k)$ using the spectral window parameter $M(1)$ .
7	Cross-amplitude spectrum estimate at $F(k)$ using the spectral window parameter $M(1)$ .
8	Phase spectrum estimate at $F(k)$ using the spectral window parameter $M(1)$ .
9	Coherence estimate at $F(k)$ using the spectral window parameter $M(1)$ .
.	.
.	.
.	.
	$NM * 7 + 2$ Coherence estimate at $F(k)$ using the spectral window parameter $M(NM)$ .

where  $k = 1, \dots, NF$ .

**LDCSM** — Leading dimension of **CSM** exactly as specified in the dimension statement of the calling program. (Input)

**LDCSM** must be greater than or equal to  $NF$ .

### Comments

- Automatic workspace storage is

CSSWD  $10N + 15$  units, or  
DCSSWD  $20N + 30$  units.

Workspace may be explicitly provided, if desired, by use of **C2SWD/DC2SWD**. The reference is

CALL C2SWD (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD,  
IFSCAL, NF, F, TINT, ISWVER, NM, M, CPM,  
LDCPM, CSM, LDCSM, CX, COEF, WFFTC, CPY)

The additional arguments are as follows:

**CX** — Complex work vector of length  $N$ . (Output)

**COEF** — Complex work vector of length  $N$ . (Output)



**WFFTC** — Vector of length  $4N + 15$ .

**CPY** — Vector of length  $2N$ .

2. The centered and padded time series are defined by

$$CX(j) = X(j)XCNTR \quad \text{for } j = 1, \dots, \text{NOBS}$$

$$CX(j) = 0 \quad \text{for } j = \text{NOBS} + 1, \dots, N$$

and

$$CY(j) = Y(j)YCNTR \quad \text{for } j = 1, \dots, \text{NOBS}$$

$$CY(j) = 0 \quad \text{for } j = \text{NOBS} + 1, \dots, N$$

where  $N = \text{NOBS} + \text{NPAD}$ .

3. The normalized cross-spectral density estimate is obtained by dividing the nonnormalized cross-spectral density estimate in matrix CSM by the product of the estimated standard deviation of X and the estimated standard deviation of Y.

### Algorithm

Routine CSSWD estimates the nonnormalized cross-spectral density function of two jointly stationary time series using a spectral window given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  and  $\{Y_t\}$  for  $t = 1, 2, \dots, n$ .

Let

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

represent the centered and padded data where  $N = \text{NOBS} + \text{NPAD}$ ,

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu}_X & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_X = XCNTR$$

is determined by

$$\hat{\mu}_X = \begin{cases} \mu_X & \mu_X \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t & \mu_X \text{ unknown} \end{cases}$$

Similarly, let

$$\{\tilde{Y}_t\} \text{ for } t = 1, \dots, N$$

represent the centered and padded data where

$$\tilde{Y}_t = \begin{cases} Y_t - \hat{\mu}_Y & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_Y = \text{YCNTR}$$

is determined by

$$\hat{\mu}_Y = \begin{cases} \mu_Y & \mu_Y \text{ known} \\ \frac{1}{n} \sum_{t=1}^n Y_t & \mu_Y \text{ unknown} \end{cases}$$

The modified periodogram of

$$\{\tilde{X}_t\} \text{ for } t = 1, \dots, N$$

is estimated by

$$I_{n,N,\tilde{X}}(\omega_k) = A_{\tilde{X}}^2(\omega_k) + B_{\tilde{X}}^2(\omega_k)$$

where

$$A_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \cos(\omega_k t)$$

and

$$B_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \sin(\omega_k t)$$

represent the

$$\tilde{X}_t$$

cosine and sine transforms, respectively, and  $K$  is the scale factor equal to  $1/(2\pi n)$ . The modified periodogram of

$$\{\tilde{Y}_t\} \text{ for } t = 1, \dots, N$$

is estimated by

$$I_{n,N,\tilde{Y}}(\omega_k) = A_{\tilde{Y}}^2(\omega_k) + B_{\tilde{Y}}^2(\omega_k)$$

where

$$A_{\tilde{Y}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{Y}_t \cos(\omega_k t)$$

and

$$B_{\tilde{Y}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{Y}_t \sin(\omega_k t)$$

represent the

$$\tilde{Y}_t$$

cosine and sine transforms, respectively. Since the periodogram is an even function of the frequency, it is sufficient to estimate the periodogram at the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

The routine `PEFFT` (page 723) is used to compute the modified periodograms of both

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

The computational formula for the cross periodogram is given by

$$I_{n,N,\tilde{X}\tilde{Y}}(\omega_k) = \Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} + i\Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\}$$

where

$$\Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} = A_{\tilde{X}}(\omega_k)A_{\tilde{Y}}(\omega_k) + B_{\tilde{X}}(\omega_k)B_{\tilde{Y}}(\omega_k)$$

and

$$\Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} = A_{\tilde{X}}(\omega_k)B_{\tilde{Y}}(\omega_k) - B_{\tilde{X}}(\omega_k)A_{\tilde{Y}}(\omega_k)$$

The routine `CEFFT` (page 750) is used to compute the modified cross periodogram between

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

The nonnormalized spectral density of  $X_t$  is estimated by

$$\hat{h}_X(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} I_{n,N,\tilde{X}}(\omega_k) W_n(\omega - \omega_k)$$

and the nonnormalized spectral density of  $Y_t$  is estimated by

$$\hat{h}_Y(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} I_{n,N,\tilde{Y}}(\omega_k) W_n(\omega - \omega_k)$$

where the spectral window  $W_n(\theta)$  is specified by argument `ISWVER`. The following spectral windows  $W_n(\theta)$  are available.

#### Modified Bartlett

$$W_n(\theta) = \frac{1}{2\pi M} \left\{ \frac{\sin(M\theta/2)}{\sin(\theta/2)} \right\}^2 = F_M(\theta)$$

where  $F_M(\theta)$  corresponds to the Fejér kernel of order  $M$ .

**Daniell**

$$W_n(\theta) = \begin{cases} M / 2\pi & -\pi / M \leq \theta \leq \pi / M \\ 0 & \text{otherwise} \end{cases}$$

**Tukey**

$$W_n(\theta) = aD_M\left(\theta - \frac{\pi}{M}\right) + (1 - 2a)D_M(\theta) + aD_M\left(\theta + \frac{\pi}{M}\right) \quad 0 < a \leq 0.25$$

where  $D_M(\theta)$  represents the Dirichlet kernel. The Tukey-Hamming window is obtained when  $a = 0.23$ , and the Tukey-Hanning window is obtained when  $a = 0.25$ .

**Parzen**

$$W_n(\theta) = \frac{6\pi}{M} [F_{M/2}(\theta)]^2 \left\{ 1 - \frac{2}{3} \sin^2(\theta / 2) \right\}$$

where  $M$  is even. If  $M$  is odd, then  $M + 1$  is used instead of  $M$  in the above formula.

**Bartlett-Priestley**

$$W_n(\theta) = \begin{cases} \frac{3M}{4\pi} \left\{ 1 - \left( \frac{M\theta}{\pi} \right)^2 \right\} & |\theta| \leq \pi / M \\ 0 & |\theta| > \pi / M \end{cases}$$

The argument `NM` specifies the number of window parameters  $M$  and, hence, corresponds to the number of spectral density estimates to be computed for a given spectral window. Note that the same spectral window  $W_n(\theta)$  and set of parameters  $M$  are used to obtain both

$$\hat{h}_X(\omega) \text{ and } \hat{h}_Y(\omega)$$

The above spectral density formulas assume the data  $\{X_t\}$  and  $\{Y_t\}$  correspond to a realization of a bivariate discrete-parameter stationary process observed consecutively in time. In this case, the observations are equally spaced in time with interval  $\Delta t = \text{TINT}$  equal to one. However, if the data correspond to a realization of a bivariate continuous-parameter stationary process recorded at equal time intervals, then the spectral density estimates must be adjusted for the effect of aliasing. In general, the estimate of  $h_X(\omega)$  is given by

$$\hat{h}_X(\omega) = \Delta t \hat{h}_X(\omega), \quad |\omega| \leq \pi / \Delta t$$

and the estimate of  $h_Y(\omega)$  is given by

$$\hat{h}_Y(\omega) = \Delta t \hat{h}_Y(\omega), \quad |\omega| \leq \pi / \Delta t$$

The nonnormalized spectral density is estimated over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale specified by the argument `IFSCAL` but are transformed to the scale of radians per unit time for computational purposes. The frequency  $\omega$  of the desired spectral estimate is assumed to be input in a form already adjusted for the time interval  $\Delta t$ .

The cross-spectral density function is complex-valued in general and may be written in the following form:

$$h_{XY}(\omega) = c_{XY}(\omega) - iq_{XY}(\omega)$$

The *cospectrum* is estimated by

$$\hat{c}_{XY}(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} \Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} W_n(\omega - \omega_k)$$

and the *quadrature spectrum* is estimated by

$$\hat{q}_{XY}(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} \Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} W_n(\omega - \omega_k)$$

Note that the same spectral window  $W_n(\theta)$  and window parameter  $M$  used to derive

$$\hat{h}_X(\omega) \text{ and } \hat{h}_Y(\omega)$$

are also used to compute

$$\hat{h}_{XY}(\omega)$$

The nonnormalized cross-spectral density estimate is computed over the same set of frequencies as the nonnormalized spectral density estimates with a similar adjustment for  $\Delta t$ .

An equivalent representation of  $h_{XY}(\omega)$  is the *polar form* defined by

$$h_{XY}(\omega) = \alpha_{XY}(\omega) e^{i\phi_{XY}(\omega)}$$

The *cross-amplitude spectrum* is estimated by

$$\hat{\alpha}_{XY}(\omega) = \{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)\}^{1/2}$$

and the *phase spectrum* is estimated by

$$\hat{\phi}_{XY}(\omega) = \tan^{-1}\{-\hat{q}_{XY}(\omega) / \hat{c}_{XY}(\omega)\}$$

Finally, the *coherency spectrum* is estimated by

$$|\hat{w}_{XY}(\omega)| = \left\{ \frac{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)}{\hat{h}_X(\omega)\hat{h}_Y(\omega)} \right\}^{1/2}$$

The *coherence* or squared coherency is output.

### Example

Consider the Robinson Multichannel Time Series Data (Robinson 1967, page 204) where  $X$  is the Wölfer sunspot number and  $Y$  is the northern light activity for the time period from 1770 through 1869. Application of routine CSSWD to these data produces the following results:

```

INTEGER      IPRINT, LDCPM, LDCSM, LDRDAT, N, NDRDAT, NF, NM,
&            NOBS, NPAD
PARAMETER    (IPRINT=0, LDRDAT=100, NDRDAT=4, NF=10, NM=2,
&            NOBS=100, LDCSM=NF, NPAD=NOBS-1, N=NOBS+NPAD,
&            LDCPM=N/2+1)
C
INTEGER      I, IFSCAL, ISWVER, J, JPT, M(NM), NOUT, NRCOL, NRROW
REAL         ASIN, CPM(LDCPM,10), CSM(LDCSM,NM*7+2), F(NF), FLOAT,
&            PI, RDATA(LDRDAT,NDRDAT), SSUM, TINT, X(NOBS), XCNTR,
&            Y(NOBS), YCNTR
CHARACTER    CLABEL1(3)*9, CLABEL2(6)*16, FMT*7, RLABEL(1)*6,
&            TITLE*80
INTRINSIC    ASIN, FLOAT
EXTERNAL     CSSWD, GDATA, SSUM, UMACH, WRRRL
C
EQUIVALENCE (X(1), RDATA(1,2)), (Y(1), RDATA(1,3))
C
DATA FMT/'(F10.4)'/
DATA CLABEL1/' k', 'Frequency', 'Period'/
DATA CLABEL2/'%/ k', '%/Cospectrum', '%/Quadrature',
&            'Cross%/Amplitude', '%/Phase', '%/Coherence'/
DATA RLABEL/'NUMBER'/
C
                                Initialization
CALL UMACH (2, NOUT)
PI = 2.0*ASIN(1.0)
DO 10 I=1, NF
    F(I) = PI*FLOAT(I)/FLOAT(NF)
10 CONTINUE
C
                                Robinson Data
CALL GDATA (8, 0, NRROW, NRCOL, RDATA, LDRDAT, NDRDAT)
C
                                Center on arithmetic means
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
YCNTR = SSUM(NOBS,Y,1)/FLOAT(NOBS)
C
                                Frequency in radians per unit time
IFSCAL = 0
C
                                Spectral window parameters
M(1) = 10
M(2) = 30
C
                                Time interval for discrete data
TINT = 1.0
C
                                Compute cross-spectral density
                                using the Parzen window
C
ISWVER = 5
CALL CSSWD (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL, NF,

```

```

&          F, TINT, ISWVER, NM, M, CPM, LDCPM, CSM, LDCSM)
C          Print results
TITLE = 'Cross-Spectral Analysis Using Parzen Window'
CALL WRRRL (TITLE, NF, 2, CSM, LDCSM, 0, FMT, RLABEL, CLABEL1)
DO 20 J=1, NM
  JPT = 7*(J-1) + 5
  TITLE = '%/Results of the Cross-Spectral Analysis With '//
&         'Spectral Window Parameter M = '
  WRITE (TITLE(77:78), '(I2)') M(J)
  CALL WRRRL (TITLE, NF, 5, CSM(1,JPT), LDCSM, 0, FMT, RLABEL,
&            CLABEL2)
20 CONTINUE
C
END

```

### Output

Cross-Spectral Analysis Using Parzen Window

k	Frequency	Period
1	0.3142	20.0000
2	0.6283	10.0000
3	0.9425	6.6667
4	1.2566	5.0000
5	1.5708	4.0000
6	1.8850	3.3333
7	2.1991	2.8571
8	2.5133	2.5000
9	2.8274	2.2222
10	3.1416	2.0000

Results of the Cross-Spectral Analysis With Spectral Window Parameter M = 10

Cross					
k	Cospectrum	Quadrature	Amplitude	Phase	Coherence
1	463.5888	-65.9763	468.2600	0.1414	0.2570
2	286.5450	-75.0209	296.2029	0.2561	0.1710
3	150.1073	-57.8263	160.8604	0.3677	0.1438
4	52.9840	-32.3642	62.0866	0.5483	0.0998
5	21.5435	-15.0888	26.3020	0.6110	0.0794
6	21.4228	-9.8188	23.5658	0.4298	0.1716
7	15.7005	-5.3704	16.5936	0.3296	0.2112
8	8.0118	-1.8887	8.2314	0.2315	0.1272
9	2.7682	0.2007	2.7754	-0.0724	0.0446
10	0.5777	0.1008	0.5864	-0.1727	0.0091

Results of the Cross-Spectral Analysis With Spectral Window Parameter M = 30

Cross					
k	Cospectrum	Quadrature	Amplitude	Phase	Coherence
1	169.7542	-193.4384	257.3615	0.8505	0.1620
2	452.6187	32.3813	453.7755	-0.0714	0.2213
3	94.5221	-90.8159	131.0800	0.7654	0.2629
4	-0.2096	-6.1127	6.1163	1.6051	0.0019
5	27.4711	-22.1946	35.3166	0.6796	0.2492
6	29.1329	-4.0128	29.4080	0.1369	0.3170
7	11.2058	-9.3403	14.5881	0.6948	0.2594
8	8.0017	0.8813	8.0501	-0.1097	0.1928
9	-0.4199	2.2893	2.3275	-1.7522	0.0468
10	0.5570	-1.0767	1.2123	1.0934	0.0678

---

## CSSWP/DCSSWP (Single/Double precision)

Estimate the nonnormalized cross-spectral density of two stationary time series using a spectral window given the spectral densities and cross periodogram.

### Usage

```
CALL CSSWP (N, SX, SY, CPREAL, CPIMAG, NF, F, ISWVER, M,  
           COSPEC, QUADRA, CRAMPL, PHASE, COHERE)
```

### Arguments

**N** — Number of observations in each of the appropriately centered and padded time series X and Y. (Input)

N must be greater than or equal to two.

**SX** — Vector of length NF containing the estimate of the spectral density of the first time series X. (Input)

**SY** — Vector of length NF containing the estimate of the spectral density of the second time series Y. (Input)

**CPREAL** — Vector of length  $\lfloor N/2 \rfloor + 1$  containing the real part of the cross periodogram between X and Y. (Input)

The real part of the cross periodogram evaluated at (angular) frequency  $w_k = 2\pi k/N$  is given by  $CPREAL(k + 1)$ ,  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ .

**CPIMAG** — Vector of length  $\lfloor N/2 \rfloor + 1$  containing the imaginary part of the cross periodogram between X and Y. (Input)

The imaginary part of the cross periodogram evaluated at (angular) frequency  $w_k = 2\pi k/N$  is given by  $CPIMAG(k + 1)$ ,  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ .

**NF** — Number of (angular) frequencies. (Input) NF must be greater than or equal to one.

**F** — Vector of length NF containing the (angular) frequencies at which the spectral and cross-spectral densities are estimated. (Input)

The elements of F must be in the range  $(-\pi, \pi)$  inclusive.

**ISWVER** — Option for version of the spectral window. (Input)

#### ISWVER Action

- |   |                    |
|---|--------------------|
| 1 | Modified Bartlett  |
| 2 | Daniell            |
| 3 | Tukey-Hamming      |
| 4 | Tukey-Hanning      |
| 5 | Parzen             |
| 6 | Bartlett-Priestley |

Refer to the “Algorithm” section for further details.



***M*** — Spectral window parameter. (Input)

*M* must be greater than or equal to one and less than *N*. For the Parzen spectral window (*ISWVER* = 5), the spectral window parameter *M* must be even.

***COSPEC*** — Vector of length *NF* containing the estimate of the cospectrum. (Output)

***QUADRA*** — Vector of length *NF* containing the estimate of the quadrature spectrum. (Output)

***CRAMPL*** — Vector of length *NF* containing the estimate of the cross-amplitude spectrum. (Output)

***PHASE*** — Vector of length *NF* containing the estimate of the phase spectrum. (Output)

***COHERE*** — Vector of length *NF* containing the estimate of the coherence or squared coherency. (Output)

### Comments

1. The periodograms of *X* and *Y* and cross periodogram between *X* and *Y* may be computed using the routine *CPFFT* (page 750). The spectral densities of *X* and *Y* may then be estimated using any of the routines *SSWD* (page 729), *SWED* (page 741), *SSWP* (page 736), or *SWEP* (page 747). Thus, different window types and/or weight sequences may be used to estimate the spectral and cross-spectral densities given either the series or their periodograms. Note that use of the modified periodograms and modified cross periodogram ensures that the scale of the spectral and cross-spectral densities and their estimates is equivalent.
2. The time sampling interval, *TINT*, is assumed to be equal to one. This assumption is appropriate for discrete parameter processes. The adjustment for continuous parameter processes (*TINT* > 0.0) involves multiplication of the frequency vector *F* by 1/*TINT* and multiplication of the spectral and cross-spectral density estimates by *TINT*.
3. To convert the frequency scale from radians per unit time to cycles per unit time, multiply *F* by 1/(2 $\pi$ ).

### Algorithm

Routine *CSSWP* estimates the nonnormalized cross-spectral density function of two jointly stationary time series using a spectral window given the modified cross-periodogram and spectral densities of the appropriately centered and padded data

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

for  $t = 1, \dots, N$ .

The routine CPFFFT (page 750) may be used to compute the modified periodograms

$$I_{n,N,\tilde{X}}(\omega_k) \text{ and } I_{n,N,\tilde{Y}}(\omega_k)$$

and cross periodogram

$$I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)$$

over the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

(Here,  $\lfloor a \rfloor$  means the greatest integer less than or equal to  $a$ .) Either routine SSWP (page 736) or routine SWEP (page 747) may be applied to the periodograms to obtain nonnormalized spectral density estimates

$$\hat{h}_X(\omega) \text{ and } \hat{h}_Y(\omega)$$

over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale of radians per unit time. The time sampling interval  $\Delta t$  is assumed to be equal to one. Note that the spectral window or weight sequence used to compute

$$\hat{h}_X(\omega)$$

may differ from that used to compute

$$\hat{h}_Y(\omega)$$

The cross-spectral density function is complex-valued in general and may be written as

$$h_{XY}(\omega) = c_{XY}(\omega) - iq_{XY}(\omega)$$

The *cospectrum* is estimated by

$$\hat{c}_{XY}(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} \Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} W_n(\omega - \omega_k)$$

and the *quadrature spectrum* is estimated by

$$\hat{q}_{XY}(\omega) = \frac{2\pi}{N} \sum_{k=-\lfloor N/2 \rfloor}^{\lfloor N/2 \rfloor} \Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} W_n(\omega - \omega_k)$$

where the spectral window  $W_n(\theta)$  is specified by argument ISWVER. The following spectral windows  $W_n(\theta)$  are available.

### Modified Bartlett

$$W_n(\theta) = \frac{1}{2\pi M} \left\{ \frac{\sin(M\theta/2)}{\sin(\theta/2)} \right\}^2 = F_M(\theta)$$

where  $F_M(\theta)$  corresponds to the Fejér kernel of order  $M$ .

### Daniell

$$W_n(\theta) = \begin{cases} M/2\pi & -\pi/M \leq \theta \leq \pi/M \\ 0 & \text{otherwise} \end{cases}$$

### Tukey

$$W_n(\theta) = aD_M\left(\theta - \frac{\pi}{M}\right) + (1-2a)D_M(\theta) + aD_M\left(\theta + \frac{\pi}{M}\right), \quad 0 < a \leq 0.25$$

where  $D_M(\theta)$  represents the Dirichlet kernel. The Tukey-Hamming window is obtained when  $a = 0.23$ , and the Tukey-Hanning window is obtained when  $a = 0.25$ .

### Parzen

$$W_n(\theta) = \frac{6\pi}{M} [F_{M/2}(\theta)]^2 \left\{ 1 - \frac{2}{3} \sin^2(\theta/2) \right\}$$

where  $M$  is even. If  $M$  is odd, then  $M + 1$  is used instead of  $M$  in the above formula.

### Bartlett-Priestley

$$W_n(\theta) = \begin{cases} \frac{3M}{4\pi} \left\{ 1 - \left( \frac{M\theta}{\pi} \right)^2 \right\} & |\theta| \leq \pi/M \\ 0 & |\theta| > \pi/M \end{cases}$$

Only one window parameter  $M$  may be specified so that only one estimate of  $\hat{h}_{XY}(\omega)$  is computed. The nonnormalized cross-spectral density estimate is computed over the same set of frequencies as the nonnormalized spectral density estimates discussed above. However, the particular spectral window used to compute

$$\hat{h}_{XY}(\omega)$$

need not correspond to either the spectral window or the weight sequence used to compute either

$$\hat{h}_X(\omega) \text{ or } \hat{h}_Y(\omega)$$

An equivalent representation of  $h_{XY}(\omega)$  is the *polar form* defined by

$$h_{XY}(\omega) = \alpha_{XY}(\omega)e^{i\phi_{XY}(\omega)}$$

The *cross-amplitude* spectrum is estimated by

$$\hat{\alpha}_{XY}(\omega) = \{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)\}^{1/2}$$

and the *phase spectrum* is estimated by

$$\hat{\phi}_{XY}(\omega) = \tan^{-1}\{-\hat{q}_{XY}(\omega) / \hat{c}_{XY}(\omega)\}$$

Finally, the *coherency spectrum* is estimated by

$$|\hat{w}_{XY}(\omega)| = \left\{ \frac{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)}{\hat{h}_X(\omega)\hat{h}_Y(\omega)} \right\}^{1/2}$$

The *coherence* or squared coherency is output.

### Example

Consider the Robinson Multichannel Time Series Data (Robinson 1967, page 204) where  $X$  is the Wölfer sunspot number and  $Y$  is the northern light activity for the years 1770 through 1869. Application of routine CSSWP to these data produces the following results.

```

INTEGER      IPRINT, LDCPM, LDCSM, LDRDAT, N, NDRDAT, NF, NM,
&            NOBS, NPAD
PARAMETER    (IPRINT=0, LDRDAT=100, NDRDAT=4, NF=10, NM=2,
&            NOBS=100, LDCSM=NF, NPAD=NOBS-1, N=NOBS+NPAD,
&            LDCPM=N/2+1)
C
INTEGER      I, IFSCAL, IPVER, ISWVER, J, JPT, JST, M(NM), NRCOL,
&            NRROW
REAL         ASIN, COHERE(NF), COSPEC(NF), CPIMAG(LDCPM),
&            CPM(LDCPM,10), CPREAL(LDCPM), CRAMPL(NF),
&            CSM(LDCSM,7*NM+2), F(NF), FLOAT, P(NF), PHASE(NF),
&            PI, PX(LDCPM), PY(LDCPM), QUADRA(NF),
&            RDATA(LDRDAT,NDRDAT), SSUM, SX(NF), SY(NF), X(NOBS),
&            XCNTR, Y(NOBS), YCNTR
CHARACTER    CLABEL1(3)*9, CLABEL2(6)*16, FMT*8, RLABEL(1)*6,
&            TITLE*80
INTRINSIC    ASIN, FLOAT
EXTERNAL     CPFFT, CSSWP, GDATA, SCOPY, SSUM, SSWP, WRRRL
C
EQUIVALENCE (X(1), RDATA(1,2)), (Y(1), RDATA(1,3))
EQUIVALENCE (PX(1), CPM(1,3)), (PY(1), CPM(1,6))
EQUIVALENCE (CPREAL(1), CPM(1,9)), (CPIMAG(1), CPM(1,10))
EQUIVALENCE (CSM(1,1), F(1)), (CSM(1,2), P(1))
C
DATA FMT/'(F12.4)'/
DATA CLABEL1/' k', 'Frequency', 'Period'/
DATA CLABEL2/'%/ k', '%/Cospectrum', '%/Quadrature',
&            'Cross%/Amplitude', '%/Phase', '%/Coherence'/
DATA RLABEL/'NUMBER'/

```

```

C                                     Initialization
    PI = 2.0*ASIN(1.0)
    DO 10 I=1, NF
        F(I) = PI*FLOAT(I)/FLOAT(NF)
        P(I) = 2.0*FLOAT(NF)/FLOAT(I)
10 CONTINUE
C                                     Robinson Data
    CALL GDATA (8, 0, NRROW, NRCOL, RDATA, LDRDAT, NDRDAT)
C                                     Center on arithmetic means
    XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
    YCNTR = SSUM(NOBS,Y,1)/FLOAT(NOBS)
C                                     Frequency in radians per unit time
    IFSCAL = 0
C                                     Modified periodogram version
    IPVER = 1
C                                     Compute cross periodogram
    CALL CPFFT (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL,
&              IPVER, CPM, LDCPM)
C                                     Spectral window parameters
    M(1) = 10
    M(2) = 30
C                                     Compute cross-spectral density
C                                     using the Parzen window
C
C                                     Print frequency and period
    TITLE = 'Cross-Spectral Analysis Using Parzen Window'
    CALL WRRRL (TITLE, NF, 2, CSM, LDCSM, 0, FMT, RLABEL, CLABEL1)
    ISWVER = 5
    DO 20 J=1, NM
C                                     Estimate the spectral densities
        CALL SSWP (N, PX, NF, F, ISWVER, M(J), SX)
        CALL SSWP (N, PY, NF, F, ISWVER, M(J), SY)
C                                     Estimate the cross-spectral density
&      CALL CSSWP (N, SX, SY, CPREAL, CPIMAG, NF, F, ISWVER, M(J),
        COSPEC, QUADRA, CRAMPL, PHASE, COHERE)
C                                     Copy results to output matrices
        JPT = 7*(J-1) + 2
        JST = 7*(J-1) + 5
        CALL SCOPY (NF, SX, 1, CSM(1,JPT+1), 1)
        CALL SCOPY (NF, SY, 1, CSM(1,JPT+2), 1)
        CALL SCOPY (NF, COSPEC, 1, CSM(1,JPT+3), 1)
        CALL SCOPY (NF, QUADRA, 1, CSM(1,JPT+4), 1)
        CALL SCOPY (NF, CRAMPL, 1, CSM(1,JPT+5), 1)
        CALL SCOPY (NF, PHASE, 1, CSM(1,JPT+6), 1)
        CALL SCOPY (NF, COHERE, 1, CSM(1,JPT+7), 1)
C                                     Print results
        TITLE = '%/Results of the Cross-Spectral Analysis With '//
&              'Spectral Window Parameter M = '
        WRITE (TITLE(77:78), '(I2)') M(J)
        CALL WRRRL (TITLE, NF, 5, CSM(1,JST), LDCSM, 0, FMT, RLABEL,
&              CLABEL2)
20 CONTINUE
C
    END

```

### Output

```

Cross-Spectral Analysis Using Parzen Window
      k      Frequency      Period

```

1	0.3142	20.0000
2	0.6283	10.0000
3	0.9425	6.6667
4	1.2566	5.0000
5	1.5708	4.0000
6	1.8850	3.3333
7	2.1991	2.8571
8	2.5133	2.5000
9	2.8274	2.2222
10	3.1416	2.0000

Results of the Cross-Spectral Analysis With Spectral Window Parameter M = 10

Cross					
k	Cospectrum	Quadrature	Amplitude	Phase	Coherence
1	463.5888	-65.9763	468.2600	0.1414	0.2570
2	286.5450	-75.0209	296.2029	0.2561	0.1710
3	150.1073	-57.8263	160.8604	0.3677	0.1438
4	52.9840	-32.3642	62.0866	0.5483	0.0998
5	21.5435	-15.0888	26.3020	0.6110	0.0794
6	21.4228	-9.8188	23.5658	0.4298	0.1716
7	15.7005	-5.3704	16.5936	0.3296	0.2112
8	8.0118	-1.8887	8.2314	0.2315	0.1272
9	2.7682	0.2007	2.7754	-0.0724	0.0446
10	0.5777	0.1008	0.5864	-0.1727	0.0091

Results of the Cross-Spectral Analysis With Spectral Window Parameter M = 30

Cross					
k	Cospectrum	Quadrature	Amplitude	Phase	Coherence
1	169.7542	-193.4384	257.3615	0.8505	0.1620
2	452.6187	32.3813	453.7755	-0.0714	0.2213
3	94.5221	-90.8159	131.0800	0.7654	0.2629
4	-0.2096	-6.1127	6.1163	1.6051	0.0019
5	27.4711	-22.1946	35.3166	0.6796	0.2492
6	29.1329	-4.0128	29.4080	0.1369	0.3170
7	11.2058	-9.3403	14.5881	0.6948	0.2594
8	8.0017	0.8813	8.0501	-0.1097	0.1928
9	-0.4199	2.2893	2.3275	-1.7522	0.0468
10	0.5570	-1.0767	1.2123	1.0934	0.0678

---

## CSWED/DCSWED (Single/Double precision)

Estimate the nonnormalized cross-spectral density of two stationary time series using a weighted cross periodogram given the time series data.

### Usage

CALL CSWED (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL, NF, F, TINT, NWT, WT, CPM, LDCPM, CSM, LDCSM)

### Arguments

**NOBS** — Number of observations in each stationary time series X and Y. (Input)  
NOBS must be greater than or equal to two.

$X$  — Vector of length NOBS containing the first stationary time series. (Input)

$Y$  — Vector of length NOBS containing the second stationary time series. (Input)

**IPRINT** — Printing option. (Input)

**IPRINT Action**

0 No printing is performed.

1 Print the periodogram, cosine and sine transformations of each centered and padded time series, the real and imaginary components of the cross periodogram, and the cross-spectral density estimate based on a specified weight sequence.

**XCNTR** — Constant used to center the time series  $x$ . (Input)

**YCNTR** — Constant used to center the time series  $y$ . (Input)

**NPAD** — Number of zeroes used to pad each centered time series. (Input)

NPAD must be greater than or equal to zero. The length of each centered and padded time series is  $N = \text{NOBS} + \text{NPAD}$ .

**IFSCAL** — Option for frequency scale. (Input)

**IFSCAL Action**

0 Frequency in radians per unit time.

1 Frequency in cycles per unit time.

**NF** — Number of frequencies at which to evaluate the cross-spectral density estimate. (Input)

$F$  — Vector of length NF containing the frequencies at which to evaluate the cross-spectral density estimate. (Input)

The units of  $F$  correspond to the scale specified by **IFSCAL**. The elements of  $F$  must be in the range  $(-\pi/\text{TINT}, \pi/\text{TINT})$  inclusive, for **IFSCAL** = 0 and  $(-1/(2 * \text{TINT}), 1/(2 * \text{TINT}))$  inclusive, for **IFSCAL** = 1.

**TINT** — Time interval at which the series are sampled. (Input)

For a discrete parameter process, usually **TINT** = 1.0. For a continuous parameter process, **TINT** > 0.0. **TINT** is used to adjust the cross-spectral density estimate.

**NWT** — Number of weights. (Input)

NWT must be greater than or equal to one.

$WT$  — Vector of length NWT containing the weights used to smooth the periodogram. (Input)

The actual weights are the values in  $WT$  normalized to sum to 1 with the current periodogram ordinate taking the middle weight for NWT odd or the weight to the right of the middle for NWT even.

**CPM** —  $(\lfloor N/2 \rfloor + 1)$  by 10 matrix containing a summarization of the cross periodogram analysis. (Output)

For  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ , the  $(k + 1)$ -st element of the  $j$ -th column of CPM is defined as

Col.	Description
1	Frequency, $\omega_k$ where $\omega_k = 2\pi k/N$ for IFSCAL = 0 or $\omega_k = k/N$ for IFSCAL = 1
2	Period, $p_k$ where $p_k = 2\pi/\omega_k$ for IFSCAL = 0 and $p_k = 1/\omega_k$ for IFSCAL = 1. If $\omega_k = 0$ , $p_k$ is set to missing.
3	X periodogram ordinate, $I_X(\omega_k)$
4	X cosine transformation coefficient, $A_X(\omega_k)$
5	X sine transformation coefficient, $B_X(\omega_k)$
6	Y periodogram ordinate, $I_Y(\omega_k)$
7	Y cosine transformation coefficient, $A_Y(\omega_k)$
8	Y sine transformation coefficient, $B_Y(\omega_k)$
9	Real part of the XY cross periodogram ordinate $I_{XY}(\omega_k)$ .
10	Imaginary part of the XY cross periodogram ordinate $I_{XY}(\omega_k)$ .

**LDCPM** — Leading dimension of CPM exactly as specified in the dimension statement of the calling program. (Input)  
LDCPM must be greater than or equal to  $\lfloor N/2 \rfloor + 1$ .

**CSM** — NF by 9 matrix containing a summarization of the cross-spectral analysis. (Output)

The  $k$ -th element of the  $j$ -th column of CSM is defined as

Col.	Description
1	Frequency, $F(k)$ .
2	Period, $p_k$ where $p_k = 2\pi/F(k)$ for IFSCAL = 0 and $p_k = 1/F(k)$ for IFSCAL = 1. If $F(k) = 0$ , $p_k$ is set to missing.
3	X spectral density estimate at $F(k)$ using the specified relative weights contained in WT.
4	Y spectral density estimate at $F(k)$ using the specified relative weights contained in WT.
5	Co-spectrum estimate at $F(k)$ using the specified relative weights contained in WT.
6	Quadrature spectrum estimate at $F(k)$ using the specified relative weights contained in WT.
7	Cross-amplitude spectrum estimate at $F(k)$ .
8	Phase spectrum estimate at $F(k)$ .
9	Coherence estimate at $F(k)$ .

where  $k = 1, \dots, NF$ .

**LDCSM** — Leading dimension of CSM exactly as specified in the dimension statement of the calling program. (Input)  
LDCSM must be greater than or equal to NF.



## Comments

1. Automatic workspace storage is

CSWED  $10N + 15$  units, or  
DCSWED  $20N + 30$  units.

Workspace may be explicitly provided, if desired, by use of C2WED/DC2WED. The reference is

```
CALL C2WED (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD,
            IFSCAL, NF, F, TINT, NWT, WT, CPM,
            LDPCM, CSM, LDCSM, CWK, COEFWK, WFFTC,
            CPY)
```

The additional arguments are as follows:

**CWK** — Complex work vector of length  $N$ . (Output)

**COEFWK** — Complex work vector of length  $N$ . (Output)

**WFFTC** — Vector of length  $4N + 15$ .

**CPY** — Vector of length  $2N$ .

2. The normalized cross-spectral density estimate is obtained by dividing the nonnormalized cross-spectral density estimate in matrix CSM by the product of the estimated standard deviation of  $X$  and the estimated standard deviation of  $Y$ .

## Algorithm

Routine CSWED estimates the nonnormalized cross-spectral density function of two jointly stationary time series using a fixed sequence of weights given a sample of  $n = \text{NOBS}$  observations  $\{X_t\}$  and  $\{Y_t\}$  for  $t = 1, 2, \dots, n$ . Let

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  represent the centered and padded data where  $N = \text{NOBS} + \text{NPAD}$ ,

$$\tilde{X}_t = \begin{cases} X_t - \hat{\mu}_X & t = 1, \dots, n \\ 0 & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_X = \text{XCNTR}$$

is determined by

$$\hat{\mu}_X = \begin{cases} \mu_X, & \mu_X \text{ known} \\ \frac{1}{n} \sum_{t=1}^n X_t, & \mu_X \text{ unknown} \end{cases}$$

Similarly, let

$$\{\tilde{Y}_t\}$$

for  $t = 1, \dots, N$  represent the centered and padded data where

$$\tilde{Y}_t = \begin{cases} Y_t - \hat{\mu}_Y, & t = 1, \dots, n \\ 0, & t = (n+1), \dots, N \end{cases}$$

and

$$\hat{\mu}_Y = \text{YCNTR}$$

is determined by

$$\hat{\mu}_Y = \begin{cases} \mu_Y, & \mu_Y \text{ known} \\ \frac{1}{n} \sum_{t=1}^n Y_t, & \mu_Y \text{ unknown} \end{cases}$$

The modified periodogram of

$$\{\tilde{X}_t\}$$

for  $t = 1, \dots, N$  is estimated by

$$I_{n,N,\tilde{X}}(\omega_k) = A_{\tilde{X}}^2(\omega_k) + B_{\tilde{X}}^2(\omega_k)$$

where

$$A_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \cos(\omega_k t)$$

and

$$B_{\tilde{X}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{X}_t \sin(\omega_k t)$$

represent the

$$\tilde{X}_t$$

cosine and sine transforms, respectively, and  $K$  is the scale factor equal to  $1/(2\pi n)$ . The modified periodogram of  $\{Y_t\}$  for  $t = 1, \dots, N$  is estimated by

$$I_{n,N,\tilde{Y}}(\omega_k) = A_{\tilde{Y}}^2(\omega_k) + B_{\tilde{Y}}^2(\omega_k)$$

where

$$A_{\tilde{Y}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{Y}_t \cos(\omega_k t)$$

and

$$B_{\tilde{Y}}(\omega_k) = K^{1/2} \sum_{t=1}^N \tilde{Y}_t \sin(\omega_k t)$$

represent the

$$\tilde{Y}_t$$

cosine and sine transforms, respectively. Since the periodogram is an even function of the frequency, it is sufficient to estimate the periodogram at the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

(Here,  $\lfloor a \rfloor$  means the greatest integer less than or equal to  $a$ ). The routine `PEFFT` (page 723) is used to compute the modified periodograms of both

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

The computational formula for the cross periodogram is given by

$$I_{n,N,\tilde{X}\tilde{Y}}(\omega_k) = \Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} + i\Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\}$$

where

$$\Re\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} = A_{\tilde{X}}(\omega_k)A_{\tilde{Y}}(\omega_k) + B_{\tilde{X}}(\omega_k)B_{\tilde{Y}}(\omega_k)$$

and

$$\Im\{I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)\} = A_{\tilde{X}}(\omega_k)B_{\tilde{Y}}(\omega_k) - B_{\tilde{X}}(\omega_k)A_{\tilde{Y}}(\omega_k)$$

The routine `CPFFT` (page 750) is used to compute the modified cross periodogram between

$$\{\tilde{X}_t\} \text{ and } \{\tilde{Y}_t\}$$

The nonnormalized spectral density of  $X_t$  is estimated by

$$\hat{h}_X(\omega) = \sum_j \omega_j I_{n,N,\tilde{X}}(\omega_{k,j})$$

and the nonnormalized spectral density of  $Y_t$  is estimated by

$$\hat{h}_Y(\omega) = \sum_j \omega_j I_{n,N,\tilde{Y}}(\omega_{k,j})$$

where

$$\omega_{k,j} = \frac{2\pi\{k(\omega) + j\}}{N}$$

and  $k(\omega)$  is the integer such that  $\omega_{k,0}$  is closest to  $\omega$ . The sequence of  $m = \text{NWT}$  weights  $\{w_j\}$  for  $j = -\lfloor m/2 \rfloor, \dots, (m - \lfloor m/2 \rfloor - 1)$  satisfies  $\sum_j w_j = 1$ . These weights are fixed in the sense that they do not depend on the frequency  $\omega$  at which to estimate the spectral density. Usually,  $m$  is odd with the weights symmetric about the middle weight  $w_0$ . If  $m$  is even, the weight to the right of the middle is considered  $w_0$ . The argument `WT` may contain relative weights since they are normalized to sum to one in the actual computations. The above spectral density formulas assume the data  $\{X_t\}$  and  $\{Y_t\}$  correspond to a realization of a bivariate discrete-parameter stationary process observed consecutively in time. In this case, the observations are equally spaced in time with interval  $\Delta t = \text{TINT}$  equivalent to one. However, if the data correspond to a realization of a bivariate continuous-parameter stationary process recorded at equal time intervals, then the spectral density estimates must be adjusted for the effect of aliasing. In general, the estimate of  $h_X(\omega)$  is given by

$$\hat{h}_X(\omega) = \Delta t \hat{h}_X(\omega), \quad |\omega| \leq \pi / \Delta t$$

and the estimate of  $h_Y(\omega)$  is given by

$$\hat{h}_Y(\omega) = \Delta t \hat{h}_Y(\omega), \quad |\omega| \leq \pi / \Delta t.$$

The nonnormalized spectral density is estimated over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = \text{NF}$ . These frequencies are in the scale specified by the argument `IFSCAL` but are transformed to the scale of radians per unit time for computational purposes. The frequency  $\omega$  of the desired spectral estimate is assumed to be input in a form already adjusted for the time interval  $\Delta t$ . The cross-spectral density function is complex-valued in general and may be written as

$$h_{XY}(\omega) = c_{XY}(\omega) - iq_{XY}(\omega)$$

The *cospectrum* is estimated by

$$\hat{c}_{XY}(\omega) = \sum_j w_j \Re \left\{ I_{n,N,\tilde{X}\tilde{Y}}(\omega_{k,j}) \right\}$$

and the *quadrature spectrum* is estimated by

$$\hat{q}_{XY}(\omega) = \sum_j w_j \Im \left\{ I_{n,N,\tilde{X}\tilde{Y}}(\omega_{k,j}) \right\}$$

Note that the same sequence of weights  $\{w_j\}$  used to estimate

$$\hat{h}_X(\omega) \text{ and } \hat{h}_Y(\omega)$$

is used to estimate

$$\hat{c}_{XY}(\omega) \text{ and } \hat{q}_{XY}(\omega)$$

The nonnormalized cross-spectral density estimate is computed over the same set of frequencies as the nonnormalized spectral density estimates discussed above with a similar adjustment for  $\Delta t$ . An equivalent representation of  $h_{XY}(\omega)$  is the polar form defined by

$$h_{XY}(\omega) = \alpha_{XY}(\omega)e^{i\phi_{XY}(\omega)}$$

The *cross-amplitude* spectrum is estimated by

$$\hat{\alpha}_{XY}(\omega) = \left\{ \hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega) \right\}^{1/2}$$

and the *phase spectrum* is estimated by

$$\hat{\phi}_{XY}(\omega) = \tan^{-1} \left\{ -\hat{q}_{XY}(\omega) / \hat{c}_{XY}(\omega) \right\}$$

Finally, the *coherency spectrum* is estimated by

$$|\hat{w}_{XY}(\omega)| = \left\{ \frac{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)}{\hat{h}_X(\omega)\hat{h}_Y(\omega)} \right\}^{1/2}$$

The *coherence* or squared coherency is output.

### Example

Consider the Robinson Multichannel Time Series Data (Robinson 1967, page 204) where  $X$  is the Wölfer sunspot number and  $Y$  is the northern light activity for the years 1770 through 1869. Application of routine CSWED to these data produces the following results.

```

INTEGER      IPRINT, LDCPM, LDCSM, LDRDAT, N, NDRDAT, NF, NOBS,
&            NPAD, NWT
PARAMETER    (IPRINT=0, LDRDAT=100, NDRDAT=4, NF=10, NOBS=100,
&            NWT=7, LDCSM=NF, NPAD=NOBS-1, N=NPAD+NOBS,
&            LDCPM=N/2+1)
C
INTEGER      I, IFSCAL, NRCOL, NRROW
REAL         ASIN, CPM(LDCPM,10), CSM(LDCSM,9), F(NF), FLOAT, PI,
&            RDATA(LDRDAT,NDRDAT), SSUM, TINT, WT(NWT), X(NOBS),
&            XCNTR, Y(NOBS), YCNTR
CHARACTER    CLABEL1(5)*24, CLABEL2(6)*16, FMT*7, RLABEL(1)*6,
&            TITLE1*32, TITLE2*40
INTRINSIC    ASIN, FLOAT
EXTERNAL     CSWED, GDATA, SSUM, WRRRL
C
EQUIVALENCE (X(1), RDATA(1,2)), (Y(1), RDATA(1,3))
C
DATA WT/1.0, 2.0, 3.0, 4.0, 3.0, 2.0, 1.0/
DATA FMT/'(F12.4)'/
DATA CLABEL1/'%/%/ k', '%/%/Frequency', '%/%/Period',
&            'Spectral%/Estimate%/of X', 'Spectral%/Estimate%/of Y'/
DATA CLABEL2/'%/ k', '%/Cospectrum', '%/Quadrature',
&            'Cross%/Amplitude', '%/Phase', '%/Coherence'/
DATA RLABEL/'NUMBER'/
DATA TITLE1/'Results of the Spectral Analyses'/

```

```

DATA TITLE2/'%/Results of the Cross-Spectral Analysis'/
C                               Initialization
PI = 2.0*ASIN(1.0)
DO 10 I=1, NF
    F(I) = PI*FLOAT(I)/FLOAT(NF)
10 CONTINUE
C                               Robinson data
CALL GDATA (8, 0, NRROW, NRCOL, RDATA, LDRDAT, NDRDAT)
C                               Center on arithmetic means
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
YCNTR = SSUM(NOBS,Y,1)/FLOAT(NOBS)
C                               Frequency in radians per unit time
IFSCAL = 0
C                               Time interval for discrete data
TINT = 1.0
C                               Compute the cross periodogram
CALL CSWED (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL, NF,
&          F, TINT, NWT, WT, CPM, LDCPM, CSM, LDCSM)
C                               Print results
CALL WRRRL (TITLE1, NF, 4, CSM, LDCSM, 0, FMT, RLABEL, CLABEL1)
CALL WRRRL (TITLE2, NF, 5, CSM(1,5), LDCSM, 0, FMT, RLABEL,
&          CLABEL2)
C
END

```

### Output

#### Results of the Spectral Analyses

k	Frequency	Period	Spectral Estimate of X	Spectral Estimate of Y
1	0.3142	20.0000	116.9550	1315.8370
2	0.6283	10.0000	1206.6086	1005.1219
3	0.9425	6.6667	84.8369	317.2589
4	1.2566	5.0000	55.2120	270.2111
5	1.5708	4.0000	46.5748	115.6768
6	1.8850	3.3333	12.4050	250.0125
7	2.1991	2.8571	7.0934	82.6773
8	2.5133	2.5000	3.4091	62.3267
9	2.8274	2.2222	5.6828	12.8970
10	3.1416	2.0000	4.0346	17.6441

#### Results of the Cross-Spectral Analysis

k	Cospectrum	Quadrature	Amplitude	Phase	Coherence
1	94.0531	-254.0125	270.8659	1.2162	0.4767
2	702.5118	21.9823	702.8557	-0.0313	0.4073
3	70.2379	-31.4431	76.9547	0.4209	0.2200
4	-1.8715	-36.1639	36.2123	1.6225	0.0879
5	36.6366	-18.5925	41.0843	0.4696	0.3133
6	32.7071	-6.6569	33.3776	0.2008	0.3592
7	9.4887	-9.1692	13.1950	0.7683	0.2969
8	9.2534	-0.3000	9.2583	0.0324	0.4034
9	-0.5568	2.9455	2.9977	-1.7576	0.1226
10	1.7640	-1.8321	2.5433	0.8043	0.0909

---

## CSWEP/DCSWEP (Single/Double precision)

Estimate the nonnormalized cross-spectral density of two stationary time series using a weighted cross periodogram given the spectral densities and cross periodogram.

### Usage

```
CALL CSWEP (N, SX, SY, CPREAL, CPIMAG, NF, F, NWT, WT,  
           COSPEC, QUADRA, CRAMPL, PHASE, COHERE)
```

### Arguments

**N** — Number of observations in each of the appropriately centered and padded time series *X* and *Y*. (Input)

*N* must be greater than or equal to two.

**SX** — Vector of length *NF* containing the estimate of the spectral density of the first time series *X*. (Input)

**SY** — Vector of length *NF* containing the estimate of the spectral density of the second time series *Y*. (Input)

**CPREAL** — Vector of length  $\lfloor N/2 \rfloor + 1$  containing the real part of the cross periodogram between *X* and *Y*. (Input)

The real part of the cross periodogram evaluated at (angular) frequency  $\omega_k = 2\pi k/N$  is given by  $CPREAL(k + 1)$ ,  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ .

**CPIMAG** — Vector of length  $\lfloor N/2 \rfloor + 1$  containing the imaginary part of the cross periodogram between *X* and *Y*. (Input)

The imaginary part of the cross periodogram evaluated at (angular) frequency  $\omega_k = 2\pi k/N$  is given by  $CPIMAG(k + 1)$ ,  $k = 0, 1, \dots, \lfloor N/2 \rfloor$ .

**NF** — Number of (angular) frequencies. (Input)

*F* must be greater than or equal to one.

**F** — Vector of length *NF* containing the (angular) frequencies at which the spectral density is estimated. (Input)

The elements of *F* must be in the range  $(-\pi, \pi)$  inclusive.

**NWT** — Number of weights. (Input)

*NWT* must be greater than or equal to one.

**WT** — Vector of length *NWT* containing the weights used to smooth the periodogram. (Input)

The actual weights are the values in *WT* normalized to sum to 1 with the current periodogram ordinate taking the middle weight for *NWT* odd or the weight to the right of the middle for *NWT* even.

**COSPEC** — Vector of length *NF* containing the estimate of the cospectrum. (Output)

**QUADRA** — Vector of length  $NF$  containing the estimate of the quadrature spectrum. (Output)

**CRAMPL** — Vector of length  $NF$  containing the estimate of the cross-amplitude spectrum. (Output)

**PHASE** — Vector of length  $NF$  containing the estimate of the phase spectrum. (Output)

**COHERE** — Vector of length  $NF$  containing the estimate of the coherence. (Output)

### Comments

1. The periodograms of  $X$  and  $Y$  and cross periodogram between  $X$  and  $Y$  may be computed via the routine `CPFFFT` (page 750). The spectral densities of  $X$  and  $Y$  may then be estimated using any of the routines `SSWD` (page 729), `SWED` (page 741), `SSWP` (page 736), or `SWEP` (page 747). Thus, different window types and/or weight sequences may be used to estimate the spectral and cross-spectral densities given either the series or their periodograms. Note that use of the modified periodograms and modified cross periodogram ensures that the scales of the spectral and cross-spectral densities and their estimates are equivalent.
2. The time sampling interval,  $TINT$ , is assumed to be equal to one. This assumption is appropriate for discrete parameter processes. The adjustment for continuous parameter processes ( $TINT > 0.0$ ) involves multiplication of the frequency vector  $F$  by  $1/TINT$  and multiplication of the spectral and cross-spectral density estimates by  $TINT$ .
3. To convert the frequency scale from radians per unit time to cycles per unit time, multiply  $F$  by  $1/(2\pi)$ .

### Algorithm

Routine `CSWEP` estimates the nonnormalized cross-spectral density function of two jointly stationary time series using a fixed sequence of weights given the modified cross-periodogram and spectral densities of the appropriately centered and padded data

$$\{\tilde{X}_t\}$$

and

$$\{\tilde{Y}_t\}$$

for  $t = 1, \dots, N$ . The routine `CPFFFT` (page 750) may be used to compute the modified periodograms

$$I_{n,N,\tilde{X}}(\omega_k) \text{ and } I_{n,N,\tilde{Y}}(\omega_k)$$

and cross-periodogram



$$I_{n,N,\tilde{X}\tilde{Y}}(\omega_k)$$

over the discrete set of nonnegative frequencies

$$\omega_k = \frac{2\pi k}{N}, \quad k = 0, 1, \dots, \lfloor N/2 \rfloor$$

(Here,  $\lfloor a \rfloor$  means the greatest integer less than or equal to  $a$ .) Either routine *SSWP* (page 736) or routine *SWEF* (page 747) may be applied to the periodograms to obtain nonnormalized spectral density estimates

$$\hat{h}_X(\omega) \text{ and } \hat{h}_Y(\omega)$$

over the set of frequencies

$$\omega = f_i, \quad i = 1, \dots, n_f$$

where  $n_f = NF$ . These frequencies are in the scale of radians per unit time. The time sampling interval  $\Delta t$  is assumed to be equal to one. Note that the spectral window or weight sequence used to compute

$$\hat{h}_X(\omega)$$

may differ from that used to compute

$$\hat{h}_Y(\omega)$$

The cross-spectral density function is complex-valued in general and may be written as

$$h_{XY}(\omega) = c_{XY}(\omega) - iq_{XY}(\omega)$$

The *cospectrum* is estimated by

$$\hat{c}_{XY}(\omega) = \sum_j w_j \Re \{ I_{n,N,\tilde{X}\tilde{Y}}(\omega_{k,j}) \}$$

and the *quadrature spectrum* is estimated by

$$\hat{q}_{XY}(\omega) = \sum_j w_j \Im \{ I_{n,N,\tilde{X}\tilde{Y}}(\omega_{k,j}) \}$$

where

$$\omega_{k,j} = \frac{2\pi\{k(\omega) + j\}}{N}$$

and  $k(\omega)$  is the integer such that  $\omega_{k,0}$  is closest to  $\omega$ . The sequence of  $m = NWT$  weights  $\{w_j\}$  for  $j = -\lfloor m/2 \rfloor, \dots, (m - \lfloor m/2 \rfloor - 1)$  satisfies  $\sum_j w_j = 1$ . These weights are fixed in the sense that they do not depend on the frequency  $\omega$  at which to estimate  $h_{XY}(\omega)$ . Usually,  $m$  is odd with the weights symmetric about the middle weight  $w_0$ . If  $m$  is even, the weight to the right of the middle is

considered  $w_0$ . The argument WT may contain relative weights since they are normalized to sum to one in the actual computations. The nonnormalized cross-spectral density estimate is computed over the same set of frequencies as the nonnormalized spectral density estimates. However, the particular weight sequence used to compute

$$\hat{h}_{XY}(\omega)$$

need not correspond to either the weight sequence or spectral window used to compute either

$$\hat{h}_X(\omega) \text{ or } \hat{h}_Y(\omega)$$

An equivalent representation of  $h_{XY}(\omega)$  is the *polar form* defined by

$$h_{XY}(\omega) = \alpha_{XY}(\omega)e^{i\phi_{XY}(\omega)}$$

The *cross-amplitude* spectrum is estimated by

$$\hat{\alpha}_{XY}(\omega) = \{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)\}^{1/2}$$

and the *phase spectrum* is estimated by

$$\hat{\phi}_{XY}(\omega) = \tan^{-1}\{-\hat{q}_{XY}(\omega) / \hat{c}_{XY}(\omega)\}$$

Finally, the *coherency spectrum* is estimated by

$$|\hat{w}_{XY}(\omega)| = \left\{ \frac{\hat{c}_{XY}^2(\omega) + \hat{q}_{XY}^2(\omega)}{\hat{h}_X(\omega)\hat{h}_Y(\omega)} \right\}^{1/2}$$

The coherence or squared coherency is output.

### Example

Consider the Robinson Multichannel Time Series Data (Robinson 1967, page 204) where  $X$  is the Wölfer sunspot number and  $Y$  is the northern light activity for the years 1770 through 1869. Application of routine CSWEP to these data produces the following results.

```

C
INTEGER      IPRINT, LDCPM, LDCSM, LDRDAT, N, NDRDAT, NF, NOBS,
&            NPAD, NWT
PARAMETER    (IPRINT=0, LDRDAT=100, NDRDAT=4, NF=10, NOBS=100,
&            NWT=7, LDCSM=NF, NPAD=NOBS-1, N=NOBS+NPAD,
&            LDCPM=N/2+1)

INTEGER      I, IFSCAL, IPVER, NROW, NVAR
REAL         ASIN, COHERE(NF), COSPEC(NF), CPIMAG(LDCPM),
&            CPM(LDCPM,10), CPREAL(LDCPM), CRAMPL(NF),
&            CSM(LDCSM,9), F(NF), FLOAT, PHASE(NF), PI, PX(LDCPM),
&            PY(LDCPM), QUADRA(NF), RDATA(LDRDAT,NDRDAT), SSUM,
&            SX(NF), SY(NF), WT(NWT), X(NOBS), XCNTR, Y(NOBS),
&            YCNTR
CHARACTER    CLABEL1(5)*24, CLABEL2(6)*16, FMT*8, RLABEL(1)*6,
```

```

&          TITLE1*32, TITLE2*40
INTRINSIC  ASIN, FLOAT
EXTERNAL  CPFFFT, CSWEP, GDATA, SCOPY, SSUM, SWEP, WRRRL
C
EQUIVALENCE (X(1), RDATA(1,2)), (Y(1), RDATA(1,3))
EQUIVALENCE (PX(1), CPM(1,3)), (PY(1), CPM(1,6))
EQUIVALENCE (CPREAL(1), CPM(1,9)), (CPIMAG(1), CPM(1,10))
C
DATA WT/1.0, 2.0, 3.0, 4.0, 3.0, 2.0, 1.0/
DATA FMT/'(F12.4)'/
DATA CLABEL1/'%/ k', '%/Frequency', '%/Period',
& 'Spectral%/Estimate%/of X', 'Spectral%/Estimate%/of Y'/
DATA CLABEL2/'%/ k', '%/Cospectrum', '%/Quadrature',
& 'Cross%/Amplitude', '%/Phase', '%/Coherence'/
DATA RLABEL/'NUMBER'/
DATA TITLE1/'Results of the Spectral Analyses'/
DATA TITLE2/'%/Results of the Cross-Spectral Analysis'/
C
                                Initialization
PI = 2.0*ASIN(1.0)
DO 10 I=1, NF
    F(I) = PI*FLOAT(I)/FLOAT(NF)
    CALL SCOPY (NF, F, 1, CSM(1,1), 1)
    CSM(I,2) = 2.0*FLOAT(NF)/FLOAT(I)
10 CONTINUE
C
                                Robinson data
CALL GDATA (8, 0, NROW, NVAR, RDATA, LDRDAT, NDRDAT)
C
                                Center on arithmetic means
XCNTR = SSUM(NOBS,X,1)/FLOAT(NOBS)
YCNTR = SSUM(NOBS,Y,1)/FLOAT(NOBS)
C
                                Frequency in radians per unit time
IFSCAL = 0
C
                                Modified periodogram version
IPVER = 1
C
                                Compute the cross periodogram
CALL CPFFFT (NOBS, X, Y, IPRINT, XCNTR, YCNTR, NPAD, IFSCAL,
& IPVER, CPM, LDCPM)
C
                                Estimate the spectral densities
CALL SWEP (N, PX, NF, F, NWT, WT, SX)
CALL SWEP (N, PY, NF, F, NWT, WT, SY)
C
                                Estimate the cross-spectral density
CALL CSWEP (N, SX, SY, CPREAL, CPIMAG, NF, F, NWT, WT, COSPEC,
& QUADRA, CRAMPL, PHASE, COHERE)
C
                                Print results
C
C
C
                                Copy results to output matrices
CALL SCOPY (NF, SX, 1, CSM(1,3), 1)
CALL SCOPY (NF, SY, 1, CSM(1,4), 1)
CALL SCOPY (NF, COSPEC, 1, CSM(1,5), 1)
CALL SCOPY (NF, QUADRA, 1, CSM(1,6), 1)
CALL SCOPY (NF, CRAMPL, 1, CSM(1,7), 1)
CALL SCOPY (NF, PHASE, 1, CSM(1,8), 1)
CALL SCOPY (NF, COHERE, 1, CSM(1,9), 1)
C
                                Call printing routines
CALL WRRRL (TITLE1, NF, 4, CSM, NF, 0, FMT, RLABEL, CLABEL1)
CALL WRRRL (TITLE2, NF, 5, CSM(1,5), NF, 0, FMT, RLABEL, CLABEL2)
C
END

```

## Output

### Results of the Spectral Analyses

k	Frequency	Period	Spectral Estimate of X	Spectral Estimate of Y
1	0.3142	20.0000	116.9550	1315.8370
2	0.6283	10.0000	1206.6086	1005.1219
3	0.9425	6.6667	84.8369	317.2589
4	1.2566	5.0000	55.2120	270.2111
5	1.5708	4.0000	46.5748	115.6768
6	1.8850	3.3333	12.4050	250.0125
7	2.1991	2.8571	7.0934	82.6773
8	2.5133	2.5000	3.4091	62.3267
9	2.8274	2.2222	5.6828	12.8970
10	3.1416	2.0000	4.0346	17.6441

### Results of the Cross-Spectral Analysis

k	Cospectrum	Quadrature	Cross Amplitude	Phase	Coherence
1	94.0531	-254.0125	270.8659	1.2162	0.4767
2	702.5118	21.9823	702.8557	-0.0313	0.4073
3	70.2379	-31.4431	76.9547	0.4209	0.2200
4	-1.8715	-36.1639	36.2123	1.6225	0.0879
5	36.6366	-18.5925	41.0843	0.4696	0.3133
6	32.7071	-6.6569	33.3776	0.2008	0.3592
7	9.4887	-9.1692	13.1950	0.7683	0.2969
8	9.2534	-0.3000	9.2583	0.0324	0.4034
9	-0.5568	2.9455	2.9977	-1.7576	0.1226
10	1.7640	-1.8321	2.5433	0.8043	0.0909

# Chapter 9: Covariance Structures and Factor Analysis

---

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---

## Usage Notes

Notation that is consistently used throughout this chapter is given in the following table. The FORTRAN equivalent of the symbols used are also given.

**Notation Used**

Symbol	FORTRAN Symbol	Meaning
$p$	NVAR	Number of variables in the observed variables
$k$	NF	Number of factors
$\Sigma$	COV	Population (or sample) covariance (correlation) matrix
$A$	A	Unrotated factor loadings
$B$	B	Rotated factor loadings
$T$	T	Factor rotation matrix
	TI	Image transformation matrix
$\beta$	SCOEf	Factor score coefficients

The routines in this chapter can generally be used for one or more of several purposes. Among these purposes are the following:

1. Data description: The information in the data is summarized by the factor loadings or by the eigenvectors and eigenvalues.
2. Data reduction: The information in a multivariate sample is reduced to a much smaller number of factors or principal components.
3. Variable clustering: The principal component coefficients or factor loadings lead to a grouping (clustering) of the variables.
4. Model building: Linear models relating the variables to the factors or principal components are estimated. Hypothesis tests may be used to obtain parsimonious and/or other descriptions of the data.

### Principal Components

The idea in principal components is to find a small number of linear combinations of the original variables that maximize the variance accounted for in the original data. This amounts to an eigensystem analysis of the covariance (or correlation) matrix. In addition to the eigensystem analysis, routine PRINC (page 793) computes standard errors for the eigenvalues. Correlations of the original variables with the principal component scores are also computed.

The computation of common principal components via routine KPRIN (page 797) is equivalent to finding the “eigenvectors” that best simultaneously

diagonalize one or more variance-covariance matrices. For only one input variance-covariance matrix, the vectors computed actually are the eigenvectors of the matrix.

## Factor Analysis

Factor analysis and principal component analysis, while quite different in assumptions, often serve the same ends. Unlike principal components in which linear combinations yielding the highest possible variances are obtained, factor analysis generally obtains linear combinations of the observed variables according to a model relating the observed variables to hypothesized underlying factors, plus a random error term called the unique error or uniqueness. In factor analysis, the unique errors associated with each variable are usually assumed to be independent of the factors. In addition, in the common factor model, the unique errors are assumed to be mutually independent. The factor analysis model is

$$x - \mu = \Lambda f + e$$

where  $x$  is the  $p$  vector of observed variables,  $\mu$  is the  $p$  vector of variable means,  $\Lambda$  is the  $p \times k$  matrix of factor loadings,  $f$  is the  $k$  vector of hypothesized underlying random factors, and  $e$  is the  $p$  vector of hypothesized unique random errors.

Because much of the computation in factor analysis was originally done by hand or was expensive on early computers, quick (but dirty) algorithms that made the calculations possible were developed. One result is the many factor extraction methods available today. Generally speaking, in the exploratory or model building phase of a factor analysis, a method of factor extraction that is not computationally intensive (such as principal components, principal factor, or image analysis) is used. If desired, a computationally intensive method is then used to obtain (what is hoped will be) the final factors.

In exploratory factor analysis, the unrotated factor loadings obtained from the factor extraction are generally transformed (rotated) to simplify the interpretation of the factors. Rotation is possible because of the overparameterization in the factor analysis model. The method used for rotation may result in factors that are independent (orthogonal rotations) or correlated (oblique rotations). Prior information may be available (or hypothesized) in which case a Procrustes rotation could be used. When no prior information is available, an analytic rotation can be performed.

Once the factor loadings have been extracted and rotated (if desired), estimates for the hypothesized underlying factors can be computed. First, one of several available methods in routine `FCOEF` (page 833) is used to compute the factor score coefficients. Routine `FSCOR` (page 838) is then called with these factor score coefficients to compute the factor scores.

The steps generally used in a factor analysis are summarized as follows:

## Steps in a Factor Analysis

### Step 1

Calculate Covariance (Correlation) Matrix IMSL routine CORVC (page 314)
--

### Step 2

Initial Factor Extraction FACTR, page 801
--

### Step 3

Factor Rotation

Orthogonal		Oblique	
No Prior Info. FROTA, page 809	Prior Info. FOPCS, page 812	No Prior Info. FPRMX, page 818 FDOBL, page 815 FHARR, page 822 FGCRF, page 825	Prior Info. FPRMX, page 818

### Step 3a

Image Analysis FIMAG, page 829
-----------------------------------

### Step 3b

Factor Variances FRVAR, page 831
-------------------------------------

### Step 4

Factor Coefficients FCOEF, page 833
--

### Step 5

Factor Scores FSCOR, page 838
----------------------------------



## Independence of Sets of Variables and Canonical Correlation Analysis

Routine `MVIND` (page 842) computes the Wilks likelihood-ratio test of independence among several sets of variables. Routines `CANCR` (page 844) and `CANVC` (page 857) compute some other tests of independence between exactly two sets of variables. Routine `CANCR` uses the raw data as input while `CANVC` uses the sample variance-covariance matrix. Furthermore, `CANCR` and `CANVC` perform a canonical correlation analysis. Since `CANCR` uses a better algorithm in terms of numerical stability (it does not compute the covariance matrix), `CANCR` should be used if possible. However, if the raw data is not available, or if there is too much data for all of it to reside in memory at the same time, or if multiple canonical correlation analyses are to be performed based on the same pre-computed sample variance-covariance matrix, then the use of `CANVC` may be necessary. Canonical correlation analysis is useful for characterizing the independent linear statistical relationships that exist between the two sets of variables. This involves computing linear combinations of the variables in the two separate sets and their associated correlation. The coefficients of the variables in the linear combinations are called the “canonical coefficients,” and the correlations are called “canonical correlations.” Evaluation of the linear combinations using the canonical coefficients gives the “canonical scores.” Routine `CANCR` computes the canonical scores for the observed data. Routine `FSCOR` can be used to compute the canonical scores for new data or for the observed data if `CANVC` is used.

---

## PRINC/DPRINC (Single/Double precision)

Compute principal components from a variance-covariance matrix or a correlation matrix.

### Usage

```
CALL PRINC (NDF, NVAR, COV, LDCOV, ICOV, EVAL, PCT, STD,  
           EVEC, LDEVEC, A, LDA)
```

### Arguments

***NDF*** — Number of degrees of freedom in `COV`. (Input)

If `NDF` is less than or equal to 0, 100 degrees of freedom are assumed.

***NVAR*** — Order of matrix `COV`. (Input)

***COV*** — `NVAR` by `NVAR` matrix containing the covariance or correlation matrix. (Input)

Only the upper triangular part of `COV` is referenced.

***LDCOV*** — Leading dimension of `COV` exactly as specified in the dimension statement in the calling program. (Input)

**ICOV** — Covariance/Correlation matrix option parameter. (Input)  
ICOV = 0 means that a covariance matrix is input. Otherwise, a correlation matrix is input.

**EVAL** — Vector of length NVAR containing the eigenvalues from matrix COV ordered from largest to smallest. (Output)

**PCT** — Vector of length NVAR containing the cumulative percent of the total variance explained by each principal component. (Output)

**STD** — Vector of length NVAR containing the estimated asymptotic standard errors of the eigenvalues. (Output)

**EVEC** — NVAR by NVAR matrix containing the eigenvectors of COV, stored columnwise. (Output)  
Each vector is normalized to have Euclidean length equal to the value one. Also, the sign of each vector is set so that the largest component in magnitude (the first of the largest if there are ties) is made positive.

**LDEVEC** — Leading dimension of EVEC exactly as specified in the dimension statement in the calling program. (Input)

**A** — NVAR by NVAR matrix containing the correlations of the principal components (the columns) with the observed/standardized variables (the rows). (Output)  
If ICOV = 0, then the correlations are with the observed variables. Otherwise, the correlations are with the standardized (to a variance of 1.0) variables. In the principal component model for factor analysis, matrix A is the matrix of unrotated factor loadings.

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

### Comments

#### Informational errors

Type	Code	
3	1	Because NDF is zero or less, 100 degrees of freedom will be used.
3	2	One or more eigenvalues much less than zero are computed. The matrix COV is not nonnegative definite. In order to continue computations of STD and A, these eigenvalues are treated as zero.

### Algorithm

Routine PRINC finds the principal components of a set of variables from a sample covariance or correlation matrix. The characteristic roots, characteristic vectors, standard errors for the characteristic roots, and the correlations of the principal component scores with the original variables are computed. Principal components obtained from correlation matrices are the same as principal components obtained from standardized (to unit variance) variables.

The principal component scores are the elements of the vector  $y = \Gamma^T x$  where  $\Gamma$  is the matrix whose columns are the characteristic vectors (eigenvectors) of the sample covariance (or correlation) matrix and  $x$  is the vector of observed (or standardized) random variables. The variances of the principal component scores are the characteristic roots (eigenvalues) of the the covariance (correlation) matrix.

Asymptotic variances for the characteristic roots were first obtained by Girshick (1939) and are given more recently by Kendall, Stuart, and Ord (1983, page 331). These variances are computed either for covariance matrices ( $ICOV = 0$ ) or for correlation matrices ( $ICOV \neq 0$ ).

The correlations of the principal components with the observed (or standardized) variables are given in the matrix A. When the principal components are obtained from a correlation matrix, A is the same as the matrix of unrotated factor loadings obtained for the principal components model for factor analysis.

### Example

Principal components are computed for a nine-variable matrix.

```

C      INTEGER      ICOV, LDA, LDICOV, LDEVEC, NDF, NVAR
PARAMETER (ICOV=1, LDA=9, LDICOV=9, LDEVEC=9, NDF=100, NVAR=9)
C
C      REAL          A(LDA,NVAR), COV(LDICOV,NVAR), EVAL(NVAR),
&      EVEC(LDEVEC,NVAR), PCT(NVAR), STD(NVAR)
C      EXTERNAL      PRINC, WRRRN
C
DATA COV/
& 1.000, 0.523, 0.395, 0.471, 0.346, 0.426, 0.576, 0.434, 0.639,
& 0.523, 1.000, 0.479, 0.506, 0.418, 0.462, 0.547, 0.283, 0.645,
& 0.395, 0.479, 1.000, 0.355, 0.270, 0.254, 0.452, 0.219, 0.504,
& 0.471, 0.506, 0.355, 1.000, 0.691, 0.791, 0.443, 0.285, 0.505,
& 0.346, 0.418, 0.270, 0.691, 1.000, 0.679, 0.383, 0.149, 0.409,
& 0.426, 0.462, 0.254, 0.791, 0.679, 1.000, 0.372, 0.314, 0.472,
& 0.576, 0.547, 0.452, 0.443, 0.383, 0.372, 1.000, 0.385, 0.680,
& 0.434, 0.283, 0.219, 0.285, 0.149, 0.314, 0.385, 1.000, 0.470,
& 0.639, 0.645, 0.504, 0.505, 0.409, 0.472, 0.680, 0.470, 1.000/
C
CALL PRINC (NDF, NVAR, COV, LDICOV, ICOV, EVAL, PCT, STD, EVEC,
&          LDEVEC, A, LDA)
C
CALL WRRRN ('EVAL', 1, NVAR, EVAL, 1, 0)
CALL WRRRN ('PCT', 1, NVAR, PCT, 1, 0)
CALL WRRRN ('STD', 1, NVAR, STD, 1, 0)
CALL WRRRN ('EVEC', NVAR, NVAR, EVEC, LDEVEC, 0)
CALL WRRRN ('A', NVAR, NVAR, A, LDA, 0)
END

```

### Output

				EVAL				
1	2	3	4	5	6	7	8	9
4.677	1.264	0.844	0.555	0.447	0.429	0.310	0.277	0.196

PCT								
1	2	3	4	5	6	7	8	9
0.520	0.660	0.754	0.816	0.865	0.913	0.947	0.978	1.000

STD							
1	2	3	4	5	6	7	8
0.6498	0.1771	0.0986	0.0879	0.0882	0.0890	0.0944	0.0994

9  
0.1113

EVEC								
	1	2	3	4	5	6	7	8
1	0.3462	-0.2354	0.1386	-0.3317	-0.1088	0.7974	0.1735	-0.1240
2	0.3526	-0.1108	-0.2795	-0.2161	0.7664	-0.2002	0.1386	-0.3032
3	0.2754	-0.2697	-0.5585	0.6939	-0.1531	0.1511	0.0099	-0.0406
4	0.3664	0.4031	0.0406	0.1196	0.0017	0.1152	-0.4022	-0.1178
5	0.3144	0.5022	-0.0733	-0.0207	-0.2804	-0.1796	0.7295	0.0075
6	0.3455	0.4553	0.1825	0.1114	0.1202	0.0696	-0.3742	0.0925
7	0.3487	-0.2714	-0.0725	-0.3545	-0.5242	-0.4355	-0.2854	-0.3408
8	0.2407	-0.3159	0.7383	0.4329	0.0861	-0.1969	0.1862	-0.1623
9	0.3847	-0.2533	-0.0078	-0.1468	0.0459	-0.1498	-0.0251	0.8521

9  
1 -0.0488  
2 -0.0079  
3 -0.0997  
4 0.7060  
5 0.0046  
6 -0.6780  
7 -0.1089  
8 0.0505  
9 0.1225

A								
	1	2	3	4	5	6	7	8
1	0.7487	-0.2646	0.1274	-0.2471	-0.0728	0.5224	0.0966	-0.0652
2	0.7625	-0.1245	-0.2568	-0.1610	0.5124	-0.1312	0.0772	-0.1596
3	0.5956	-0.3032	-0.5133	0.5170	-0.1024	0.0990	0.0055	-0.0214
4	0.7923	0.4532	0.0373	0.0891	0.0012	0.0755	-0.2240	-0.0620
5	0.6799	0.5646	-0.0674	-0.0154	-0.1875	-0.1177	0.4063	0.0039
6	0.7472	0.5119	0.1677	0.0830	0.0804	0.0456	-0.2084	0.0487
7	0.7542	-0.3051	-0.0666	-0.2641	-0.3505	-0.2853	-0.1589	-0.1794
8	0.5206	-0.3552	0.6784	0.3225	0.0576	-0.1290	0.1037	-0.0854
9	0.8319	-0.2848	-0.0072	-0.1094	0.0307	-0.0981	-0.0140	0.4485

9  
1 -0.0216  
2 -0.0035  
3 -0.0442  
4 0.3127  
5 0.0021  
6 -0.3003  
7 -0.0482  
8 0.0224  
9 0.0543

---

## KPRIN/DKPRIN (Single/Double precision)

Maximum likelihood or least-squares estimates for principal components from one or more matrices.

### Usage

```
CALL KPRIN (NVAR, NMAT, COV, LDICOV, ANI, IMETH, EVEC,  
           LDEVEC)
```

### Arguments

**NVAR** — Number of variables in each matrix. (Input)  
NVAR must be 2 or greater.

**NMAT** — Number of matrices. (Input)

**COV** — NVAR by NVAR by NMAT array containing the NMAT covariance or correlation matrices. (Input)  
Only the upper triangular elements of each matrix are referenced.

**LDICOV** — Leading and second dimensions of COV exactly as specified in the dimension statement of the calling program. (Input)  
The first two dimensions of COV must be equal.

**ANI** — Vector of length NMAT containing the number of observations in each of the covariance matrices. (Input)  
For least-squares estimation, the square root of ANI(I) is the weight to be used for the I-th covariance matrix. Since the elements of ANI are used as weights, they need not be integers.

**IMETH** — Method to be used for extracting the estimated principal components. (Input)  
For IMETH = 0, maximum likelihood estimation is used. For IMETH = 1, least-squares estimation is used.

**EVEC** — NVAR by NVAR matrix containing the estimated principal components. (Output)  
Each column of EVEC contains a principal component vector (an “eigenvector”). The ordering of the eigenvectors is such that the sum of the corresponding eigenvalues are ordered from largest to smallest. Each vector is normalized to have Euclidean length equal to the value one.

**LDEVEC** — Leading dimension of EVEC exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Automatic workspace usage is

$$\text{KPRIN } \text{NVAR}^2 * (\text{NMAT} + 1) + 2 * \text{NVAR} + 4 * \text{NMAT} + \max(3 * \text{NMAT}, \text{NVAR}^2) \text{ units, or}$$

DKPRIN  $2 * (NVAR^2 * (NMAT + 1) + 2 * NVAR + 4 * NMAT + \max(3 * NMAT, NVAR^2))$  units.

Workspace may be explicitly provided, if desired, by use of K2RIN/DK2RIN. The reference is

```
CALL K2RIN (NVAR, NMAT, COV, LDICOV, ANI, IMETH,
           EVEC, LDEVEC, H, AUX, BOLD, G, T)
```

The additional arguments are as follows:

**H** — Work vector of length  $NVAR^2 * NMAT$ . On return from K2RIN, H may be treated as an array dimensioned as H(NVAR, NVAR, NMAT). Each NVAR by NVAR matrix in H is computed as  $(EVEC)^T * COV(I) * EVEC$ , i.e., H contains the “eigenvalues” and the “residuals” for each covariance matrix. Here, COV(I) is the I-th covariance matrix.

**AUX** — Work vector of length  $\max(3 * NMAT, NVAR^2)$ .

**BOLD** — Work vector of length  $NVAR^2$ .

**G** — Work vector of length  $2 * NVAR$ .

**T** — Work vector of length  $4 * NMAT$ .

2. Informational errors

Type	Code	
3	1	Convergence did not occur within 50 iterations. Convergence is assumed.
4	2	An input matrix is singular. Singular input matrices are not allowed in maximum likelihood estimation.

3. If user specified initial estimates for EVEC are desired (and argument error checking is not needed), then the routine K3RIN (DK3RIN) may be used. The calling sequence is

```
CALL K3RIN (NVAR, NMAT, COV, LDICOV, ANI, IMETH,
           EVEC, LDEVEC, AUX, BOLD, G, T)
```

On input, EVEC contains the initial estimates of the common principal components (EVEC must be an orthogonal matrix). On output, COV contains the NMAT matrices  $(EVEC)^T * COV(I) * EVEC$ . The user should be wary of stationary points in the likelihood if K3RIN is used.

### Algorithm

Routine KPRIN is the IMSL version of the F-G diagonalization routine of Flury and Constantine (1985) with modifications as discussed by Clarkson (1988a, 1988b). Let  $k = NMAT$ . Routine KPRIN computes the common principal components of  $k \geq 1$  covariance (or correlation) matrices using either a least-squares or a maximum likelihood criterion. Computing common principal components is equivalent to finding the “eigenvectors” that best simultaneously diagonalize  $k$  symmetric matrices. (Note that when  $k = 1$ , both least-squares and

maximum likelihood estimation yield the eigenvectors of the input matrix.) See Flury (1988) for applications of common principal components.

The algorithm proceeds by accumulating simple rotations as follows: Initial estimates of the diagonalizing principal components are found as the eigenvectors of the summed covariance matrices (unless `K3RIN` is used, see Comment 3 above). The covariance matrices are then pre- and post-multiplied by the initial estimates to obtain approximately diagonal matrices. Let

$$w_l^{ij}$$

denote the  $l$ -th  $2 \times 2$  matrix obtained from the  $(i, j)$ ,  $(i, i)$ , and  $(j, j)$  elements of  $S_l$ , where  $S_l$  is the  $l$ -th covariance matrix in `COV`. Then, for each  $i$  and  $j$ , a Jacobi rotation is found and applied such that the least-squares or maximum likelihood criterion is optimized over all  $k$  matrices in

$$w_l^{ij}$$

An iteration consists of computing and applying a Jacobi rotation for all  $p(p - 1)/2$  possible off-diagonal elements  $(i, j)$  where  $p = \text{NVAR}$ . A maximum of 50 iterations are allowed before convergence. Convergence is assumed when the maximum change in the any element in the eigenvectors from one iteration to the next is less than 0.0001.

Let  $\Gamma$  denote the current estimates of the optimizing principal components. Then, maximizing the multivariate normal likelihood is equivalent to minimizing the criterion

$$L = \prod_{i=1}^k \left( \frac{\det \hat{\Sigma}_i}{\det S_i} \right)^{n_i}$$

where  $S_i$  is the  $i$ -th covariance matrix,  $n_i$  is its degrees of freedom, and

$$\hat{\Sigma}_i$$

is the estimate of the covariance matrix under the common principal components model.

During each Jacobi iteration, an optimal orthogonal matrix  $T_{ij}$  is found that rotates the two vectors in columns  $i$  and  $j$  of  $\Gamma$ . When restricted to  $T_{ij}$ , the criterion above becomes

$$L = \prod_{i=1}^k \left( \frac{\det(\text{diag}(T_{ij}^T \Gamma^T S_i \Gamma T_{ij}))}{\det(S_i)} \right)^{n_i}$$

$\Gamma$  is updated as  $\Gamma T_{ij}$ . When convergence has been reached (the maximum change in  $\Gamma$  is less than 0.0001),  $\Gamma$  contains the optimizing principal components. Initially,  $\Gamma$  is taken as the eigenvectors of the matrix  $\sum_i S_i$ .

In least-squares estimation, the matrices  $T_{ij}$  are found such that the sum of the squared off-diagonal elements in the resulting “diagonalized” matrices are minimized. That is,  $T_{ij}$  is found to minimize

$$v_{ij}^T v_{ij}$$

where  $v_{ij}$  is the vector of length  $k$  containing the off-diagonal elements in the matrices

$$w_l^{ij}$$

See Flury and Gautschi (1986) for further details on the general algorithm, especially in maximum likelihood estimation. See Clarkson (1988b) for details of the least-squares algorithm.

If the “residual” matrices  $\Gamma^T S_l \Gamma$  are desired, they may be obtained in the work vector  $H$  returned from `K2RIN` or from the matrix `COV` returned from `K3RIN`. If the least-squares criterion is needed, it is easily computed as the sum of the squared off-diagonal elements in  $H$  (or `COV`). To compute the likelihood ratio criterion, the eigenvalues of each matrix in `COV` first need to be computed. Denote the eigenvalues from the  $l$ -th matrix by  $\lambda_{lj}$ , and let

$$\hat{\lambda}_{lj}$$

be the eigenvalues obtained under the common principal component model (and returned as the diagonal elements of  $H$  or, from `K3RIN`, `COV`). Then, the log-likelihood-ratio statistic for testing,

$$H_0: \Gamma^T \sum_l \Gamma$$

is diagonal,  $l = 1, \dots, k$ , is computed as:

$$\ell = \sum_{l=1}^k \sum_{j=1}^p n_l \log \frac{\hat{\lambda}_{lj}}{\lambda_{lj}}$$

The distribution of  $\ell$  under  $H_0$  is asymptotically  $\chi^2$  with  $(k-1)p(p-1)/2$  degrees of freedom (see Flury 1984).

### Example

The following example is taken from Flury and Constantine (1985). It involves two 4 by 4 covariance matrices. The two covariance matrices are given as:

$$\begin{pmatrix} 1.3 & -0.3 & -0.6 & 0 \\ -0.3 & 2.1 & 0 & -0.6 \\ -0.6 & 0 & 2.9 & -0.3 \\ 0 & -0.6 & -0.3 & 3.7 \end{pmatrix}$$



$$\begin{pmatrix} 1 & 0 & 0 & 0 \\ 0 & 2 & 0 & 0 \\ 0 & 0 & 3 & 0 \\ 0 & 0 & 0 & 4 \end{pmatrix}$$

```

C                                     SPECIFICATIONS FOR PARAMETERS
INTEGER      IMETH, LDICOV, LDEVEC, NGROUP, NVAR
PARAMETER    (IMETH=0, LDICOV=4, LDEVEC=4, NGROUP=2, NVAR=4)
C
REAL         ANI(NGROUP), COV(LDICOV,LDICOV,NGROUP),
&            EVEC(LDEVEC,NVAR)
EXTERNAL     KPRIN, WRRRN
C
DATA COV/1.3, -0.3, -0.6, 0, -0.3, 2.1, 0, -0.6, -0.6, 0, 2.9,
&      -0.3, 0, -0.6, -0.3, 3.7, 1, 0, 0, 0, 2, 0, 0, 0, 3,
&      0, 0, 0, 0, 4/
C
ANI(1) = 1
ANI(2) = 1
C
CALL KPRIN (NVAR, NGROUP, COV, LDICOV, ANI, IMETH, EVEC, LDEVEC)
C
CALL WRRRN ('EVEC', NVAR, NVAR, EVEC, LDEVEC, 0)
C
END

```

### Output

	EVEC			
	1	2	3	4
1	0.9743	-0.1581	-0.1581	0.0257
2	0.1581	0.9743	-0.0257	-0.1581
3	0.1581	-0.0257	0.9743	-0.1581
4	0.0257	0.1581	0.1581	0.9743

---

## FACTR/DFACTR (Single/Double precision)

Extract initial factor loading estimates in factor analysis.

### Usage

```
CALL FACTR (NVAR, COV, LDICOV, NF, IMTH, NDF, INIT, MAXIT,
            MAXSTP, EPS, EPSE, IPRINT, UNIQ, A, LDA, EVAL,
            STAT, DER)
```

### Arguments

**NVAR** — Number of variables. (Input)

**COV** — NVAR by NVAR matrix containing the variance-covariance or correlation matrix. (Input)

**LDICOV** — Leading dimension of COV exactly as specified in the dimension statement in the calling program. (Input)

**NF** — Number of factors in the model. (Input)

**IMTH** — Method used to obtain the estimates. (Input)

**IMTH Method**

- 0 Principal component (principal component model) or principal factor (common factor model). If **INIT** = 1 and **UNIQ** contains zeros, then this option results in the principal component method. Otherwise, the principal factor method is used.
- 1 Unweighted least squares (common factor model).
- 2 Generalized least squares (common factor model).
- 3 Maximum likelihood (common factor model).
- 4 Image factor analysis (common factor model).
- 5 Alpha factor analysis (common factor model).

**NDF** — Number of degrees of freedom in **COV**. (Input)

**NDF** is not required when **IMTH** = 0, 1, or 4. **NDF** defaults to 100 if **NDF** = 0.

**INIT** — Method used to obtain initial estimates of the unique variances. (Input)

**INIT Method**

- 0 Initial estimates are taken as the constant  $1 - \text{NF}/(2 * \text{NVAR})$  divided by the diagonal elements of the inverse of **COV**.
- 1 Initial estimates are input in vector **UNIQ**.

**MAXIT** — Maximum number of iterations in the iterative procedure. (Input)

Typical for methods 1 to 3 is 30, while 60 is typical for method 5. **MAXIT** is not referenced when **IMTH** = 0 or 4.

**MAXSTP** — Maximum number of step halvings allowed during any one iteration. (Input)

Typical is 8. **MAXSTP** is not referenced when **IMTH** = 0, 4, or 5.

**EPS** — Convergence criterion used to terminate the iterations. (Input)

For methods 1 to 3, convergence is assumed when the relative change in the criterion is less than **EPS**. For method 5, convergence is assumed when the maximum change (relative to the variance) of a uniqueness is less than **EPS**. **EPS** is not referenced when **IMTH** = 0 or 4. **EPS** = 0.0001 is typical.

**EPSE** — Convergence criterion used to switch to exact second derivatives. (Input)

When the largest relative change in the unique standard deviation vector is less than **EPSE**, exact second derivative vectors are used. Typical is 0.1. **EPSE** is not referenced when **IMTH** = 0, 4, or 5.

**IPRINT** — Printing option. (Input)

If **IPRINT** = 0, then no printing is performed. If **IPRINT** = 1, then printing of the final results is performed. If **IPRINT** = 2, then printing of an iteration summary and the final results is performed.

**UNIQ** — Vector of length *NVAR* containing the unique variances. (Input/Output, if *INIT* = 1; output, otherwise)

If *INIT* = 1, **UNIQ** contains the initial estimates of these variances on input. On output, **UNIQ** contains the estimated unique variances. For *IMTH* = 0, the unique variances are assumed to be known and are not changed from the input values when *INIT* = 1.

**A** — *NVAR* by *NF* matrix of unrotated factor loadings. (Output)

**LDA** — Leading dimension of **A** exactly as specified in the dimension statement of the calling program. (Input)

**EVAL** — Vector of length *NVAR* containing the eigenvalues of the matrix from which the factors were extracted. (Output)

If *IMTH* = 5, then the first *NF* positions of **EVAL** contain the **ALPHA** coefficients. Note that **EVAL** does not usually contain eigenvalues for matrix **COV**.

**STAT** — Vector of length 6 containing some output statistics. (Output)

<b>I</b>	<b>STAT(I)</b>
1	Value of the function minimum.
2	Tucker reliability coefficient.
3	Chi-squared test statistic for testing that <i>NF</i> common factors are adequate for the data.
4	Degrees of freedom in chi-squared. This is computed as $((NVAR - NF)^2 - NVAR - NF)/2$ .
5	Probability of a greater chi-squared statistic.
6	Number of iterations.

**STAT** is not used when *IMTH* = 0, 4, or 5.

**DER** — Vector of length *NVAR* containing the parameter updates when convergence was reached (or the iterations terminated). (Output)

### Comments

1. Automatic workspace usage is

**FACTR**  $3 * NVAR + 3 * NVAR^2$  units, or

**DFACTR**  $5 * NVAR + 6 * NVAR^2$  units.

Workspace may be explicitly provided, if desired, by use of **F2CTR/DF2CTR**. The reference is

```
CALL F2CTR (NVAR, COV, LDCOV, NF, IMTH, NDF, INIT,
            MAXIT, MAXSTP, EPS, EPSE, IPRINT, UNIQ,
            A, LDA, EVAL, STAT, DER, IS, COVI, WK,
            OLD, EVEC, HESS)
```

The additional arguments are as follows:

**IS** — Integer work vector of length equal to *NVAR*.

**COVI** — Real work vector of length equal to  $NVAR^2$ .

**WK** — Real work vector of length equal to `NVAR`.

**OLD** — Real work vector of length equal to `NVAR`.

**EVEC** — Real work vector of length equal to `NVAR`<sup>2</sup>.

**HESS** — Real work vector of length equal to `NVAR`<sup>2</sup>.

2. Informational errors

Type	Code	
3	1	Too many iterations. Convergence is assumed.
3	2	Too many step halvings. Convergence is assumed.
3	4	There are no degrees of freedom for the significance testing.

### Algorithm

Routine `FACTR` computes unrotated factor loadings in exploratory factor analysis models. Models available in `FACTR` are the principal component model for factor analysis and the common factor model with additions to the common factor model in alpha factor analysis and image analysis. Methods of estimation include principal components, principal factor, image analysis, unweighted least squares, generalized least squares, and maximum likelihood.

In the factor analysis model used for factor extraction, the basic model is given as  $\Sigma = \Lambda\Lambda^T + \Psi$  where  $\Sigma$  is the  $p \times p$  population covariance matrix,  $\Lambda$  is the  $p \times k$  matrix of factor loadings relating the factors  $f$  to the observed variables  $x$ , and  $\Psi$  is the  $p \times p$  matrix of covariances of the unique errors  $e$ . Here,  $p = \text{NVAR}$  and  $k = \text{NF}$ . The relationship between the factors, the unique errors, and the observed variables is given as  $x = \Lambda f + e$  where, in addition, it is assumed that the expected values of  $e$ ,  $f$ , and  $x$  are zero. (The sample means can be subtracted from  $x$  if the expected value of  $x$  is not zero.) It is also assumed that each factor has unit variance, the factors are independent of each other, and that the factors and the unique errors are mutually independent. In the common factor model, the elements of the vector of unique errors  $e$  are also assumed to be independent of one another so that the matrix  $\Psi$  is diagonal. This is not the case in the principal component model in which the errors may be correlated.

Further differences between the various methods concern the criterion that is optimized and the amount of computer effort required to obtain estimates. Generally speaking, the least-squares and maximum likelihood methods, which use iterative algorithms, require the most computer time with the principal factor, principal component and the image methods requiring much less time since the algorithms in these methods are not iterative. The algorithm in alpha factor analysis is also iterative, but the estimates in this method generally require somewhat less computer effort than the least-squares and maximum likelihood estimates. In all algorithms, one eigensystem analysis is required on each iteration.

## The Principal Component and Principal Factor Methods

Both the principal component and the principal factor methods compute the factor loading estimates as

$$\hat{\Gamma}\hat{\Delta}^{-1/2}$$

where  $\Gamma$  and the diagonal matrix  $\Delta$  are the eigenvectors and eigenvalues of a matrix. In the principal component model, the eigensystem analysis is performed on the sample covariance (correlation) matrix  $S$  while in the principal factor model the matrix  $(S - \Psi)$  is used. If the unique error variances  $\Psi$  are not known (i.e., if `INIT = 0`) in the principal factor model, then `FACTR` obtains estimates for them as discussed in Comment 3. If the principal components model is to be used, then the `INIT = 1` option should be set, and the vector `UNIQ` should be set so that all elements are zero. If `UNIQ` is not set, principal factor model estimates are computed.

The basic idea in the principal component method is to find factors that maximize the variance in the original data that is explained by the factors. Because this method allows the unique errors to be correlated, some factor analysts insist that the principal component method is not a factor analytic method. Usually however, the estimates obtained via the principal component model and other models in factor analysis will be quite similar.

It should be noted that both the principal component and the principal factor methods give different results when the correlation matrix is used in place of the covariance matrix. Indeed, any rescaling of the sample covariance matrix can lead to different estimates with either of these methods. A further difficulty with the principal factor method is the problem of estimating the unique error variances. Theoretically, these must be known in advance and passed to `FACTR` through `UNIQ`. In practice, the estimates of these parameters produced via the `INIT = 0` option in `FACTR` are often used. In either case, the resulting adjusted covariance (correlation) matrix

$$(S - \hat{\Psi})$$

may not yield the `NF` positive eigenvalues required for `NF` factors to be obtained. If this occurs, the user must either lower the number of factors to be estimated or give new unique error variance values.

## The Least-Squares and Maximum Likelihood Methods

Unlike the previous two methods, the algorithm used to compute estimates in this section is iterative (see Joreskog 1977). As with the principal factor model, the user may either initialize `UNIQ` or allow `FACTR` to compute initial estimates for the unique error variances. Unlike the principal factor method, `FACTR` then optimizes the criterion function with respect to both  $\Psi$  and  $\Gamma$ . (In the principal factor method,  $\Psi$  is assumed to be known. Given  $\Psi$ , estimates for  $\Lambda$  may be obtained.)

The major differences between the methods discussed in this section are in the criterion function that is optimized. Let  $S$  denote the sample covariance (correlation) matrix, and let  $\Sigma$  denote the covariance matrix that is to be estimated by the factor model. In the unweighted least-squares method, also called the iterated principal factor method or the minres method (see Harman 1976, page 177), the function minimized is the sum of the squared differences between  $S$  and  $\Sigma$ . This is written as  $\Phi_{ul} = .5 \text{ trace}((S - \Sigma)^2)$ .

Generalized least-squares and maximum likelihood estimates are asymptotically equivalent methods. Maximum likelihood estimates maximize the (normal theory) likelihood  $\{\Phi_{ml} = \text{trace}(\Sigma^{-1}S) - \log(|\Sigma^{-1}S|)\}$  while generalized least squares optimizes the function  $\Phi_{gs} = \text{trace}(\Sigma S^{-1} - I)^2$ .

In all three methods, a two-stage optimization procedure is used. This proceeds by first solving the likelihood equations for  $\Lambda$  in terms of  $\Psi$  and substituting the solution into the likelihood. This gives a criterion  $\phi(\Psi, \Lambda(\Psi))$ , which is optimized with respect to  $\Psi$ . In the second stage, the estimates

$$\hat{\Lambda}$$

are obtained from the estimates for  $\Psi$ .

The generalized least-squares and the maximum likelihood methods allow for the computation of a statistic (STAT(3)) for testing that  $NF$  common factors are adequate to fit the model. This is a chi-squared test that all remaining parameters associated with additional factors are zero. If the probability of a larger chi-squared is small (see STAT(5)) so that the null hypothesis is rejected, then additional factors are needed (although these factors may not be of any practical importance). Failure to reject does not legitimize the model. The statistic STAT(3) is a likelihood ratio statistic in maximum likelihood estimation. As such, it asymptotically follows a chi-squared distribution with degrees of freedom given in STAT(4).

The Tucker and Lewis (1973) reliability coefficient,  $\rho$ , is returned in STAT(2) when the maximum likelihood or generalized least-squares methods are used. This coefficient is an estimate of the ratio of explained to the total variation in the data. It is computed as follows:

$$\rho = \frac{mM_o - mM_k}{mM_o - 1}$$

$$m = d - \frac{2p + 5}{6} - \frac{2k}{6}$$

$$M_o = \frac{-\ln(|S|)}{p(p-1)/2}$$

$$M_k = \frac{\phi}{((p-k)^2 - p - k)/2}$$

where  $|S|$  is determinant of COV,  $p = \text{NVAR}$ ,  $k = \text{NVAR}$ ,  $\phi$  is the optimized criterion, and  $d = \text{NDF}$ .

### Image Analysis

The term “image analysis” is used here to denote the noniterative image method of Kaiser (1963). It is not the image factor analysis discussed by Harman (1976, page 226). The image method (as well as the alpha factor analysis method) begins with the notion that only a finite number from an infinite number of possible variables have been measured. The image factor pattern is calculated under the assumption that the ratio of the number of factors to the number of observed variables is near zero so that a very good estimate for the unique error variances (for standardized variables) is given as one minus the squared multiple correlation of the variable under consideration with all variables in the covariance matrix.

First, the matrix  $D^2 = (\text{diag}(S^{-1}))^{-1}$  is computed where the operator “diag” results in a matrix consisting of the diagonal elements of its argument, and  $S$  is the sample covariance (correlation) matrix. Then, the eigenvalues  $\Lambda$  and eigenvectors  $\Gamma$  of the matrix  $D^{-1} S D^{-1}$  are computed. Finally, the unrotated image factor pattern matrix is computed as  $\mathbb{A} = D\Gamma[(\Lambda - I)^2 \Lambda^{-1}]^{1/2}$ .

### Alpha Factor Analysis

The alpha factor analysis method of Kaiser and Caffrey (1965) finds factor-loading estimates to maximize the correlation between the factors and the complete universe of variables of interest. The basic idea in this method is as follows: only a finite number of variables out of a much larger set of possible variables is observed. The population factors are linearly related to this larger set while the observed factors are linearly related to the observed variables. Let  $f$  denote the factors obtainable from a finite set of observed random variables, and let  $\xi$  denote the factors obtainable from the universe of observable variables. Then, the alpha method attempts to find factor-loading estimates so as to maximize the correlation between  $f$  and  $\xi$ . In order to obtain these estimates, the iterative algorithm of Kaiser and Caffrey (1965) is used.

### Comments

1. FACTR makes no attempt to solve for NF, the number of factors. In general, if NF is not known in advance, several different values of NF should be used, and the most reasonable value kept in the final solution.
2. The iterative methods are generally thought to be superior from a theoretical point of view but, in practice, often lead to solutions which differ little from the noniterative methods. For this reason, it is usually suggested that a non-iterative method be used in the initial stages of the factor analysis, and that the iterative methods be used when issues such as the number of factors have been resolved.

3. Initial estimates for the unique variances are input when `INIT = 1`. If the iterative methods fail for these values, new initial estimates should be tried. These may be obtained by use of another factoring method (use the final estimates from the new method as initial estimates in the old method).

Another alternative is to let `FACTR` compute initial estimates of the unique error variances. When `INIT = 0`, the initial estimates are taken as a constant

$$\left(1 - \frac{k}{2p}\right)$$

divided by the diagonal elements of the

$$\hat{\Sigma}^{-1}$$

matrix. When the correlation matrix is factor analyzed, this is a constant times one minus the squared multiple correlation coefficient.

### Example

The following data were originally analyzed by Emmett (1949). There are 211 observations on 9 variables. Following Lawley and Maxwell (1971), three factors will be obtained by the method of maximum likelihood.

```

INTEGER      IMTH, INIT, IPRINT, LDA, LDICOV, MAXIT, MAXSTP, NDF,
&            NF, NVAR
REAL         EPS, EPSE
PARAMETER   (EPS=0.000001, EPSE=0.01, IMTH=3, INIT=0, IPRINT=1,
&            LDA=9, LDICOV=9, MAXIT=30, MAXSTP=10, NDF=210, NF=3,
&            NVAR=9)
C
REAL         A(LDA,NF), COV(LDICOV,NVAR), DER(NVAR), EVAL(NVAR),
&            STAT(6), UNIQ(NVAR)
EXTERNAL    FACTR
C
DATA COV/
& 1.000, 0.523, 0.395, 0.471, 0.346, 0.426, 0.576, 0.434, 0.639,
& 0.523, 1.000, 0.479, 0.506, 0.418, 0.462, 0.547, 0.283, 0.645,
& 0.395, 0.479, 1.000, 0.355, 0.270, 0.254, 0.452, 0.219, 0.504,
& 0.471, 0.506, 0.355, 1.000, 0.691, 0.791, 0.443, 0.285, 0.505,
& 0.346, 0.418, 0.270, 0.691, 1.000, 0.679, 0.383, 0.149, 0.409,
& 0.426, 0.462, 0.254, 0.791, 0.679, 1.000, 0.372, 0.314, 0.472,
& 0.576, 0.547, 0.452, 0.443, 0.383, 0.372, 1.000, 0.385, 0.680,
& 0.434, 0.283, 0.219, 0.285, 0.149, 0.314, 0.385, 1.000, 0.470,
& 0.639, 0.645, 0.504, 0.505, 0.409, 0.472, 0.680, 0.470, 1.000/
C
CALL FACTR (NVAR, COV, LDICOV, NF, IMTH, NDF, INIT, MAXIT,
&           MAXSTP, EPS, EPSE, IPRINT, UNIQ, A, LDA, EVAL, STAT,
&           DER)
END

```



## Output

```
Unique Error Variances
  1      2      3      4      5      6      7      8
0.4505  0.4271  0.6166  0.2123  0.3805  0.1769  0.3995  0.4615

  9
0.2309
```

```
Unrotated Loadings
  1      2      3
1  0.6642 -0.3209  0.0735
2  0.6888 -0.2471 -0.1933
3  0.4926 -0.3022 -0.2224
4  0.8372  0.2924 -0.0354
5  0.7050  0.3148 -0.1528
6  0.8187  0.3767  0.1045
7  0.6615 -0.3960 -0.0777
8  0.4579 -0.2955  0.4913
9  0.7657 -0.4274 -0.0117
```

```
Eigenvalues
  1      2      3      4      5      6      7      8      9
0.063  0.229  0.541  0.865  0.894  0.974  1.080  1.117  1.140
```

```
STAT
  1      2      3      4      5
0.0350  1.0000  7.1494  12.0000  0.8476
```

```
  6
5.0000
```

```
Final Parameter Updates
  1      2      3      4      5
2.02042E-07  2.95010E-07  1.80908E-07  6.38808E-08  2.00809E-07

  6      7      8      9
1.48762E-07  1.73797E-08  3.95484E-07  1.42415E-07
```

---

## FROTA/DFROTA (Single/Double precision)

Compute an orthogonal rotation of a factor loading matrix using a generalized orthomax criterion, including quartimax, varimax, and equamax rotations.

### Usage

```
CALL FROTA (NVAR, NF, A, LDA, NORM, MAXIT, W, EPS, B, LDB,
            T, LDT)
```

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of **A** exactly as specified in the dimension statement in the calling program. (Input)

**NORM** — Row normalization option. (Input)

If **NORM** = 1, then row (i.e., Kaiser) normalization is performed. Otherwise, row normalization is not performed.

**MAXIT** — Maximum number of iterations. (Input)

**MAXIT** = 30 is typical. **MAXIT** ≤ 30 defaults to 30 iterations.

**W** — Nonnegative constant used to define the rotation. (Input)

**W** = 0.0 results in quartimax rotations, **W** = 1.0 results in varimax rotations, and **W** = **NF**/2.0 results in equamax rotations. Other nonnegative values of **W** may also be used, but the best values for **W** are in the range (0.0, 5 \* **NF**).

**EPS** — Convergence constant. (Input)

When the relative change in the criterion function is less than **EPS** from one iteration to the next, convergence is assumed. **EPS** = 0.0001 is typical. **EPS** ≤ 0.0 defaults to 0.0001.

**B** — **NVAR** by **NF** matrix of rotated factor loadings. (Output)

If **A** is not needed, **A** and **B** may share the same storage locations.

**LDB** — Leading dimension of **B** exactly as specified in the dimension statement in the calling program. (Input)

**T** — **NF** by **NF** matrix containing the rotation transformation matrix. (Output)

**LDT** — Leading dimension of **T** exactly as specified in the dimension statement in the calling program. (Input)

### Comments

Automatic workspace usage is

**FROTA** **NVAR** units, or  
**DFROTA** 2 \* **NVAR** units.

Workspace may be explicitly provided, if desired, by use of **F2OTA**/**DF2OTA**. The reference is

```
CALL F2OTA (NVAR, NF, A, LDA, NORM, MAXIT, W, EPS, B, LDB,  
           T, LDT, WORK)
```

The additional argument is

**WORK** — Real work vector of length equal to **NVAR**.

### Algorithm

Routine **FROTA** performs an orthogonal rotation according to an orthomax criterion. In this analytic method of rotation, the criterion function

$$Q = \sum_i \sum_r \lambda_{ir}^4 - \frac{\gamma}{p} \sum_r \left[ \sum_i \lambda_{ir}^2 \right]^2$$

is minimized by finding an orthogonal rotation matrix  $T$  such that  $(\lambda_{ij}) = \Lambda = AT$  where  $A$  is the matrix of unrotated factor loadings. Here,  $\gamma \geq 0$  is a user-specified constant ( $W$ ) yielding a family of rotations, and  $p$  is the number of variables.

Kaiser (row) normalization can be performed on the factor loadings prior to rotation via the option parameter `NORM`. In Kaiser normalization, the rows of  $A$  are first “normalized” by dividing each row by the square root of the sum of its squared elements (Harman 1976). After the rotation is complete, each row of  $B$  is “denormalized” by multiplication by its initial normalizing constant.

The method for optimizing  $Q$  proceeds by accumulating simple rotations where a simple rotation is defined to be one in which  $Q$  is optimized for two columns in  $\Lambda$  and for which the requirement that  $T$  be orthogonal is satisfied. A single iteration is defined to be such that each of the  $NF(NF - 1)/2$  possible simple rotations is performed where  $NF$  is the number of factors. When the relative change in  $Q$  from one iteration to the next is less than `EPS` (the user-specified convergence criterion), the algorithm stops. `EPS = 0.0001` is usually sufficient. Alternatively, the algorithm stops when the user-specified maximum number of iterations, `MAXIT`, is reached. `MAXIT = 30` is usually sufficient.

The parameter in the rotation,  $\gamma$ , is used to provide a family of rotations. When  $\gamma = 0.0$ , a direct quartimax rotation results. Other values of  $\gamma$  yield other rotations.

### Example

The example is taken from Emmett (1949) and involves factors derived from nine variables. In this example, the varimax method is chosen with row normalization by using `W = 1.0` and `NORM = 1`, respectively. The results correspond to those given by Lawley and Maxwell (1971, page 84).

```

C      INTEGER   LDA, LDB, LDT, MAXIT, NF, NORM, NVAR
      REAL      EPS, W
      PARAMETER (EPS=0.0, LDA=9, LDB=9, LDT=3, MAXIT=30, NF=3,
&              NORM=1, NVAR=9, W=1.0)
C
      REAL      A(LDA,NF), B(LDB,NF), T(LDT,NF)
      EXTERNAL  FROTA, WRRRN
C
      DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&          .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&          -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&          .1045, -.0778, .4914, -.0117/
C
      CALL FROTA (NVAR, NF, A, LDA, NORM, MAXIT, W, EPS, B, LDB, T,
&              LDT)
C
      CALL WRRRN ('B', NVAR, NF, B, LDB, 0)
      CALL WRRRN ('T', NF, NF, T, LDT, 0)

```

END

### Output

	B		
	1	2	3
1	0.2638	-0.5734	0.3888
2	0.3423	-0.6610	0.1370
3	0.1625	-0.5943	0.0622
4	0.8124	-0.3197	0.1594
5	0.7356	-0.2800	0.0036
6	0.8510	-0.1890	0.2513
7	0.2164	-0.6906	0.2768
8	0.1144	-0.2431	0.6828
9	0.2687	-0.7431	0.3804

	T		
	1	2	3
1	0.7307	-0.5939	0.3367
2	0.6816	0.6623	-0.3112
3	-0.0382	0.4569	0.8887

---

## FOPCS/DFOPCS (Single/Double precision)

Compute an orthogonal Procrustes rotation of a factor-loading matrix using a target matrix.

### Usage

CALL FOPCS (NVAR, NF, A, LDA, X, LDX, B, LDB, T, LDT)

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**X** — NVAR by NF target matrix of the rotation. (Input)

**LDX** — Leading dimension of X exactly as specified in the dimension statement in the calling program. (Input)

**B** — NVAR by NF matrix of rotated factor loadings. (Output)

**LDB** — Leading dimension of B exactly as specified in the dimension statement in the calling program. (Input)

**T** — NF by NF factor rotation matrix. (Output)

**LDT** — Leading dimension of T exactly as specified in the dimension statement in the calling program. (Input)

## Comments

1. Automatic workspace usage is

FOPCS (NF \* (3 \* NF + 4)) - 1 units, or

DFOPCS (NF \* (6 \* NF + 8)) - 2 units.

Workspace may be explicitly provided, if desired, by use of F2PCS/DF2PCS. The reference is

```
CALL F2PCS (NVAR, NF, A, LDA, X, LDX, B, LDB, T,  
           LDT, WK, S)
```

The additional arguments are as follows:

**WK** — Work vector of length  $NF * (2 * NF + 3) - 1$ .

**S** — Work vector of length  $NF * (NF + 1)$ .

2. Informational errors

Type	Code	
------	------	--

4	1	NF = 1. No rotation is possible.
---	---	----------------------------------

4	2	The rank of $A^T * X$ is less than NF.
---	---	--

3. The target matrix is a hypothesized rotated factor loading matrix with loadings chosen (based on prior knowledge) to enhance interpretability. A simple structure solution will have most of the elements in  $X$  near zero or one (for correlation matrix loadings).
4. This routine may also be used to refine a solution obtained by analytic rotation in routine FROTA (page 809). Choose the target matrix so that it closely resembles the analytic solution but modified to have a simple structure.

## Algorithm

Routine FOPCS performs orthogonal Procrustes rotation according to a method proposed by Schöneman (1966). Let  $k = NF$  denote the number of factors,  $p = NVAR$  denote the number of variables,  $A$  denote the  $p \times k$  matrix of unrotated factor loadings,  $T$  denote the  $k \times k$  orthogonal rotation matrix (orthogonality requires that  $T^T T$  be a  $k \times k$  identity matrix), and let  $X$  denote the target matrix. The basic idea in orthogonal Procrustes rotation is to find an orthogonal rotation matrix  $T$  such that  $B = AT$  and  $T$  provides a least-squares fit between the target matrix  $X$  and the rotated loading matrix  $B$ . Schöneman's algorithm proceeds by finding the singular value decomposition of the matrix  $A^T X = U\Sigma V^T$ . The rotation matrix is computed as  $T = UV^T$ .

## Example

The following example is taken from Harman (1976, page 355). It involves the orthogonal Procrustes rotation of an  $8 \times 2$  unrotated factor loading matrix. The original variables are measures of physical features ("lankiness" and

“stockiness”). The target matrix  $X$  is also printed. Note that because different methods are used, Harman (1976) gets slightly different results.

```

INTEGER    LDA, LDB, LDT, LDX, NF, NVAR
PARAMETER  (LDA=8, LDB=8, LDT=2, LDX=8, NF=2, NVAR=8)
C
REAL       A(LDA,NF), B(LDB,NF), T(LDT,NF), X(LDX,NF)
EXTERNAL   FOPCS, WRRRN
C
DATA A/0.856, 0.848, 0.808, 0.831, 0.750, 0.631, 0.569, 0.607,
&      -0.324, -0.412, -0.409, -0.342, 0.571, 0.492, 0.510, 0.351/
DATA X/0.9, 0.9, 0.9, 0.9, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,
&      0.0, 0.9, 0.9, 0.9, 0.9/
C
CALL FOPCS (NVAR, NF, A, LDA, X, LDX, B, LDB, T, LDT)
C
CALL WRRRN ('A', NVAR, NF, A, LDA, 0)
CALL WRRRN ('X', NVAR, NF, X, LDX, 0)
CALL WRRRN ('B', NVAR, NF, B, LDB, 0)
CALL WRRRN ('T', NF, NF, T, LDT, 0)
END

```

### Output

```

      A
      1      2
1  0.8560  -0.3240
2  0.8480  -0.4120
3  0.8080  -0.4090
4  0.8310  -0.3420
5  0.7500   0.5710
6  0.6310   0.4920
7  0.5690   0.5100
8  0.6070   0.3510

```

```

      X
      1      2
1  0.9000   0.0000
2  0.9000   0.0000
3  0.9000   0.0000
4  0.9000   0.0000
5  0.0000   0.9000
6  0.0000   0.9000
7  0.0000   0.9000
8  0.0000   0.9000

```

```

      B
      1      2
1  0.8763   0.264
2  0.9235   0.1896
3  0.8900   0.1677
4  0.8674   0.2348
5  0.2471   0.9096
6  0.2009   0.7745
7  0.1407   0.7510
8  0.2677   0.6481

```

```

      T
      1      2

```

```
1  0.7932  0.6090
2 -0.6090  0.7932
```

---

## FDOBL/DFDOBL (Single/Double precision)

Compute a direct oblimin rotation of a factor loading matrix.

### Usage

```
CALL FDOBL (NVAR, NF, A, LDA, NORM, W, MAXIT, EPS, B, LDB,
           T, LDT, FCOR, LDFCOR)
```

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** —  $NVAR$  by  $NF$  matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of **A** exactly as specified in the dimension statement in the calling program. (Input)

**NORM** — Row normalization option. (Input)

If **NORM** = 1, then row (i.e., Kaiser) normalization is performed. Otherwise, row normalization is not performed.

**W** — Nonpositive constant used to define the rotation. (Input)

**MAXIT** — Maximum number of iterations. (Input)

**MAXIT** = 30 is typical. **MAXIT** = 0 defaults to 30 iterations.

**EPS** — Convergence constant. (Input)

When the relative change in the criterion function is less than **EPS** from one iteration to the next, convergence is assumed. **EPS** = 0.0001 is typical. **EPS** = 0 defaults to 0.0001.

**B** —  $NVAR$  by  $NF$  matrix of rotated factor loadings. (Output)

If **A** is not needed, **A** and **B** may share the same storage locations.

**LDB** — Leading dimension of **B** exactly as specified in the dimension statement in the calling program. (Input)

**T** —  $NF$  by  $NF$  matrix containing the rotation transformation matrix. (Output)

**LDT** — Leading dimension of **T** exactly as specified in the dimension statement in the calling program. (Input)

**FCOR** —  $NF$  by  $NF$  matrix of factor correlations. (Output)

**LDFCOR** — Leading dimension of **FCOR** exactly as specified in the dimension statement in the calling program. (Input)

## Comments

1. Automatic workspace usage is

FDOBL  $NF + 2 * NVAR$  units, or  
DFDOBL  $2 * NF + 4 * NVAR$  units.

Workspace may be explicitly provided, if desired, by use of  
F2OBL/DF2OBL. The reference is

```
CALL F2OBL (NVAR, NF, A, LDA, NORM, W, MAXIT, EPS,  
           B, LDB, T, LDT, FCOR, LDFCOR, WK1, WK2,  
           WK3)
```

The additional arguments are as follows:

**WK1** — Real work vector of length equal to NVAR.

**WK2** — Real work vector of length equal to NF.

**WK3** — Real work vector of length equal to NVAR.

2. Informational errors

Type	Code	
3	1	The algorithm did not converge within MAXIT iterations.
4	1	NF = 1. No rotation is possible.

3. The parameter *w* determines the type of direct OBLIMIN rotation to be performed. In general, *w* must be negative. *w* = 0.0 yields direct quartimin rotation. As *w* approaches negative infinity, the orthogonality among the factors will increase.

## Algorithm

Routine FDOBL performs direct oblimin rotation. In this analytic method of rotation, the criterion function

$$Q = \sum_{r \neq s} \left[ \sum_i \lambda_{ir}^2 \lambda_{is}^2 - \frac{\gamma}{p} \sum_i \lambda_{ir}^2 \sum_i \lambda_{is}^2 \right]$$

is minimized by finding a rotation matrix *T* such that  $(\lambda_{ir}) = \Lambda = AT$  and  $(T^T T)^{-1}$  is a correlation matrix. Here,  $\gamma \leq 0$  is a user-specified constant (*W*) yielding a family of rotations, and *p* is the number of variables. The rotation is said to be direct because it minimizes *Q* with respect to the factor loadings directly, ignoring the reference structure.

Kaiser normalization can be performed on the factor loadings prior to rotation via the option parameter NORM. In Kaiser normalization (see Harman 1976), the rows of *A* are first “normalized” by dividing each row by the square root of the sum of its squared elements. After the rotation is complete, each row of *B* is “denormalized” by multiplication by its initial normalizing constant.



The method for optimizing  $Q$  is essentially the method first proposed by Jennrich and Sampson (1966). It proceeds by accumulating simple rotations where a simple rotation is defined to be one in which  $Q$  is optimized for a given factor in the plane of a second factor, and for which the requirement that

$(T^T T)^{-1}$  be a correlation matrix is satisfied. An iteration is defined to be such that each of the  $NF(NF - 1)$  possible simple rotations is performed, where  $NF$  is the number of factors. When the relative change in  $Q$  from one iteration to the next is less than  $EPS$  (the user-specified convergence criterion), the algorithm stops.  $EPS = .0001$  is usually sufficient. Alternatively, the algorithm stops when the user-specified maximum number of iterations,  $MAXIT$ , is reached.  $MAXIT = 30$  is usually sufficient.

The parameter in the rotation,  $\gamma$ , is used to provide a family of rotations. Harman (1976) recommends that  $\gamma$  be strictly less than or equal to zero. When  $\gamma = 0.0$ , a direct quartimin rotation results. Other values of  $\gamma$  yield other rotations. Harman (1976) suggests that the direct quartimin rotations yield the most highly correlated factors while more orthogonal factors result as  $\gamma$  approaches  $-\infty$ .

### Example

The example is a continuation of the example given in routine FACTR. It involves factors derived from nine variables and uses  $\gamma = -1$ .

```

INTEGER   LDA, LDB, LDFCOR, LDT, MAXIT, NF, NORM, NVAR
REAL      EPS, W
PARAMETER (EPS=0.00001, LDA=9, LDB=9, LDFCOR=3, LDT=3,
&
MAXIT=30, NF=3, NORM=1, NVAR=9, W=-1.0)
C
REAL      A(LDA,NF), B(LDB,NF), FCOR(LDFCOR,NF), T(LDT,NF)
EXTERNAL  FDOBL, WRRRN
C
DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&
.7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&
-.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&
.1045, -.0778, .4914, -.0117/
C
CALL FDOBL (NVAR, NF, A, LDA, NORM, W, MAXIT, EPS, B, LDB, T,
&
LDT, FCOR, LDFCOR)
C
CALL WRRRN ('B', NVAR, NF, B, LDA, 0)
CALL WRRRN ('T', NF, NF, T, LDT, 0)
CALL WRRRN ('FCOR', NF, NF, FCOR, LDFCOR, 0)
END

```

### Output

```

      B
      1      2      3
1  0.1127 -0.5145  0.2917
2  0.1847 -0.6602 -0.0019
3  0.0128 -0.6354 -0.0585
4  0.7797 -0.1751  0.0598
5  0.7147 -0.1813 -0.0959

```

```

6  0.8520  0.0038  0.1820
7  0.0355 -0.6845  0.1509
8  0.0276 -0.0941  0.6824
9  0.0729 -0.7100  0.2493

```

```

      T
      1      2      3
1  0.611  -0.462  0.203
2  0.923   0.813 -0.249
3  0.042   0.728  1.050

```

```

      FCOR
      1      2      3
1  1.000  -0.427  0.217
2 -0.427   1.000 -0.411
3  0.217  -0.411  1.000

```

---

## FPRMX/DFPRMX (Single/Double precision)

Compute an oblique Promax or Procrustes rotation of a factor loading matrix using a target matrix, including pivot and power vector options.

### Usage

```
CALL FPRMX (NVAR, NF, A, LDA, IMTH, NORM, W, MAXIT, EPS, F,
           X, LDX, B, LDB, T, LDT, FCOR, LDFCOR)
```

### Arguments

**NVAR** — Number of variables. (Input)  
 NVAR must be greater than or equal to 2.

**NF** — Number of factors. (Input)  
 NF must be greater than or equal to 2.

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**IMTH** — Method used for rotation. (Input)

#### **IMTH** Method

- 1 The Promax method.
- 2 The pivotal Promax method.
- 3 Oblique Procrustes method.

**NORM** — Normalization option parameter. (Input)  
 NORM = 0 indicates that no row (Kaiser) normalization is to be performed in the orthomax orthogonal rotation. Otherwise, row normalization is performed. Not used when IMTH = 3.

**W** — Constant used to define the orthomax orthogonal rotation. (Input)  
Values for *w* are discussed in the Comments. *w* must be nonnegative. Not used if *IMTH* = 3.

**MAXIT** — Maximum number of iterations. (Input)  
Thirty is typical. Not used if *IMTH* = 3.

**EPS** — Convergence constant for the orthogonal rotation. (Input)  
When the relative change in the orthomax criterion function is less than *EPS* from one iteration to the next, convergence is assumed. *EPS* = 0.0001 is typical. *EPS* nonpositive defaults to *EPS* = 0.0001.

**F** — Vector of length *NF* containing the power vector or the pivot constants depending upon whether *IMTH* = 1 or *IMTH* = 2, respectively. (Input)  
Not used if *IMTH* = 3.

**X** — *NVAR* by *NF* target matrix for the rotation. (Output, if *IMTH* = 1 or 2; input, if *IMTH* = 3)  
For *IMTH* = 1 or 2, *X* is the target matrix derived from the orthomax rotation. For *IMTH* = 3, *X* is input.

**LDX** — Leading dimension of *X* exactly as specified in the dimension statement in the calling program. (Input)

**B** — *NVAR* by *NF* matrix of rotated factor loadings. (Output)

**LDB** — Leading dimension of *B* exactly as specified in the dimension statement in the calling program. (Input)

**T** — *NF* by *NF* factor rotation matrix. (Output)

**LDT** — Leading dimension of *T* exactly as specified in the dimension statement in the calling program. (Input)

**FCOR** — *NF* by *NF* matrix of factor correlations. (Output)

**LDFCOR** — Leading dimension of *FCOR* exactly as specified in the dimension statement in the calling program. (Input)

## Comments

1. Automatic workspace usage is

*FPRMX* (*NVAR* + 4) \* *NF* units, or  
*DFPRMX* (2 \* *NVAR* + 7) \* *NF* units.

Workspace may be explicitly provided, if desired, by use of *F2RMX/DF2RMX*. The reference is

```
CALL F2RMX (NVAR, NF, A, LDA, IMTH, NORM, W, MAXIT,  
           EPS, F, X, LDX, B, LDB, T, LDT, FCOR,  
           LDFCOR, QR, QRAUX, IPVT, WORK)
```

The additional arguments are as follows:

**QR** — Work vector of length *NVAR* \* *NF*.

**QRAUX** — Work vector of length  $NF$ .

**IPVT** — Work vector of length  $NF$ .

**WORK** — Work vector of length  $2 * NF$ .

2. Arguments  $w$ ,  $EPS$ , and  $NORM$  are input arguments to routine `FROTA` (page 809) when  $IMTH = 1$  or  $2$ . (They are not used when  $IMTH = 3$ .) See `FROTA` for common values of  $w$ . Generally,  $w$  can be any positive real number, but the best values lie in the range  $(1.0, 5.0 * NF)$ . Generally, the variances accounted for by the factors approach the same value as  $w$  increases.
3. For  $IMTH = 1$ , all  $F(j)$  should be greater than  $1.0$ , typically  $4.0$ . Generally, the larger the values of  $F(j)$ , the more oblique the solution will be. For  $IMTH = 2$ ,  $F(j)$  should be in the interval  $(0.0, 1.0)$ .
4. When  $IMTH = 3$ , the target matrix,  $x$ , is a hypothesized rotated factor loading matrix based upon prior knowledge with loadings chosen to enhance interpretability. A simple structure solution will have most of the weights  $x(i, j)$  either zero or large in magnitude. Note that the two options  $IMTH = 1$  or  $2$  attempt to achieve this simple structure based upon an initial orthogonal rotation.

### Algorithm

Routine `FPRMX` performs oblique rotations via the Promax, the pivotal Promax, or the oblique Procrustes methods. In all of these methods, a target matrix  $X$  is first either computed or specified by the user. The differences in the methods relate to how the target matrix is first obtained.

Given a  $p \times k$  target matrix,  $X$ , and a  $p \times k$  orthogonal matrix of unrotated factor loadings,  $A$ , compute the rotation matrix  $T$  as follows: First regress each column of  $A$  on  $X$  yielding a  $k \times k$  matrix  $\beta$ . Then, let  $\gamma = \text{diag}(\beta^T \beta)$  where  $\text{diag}$  denotes the diagonal matrix obtained from the diagonal of the square matrix. Standardize  $\beta$  to obtain  $T = \gamma^{-1/2} \beta$ . The rotated loadings are computed as  $B = AT$  while the factor correlations can be computed as the inverse of the  $T^T T$  matrix.

In the Promax method, the unrotated factor loadings are first rotated according to an orthomax criterion via routine `FROTA` (page 809). The target matrix  $X$  is taken as the elements of the  $B$  raised to a power greater than one but retaining the same sign as the original loadings. In `FPRMX`, column  $i$  of the rotated matrix  $B$  is raised to the power  $F(i)$ . A power of four is commonly used. Generally, the larger the power, the more oblique the solution.

In the pivotal Promax method, the unrotated matrix is first rotated to an orthomax orthogonal solution as in the Promax case. Then, rather than raising the  $i$ -th column in  $B$  to the power  $F(i)$ , the elements  $x_{ij}$  of  $X$  are obtained from the elements  $b_{ij}$  of  $B$  by raising the  $ij$  element of  $B$  to the power  $F(i)/b_{ij}$ . This has

the effects of greatly increasing in  $X$  those elements in  $B$  that are greater in magnitude than the pivot elements  $F(i)$ , and of greatly decreasing those elements that are less than  $F(i)$ .

In the oblique Procrustes method, the elements of  $X$  are specified by the user as input to the `FPRMX` routine. No orthogonal rotation is performed in the oblique Procrustes method.

### Example

The following example is a continuation of the example in the `FACTR` (page 801) procedure. It involves nine variables and three factors. The pivotal Promax method is illustrated.

```

INTEGER      IMTH, LDA, LDB, LDFCOR, LDT, LDX, MAXIT, NF, NORM,
&           NVAR
REAL        EPS, W
PARAMETER    (EPS=0.0, IMTH=2, LDA=9, LDB=9, LDFCOR=3, LDT=3,
&           LDX=9, MAXIT=30, NF=3, NORM=1, NVAR=9, W=1.0)
C
REAL        A(LDA,NF), B(LDB,NF), F(NF), FCOR(LDFCOR,NF),
&           T(LDT,NF), X(LDX,NF)
EXTERNAL     FPRMX, WRRRN
C
DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&     .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&     -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&     .1045, -.0778, .4914, -.0117/
C
DATA F/0.5, 0.5, 0.5/
C
CALL FPRMX (NVAR, NF, A, LDA, IMTH, NORM, W, MAXIT, EPS, F, X,
&         LDX, B, LDB, T, LDT, FCOR, LDFCOR)
C
CALL WRRRN ('X', NVAR, NF, X, LDX, 0)
CALL WRRRN ('B', NVAR, NF, B, LDB, 0)
CALL WRRRN ('T', NF, NF, T, LDT, 0)
CALL WRRRN ('FCOR', NF, NF, FCOR, LDFCOR, 0)
END

```

### Output

```

          X
          1      2      3
1  0.0800 -0.6157  0.2967
2  0.2089 -0.7311  0.0007
3  0.0037 -0.6454  0.0000
4  0.8800 -0.1681  0.0032
5  0.8116 -0.1030  0.0000
6  0.9096 -0.0122  0.0640
7  0.0291 -0.7649  0.0982
8  0.0001 -0.0546  0.7563
9  0.0866 -0.8189  0.2807

          B
          1      2      3
1  0.0997 -0.5089  0.3038
2  0.1900 -0.6463  0.0077

```

3	0.0163	-0.6270	-0.0421
4	0.7991	-0.1469	0.0285
5	0.7408	-0.1531	-0.1256
6	0.8668	0.0308	0.1436
7	0.0280	-0.6777	0.1699
8	-0.0094	-0.1017	0.6911
9	0.0611	-0.7031	0.2683

T			
	1	2	3
1	0.617	-0.439	0.189
2	0.963	0.839	-0.318
3	-0.015	0.707	1.039

FCOR			
	1	2	3
1	1.000	-0.464	0.316
2	-0.464	1.000	-0.395
3	0.316	-0.395	1.000

---

## FHARR/DFHARR (Single/Double precision)

Compute an oblique rotation of an unrotated factor loading matrix using the Harris-Kaiser method.

### Usage

```
CALL FHARR (NVAR, NF, A, LDA, NORM, MAXIT, W, C, EPS,
           SCALE, B, LDB, T, LDT, FCOR, LDFCOR)
```

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**NORM** — Row normalization option. (Input)

If **NORM** = 1, then row (i.e., Kaiser) normalization is performed. Otherwise, row normalization is not performed.

**MAXIT** — Maximum number of iterations. (Input)

A typical value is 30.

**W** — Constant used to define the rotation. (Input)

The value of **w** must be nonnegative. See Comments.

**C** — Constant between zero and one used to define the rotation. (Input)

See Comments.

**EPS** — Convergence constant for the rotation angle. (Input)  
EPS = 0.0001 is typical. If EPS is less than or equal to 0.0, then EPS = 0.0001 is used.

**SCALE** — Vector of length NVAR containing a scaling vector. (Input)  
All elements in SCALE should be set to one if principal components or unweighted least squares was used to obtain the unrotated factor loadings. The elements of SCALE should be set to the unique error variances (vector UNIQ in subroutine FACTR) if the principal factor, generalized least squares, maximum likelihood, or the image method was used. Finally, in alpha factor analysis, the elements of SCALE should be set to the communalities (one minus the uniquenesses in standardized data).

**B** — NVAR by NF matrix containing the rotated factor loadings. (Output)

**LDB** — Leading dimension of B exactly as specified in the dimension statement in the calling program. (Input)

**T** — NF by NF factor rotation matrix. (Output)

**LDT** — Leading dimension of T exactly as specified in the dimension statement in the calling program. (Input)

**FCOR** — NF by NF matrix containing the factor correlations. (Output)

**LDFCOR** — Leading dimension of FCOR exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Automatic workspace usage is

FHARR 2 \* NF + NVAR units, or  
DFHARR 4 \* NF + 2 \* NVAR units.

Workspace may be explicitly provided, if desired, by use of F2ARR/DF2ARR. The reference is

```
CALL F2ARR (NVAR, NF, A, LDA, NORM, MAXIT, W, C,  
           EPS, SCALE, B, LDB, T, LDT, FCOR,  
           LDFCOR, RWK1, RWK2)
```

The additional arguments are as follows:

**RWK1** — Real work vector of length equal to 2 \* NF.

**RWK2** — Real work vector of length equal to NVAR.

2. Argument C must be between 0.0 and 1.0. The larger C is, the more orthogonal the rotated factors are. Rarely, should C be greater than 0.5.
3. Arguments W, EPS, and NORM are arguments to routine FROTA (page 809). See FROTA for common values of W in orthogonal rotations. For FHARR, the best values of W are in the range (0.0, 5.0 \* NF). Generally, the variances of the factors converge to the same value as W increases.

## Algorithm

Routine FHARR performs an oblique analytic rotation of unrotated factor loadings via a method proposed by Harris and Kaiser (1964). In this method of rotation, the eigenvectors obtained from the factor extraction are weighted by a factor  $\Delta^{c/2}$  where  $\Delta$  is the diagonal matrix of eigenvalues obtained in the factor extraction and  $c$  is a specified constant. These transformed eigenvectors are then rotated according to an orthomax criterion.

The transformation used to obtain the weighted eigenvectors,  $\Gamma^*$ , from the unrotated loadings,  $A$ , is given as  $\Gamma^* = \Psi^{-1/2} A \Delta^{(c-1)/2}$  where  $\Psi$  is the matrix of unique error variances output by routine FACTR (page 801). The matrix should be set to an identity matrix if the principal component, unweighted least squares, or alpha factor analysis method is used in routine FACTR to obtain the unrotated factor loadings (IMTH = 0,1, or 5). This is required because in these methods of factor analysis, the eigenvectors are not premultiplied by a diagonal matrix when obtaining the unrotated factor loadings.

After  $\Gamma^*$  has been computed, it is rotated according to a user-selected orthomax criterion. The member of the orthomax family to be used is selected via a constant  $w$ . (See the description of routine FROTA, page 809.) Because  $\Gamma^*$  is used in place of  $A$  (the unrotated factor loadings in routine FROTA), the matrix resulting from the rotation is (after standardizing by preand postmultiplication by the diagonal matrices  $U^{-1}$  and  $\Delta^{1-c}$ ) a matrix of obliquely rotated loadings.

Note that the effect of  $w$  is less pronounced than the effect of  $c$ . Using  $c = 1.0$  yields an orthogonal orthomax rotation while  $c = 0.0$  yields the most oblique factors. A common choice for  $c$  is given by  $c = 0.5$ . One good choice for  $w$  is 1.0.  $w = 1.0$  yields a varimax rotation on the weighted eigenvectors.

## Example

The example is a continuation of the example in routine FACTR (page 801). It involves 9 variables. A rotation with row normalization and 3 factors is performed.

```
INTEGER      LDA, LDB, LDFCOR, LDT, MAXIT, NF, NORM, NVAR
REAL         C, EPS, W
PARAMETER   (C=0.5, EPS=0.0001, LDA=9, LDB=9, LDFCOR=3, LDT=3,
&           MAXIT=30, NF=3, NORM=1, NVAR=9, W=1.0)
C
REAL         A(LDA,NF), B(LDB,NF), FCOR(LDFCOR,NF), SCALE(NVAR),
&           T(LDT,NF)
EXTERNAL    FHARR, WRRRN
C
DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&     .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&     -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&     .1045, -.0778, .4914, -.0117/
C
DATA SCALE/.4505, .4271, .6165, .2123, .3805, .1769, .3995,
&     .4616, .2309/
C
```



```

CALL FHARR (NVAR, NF, A, LDA, NORM, MAXIT, W, C, EPS, SCALE, B,
&          LDB, T, LDT, FCOR, LDFCOR)
C
CALL WRRRN ('B', NVAR, NF, B, LDB, 0)
CALL WRRRN ('T', NF, NF, T, LDT, 0)
CALL WRRRN ('FCOR', NF, NF, FCOR, LDFCOR, 0)
END

```

### Output

```

      B
      1      2      3
1  0.1542 -0.5103  0.2749
2  0.2470 -0.6477 -0.0233
3  0.0744 -0.6185 -0.0750
4  0.7934 -0.1897  0.0363
5  0.7329 -0.1909 -0.1175
6  0.8456 -0.0194  0.1610
7  0.0966 -0.6713  0.1320
8  0.0198 -0.1067  0.6773
9  0.1340 -0.6991  0.2285

```

```

      T
      1      2      3
1  0.649 -0.469  0.175
2  0.850  0.777 -0.249
3 -0.053  0.687  1.065

```

```

      FCOR
      1      2      3
1  1.000 -0.335  0.250
2 -0.335  1.000 -0.413
3  0.250 -0.413  1.000

```

---

## FGCRF/DFGCRF (Single/Double precision)

Compute direct oblique rotation according to a generalized fourth-degree polynomial criterion.

### Usage

```

CALL FGCRF (NVAR, NF, A, LDA, NORM, W, MAXIT, EPS, B, LDB,
            T, LDT, FCOR, LDFCOR)

```

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**NORM** — Row normalization option. (Input)

If  $NORM = 1$ , then row (i.e., Kaiser) normalization is performed. If  $NORM = 0$ , row normalization is not performed.

**W** — Vector of length 4 containing the constants  $\omega_1, \omega_2, \omega_3, \omega_4$  necessary to define the rotation. (Input)

Some common rotations are

Rotation	w(1)	w(2)	w(3)	w(4)
Quartimin	0	1	0	-1
Covarimin	$-1/NVAR$	1	$1/NVAR$	-1
Oblimin	$-\gamma/NVAR$	1	$\gamma/NVAR$	-1
Crawford-Ferguson	0	$K_1$	$K_2$	$-K_1 - K_2$

where  $K_1, K_2$ , and  $\gamma$  are constants (determined by the user).

**MAXIT** — Maximum number of iterations. (Input)

$MAXIT = 30$  is typical.  $MAXIT \leq 30$  defaults to 30 iterations.

**EPS** — Convergence constant. (Input)

When the relative change in the criterion function is less than  $EPS$  from one iteration to the next, convergence is assumed.  $EPS = 0.0001$  is typical.  $EPS \leq 0.0$  defaults to 0.0001.

**B** —  $NVAR$  by  $NF$  matrix of rotated factor loadings. (Output)

If **A** is not needed, **A** and **B** can share the same storage locations.

**LDB** — Leading dimension of **B** exactly as specified in the dimension statement in the calling program. (Input)

**T** —  $NF$  by  $NF$  matrix containing the rotation transformation matrix. (Output)

**LDT** — Leading dimension of **T** exactly as specified in the dimension statement in the calling program. (Input)

**FCOR** —  $NF$  by  $NF$  matrix of factor correlations. (Output)

**LDFCOR** — Leading dimension of **FCOR** exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Automatic workspace usage is

$FGCRF \quad NVAR * (NF + 2) + NF^2$  units, or

$DFGCRF \quad 2 * (NVAR * (NF + 2) + NF^2)$  units.

Workspace may be explicitly provided, if desired, by use of

$F2CRF/DF2CRF$ . The reference is

CALL F2CRF (NVAR, NF, A, LDA, NORM, W, MAXIT, EPS,  
 B, LDB, T, LDT, FCOR, LDFCOR, RWK1,  
 RWK2, RWK3)

The additional arguments are as follows:

**RWK1** — Work vector of length NVAR.

**RWK2** — Work vector of length NVAR \* (NF + 1).

**RWK3** — Work vector of length NF<sup>2</sup>.

2. Informational error

Type	Code	
3	1	The algorithm did not converge within MAXIT iterations.

**Algorithm**

Routine FGCRF performs direct oblique factor rotation for an arbitrary fourth-degree polynomial criterion function. Let  $p = \text{NVAR}$  denote the number of variables, and let  $k = \text{NF}$  denote the number of factors. Then, the criterion function

$$Q = \omega_1 \left( \sum_{i=1}^p \sum_{r=1}^k \lambda_{ir}^2 \right)^2 + \omega_2 \sum_{i=1}^p \left( \sum_{r=1}^k \lambda_{ir}^2 \right)^2 + \omega_3 \sum_{r=1}^k \left( \sum_{i=1}^p \lambda_{ir}^2 \right)^2 + \omega_4 \sum_{i=1}^p \sum_{r=1}^k \lambda_{ir}^4$$

is minimized by finding a rotation matrix  $T$  such that  $(\lambda_{ij}) = \Lambda = AT$  and  $T^{-1} (T^{-1})^T$  is a correlation matrix. Here,  $\omega_i = w(i)$ ,  $i = 1, \dots, 4$  are user specified constants. The rotation is said to be direct because it minimizes  $Q$  with respect to the factor loadings directly, ignoring the reference structure (see, e.g., Harman, 1976).

Kaiser normalization (Harman, 1976) is specified when option parameter  $\text{NORM} = 1$ . When Kaiser normalization is performed, the rows of  $A$  are first “normalized” by dividing each row by the square root of the sum of its squared elements. The rotation is then performed. The rows of  $B$  are then “denormalized” by multiplying each row by the initial row normalizing constant.

The criterion function  $Q$  was first proposed by Jennrich (1973). It generalizes the oblimin criterion function and the criterion function proposed by Crawford and Ferguson (1970) to an arbitrary fourth degree criterion.  $Q$  is optimized by accumulating simple rotations where a simple rotation is defined to be an optimal factor rotation (with respect to  $Q$ ) for two columns of  $\Lambda$ , and for which the requirement that  $T^{-1} (T^{-1})^T$  be a correlation matrix is satisfied. FGCRF

determines the optimal simple rotation by finding the roots of a cubic polynomial equation. The details are contained in Clarkson and Jennrich (1988).

**Table 1: Specific Criteria in the General Symmetric Family**

Criterion	$\omega_1$	$\omega_2$	$\omega_3$	$\omega_4$
Quartimin	0	1	0	-1
Covarimin	$-1/p$	1	$1/p$	-1
Oblimin	$-\gamma/p$	1	$\gamma/p$	-1
Crawford-Ferguson	0	$K_1$	$K_2$	$-K_1 - K_2$

An iteration is complete after all possible  $k(k - 1)$  simple rotations have been performed. When the relative change in  $Q$  from one iteration to the next is less than  $EPS$ , the algorithm stops.  $EPS = .0001$  is usually sufficient. Alternatively, the algorithm stops when the user specified maximum number of iterations,  $MAXIT$ , is reached.  $MAXIT = 30$  is typical.

### Notes

The parameters in the rotation,  $\omega_1$ , provide for a two-dimensional family of rotations. When  $\omega_1 = -\gamma/p$ ,  $\omega_2 = 1$ ,  $\omega_3 = \gamma/p$ , and  $\omega_4 = -1$ , then a direct oblimin rotation with parameter  $\gamma$  is performed. Direct oblimin rotations are also performed by routine  $FDOBL$  (page 815), which is somewhat faster. For  $\omega_1 = 0$ ,  $\omega_2 = K_1$ ,  $\omega_3 = K_2$ , and  $\omega_4 = -(K_1 + K_2)$  direct Crawford-Ferguson rotation with parameters  $K_1$  and  $K_2$  results (see Crawford and Ferguson 1970, or Clarkson and Jennrich 1988). Other values of  $\omega$  yield other rotations. Common values for  $\omega$  are as in Table 1.

### Example

The example is a continuation of the example in routine  $FACTR$  (page 801). It involves nine variables. A Crawford-Ferguson rotation with row normalization and 3 factors is performed.

```

INTEGER   LDA, LDB, LDFCOR, LDT, MAXIT, NF, NORM, NVAR
REAL      EPS
PARAMETER (EPS=0.0001, LDA=9, LDB=9, LDFCOR=3, LDT=3, MAXIT=30,
&          NF=3, NORM=1, NVAR=9)
C
REAL      A(LDA,NF), B(LDB,NF), FCOR(LDFCOR,NF), T(LDT,NF), W(4)
EXTERNAL  FGCRF, WRRRN
C
DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&      .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&      -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&      .1045, -.0778, .4914, -.0117/
DATA W/0.0, 7.0, 1.0, -8.0/
C

```

```

CALL FGCRF (NVAR, NF, A, LDA, NORM, W, MAXIT, EPS, B, LDB, T,
&          LDT, FCOR, LDFCOR)
C
CALL WRRRN ('B', NVAR, NF, B, LDA, 0)
CALL WRRRN ('T', NF, NF, T, LDT, 0)
CALL WRRRN ('FCOR', NF, NF, FCOR, LDFCOR, 0)
END

```

### Output

```

      B
      1      2      3
1  0.1156 -0.3875  0.3992
2  0.2164 -0.5829  0.0924
3  0.0426 -0.5858  0.0263
4  0.8052 -0.0903  0.0887
5  0.7497 -0.1370 -0.0838
6  0.8638  0.1047  0.1990
7  0.0529 -0.5792  0.2670
8 -0.0166  0.0776  0.7748
9  0.0854 -0.5765  0.3803

```

```

      T
      1      2      3
1  0.633 -0.327  0.290
2  0.935  0.738 -0.399
3 -0.061  0.907  1.066

```

```

      FCOR
      1      2      3
1  1.000 -0.434  0.366
2 -0.434  1.000 -0.498
3  0.366 -0.498  1.000

```

---

## FIMAG/DFIMAG (Single/Double precision)

Compute the image transformation matrix.

### Usage

```
CALL FIMAG (NF, T, LDT, TI, LDTI)
```

### Arguments

*NF* — Number of factors. (Input)

*T* — *NF* by *NF* transformation matrix. (Input)

*LDT* — Leading dimension of *T* exactly as specified in the dimension statement in the calling program. (Input)

*TI* — *NF* by *NF* image transformation matrix. (Output)

*LDTI* — Leading dimension of *TI* exactly as specified in the dimension statement in the calling program. (Input)

## Comments

1. Automatic workspace usage is

FIMAG  $2 * NF + NF(NF - 1)/2$  units, or

DFIMAG  $3 * NF + NF(NF - 1)$  units.

Workspace may be explicitly provided, if desired, by use of F2MAG/DF2MAG. The reference is

```
CALL F2MAG (NF, T, LDT, TI, LD TI, RWK, IWK)
```

The additional arguments are as follows:

**RWK** — Real work vector of length  $NF + NF(NF - 1)/2$ .

**IWK** — Integer work vector of length  $NF$ .

2. Informational Error

Type	Code
------	------

3	1	T is ill-conditioned. The solution may not be accurate.
---	---	---

## Algorithm

Routine FIMAG computes the image transformation matrix **TI** from the factor rotation matrix ( $T$ ). The image transformation matrix takes the unrotated factor loadings into the factor structure matrix when the unrotated loadings are computed from a correlation matrix. It is computed as the inverse of the transpose of the factor rotation matrix  $T$ . When orthogonal rotations are used,  $(T^T)^{-1} = T$  so there is no reason to compute the image transformation matrix.

## Example

This example is a continuation of the example contained in the manual document for routine FROTA (page 809). The image transformation matrix is obtained from the orthogonal rotation matrix. Some small differences between the matrix **TI** when compared with the matrix **T** computed via routine FROTA can be seen. These differences are because of roundoff error since for orthogonal rotations, the image transformation matrix is the same as the rotation matrix.

```
INTEGER      LDT, LD TI, NF
PARAMETER   (LDT=3, LD TI=3, NF=3)
C
REAL        T(LDT,NF), TI(LD TI,NF)
EXTERNAL    FIMAG, WRRRN
C
DATA T/.7307, .6816, -.0382, -.5939, .6623, .4569, .3367,
&        -.3112, .8887/
C
CALL FIMAG (NF, T, LDT, TI, LD TI)
C
CALL WRRRN ('TI', NF, NF, TI, LD TI, 0)
END
```

<b>Output</b>			
	T1		
	1	2	3
1	0.7307	-0.5938	0.3367
2	0.6816	0.6622	-0.3112
3	-0.0382	0.4569	0.8887

---

## FRVAR/DFRVAR (Single/Double precision)

Compute the factor structure and the variance explained by each factor.

### Usage

CALL FRVAR (NVAR, NF, A, LDA, T, LDT, VAR, S, LDS, FVAR)

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**T** — NF by NF factor rotation matrix. (Input)

**LDT** — Leading dimension of T exactly as specified in the dimension statement in the calling program. (Input)

**VAR** — Vector of length NVAR containing the variances of the original variables. (Input)

If standardized variables were used (i.e., the loadings are from a correlation matrix), then set VAR(1) to any negative number. In this case, VAR may be dimensioned of length one.

**S** — NVAR by NF factor structure matrix. (Output)

**LDS** — Leading dimension of S exactly as specified in the dimension statement in the calling program. (Input)

**FVAR** — Vector of length NF containing the variance accounted for by each of the NF rotated factors. (Output)

### Comments

Automatic workspace usage is

FRVAR  $NF^2 + NF * (1 + NVAR) + NF$  units, or

DFRVAR  $2 * NF^2 + 2 * NF * (1 + NVAR) + NF$  units.

Workspace may be explicitly provided, if desired, by use of F2VAR/DF2VAR. The reference is

```
CALL F2VAR (NVAR, NF, A, LDA, T, LDT, VAR, S, LDS, FVAR,
           TINV, WK, IWK)
```

The additional arguments are as follows:

**TINV** — Work vector of length  $NF^2$ .

**WK** — Work vector of  $NF * (1 + NVAR)$ .

**IWK** — Work vector of length  $NF$ .

### Algorithm

Routine FRVAR computes the factor structure matrix (the matrix of correlations between the observed variables and the hypothesized factors) and the variance explained by each of the factors (for orthogonal rotations). For oblique rotations, FRVAR computes a measure of the importance of the factors, the sum of the squared elements in each column.

Let  $\Delta$  denote the diagonal matrix containing the elements of the vector VAR along its diagonal. The estimated factor structure matrix  $S$  is computed as

$$S = \Delta^{-\frac{1}{2}} A(T^{-1})^T$$

while the elements of FVAR are computed as the diagonal elements of

$$S^T \Delta^{\frac{1}{2}} AT$$

If the factors were obtained from a correlation matrix (or the factor variances for standardized variables are desired), then the elements of the vector VAR should either all be 1.0, or the first element of VAR should be set to any negative number. In either case, variances of 1.0 are used.

The user should be careful to input the unrotated loadings. When obliquely rotated loadings are input, the output vector FVAR contains a measure of each factors importance, but it does not contain the variance of each factor.

### Example

The following example illustrates the use of routine FRVAR when the structure and an index of factor importance for obliquely rotated loadings (obtained from routine FDOBL, page 815) are desired. Note in this example that the elements of FVAR are not variances since the rotation is oblique.

```

C      INTEGER    LDA, LDS, LDT, NF, NVAR
      PARAMETER  (LDA=9, LDS=9, LDT=3, NF=3, NVAR=9)
C
      REAL       A(LDA,NF), FVAR(NF), S(LDS,NF), T(LDT,NF), VAR(NVAR)
      EXTERNAL   FRVAR, WRRRN
C
      DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&          .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&          -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&          .1045, -.0778, .4914, -.0117/
C
      DATA T/0.611, 0.923, 0.042, -0.462, 0.813, 0.728, 0.203, -0.249,
&          1.050/
```



```

C      DATA VAR/9*1.0/
C
C      CALL FRVAR (NVAR, NF, A, LDA, T, LDT, VAR, S, LDS, FVAR)
C
C      CALL WRRRN ('S', NVAR, NF, S, LDS, 0)
C      CALL WRRRN ('FVAR', 1, NF, FVAR, 1, 0)
C      END

```

### Output

```

      S
      1      2      3
1  0.3958 -0.6825  0.5274
2  0.4662 -0.7385  0.3093
3  0.2715 -0.6171  0.2052
4  0.8673 -0.5328  0.3010
5  0.7712 -0.4473  0.1338
6  0.8897 -0.4348  0.3654
7  0.3606 -0.7618  0.4397
8  0.2160 -0.3860  0.7270
9  0.4303 -0.8437  0.5566

```

```

      FVAR
      1      2      3
2.170  2.559  0.915

```

---

## FCOEF/DFCOEF (Single/Double precision)

Compute a matrix of factor score coefficients for input to the routine FSCOR.

### Usage

```

CALL FCOEF (NVAR, NF, A, LDA, IMTH, COV, LDCOV, T,
           LDT, SCOEF, LDSCOE)

```

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF matrix of unrotated factor loadings. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**IMTH** — Method to be used to obtain the factor scores. (Input)

IMTH	Method
1	Regression method
2	Least squares method
3	Bartlett method
4	Anderson and Rubin method
5	Image score for image analysis

See the Comments for a table of the methods that are appropriate for a given type of factor extraction and rotation.

**COV** — The variance-covariance or correlation matrix of order *NVAR* from which the factor loadings were obtained. (Input)

*COV* is not used and may be dimensioned of length 1 if *IMTH* = 2 or 5.

**LDCOV** — Leading dimension of *COV* exactly as specified in the dimension statement in the calling program. (Input)

**T** — *NF* by *NF* factor rotation matrix or transformation matrix. (Input)

If the image method is being used, then routine *FIMAG* (page 829) needs to be called after the rotation routine to obtain the image transformation matrix. *TI* is then input for *T* in *FCOEF*. If factor score coefficients for the unrotated loadings are desired, *T* should be set to the identity matrix prior to calling *FCOEF*.

**LDT** — Leading dimension of *T* exactly as specified in the dimension statement in the calling program. (Input)

**SCOEF** — *NVAR* by *NF* factor score coefficient matrix. (Output)

**LDSCOE** — Leading dimension of *SCOEF* exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1 Automatic workspace usage for *FCOEF* depends on *IMTH*.

<b>IMTH</b>	<b>Usage</b>
1	$NVAR * (NVAR + NF) + NF + 2$ units
2,3	$NF * (NVAR + NF) + NVAR + NF + 1$ units
4	$NVAR * (NVAR + 2 * NF + 1) + NF * (NF + 1)$ units
5	$NVAR * NF + 4$ units

Workspace for *DFCOEF* is twice the workspace required for *FCOEF*.  
Workspace may be explicitly provided, if desired, by use of  
*F2OEF/DF2OEF*. The reference is

```
CALL F2OEF (NVAR, NF, A, LDA, IMTH, COV, LDCOV, T,  
           LDT, SCOEF, LDSCOE, B, RWK1, S, UNIQ,  
           RWK2)
```

The additional arguments are as follows:

**B** — Real work vector of length  $2 * NVAR * NF$  if *IMTH* = 4, and of length *NVAR* \* *NF* otherwise.

**RWK1** — Real work vector of length  $NVAR^2$  if *IMTH* = 1 or 4, and of length  $NF^2$  if *IMTH* = 2 or 3. Otherwise, *RWK1* is of length 1.

**S** — Real work vector of length  $NF^2$  if *IMTH* = 4. Otherwise, *S* is dimensioned of length 1.

**UNIQ** — Real work vector of length *NVAR* if *IMTH* = 2, 3, or 4. Otherwise, *UNIQ* is dimensioned of length 1.

**RWK2** — Real work vector of length *NF* if *IMTH* is not 5. If *IMTH* = 5, then *RWK2* is of length 1.

- The method used for computing the factor score coefficients depends both upon the method used to extract the factor loadings in routine *FACTR* (page 801) and whether the factor loadings were orthogonally or obliquely rotated. In the following table, the numbers in parentheses refer to *IMTH* in routine *FACTR* and the numbers in the cells refer to *IMTH* in *F*COEF.

<b>FACTR Method (IMTH)</b>	<b>No Rotation</b>	<b>Orthogonal Rotation</b>	<b>Oblique Rotation</b>
Component (0)	1, 2	1, 2	1, 2
Image (4)	5	5	5
Common Factor			
ULS (1)	1, 2, 3, 4	1, 2, 3, 4	1, 2, 3, 4
GLS (2)	1, 2, 3, 4	1, 2, 3, 4	1, 2, 3, 4
ML (3)	1, 2, 3, 4	1, 2, 3, 4	1, 2, 3, 4
Alpha (5)	1, 2, 3, 4	1, 2, 3, 4	1, 2, 3, 4

### Algorithm

Routine *F*COEF computes factor score coefficients that may subsequently be used in computing the factor scores in routine *F*SCOR (page 838). Five options for computing the coefficients are available according to the input parameter *IMTH*. The method that should be used depends upon the method used in extracting the factor loadings. See the Comments section above for values to use for *IMTH* when various methods of factor extraction are used.

Let *S* denote the covariance (or correlation) matrix from which the factors were obtained, let  $\beta$  denote the factor score coefficients, let  $U^2 = \text{diag}(S - AA^T)$  denote the unique error variances, and let  $B = AT$  denote the rotated factor loadings (if coefficients for the unrotated loadings are desired, then  $B = A$ ). The various methods for computing the factor score coefficients are discussed in detail in Harman (1976, Chapter 16) and are given as follows:

- The regression method may be used with any method of factor extraction and rotation (but not with image analysis). The coefficients are computed as follows:

$$\hat{\beta} = S^{-1}B(T^T T)^{-1}$$

- The least-squares method may also be used with any method of factor extraction and rotation (but not in image analysis). The factor score coefficients are computed as

$$\hat{\beta} = B(B^T B)^{-1}$$

Note that estimated coefficients in the least-squares method yield different factor scores depending upon the scale of the observed variables. In particular, factor scores computed from standardized data (i.e., for the correlation matrix) will be different from factor scores computed from the raw data (i.e., from a covariance matrix). Generally, the differences will not be great. These differences are not observed in any of the other methods.

3. The Bartlett (1937) method may be used with common factor models only. The coefficients are computed as

$$\hat{\beta} = U^{-2} B(B^T U^{-2} B)^{-1}$$

4. The Anderson and Rubin (1956) method may also be used with common factor models only. It is a modification of the Bartlett method where the modification is used to insure that the factors obtained are orthogonal. The factor score coefficients are computed as

$$\hat{\beta} = U^{-2} B(B^T U^{-2} S U^{-2} B)^{-\frac{1}{2}}$$

5. The image method is appropriate for image analysis. In this method, the coefficients are computed as

$$\hat{\beta} = B_I T_I = A(T^T)^{-1} (T^T)^{-1}$$

where  $B_I$  is the image score coefficient matrix, and  $T_I$  is the image transformation matrix (the matrix `TI` in routine `FIMAG`, page 829).

Harman (1976, pages 385-387) discusses choosing a method for computing factor score coefficients. According to Harman, the most desirable properties of any of the methods can be summarized as follows.

- Validity—The estimated factor scores should have high correlation with the population factor scores.
- Orthogonality—The estimated factor scores should not correlate highly with one another.
- Univocal—The estimated factor scores should correlate only with the corresponding true factor scores.

With these criteria in mind, Harman states that:

1. The regression method yields factor scores which usually have the highest correlation with the true factor scores.
2. The Bartlett and least-squares methods are univocal but not orthogonal.
3. The Anderson and Rubin method is orthogonal but not univocal.
4. Univocality is of more significance than orthogonality.

## Example

In the following example, the regression method is used to obtain estimated factor score coefficients for a 9-variable problem with 3 factors. An oblique rotation method was used with the maximum likelihood common factor model to obtain the factor loadings. Routine FDOBL (page 815) was used to obtain the oblique factor loadings.

```
INTEGER    IMTH, LDA, LDCOV, LDSCOE, LDT, NF, NVAR
PARAMETER  (IMTH=1, LDA=9, LDCOV=9, LDSCOE=9, LDT=3, NF=3,
&          NVAR=9)
C
REAL       A(LDA,NF), COV(LDCOV,NVAR), SCOEFLDSCOE,NF),
&          T(LDT,NF)
EXTERNAL   FCOEF, WRRRN
C
DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&      .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&      -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&      .1045, -.0778, .4914, -.0117/
C
DATA T/0.611, 0.923, 0.042, -0.462, 0.813, 0.728, 0.203, -0.249,
&      1.050/
C
DATA COV/1.000, 0.523, 0.395, 0.471, 0.346, 0.426, 0.576, 0.434,
&      0.639, 0.523, 1.000, 0.479, 0.506, 0.418, 0.462, 0.547,
&      0.283, 0.645, 0.395, 0.479, 1.000, 0.355, 0.270, 0.254,
&      0.452, 0.219, 0.504, 0.471, 0.506, 0.355, 1.000, 0.691,
&      0.791, 0.443, 0.285, 0.505, 0.346, 0.418, 0.270, 0.691,
&      1.000, 0.679, 0.383, 0.149, 0.409, 0.426, 0.462, 0.254,
&      0.791, 0.679, 1.000, 0.372, 0.314, 0.472, 0.576, 0.547,
&      0.452, 0.443, 0.383, 0.372, 1.000, 0.385, 0.680, 0.434,
&      0.283, 0.219, 0.285, 0.149, 0.314, 0.385, 1.000, 0.470,
&      0.639, 0.645, 0.504, 0.505, 0.409, 0.472, 0.680, 0.470,
&      1.000/
C
CALL FCOEF (NVAR, NF, A, LDA, IMTH, COV, LDCOV, T, LDT, SCOEFLDSCOE),
&          LDSCOE)
C
CALL WRRRN ('SCOEFL', NVAR, NF, SCOEFLDSCOE, 0)
END
```

## Output

```
          SCOEFL
          1          2          3
1 -0.0102 -0.1350  0.1781
2  0.0269 -0.2191 -0.0825
3 -0.0080 -0.1536 -0.0791
4  0.3788 -0.0597 -0.0596
5  0.2067 -0.0554 -0.1768
6  0.4885  0.1103  0.2084
7 -0.0258 -0.2317  0.0612
8 -0.0474  0.0345  0.5269
9 -0.0431 -0.3967  0.2507
```

---

## FSCOR/DFSCOR (Single/Double precision)

Compute a set of factor scores given the factor score coefficient matrix.

### Usage

```
CALL FSCOR (NVAR, NF, SCOE, LDSCOE, NOBS, X, LDX, XBAR,  
           STD, SCOR, LDSCOR)
```

### Arguments

**NVAR** — Number of variables. (Input)

**NF** — Number of factors. (Input)

**SCOE** — NVAR by NF matrix containing the factor score coefficients as output from routine FCOEF (page 833). (Input)

**LDSCOE** — Leading dimension of SCOE exactly as specified in the dimension statement in the calling program. (Input)

**NOBS** — Number of observations for which factor scores are to be computed. (Input)

**X** — NOBS by NVAR data matrix for which factor scores are to be computed. (Input)

**LDX** — Leading dimension of X exactly as specified in the dimension statement in the calling program. (Input)

**XBAR** — Vector of length NVAR containing the means of the NVAR variables. (Input)

**STD** — Vector of length NVAR containing the standard deviations of the NVAR variables. (Input)

If STD(1) is not positive, then it is assumed that the factor score coefficients are from a covariance matrix and the observed variables are not standardized to unit variance.

**SCOR** — NOBS by NF matrix containing the factor scores. (Output)  
If X is not needed, X and SCOR can share the same memory locations.

**LDSCOR** — Leading dimension of SCOR exactly as specified in the dimension statement in the calling program. (Input)

### Comments

Automatic workspace usage is

```
FSCOR NVAR units, or  
DFSCOR 2 * NVAR units.
```

Workspace may be explicitly provided, if desired, by use of F2COR/DF2COR. The reference is

```
CALL F2COR (NVAR, NF, SCOE, LDSCOE, NOBS, X, LDX, XBAR,
           STD, SCOR, LDSCOR, WK)
```

The additional argument is

**WK** — Work vector of length NVAR.

### Algorithm

Routine FSCOR computes the factor scores from the factor score coefficient matrix. In FSCOR, the data are input as originally observed, and standardization is performed as required according to the value of STD(1). When the factor loadings are computed from the correlation matrix, the observed data must be standardized to a mean of zero and a variance of one prior to computing the factor scores. This requires that STD contain the observed standard deviations of the observed data and that XBAR contain the means. On the other hand, if the factor loadings are computed from the covariance matrix, then the observed data must be standardized to a mean of zero, but the variance must be left unchanged in computing the factor scores. In this case, STD(1) must be negative or zero.

After standardizing the observed data, the factor scores are computed as the product of the factor score coefficient matrix times the standardized data. If factor scores are computed from the same data from which the covariance matrix was computed, then the sample variance (using weights and frequencies as required) of the resulting factor scores will be 1.0.

### Example

The following example is a continuation of the example given in the manual document for routine FACTR (page 801). The rotated loadings are those obtained from the manual document for routine FROTA (page 809), and the factor score coefficients are as described in the manual document for routine FCOEF (page 833).

```

C      INTEGER      LDSCOE, LDSCOR, LDX, NF, NOBS, NVAR
C      PARAMETER   (LDSCOE=2, LDSCOR=5, LDX=5, NF=1, NOBS=5, NVAR=2)
C
C      REAL        SCOE(NVAR,NF), SCOR(LDSCOR,NF), STD(NVAR),
C      &           X(LDX,NVAR), XBAR(NVAR)
C      EXTERNAL    FSCOR, WRRRN
C
C      DATA X/40.0, 60.0, 30.0, 15.0, 45.0, 3.0, 9.0, 2.0, 0.0, 4.0/
C      DATA SCOE/0.33563, 0.33562/
C      DATA XBAR/38.0, 3.6/, STD/16.80774, 3.361547/
C
C      CALL FSCOR (NVAR, NF, SCOE, LDSCOE, NOBS, X, LDX, XBAR, STD,
C      &           SCOR, LDSCOR)
C
C      CALL WRRRN ('Factor Scores', NOBS, NF, SCOR, LDSCOR, 0)
C      END

```

### Output

```
Factor Scores
1 -0.0200
```

2 0.9785  
3 -0.3195  
4 -0.8187  
5 0.1797

---

## FRESI/DFRESI (Single/Double precision)

Compute communalities and the standardized factor residual correlation matrix.

### Usage

CALL FRESI (NVAR, COV, LDCOV, NF, A, LDA, Y, RESID, LDRESI)

### Arguments

**NVAR** — Number of variables. (Input)

**COV** — NVAR by NVAR matrix containing the variance-covariance or correlation matrix. (Input)

Only the upper triangular part of COV is referenced.

**LDCOV** — Leading dimension of COV exactly as specified in the dimension statement in the calling program. (Input)

**NF** — Number of factors. (Input)

**A** — NVAR by NF orthogonal factor-loading matrix. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**Y** — Vector of length NVAR containing the communalities. (Output)

**RESID** — NVAR by NVAR matrix containing the normalized residual variance-covariance or correlation matrix. (Output)

**LDRESI** — Leading dimension of RESID exactly as specified in the dimension statement in the calling program. (Input)

### Algorithm

Routine FRESI computes the communalities and a standardized residual covariance/correlation matrix for input covariance/correlation matrix COV. The user must also input the orthogonal (unrotated) factor loadings, A, obtained from the matrix COV. Let  $a_i$  denote the  $i$ -th row of matrix A. Then, the communalities are given as

$$y_i = a_i a_i^T$$

where  $y_i$  is the  $i$ -th communality. The residual covariance/correlation matrix is given by

$$r_{ij} = s_{ij} - a_i a_j^T$$



where  $s_{ij}$  denotes an element of the covariance/correlation matrix and  $R = (r_{ij})$  denotes the residual matrix. Standardization is performed by dividing the  $r_{ij}$  by

$$\sqrt{u_i u_j}$$

where  $u_i = s_{ii} - y_i$  is the unique error variance for the  $i$ -th variable. If  $u_i$  is zero (or slightly less than zero due to roundoff error),  $u_i = 1.0$  is assumed and division by zero is avoided.

### Example

The following example computes the residual correlation matrix with communalities in a 9-factor problem. The resulting residual correlations do not seem to exhibit any pattern.

```

INTEGER      LDA, LDCOV, LDRESI, NF, NVAR
PARAMETER    (LDA=9, LDCOV=9, LDRESI=9, NF=3, NVAR=9)
C
REAL         A(9,3), COV(9,9), RESID(9,9), Y(9)
EXTERNAL     FRESI, WRRRN
C
DATA COV/1.000, 0.523, 0.395, 0.471, 0.346, 0.426, 0.576, 0.434,
&      0.639, 0.523, 1.000, 0.479, 0.506, 0.418, 0.462, 0.547,
&      0.283, 0.645, 0.395, 0.479, 1.000, 0.355, 0.270, 0.254,
&      0.452, 0.219, 0.504, 0.471, 0.506, 0.355, 1.000, 0.691,
&      0.791, 0.443, 0.285, 0.505, 0.346, 0.418, 0.270, 0.691,
&      1.000, 0.679, 0.383, 0.149, 0.409, 0.426, 0.462, 0.254,
&      0.791, 0.679, 1.000, 0.372, 0.314, 0.472, 0.576, 0.547,
&      0.452, 0.443, 0.383, 0.372, 1.000, 0.385, 0.680, 0.434,
&      0.283, 0.219, 0.285, 0.149, 0.314, 0.385, 1.000, 0.470,
&      0.639, 0.645, 0.504, 0.505, 0.409, 0.472, 0.680, 0.470,
&      1.000/
C
DATA A/.6642, .6888, .4926, .8372, .7050, .8187, .6615, .4579,
&      .7657, -.3209, -.2471, -.3022, .2924, .3148, .3767, -.3960,
&      -.2955, -.4274, .0735, -.1933, -.2224, -.0354, -.1528,
&      .1045, -.0778, .4914, -.0117/
C
CALL FRESI (NVAR, COV, LDCOV, NF, A, LDA, Y, RESID, LDRESI)
C
CALL WRRRN ('Communalities', 1, NVAR, Y, 1, 0)
CALL WRRRN ('Residuals', NVAR, NVAR, RESID, LDRESI, 0)
END

```

### Output

		Communalities								
		1	2	3	4	5	6	7	8	
0.5495										
	0.5729									
		0.3834								
			0.7877							
				0.6195						
					0.8231					
						0.6005				
							0.5385			
								0.7691		
		Residuals								
		1	2	3	4	5	6	7	8	9
1	1.000	0.001	-0.024	0.037	-0.024	-0.016	0.036	-0.002	-0.018	
2	0.001	1.000	0.043	-0.017	-0.048	0.041	-0.052	-0.023	0.031	

3	-0.024	0.043	1.000	0.064	-0.033	-0.037	-0.022	0.025	-0.013
4	0.037	-0.017	0.064	1.000	0.012	-0.004	0.008	0.017	-0.052
5	-0.024	-0.048	-0.033	0.012	1.000	-0.003	0.075	-0.014	0.007
6	-0.016	0.041	-0.037	-0.004	-0.003	1.000	-0.046	-0.003	0.036
7	0.036	-0.052	-0.022	0.008	0.075	-0.046	1.000	0.008	0.011
8	-0.002	-0.023	0.025	0.017	-0.014	-0.003	0.008	1.000	-0.004
9	-0.018	0.031	-0.013	-0.052	0.007	0.036	0.011	-0.004	1.000

---

## MVIND/DMVIND (Single/Double precision)

Compute a test for the independence of  $k$  sets of multivariate normal variables.

### Usage

CALL MVIND (NDF, NVAR, COV, LDCOV, NGROUP, NVSET, STAT)

### Arguments

**NDF** — Number of degrees of freedom in COV. (Input)

**NVAR** — Number of variables in the covariance matrix. (Input)

**COV** — NVAR by NVAR variance-covariance matrix. (Input)

**LDCOV** — Leading dimension of COV exactly as specified in the dimension statement in the calling program. (Input)

**NGROUP** — Number of sets of variables to be tested for independence. (Input)

**NVSET** — Index vector of length NGROUP. (Input)

NVSET( $i$ ) gives the number of variables in the  $i$ -th set of variables. The first NVSET(1) variables in COV define the first set of covariates, the next NVSET(2) variables define the second set of covariates, etc.

**STAT** — Vector of length 4 containing the output statistics. (Output)

**I**        **STAT(I)**

1        Statistic  $V$  for testing the hypothesis of independence of the NGROUP sets of variables.

2        Chi-squared statistic associated with  $V$ .

3        Degrees of freedom for STAT(2).

4        Probability of exceeding STAT(2) under the null hypothesis of independence.

### Comments

- Automatic workspace usage is

MVIND NVAR \* (NVAR + 2) units, or

DMVIND NVAR \* (2 \* NVAR + 3) units.

Workspace may be explicitly provided, if desired, by use of M2IND/DM2IND. The reference is

CALL M2IND (NDF, NVAR, COV, LD COV, NGROUP, NVSET,  
STAT, FAC, WK, IPVT)

The additional arguments are as follows:

**FAC** — Work vector of length  $NVAR^2$ .

**WK** — Work vector of length  $NVAR$ .

**IPVT** — Work vector of length  $NVAR$ .

2. Informational errors

Type	Code	
4	1	A covariance matrix for a subset of the variables is singular.
4	2	The covariance matrix for all variables is singular.

**Algorithm**

Routine MVIND computes a likelihood ratio test statistic proposed by Wilks (1935) for testing the independence of  $NGROUP$  sets of multivariate normal variates. The likelihood ratio statistic is computed as the ratio of the determinant  $|S|$  of the sample covariance matrix to the product of the determinants  $|S_1| \dots |S_k|$  of the covariance matrices of each of the  $k = NGROUP$  sets of variates. An asymptotic chi-squared statistic obtained from the likelihood ratio, along with corresponding  $p$ -value, is computed according to formulas given by Morrison (1976, pages 258-259). The chi-squared statistic is computed as:

$$\chi^2 = -\frac{n}{C} \ln(V)$$

where  $n = NDF$ ,

$$V = \frac{|S|}{|S_{11}| \cdots |S_{kk}|}$$

$$C^{-1} = 1 - \frac{2\sigma_2 + 3\sigma_3}{6n\sigma_2}$$

$$\sigma_2 = \left( \sum_{i=1}^k p_i \right)^2 - \sum_{i=1}^k p_i^2$$

$$\sigma_3 = \left( \sum_{i=1}^k p_i \right)^3 - \sum_{i=1}^k p_i^3$$

where  $|S_{ii}|$  is the determinant of the  $i$ -th covariance matrix,  $k = \text{NGROUP}$ , and  $p_i = \text{NVSET}(i)$ , and  $|S|$  is the determinant of COV.

Because determinants appear in both the numerator and denominator of the likelihood ratio, the test statistic is unchanged when correlation matrices are substituted for covariance matrices as input to MVIND.

In using MVIND, the covariance matrix must first be computed (possibly via routine CORVC, page 314). The covariance matrix may then need to be rearranged (possible via routine RORDM, page 1268) so that the NVSET(1) variables in the first set correspond to the first NVSET(1) columns (and rows) of the covariance matrix, with the next NVSET(2) columns and rows containing the variables for the second set of variables, etc. With this special arrangement of the covariance matrix, routine MVIND may then be called.

### Example

The example is taken from Morrison (1976, page 258). It involves two sets of covariates, with each set having two covariates. The null hypothesis of no relationship is rejected.

```

INTEGER      LDCOV, NDF, NGROUP, NVAR
PARAMETER    (NDF=932, NGROUP=2, NVAR=4, LDCOV=NVAR)
C
INTEGER      NOUT, NVSET(NGROUP)
REAL         COV(NVAR,NVAR), STAT(4)
EXTERNAL     MVIND, UMACH
C
DATA COV/1.00, 0.45, -0.19, 0.43, 0.45, 1.00, -0.02, 0.62,
&      -0.19, -0.02, 1.00, -0.29, 0.43, 0.62, -0.29, 1.00/
C
DATA NVSET/2, 2/
C
CALL MVIND (NDF, NVAR, COV, LDCOV, NGROUP, NVSET, STAT)
C
CALL UMACH (2, NOUT)
WRITE (NOUT,99999) STAT
99999 FORMAT (' Likelihood ratio ..... ', F12.4, /, ' ',
&           'Chi-squared ..... ', F9.1, /, ' Degrees of '
&           ', 'freedom ..... ', F9.1, /, ' p-value ',
&           '..... ', F12.4)
END

```

### Output

```

Likelihood ratio .....      0.5497
Chi-squared .....          556.2
Degrees of freedom .....     4.0
p-value .....              0.0000

```

---

## CANCR/DCANCR (Single/Double precision)

Perform canonical correlation analysis from a data matrix.

## Usage

```
CALL CANCR (NOBS, NVAR1, NVAR2, NCOL, X, LDX, IFRQ,  
            IWT, IND1, IND2, TOL, IPRINT, XX, LDXX,  
            CORR, LDCORR, COEF1, LDCOF1, COEF2, LDCOF2,  
            COEFR1, LDCFR1, COEFR2, LDCFR2, STAT, LDSTAT)
```

## Arguments

**NOBS** — Number of observations. (Input)

**NVAR1** — Number of variables in group 1. (Input)

**NVAR2** — Number of variables in group 2. (Input)

**NCOL** — Number of columns in  $X$ . (Input)

**X** —  $NOBS$  by  $NVAR1 + NVAR2 + m$  data matrix where  $m$  is 0, 1, or 2 depending on whether any columns of  $X$  correspond to frequencies or weights. (Input)  
Each row of  $X$  contains an observation of the  $NVAR1 + NVAR2$  variables for which canonical correlations are desired (plus a weight and/or a frequency variable if  $IFRQ$  and/or  $IWT$  (see below) are not zero). If both  $IWT$  and  $IFRQ$  are zero,  $m$  is 0; 1, if one of  $IFRQ$  or  $IWT$  is positive; and 2, otherwise.  $X$  may not have any missing values (NaN, not a number).

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

**IFRQ** — Frequency option. (Input)

If  $IFRQ = 0$ , then all frequencies are 1. If  $IFRQ$  is positive, then column number  $IFRQ$  of  $X$  contains the nonnegative frequencies.

**IWT** — Weighting option. (Input)

If  $IWT = 0$ , then there is no weighting, i.e., all weights are 1. If  $IWT$  is positive, then column number  $IWT$  of  $X$  contains the nonnegative weights.

**IND1** — Vector of length  $NVAR1$  containing the column numbers in  $X$  of the group 1 variables. (Input)

**IND2** — Vector of length  $NVAR2$  containing the column numbers in  $X$  of the group 2 variables. (Input)

**TOL** — Constant used for determining linear dependence. (Input)

If the squared multiple correlation coefficient of a variable with its predecessors in  $IND1$  (or  $IND2$ ) is greater than  $1 - TOL$ , then the variable is considered to be linearly dependent upon the previous variables; it is excluded from the analysis.  $TOL = .001$  is a typical value.  $TOL$  must be in the exclusive range of 0.0 to 1.0.

**IPRINT** — Printing option. (Input)

### **IPRINT Action**

- |   |   |
|---|---|
| 0 | No printing.  |
| 1 | Print CORR, COEF1, COEF2, COEFR1, COEFR2, and STAT. |
| 2 | Print all output.                                   |

**XX** — NOBS by NVAR1 + NVAR2 +  $m$  matrix containing the canonical scores.

(Output)

$m$  is defined in the description for  $x$ .  $x$  and  $xx$  may occupy the same storage locations. Canonical scores are returned in the first NVAR1 + NVAR2 columns of  $xx$ . Scores for the NVAR1 variables come first. If one of IFRQ or IWT are not zero, then the last column of  $xx$  contains the weight or frequency. If both IFRQ and IWT are not zero, then the frequencies and weights are in the second to last and last column of  $xx$ , respectively.

**LDXX** — Leading dimension of  $xx$  exactly as specified in the dimension statement in the calling program. (Input)

**CORR** — NV by 6 matrix of output statistics. (Output)

NV is the minimum of NVAR1 and NVAR2. CORR has the following statistics.

Col.	Statistic
1	Canonical correlations sorted from the largest to the smallest.
2	Wilks' lambda for testing that the current and all smaller canonical correlations are zero.
3	Rao's $F$ corresponding to Wilks' lambda. If the canonical correlation is greater than 0.99999, then $F$ is set to 9999.99.
4	Numerator degrees of freedom for $F$ .
5	Denominator degrees of freedom for $F$ .
6	Probability of a larger $F$ statistic.

If an  $F$  statistic is negative, then CORR( $i$ , 6) is set to one. If either CORR( $i$ , 4) or CORR( $i$ , 5) is not positive, then CORR( $i$ , 6) is set to the missing value code (NaN).

**LDCORR** — Leading dimension of CORR exactly as specified in the dimension statement in the calling program. (Input)

**COEF1** — NVAR1 by NVAR1 matrix containing the group 1 canonical coefficients. (Output)

The columns of COEF1 contain the vectors of canonical coefficients for group 1.

**LDCOF1** — Leading dimension of COEF1 exactly as specified in the dimension statement in the calling program. (Input)

**COEF2** — NVAR2 by NVAR2 matrix containing the group 2 canonical coefficients. (Output)

The columns of COEF2 contain the vectors of canonical coefficients for group 2.

**LDCOF2** — Leading dimension of COEF2 exactly as specified in the dimension statement in the calling program. (Input)

**COEFR1** — NVAR1 by NV matrix containing the correlations between the group 1 variables and the group 1 canonical scores. (Output)

NV is the minimum of NVAR1 and NVAR2.

**LDCFR1** — Leading dimension of COEFR1 exactly as specified in the dimension statement in the calling program. (Input)

**COEFR2** — NVAR2 by NV matrix containing the correlations between the group 2 variables and the group 2 canonical scores. (Output)

NV is the minimum of NVAR1 and NVAR2.

**LDCFR2** — Leading dimension of COEFR2 exactly as specified in the dimension statement in the calling program. (Input)

**STAT** — 15 by NVAR1 + NVAR2 matrix containing statistics on all of the variables. (Output)

The first NVAR1 columns of STAT correspond to the group one variables with the last NVAR2 columns corresponding to the group two variables.

Row	Statistic
1	Means
2	Variances
3	Standard deviations
4	Coefficients of skewness
5	Coefficients of excess (kurtosis)
6	Minima
7	Maxima
8	Ranges
9	Coefficients of variation, when defined, 0.0 otherwise
10	Numbers of nonmissing observations
11	Lower endpoints of 95% confidence interval for the means
12	Upper endpoints of 95% confidence interval for the means
13	Lower endpoints of 95% confidence interval for the variances
14	Upper endpoints of 95% confidence interval for the variances
15	Sums of the weights if IWT greater than zero, 0.0 otherwise

**LDSTAT** — Leading dimension of STAT exactly as specified in the dimension statement in the calling program. (Input)

### Comments

- Automatic workspace usage is

CANCR  $\text{NVAR1}^2 + \text{NVAR2}^2 + \text{NVAR1} + \text{NVAR2} + 1 + \max(\text{NOBS}, 2 * (\text{NVAR1} + \text{NVAR2})) + (\max(\text{NVAR1}, \text{NVAR2}))^2 + 3 * \max(\text{NVAR1}, \text{NVAR2})$  units, or

DCANCR  $2 * (\text{NVAR1}^2 + \text{NVAR2}^2) + \text{NVAR1} + \text{NVAR2} + 2 * (\max(\text{NOBS}, 2 * (\text{NVAR1} + \text{NVAR2}))) + 2 * (\max(\text{NVAR1}, \text{NVAR2}))^2 + 6 * \max(\text{NVAR1}, \text{NVAR2})$  units.

Workspace may be explicitly provided, if desired, by use of C2NCR/DC2NCR. The reference is

```
CALL C2NCR (NOBS, NVAR1, NVAR2, NCOL, X, LDX, IFRQ,
            IWT, IND1, IND2, TOL, IPRINT, XX, LDXX,
            CORR, LDCORR, COEF1, LDCOF1, COEF2,
            LDCOF2, COEFR1, LDCFR1, COEFR2, LDCFR2,
            STAT, LDSTAT, R, S, IND, WORK, WKA, WK)
```

The additional arguments are as follows:

**R** — Work vector of length  $NVAR1^2$ .

**S** — Work vector of length  $NVAR2^2$ .

**IND** — Work vector of length  $NVAR1 + NVAR2 + 2$ .

**WORK** — Work vector of length  $\max(\text{NOBS}, 2 * (NVAR1 + NVAR2))$

**WKA** — Work vector of length  $(\max(NVAR1, NVAR2))^2$ .

**WK** — Work vector of length  $3 * \max(NVAR1, NVAR2) - 1$ .

2. Informational errors

Type	Code	
3	1	The standardized cross covariance matrix is not of full rank or is very ill-conditioned. Small canonical correlations may not be accurate.
3	2	One or more variables is linearly dependent upon the preceding variables in its group.
4	3	The sum of the frequencies is equal to zero. The sum of the frequencies must be positive.
4	4	The sum of the weights is equal to zero. The sum of the weights must be positive.

**Algorithm**

Routine CANCR computes the canonical correlations, the canonical coefficients, the canonical scores, Wilks' lambda for testing the independence of two sets of variates, and a series of Bartlett's tests of the hypothesis that the  $k$ -th largest and all larger canonical correlations are simultaneously zero. A matrix of observations is used in these computations.

Let  $x_{ij}$  denote the  $j$ -th variable on the  $i$ -th observation,  $w_i$  denote the observation weight,  $f_i$  denote the observation frequency,  $\Gamma_{11}$  denote the upper triangular Cholesky ( $R^T R$ ) factorization of the sample covariance matrix of the group 1 variables,  $\Gamma_{22}$  denote the upper triangular Cholesky ( $R^T R$ ) factorization of the group 2 variables sample covariance matrix, and

$$\Gamma_{12} = (\Gamma_{11})^{-1} \hat{\Sigma}_{12} (\Gamma_{22})^{-T}$$

where

$$\hat{\Sigma}_{12}$$

is the sample estimate of the matrix of covariances between the group 1 and the group 2 variables. Then, the computational procedure in obtaining the canonical correlations is as follows:



1. The weighted mean of each variable is computed via the standard formula (see UVSTA, page 16). The means are then subtracted from the observations.

2. Each element in the  $i$ -th row of  $x$  is multiplied by

$$\sqrt{(w_i f_i)}$$

3. Gram-Schmidt orthogonalization is used on  $x$  to obtain  $Y_1$  and  $Y_2$ , where  $Y_1$  and  $Y_2$  are the results of the Gram-Schmidt orthogonalization of the group 1 and the group 2 variables, respectively. The matrices  $\Gamma_{11}$  and  $\Gamma_{22}$  are obtained as a by-product of the orthogonalization. Compute

$$\Gamma_{12} = Y_1^T Y_2$$

4. The canonical correlations are obtained as the singular values of the matrix  $\Gamma_{12}$ . Denote the left and right orthogonal matrices obtained as a by-product of this decomposition by  $L$  and  $R$ , respectively.
5. The canonical coefficients are obtained from  $L$  and  $R$  by multiplying  $L$  and  $R$  by the inverses of  $\Gamma_{11}$  and  $\Gamma_{22}$ , respectively (see Golub 1969).
6. The correlations of the original variables with the canonical variables are obtained by multiplying  $L$  and  $R$  by  $\Gamma_{11}$  and  $\Gamma_{22}$ , respectively.
7. The canonical scores are obtained by multiplying the matrices  $Y_1$  and  $Y_2$  by the matrices  $L$  and  $R$ , respectively, and then dividing each row of  $Y_1$  and  $Y_2$  by

$$\sqrt{(w_i f_i)}$$

8. Wilks' lambda, the Bartlett's tests, Rao's  $F$  corresponding to these tests, the numerator and denominator degrees of freedom of  $F$ , and the significance level of  $F$  are computed as in Rao (1973, page 556). Bartlett's tests are computed as

$$\Lambda_i = \prod_{j=1}^q (1 - \rho_j^2)$$

where  $q = \text{NVAR2}$  is the number of canonical correlations, the canonical correlations are ordered from largest to smallest, and  $\rho_j$  denotes the  $j$ -th largest canonical correlation. Wilks' lambda is given as  $\Lambda_1$ . The degrees of freedom in the numerator of the corresponding Rao's  $F$  statistic is given as

$$d_1 = pu$$

where  $p = v_1 - i + 1$ ,  $u = v_2 - i + 1$ ,  $v_1 = \text{NVAR2}$ , and  $v_2 = \text{NVAR1}$ . Let

$$m = t - \frac{p + u + 1}{2}$$

where  $t$  is the degrees of freedom in  $\text{COV}(\sum_i f_i - 1)$ , and let

$$s = \sqrt{\frac{p^2 u^2 - 4}{p^2 + u^2 - 5}}$$

if  $p^2 + u^2 - 5 \neq 0$ , and let  $s = 2$  otherwise. Then, Rao's  $F$  corresponding to Bartlett's test is computed as

$$F_i = \frac{1 - \Lambda_i^{\frac{1}{s}}}{\Lambda_i^{\frac{1}{s}}} (ms - pu/2 + 1) / pu$$

Rao's  $F$  has numerator degrees of freedom  $d_2 = ms - pu/2 + 1$ . The significance level of  $F$  is obtained from the standard  $F$  distribution.

### Example 1

The following example is taken from Levin and Marascuilo (1983), pages 191–197. It is examining the relationship between the performance of individuals in a sociology course and predictor variables. The measures of performance in the sociology course are two midterms examinations, a final examination, and a course evaluation, the predictor variables are social class, sex, grade point average, college board test score, whether the student has previously taken a course in sociology, and the student's score on a pretest.

```

C
INTEGER   IFRQ, IPRINT, IWT, LDCFR1, LDCFR2, LDCOF1, LDCOF2,
&         LDCORR, LDSTAT, LDX, LDXX, NCOL, NOBS, NV, NVAR1,
&         NVAR2
REAL      TOL
PARAMETER (IFRQ=0, IPRINT=1, IWT=0, LDSTAT=15, NCOL=10,
&         NOBS=40, NVAR1=6, NVAR2=4, TOL=0.0001, LDCFR1=NVAR1,
&         LDCFR2=NVAR2, LDCOF1=NVAR1, LDCOF2=NVAR2, LDX=NOBS,
&         LDXX=NOBS, NV=NVAR2, LDCORR=NV)

C
INTEGER   IND1(NVAR1), IND2(NVAR2)
REAL      COEF1(LDCOF1, NVAR1), COEF2(LDCOF2, NVAR2),
&         COEFR1(LDCFR1, NV), COEFR2(LDCFR2, NV), CORR(LDCORR, 6),
&         STAT(LDSTAT, NVAR1+NVAR2), X(LDX, NCOL), XX(LDXX, NCOL)
CHARACTER FMT*35, NUMBER*6, XLAB(11)*25
EXTERNAL  CANCR, WRRRL

C
DATA IND1/1, 2, 3, 4, 5, 6/, IND2/7, 8, 9, 10/
DATA (X(I,1), I=1, NOBS)/3*2.0, 3.0, 2.0, 3.0, 1.0, 2.0, 3.0,
& 2*2.0, 3.0, 1.0, 4*2.0, 3.0, 3*2.0, 1.0, 3*2.0, 1.0, 2.0,
& 1.0, 2.0, 3.0, 2*2.0, 2*1.0, 2.0, 3.0, 1.0, 2.0, 3.0, 1.0/
DATA (X(I,2), I=1, NOBS)/6*1.0, 0.0, 2*1.0, 3*0.0, 3*1.0, 3*0.0,
& 1.0, 0.0, 3*1.0, 3*0.0, 4*1.0, 0.0, 8*1.0, 0.0/
DATA (X(I,3), I=1, NOBS)/3.55, 2.70, 3.50, 2.91, 3.10, 3.49, 3.17,
& 3.57, 3.76, 3.81, 3.60, 3.10, 3.08, 3.50, 3.43, 3.39, 3.76,
& 3.71, 3.00, 3.47, 3.69, 3.24, 3.46, 3.39, 3.90, 2.76, 2.70,
& 3.77, 4.00, 3.40, 3.09, 3.80, 3.28, 3.70, 3.42, 3.09, 3.70,

```

```

&      2.69, 3.40, 2.95/
DATA (X(I,4),I=1,NOBS)/410.0, 390.0, 510.0, 430.0, 600.0,
&      2*610.0, 560.0, 700.0, 460.0, 590.0, 500.0, 410.0, 470.0,
&      210.0, 610.0, 510.0, 600.0, 470.0, 460.0, 800.0, 610.0,
&      490.0, 470.0, 610.0, 580.0, 410.0, 630.0, 790.0, 490.0,
&      400.0, 2*610.0, 500.0, 430.0, 540.0, 610.0, 400.0, 390.0,
&      490.0/
DATA (X(I,5),I=1,NOBS)/8*0.0, 4*1.0, 0.0, 2*1.0, 0.0, 1.0, 0.0,
&      1.0, 0.0, 1.0, 3*0.0, 1.0, 2*0.0, 2*1.0, 2*0.0, 4*1.0,
&      5*0.0/
DATA (X(I,6),I=1,NOBS)/17.0, 20.0, 22.0, 13.0, 16.0, 28.0, 14.0,
&      10.0, 28.0, 30.0, 28.0, 15.0, 24.0, 15.0, 26.0, 16.0, 25.0,
&      3.0, 5.0, 16.0, 28.0, 13.0, 9.0, 13.0, 30.0, 10.0, 13.0,
&      8.0, 29.0, 17.0, 15.0, 16.0, 13.0, 30.0, 2*17.0, 25.0,
&      10.0, 23.0, 18.0/
DATA (X(I,7),I=1,NOBS)/43.0, 50.0, 47.0, 24.0, 47.0, 57.0,
&      2*42.0, 69.0, 48.0, 59.0, 21.0, 52.0, 2*35.0, 59.0, 68.0,
&      38.0, 45.0, 37.0, 54.0, 45.0, 31.0, 39.0, 67.0, 30.0, 19.0,
&      71.0, 80.0, 47.0, 46.0, 59.0, 48.0, 68.0, 43.0, 31.0, 64.0,
&      19.0, 43.0, 20.0/
DATA (X(I,8),I=1,NOBS)/61.0, 47.0, 79.0, 40.0, 60.0, 59.0, 61.0,
&      79.0, 83.0, 67.0, 74.0, 40.0, 71.0, 40.0, 57.0, 58.0, 66.0,
&      58.0, 24.0, 48.0, 100.0, 83.0, 70.0, 48.0, 85.0, 14.0,
&      55.0, 100.0, 94.0, 45.0, 58.0, 90.0, 84.0, 81.0, 49.0,
&      54.0, 87.0, 36.0, 51.0, 59.0/
DATA (X(I,9),I=1,NOBS)/129.0, 60.0, 119.0, 100.0, 79.0, 99.0,
&      92.0, 107.0, 156.0, 110.0, 116.0, 49.0, 107.0, 125.0, 64.0,
&      100.0, 138.0, 63.0, 82.0, 73.0, 132.0, 87.0, 89.0, 99.0,
&      119.0, 100.0, 84.0, 166.0, 111.0, 110.0, 93.0, 141.0, 99.0,
&      114.0, 96.0, 39.0, 149.0, 53.0, 39.0, 91.0/
DATA (X(I,10),I=1,NOBS)/3.0, 3*1.0, 2.0, 1.0, 3.0, 2.0, 4*1.0,
&      5.0, 1.0, 5.0, 1.0, 2.0, 1.0, 2*3.0, 3*2.0, 1.0, 2.0, 1.0,
&      2.0, 3.0, 2.0, 2*1.0, 2*2.0, 5.0, 2*1.0, 4.0, 3.0, 2*1.0/
C
DATA XLAB/' ', 'Social%/Class', '%/Sex', '%/GPA',
&      'College%/Boards', 'H.S.%/Soc.', 'Pretest%/Score',
&      '%/Exam 1', '%/Exam 2', 'Final%/Exam', 'Course%/Eval.'/
C
DATA NUMBER/'NUMBER'/, FMT/'(2W3.1,W5.3,W4.1,W3.1,4W5.1,W3.1)'/
C
CALL WRRRL ('First 10 Observations', 10, NCOL, X, LDX, 0, FMT,
&      NUMBER, XLAB)
C
CALL CANCR (NOBS, NVAR1, NVAR2, NCOL, X, LDX, IFRQ, IWT, IND1,
&      IND2, TOL, IPRINT, XX, LDXX, CORR, LDCORR, COEF1,
&      LDCOF1, COEF2, LDCOF2, COEFR1, LDCFR1, COEFR2,
&      LDCFR2, STAT, LDSTAT)
C
END

```

### Output

	First 10 Observations									
	Social Class	Sex	GPA	College Boards	H.S. Soc.	Pretest Score	Exam 1	Exam 2	Final Exam	Course Eval
1	2	1	3.55	410	0	17	43	61	129	3
2	2	1	2.70	390	0	20	50	47	60	1
3	2	1	3.50	510	0	22	47	79	119	1
4	3	1	2.91	430	0	13	24	40	100	1
5	2	1	3.10	600	0	16	47	60	79	2

6	3	1	3.49	610	0	28	57	59	99	1
7	1	0	3.17	610	0	14	42	61	92	3
8	2	1	3.57	560	0	10	42	79	107	2
9	3	1	3.76	700	1	28	69	83	156	1
10	2	0	3.81	460	1	30	48	67	110	1

\*\*\* Canonical Correlations Statistics \*\*\*

	Canonical Correlations	Wilks	Lambda	Raos F	Num. df	Denom. df	Prob. of Larger F
1	0.9242		0.0612	5.412	24	105.9	0.0000
2	0.7184		0.4201	2.116	15	86.0	0.0162
3	0.2893		0.8683	0.586	8	64.0	0.7861
4	0.2290		0.9476	0.609	3	33.0	0.6142

Group One Canonical Coefficients

	1	2	3	4	5	6
1	-0.622	1.158	-0.285	-0.179	0.601	-0.423
2	0.558	-0.739	0.231	-1.278	1.391	-0.024
3	1.796	-0.432	0.765	0.185	-0.643	-3.314
4	0.002	0.006	0.004	-0.002	0.000	0.006
5	-0.059	-0.043	-0.456	1.671	1.463	0.774
6	0.031	0.018	-0.121	-0.058	-0.042	0.056

Group Two Canonical Coefficients

	1	2	3	4
1	0.0233	-0.0365	0.0845	-0.0176
2	0.0257	-0.0057	-0.0352	0.0555
3	0.0073	0.0110	-0.0259	-0.0341
4	0.1034	0.8089	0.2828	0.0260

Correlations Between the Group One Variables and the Group One Canonical Scores

	1	2	3	4
1	-0.3685	0.6795	-0.2291	-0.1854
2	0.2157	-0.3252	0.0521	-0.5985
3	0.8153	0.2770	-0.0692	0.2123
4	0.6144	0.5681	0.4151	-0.0050
5	0.4661	0.0603	-0.3034	0.6530
6	0.5461	0.1768	-0.7915	-0.1375

Correlations Between the Group Two Variables and the Group Two Canonical Scores

	1	2	3	4
1	0.8713	-0.2406	0.3864	-0.1835
2	0.9174	-0.0557	-0.2068	0.3355
3	0.7707	0.0293	-0.3146	-0.5533
4	0.3490	0.8765	0.3077	0.1240

\*\*\* Statistics for Group One Variables \*\*\*  
Univariate Statistics from UVSTA

Variable	Mean	Variance	Std. Dev.	Skewness	Kurtosis
1	1.9750	0.4353	0.6597	0.02476	-0.6452
2	0.6750	0.2250	0.4743	-0.74726	-1.4416
3	3.3758	0.1247	0.3532	-0.37911	-0.7521
4	524.2499	13148.1377	114.6653	0.09897	0.6494
5	0.4000	0.2462	0.4961	0.40825	-1.8333
6	18.1250	55.1378	7.4255	0.10633	-0.9358

Variable	Minimum	Maximum	Range	Coef. Var.	Count
1	1.0000	3.0000	2.0000	0.3340	40.0000
2	0.0000	1.0000	1.0000	0.7027	40.0000
3	2.6900	4.0000	1.3100	0.1046	40.0000
4	210.0000	800.0000	590.0000	0.2187	40.0000
5	0.0000	1.0000	1.0000	1.2403	40.0000
6	3.0000	30.0000	27.0000	0.4097	40.0000

Variable	Lower CLM	Upper CLM	Lower CLV	Upper CLV
1	1.7640	2.1860	0.29207	0.7176
2	0.5233	0.8267	0.15098	0.3710
3	3.2628	3.4887	0.08369	0.2056
4	487.5782	560.9217	8822.72168	21677.9590
5	0.2413	0.5587	0.16518	0.4058
6	15.7502	20.4998	36.99883	90.9083

\*\*\* Statistics for Group Two Variables \*\*\*  
Univariate Statistics from UVSTA

Variable	Mean	Variance	Std. Dev.	Skewness	Kurtosis
1	46.0500	237.0231	15.3956	0.08762	-0.5505
2	62.8750	403.4967	20.0872	-0.10762	-0.3642
3	99.4750	919.4864	30.3230	-0.03483	-0.2533
4	1.9500	1.4333	1.1972	1.27704	0.8407

Variable	Minimum	Maximum	Range	Coef. Var.	Count
1	19.0000	80.0000	61.0000	0.3343	40.0000
2	14.0000	100.0000	86.0000	0.3195	40.0000
3	39.0000	166.0000	127.0000	0.3048	40.0000
4	1.0000	5.0000	4.0000	0.6140	40.0000

Variable	Lower CLM	Upper CLM	Lower CLV	Upper CLV
1	41.1263	50.9737	159.0483	390.7912
2	56.4508	69.2992	270.7562	665.2642
3	89.7772	109.1728	616.9979	1516.0009
4	1.5671	2.3329	0.9618	2.3632

## Example 2

Correspondence analysis is an interesting application of canonical correlation in the analysis of contingency tables. The example is taken from Kendall and Stuart (1979, pages 595–599) and involves finding the optimal scores for the values of two categorical variables to maximize the correlation between the two variables. The contingency table is given below, along with the more traditional matrix  $X$  of “observations” for which canonical correlations are desired.

$$\begin{pmatrix} 821 & 112 & 85 & 35 \\ 116 & 494 & 145 & 27 \\ 72 & 151 & 583 & 87 \\ 43 & 34 & 106 & 331 \end{pmatrix}$$

The data matrix  $X$  is given as:

Group 1 Var.				Group 2 Var.				Frequencies
1	0	0	0	1	0	0	0	821
1	0	0	0	0	1	0	0	112
1	0	0	0	0	0	1	0	85
1	0	0	0	0	0	0	1	35
0	1	0	0	1	0	0	0	116
0	1	0	0	0	1	0	0	494
0	1	0	0	0	0	1	0	145
0	1	0	0	0	0	0	1	27
0	0	1	0	1	0	0	0	72
0	0	1	0	0	1	0	0	151
0	0	1	0	0	0	1	0	583
0	0	1	0	0	0	0	1	87
0	0	0	1	1	0	0	0	43
0	0	0	1	0	1	0	0	34
0	0	0	1	0	0	1	0	106
0	0	0	1	0	0	0	1	331

For this table, the optimal correlation turns out to be 0.70 when scores of 2.67, 1.34, 0.62, and 0.00 (see Column 1 of COEF1) are assigned to the variable 1 categories, and scores of 2.72, 1.37, 0.68, and 0.00 are assigned to the variable 2 categories. These scores are obtained as the canonical scores when canonical correlations are computed between the the row and column variable indicator variables (variables 1-4 and variables 5-8 in x, respectively). The warning error appears in the output because the covariance matrix is not of full rank (indeed, neither the group 1 or the group 2 covariance matrices are of full rank).

```

      INTEGER      IFRQ, IPRINT, IWT, LDCFR1, LDCFR2, LDCOF1, LDCOF2,
&                LDCORR, LDSTAT, LDX, LDXX, NCOL, NOBS, NV, NVAR1,
&                NVAR2
      REAL         TOL
      PARAMETER   (IFRQ=9, IPRINT=2, IWT=0, LDCFR1=4, LDCFR2=4,
&                LDCOF1=4, LDCOF2=4, LDCORR=4, LDSTAT=15, LDX=16,
&                LDXX=16, NCOL=9, NOBS=16, NV=4, NVAR1=4, NVAR2=4,
&                TOL=0.0001)
C
      INTEGER      IND1(NVAR1), IND2(NVAR2)
      REAL         COEF1(LDCOF1,NVAR1), COEF2(LDCOF2,NVAR2),
&                COEFR1(LDCFR1,NV), COEFR2(LDCFR2,NV), CORR(LDCORR,6),
&                STAT(LDSTAT,8), X(LDX,NCOL), XX(LDXX,NCOL)
      EXTERNAL    CANCR
C
      DATA IND1/1, 2, 3, 4/, IND2/5, 6, 7, 8/
      DATA X/4*1.0, 16*0.0, 4*1.0, 16*0.0, 4*1.0, 16*0.0, 5*1.0,
&         3*0.0, 1.0, 3*0.0, 1.0, 3*0.0, 1.0, 4*0.0, 1.0, 3*0.0, 1.0,
&         3*0.0, 1.0, 3*0.0, 1.0, 4*0.0, 1.0, 3*0.0, 1.0, 3*0.0, 1.0,
&         3*0.0, 1.0, 4*0.0, 1.0, 3*0.0, 1.0, 3*0.0, 1.0, 3*0.0, 1.0,

```

```

&      821.0, 112.0, 85.0, 35.0, 116.0, 494.0, 145.0, 27.0, 72.0,
&      151.0, 583.0, 87.0, 43.0, 34.0, 106.0, 331.0/
C
CALL CANCR (NOBS, NVAR1, NVAR2, NCOL, X, LDX, IFRQ, IWT, IND1,
&          IND2, TOL, IPRINT, XX, LDXX, CORR, LDCORR, COEF1,
&          LDCOF1, COEF2, LDCOF2, COEFR1, LDCFR1, COEFR2,
&          LDCFR2, STAT, LDSTAT)
C
END

```

### Output

\*\*\* WARNING ERROR 2 from C2NCR. One or more Group 1 variables is linearly  
\*\*\* dependent on the preceding variables in Group 1.

Here is a traceback of subprogram calls in reverse order:

Routine name	Error type	Error code
C2NCR	6	2 (Called internally)
CANCR	0	0
USER	0	0

\*\*\* WARNING ERROR 3 from C2NCR. One or more Group 2 variables is linearly  
\*\*\* dependent on the preceding variables in Group 2.

Here is a traceback of subprogram calls in reverse order:

Routine name	Error type	Error code
C2NCR	6	3 (Called internally)
CANCR	0	0
USER	0	0

### \*\*\* Canonical Correlations Statistics \*\*\*

Canonical Correlations	Wilks	Lambda	Raos F	Num. df	Denom. df	Prob. of Larger F
1	0.6965	0.2734	615.925	9	7875.7	0.0000
2	0.5883	0.5310	602.598	4	6474.0	0.0000
3	0.4336	0.8120	749.823	1	3238.0	0.0000
4	0.0000	0.0000	0.000	0	0.0	0.0000

### Group One Canonical Coefficients

	1	2	3	4
1	2.670	1.100	1.023	0.000
2	1.341	2.905	-0.460	0.000
3	0.624	2.222	2.147	0.000
4	0.000	0.000	0.000	0.000

### Group Two Canonical Coefficients

	1	2	3	4
1	2.715	1.164	1.053	0.000
2	1.366	2.972	-0.393	0.000
3	0.676	2.250	2.182	0.000
4	0.000	0.000	0.000	0.000

### Correlations Between the Group One Variables and the Group One Canonical Scores

	1	2	3	4
1	0.9068	-0.3954	0.1459	0.0000
2	-0.0121	0.6965	-0.7175	0.0000
3	-0.4555	0.3404	0.8226	0.0000

4 0.0000 0.0000 0.0000 0.0000

Correlations Between the Group Two Variables  
and the Group Two Canonical Scores

	1	2	3	4
1	0.9072	-0.3997	0.1310	0.0000
2	-0.0227	0.6995	-0.7143	0.0000
3	-0.4590	0.3205	0.8287	0.0000
4	0.0000	0.0000	0.0000	0.0000

\*\*\* Statistics for Group One Variables \*\*\*  
Univariate Statistics from UVSTA

Variable	Mean	Variance	Std. Dev.	Skewness	Kurtosis
1	0.3248	0.2194	0.4684	0.7482	-1.4401
2	0.2412	0.1831	0.4279	1.2098	-0.5363
3	0.2754	0.1996	0.4468	1.0053	-0.9894
4	0.1585	0.0000	0.0000	1.8697	1.4958

Variable	Minimum	Maximum	Range	Coef. Var.	Count
1	0.0000	1.0000	1.0000	1.4420	3242.0000
2	0.0000	1.0000	1.0000	1.7739	3242.0000
3	0.0000	1.0000	1.0000	1.6221	3242.0000
4	0.0000	1.0000	1.0000	2.3041	3242.0000

Variable	Lower CLM	Upper CLM	Lower CLV	Upper CLV
1	0.3087	0.3409	0.2091	0.2305
2	0.2265	0.2559	0.1745	0.1923
3	0.2601	0.2908	0.1903	0.2097
4	0.1460	0.1711	0.1272	0.1402

Canonical Scores for Group One

	1	2	3	4
1	1.307	-0.570	0.210	0.000
2	1.307	-0.570	0.210	0.000
3	1.307	-0.570	0.210	0.000
4	1.307	-0.570	0.210	0.000
5	-0.021	1.235	-1.272	0.000
6	-0.021	1.235	-1.272	0.000
7	-0.021	1.235	-1.272	0.000
8	-0.021	1.235	-1.272	0.000
9	-0.739	0.552	1.334	0.000
10	-0.739	0.552	1.334	0.000
11	-0.739	0.552	1.334	0.000
12	-0.739	0.552	1.334	0.000
13	-1.362	-1.670	-0.813	0.000
14	-1.362	-1.670	-0.813	0.000
15	-1.362	-1.670	-0.813	0.000
16	-1.362	-1.670	-0.813	0.000

\*\*\* Statistics for Group Two Variables \*\*\*  
Univariate Statistics from UVSTA

Variable	Mean	Variance	Std. Dev.	Skewness	Kurtosis
1	0.3245	0.2193	0.4683	0.7497	-1.4379
2	0.2440	0.1845	0.4296	1.1922	-0.5787
3	0.2835	0.2032	0.4508	0.9609	-1.0766
4	0.1481	0.0000	0.0000	1.9819	1.9280

Variable	Minimum	Maximum	Range	Coef. Var.	Count
----------	---------	---------	-------	------------	-------



1	0.0000	1.0000	1.0000	1.4430	3242.0000
2	0.0000	1.0000	1.0000	1.7606	3242.0000
3	0.0000	1.0000	1.0000	1.5901	3242.0000
4	0.0000	1.0000	1.0000	2.3992	3242.0000
Variable	Lower CLM	Upper CLM	Lower CLV	Upper CLV	
1	0.3084	0.3406	0.2090	0.2303	
2	0.2292	0.2588	0.1758	0.1938	
3	0.2679	0.2990	0.1936	0.2134	
4	0.1358	0.1603	0.1203	0.1326	

Canonical Scores for Group Two

	1	2	3	4
1	1.309	-0.577	0.189	0.000
2	-0.040	1.231	-1.257	0.000
3	-0.730	0.509	1.317	0.000
4	-1.406	-1.740	-0.864	0.000
5	1.309	-0.577	0.189	0.000
6	-0.040	1.231	-1.257	0.000
7	-0.730	0.509	1.317	0.000
8	-1.406	-1.740	-0.864	0.000
9	1.309	-0.577	0.189	0.000
10	-0.040	1.231	-1.257	0.000
11	-0.730	0.509	1.317	0.000
12	-1.406	-1.740	-0.864	0.000
13	1.309	-0.577	0.189	0.000
14	-0.040	1.231	-1.257	0.000
15	-0.730	0.509	1.317	0.000
16	-1.406	-1.740	-0.864	0.000

\*\*\* WARNING ERROR 1 from CANCR. The standardized cross covariance matrix is not of full rank or is very ill-conditioned. Small canonical correlations may not be accurate.

---

## CANVC/DCANVC (Single/Double precision)

Perform canonical correlation analysis from a variance-covariance matrix or a correlation matrix.

### Usage

CALL CANVC (NDF, NVAR1, NVAR2, COV, LDCOV, IND1, IND2, IPRINT, CORR, LDCORR, COEF1, LDCOF1, COEF2, LDCOF2, COEFR1, LDCFR1, COEFR2, LDCFR2)

### Arguments

**NDF** — Number of degrees of freedom in the covariance or correlation matrix. (Input)

If NDF is unknown, an estimate of NDF = 100 is suggested in which case the last four columns of CORR are meaningless.

**NVAR1** — Number of variables in group 1. (Input)

**NVAR2** — Number of variables in group 2. (Input)

**COV** —  $NVAR1 + NVAR2$  by  $NVAR1 + NVAR2$  matrix containing the covariance or correlation matrix. (Input)

Routines **COVPL** (page 322), **RBCOV** (page 331), or **CORVC** (page 314) may be used to calculate **COV** from a data matrix. **COV** must be nonnegative definite within a tolerance of  $100.0 * AMACH(4)$ . Only the upper triangle of **COV** is referenced.

**LDCOV** — Leading dimension of **COV** exactly as specified in the dimension statement in the calling program. (Input)

**IND1** — Vector of length  $NVAR1$  containing the column and row numbers in **COV** for the group 1 variables. (Input)

**IND2** — Vector of length  $NVAR2$  containing the column and row numbers in **COV** for the group 2 variables. (Input)

**IPRINT** — Printing option. (Input)

**IPRINT Action**

0 No printing.

1 Printing of **CORR**, **COEF1**, **COEF2**, **COEFR1**, and **COEFR2** is performed.

**CORR** —  $NV$  by 6 matrix containing the output statistics. (Output)

$NV$  is the minimum of  $NVAR1$  and  $NVAR2$ .

**Col. Statistic**

1 Canonical correlations sorted from the largest to the smallest.

2 Wilks' lambda for testing that the current and all smaller canonical correlations are zero.

3 Rao's  $F$  corresponding to Wilks' lambda. If the canonical correlation is greater than 0.99999,  $F$  is set to 9999.99.

4 Numerator degrees of freedom for the  $F$ .

5 Denominator degrees of freedom for the  $F$ .

6 Probability of a larger  $F$  statistic.

If an  $F$  statistic is negative, then **CORR**( $i$ , 6) is set to one. If either **CORR**( $i$ , 4) or **CORR**( $i$ , 5) is not positive, then **CORR**( $i$ , 6) is set to the missing value code (NaN).

**LDCORR** — Leading dimension of **CORR** exactly as specified in the dimension statement in the calling program. (Input)

**COEF1** —  $NVAR1$  by  $NVAR1$  matrix containing the group 1 canonical coefficients. (Output)

The columns of **COEF1** contain the vectors of canonical coefficients for group 1.

**LDCOF1** — Leading dimension of **COEF1** exactly as specified in the dimension statement in the calling program. (Input)

**COEF2** —  $NVAR2$  by  $NVAR2$  matrix containing the group 2 canonical coefficients. (Output)

The columns of **COEF2** contain the vectors of canonical coefficients for group 2.

**LDCOF2** — Leading dimension of **COEF2** exactly as specified in the dimension statement in the calling program. (Input)

**COEFR1** — NVAR1 by NV matrix containing the correlations between the group 1 variables and the group 1 canonical scores. (Output)  
 NV is the minimum of NVAR1 and NVAR2.

**LDCFR1** — Leading dimension of COEFR1 exactly as specified in the dimension statement in the calling program. (Input)

**COEFR2** — NVAR2 by NV matrix containing the correlations between the group 2 variables and the group 2 canonical scores. (Output)  
 NV is the minimum of NVAR1 and NVAR2.

**LDCFR2** — Leading dimension of COEFR2 exactly as specified in the dimension statement in the calling program. (Input)

### Comments

- Automatic workspace usage is

CANVC  $\text{NVAR1}^2 + \text{NVAR2}^2 + (\text{NVAR1} + \text{NVAR2})^2 + \text{NVAR1} + \text{NVAR2} + 3$   
 $\times \max(\text{NVAR1}, \text{NVAR2})$  units, or

DCANVC  $2 * (\text{NVAR1}^2 + \text{NVAR2}^2 + (\text{NVAR1} + \text{NVAR2})^2 + \text{NVAR1} + \text{NVAR2}$   
 $+ 3 * \max(\text{NVAR1}, \text{NVAR2}))$  units.

Workspace may be explicitly provided, if desired, by use of  
 C2NVC/DC2NVC. The reference is

```
CALL C2NVC (NDF, NVAR1, NVAR2, COV, LD COV, IND1,
            IND2, IPRINT, CORR, LDCORR, COEF1,
            LD COF1, COEF2, LD COF2, COEFR1, LDCFR1,
            COEFR2, LDCFR2, R, S, STD1, STD2, WKA,
            WK)
```

The additional arguments are as follows:

**R** — Work vector of length  $\text{NVAR1}^2$ .

**S** — Work vector of length  $\text{NVAR2}^2$ .

**STD1** — Work vector of length NVAR1.

**STD2** — Work vector of length NVAR2.

**WKA** — Work vector of length  $(\text{NVAR1} + \text{NVAR2})^2$ .

**WK** — Work vector of length  $3 * \max(\text{NVAR1}, \text{NVAR2})$ .

- Informational errors

Type	Code	
3	1	The standardized cross covariance matrix is not of full rank or is very ill-conditioned. Small canonical correlations may not be accurate.
4	2	COV is not nonnegative definite.

## Algorithm

Routine `CANVC` computes the canonical correlations, the canonical coefficients, Wilks' lambda (for testing the independence of two sets of variates), and a series of tests due to Bartlett for testing that all canonical correlations greater than or equal to the  $k$ -th largest are simultaneously zero. The covariance matrix is used in these computations.

The group 1 variables covariance matrix is first extracted from `COV` and placed in the matrix  $S_{11}$ . Similarly, the group 2 variables covariance matrix is placed in  $S_{22}$ . The "standardized" cross covariance matrix is then computed as:

$$C = \left( S_{11}^{-\frac{1}{2}} \right)^T S_{12} \left( S_{22}^{-\frac{1}{2}} \right)$$

where  $S_{12}$  is the  $NVAR1 \times NVAR2$  matrix of covariances between the group 1 and group 2 variables, and  $S^{1/2}$  denotes the upper triangular Cholesky ( $R^T R$ ) factorization of  $S$ . In the computation of  $C$  and in the following, it is assumed that  $NVAR1$  is greater than  $NVAR2$ . The group 1 and group 2 variables should be interchanged in the following if this is not the case.

The canonical correlations are computed as the singular values of the matrix  $C$ . The canonical coefficients are obtained from the left and right orthogonal matrices resulting from the singular value decomposition of  $C$ . In particular, for  $\Gamma_1 = \text{COEF1}$ .

$$\Gamma_1 = \left( S_{11}^{-\frac{1}{2}} \right) L$$

where  $L$  is the left orthogonal matrix from the singular value decomposition.

Similarly, the correlations between the original variables and the canonical variables,  $R_1 = \text{COEFR1}$ , are obtained for the group 1 variables as:

$$R_1 = \Delta_{11}^{-\frac{1}{2}} \left( S_{11}^{\frac{1}{2}} \right)^T L$$

where  $\Delta_{11}$  is a diagonal matrix containing the diagonal of  $S_{11}$  along its diagonal.

Wilks' lambda, the Bartlett's tests, Rao's  $F$  corresponding to these tests, the numerator and denominator degrees of freedom of  $F$ , and the significance level of  $F$  are computed as in Rao (1973, page 556). Bartlett's tests are computed as

$$\Lambda_i = \prod_{j=i}^q (1 - \rho_j^2)$$

where  $q = NVAR2$  is the number of canonical correlations, the canonical correlations are ordered from largest to smallest, and  $\rho_j$  denotes the  $j$ -th largest canonical correlation. Wilks' lambda is given as  $\Lambda_1$ . The degrees of freedom in the numerator of the corresponding Rao's  $F$  statistic is given as

$$d_1 = pu$$

where  $p = v_1 - i + 1$ ,  $u = v_2 - i + 1$ ,  $v_1 = \text{NVAR2}$ , and  $v_2 = \text{NVAR1}$ . Let

$$m = t - \frac{p + u + 1}{2}$$

where  $t$  is the degrees of freedom in COV, and let

$$s = \sqrt{\frac{p^2 u^2 - 4}{p^2 + u^2 - 5}}$$

if  $p^2 + u^2 - 5 \neq 0$ , and let  $s = 2$  otherwise. Then, Rao's  $F$  corresponding to Bartlett's test is computed as

$$F_i = \frac{1 - \Lambda_i^{\frac{1}{s}}}{\Lambda_i^{\frac{1}{s}}} \frac{ms - pu / 2 + 1}{pu}$$

Rao's  $F$  has numerator degrees of freedom  $d_2 = ms - pu/2 + 1$ . The significance level of  $F$  is obtained from the standard  $F$  distribution.

### Example

The following example is taken from Van de Geer (1971). There are six group 1 variables and two group 2 variables. The maximum correlation turns out to be 0.609.

```

INTEGER      IPRINT, LDCFR1, LDCFR2, LDCOF1, LDCOF2, LDCORR,
&            LDCOV, NDF, NV, NVAR1, NVAR2
PARAMETER    (IPRINT=1, LDCFR1=6, LDCFR2=2, LDCOF1=6, LDCOF2=2,
&            LDCORR=2, LDCOV=8, NDF=100, NV=2, NVAR1=6, NVAR2=2)
C
INTEGER      IND1(NVAR1), IND2(NVAR2)
REAL         COEF1(NVAR1,NVAR1), COEF2(NVAR2,NVAR2),
&            COEFR1(NVAR1,NVAR2), COEFR2(NVAR2,NVAR2),
&            CORR(NVAR2,NVAR1), COV(LDCOV,NVAR1+NVAR2)
EXTERNAL    CANVC
C
DATA COV/1.0000, 0.1839, 0.0489, 0.0186, 0.0782, 0.1147, 0.2137,
&    0.2742, 0.1839, 1.0000, 0.2220, 0.1861, 0.3355, 0.1021,
&    0.4105, 0.4043, 0.0489, 0.2220, 1.0000, 0.2707, 0.2302,
&    0.0931, 0.3240, 0.4047, 0.0186, 0.1861, 0.2707, 1.0000,
&    0.2950, -0.0438, 0.2930, 0.2407, 0.0782, 0.3355, 0.2302,
&    0.2950, 1.0000, 0.2087, 0.2995, 0.2863, 0.1147, 0.1021,
&    0.0931, -0.0438, 0.2087, 1.0000, 0.0760, 0.0702, 0.2137,
&    0.4105, 0.3240, 0.2930, 0.2995, 0.0760, 1.0000, 0.6247,
&    0.2742, 0.4043, 0.4047, 0.2407, 0.2863, 0.0702, 0.6247,
&    1.0000/
C
DATA IND1/1, 2, 3, 4, 5, 6/, IND2/7, 8/
C
CALL CANVC (NDF, NVAR1, NVAR2, COV, LDCOV, IND1, IND2, IPRINT,
&          CORR, LDCORR, COEF1, LDCOF1, COEF2, LDCOF2, COEFR1,
&          LDCFR1, COEFR2, LDCFR2)

```

C  
C

END

### Output

#### \*\*\* Canonical Correlations Statistics \*\*\*

	Canonical Correlations	Wilks	Lambda	Raos F	Num. df	Denom. df	Prob. of Larger F
1	0.6093		0.6159	4.250	12	186	0.0000
2	0.1431		0.9795	0.393	5	94	0.8524

#### Group One Canonical Coefficients

	1	2	3	4	5	6
1	0.326	0.411	-0.799	0.358	-0.032	0.053
2	0.481	-0.340	-0.083	-0.766	-0.484	-0.139
3	0.456	0.718	0.625	0.134	-0.056	0.038
4	0.202	-0.689	0.060	0.732	-0.335	0.080
5	0.184	-0.125	-0.064	-0.045	1.079	-0.225
6	-0.027	-0.174	0.054	-0.086	-0.021	1.017

#### Group Two Canonical Coefficients

	1	2
1	0.464	1.194
2	0.642	-1.108

#### Correlations Between the Group One Variables and the Group One Canonical Scores

	1	2
1	0.4517	0.3408
2	0.7388	-0.2932
3	0.6733	0.4313
4	0.4769	-0.5799
5	0.5299	-0.2811
6	0.1319	-0.0903

#### Correlations Between the Group Two Variables and the Group Two Canonical Scores

	1	2
1	0.8653	0.5013
2	0.9320	-0.3625

# Chapter 10: Discriminant Analysis

---

## Routines

<b>10.1. Parametric Discrimination</b>		
Linear and quadratic discrimination .....	DSCRM	863
Fisher discriminant scores .....	DMSCR	876
<b>10.2. Nonparametric Discrimination</b>		
Nearest neighbor discrimination .....	NNBRD	880

---

## Usage Notes

The routine `DSCRM` (page 863) allows linear or quadratic discrimination and the use of either reclassification, split sample, or the leaving-out-one methods in order to evaluate the rule. Moreover, `DSCRM` can be executed in an online mode, that is, one or more observations can be added to the rule during each invocation of `DSCRM`.

The mean vectors for each group of observations and an estimate of the common covariance matrix for all groups are input to `DMSCR` (page 876). These estimates can be computed via routine `DSCRM`. Output from `DMSCR` are linear combinations of the observations, which at most separate the groups. These linear combinations may subsequently be used for discriminating between the groups. Their use in graphically displaying differences between the groups is possibly more important, however.

Nearest neighbor discrimination is performed in routine `NNBRD` (page 880). In this routine, the user can set the number of nearest neighbors to be used in the discrimination and the threshold for classification. Split samples can also be used.

---

## DSCRM/DDSCRM (Single/Double precision)

Perform a linear or a quadratic discriminant function analysis among several known groups.

## Usage

CALL DSCRM (IDO, NROW, NVAR, NCOL, X, LDX, IND, IFRQ, IWT, IGRP, NGROUP, IMTH, IPRINT, PRIOR, NI, XMEAN, LDXMEA, COV, LDCOV, COEF, LDCOEF, ICLASS, PROB, LDPROB, CLASS, LDCLAS, D2, LDD2, STAT, NRMISS)

## Arguments

**IDO** — Processing option. (Input)

IDO	Action
0	This is the only invocation of DSCRM; all the data are input at once.
1	This is the first invocation of DSCRM with this data, additional calls will be made. Initialization and updating for the NROW observations are performed.
2	This is an intermediate invocation of DSCRM; updating for the NROW observations is performed.
3	All statistics are updated for the NROW observations. The discriminant functions and other statistics are computed.
4	The discriminant functions are used to classify each of the NROW observations in X.
5	The covariance matrices are computed, and workspace is released. No further calls to DSCRM with IDO greater than 1 should be made without first calling DSCRM with IDO = 1.
6	Workspace is released. No further calls to DSCRM with IDO greater than 1 should be made without first calling DSCRM with IDO = 1. This option is not required if a call has been made with IDO = 5 or if workspace is explicitly provided by use of D2CRM.

See Comments 5 and 6 for further information.

**NROW** — The absolute value of NROW is the number of rows of X that contain an observation. (Input)

If NROW is negative, the observations are deleted from the discriminant statistics.

If NROW is positive, they are added.

**NVAR** — Number of variables to be used in the discrimination. (Input)

**NCOL** — Number of columns in matrix X. (Input)

**X** — |NROW| by NVAR + *m* matrix containing the data to be used on this call.

(Input, if |NROW| > 0; X is not referenced otherwise)

*m* is 1, 2, or 3 depending upon whether any columns in X contain frequencies or weights. One column in X must contain the group number for each observation.

Group numbers must be 1.0, 2, 0, ..., NGROUP. If present, IFRQ gives the column containing the frequencies, while IWT gives the column in X containing the weights.

**LDX** — Leading dimension of X exactly as specified in the dimension statement in the calling program. (Input)



**IND** — Vector of length *NVAR* containing the column numbers in *X* to be used in the discrimination. (Input)

**IFRQ** — Frequency option. (Input)

*IFRQ* = 0 means that all frequencies are 1.0. Positive *IFRQ* indicates that column number *IFRQ* of *X* contains the frequencies. All frequencies should be integer values. If this is not the case, the *NINT* (nearest integer) function is used to obtain integer frequencies.

**IWT** — Weighting option. (Input)

*IWT* = 0 means that all weights are 1.0. Positive *IWT* means that column *IWT* of *X* contains the weights. Negative weights are not allowed.

**IGRP** — Column number in *X* containing the group numbers. (Input)

The group numbers must be 1.0, 2.0, ..., *NGROUP* for an observation to be used in the discriminant functions. An observation will be classified regardless of its group number when the reclassification method is specified.

**NGROUP** — Number of groups in the data. (Input)

**IMTH** — Option parameter giving the method of discrimination. (Input)

*IMTH* determines whether linear or quadratic discrimination is used whether the group covariance matrices are computed (the pooled covariance matrix is always computed) and whether the leaving-out-one or the reclassification method is used to classify each observation.

<b>IMTH</b>	<b>Discrim.</b>	<b>Covariance</b>	<b>Classification</b>
1	Linear	All	Reclassification
2	Quadratic	All	Reclassification
3	Linear	Pooled only	Reclassification
4	Linear	All	Leaving-out-one
5	Quadratic	All	Leaving-out-one
6	Linear	Pooled only	Leaving-out-one

In the leaving-out-one method of classification, the posterior probabilities are adjusted so as to eliminate the effect of the observation from the sample statistics prior to its classification. In the reclassification method, the effect of the observation is not eliminated from the classification function. Calls to *DSCRM* with *IMTH* = 1, 2, 4, or 5 can be intermixed, as can calls to *DSCRM* with *IMTH* = 3 or 6. Calls to *DSCRM* with *IMTH* = 1, 2, 4, or 5 cannot be intermixed with calls to *DSCRM* with *IMTH* = 3 or 6 without first calling *DSCRM* with *IDO* = 1 (or 0).

**IPRINT** — Printing option. (Input)

For the given combination of *IDO* and *IPRINT*, the following arrays are printed.

<b>IPRINT</b>	<b>IDO</b>	<b>Printing</b>
0	Any	None
1 or 2	0	PRIOR, NI, XMEAN, COV, COEF, ICLASS, PROB, CLASS, D2, STAT, NRMISS

<b>IPRINT</b>	<b>IDO</b>	<b>Printing</b>
1 or 2	1 or 2	None
1 or 2	3	PRIOR, NI, XMEAN, COEF, D2, STAT, NRMISS
1	4	None
2	4	ICLASS, PROB
1 or 2	5	COV, CLASS
1 or 2	6	None

Note that the only change from `IPRINT = 1` to `IPRINT = 2` is the printing when `IDO = 4`. Also, note that `PRIOR` is printed even though it may be input only.

**PRIOR** — Vector of length `NGROUP` containing the prior probabilities for each group. (Input, if `PRIOR(1)` is not `-1.0` and `IDO` is 0 or 3; input/output, if `PRIOR(1)` is `-1.0` and `IDO` is 0 or 3; input, if `IDO` is 4; not referenced if `IDO` is 1, 2, 5, or 6)

If `PRIOR(1)` is not `-1.0`, then the elements of `PRIOR` should sum to 1.0.

Proportional priors can be selected by setting `PRIOR(1) = -1.0`. In this case, the prior probabilities will be proportional to the sample size in each group, and the elements of `PRIOR` will contain the proportional prior probabilities after the first call with `IDO = 0` or 3.

**NI** — Vector of length `NGROUP`. (Input, for `IDO = 3, 4, or 5`; input/output, for `IDO = 2`; output, for `IDO = 0 or 1`; not referenced if `IDO = 6`)

The  $i$ -th element of `NI` contains the number of observations in group  $i$ .

**XMEAN** — `NGROUP` by `NVAR` matrix. (Input, for `IDO = 3, 4, or 5`; input/output, for `IDO = 2`; output, for `IDO = 0 or 1`; not referenced if `IDO = 6`)

The  $i$ -th row of `XMEAN` contains the group  $i$  variable means.

**LDXMEA** — Leading dimension of `XMEAN` exactly as specified in the dimension statement in the calling program. (Input)

**COV** — `NVAR` by `NVAR` by  $g$  matrix of covariances. (Output, for `IDO = 0 or 1`; input/output, for `IDO = 2, 3, or 5`; input, for `IDO = 4`; not referenced if `IDO = 6`)  $g = \text{NGROUP} + 1$  when `IMTH = 1, 2, 4, or 5`, and  $g = 1$  otherwise. When `IMTH = 3 or 6`, the within-group covariance matrices are not computed. Regardless of the value of `IMTH`, the pooled covariance matrix is always computed and saved as the  $g$ -th covariance matrix in `COV`.

**LDCOV** — Leading and second dimensions of `COV` exactly as specified in the dimension statement of the calling program. (Input)

The first two dimensions of `COV` must be equal.

**COEF** — `NGROUP` by `NVAR + 1` matrix containing the linear discriminant function coefficients. (Output, if `IDO = 0 or 3`; input, if `IDO = 4`; not referenced if `IDO = 1, 2, 5, or 6`)

The first column of `COEF` contains the constant term, and the remaining columns contain the variable coefficients. Row  $i$  of `COEF` corresponds to group  $i$ . `COEF` is always computed as the linear discriminant function coefficients even when quadratic discrimination is specified.

**LD<sub>COEF</sub>** — Leading dimension of **COEF** exactly as specified in the dimension statement in the calling program. (Input)

**ICLASS** — Vector of length **|NROW|** containing the group to which the observation was classified. (Output, if **IDO** = 0 or 4; not referenced otherwise)

If an observation has an invalid group number, frequency, or weight when the leaving-out-one method has been specified, then the observation is not classified and the corresponding elements of **ICLASS** and **PROB** are set to zero.

**PROB** — **|NROW|** by **NGROUP** matrix containing the posterior probabilities for each observation. (Output, if **IDO** = 0 or 4; not referenced otherwise)

**LD<sub>PROB</sub>** — Leading dimension of **PROB** exactly as specified in the dimension statement in the calling program. (Input)

**CLASS** — **NGROUP** by **NGROUP** matrix containing the classification table. (Output, if **IDO** = 0 or 1, input/output, if **IDO** = 4; not referenced otherwise)

Each observation that is classified and has a group number equal to 1.0, 2.0, ..., **NGROUP** is entered into the table. The rows of the table correspond to the known group membership. The columns refer to the group to which the observation was classified. Classification results accumulate with each call to **DSCRM** with **IDO** = 4. For example, if 2 calls with **IDO** = 4 are made, then the elements in **CLASS** sum to the total number of valid observations in the 2 calls.

**LD<sub>CLAS</sub>** — Leading dimension of **CLASS** exactly as specified in the dimension statement in the calling program. (Input)

**D<sub>2</sub>** — **NGROUP** by **NGROUP** matrix containing the Mahalanobis distances

$$D_{ij}^2$$

between the group means. (Output, when **IDO** = 0 or 3; not referenced otherwise)  
For linear discrimination, the Mahalanobis distance is computed using the pooled covariance matrix. Otherwise, the Mahalanobis distance

$$D_{ij}^2$$

between group means *i* and *j* is computed using the within covariance matrix for group *i* in place of the pooled covariance matrix.

**LD<sub>D2</sub>** — Leading dimension of **D<sub>2</sub>** exactly as specified in the dimension statement of the calling program. (Input)

**STAT** — Vector of length  $4 + 2 * (\text{NGROUP} + 1)$  containing statistics of interest. (Input/ Output, if **IDO** = 3 or 5; output, if **IDO** = 0 or 1; not referenced otherwise)

The first element of **STAT** is the sum of the degrees of freedom for the within-covariance matrices. The second, third and fourth elements of **STAT** correspond to the chi-squared statistic, its degrees of freedom, and the probability of a greater chi-squared, respectively, of a test of the homogeneity of the within-covariance matrices (not computed if **IMTH** = 3 or 6). The 5-th through

5 + NGROUP elements of STAT contain the log of the determinants of each group's covariance matrix (not computed if IMTH = 3 or 6) and of the pooled covariance matrix (element 5 + NGROUP). Finally, the last NGROUP + 1 elements of STAT contain the sum of the weights within each group and, in the last position, the sum of the weights in all groups.

**NRMISS** — Number of rows of data encountered in calls to DSCRM containing missing values (NaN) for the classification, group, weight, and/or frequency variables. (Output, if IDO = 0 or 1; input/output, if IDO = 2 or 3, not referenced otherwise)

If a row of data contains a missing value (NaN) for any of these variables, that row is excluded from the computations.

### Comments

1. Automatic workspace usage is

DSCRM (NGROUP + 3) \* NVAR units if IMTH is not 3 or 6. Otherwise, 3 \* NVAR units.

DDSCRM 2 \* (NGROUP + 3) \* NVAR units if IMTH is not 3 or 6. Otherwise, 6 \* NVAR units.

Workspace may be explicitly provided, if desired, by use of D2CRM/DD2CRM. The reference is

```
CALL D2CRM (IDO, NROW, NVAR, NCOL, X, LDX, IND,
            IFRQ, IWT, IGRP, NGROUP, IMTH, IPRINT,
            PRIOR, NI, XMEAN, LDXMEA, COV, LDCOV,
            COEF, LDCOEF, ICLASS, PROB, LDPROB,
            CLASS, LDCLAS, D2, LDD2, STAT, NRMISS,
            D, OB, OBI)
```

The additional arguments are as follows:

**D** — Work vector of length equal to (NGROUP + 1) \* NVAR if IMTH is not 3 or 6, and of length NVAR otherwise.

**OB** — Work vector of length equal to NVAR.

**OBI** — Work vector of length equal to NVAR.

2. Informational errors

Type	Code	
3	1	A row of the data matrix X has an invalid group number.
4	2	The variance-covariance matrix for a group is singular.
4	3	The pooled variance-covariance matrix is singular.
3	4	The variance-covariance matrix for a group is singular. STAT(2) cannot be computed. STAT(2) and STAT(4) are set to the missing value code (NaN).
3	5	An element of PRIOR is less than or equal to $10^{-20}$ .

- |   |   |   |
|---|---|---|
| 3 | 6 | The leaving-out-one method is specified, but this observation does not have a valid weight, or it does not have a valid frequency. This observation is ignored. |
| 3 | 7 | The leaving-out-one method is specified, but this observation does not have a valid group number. This observation is ignored.                                  |
3. Common choices for the Bayesian prior probabilities are given by:  
 $PRIOR(i) = 1.0/NGROUP$  (equal prior probabilities)  
 $PRIOR(i) = NI(i)/NOBS$  (proportional prior probabilities)  
 $PRIOR(i) =$  Past history or subjective judgement  
 In all cases, the prior probabilities should sum to 1.0.
  4. Two passes of the data are made. In the first pass, the statistics required to compute the discriminant functions are obtained ( $IDO = 1, 2,$  and  $3$ ). In the second pass, the discriminant functions are used to classify the observations. When  $IDO = 0$ , all of the data are memory resident, and both passes are made in one call to `DSCRM`. When  $IDO > 0$  and workspace is not explicitly provided by use of `D2CRM`, a third call to `DSCRM` involving no data is required with  $IDO = 5$  or  $6$ .
  5. Here are a few rules and guidelines for the correct value of  $IDO$  in a series of calls.
    - (1) Calls with  $IDO = 0$  or  $1$  may be made at any time. These calls destroy all statistics from previous calls.
    - (2)  $IDO$  may not be  $2, 3, 4, 5,$  or  $6$ 
      - (a) immediately after a call where  $IDO$  was  $0$ ,
      - (b) before a call with  $IDO = 1$  has been made, or
      - (c) immediately after a call with  $IDO = 5$  or  $6$  has been made.
    - (3)  $IDO$  may not be  $4$  or  $5$  before a call with  $IDO = 3$  has been made.
    - (4) Each series of calls to `DSCRM` which begins with  $IDO = 1$  should end with  $IDO = 5$  or  $6$  to ensure the proper release of workspace.

This is a valid sequence of  $IDOs$ :  
 $0, 1, 2, 3, 4, 5, 1, 3, 4, 3, 5, 1, 6, 1, 2, 0, 0, 1, 3, 5.$
  6. Unlike many routines using the parameter  $IDO$ , because of the workspace allocation and saved variables, neither `DSCRM` or `D2CRM` can be called with  $IDO$  greater than  $1$  in consecutive invocations with more than one dataset.

### Algorithm

Routine `DSCRM` performs discriminant function analysis using either linear or quadratic discrimination. The output from `DSCRM` includes a measure of distance between the groups, a table summarizing the classification results, a matrix containing the posterior probabilities of group membership for each observation,

and the within-sample means and covariance matrices. The linear discriminant function coefficients are also computed.

All observations can be input during one call to `DSCRM`, a method of operation that has the advantage of simplicity. Alternatively, one or more rows of observations can be input during separate calls. This method does not require that all observations be memory resident, a significant advantage with large data sets. Note, however, that `DSCRM` requires two passes of the data. During the first pass the discriminant functions are computed while in the second pass, the observations are classified. Thus, with the second method of operation, the data will usually need to be input into `DSCRM` twice.

Because both methods result in the same operations being performed, the algorithm for `DSCRM` is discussed as if only a few observations are input during each call. The operations performed during each call to `DSCRM` depend upon the `IDO` parameter. `IDO = 0` should be used if all observations are to be input at one time.

The `IDO = 1` step is the initialization step. The variables `XMEAN`, `CLASS`, and `COV` are initialized to zero, and other program parameters are set. After this call, all subroutine arguments except `IDO`, `NROW`, `X`, `LDX` and `IMTH` should not be changed by the user except via another call to `DSCRM` with `IDO = 0` or `IDO = 1`. `IMTH` can be changed from one call to the next within the two sets  $\{1, 2, 4, 5\}$  or  $\{3, 6\}$  but not between these sets when `IDO > 1`. That is, do not call `DSCRM` with `IMTH = 1` in one call and `IMTH = 3` in another call without first calling `DSCRM` with `IDO = 1`.

After initialization has been performed in the `IDO = 1` step, the within-group means are updated for all valid observations in `X`. Observations with invalid group numbers are ignored, as are observations with missing values. The *LU* factorization of the covariance matrices are updated by adding (or deleting) observations via Givens rotations.

The `IDO = 2` step is used solely for adding or deleting observations from the model as in the above paragraph.

The `IDO = 3` step begins by adding all observations in `X` to the means and the factorizations of the covariance matrices. It continues by computing some statistics of interest: the linear discriminant functions, the prior probabilities (if `PRIOR(1) = -1.0`), the log of the determinant of each of the covariance matrices, a test statistic for testing that all of the within-group covariance matrices are equal, and a matrix of Mahalanobis distances between the groups. The matrix of Mahalanobis distances is computed via the pooled covariance matrix when linear discrimination is specified, the row covariance matrix is used when the discrimination is quadratic.

Covariance matrices are defined as follows. Let  $N_i$  denote the sum of the frequencies of the observations in group  $i$ , and let  $M_i$  denote the number of observations in group  $i$ . Then, if  $S_i$  denotes the within-group  $i$  covariance matrix,

$$S_i = \frac{1}{N_i - 1} \sum_{j=1}^{M_i} w_j f_j (x_j - \bar{x})(x_j - \bar{x})^T$$

where  $w_j$  is the weight of the  $j$ -th observation in group  $i$ ,  $f_j$  is its frequency,  $x_j$  is the  $j$ -th observation column vector (in group  $i$ ), and  $\bar{x}$  denotes the mean vector of the observations in group  $i$ . The mean vectors are computed as

$$\bar{x} = \frac{1}{W_i} \sum_{j=1}^{M_i} w_j f_j x_j$$

where

$$W_i = \sum_{j=1}^{M_i} w_j f_j$$

Given the means and the covariance matrices, the linear discriminant function for group  $i$  is computed as:

$$z_i = \ln(p_i) - 0.5 \bar{x}_i^T S_p^{-1} \bar{x}_i + x^T S_p^{-1} \bar{x}_i$$

where  $\ln(p_i)$  is the natural log of the prior probability for the  $i$ -th group,  $x$  is the observation to be classified, and  $S_p$  denotes the pooled covariance matrix.

Let  $S$  denote either the pooled covariance matrix or one of the within-group covariance matrices  $S_i$ . ( $S$  will be the pooled covariance matrix in linear discrimination, and  $S_i$  otherwise.) The Mahalanobis distance between group  $i$  and group  $j$  is computed as:

$$D_{ij}^2 = (\bar{x}_i - \bar{x}_j)^T S^{-1} (\bar{x}_i - \bar{x}_j)$$

Finally, the asymptotic chi-squared test for the equality of covariance matrices is computed as follows (Morrison 1976, page 252):

$$\gamma = C^{-1} \sum_{i=1}^k n_i \left\{ \ln(|S_p|) - \ln(|S_i|) \right\}$$

where  $n_i$  is the number of degrees of freedom in the  $i$ -th sample covariance matrix,  $k$  is the number of groups, and

$$C^{-1} = 1 - \frac{2p^2 + 3p - 1}{6(p+1)(k-1)} \left( \sum_{i=1}^k \frac{1}{n_i} - \frac{1}{\sum_j n_j} \right)$$

where  $p$  is the number of variables.

When  $\text{IDO} = 4$ , the estimated posterior probability of each observation  $x$  belonging to group  $i$  is computed using the prior probabilities and the sample mean vectors and estimated covariance matrices under a multivariate normal assumption. Under quadratic discrimination, the within-group covariance

matrices are used to compute the estimated posterior probabilities. The estimated posterior probability of an observation  $x$  belonging to group  $i$  is

$$\hat{q}_i(x) = \frac{e^{-\frac{1}{2}D_i^2(x)}}{\sum_{j=1}^k e^{-\frac{1}{2}D_j^2(x)}}$$

where

$$D_i^2(x) = \begin{cases} (x - \bar{x}_i)^T S_i^{-1} (x - \bar{x}_i) + \ln|S_i| - 2 \ln(p_i) & \text{IMTH} = 1 \text{ or } 2 \\ (x - \bar{x}_i)^T S_p^{-1} (x - \bar{x}_i) - 2 \ln(p_i) & \text{IMTH} = 3 \end{cases}$$

For the leaving-out-one method of classification (IMTH = 4, 5, and 6), the sample mean vector and sample covariance matrices in the formula for

$$D_i^2(x)$$

are adjusted so as to remove the observation  $x$  from their computation. For linear discrimination (IMTH = 1, 2, 4, and 6), the linear discriminant function coefficients are actually used to compute the same posterior probabilities.

Using the posterior probabilities, each observations in  $x$  is classified into a group; the result is tabulated in the matrix CLASS and saved in the vector ICLASS. CLASS is not altered at this stage if  $x(i, \text{IGRP})$  contains a group number that is out of range. If the reclassification method is specified, then all observations with no missing values in the NVAR classification variables are classified. When the leaving-out-one method is used, observations with invalid group numbers, weights, frequencies or classification variables are not classified. Regardless of the frequency, a 1 is added (or subtracted) from CLASS for each row of  $x$  that is classified and contains a valid group number.

When IMTH > 3, adjustment is made to the posterior probabilities to remove the effect of the observation in the classification rule. In this adjustment, each observation is presumed to have a weight of  $x(i, \text{IWT})$ , if IWT > 0 and a frequency of 1.0. See Lachenbruch (1975, page 36) for the required adjustment.

Finally, when IDO = 5, the covariance matrices are computed from their LU factorizations.

### Example 1

The following example uses linear discrimination with equal prior probabilities on Fisher's (1936) iris data. This example illustrates the execution of DSCRM when one call is made.

```

INTEGER      IDO, IFRQ, IGRP, IMTH, IPRINT, IWT, LDCLAS, LDCOEF,
&            LDCOV, LDD2, LDPROB, LDX, LDXMEA, NCOL, NGROUP, NROW,
&            NVAR
PARAMETER    (IDO=0, IFRQ=0, IGRP=1, IMTH=3, IPRINT=1, IWT=0,
&            LDCOV=4, NCOL=5, NGROUP=3, NROW=150, NVAR=4,
&            LDCLAS=NGROUP, LDCOEF=NGROUP, LDD2=NGROUP,
&            LDPROB=NROW, LDX=NROW, LDXMEA=NGROUP)

```

C



```

      INTEGER      ICLASS(NROW), IND(4), NI(NGROUP), NOBS, NRMISS, NV
      REAL         CLASS(LDCLAS,NGROUP), COEF(LDCOEF,NVAR+1),
&                COV(LDCOV,LDCOV,1), D2(LDD2,NGROUP), PRIOR(3),
&                PROB(LDPROB,NGROUP), STAT(6+2*NGROUP), X(LDX,5),
&                XMEAN(LDXMEA,NVAR)
      EXTERNAL    DSCRM, GDATA
C
      DATA IND/2, 3, 4, 5/, PRIOR/0.3333333, 0.3333333, 0.3333333/
C
      CALL GDATA (3, 0, NOBS, NV, X, LDX, 5)
C
      CALL DSCRM (IDO, NROW, NVAR, NCOL, X, LDX, IND, IFRQ, IWT, IGRP,
&              NGROUP, IMTH, IPRINT, PRIOR, NI, XMEAN, LDXMEA, COV,
&              LDCOV, COEF, LDCOEF, ICLASS, PROB, LDPROB, CLASS,
&              LDCLAS, D2, LDD2, STAT, NRMISS)
C
      END

```

### Output

PRIOR, the prior probabilities

1	2	3
0.3333	0.3333	0.3333

NI, the number in each group

1	2	3
50	50	50

XMEAN, the group means

	1	2	3	4
1	5.006	3.428	1.462	0.246
2	5.936	2.770	4.260	1.326
3	6.588	2.974	5.552	2.026

The pooled within-groups covariance matrix

	1	2	3	4
1	0.2650	0.0927	0.1675	0.0384
2	0.0927	0.1154	0.0552	0.0327
3	0.1675	0.0552	0.1852	0.0427
4	0.0384	0.0327	0.0427	0.0419

COEF, the discriminant function coefficients

	1	2	3	4	5
1	-86.3	23.5	23.6	-16.4	-17.4
2	-72.9	15.7	7.1	5.2	6.4
3	-104.4	12.4	3.7	12.8	21.1

ICLASS, the classifications

Obs.	Class
1	1
2	1
3	1
4	1
5	1
6	1

```

      .
      .
      .
145   3
146   3
147   3
148   3
149   3
150   3

PROB, the posterior probabilities
      1      2      3
  1  1.000  0.000  0.000
  2  1.000  0.000  0.000
  3  1.000  0.000  0.000
  4  1.000  0.000  0.000
  5  1.000  0.000  0.000
  6  1.000  0.000  0.000
      .
      .
145  0.000  0.000  1.000
146  0.000  0.000  1.000
147  0.000  0.006  0.994
148  0.000  0.003  0.997
149  0.000  0.000  1.000
150  0.000  0.018  0.982

CLASS, the classification table
      1      2      3
  1  50.00  0.00  0.00
  2   0.00  48.00  2.00
  3   0.00   1.00  49.00

D2, the distances between group means
      1      2      3
  1   0.0   89.9  179.4
  2  89.9   0.0   17.2
  3 179.4  17.2   0.0

      1      2      3      4      5      STAT
147.0   NaN   NaN   NaN   NaN   NaN   6
      7      8      9     10
      NaN  -10.0  50.0  50.0

11     12
50.0  150.0

NRMISS, number of missing observations = 0

```

### Example 2

Continuing with Fisher's iris data, the following example computes the quadratic discriminant functions using values of `IDO > 0`. In the first loop, all observations are added to the functions, two observations at a time. In the second loop, each of three observations is classified, one by one, using the leaving-out-one method. Output for statistics that are identical to those reported in the first example are not printed here.

```

    INTEGER   IFRQ, IGRP, IMTH, IWT, LDCLAS, LDCOEF, LDCOV, LDD2,
&            LDPROB, LDX, LDXMEA, NCOL, NGROUP, NROW, NVAR
    PARAMETER (IFRQ=0, IGRP=1, IMTH=2, IWT=0, LDPROB=10, LDX=150,
&            NCOL=5, NGROUP=3, NROW=1, NVAR=4, LDCLAS=NGROUP,
&            LDCOEF=NGROUP, LDCOV=NVAR, LDD2=NGROUP, LDXMEA=NGROUP)
C
    INTEGER   I, ICLASS(LDPROB), IDO, IND(4), IPRINT, NI(NGROUP),
&            NOBS, NRMISS, NV
    REAL      CLASS(LDCLAS,NGROUP), COEF(LDCOE, NVAR+1),
&            COV(LDCOV, LDCOV, NGROUP+1), D2(LDD2, NGROUP), PRIOR(3),
&            PROB(LDPROB, NGROUP), STAT(6+2*NGROUP), X(LDX, 5),
&            XMEAN(LDXMEA, NVAR)
    EXTERNAL  DSCRM, GDATA, WRRRN
C
    DATA IND/2, 3, 4, 5/, PRIOR/0.3333333, 0.3333333, 0.3333333/
C
    CALL GDATA (3, 0, NOBS, NV, X, LDX, 5)
C
    IPRINT = 0
    IDO = 1
    CALL DSCRM (IDO, 0, NVAR, NCOL, X, LDX, IND, IFRQ, IWT, IGRP,
&            NGROUP, IMTH, IPRINT, PRIOR, NI, XMEAN, LDXMEA, COV,
&            LDCOV, COEF, LDCE, ICLASS, PROB, LDPROB, CLASS,
&            LDCLAS, D2, LDD2, STAT, NRMISS)
C
    Add the observations
    IDO = 2
    DO 10 I=1, NOBS
        CALL DSCRM (IDO, NROW, NVAR, NCOL, X(I,1), LDX, IND, IFRQ,
&            IWT, IGRP, NGROUP, IMTH, IPRINT, PRIOR, NI,
&            XMEAN, LDXMEA, COV, LDCOV, COEF, LDCE, ICLASS,
&            PROB, LDPROB, CLASS, LDCLAS, D2, LDD2, STAT,
&            NRMISS)
10 CONTINUE
C
    Summarize the statistics
    IDO = 3
    CALL DSCRM (IDO, 0, NVAR, NCOL, X, LDX, IND, IFRQ, IWT, IGRP,
&            NGROUP, IMTH, IPRINT, PRIOR, NI, XMEAN, LDXMEA, COV,
&            LDCOV, COEF, LDCE, ICLASS, PROB, LDPROB, CLASS,
&            LDCLAS, D2, LDD2, STAT, NRMISS)
C
    Classify the first three observations
    IPRINT = 2
    IDO = 4
    DO 20 I=1, 3
        CALL DSCRM (IDO, NROW, NVAR, NCOL, X(I,1), LDX, IND, IFRQ,
&            IWT, IGRP, NGROUP, IMTH, IPRINT, PRIOR, NI,
&            XMEAN, LDXMEA, COV, LDCOV, COEF, LDCE,
&            ICLASS(I), PROB(I,1), LDPROB, CLASS, LDCLAS, D2,
&            LDD2, STAT, NRMISS)
20 CONTINUE
C
    Release Workspace
    IDO = 6
    CALL DSCRM (IDO, 0, NVAR, NCOL, X, LDX, IND, IFRQ, IWT, IGRP,
&            NGROUP, IMTH, IPRINT, PRIOR, NI, XMEAN, LDXMEA, COV,
&            LDCOV, COEF, LDCE, ICLASS, PROB, LDPROB, CLASS,
&            LDCLAS, D2, LDD2, STAT, NRMISS)
C
    END

```

## Output

ICLASS, the classifications

```
Obs.  Class
1     1
```

PROB, the posterior probabilities

```
      1      2      3
1.000  0.000  0.000
```

ICLASS, the classifications

```
Obs.  Class
1     1
```

PROB, the posterior probabilities

```
      1      2      3
1.000  0.000  0.000
```

ICLASS, the classifications

```
Obs.  Class
1     1
```

PROB, the posterior probabilities

```
      1      2      3
1.000  0.000  0.000
```

---

## DMSCR/DDMSCR (Single/Double precision)

Use Fisher's linear discriminant analysis method to reduce the number of variables.

### Usage

```
CALL DMSCR (NGROUP, NVAR, XMEAN, LDXMEA, SUMWT, COV, LDCOV,
            NNV, EVAL, COEF, LDCOEF, CMEAN, LDCMEA)
```

### Arguments

**NGROUP** — Number of groups. (Input)

**NVAR** — Number of variables. (Input)

**XMEAN** —  $NGROUP$  by  $NVAR$  matrix containing the means of the variables in each group. (Input)

**LDXMEA** — Leading dimension of **XMEAN** exactly as specified in the dimension statement in the calling program. (Input)

**SUMWT** — Vector of length  $NGROUP$  containing the sum of the weights of the observations in each group. (Input)

**COV** —  $NVAR$  by  $NVAR$  matrix containing the pooled within-groups variance-covariance matrix  $S_p$ . (Input)

**LDCOV** — Leading dimension of COV exactly as specified in the dimension statement in the calling program. (Input)

**NNV** — Number of eigenvectors extracted from

$$S_p^{-1} S_b$$

the standardized between-groups variance-covariance matrix. (Output)  
 $S_p$  is the pooled within-groups variance-covariance matrix, and  $S_b$  is the between-groups variance-covariance matrix. NNV is usually the minimum of NVAR and NGROUP - 1, but it may be smaller if any row of XMEAN or COV is a linear combination of the other rows.

**EVAL** — Vector of length NNV containing the eigenvalues extracted from the standardized between-means variance-covariance matrix, in descending order. (Output)

NNV is less than or equal to the minimum of NVAR and (NGROUP - 1).

**COEF** — NVAR by NNV matrix of eigenvectors from the standardized between-means variance-covariance matrix. (Output)

The eigenvector coefficients have been standardized such that the canonical scores can be obtained directly by multiplication of the original data by COEF.

**LDCOEF** — Leading dimension of COEF exactly as specified in the dimension statement in the calling program. (Input)

**CMEAN** — NGROUP by NNV matrix of group means of the canonical variables. (Output)

**LDCMEA** — Leading dimension of CMEAN exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Automatic workspace usage is

DMSCR 3 \* NVAR \* (NVAR + 1) units, or  
DDMSCR 6 \* NVAR \* (NVAR + 1) units.

Workspace may be explicitly provided, if desired, by use of  
D2SCR/DD2SCR. The reference is

```
CALL D2SCR (NGROUP, NVAR, XMEAN, LDXMEA, SUMWT, COV,  
           LDCOV, NNV, EVAL, COEF, LDCOEF,  
           CMEAN, LDCMEA, BCOV, EVAL2, EVEC, WKR,  
           WK)
```

The additional arguments are as follows:

**BCOV** — Work array of length NVAR \* NVAR.

**EVAL2** — Work array of length NVAR.

**EVEC** — Work array of length NVAR \* NVAR.

**WKR** — Work array of length NVAR \* NVAR.

**WK** — Work array of length  $2 * NVAR$ .

2. IMSL routine `DSCRM` (page 863) may be used to calculate the input arrays for this routine from the original data.

### Algorithm

Routine `DMSCR` is a natural generalization of R.A. Fisher's linear discrimination procedure for two groups. This method of discrimination obtains those linear combinations of the observed random variables that maximize the between-groups variation relative to the within-groups variation. Denote the first of these linear combinations by

$$z_1 = \beta_1^T x$$

where  $\beta_1$  is a column vector of coefficients of length  $NVAR$  and  $x$  is an observation to be classified. On the basis of one linear combination, the discriminant rule assigns the observation,  $z$ , to a group (characterized by the group mean) by minimizing the Euclidean distance between  $z$  and the group mean.

To obtain  $\beta_1$  (see, e.g., Tatsuoka 1971, page 158), let  $S_p$  denote the pooled within-groups covariance matrix ( $S_p$  is defined and can be computed via routine `DSCRM`, page 863) and let  $S_b$  denote the between-groups covariance matrix defined by

$$S_b = \sum_{i=1}^g w_i (\bar{x}_i - \bar{x})(\bar{x}_i - \bar{x})^T / (N - g)$$

where  $g$  is the number of groups,

$$\bar{x}_i$$

is the mean vector for the  $i$ -th group of observations,  $\bar{x}$  denotes the vector of means over all observations,  $w_i$  is the sum of the weights times the frequencies as input in `SUMWT` and as used in the computation of

$$\bar{x}_i$$

and  $N$  is the total number of observations used in computing `COV`. Then,  $\beta_1$ , such that

$$\beta_1^T S_p \beta_1 = 1$$

can be computed as the maximum of

$$\Psi = \beta_1^T S_b \beta_1$$

This yields  $\beta_1$  as the eigenvector associated with the largest eigenvalue from

$$S_p^{-1} S_b$$

Generally,

$$S_p^{-1} S_b$$

has rank  $m$ , where  $m = \min(g - 1, p)$  and  $p = \text{NVAR}$ .

$$S_p^{-1} S_b$$

has  $m$  such eigenvectors, and the matrix COEF is obtained as  $(\beta_1, \beta_2, \dots, \beta_m)$ , where each  $\beta_i$  is an eigenvector.

The matrix CMEAN is taken as the within-group means vector of the linear combinations  $z_i$  defined by the  $\beta$ 's. For each observation  $x$ , scores

$$z_i = \beta_i^T x$$

can be computed, because of the restriction on  $\beta_i$ , the sample variance of the  $z_i$  is 1.0. The observation is classified into the group (as specified by the group mean of the  $z_i$ 's) to which, on the basis of the  $z_i$ , the Euclidean distance is the least.

Note that the linear combinations  $z_i$  have meaning even when discrimination is not desired. The linear combination of the observed variables that most separates the  $g$  groups is  $z_1$ ;  $z_2$ , giving the second highest such separation orthogonal to the first, and so on. Thus, a plot of the mean vectors of the first two variables gives a good two-dimensional summarization of the relationships between the groups.

### Example

The following example illustrates a typical sequence. Fisher's iris data is used. (See routine GDATA, page 1302). Routine DSCRM is first used to perform a discriminant analysis based on all the variables. COV, XMEAN, and NI are obtained from DSCRM. Function DMSCR, which uses these arrays, is then called.

```

C      INTEGER      IDO, IFRQ, IGRP, IMTH, IPRINT, IWT, LDCLAS, LDCMEA,
&      LDCO, LDCOEF, LDCOV, LDD2, LDPROB, LDX, LDXMEA, NCOL,
&      NGROUP, NROW, NVAR
C      PARAMETER    (IDO=0, IFRQ=0, IGRP=1, IMTH=3, IPRINT=0, IWT=0,
&      LDCOV=4, NCOL=5, NGROUP=3, NROW=150, NVAR=4,
&      LDCLAS=NGROUP, LDCMEA=NGROUP, LDCO=NGROUP,
&      LDCOEF=NVAR, LDD2=NGROUP, LDPROB=NROW, LDX=NROW,
&      LDXMEA=NGROUP)
C
C      INTEGER      ICLASS(NROW), IND(4), NI(NGROUP), NNV, NOBS, NOUT,
&      NRMIS, NV
C      REAL         CLASS(LDCLAS,NGROUP), CMEAN(LDCMEA,NGROUP-1),
&      CO(LDCO,NVAR+1), COEF(LDCOEF,NGROUP-1),
&      COV(LDCOV,LDCOV,1), D2(LDD2,NGROUP), EVAL(NGROUP-1),
&      PRIOR(3), PROB(LDPROB,NGROUP), REAL,
&      STAT(6+2*NGROUP), SUMWT(NGROUP), X(LDX,5),
&      XMEAN(LDXMEA,NVAR)
C      INTRINSIC    REAL
C      EXTERNAL     DMSCR, DSCRM, GDATA, UMACH, WRRRN
C

```

```

DATA IND/2, 3, 4, 5/, PRIOR/0.3333333, 0.3333333, 0.3333333/
C
CALL GDATA (3, 0, NOBS, NV, X, LDX, 5)
C
CALL DSCRM (IDO, NROW, NVAR, NCOL, X, LDX, IND, IFRQ, IWT, IGRP,
&          NGROUP, IMTH, IPRINT, PRIOR, NI, XMEAN, LDXMEA, COV,
&          LDCOV, CO, LDCO, ICLASS, PROB, LDPROB, CLASS,
&          LDCLAS, D2, LDD2, STAT, NRMISS)
C
SUMWT(1) = STAT(6+NGROUP)
SUMWT(2) = STAT(7+NGROUP)
SUMWT(3) = STAT(8+NGROUP)
C
CALL DMSCR (NGROUP, NVAR, XMEAN, LDXMEA, SUMWT, COV, LDCOV, NNV,
&          EVAL, COEF, LDcoef, CMEAN, LDCMEA)
C
CALL UMACH (2, NOUT)
WRITE (NOUT, '( '' NNV = '' ,I1) ' ) NNV
CALL WRRRN ('EVAL', 1, NNV, EVAL, 1, 0)
CALL WRRRN ('COEF', NVAR, NNV, COEF, LDcoef, 0)
CALL WRRRN ('CMEAN', NGROUP, NNV, CMEAN, LDCMEA, 0)
END

```

### Output

NNV = 2

EVAL	
1	2
32.19	0.29

COEF		
	1	2
1	-0.829	0.024
2	-1.534	2.165
3	2.201	-0.932
4	2.810	2.839

CMEAN		
	1	2
1	-5.502	6.877
2	3.930	5.934
3	7.888	7.174

---

## NNBRD/DNNBRD (Single/Double precision)

Perform  $k$  nearest neighbor discrimination.

### Arguments

```

CALL NNBRD (NROW, NVAR, NCOL, X, LDX, K, IND, IGRP, NGROUP,
NRULE, NCLASS, METRIC, PRIOR, THRESH, PART,
IDISCR, NI, ICLASS, PROB, LDPROB, CLASS,
LDCLAS)

```



## Arguments

**NROW** — Number of rows of  $X$  that contain an observation. (Input)

**NVAR** — Number of variables to be used in the discrimination. (Input)

**NCOL** — Number of columns in matrix  $X$ . (Input)

**X** —  $NROW$  by  $NVAR + 1$  matrix containing the data to be used on this call.  
(Input/Output)

One column in  $X$  must contain the group number for each observation. On output,  $X$  is sorted into a  $k$ - $d$  tree. The first  $NRULE + NCLASS$  rows of  $X$  must not contain missing values in the columns specified by **IND** and **IGRP**.

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

**K** — Number of nearest neighbors to be used in the discriminant rule. (Input)

**IND** — Vector of length  $NVAR$  containing the column numbers in  $X$  to be used in the discrimination. (Input)

**IGRP** — Column number in  $X$  containing the group numbers. (Input)

The group numbers must be 1.0, 2.0, ...,  $NGROUP$  for an observation to be used in the discriminant functions. (Note, however, that the nearest integer (**NINT**) function is used to obtain the group numbers.)

**NGROUP** — Number of groups in the data. (Input)

**NRULE** — Number of observations in  $X$  to be used in the discriminant rule.  
(Input)

The first  $|NRULE|$  observations in  $X$  are used as the set defining the rule. If **NRULE** is positive, then the **NRULE** observations defining the rule are classified. If **NRULE** is negative, the **NRULE** observations defining the rule are not classified.

**NCLASS** — Number of observations in  $X$  to classify. (Input)

**NCLASS** is the number of observations in a second sample that may be used to test the rule formed from the first **NRULE** observations. If present, this sample is in rows  $NRULE + 1$  through  $NRULE + NCLASS$  of  $X$ .

**METRIC** — Metric to be used in computing the  $k$  nearest neighbors. (Input)

### **METRIC** Metric used

0      Euclidean distance

1       $L_1$  norm

2       $L_\infty$  norm

**PRIOR** — Vector of length  $NGROUP$  containing the prior probabilities for each group. (Input, if **PRIOR(1)** is not  $-1.0$ ; input/output, if **PRIOR(1)** is  $-1.0$ )

If **PRIOR(1)** is not  $-1.0$ , then the elements of **PRIOR** should sum to 1.0.

Proportional priors can be selected by setting **PRIOR(1)** =  $-1.0$ . In this case, the prior probabilities will be proportional to the sample size in each group based upon the first **NRULE** observations, and the elements of **PRIOR** will contain the proportional prior probabilities on return from **NNBRD**.

**THRESH** — Threshold for the posterior probabilities. (Input)

If the maximum posterior probability is less than **THRESH**, the observation is classified into group **NGROUP** + 1 (the group “other”).

**PART** — Vector of length **NRULE** containing the values to be used in the partition of **x** for the *k-d* tree. (Output)

**IDISCR** — Vector of length **NRULE** containing the element number in **IND** that points to the column of **x** to be used as the discriminator in the *k-d* tree. (Output)  
**IDISCR**(*i*) = 0 if the observation is a terminal node. **IND**(**IDISCR**(*i*)) is the column number in **x** to be used as the discriminator.

**NI** — Vector of length **NGROUP** containing the number of observations in each group. (Output)

**ICLASS** — Vector of length *m* containing the group to which the observation was classified. (Output)

If **NRULE** > 0, *m* = **NRULE** + **NCLASS**; otherwise, *m* = **NCLASS**. The *i*-th element in **ICLASS** corresponds to to *i*-th row in the sorted matrix **x**.

**PROB** — *m* by **NGROUP** matrix containing the posterior probabilities for each observation. (Output)

The *i*-th row in **PROB** corresponds to the *i*-th row in the in the sorted matrix **x**.

**LDPROB** — Leading dimension of **PROB** exactly as specified in the dimension statement in the calling program. (Input)

**CLASS** — **NGROUP** by **NGROUP** + 1 matrix containing the classification table. (Output)

Each observation that is classified and has a group number equal to 1.0, 2.0, ..., **NGROUP** is entered into the table. The rows of the table correspond to the known group membership. The columns refer to the group to which the observation was classified. Column **NGROUP** + 1 refers to the column “other” (see **THRESH**).

**LDCLAS** — Leading dimension of **CLASS** exactly as specified in the dimension statement in the calling program. (Input)

## Comments

Automatic workspace usage is

NNBRD  $(2 * \text{NVAR} + 3) * (\log_2(\text{NROW}) + 3) + (2 * \text{NROW}) + (2 * \text{K} + 2) + \text{NVAR}$   
units, or

DNNBRD  $(4 * \text{NVAR} + 3) * (\log_2(\text{NROW}) + 3) + (4 * \text{NROW}) + (3 * \text{K} + 2) + 2 * \text{NVAR}$   
units.

Workspace may be explicitly provided, if desired, by use of **N2BRD**/**DN2BRD**. The reference is

```
CALL N2BRD (NROW, NVAR, NCOL, X, LDX, K, IND, IGRP, NGROUP,
           NRULE, NCLASS, METRIC, PRIOR, THRESH, PART,
           IDISCR, NI, ICLASS, PROB, LDPROB, CLASS,
```

LDCLAS, WK, IWK, ILOW, IHIGH, ISIDE, BNDL,  
BNDH, XKEY, IPQR, PQD)

The additional arguments are as follows:

**WK** — Work vector of length NROW.

**IWK** — Work vector of length NROW.

**ILOW** — Work vector of length  $\log_2(\text{NROW}) + 3$ .

**IHIGH** — Work vector of length  $\log_2(\text{NROW}) + 3$ .

**ISIDE** — Work vector of length  $\log_2(\text{NROW}) + 3$ .

**BNDL** — Work vector of length  $\text{NVAR} * (\log_2(\text{NROW}) + 3)$ .

**BNDH** — Work vector of length  $\text{NVAR} * (\log_2(\text{NROW}) + 3)$ .

**XKEY** — Work vector of length NVAR.

**IPQR** — Work vector of length  $\kappa + 1$ .

**PQD** — Work vector of length  $\kappa + 1$ .

### Algorithm

Routine NNBRD performs  $k$ -th nearest neighbor discriminant function analysis. The  $k$ - $d$  tree algorithm of Friedman, Bentley, and Finkel (1977) is used to find the nearest neighbors. Consult this reference for a discussion of  $k$ - $d$  trees and how one goes about finding nearest neighbors in them.

In NNBRD, the  $k$  nearest neighbors of any observation used in forming the rule (i.e., one of the first NRULE observations in  $\mathbf{x}$ ), do not include the observation. Let  $k_i (i = 1, \dots, \text{NGROUP})$  denote the number of nearest neighbors found from each of the groups for a given observation ( $\sum_i k_i = k$ ); let  $p_i = \text{PRIOR}(i) (\sum_i p_i = 1)$ ; and let

$$\hat{\theta}_i$$

denote the estimated posterior probability of membership in group  $i$ . Compute

$$\hat{\theta}_i \text{ as } \hat{\theta}_i = \frac{k_i p_i / n_i}{\sum_{j=1}^g k_j p_j / n_j}$$

where  $g = \text{NGROUP}$ . (If  $n_j = 0$  for some  $j$ , the associated term in the denominator is excluded and

$$\hat{\theta}_j$$

is set to 0.0.)

Let  $m$  denote the index of the maximum

$$\hat{\theta}_i$$

and  $\phi = \text{THRESH}$ . Then if

$$\hat{\theta}_m > \phi$$

the observation is classified into group  $m$ . If

$$\hat{\theta}_m \leq \phi$$

or if the maximum  $\hat{\theta}$  is not unique, then the observation is not classified into any group and ICLASS is set to zero.

Three metrics are available in NNBRD for finding the nearest neighbors. These are Euclidean ( $L_2$ ) distance,  $L_1$  norm, and  $L_\infty$  norm. In order to use Mahalanobis distance,  $x^T \Sigma^{-1} x$ , a transformation  $y = \Sigma^{-1/2} x$  is first needed so that  $\text{Var}(y) = I$ . These transformations can be accomplished by use of the mathematical routines. The  $L_2$  norm would then be used with  $y$  as input to obtain the Mahalanobis metric.

### Example

Fisher's iris data are used to illustrate routine NNBRD. The data consist of three types of iris. NNBRD is called with  $k = 5$  and Euclidean distance as the metric. The results show a clear separation of the groups.

```

INTEGER      IGRP, K, LDCLAS, LDPROB, LDX, METRIC, NCLASS, NCOL,
&            NGROUP, NROW, NRULE, NVAR
REAL         THRESH
PARAMETER   (IGRP=1, K=5, LDCLAS=3, LDPROB=150, LDX=150,
&            METRIC=0, NCLASS=0, NCOL=5, NGROUP=3, NROW=150,
&            NRULE=150, NVAR=4, THRESH=0.10)
C
INTEGER      ICLASS(NROW), IDISCR(NROW), IND(NVAR), NI(NGROUP),
&            NRA, NRB
REAL         CLASS(LDCLAS,NGROUP+1), PART(NRULE), PRIOR(NGROUP),
&            PROB(LDPROB,NGROUP), X(LDX,NCOL)
EXTERNAL    GDATA, NNBRD, WRIRN, WRRRN
C
DATA IND/2, 3, 4, 5/
C
CALL GDATA (3, 0, NRA, NRB, X, 150, 5)
C
PRIOR(1) = -1.0
CALL NNBRD (NROW, NVAR, NCOL, X, LDX, K, IND, IGRP, NGROUP,
&            NRULE, NCLASS, METRIC, PRIOR, THRESH, PART, IDISCR,
&            NI, ICLASS, PROB, LDPROB, CLASS, LDCLAS)
CALL WRRRN ('The first 10 rows of X', 10, NCOL, X, LDX, 0)
CALL WRRRN ('PRIOR', 1, NGROUP, PRIOR, 1, 0)
CALL WRRRN ('The first 10 elements of PART', 1, 10, PART, 1, 0)
CALL WRIRN ('The first 10 elements of IDISCR', 1, 10, IDISCR, 1,
&            0)
CALL WRIRN ('NI', 1, NGROUP, NI, 1, 0)
CALL WRIRN ('The first 10 elements of ICLASS', 1, 10, ICLASS, 1,
&            0)
CALL WRRRN ('The first 10 rows of PROB', 10, NGROUP, PROB,
&            LDPROB, 0)
CALL WRRRN ('CLASS', NGROUP, NGROUP, CLASS, LDCLAS, 0)

```

C

END

### Output

The first 10 rows of X

	1	2	3	4	5
1	1.000	4.500	2.300	1.300	0.300
2	1.000	4.400	2.900	1.400	0.200
3	1.000	4.800	3.000	1.400	0.300
4	1.000	4.400	3.000	1.300	0.200
5	1.000	4.800	3.000	1.400	0.100
6	1.000	4.300	3.000	1.100	0.100
7	1.000	4.600	3.100	1.500	0.200
8	1.000	4.900	3.100	1.500	0.100
9	1.000	4.900	3.000	1.400	0.200
10	1.000	4.900	3.100	1.500	0.200

PRIOR

	1	2	3
0.3333	0.3333	0.3333	

The first 10 elements of PART

	1	2	3	4	5	6	7	8	9	10
0.000	0.000	3.000	0.000	3.000	0.000	0.000	4.900	0.000	3.100	

The first 10 elements of IDISCR

1	2	3	4	5	6	7	8	9	10
0	0	2	0	2	0	0	1	0	2

NI

	1	2	3
50	50	50	

The first 10 elements of ICLASS

1	2	3	4	5	6	7	8	9	10
1	1	1	1	1	1	1	1	1	1

The first 10 rows of PROB

	1	2	3
1	1.000	0.000	0.000
2	1.000	0.000	0.000
3	1.000	0.000	0.000
4	1.000	0.000	0.000
5	1.000	0.000	0.000
6	1.000	0.000	0.000
7	1.000	0.000	0.000
8	1.000	0.000	0.000
9	1.000	0.000	0.000
10	1.000	0.000	0.000

CLASS

	1	2	3
1	50.00	0.00	0.00
2	0.00	47.00	3.00
3	0.00	2.00	48.00

# Chapter 11: Cluster Analysis

---

## Routines

11.1.	<b>Hierarchical Cluster Analysis</b>		
	Compute distance or similarity matrix .....	CDIST	889
	Hierarchical cluster analysis.....	CLINK	892
	Retrieve cluster numbers in hierarchical cluster analysis ...	CNUMB	897
11.2.	<b>K-means Cluster Analysis</b>		
	The basic K-means algorithm .....	KMEAN	900

---

## Usage Notes

The routines described in this chapter perform various forms of hierarchical or *K*-means cluster analysis. By appropriate manipulation of the input data, either variables or cases may be clustered. Additionally, for hierarchical clustering, similarity or dissimilarity (distance) matrices created by routines not included in this chapter can be clustered. Hartigan (1975) and Anderberg (1973) are general references that may be used in this chapter.

The first step in agglomerative hierarchical cluster analysis is to compute the distance between each observation (or variable). Initially, each observation (variable) is treated as a cluster. The two clusters that are closest to one another in distance are merged, and the distance of the new cluster from all other clusters is computed. This process continues until only one cluster remains. No attempt at finding an optimal clustering (in the sense of minimizing some criterion) is made.

The usual steps in a hierarchical cluster analysis might proceed as follows:

1. Routine CDIST (page 889) is used to compute a distance (or possibly a similarity) matrix from the input data matrix. A scaled matrix of Euclidean distances is a common choice for a distance matrix, while a correlation matrix is a common choice for a similarity matrix. If a correlation matrix is to be used, many of the routines described in Chapter 3, “Correlation”, may also be used to compute the correlation

measures for the matrix. In particular, routine `CORVC` (page 314) from this chapter may be used.

2. Once the distance matrix has been computed, routine `CLINK` (page 892) is used to perform the agglomerative hierarchical cluster analysis using either single, complete, average, or Ward's linkage.
3. The results obtained from `CLINK` are examined, and if desired, the number of clusters is selected. Routine `TREEP` (page 1098) in Chapter 16, "Line Printer Graphics," may be used to print the cluster tree. This tree may aid in selecting the number of clusters, assuming that such a number is desired. Based upon the number of clusters selected, routine `CNUMB` (page 897) is used to obtain the cluster number of each of the clustered observations (or variables).
4. Routines described in Chapter 1, "Basic Statistics," and other chapters in the `IMSL STAT/LIBRARY` are used to obtain descriptive and other statistics to evaluate the clustering.

Because routine `CDIST` produces similarity and distance matrices for either rows or columns, it is easy to cluster either observations or variables. Optionally, the user may wish to cluster a correlation matrix obtained from one of the routines in the correlation chapter or to input a matrix of similarities (or dissimilarities) obtained via experimentation. The objects within such matrices may be clustered directly in routine `CLINK`.

Basic *K*-means clustering attempts to find a clustering that minimizes the within-cluster sums of squares. In this method of clustering the data, matrix  $x$  is grouped so that each observation (row in  $x$ ) is assigned to one of a fixed number,  $K$ , of clusters. The sum of the squared difference of each observation about its assigned clusters mean is used as the criterion for assignment. In the basic algorithm, observations are transferred from one cluster to another when doing so decreases the within-cluster sums of squared differences. When, in a pass through the entire data set, no transfer occurs, the algorithm stops. Routine `KMEAN` (page 900) is one implementation of the basic algorithm.

The usual course of events in *K*-means cluster analysis might be to use routine `KMEAN` to obtain the optimal clustering. The clustering is then evaluated via routines described in Chapter 1, "Basic Statistics," and/or other chapters in the `IMSL STAT/LIBRARY`. Often, *K*-means clustering with more than one value for  $K$  is performed, and the value of  $K$  that best fits the data is used.

Clustering can be performed either on observations or on variables. The discussion of the routine `KMEAN` assumes the clustering is to be performed on the observations, which correspond to the rows of the input data matrix. If variables, rather than observations, are to be clustered using `KMEAN`, the data matrix should first be transposed (possibly using routine `TRNRR` (`IMSL MATH/LIBRARY`)). In the documentation for `KMEAN`, the words "observation" and "variable" would then be exchanged.

---

## CDIST/DCDIST (Single/Double precision)

Compute a matrix of dissimilarities (or similarities) between the columns (or rows) of a matrix.

### Usage

```
CALL CDIST (NROW, NCOL, X, LDX, NDSTM, IND, IMETH, IROW,  
           ISCALE, DIST, LDDIST)
```

### Arguments

**NROW** — Number of rows in the matrix. (Input)

**NCOL** — Number of columns in the matrix. (Input)

**X** — NROW by NCOL matrix containing the data. (Input)

**LDX** — Leading dimension of X exactly as specified in the dimension statement in the calling program. (Input)

**NDSTM** — Number of rows (columns, if IROW = 1) to be used in computing the distance measure between the columns (rows). (Input)

**IND** — Vector of length NDSTM containing the indices of the rows (columns, if IROW = 1) to be used in computing the distance measure. (Input)  
If IND(1) = 0; the first NDSTM rows (columns) are used.

**IMETH** — Method to be used in computing the dissimilarities or similarities. (Input)

#### **IMETH** Method

- 0 Euclidean distance ( $L_2$  norm)
- 1 Sum of the absolute differences ( $L_1$  norm)
- 2 Maximum difference ( $L_\infty$  norm)
- 3 Mahalanobis distance
- 4 Absolute value of the cosine of the angle between the vectors
- 5 Angle in radians ( $0, \pi$ ) between the lines through the origin defined by the vectors
- 6 Correlation coefficient
- 7 Absolute value of the correlation coefficient
- 8 Number of exact matches

The algorithm section of the manual document has a more detailed description of each measure.

**IROW** — Row or columns option. (Input)

If IROW = 1, distances are computed between the NROW rows of X. Otherwise, distances between the NCOL columns of X are computed.

**ISCALE** — Scaling option. (Input)

ISCALE is not used for methods 3 through 8.



### **ISCALE Scaling Performed**

- 0 No scaling is performed.
- 1 Scale each column (row, if *IROW* = 1) by the standard deviation of the column (row).
- 2 Scale each column (row, if *IROW* = 1) by the range of the column (row).

**DIST** — *m* by *m* matrix containing the computed dissimilarities or similarities, where *m* = *NROW* if *IROW* = 1 and *m* = *NCOL* otherwise. (Output)

**LDDIST** — Leading dimension of **DIST** exactly as specified in the dimension statement in the calling program. (Input)

### **Comments**

1. Automatic workspace usage is

<b>IMETH</b>	<b>IROW</b>	<b>Workspace in CDIST</b>
0, 1, 2	0, 1	4 * NDSTM units
3	0, 1	4 * NDSTM + NDSTM * NDSTM units
4, 5	0	3 * NDSTM + NCOL units
4, 5	1	3 * NDSTM + NROW units
6, 7	0	4 * NDSTM + 2 * NCOL units
6, 7	1	4 * NDSTM + 2 * NROW units
8	0, 1	NDSTM units

The routine **DCDIST** requires twice the workspace required for **CDIST**. Workspace may be explicitly provided, if desired, by use of **C2IST/DC2IST**. The reference is

```
CALL C2IST (NROW, NCOL, X, LDX, NDSTM, IND, IMETH,  
           IROW, ISCALE, DIST, LDDIST, X1, X2,  
           SCALE, WK, IND1)
```

The additional arguments are as follows:

**X1** — Work vector of length **NDSTM**. Not used if **IMETH** = 8.

**X2** — Work vector of length **NDSTM**. Not used if **IMETH** = 8.

**SCALE** — Work vector of length **NDSTM** if **IMETH** is less than 4; of length **NCOL** or **NROW** when **IROW** is 0 or 1, respectively, and **IMETH** is 4 or 5; and of length 2 \* **NCOL** or 2 \* **NROW** when **IROW** is 0 or 1 and **IMETH** is 6 or 7. **SCALE** is not used when **IMETH** is 8.

**WK** — Work vector of length **NDSTM \* NDSTM** when **IMETH** is 3, or of length **NDSTM** when **IMETH** = 6 or 7. Not used otherwise.

**INDI** — Integer work vector of length **NDSTM**.

2. Informational error

Type	Code	
3	3	A variable is numerically linearly dependent on the previous variables when <b>IMETH</b> is 3. The variable detected as being linearly dependent is omitted from the distance measure.

## Algorithm

Routine `CDIST` computes an upper triangular matrix (excluding the diagonal) of dissimilarities (or similarities) between the columns or rows of a matrix. Nine different distance measures can be computed. For the first three measures, three different scaling options can be employed. Output from `CDIST` is generally used as input to clustering or multidimensional scaling routines.

The following discussion assumes that the distance measure is being computed between the columns of the matrix, i.e., that `IROW` is not 1. If distances between the rows of the matrix are desired, set `IROW` to 1.

For `IMETH` = 0 to 2, each row of `X` is first scaled according to the value of `ISCALE`. The scaling parameters are obtained from the values in the row scaled as either the standard deviation of the row or the row range; the standard deviation is computed from the unbiased estimate of the variance. If `ISCALE` is 0, no scaling is performed, and the parameters in the following discussion are all 1.0. Once the scaling value (if any) has been computed, the distance between column  $i$  and column  $j$  is computed via the difference vector  $z_k = (x_k - y_k)/s_k$ ,  $i = 1, \dots, \text{NDSTM}$ , where  $x_k$  denotes the  $k$ -th element in the  $i$ -th column, and  $y_k$  denotes the corresponding element in the  $j$ -th column. For given  $z_i$ , the metrics 0 to 2 are defined as:

### **IMETH Metric**

- |   |   |                    |
|---|---|--------------------|
| 0 | $\sqrt{\left(\sum_{i=1}^{\text{NDSTM}} z_i^2\right)}$ | Euclidean distance |
| 1 | $\sum_{i=1}^{\text{NDSTM}}  z_i $                     | $L_1$ norm         |
| 2 | $\max_i  z_i $  | $L_\infty$ norm    |

Distance measures corresponding to `IMETH` = 3 to 8 do not allow for scaling. These measures are defined via the column vectors  $X = (x_i)$ ,  $Y = (y_i)$ , and  $Z = (x_i - y_i)$  as follows:

### **IMETH Metric**

- |   |  |  |
|---|--|--|
| 3 | $Z' \hat{\Sigma}^{-1} Z$                             | Mahalanobis distance, where $\hat{\Sigma}$ is the usual unbiased sample estimate of the covariance matrix of the rows. |
| 4 | $\cos(\theta) = X^T Y / (\sqrt{X^T X} \sqrt{Y^T Y})$ | the dot product of $X$ and $Y$ divided by the length of $X$ times the length of $Y$ .                                  |
| 5 | $\theta$   | where $\theta$ is defined in 4.  |
| 6 | $\rho$   | the usual (centered) estimate of the correlation between $X$ and $Y$ .   |

- 7 The absolute value of  $\rho$  (where  $\rho$  is defined in 6).
- 8 The number of times  $x_i = y_i$ , where  $x_i$  and  $y_i$  are elements of  $X$  and  $Y$ .

For the Mahalanobis distance, any variable used in computing the distance measure that is (numerically) linearly dependent upon the previous variables in the IND vector is omitted from the distance measure.

### Example

The following example illustrates the use of CDIST for computing the Euclidean distance between the rows of a matrix.

```

INTEGER      IMETH, IROW, ISCALE, LDDIST, LDX, NCOL, NDSTM, NROW
PARAMETER   (IMETH=0, IROW=1, ISCALE=0, NCOL=2, NDSTM=2, NROW=4,
&           LDDIST=NROW, LDX=NROW)
C
INTEGER      IND(1)
REAL         DIST(LDDIST,NROW), X(NROW,NCOL)
EXTERNAL     CDIST, WRRRN
C
DATA IND/0/
DATA X/1, 1, 1, 1, 1, 0, -1, 2/
DATA DIST/16*0.0/
C
CALL WRRRN ('X', NROW, NCOL, X, LDX, 0)           Print input matrix
C
CALL CDIST (NROW, NCOL, X, LDX, NDSTM, IND, IMETH, IROW, ISCALE,
&          DIST, LDDIST)
C
CALL WRRRN ('DIST', NROW, NROW, DIST, LDDIST, 0) Print distance matrix
C
END

```

### Output

```

      X
      1      2
1  1.000  1.000
2  1.000  0.000
3  1.000 -1.000
4  1.000  2.000

      DIST
      1      2      3      4
1  0.000  1.000  2.000  1.000
2  0.000  0.000  1.000  2.000
3  0.000  0.000  0.000  3.000
4  0.000  0.000  0.000  0.000

```

---

## CLINK/DCLINK (Single/Double precision)

Perform a hierarchical cluster analysis given a distance matrix.

## Usage

CALL CLINK (NPT, IMETH, IDIST, DIST, LDDIST, CLEVEL,  
ICLSON, ICRSON)

## Arguments

**NPT** — Number of data points to be clustered. (Input)

**IMETH** — Option giving the method to be used for clustering. (Input)

### **IMETH** Method

- 0 Single linkage (minimum distance)
- 1 Complete linkage (maximum distance)
- 2 Average distance within (average distance between objects within the merged cluster)
- 3 Average distance between (average distance between objects in the two clusters)
- 4 Ward's method (minimize the within-cluster sums of squares). For Ward's method, the elements of **DIST** are assumed to be Euclidean distances.

**IDIST** — Option giving the type of transformation to be applied to the measures in **DIST**. (Input)

### **IDIST** Transformation

- 0 No transformation is required. The elements of **DIST** are distances.
- 1 Convert similarities to distances by multiplication by  $-1.0$ .
- 2 Convert similarities (usually correlations) to distances by taking the reciprocal of the absolute value.

**DIST** —  $NPT$  by  $NPT$  matrix containing the distance (or similarity) matrix. (Input/Output)

**DIST** is a symmetric matrix. On input, only the upper triangular part needs to be present. The routine **CLINK** saves the upper triangular part of **DIST** in the lower triangle. On return from **CLINK**, the upper triangular part of **DIST** is restored, and the matrix has been made symmetric.

**LDDIST** — Leading dimension of **DIST** exactly as specified in the dimension statement in the calling program. (Input)

**CLEVEL** — Vector of length  $NPT - 1$  containing the level at which the clusters are joined. (Output)

**CLEVEL**( $k$ ) contains the distance (or similarity) level at which cluster  $NPT + k$  was formed. If the original data in **DIST** was transformed via the option parameter **IDIST**, the inverse transformation is applied to the values in **CLEVEL** prior to exit from **CLINK**.

**ICLSON** — Vector of length  $NPT - 1$  containing the left sons of each merged cluster. (Output)

Cluster  $NPT + k$  is formed by merging clusters **ICLSON**( $k$ ) and **ICRSN**( $k$ ).

**ICRSON** — Vector of length  $NPT - 1$  containing the right sons of each merged cluster. (Output)

Cluster  $NPT + k$  is formed by merging clusters  $ICLSON(k)$  and  $ICRSON(k)$ .

### Comments

1. Automatic workspace usage is

**CLINK**  $2 * NPT$  units if  $IMETH = 0$  or  $1$ , and  $4 * NPT$  units otherwise,  
or

**DCLINK**  $2 * NPT$  units if  $IMETH = 0$  or  $1$ , and  $6 * NPT$  units otherwise.

Workspace may be explicitly provided, if desired, by use of **C2INK/DC2INK**. The reference is

```
CALL C2INK (NPT, IMETH, IDIST, DIST, LDDIST, CLEVEL,  
           ICLSON, ICRSON, IPTR, ICLUS, CWT, CSUM)
```

The additional arguments are as follows:

**IPTR** — Integer work vector of length  $NPT$ .

**ICLUS** — Integer work vector of length  $NPT$ .

**CWT** — Work vector of length  $NPT$ . Not used if  $IMETH = 0$  or  $1$ .

**CSUM** — Work vector of length  $NPT$ . Not used if  $IMETH = 0$  or  $1$ .

2. The clusters corresponding to the original data points are numbered from  $1$  to  $NPT$ . The  $NPT - 1$  clusters formed by merging clusters are numbered  $NPT + 1$  to  $NPT + (NPT - 1)$ .
3. Raw correlations, if used as similarities, should be made positive and transformed to a distance measure. One such transformation can be performed by specifying  $IDIST = 2$  in **CLINK**.
4. The user may cluster either variables or observations in **CLINK** since a dissimilarity matrix, not the original data, is used. Routine **CDIST** (page 889) may be used to compute the matrix **DIST**.
5. Routine **TREEP** (page 1098) in the graphics chapter can be used to obtain a line printer plot of the clustering tree. Routine **CNUMB** (page 897) can be used to obtain the cluster number assigned to each of the original clusters when a specified number of clusters is desired.

### Algorithm

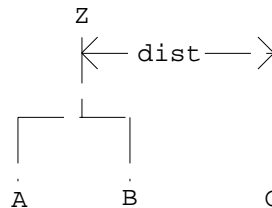
Routine **CLINK** performs hierarchical cluster analysis based upon a distance matrix, or by appropriate use of the **IDIST** option, based upon a similarity matrix. Only the upper triangular part of the matrix needs to be input to **CLINK**.

Hierarchical clustering in **CLINK** proceeds as follows. Initially, each data point is considered to be a cluster, numbered  $1$  to  $n = NPT$ .

1. If the data matrix contains similarities, they are converted to distances by the method specified in **IDIST**. Set  $k = 1$ .

2. A search is made of the distance matrix to find the two closest clusters. These clusters are merged to form a new cluster, numbered  $n + k$ . The cluster numbers of the two clusters joined at this stage are saved in `ICRSON` and `ICLSON`, and the distance measure between the two clusters is stored in `CLEVEL`.
3. Based upon the method of clustering, updating of the distance measure in the row and column of `DIST` corresponding to the new cluster is performed.
4. Set  $k = k + 1$ . If  $k < n$ , go to Step 2.

The five methods differ primarily in how the distance matrix is updated after two clusters have been joined. The `IMETH` option parameter specifies how the distance of the cluster just merged with each of the remaining clusters will be updated. Routine `CLINK` allows five methods of computing the distances. To understand these measures, suppose in the following discussion that clusters “A” and “B” have just been joined to form cluster “Z”, and interest is in computing the distance of Z with another cluster called “C”.



**IMETH Method**

- 0 This is the single linkage method. The distance from Z to C is the minimum of the distances (A to C, B to C).
- 1 This is the complete linkage method. The distance from Z to C is the maximum of the distances (A to C, B to C).
- 2 This is the average-distance-within-clusters method. The distance from Z to C is the average distance of all objects that would be within the cluster formed by merging clusters Z and C. This average may be computed according to formulas given by Anderberg (1973, page 139).
- 3 This is the average-distance-between-clusters method. The distance from Z to C is the average distance of objects within cluster Z to objects within cluster C. This average may be computed according to methods given by Anderberg (1973, page 140).
- 4 This is Ward’s method. Clusters are formed so as to minimize the increase in the within-cluster sums of squares. The distance between two clusters is the increase in these sums of squares if the two clusters were merged. A method for computing this distance from a squared Euclidean distance matrix is given by Anderberg (1973, pages 142–145).

In general, single linkage will yield long thin clusters while complete linkage will yield clusters that are more spherical. Average linkage and Ward's linkage tend to yield clusters that are similar to those obtained with complete linkage.

Routine CLINK produces a unique representation of the binary cluster tree via the following three conventions; the fact that the tree is unique should aid in interpreting the clusters. First, when two clusters are joined and each cluster contains two or more data points, the cluster that was initially formed with the smallest level (in CLEVEL) becomes the left son. Second, when a cluster containing more than one data point is joined with a cluster containing a single data point, the cluster with the single data point becomes the right son. Finally, when two clusters containing only one object are joined, the cluster with the smallest cluster number becomes the right son.

### Example

In the following example, the average distance within clusters method is used to perform a hierarchical cluster analysis of the Fisher iris data. Routine GDATA (page 1302) is first used to obtain the Fisher iris data. The example is typical in that after the program obtains the data, routine CDIST (page 889) computes the distance matrix (DIST) prior to calling CLINK.

```

      INTEGER      IDATA, IDIST, IMETH, IMTH, IPRINT, IROW, ISCALE,
&                LDDIST, LDX, NCOL, NPT, NROW, NVAR
      PARAMETER   (IDATA=3, IDIST=0, IMETH=2, IMTH=0, IPRINT=0, IROW=1,
&                ISCALE=1, NCOL=5, NROW=150, NVAR=4, LDX=NROW,
&                NPT=NROW, LDDIST=LDX)
C
      INTEGER      I, ICLSON(NROW-1), ICRSON(NROW-1), IND(4), NOUT,
&                NXCOL, NXROW
      REAL         CLEVEL(NROW-1), DIST(LDDIST,LDDIST), X(LDX,NCOL)
      EXTERNAL     CDIST, CLINK, GDATA, UMACH
C
      DATA IND/2, 3, 4, 5/
C
      CALL GDATA (IDATA, IPRINT, NXROW, NXCOL, X, LDX, NCOL)
C
      CALL CDIST (NROW, NCOL, X, LDX, NVAR, IND, IMTH, IROW, ISCALE,
&                DIST, LDDIST)
C
      CALL CLINK (NPT, IMETH, IDIST, DIST, LDDIST, CLEVEL, ICLSON,
&                ICRSON)
C
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99996) (I,I=1,149,15)
      WRITE (NOUT,99997) (CLEVEL(I),I=1,149,15)
      WRITE (NOUT,99998) (ICLSON(I),I=1,149,15)
      WRITE (NOUT,99999) (ICRSON(I),I=1,149,15)
C
99996 FORMAT (' OBS # ', 10I6)
99997 FORMAT (' CLEVEL', 10F6.2)
99998 FORMAT (' ICLSON', 10I6)
99999 FORMAT (' ICRSON', 10I6)
C
      END

```

		<b>Output</b>								
OBS #	1	16	31	46	61	76	91	106	121	136
CLEVEL	0.00	0.17	0.23	0.27	0.31	0.37	0.41	0.48	0.60	0.78
ICLSON	143	153	17	140	53	198	186	218	261	249
ICRSON	102	29	6	113	51	91	212	243	266	262

## CNUMB

Compute cluster membership for a hierarchical cluster tree.

### Usage

```
CALL CNUMB (NODE, ICLSON, ICRSON, K, ICLUS, NCLUS)
```

### Arguments

**NODE** — Number of data points clustered. (Input)

**ICLSON** — Vector of length  $NODE - 1$  containing the left son cluster numbers. (Input)

Cluster  $NODE + I$  is formed by merging clusters  $ICLSON(I)$  and  $ICRSON(I)$ .

**ICRSON** — Vector of length  $NODE - 1$  containing the right son cluster numbers. (Input)

Cluster  $NODE + I$  is formed by merging clusters  $ICLSON(I)$  and  $ICRSON(I)$ .

**K** — Desired number of clusters. (Input)

**ICLUS** — Vector of length  $NODE$  containing the cluster membership of each observation. (Output)

Observation  $I$  is in cluster  $ICLUS(I)$  when  $K$  clusters are specified.

**NCLUS** — Vector of length  $K$  containing the number of observations in each cluster. (Output)

### Comments

1. Automatic workspace usage is

$CNUMB \ 2 * NODE$  units.

Workspace may be explicitly provided, if desired, by use of **C2UMB**. The reference is

```
CALL C2UMB (NODE, ICLSON, ICRSON, K, ICLUS, NCLUS,
            IPT)
```

The additional argument is

**IPT** — Work vector of length  $2 * NODE$ .

2. Informational errors

Type    Code



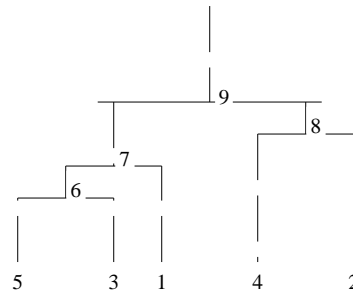
- 4      1      The tree structure specified by ICLSON and ICRSON is invalid because an attempt to assign an observation to more than one cluster is being made.
- 4      2      The tree structure specified by ICLSON and ICRSON is incorrect because an observation is not assigned to a cluster.

### Algorithm

Given a fixed number of clusters ( $K$ ) and the cluster tree (vectors ICRSON and ICLSON) produced by the hierarchical clustering algorithm (see routine CLINK, page 892), routine CNUMB determines the cluster membership of each observation. The routine CNUMB first determines the root nodes for the  $K$  distinct subtrees forming the  $K$  clusters and then traverses each subtree to determine the cluster membership of each observation. The routine CNUMB also returns the number of observations found in each cluster.

### Example 1

In the following example, cluster membership for  $K = 2$  clusters is found for the displayed cluster tree. The output vector ICLUS contains the cluster numbers for each observation.



```

INTEGER      K, NODE
PARAMETER    (K=2, NODE=5)
C
INTEGER      ICLSON(NODE-1), ICLUS(NODE), ICRSON(NODE-1), NCLUS(K)
EXTERNAL     CNUMB, WRIRN
C
DATA ICLSON/5, 6, 4, 7/
DATA ICRSON/3, 1, 2, 8/
C
                                Compute cluster membership
CALL CNUMB (NODE, ICLSON, ICRSON, K, ICLUS, NCLUS)
C
                                Print output
CALL WRIRN ('ICLUS', 1, NODE, ICLUS, 1, 0)
CALL WRIRN ('NCLUS', 1, K, NCLUS, 1, 0)
C
END
  
```

### Output

```

      ICLUS
1  2  3  4  5
1  2  1  2  1
  
```

```

1      NCLUS
2
3      2

```

## Example 2

This example illustrates the typical usage of CNUMB. The Fisher iris data (see routine GDATA, page 1302) is clustered. First the distance between the irises are computed using routine CDIST (page 889). The resulting distance matrix is then clustered using routine CLINK (page 892). The cluster membership for 5 clusters is then obtained via routine CNUMB using the output from CLINK. The need for 5 clusters can be obtained either by theoretical means or by examining a cluster tree. Because the cluster tree is too large to be included in this example, the call to routine TREEP (page 1098) that would ordinarily print the cluster tree has been commented in the example code. The cluster membership for each of the iris observations is printed.

```

      INTEGER      IDATA, IDIST, IMETHD, IMTH, IPRINT, IROW, ISCALE, K,
&                LDDIST, LDX, NCOL, NODE, NODEX, NROW, NVAR
      PARAMETER    (IDATA=3, IDIST=0, IMETHD=0, IMTH=1, IPRINT=0,
&                IROW=1, ISCALE=0, K=5, LDDIST=150, LDX=150, NCOL=5,
&                NODE=150, NODEX=5, NROW=150, NVAR=4)
C
      INTEGER      I, ICLSON(NROW-1), ICLUS(NODE), ICRSON(NROW-1),
&                IMETH, IND(4), IROOT, J, NCLUS(K), NFILL, NSCALE,
&                NXCOL, NXROW
      REAL         AMAX1, CLEVEL(NROW-1), DIST(LDDIST,LDDIST), RNUNF,
&                SCALE(2), X(LDX,NCOL)
      CHARACTER    NODENM(NODE)*7
      INTRINSIC    AMAX1
      EXTERNAL     CDIST, CLINK, CNUMB, GDATA, RNSET, RNUNF, SCOPY, WRIRN
C
      DATA IND/2, 3, 4, 5/
      DATA IMETH, IROOT, NSCALE, NFILL/1, 0, 1, 1/
      DATA SCALE/0.0, 3.5/
C
      CALL GDATA (IDATA, IPRINT, NXROW, NXCOL, X, LDX, NCOL)
C
      CALL CDIST (NROW, NCOL, X, LDX, NVAR, IND, IMETHD, IROW, ISCALE,
&                DIST, LDDIST)
C
      CALL RNSET (4)
C
      DO 20 I=1, NODE
      DO 10 J=I + 1, NODE
      DIST(I,J) = AMAX1(0.0, DIST(I,J)+(0.001*RNUNF()))
10    CONTINUE
      DIST(I,I) = 0.0
      CALL SCOPY (I-1, DIST(1,I), 1, DIST(I,1), LDDIST)
20  CONTINUE
C
      CALL CLINK (NODE, IMETHD, IDIST, DIST, LDDIST, CLEVEL, ICLSON,
&                ICRSON)
C
      NODENM(1) = 'DEFAULT'

```

```

C     CALL TREEP (NODE, ICLSON, ICRSON,
C     IMTH, CLEVEL, IROOT, NSCALE,
C     &           NFFILL, SCALE, NODENM)
C                                     Compute membership for 5 clusters
C     CALL CNUMB (NODE, ICLSON, ICRSON, K, ICLUS, NCLUS)
C                                     Print output
C     CALL WRIRN ('ICLUS', 1, NODE, ICLUS, 1, 0)
C     CALL WRIRN ('NCLUS', 1, K, NCLUS, 1, 0)
C
C     END

```

### Output

```

                                     ICLUS
1  2  3  4  5  6  7  8  9  10 11 12 13 14 15 16 17 18 19 20
5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5

21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40
  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5  5

41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60
  5  5  5  5  5  5  5  5  5  5  2  2  2  2  2  2  2  1  2  2

61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80
  1  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2

81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99
  2  2  2  2  2  2  2  2  2  2  2  2  2  1  2  2  2  2  1

100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115
  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2

116 117 118 119 120 121 122 123 124 125 126 127 128 129 130 131
  2  2  4  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2

132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147
  4  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2

148 149 150
  2  2  2

                                     NCLUS
1  2  3  4  5
4  93 1  2  50

```

---

## KMEAN/DKMEAN (Single/Double precision)

Perform a *K*-means (centroid) cluster analysis.

### Usage

```

CALL KMEAN (NOBS, NCOL, NVAR, X, LDX, IFRQ, IWT, IND, K,
            MAXIT, CM, LDCM, SWT, LDSWT, IC, NC, WSS)

```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in  $X$ . (Input)

**NVAR** — Number of variables to be used in computing the metric. (Input)

**X** — NOBS by NCOL matrix containing the observations to be clustered. (Input)  
The only columns of  $X$  used are those indicated by **IND** and possibly **IFRQ** and/or **IWT**.

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

**IFRQ** — Frequency option. (Input)

**IFRQ** = 0 means all frequencies are 1. For positive **IFRQ**, column number **IFRQ** of  $X$  contains the nonnegative frequencies.

**IWT** — Weighting option. (Input)

**IWT** = 0 means all weights are 1. For positive **IWT**, column number **IWT** contains the nonnegative weights.

**IND** — Vector of length **NVAR** containing the columns of  $X$  to be used in computing the metric. (Input)

In the usual case in which  $X$  is the data matrix, no observation has multiple frequency, and unequal weighting is not desired, **IND** = (1, 2, 3, ..., **NVAR**).

**K** — Number of clusters. (Input)

**MAXIT** — Maximum number of iterations. (Input)

**MAXIT** = 30 is usually sufficient.

**CM** —  $K$  by **NVAR** matrix containing, on input, the cluster seeds, i.e., estimates for the cluster centers, and the cluster means on output. (Input/Output)  
The cluster seeds must be unique.

**LDCM** — Leading dimension of **CM** exactly as specified in the dimension statement in the calling program. (Input)

**SWT** —  $K$  by **NVAR** matrix containing the sum of the weights used to compute each cluster mean. (Output)

Missing observations are excluded from **SWT**.

**LDSWT** — Leading dimension of **SWT** exactly as specified in the dimension statement in the calling program. (Input)

**IC** — Vector of length NOBS containing the cluster membership for each observation. (Output)

**NC** — Vector of length  $K$  containing the number of observations in each cluster. (Output)

**WSS** — Vector of length  $K$  containing the within sum of squares for each cluster. (Output)

### Comments

1. Automatic workspace usage is

KMEAN 2 \* NOBS + 3 \* K units, or  
 DKMEAN 3 \* NOBS + 3 \* K units.

Workspace may be explicitly provided, if desired, by use of  
 K2EAN/DK2EAN. The reference is

```
CALL K2EAN (NOBS, NCOL, NVAR, X, LDX, IFRQ, IWT,
           IND, K, MAXIT, CM, LDCM, SWT, LDSWT, IC,
           NC, WSS, IC2, NCP, D, ITRAN, LIVE)
```

The additional arguments are as follows:

**IC2** — Work vector of length NOBS.

**NCP** — Work vector of length K.

**D** — Work vector of length NOBS.

**ITRAN** — Work vector of length K.

**LIVE** — Work vector of length K.

2. Informational Error

Type	Code	
3	1	Convergence did not occur within MAXIT iterations.

**Algorithm**

Routine KMEAN is an implementation of Algorithm AS 136 by Hartigan and Wong (1979). It computes *K*-means (centroid) Euclidean metric clusters for an input matrix starting with initial estimates of the *K* cluster means. Routine KMEAN allows for missing values (coded as NaN, “not a number”) and for weights and frequencies.

Let  $p = \text{NVAR}$  denote the number of variables to be used in computing the Euclidean distance between observations. The idea in *K*-means cluster analysis is to find a clustering (or grouping) of the observations so as to minimize the total within-cluster sums of squares. In this case, the total sums of squares within each cluster is computed as the sum of the centered sum of squares over all nonmissing values of each variable. That is,

$$\phi = \sum_{i=1}^K \sum_{j=1}^p \sum_{m=1}^{n_i} f_{v_{im}} w_{v_{im}} \delta_{v_{im},j} (x_{v_{im},j} - \bar{x}_{ij})^2$$

where  $v_{im}$  denotes the row index of the  $m$ -th observation in the  $i$ -th cluster in the matrix  $\mathbf{x}$ ;  $n_i$  is the number of rows of  $\mathbf{x}$  assigned to group  $i$ ;  $f$  denotes the frequency of the observation;  $w$  denotes its weight;  $\delta$  is zero if the  $j$ -th variable on observation  $v_{im}$  is missing, otherwise  $\delta$  is one; and

$$\bar{x}_{ij}$$

is the average of the nonmissing observations for variable  $j$  in group  $i$ . This method sequentially processes each observation and reassigns it to another cluster if doing so results in a decrease in the total within-cluster sums of

squares. The user is referred to Hartigan and Wong (1979) or Hartigan (1975) for the details.

### Example

This example performs *K*-means cluster analysis on Fisher's iris data, which is first obtained via routine GDATA (page 1302). The initial cluster seed for each iris type is an observation known to be in the iris type.

```

INTEGER      IFRQ, IPRINT, IWT, K, LDCM, LDSWT, LDX, MAXIT, NCOL,
&            NOBS, NV, NVAR
PARAMETER    (IFRQ=0, IPRINT=0, IWT=0, K=3, MAXIT=30, NCOL=5,
&            NOBS=150, NV=5, NVAR=4, LDCM=K, LDSWT=K, LDX=NOBS)
C
INTEGER      IC(NOBS), IND(NVAR), NC(K), NXCOL, NXROW
REAL         CM(K,NVAR), SWT(K,NVAR), WSS(K), X(NOBS,NV)
EXTERNAL     GDATA, KMEAN, SCOPY, WRIRN, WRRRN
C
DATA IND/2, 3, 4, 5/
C
CALL GDATA (3, IPRINT, NXROW, NXCOL, X, NOBS, NV)
C
                                Copy the cluster seeds into CM
CALL SCOPY (NVAR, X(1,2), LDX, CM(1,1), LDCM)
CALL SCOPY (NVAR, X(51,2), LDX, CM(2,1), LDCM)
CALL SCOPY (NVAR, X(101,2), LDX, CM(3,1), LDCM)
C
CALL KMEAN (NOBS, NCOL, NVAR, X, LDX, IFRQ, IWT, IND, K, MAXIT,
&          CM, LDCM, SWT, LDSWT, IC, NC, WSS)
C
CALL WRRRN ('CM', K, NVAR, CM, LDCM, 0)
CALL WRRRN ('SWT', K, NVAR, SWT, LDSWT, 0)
CALL WRIRN ('IC', 1, NOBS, IC, 1, 0)
CALL WRIRN ('NC', 1, K, NC, 1, 0)
CALL WRRRN ('WSS', 1, K, WSS, 1, 0)
END

```

### Output

	CM			
	1	2	3	4
1	5.006	3.428	1.462	0.246
2	5.902	2.748	4.394	1.434
3	6.850	3.074	5.742	2.071

	SWT			
	1	2	3	4
1	50.00	50.00	50.00	50.00
2	62.00	62.00	62.00	62.00
3	38.00	38.00	38.00	38.00

	IC																			
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
21	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
41	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1

```
 1  1  1  1  1  1  1  1  1  1  1  2  2  3  2  2  2  2  2  2  2
61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80
 2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  3  2  2
81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99
 2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2  2
100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115
  2  3  2  3  3  3  3  2  3  3  3  3  3  3  2  2
116 117 118 119 120 121 122 123 124 125 126 127 128 129 130 131
  3  3  3  3  2  3  2  3  2  3  3  2  2  3  3  3
132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147
  3  3  2  3  3  3  3  2  3  3  3  2  3  3  3  2
148 149 150
  3  3  2

      NC
  1    2    3
50  62  38

      WSS
  1    2    3
15.15 39.82 23.88
```

# Chapter 12: Sampling

---

## Routines

Proportions, simple random sample .....	SMPPR	906
Proportions, stratified random sample .....	SMPPS	909
Ratio or regression estimates, simple random sample .....	SMPRR	911
Ratio or regression estimates, stratified random sample....	SMPRS	918
Single-stage cluster sample .....	SMPSC	923
Simple random sample .....	SMPSR	927
Stratified random sample .....	SMPSS	930
Two-stage sample with equisized primary units.....	SMPST	933

---

## Usage Notes

The routines for inferences regarding proportions require only counts as the input data. The other routines described in this chapter require the actual data. Since the amount of data may be quite large, these routines allow for the data to be input in small quantities (or even to be deleted after it has already been passed to the subroutine). This is accomplished by means of the processing option parameter, `IDO`, and an indicator of the number of observations being passed in, `NROW`. `IDO` has the following meaning:

<code>IDO</code>	Action
0	This is the only invocation of the subroutine for this data set, and all the data are input at once.
1	This is the first invocation, and additional calls to the subroutine will be made. Initialization and updating for the data are performed.
2	This is an intermediate invocation of the subroutine, and updating for the data is performed.
3	This is the final invocation of the routine. Updating for the data and any wrap-up computations are performed.

`NROW` can be positive or negative or zero. Its absolute value is the number of sample values being input. If `NROW` is negative, it is assumed that the observations being input have already been input once and now it is desired to delete them from the analysis. When `IDO` is 3, `NROW` can be set to 0. In this case, only postprocessing is performed; no accumulation of statistics is done. This



allows input of summary statistics rather than the actual data. See Example 2 in the documentation for the routine *SMPSR* (page 927).

There are other variables used by several routines in this chapter that have a common meaning in all routines:

*Y* — The variable of interest.

*X* — The auxiliary variable.

*NSAMP* — The sample size.

*NPOP* — The population size.

*CONPER* — Confidence level.

*STAT* — Output statistics.

For stratified sampling, the following variables are often used:

*NSTRAT* — Number of strata.

*NROWS* — Vector with elements like *NROW* for strata.

*NSAMPS* — The strata sample sizes.

*NPOPS* — The population sizes for strata.

*YBARS* — The strata sample means.

*YVARS* — The strata sample variances.

---

## SMPPR/DSMPPR (Single/Double precision)

Compute statistics for inferences regarding the population proportion and total given proportion data from a simple random sample.

### Usage

```
CALL SMPPR (NINT, NSAMP, NPOP, CONPER, STAT)
```

### Arguments

*NINT* — Number of sample units in the class of interest, for the population (or subpopulation) of interest. (Input)

*NSAMP* — Number of units in the entire random sample. (Input)

*NPOP* — Number of units in the population. (Input)

*CONPER* — Confidence level for two-sided interval estimates, in percent. (Input)

A *CONPER* percent confidence interval is computed; hence, *CONPER* must be greater than or equal to 0.0 and less than 100.0. *CONPER* is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level *ONECL*, set  $CONPER = 100.0 - 2.0 * (100.0 - ONECL)$ .

**STAT** — Vector of length 10 containing the resulting statistics. (Output)  
These are

- | <b>I</b> | <b>STAT(I)</b>  |
|----------|---|
| 1        | Estimate of the proportion.   |
| 2        | Estimate of the total.  |
| 3        | Variance estimate of the proportion estimate.   |
| 4        | Variance estimate of the total estimate.  |
| 5        | Lower confidence limit for the proportion.  |
| 6        | Upper confidence limit for the proportion.  |
| 7        | Lower confidence limit for the total.   |
| 8        | Upper confidence limit for the total.   |
| 9        | Estimate (expressed as a percentage) of the coefficient of variation of the total estimate. Not defined if $NINT = 0$ .   |
| 10       | Indicator of the distribution used to approximate the hypergeometric distribution for the confidence interval calculations. If $STAT(10) = 0$ , then the normal is used. If $STAT(10) = 1$ , then the Poisson is used. If $STAT(10) = 2$ , then the binomial is used. |

### Algorithm

The routine `SMPPR` computes point and interval estimates for the population proportion and total from a simple random sample. The simplest and most common case for which this routine is appropriate is one in which the population sampled contains two or more classes, and it is desired to estimate the proportion of the population falling into a particular class (“class of interest”). The data required by `SMPPR` consist of counts of the number of sample items in the class of interest, the sample size, and the population size. If there are more than two classes in the population, some of the classes may not be of interest.

Since the hypergeometric distribution is the appropriate probability model for the sampling for proportions in a finite population without replacement, exact confidence limits could be computed using that distribution. For populations with sizes that occur in practice (more than a hundred, often in the thousands or even millions), the confidence limits can be approximated very well by use of the normal, the binomial, or the Poisson distribution. Routine `SMPPR` uses one of these distributions in setting confidence limits, following the guidelines in the table on page 58 of Cochran (1977).

### Example 1

The first example is from Cochran (1977, page 52). A simple random sample of size 200 was drawn from a list of 3042 names and addresses. Verification of the addresses in the sample showed 38 to be wrong. The objective is to estimate the total number of incorrect addresses.

INTEGER	NINT, NOUT, NPOP, NSAMP
REAL	CONPER, SQRT, STAT(10), STDP, STDT
INTRINSIC	SQRT

```

EXTERNAL  SMPPR, UMACH
C
CALL UMACH (2, NOUT)
NINT     = 38
NSAMP    = 200
NPOP     = 3042
CONPER   = 0.0
CALL SMPPR (NINT, NSAMP, NPOP, CONPER, STAT)
STDP     = SQRT(STAT(3))
STDT     = SQRT(STAT(4))
WRITE (NOUT,99999) STAT(1), STAT(2), STDP, STDT, STAT(9)
99999 FORMAT (' Estimate of proportion bad:           ', F5.3, '/',
&           ' Estimate of total bad:                 ', F5.0, '/',
&           ' Standard deviation estimate, proportion: ', F5.3, '/',
&           ' Standard deviation estimate, total:      ', F5.1, '/',
&           ' Coefficient of variation:                ', F5.1,
&           '%')
END

```

### Output

```

Estimate of proportion bad:           0.190
Estimate of total bad:                 578.
Standard deviation estimate, proportion: 0.027
Standard deviation estimate, total:      81.8
Coefficient of variation:                14.1%

```

### Example 2

The next example is also from Cochran (1977, page 68). A simple random sample of size 200 from 2000 colleges showed 120 colleges to be in favor of a certain proposal, 57 to be opposed, and 23 to have no opinion. We wish to estimate the number of colleges, out of the 2000, that favor the proposal.

```

INTEGER  NINT, NOUT, NPOP, NSAMP
REAL     CONPER, STAT(10)
EXTERNAL SMPPR, UMACH
C
CALL UMACH (2, NOUT)
NINT     = 120
NSAMP    = 200
NPOP     = 2000
CONPER   = 95.0
CALL SMPPR (NINT, NSAMP, NPOP, CONPER, STAT)
WRITE (NOUT,99999) STAT(2), STAT(7), STAT(8)
99999 FORMAT (' Estimate of number in favor:           ', F5.0, '/', ' 95% ',
&           'confidence interval: (', F5.0, ', ', F5.0, ')')
END

```

### Output

```

Estimate of number in favor:    1200.
95% confidence interval: (1066.,1334.)

```

---

## SMPPS/DSMPPS (Single/Double precision)

Compute statistics for inferences regarding the population proportion and total given proportion data from a stratified random sample.

### Usage

```
CALL SMPPS (NSTRAT, NINTS, NSAMPS, NPOPS, CONPER, PROPOR,  
           STAT)
```

### Arguments

**NSTRAT** — Number of strata into which the sample is divided. (Input)  
In the vectors of length **NSTRAT**, the elements are all ordered in the same way.

**NINTS** — Vector of length **NSTRAT** containing the observed number of units in each stratum from the class of interest. (Input)

**NSAMPS** — Vector of length **NSTRAT** containing the sample size in each stratum. (Input)

**NPOPS** — Vector of length **NSTRAT** containing the population in the strata. (Input)

If the population strata sizes are not known, estimates must be entered in their place.

**CONPER** — Confidence level for two-sided interval estimate, in percent. (Input)

A **CONPER** percent confidence interval is computed; hence, **CONPER** must be greater than or equal to 0.0 and less than 100.0. **CONPER** is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level **ONECL**, set  $CONPER = 100.0 - 2.0 * (100.0 - ONECL)$ .

**PROPOR** — Vector of length **NSTRAT** containing the within-strata proportion estimates. (Output)

**STAT** — Vector of length 10 containing the resulting statistics. (Output)  
These are

<b>I</b>	<b>STAT(I)</b>
1	Estimate of the proportion.
2	Estimate of the total.
3	Variance estimate of the proportion estimate.
4	Variance estimate of the total estimate.
5	Lower confidence limit for the proportion.
6	Upper confidence limit for the proportion.
7	Lower confidence limit for the total.
8	Upper confidence limit for the total.
9	Estimate (expressed as a percentage) of the coefficient of variation of the total estimate.

10 Variance estimate of the proportion estimate assuming that sampling was simple random instead of stratified random.

### Algorithm

Routine `SMPPS` computes point and interval estimates for the population proportion and total from a stratified random sample. If the strata are formed so that the proportions differ greatly from one stratum to the next, considerable gain in statistical efficiency can be realized by use of stratified sampling (see Cochran 1977, page 107).

Let  $N_h$  be the number in the population in the  $h$ -th stratum, let  $n_h$  be the number in the sample from the  $h$ -th stratum, let  $a_h$  be the number of the class of interest in the sample from the  $h$ -th stratum, let  $N$  be the population size ( $\sum N_h$ ), let  $p_h$  be the proportion in the  $h$ -th stratum,  $a_h/n_h$ , and let  $L$  be the number of strata. Then, the estimate of the proportion is

$$p_{st} = \sum_{h=1}^L \frac{N_h a_h}{N n_h}$$

and the estimate of the variance is

$$v(p_{st}) = \frac{1}{N^2} \sum_{h=1}^L N_h (N_h - n_h) \frac{p_h (1 - p_h)}{n_h - 1}$$

The confidence intervals are computed using a normal approximation.

### Example

This example is an artificial modification of an example used in routine `SMPPR` (page 906), which is from Cochran (1977, page 52). A list of 3042 names and addresses was built by an experienced secretary and a part-time student worker. The secretary entered 1838 names and addresses, and the student entered the remainder. Samples of size 100 were taken from the names entered by each. Verification of the addresses in the sample from the secretary's work showed 12 to be wrong, and verification of the student's sample showed 26 to be wrong. The objective is to estimate the total number of incorrect addresses.

```

C      INTEGER      NSTRAT
      PARAMETER    (NSTRAT=2)

C      INTEGER      NINTS(NSTRAT), NOUT, NPOPS(NSTRAT), NSAMPS(NSTRAT)
      REAL          CONPER, PROPOR(NSTRAT), SQRT, STAT(10), STDP, STDSRS,
&      STDT
      INTRINSIC     SQRT
      EXTERNAL      SMPPS, UMACH

C      CALL UMACH (2, NOUT)
      NINTS(1) = 12
      NINTS(2) = 26
      NSAMPS(1) = 100
      NSAMPS(2) = 100

```

```

NPOPS(1) = 1838
NPOPS(2) = 1204
CONPER   = 0.0
C
CALL SMPPS (NSTRAT, NINTS, NSAMPS, NPOPS, CONPER, PROPOR, STAT)
C
STDP     = SQRT(STAT(3))
STDT     = SQRT(STAT(4))
STDSRS   = SQRT(STAT(10))
C
WRITE (NOUT,99999) STAT(1), STAT(2), STDP, STDT, STAT(9), STDSRS
99999 FORMAT (' Estimate of proportion bad:           ', F7.3, '/',
&           ' Estimate of total bad:                ', F4.0, '/',
&           ' Standard deviation estimate, proportion: ', F7.3, '/',
&           ' Standard deviation estimate, total:      ', F5.1, '/',
&           ' Coefficient of variation:              ', F5.1, '/',
&           '%', '/', ' Std. dev. under simple random sampling: ',
&           F7.3)
END

```

### Output

```

Estimate of proportion bad:           0.175
Estimate of total bad:                534.
Standard deviation estimate, proportion: 0.025
Standard deviation estimate, total:    77.4
Coefficient of variation:              14.5%
Std. dev. under simple random sampling: 0.027

```

---

## SMPRR/DSMPRR (Single/Double precision)

Compute statistics for inferences regarding the population mean and total using ratio or regression estimation, or inferences regarding the population ratio given a simple random sample.

### Usage

```
CALL SMPRR (IDO, NROW, X, Y, NPOP, IOPT, XMEAN, COEF,
           CONPER, STAT)
```

### Arguments

**IDO** — Processing option. (Input)

#### IDO Action

- 0 This is the only invocation of SMPRR for this data set, and all the data are input at once.
- 1 This is the first invocation, and additional calls to SMPRR will be made. Initialization and updating for the data in X and Y are performed.
- 2 This is an intermediate invocation of SMPRR and updating for the data in X and Y is performed.
- 3 This is the final invocation of this routine. Updating for the data in X and Y, and wrap-up computations are performed.

**NROW** — The absolute value of **NROW** is the number of observations currently input in **X** and **Y**. (Input)

**NROW** may be positive, zero, or negative. Negative  $-\text{NROW}$  means delete the **NROW** rows of data from the analysis.

**X** — Vector of length  $|\text{NROW}|$  containing the data for the auxiliary variable in the random sample. (Input)

**Y** — Vector of length  $|\text{NROW}|$  containing the data for the variable of interest in the random sample. (Input)

The value of  $Y(I)$  corresponds to that of  $X(I)$ .

**NPOP** — Size of the population (number of pairs of elements in the sampled population). (Input)

**IOPT** — Estimation option. (Input)

**IOPT**    **Action**

- 0        Ratio estimation is used for inference about the population mean, total, and ratio.
- 1        The population mean of the auxiliary variable is not used, and only inference about the population ratio is desired.
- 2        Regression estimation with preassigned regression coefficient (in **COEF**) is used for inference about the population mean and total.
- 3        Regression estimation with estimated regression coefficient (returned in **STAT(18)**) is used for inference about the population mean and total.

**XMEAN** — Population mean of the auxiliary variable. (Input)

**XMEAN** is not used if **IOPT** = 1.

**COEF** — Reassigned regression coefficient. (Input)

**COEF** is used only when **IOPT** = 2.

**CONPER** — Confidence level for two-sided interval estimate, in percent.

(Input)

A **CONPER** percent confidence interval is computed, hence, **CONPER** must be greater than or equal to 0.0 and less than 100.0. **CONPER** is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level **ONECL**, set  $\text{CONPER} = 100.0 - 2.0 * (100.0 - \text{ONECL})$ .

**STAT** — Vector of length 20 containing the resulting statistics. (Output, if

**IDO** = 0 or 1; input/output, if **IDO** = 2 or 3)

**I**        **STAT(I)**

- 1        Estimate of the mean.
- 2        Estimate of the total.
- 3        Variance estimate of the mean estimate.
- 4        Variance estimate of the total estimate.
- 5        Lower confidence limit for the mean.
- 6        Upper confidence limit for the mean.
- 7        Lower confidence limit for the total.

- 8 Upper confidence limit for the total.
- 9 Estimate of the ratio.
- 10 Variance estimate of the estimate of the ratio. The population mean of the auxiliary variable is used in STAT(10) if the mean is known; otherwise, the sample estimate of the population mean is used.
- 11 Lower confidence limit for the ratio.
- 12 Upper confidence limit for the ratio.
- 13 Estimate (expressed as a percentage) of the coefficient of variation of the mean, total, and ratio and regression coefficient estimates that are defined, as controlled by IOPT. The standard deviation in the numerator of this quantity has been divided by the square root of the sample size. The coefficients of variation in STAT(14) and STAT(15) use the sample standard deviations without that divisor.
- 14 Estimate (expressed as a percentage) of the coefficient of variation of the auxiliary variable.
- 15 Estimate (expressed as a percentage) of the coefficient of variation of the variable of interest.
- 16 Sample mean of the auxiliary variable.
- 17 Sample mean of the variable of interest.
- 18 Estimate of the regression coefficient.
- 19 Sample size.
- 20 Number of pairs in the sample with one or both values missing.

STAT(1) through STAT(8) and STAT(13) are undefined when IOPT = 1. STAT(9) through STAT(12) are undefined when IOPT = 2 or 3. STAT(18) is defined only when IOPT = 3. The elements of STAT that are undefined due to IOPT or an error are set to NaN (not a number).

### Algorithm

Routine SMPRR computes point and interval estimates for the population mean, total, and (optionally) ratio or regression coefficient, using a simple random sample of a variable of interest and an auxiliary variable. Routine SMPRR allows various options for the estimation techniques, which are discussed in Chapters 3, 6, and 7 of Cochran (1977). Let

$$\bar{x} \text{ and } \bar{y}$$

be the sample means of the auxiliary variable and the variable of interest, respectively. Let

$$\bar{X}$$

be the population mean of the auxiliary variable. Then, the ratio estimate of the population mean is

$$\bar{y}_R = \frac{\bar{y}}{\bar{x}} \bar{X}$$

The linear regression estimate of the population mean is



$$\bar{y}_{lr} = \bar{y} + b(\bar{X} - \bar{x})$$

where  $b$  is the regression coefficient, which can be either preassigned, based on previous knowledge, or estimated from the data using least squares. The least-squares estimate of  $b$  is

$$\frac{\sum_{i=1}^n (y_i - \bar{y})(x_i - \bar{x})}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

The confidence limits for the mean and for the total are computed using the normal approximation. If the coefficient of variation of either variable exceeds 10%, then this approximation may not be very accurate.

The parameters IDO and NROW allow either all or part of the data to be brought in.

### Examples

The data for these examples come from Cochran (1977, Table 6.1, page 152). The variable of interest is the population of large U.S. cities in 1930; the auxiliary variable is the 1920 population of the same cities. There are 196 (NPOP) cities that are sampled (that is, that are in the population of interest). (Note that the word “population” is being used in two ways in this discussion.) The total 1920 population of these cities is 22,919 (XMEAN = 116.934). There are 49 cities in the sample. The data can be seen in the DATA statements in the programs below (actual values are 1000 times greater). There are no “missing data”; therefore, the sample size, STAT(19), is 49. Because the coefficient of variation is larger than 10%, SMPRR produces an informational “warning error” message in each example. When the coefficient of variation is larger than 10% (generally speaking), the confidence limits computed using the normal approximation are likely to be shorter than the actual limits at the same confidence level.

#### Example 1

In this example, ratio estimation is used, as on page 151 of Cochran (1977).

```

C      INTEGER      NROW
      PARAMETER    (NROW=49)

C      INTEGER      I, IDO, IOPT, NOUT, NPOP
      REAL          COEF, CONPER, STAT(20), X(NROW), XMEAN, Y(NROW)
      EXTERNAL     SMPRR, UMACH

C      DATA X/76., 138., 67., 29., 381., 23., 37., 120., 61., 387.,
&      93., 172., 78., 66., 60., 46., 2., 507., 179., 121., 50.,
&      44., 77., 64., 64., 56., 40., 40., 38., 136., 116., 46.,
&      243., 87., 30., 71., 256., 43., 25., 94., 43., 298., 36.,
&      161., 74., 45., 36., 50., 48./
      DATA Y/80., 143., 67., 50., 464., 48., 63., 115., 69., 459.,
&      104., 183., 106., 86., 57., 65., 50., 634., 260., 113.,

```

```

&      64., 58., 89., 63., 77., 142., 60., 64., 52., 139., 130.,
&      53., 291., 105., 111., 79., 288., 61., 57., 85., 50., 317.,
&      46., 232., 93., 53., 54., 58., 75./
DATA NPOP/196/, XMEAN/116.934/, CONPER/95./
C
      IDO = 0
      All data are input at once.
C
      Ratio estimation.
      IOPT = 0
      CALL SMPRR (IDO, NROW, X, Y, NPOP, IOPT, XMEAN, COEF, CONPER,
&              STAT)
C
      Print results
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99999) (STAT(I),I=1,17), STAT(19), STAT(20)
99999 FORMAT (/, '
&              RATIO ESTIMATION', /,
&              ' Mean estimate = ', F8.1, ' Total estimate = ',
&              F8.1, /, ' Vhat of mean = ', F8.1, ' Vhat of total '
&              ', ' = ', F8.1, /, ' Confidence limits for mean ', F8.1,
&              ', ', F8.1, /, ' Confidence limits for total ', F8.1,
&              ', ', F8.1, /, ' Ratio estimate = ', F8.3, ' Vhat of '
&              'ratio = ', F8.4, /, ' Confidence limits for ratio ',
&              F8.3, ', ', F8.3, /, ' Coefficient of variation of mean ',
&              'estimate = ', F8.1, /, ' CV of X = ', F8.1,
&              ' CV of Y = ', F8.1, /, ' Mean of X = ',
&              F8.1, ' Mean of Y = ', F8.1, /, ' Sample size '
&              ', ' = ', F8.1, ' Number missing = ', F8.1)
      END

```

### Output

```

*** WARNING ERROR 7 from SMPRR. The coefficient of variation of one or
*** both of the variables exceeds 10%. The confidence limits,
*** which are computed using a normal approximation, may not be
*** very accurate.

```

```

              RATIO ESTIMATION
Mean estimate = 144.9      Total estimate = 28397.1
Vhat of mean = 9.5      Vhat of total = 364860.1
Confidence limits for mean 138.8, 150.9
Confidence limits for total 27213.3, 29581.0
Ratio estimate = 1.239      Vhat of ratio = 0.0007
Confidence limits for ratio 1.187, 1.291
Coefficient of variation of mean estimate = 2.1
CV of X = 89.3      CV of Y = 96.3
Mean of X = 103.1      Mean of Y = 127.8
Sample size = 49.0      Number missing = 0.0

```

### Example 2

In this example, regression estimation with an estimated coefficient is used, as in Exercise 7.3 of Cochran (1977).

```

INTEGER      NROW
PARAMETER    (NROW=49)
C
INTEGER      I, IDO, IOPT, NOUT, NPOP
REAL         COEF, CONPER, STAT(20), X(NROW), XMEAN, Y(NROW)
EXTERNAL     SMPRR, UMACH
C
DATA X/76., 138., 67., 29., 381., 23., 37., 120., 61., 387.,

```

```

&      93., 172., 78., 66., 60., 46., 2., 507., 179., 121., 50.,
&      44., 77., 64., 64., 56., 40., 40., 38., 136., 116., 46.,
&      243., 87., 30., 71., 256., 43., 25., 94., 43., 298., 36.,
&      161., 74., 45., 36., 50., 48./
DATA Y/80., 143., 67., 50., 464., 48., 63., 115., 69., 459.,
&      104., 183., 106., 86., 57., 65., 50., 634., 260., 113.,
&      64., 58., 89., 63., 77., 142., 60., 64., 52., 139., 130.,
&      53., 291., 105., 111., 79., 288., 61., 57., 85., 50., 317.,
&      46., 232., 93., 53., 54., 58., 75./
DATA NPOP/196/, XMEAN/116.934/, CONPER/95./
C
C      All data are input at once.
      IDO = 0
C
C      Regression estimation, with estimated
C      coefficient (Cochran, Exercise 7.3)
      IOPT = 3
      CALL SMPRR (IDO, NROW, X, Y, NPOP, IOPT, XMEAN, COEF, CONPER,
&      STAT)
C
C      Print results
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99999) (STAT(I),I=1,8), (STAT(I),I=13,20)
99999 FORMAT (/, '          REGRESSION ESTIMATION', /,
&      ' Mean estimate = ', F8.1, '      Total estimate = ',
&      F8.1, /, ' Vhat of mean = ', F8.1, '      Vhat of total '
&      ', ' = ', F8.1, /, ' Confidence limits for mean ', F8.1,
&      ', ', F8.1, /, ' Confidence limits for total ', F8.1,
&      ', ', F8.1, /, ' Coefficient of variation of mean '
&      'estimate = ', F8.1, /, ' CV of X = ', F8.1,
&      '          CV of Y = ', F8.1, /, ' Mean of X = ',
&      F8.1, '          Mean of Y = ', F8.1, /, ' Estimated '
&      'regression coefficient = ', F8.1, /, ' Sample size = ',
&      F8.1, '          Number missing = ', F8.1)
      END

```

### Output

```

*** WARNING ERROR 7 from SMPRR. The coefficient of variation of one or
*** both of the variables exceeds 10%. The confidence limits,
*** which are computed using a normal approximation, may not be
*** very accurate.

```

```

          REGRESSION ESTIMATION
Mean estimate =    143.8      Total estimate =  28177.4
Vhat of mean   =     8.6      Vhat of total  = 329372.3
Confidence limits for mean    138.0,    149.5
Confidence limits for total  27052.6, 29302.3
Coefficient of variation of mean estimate =     2.0
CV of X =      89.3          CV of Y =     96.3
Mean of X =    103.1          Mean of Y =    127.8
Estimated regression coefficient =     1.2
Sample size =    49.0          Number missing =     0.0

```

### Example 3

In this example, regression estimation with a preassigned coefficient is used, as in Exercise 7.4 of Cochran (1977).

```

      INTEGER    NROW
      PARAMETER (NROW=49)
C

```

```

INTEGER    I, IDO, IOPT, NOUT, NPOP
REAL       COEF, CONPER, STAT(20), X(NROW), XMEAN, Y(NROW)
EXTERNAL   SMPRR, UMACH
C
DATA X/76., 138., 67., 29., 381., 23., 37., 120., 61., 387.,
& 93., 172., 78., 66., 60., 46., 2., 507., 179., 121., 50.,
& 44., 77., 64., 64., 56., 40., 40., 38., 136., 116., 46.,
& 243., 87., 30., 71., 256., 43., 25., 94., 43., 298., 36.,
& 161., 74., 45., 36., 50., 48./
DATA Y/80., 143., 67., 50., 464., 48., 63., 115., 69., 459.,
& 104., 183., 106., 86., 57., 65., 50., 634., 260., 113.,
& 64., 58., 89., 63., 77., 142., 60., 64., 52., 139., 130.,
& 53., 291., 105., 111., 79., 288., 61., 57., 85., 50., 317.,
& 46., 232., 93., 53., 54., 58., 75./
DATA NPOP/196/, XMEAN/116.934/, CONPER/95./
C
C                                     All data are input at once.
IDO = 0
C
C                                     Regression estimation, with assigned
C                                     coefficient (Cochran, Exercise 7.4)
IOPT = 2
COEF = 1.0
CALL SMPRR (IDO, NROW, X, Y, NPOP, IOPT, XMEAN, COEF, CONPER,
&          STAT)
C
C                                     Print results
CALL UMACH (2, NOUT)
WRITE (NOUT,99999) (STAT(I),I=1,8), (STAT(I),I=13,17), STAT(19),
&          STAT(20)
99999 FORMAT (/, '
REGRESSION ESTIMATION, FIXED ',
& 'COEF', /, ' Mean estimate = ', F8.1, ' Total ',
& 'estimate = ', F8.1, /, ' Vhat of mean = ', F8.1,
& ' Vhat of total = ', F8.1, /, ' Confidence limits '
& ', 'for mean ', F8.1, ', ', F8.1, /, ' Confidence limits '
& ', 'for total ', F8.1, ', ', F8.1, /, ' Coefficient of ',
& 'variation of mean estimate = ', F8.1, /, ' CV of X = '
& ', F8.1, ' CV of Y = ', F8.1, /, ' Mean of '
& ', 'X = ', F8.1, ' Mean of Y = ', F8.1, /,
& ' Sample size = ', F8.1, ' Number missing = ', F8.1)
END

```

### Output

```

*** WARNING ERROR 7 from SMPRR. The coefficient of variation of one or
*** both of the variables exceeds 10%. The confidence limits,
*** which are computed using a normal approximation, may not be
*** very accurate.

```

```

REGRESSION ESTIMATION, FIXED COEF
Mean estimate = 141.6 Total estimate = 27751.1
Vhat of mean = 12.5 Vhat of total = 481977.4
Confidence limits for mean 134.6, 148.5
Confidence limits for total 26390.4, 29111.8
Coefficient of variation of mean estimate = 2.5
CV of X = 89.3 CV of Y = 96.3
Mean of X = 103.1 Mean of Y = 127.8
Sample size = 49.0 Number missing = 0.0

```

---

## SMPRS/DSMPRS (Single/Double precision)

Compute statistics for inferences regarding the population mean and total using ratio or regression estimation given continuous data from a stratified random sample.

### Usage

```
CALL SMPRS (IDO, NSTRAT, NROWS, X, Y, NPOPS, IOPT, ITOPT,  
            XMEANS, CONPER, COEFS, XBARS, XVAR, XCVS,  
            YBARS, YVAR, YCVS, XYCOVS, NSAMPS, STAT)
```

### Arguments

**IDO** — Processing option. (Input)

<b>IDO</b>	<b>Action</b>
0	This is the only invocation of <i>SMPRS</i> for this data set, and all the data are input at once.
1	This is the first invocation, and additional calls to <i>SMPRS</i> will be made. Initialization and updating for the data in <i>X</i> and <i>Y</i> are performed.
2	This is an intermediate invocation of <i>SMPRS</i> , and updating for the data in <i>X</i> and <i>Y</i> is performed.
3	This is the final invocation of this routine. Updating for the data in <i>X</i> and <i>Y</i> and wrap-up computations are performed.

**NSTRAT** — Number of strata into which the population is divided. (Input)  
In the vectors of length *NSTRAT*, the elements are all ordered in the same way. That is, the first stratum is always the first, the second is always the second, and so on.

**NROWS** — Vector of length *NSTRAT* in which  $|NROWS(I)|$  is the number of items from the *I*-th stratum currently input in *X* and *Y*. (Input)  
Each element of *NROWS* may be positive, zero, or negative. A negative value for  $NROWS(I)$  means delete the  $-NROWS(I)$  elements of the *I*-th stratum in *X* and *Y* from the analysis.

**X** — Vector containing the data for the auxiliary variable in the stratified random sample. (Input)  
The observations within any one stratum must appear contiguously in *X*. The first  $|NROWS(1)|$  elements of *X* are from the first stratum, and so on.

**Y** — Vector containing the data for the variable of interest in the stratified random sample. (Input)  
The observations within any one stratum must appear contiguously in *Y*. The first  $|NROWS(1)|$  elements of *Y* are from the first stratum, and so on. The value of  $Y(I)$  corresponds to that of  $X(I)$ .

**NPOPS** — Vector of length *NSTRAT* containing the sizes of the population in the strata. (Input)  
The entries in *NSTRAT* must be ordered in correspondence with the ordering of

strata in the other vectors. If the population strata sizes are not known, estimates must be entered in their place.

**IOPT** — Estimation option. (Input)

**IOPT Action**

- 0 Ratio estimation used for inference about the population mean and total.
- 1 Regression estimation used with the preassigned regression coefficient(s) contained in **COEFS**.
- 2 Regression estimation used with the regression coefficient(s) estimated from the data.

**ITOPT** — Estimation technique option. (Input)

**ITOPT Action**

- 0 Separate ratio or regression estimation.
- 1 Combined ratio or regression estimation.

**XMEANS** — Vector of length **NSTRAT** containing, for each stratum, the population mean of the auxiliary variate, provided **ITOPT** = 0. (Input)  
If **ITOPT** = 1, only **XMEANS**(1) is defined and it must contain the population mean of the auxiliary variate.

**CONPER** — Confidence level for two-sided interval estimate, in percent. (Input)

A **CONPER** percent confidence interval is computed; hence, **CONPER** must be greater than or equal to 0.0 and less than 100.0. **CONPER** is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level **ONECL**, set  $CONPER = 100.0 - 2.0 * (100.0 - ONECL)$ .

**COEFS** — Vector of length **NSTRAT** containing the ratio estimates or the regression coefficients. (Input, if **IOPT** = 1; output, if **IOPT** = 0 or 2 and **IDO** = 0 or 1; input/output, if **IOPT** = 0 or 2 and **IDO** = 2 or 3)

If **IOPT** = 0, **COEFS** contains ratio estimates. When **ITOPT** = 0, **COEFS** contains the estimate of the ratio for each stratum. When **ITOPT** = 1, only **COEFS**(1) is defined and contains the combined estimate of the ratio. If **IOPT** = 1, **COEFS** contains preassigned regression coefficients. When **ITOPT** = 0, **COEFS** contains the preassigned regression coefficient for each stratum. When **ITOPT** = 1, only **COEFS**(1) is defined and contains the preassigned regression coefficient common to all strata. If **IOPT** = 2, **COEFS** contains estimated regression coefficients. When **ITOPT** = 0, **COEFS** contains the estimated regression coefficient for each stratum. When **ITOPT** = 1, only **COEFS**(1) is defined and contains the estimated regression coefficient common to all strata.

**XBARS** — Vector of length **NSTRAT** containing the strata means for the auxiliary variable. (Output, if **IDO** = 0 or 1; input/output, if **IDO** = 2 or 3.)

**XVARS** — Vector of length **NSTRAT** containing the within-strata variances of the auxiliary variable. (Output, if **IDO** = 0 or 1; input/output, if **IDO** = 2 or 3.)

**XCVS** — Vector of length `NSTRAT` containing the within-strata coefficients of variation for the auxiliary variable. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**YBARS** — Vector of length `NSTRAT` containing the strata means for the variable of interest. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**YVARS** — Vector of length `NSTRAT` containing the within-strata variances of the variable of interest. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**YCVS** — Vector of length `NSTRAT` containing the within-strata coefficients of variation for the variable of interest. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**XYCOVS** — Vector of length `NSTRAT` containing the within-strata covariances of the auxiliary variable and the variable of interest. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**NSAMPS** — Vector of length `NSTRAT` containing the number of nonmissing observations from each stratum. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**STAT** — Vector of length 12 containing the resulting statistics. (Output)  
These are

<b>I</b>	<b>STAT(I)</b>
1	Estimate of the mean.
2	Estimate of the total.
3	Variance estimate of the mean estimate.
4	Variance estimate of the total estimate.
5	Lower confidence limit for the mean.
6	Upper confidence limit for the mean.
7	Lower confidence limit for the total.
8	Upper confidence limit for the total.
9	Estimate of the coefficient of variation for the mean and total estimate.
10	Unstratified mean of the auxiliary variate.
11	Unstratified mean of the variable of interest.
12	The number of pairs in the sample that had one or both values missing.

### Algorithm

Routine `SMPRS` computes point and interval estimates for the population mean and total from a stratified random sample of a variable of interest and an auxiliary variable. Routine `SMPRS` allows for either ratio estimation, regression estimation with preassigned coefficients, or regression estimation with estimated coefficients.

This routine follows the standard methods discussed in Chapters 6 and 7 of Cochran (1977). The statistics are similar to those discussed in the documentation for routine `SMPRR` (page 911), except that they are computed from stratified data. The option parameter `IOPT` allows selection of either ratio

or regression estimation, and the parameter `ITOPT` allows selection of separate or combined estimators. “Separate” estimators means that each stratum is allowed to have different ratios or regression coefficients, while “combined” means these are assumed to be the same over all strata.

The confidence limits for the mean and for the total are computed using the normal approximation. If the coefficient of variation of either variable exceeds 10%, then this approximation may not be very accurate.

The parameters `IDO` and `NROW` allow either all or part of the data to be brought in at one time.

### Example 1

In the following example, we use a stratified sample from the data in Table 5.1 of Cochran (1977), which consists of the 1920 and the 1930 population (in 1000’s) of 64 cities in the United States. The objective is to estimate the mean and total 1930 population of the 64 cities, using a sample of size 24 of the 1920 and 1930 populations. There are two strata: the largest 16 cities and the remaining cities. We use stratified sampling with equal sample sizes. The same example is also used to illustrate routine `SMPSS` (page 930), except here we have an auxiliary variable.

In this example, separate ratio estimation is used.

```

C      INTEGER      NSTRAT
      PARAMETER    (NSTRAT=2)

C      INTEGER      I, IDO, IOPT, ITOPT, NOUT, NPOPS(NSTRAT),
&      NROWS(NSTRAT), NSAMPS(NSTRAT)
      REAL          COEFS(NSTRAT), CONPER, STAT(12), X(24),
&      XBARS(NSTRAT), XCVS(NSTRAT), XMEANS(NSTRAT),
&      XVARS(NSTRAT), XYCOVS(NSTRAT), Y(24), YBARS(NSTRAT),
&      YCVS(NSTRAT), YVARS(NSTRAT)
C      EXTERNAL    SMPRS, UMACH

C      DATA X/773., 748., 734., 577., 507., 438., 415., 401., 387.,
&      381., 324., 315., 258., 237., 235., 216., 201., 179., 136.,
&      132., 118., 118., 106., 104./
C      DATA Y/822., 781., 805., 1238., 634., 487., 442., 451., 459.,
&      464., 400., 366., 302., 291., 272., 284., 270., 260., 139.,
&      170., 154., 140., 163., 116./

C      NPOPS(1) = 16
C      NPOPS(2) = 48
C                                     All data are input at once.
C      IDO      = 0
C      NROWS(1) = 12
C      NROWS(2) = 12
C      CONPER   = 95.0
C                                     Use separate ratio estimation.
C      IOPT     = 0
C      ITOPT    = 0
C      XMEANS(1) = 521.8
C      XMEANS(2) = 165.4

```



```

C
  CALL SMPRS (IDO, NSTRAT, NROWS, X, Y, NPOPS, IOPT, ITOPT,
&            XMEANS, CONPER, COEFS, XBARS, XVARS, XCVS, YBARS,
&            YVARS, YCVS, XYCOVS, NSAMPS, STAT)
C
  PRINT results
  CALL UMACH (2, NOUT)
  WRITE (NOUT,99999) (STAT(I),I=1,9), STAT(12), COEFS
99999 FORMAT (' Mean estimate = ', F8.3, '      Total estimate = ',
&           F8.1, /, ' Vhat of mean = ', F8.5, '      Vhat of total '
&           ', ' = ', F8.1, /, ' Confidence limits for mean ', F8.3,
&           ', ', F8.3, /, ' Confidence limits for total ', F8.1,
&           ', ', F8.1, /, ' C. V.      = ', F8.2, '      Number ',
&           'missing = ', F8.1, /, ' Estimated ratios = ', 2F10.3)
END

```

### Output

```

Mean estimate = 315.511      Total estimate = 20192.7
Vhat of mean = 55.56254     Vhat of total = 227584.2
Confidence limits for mean 300.901, 330.120
Confidence limits for total 19257.7, 21127.7
C. V.      = 2.36      Number missing = 0.0
Estimated ratios = 1.225 1.255

```

### Example 2

In the following example, we use a stratified sample from the data in Table 5.1 of Cochran (1977), which consists of the 1920 and the 1930 population (in 1000's) of 64 cities in the United States. The objective is to estimate the mean and total 1930 population of the 64 cities, using a sample of size 24 of the 1920 and 1930 populations. There are two strata: the largest 16 cities and the remaining cities. We use stratified sampling with equal sample sizes. The same example is also used to illustrate routine SMPSS (page 930), except here we have an auxiliary variable.

In this example, regression estimation is used, and it is assumed that the regression equation is the same in the two strata.

```

  INTEGER    NSTRAT
  PARAMETER  (NSTRAT=2)
C
  INTEGER    I, IDO, IOPT, ITOPT, NOUT, NPOPS(NSTRAT),
&           NROWS(NSTRAT), NSAMPS(NSTRAT)
  REAL      COEFS(NSTRAT), CONPER, STAT(12), X(24),
&           XBARS(NSTRAT), XCVS(NSTRAT), XMEANS(1),
&           XVARS(NSTRAT), XYCOVS(NSTRAT), Y(24), YBARS(NSTRAT),
&           YCVS(NSTRAT), YVARS(NSTRAT)
  EXTERNAL  SMPRS, UMACH
C
  DATA X/773., 748., 734., 577., 507., 438., 415., 401., 387.,
&       381., 324., 315., 258., 237., 235., 216., 201., 179., 136.,
&       132., 118., 118., 106., 104./
  DATA Y/822., 781., 805., 1238., 634., 487., 442., 451., 459.,
&       464., 400., 366., 302., 291., 272., 284., 270., 260., 139.,
&       170., 154., 140., 163., 116./
C
  NPOPS(1) = 16

```

```

      NPOPS(2) = 48
C
      IDO      = 0
      NROWS(1) = 12
      NROWS(2) = 12
      CONPER   = 95.0
C
      IOPT     = 2
      ITOPT    = 1
      XMEANS(1) = 254.5
C
      CALL SMPRS (IDO, NSTRAT, NROWS, X, Y, NPOPS, IOPT, ITOPT,
&
      XMEANS, CONPER, COEFS, XBARS, XVAR, XCVS, YBARS,
&
      YVAR, YCVS, XYCOVS, NSAMPS, STAT)
C
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99999) (STAT(I),I=1,9), STAT(12), COEFS(1)
99999 FORMAT (' Mean estimate = ', F8.3, ' Total estimate = ',
&
      F8.1, /, ' Vhat of mean = ', F8.5, ' Vhat of total '
&
      ', ' = ', F8.1, /, ' Confidence limits for mean ', F8.3,
&
      ', ', F8.3, /, ' Confidence limits for total ', F8.1,
&
      ', ', F8.1, /, ' C. V. = ', F8.1, ' Number ',
&
      'missing = ', F8.1, /, ' Estimated combined regression ',
&
      'coefficient = ', F8.3)
      END

```

### Output

```

Mean estimate = 315.517      Total estimate = 20193.1
Vhat of mean = 54.84098     Vhat of total = 224628.6
Confidence limits for mean 301.003, 330.031
Confidence limits for total 19264.2, 21122.0
C. V. = 2.3      Number missing = 0.0
Estimated combined regression coefficient = 1.175

```

---

## SMPSC/DSMPSC (Single/Double precision)

Compute statistics for inferences regarding the population mean and total using single stage cluster sampling with continuous data.

### Usage

```

CALL SMPSC (IDO, NCLSTR, NROWS, Y, IOPT, NCLPOP, NPOP,
      SIZE, TSIZE, CONPER, CLMEAN, CLVAR, NSAMPS,
      STAT)

```

### Arguments

**IDO** — Processing option. (Input)

#### IDO Action

- 0 This is the only invocation of SMPSC for this data set, and all the data are input at once.
- 1 This is the first invocation, and additional calls to SMPSC will be made. Initialization and updating for the data in Y are performed.

- 2 This is an intermediate invocation of *SMPSC* and updating for the data in *Y* is performed.
- 3 This is the final invocation of this routine. Updating for the data in *Y* and wrap-up computations are performed.

***NCLSTR*** — Number of clusters into which the sample is divided. (Input)  
 In the vectors of length *NCLSTR*, the elements are all ordered in the same way. That is, the first cluster is always the first, the second always the second, and so on.

***NROWS*** — Vector of length *NCLSTR* in which  $|NROWS(I)|$  is the number of items from the *I*-th cluster currently input in *Y*. (Input)  
 Each element of *NROWS* may be positive, zero, or negative. A negative value for *NROWS(I)* means delete the  $-NROWS(I)$  elements of the *I*-th cluster in *Y* from the analysis.

***Y*** — Vector containing the cluster sample. (Input)  
 The observations within any one cluster must appear contiguously in *Y*. The first  $|NROWS(1)|$  elements of *Y* are from the first cluster, and so on.

***IOPT*** — Estimation option. (Input)

<b><i>IOPT</i></b>	<b>Action</b>
0	Ratio-to-size estimation is used.
1	Unbiased estimation is used.
2	Probability-proportional-to-size estimation is used and all clusters in population are of known size.
3	Probability-proportional-to-size estimation is used and the cluster sizes are known only approximately or a measure of cluster size other than the number of elements per cluster is to be used.

***NCLPOP*** — Number of clusters in the sampled population. (Input)

***NPOP*** — Number of elements in the population (sum of all the cluster sizes in the population). (Input)  
*NPOP* is not required when *IOPT* = 3.

***SIZE*** — If *IOPT* = 3, vector of length *NCLSTR* containing a measure of cluster size for each cluster in the sample. (Input)  
 The sampled cluster size measures must be ordered in correspondence with the ordering of clusters in *Y*. *SIZE* is required only when *IOPT* = 3.

***TSIZE*** — If *IOPT* = 3, measure of total size of all clusters in the population. (Input) *TSIZE* is required only when *IOPT* = 3.

***CONPER*** — Confidence level for two-sided interval estimate, in percent. (Input)

A *CONPER* percent confidence interval is computed; hence, *CONPER* must be greater than or equal to 0.0 and less than 100.0. *CONPER* is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level *ONECL*, set  $CONPER = 100.0 - 2.0 * (100.0 - ONECL)$ .

**CLMEAN** — Vector of length `NCLSTR` containing the cluster means. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**CLVAR** — Vector of length `NCLSTR` containing the within-cluster variances. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**NSAMPS** — Vector of length `NCLSTR` containing the number of nonmissing observations from each cluster. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

**STAT** — Vector of length 11 containing the resulting statistics. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2` or `3`.)

These are

<b>I</b>	<b>STAT(I)</b>
1	Estimate of the mean.
2	Estimate of the total.
3	Variance estimate of the mean estimate.
4	Variance estimate of the total estimate.
5	Lower confidence limit for the mean.
6	Upper confidence limit for the mean.
7	Lower confidence limit for the total.
8	Upper confidence limit for the total.
9	Estimate (expressed as a percentage) of the coefficient of variation of the mean and total estimate.
10	The total sample size.
11	The number of missing values.

### Algorithm

Routine `SMPSC` computes point and interval estimates for the population mean and total from a single-stage cluster sample. The routine uses the standard methods discussed in Chapters 9 and 9A of Cochran (1977). The sample means for the individual clusters are accumulated in `CLMEAN`, and the corrected sums of squares are accumulated in `CLVAR`. In the postprocessing phase, the quantities in `STAT` are computed using the cluster statistics in `CLMEAN`, `CLVAR`, and `NSAMPS`. The parameters `IDO` and `NROWS` allow either all or part of the data to be brought in at one time.

Following the notation of Cochran (1977), let  $N$  be the number of clusters in the population, let  $M_i$  be the number of elements in the  $i$ -th cluster unit, let  $M_0$  be the total number of elements in the population, let  $y_{ij}$  be the  $j$ -th element in the  $i$ -th cluster, and let  $n$  be the number of clusters in the sample. Any of three different estimators of the population total may be useful. An unbiased estimate of the total is

$$\frac{N}{n} \sum_{i=1}^n \sum_{j=1}^{M_i} y_{ij}$$

The ratio-to-size estimate is

$$M_0 \frac{\sum_{i=1}^n y_i}{\sum_{i=1}^n M_i}$$

The probability-proportional-to-size estimate is

$$\frac{M_0}{n} \sum_{i=1}^n \left( \frac{y_i}{M_i} \right)$$

The confidence limits for the mean and total are computed using the normal approximation.

### Example

In this example, we have a sample of two clusters from a population that contains 20 clusters. The sizes of the clusters in the sample are four and six, and there is a total of 100 elements in the population.

```

INTEGER      NCLSTR
PARAMETER   (NCLSTR=2)
C
  INTEGER    IDO, IOPT, NCLPOP, NOUT, NPOP, NROWS(NCLSTR),
&           NSAMPS(NCLSTR)
  REAL      CLMEAN(NCLSTR), CLVAR(NCLSTR), CONPER, SIZE(NCLSTR),
&           STAT(11), TSIZE, Y(10)
  EXTERNAL  SMPSC, UMACH
C
  DATA Y/2.7, 5.1, 4.3, 2.8, 1.9, 6.2, 4.8, 5.1, 7.2, 6.5/
C
  IOPT      = 0
  NCLPOP    = 20
  NPOP      = 100
C
  All data are input at once.
  IDO       = 0
  NROWS(1)  = 4
  NROWS(2)  = 6
  CONPER    = 95.0
  CALL SMPSC (IDO, NCLSTR, NROWS, Y, IOPT, NCLPOP, NPOP, SIZE,
&           TSIZE, CONPER, CLMEAN, CLVAR, NSAMPS, STAT)
C
  Print results
  CALL UMACH (2, NOUT)
  WRITE (NOUT,99999) STAT
99999 FORMAT (' Mean estimate = ', F8.3, '      Total estimate = ',
&           F8.1, /, ' Vhat of mean = ', F8.3, '      Vhat of total '
&           ', ' = ', F8.1, /, ' Confidence limits for mean ', F8.3,
&           ', ' , F8.3, /, ' Confidence limits for total ', F8.1,
&           ', ' , F8.1, /, ' C. V.          = ', F8.1, '%', /,
&           ' Sample size = ', F8.0, '      Number missing = ',
&           F8.0)
  END

```

<b>Output</b>			
Mean estimate =	4.660	Total estimate =	466.0
Vhat of mean =	0.504	Vhat of total =	5035.5
Confidence limits for mean	3.269,	6.051	
Confidence limits for total	326.9,	605.1	
C. V. =	15.2%		
Sample size =	10.	Number missing =	0.

---

## SMPSR/DSMPSR (Single/Double precision)

Compute statistics for inferences regarding the population mean and total, given data from a simple random sample.

### Usage

CALL SMPSR (IDO, NROW, Y, NPOP, IOPT, NSAMPO, CONPER, STAT)

### Arguments

**IDO** — Processing option. (Input)

#### IDO Action

- |   |  |
|---|--|
| 0 | This is the only invocation of SMPSR for this data set, and all the data are input at once.  |
| 1 | This is the first invocation, and additional calls to SMPSR will be made. Initialization and updating for the data in Y are performed. |
| 2 | This is an intermediate invocation of SMPSR, and updating for the data in Y is performed.  |
| 3 | This is the final invocation of this routine. Updating for the data in Y and wrap-up computations are performed.                       |

**NROW** — The absolute value of NROW is the number of rows of data currently input in Y. (Input)

NROW may be positive, zero, or negative. Negative -NROW means delete the NROW rows of data from the analysis.

**Y** — Vector of length |NROW| containing the sample data. (Input)

**NPOP** — Size of the (full) population. (Input)

**IOPT** — Subpopulation option. (Input)

If IOPT = 0, no subpopulation is assumed. If IOPT = 1, the input data come from a subpopulation (“domain of study”) of unknown size.

**NSAMPO** — Size of the sample from the full population, if a subpopulation is sampled (that is, if IOPT = 1). (Input)

**CONPER** — Confidence level for two-sided interval estimate, in percent. (Input)

A CONPER percent confidence interval is computed; hence, CONPER must be greater than or equal to 0.0 and less than 100.0. CONPER is often 90.0, 95.0, or

99.0. For a one-sided confidence interval with confidence level ONECL, set  
 $CONPER = 100.0 - 2.0 * (100.0 - ONECL)$ .

**STAT** — Vector of length 11 containing the resulting statistics. (Output, if IDO = 0 or 1; input/output, if IDO = 2 or 3.)

These are

<b>I</b>	<b>STAT(I)</b>
1	Estimate of the mean.
2	Estimate of the total.
3	Within-sample variance estimate.
4	Variance estimate of the mean estimate.
5	Variance estimate of the total estimate.
6	Lower confidence limit for the mean.
7	Upper confidence limit for the mean.
8	Lower confidence limit for the total.
9	Upper confidence limit for the total.
10	The sample size.
11	The number of missing values.

### Algorithm

Routine SMPSR computes point and interval estimates for the population mean and total from a simple random sample of one variable. The routine uses the standard methods discussed in Chapter 2 of Cochran (1977). The sample mean is accumulated in STAT(1) and the corrected sum of squares is accumulated in STAT(3). In the postprocessing phase, STAT(3) is divided by the sample size minus one, and then the other quantities in STAT are computed. The parameters IDO and NROW allow either all or part of the data to be brought in at one time.

By use of IOPT and NSAMPO, SMPSR can also be used to analyze data from a subpopulation or “domain of study”. (See Cochran 1977, pages 34–38.) In the case of a subpopulation, only the estimates relating to the subpopulation total differ from the corresponding estimates when no subpopulation is assumed. Of course, if a subpopulation is of known size, it should be considered the full population.

### Example 1

This example uses artificial data to illustrate a simple use of SMPSR to compute point and interval estimates of the population mean and total. The sample size is 15, from a population of size 150.

```

C      INTEGER      NROW
      PARAMETER    (NROW=15)

C      INTEGER      IDO, IOPT, NOUT, NPOP, NSAMPO
      REAL          CONPER, STAT(11), Y(NROW)
      EXTERNAL      SMPSR, UMACH

C      DATA Y/21., 14., 17., 22., 19., 21., 20., 15., 24., 28., 20.,

```

```

&      17., 16., 22., 19./
C
  NPOP  = 150
  CONPER = 95.0
C
  IDO = 0
C
  IOPT = 0
  CALL SMPSR (IDO, NROW, Y, NPOP, IOPT, NSAMPO, CONPER, STAT)
C
  CALL UMACH (2, NOUT)
  WRITE (NOUT,99999) STAT
99999 FORMAT (' Mean estimate = ', F8.3, '          Total estimate = ',
&            F8.1, '/', ' Within-sample variance estimate = ', F8.3, '/',
&            ' VWHAT of mean = ', F8.5, '          VWHAT of total = ',
&            F8.1, '/', ' Confidence limits for mean ', F8.3,
&            ', ', F8.3, '/', ' Confidence limits for total ', F8.1,
&            ', ', F8.1, '/', ' Sample size = ', F8.1, '          Number ',
&            'missing = ', F8.0)
  END

```

### Output

```

Mean estimate = 19.667          Total estimate = 2950.0
Within-sample variance estimate = 13.238
VWHAT of mean = 0.79429        VWHAT of total = 17871.4
Confidence limits for mean 17.755, 21.578
Confidence limits for total 2663.3, 3236.7
Sample size = 15.0          Number missing = 0.

```

### Example 2

This example is a problem of estimation in a subpopulation described on page 37 of Cochran (1977). The example illustrates how the IDO and NROW parameters can be used to allow input other than the actual data. Cochran gives only the sample total and uncorrected sum of squares, so these values are transformed to the mean and corrected sum of squares prior to input as STAT(1) and STAT(3).

```

INTEGER IDO, IOPT, NOUT, NPOP, NROW, NSAMPO
REAL CONPER, SQRT, STAT(11), Y(1)
INTRINSIC SQRT
EXTERNAL SMPSR, UMACH
C
  NPOP  = 2422
  CONPER = 95.0
C
  IDO = 0
C
  IOPT = 1
  NSAMPO = 180
C
  There are 180 items in the complete
  sample, but only a subpopulation is
  of interest.
C
  For this example, STAT is
  initialized as if the data
  have been already processed and only
  the postprocessing computations are
  to be done. There are 152 items of
  interest in the sample. The sample
  total is 343.5 and the uncorrected

```



```

C                                     sum of squares is 1491.38.
C                                     STAT(1) is initialized to the sample
C                                     mean by dividing the total by the
C                                     sample size, and STAT(3) is
C                                     initialized to the corrected sum of
C                                     squares.
C
STAT(1) = 343.5/152.0
STAT(3) = 1491.38 - 152.0*STAT(1)**2
STAT(10) = 152.0
STAT(11) = 0.0
IDO = 3
NROW = 0
CALL SMPSS (IDO, NROW, Y, NPOP, IOPT, NSAMPO, CONPER, STAT)
C                                     Print results
CALL UMACH (2, NOUT)
WRITE (NOUT,99999) STAT(2), SQRT(STAT(5))
99999 FORMAT ('      Total estimate = ', F8.1, '/', '      Standard ',
&          'deviation of the estimate = ', F8.1)
END

```

### Output

```

Total estimate = 4622.0
Standard deviation of the estimate = 375.3

```

---

## SMPSS/DSMPSS (Single/Double precision)

Compute statistics for inferences regarding the population mean and total, given data from a stratified random sample.

### Usage

```

CALL SMPSS (IDO, NSTRAT, NROWS, Y, NPOPS, IVOPT, CONPER,
            YBARS, YVARS, NSAMPS, STAT)

```

### Arguments

**IDO** — Processing option. (Input)

#### IDO Action

- 0 This is the only invocation of SMPSS for this data set, and all the data are input at once.
- 1 This is the first invocation, and additional calls to SMPSS will be made. Initialization and updating for the data in Y are performed.
- 2 This is an intermediate invocation of SMPSS, and updating for the data in Y is performed.
- 3 This is the final invocation of this routine. Updating for the data in Y and wrap-up computations are performed.

**NSTRAT** — Number of strata into which the population is divided. (Input)

In the vectors of length NSTRAT, the elements are all ordered in the same way. That is, the first stratum is always the first, the second always the second, and so on.

**NROWS** — Vector of length **NSTRAT** in which  $\text{NROWS}(I)$  is the number of items from the  $I$ -th stratum currently input in **Y**. (Input)

Each element of **NROWS** may be positive, zero, or negative. A negative value for  $\text{NROWS}(I)$  means delete the  $-\text{NROWS}(I)$  elements of the  $I$ -th stratum in **Y** from the analysis.

**Y** — Vector containing the stratified random sample. (Input)

The observations within any one stratum must appear contiguously in **Y**. The first  $\text{NROWS}(1)$  elements of **Y** are from the first stratum, and so on.

**NPOPS** — Vector of length **NSTRAT** containing the sizes of the population in the strata. (Input)

The entries must be ordered in correspondence with the ordering of strata in the other vectors. If the population strata sizes are not known, estimates must be entered in their place.

**ILOPT** — Within-stratum variance assumption indicator. (Input)

If **ILOPT** = 0, the true within-stratum variance is assumed constant, and a pooled estimate of that variance is returned in **STAT**(12). If **ILOPT** = 1, separate within-strata variance estimates are assumed.

**CONPER** — Confidence level for two-sided interval estimate, in percent. (Input)

A **CONPER** percent confidence interval is computed; hence, **CONPER** must be greater than or equal to 0.0 and less than 100.0. **CONPER** is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level **ONECL**, set  $\text{CONPER} = 100.0 - 2.0 * (100.0 - \text{ONECL})$ .

**YBARS** — Vector of length **NSTRAT** containing the strata means. (Output, if **IDO** = 0 or 1; input/output, if **IDO** = 2 or 3.)

**YVARS** — Vector of length **NSTRAT** containing the within-strata variances. (Output, if **IDO** = 0 or 1; input/output, if **IDO** = 2 or 3.)

**NSAMPS** — Vector of length **NSTRAT** containing the number of nonmissing observations from each stratum. (Output, if **IDO** = 0 or 1; input/output, if **IDO** = 2 or 3.)

**STAT** — Vector of length 13 containing the resulting statistics. (Output, if **IDO** = 0 or 1; input/output, if **IDO** = 2 or 3.)

These are

<b>I</b>	<b>STAT(I)</b>
1	Estimate of the mean.
2	Estimate of the total.
3	Variance estimate of the mean estimate.
4	Variance estimate of the total estimate.
5	Lower confidence limit for the mean.
6	Upper confidence limit for the mean.
7	Lower confidence limit for the total.
8	Upper confidence limit for the total.

- 9 Estimate of the coefficient of variation of the mean and total estimates.
- 10 Number of degrees of freedom associated with the variance estimates of the mean and total estimates. When `IVOPT = 1`, `STAT(10)` contains an effective number of degrees of freedom determined according to the Satterthwaite approximation.
- 11 Variance estimate of the mean estimate assuming that sampling was simple random instead of stratified random.
- 12 Pooled estimate of the common variance, when `IVOPT = 0`. If `IVOPT = 1`, `STAT(12)` is not defined.
- 13 The number of missing values.

### Comments

#### Information Error

Type	Code	
4	1	The population size for each stratum is equal to one.

### Algorithm

Routine `SMPSS` computes point and interval estimates for the population mean and total from a stratified random sample of one variable. The routine uses the standard methods discussed in Chapters 5 and 5A of Cochran (1977). The sample means for the individual strata are accumulated in `YBARS`, and the corrected sums of squares are accumulated in `YVARS`. In the postprocessing phase, the quantities in `STAT` are computed using the strata statistics in `YBARS`, `YVARS`, and `NSAMPS`. The parameters `IDO` and `NROWS` allow either all or part of the data to be brought in at one time.

### Example

In this example, we use a stratified sample from the data in Table 5.1 of Cochran (1977): the 1930 population (in 1000's) of 64 cities in the United States. The 64 cities are the "population", and our objective is to estimate the mean and total number of inhabitants in these 64 cities. There are two strata: the largest 16 cities and the remaining cities. We use stratified sampling with equal sample sizes. To choose the random sample, we use routine `RNSRI` (page 1241), as follows:

```

INTEGER      ISEED, NSAMP, NPOP, INDEX(12)
NSAMP = 12
NPOP = 16
ISEED = 123457
CALL RNSSET(ISEED)
CALL RNSRI(NSAMP, NPOP, INDEX)
WRITE(*, *) INDEX
NPOP = 48
CALL RNSRI(NSAMP, NPOP, INDEX)
WRITE(*, *) INDEX
END

```

This yields the population indices {2, 3, 4, 6, 8, 10, 11, 12, 13, 14, 15, 16} for the first stratum and {4, 8, 10, 11, 13, 16, 29, 30, 36, 37, 45, 46} for the second stratum. The corresponding values from Table 5.1 are encoded in the program below.

```

INTEGER      NSTRAT
PARAMETER   (NSTRAT=2)
C
INTEGER      I, IDO, IVOPT, NOUT, NPOPS(NSTRAT), NROWS(NSTRAT),
&           NSAMPS(NSTRAT)
REAL         CONPER, STAT(13), Y(24), YBARS(NSTRAT), YVARS(NSTRAT)
EXTERNAL     SMPSS, UMACH
C
DATA Y/822., 781., 805., 1238., 634., 487., 442., 451., 459.,
&      464., 400., 366., 302., 291., 272., 284., 270., 260., 139.,
&      170., 154., 140., 163., 116./
C
NPOPS(1) = 16
NPOPS(2) = 48
IVOPT    = 1
C
                                     All data are input at once.
IDO      = 0
NROWS(1) = 12
NROWS(2) = 12
CONPER   = 95.0
CALL SMPSS (IDO, NSTRAT, NROWS, Y, NPOPS, IVOPT, CONPER, YBARS,
&          YVARS, NSAMPS, STAT)
C
                                     Print results
CALL UMACH (2, NOUT)
WRITE (NOUT,99999) (STAT(I),I=1,11), STAT(13)
99999 FORMAT (' Mean estimate = ', F8.3, '          Total estimate = ',
&           F9.1, /, ' Vhat of mean = ', F8.3, '          Vhat of total '
&           ', ' = ', F9.1, /, ' Confidence limits for mean ', F8.3,
&           ', ', F8.3, /, ' Confidence limits for total ', F8.1,
&           ', ', F8.1, /, ' C. V.          = ', F8.1, '          Degrees '
&           ', 'of freedom = ', F8.1, /, ' SRS var. estimate = ',
&           F8.3, ' Number missing = ', F8.0)
END

```

### Output

```

Mean estimate = 313.167      Total estimate = 20042.7
Vhat of mean   = 264.703      Vhat of total = 1084224.6
Confidence limits for mean 279.180, 347.153
Confidence limits for total 17867.5, 22217.8
C. V.          = 5.2         Degrees of freedom = 19.6
SRS var. estimate = 1288.075 Number missing = 0.

```

---

## SMPST/DSMPST (Single/Double precision)

Compute statistics for inferences regarding the population mean and total given continuous data from a two-stage sample with equisized primary units.

## Usage

CALL SMPST (IDO, NUNSAM, NELSAM, NOBS, Y, NUNPOP, NELPOP,  
CONPER, PUMEAN, PUVAR, STAT)

## Arguments

**IDO** — Processing option. (Input)

### IDO Action

- |   |  |
|---|--|
| 0 | This is the only invocation of SMPST for this data set, and all the data are input at once.  |
| 1 | This is the first invocation, and additional calls to SMPST will be made. Initialization and updating for the data in Y are performed. |
| 2 | This is an intermediate invocation of SMPST, and updating for the data in Y is performed.  |
| 3 | This is the final invocation of this routine. Updating for the data in Y and wrap-up computations are performed.                       |

**NUNSAM** — Number of primary units into which the sample is divided. (Input)

**NELSAM** — Number of elements in the sample in each sampled primary unit. (Input)

**NOBS** — The number of observations currently input in Y. (Input)

NOBS may be positive or zero. If NOBS = 0, IDO must equal 3, and only wrap-up computations are performed.

**Y** — Vector of length NOBS containing the elements of the two-stage sample. (Input)

The elements from each primary unit must occur contiguously within Y. Since there must be an equal number from each primary unit, Y must contain no missing values.

**NUNPOP** — Number of primary units in the sampled population. (Input)

**NELPOP** — Number of elements in each primary unit in the population. (Input)

**CONPER** — Confidence level for two-sided interval estimate, in percent. (Input)

A CONPER percent confidence interval is computed; hence, CONPER must be greater than or equal to 0.0 and less than 100.0. CONPER is often 90.0, 95.0, or 99.0. For a one-sided confidence interval with confidence level ONECL, set  $CONPER = 100.0 - 2.0 * (100.0 - ONECL)$ .

**PUMEAN** — Vector of length NUNSAM containing the means of the primary units in the sample. (Output, if IDO = 0 or 1; input/output, if IDO = 2 or 3)

The estimates are ordered in correspondence with the ordering of primary units in Y.

**PUVAR** — Vector of length `NUMSAM` containing the sample variances of the primary units in the sample. (Output, if `IDO=0` or `1`; input/output, if `IDO=2` or `3`)

The estimates are ordered in correspondence with the ordering of primary units in `Y`.

**STAT** — Vector of length `9` containing the resulting statistics. (Output, if `IDO=0` or `1`; input/output, if `IDO=2` or `3`)

<b>I</b>	<b>STAT(I)</b>
1	Estimate of the mean.
2	Estimate of the total.
3	Variance of the mean estimate.
4	Variance estimate of the total estimate.
5	Lower confidence limit for the mean.
6	Upper confidence limit for the mean
7	Lower confidence limit for the total.
8	Upper confidence limit for the total.
9	Estimate (expressed as a percentage) of the coefficient of variation of the mean and total estimates.

### Algorithm

Routine `SMPST` computes point and interval estimates for the population mean and total from a two-stage sample with primary units that are all equal in size. A two-stage sample might be taken if each unit (“primary unit”) in the population can be divided into smaller units. Primary units are selected first, and then those selected are subsampled. The routine uses the standard methods discussed in Chapter 10 of Cochran (1977). The sample means for the individual primary units are accumulated in `PUMEAN`, and the corrected sums of squares are accumulated in `PUVAR`. In the postprocessing phase, the quantities in `STAT` are computed using the primary unit statistics. The parameters `IDO` and `NOBS` allow either all or part of the data to be brought in at one time.

Following the notation of Cochran (1977), let  $n$  (`NUMSAM`) be the number of primary units in the sample, let  $m$  (`NELSAM`) be the number of elements (subunits) subsampled from each primary unit, let  $N$  (`NUMPOP`) be the total number of primary units in the population, let  $M$  (`NELPOP`) be the total number of elements in each primary unit (in the population), and let  $y_{ij}$  be the  $j$ -th element in the  $i$ -th primary unit. The sample mean per subunit in the  $i$ -th primary unit is

$$\bar{y}_i = \frac{1}{m} \sum_{j=1}^m y_{ij}$$

The estimate of the population mean is

$$\bar{\bar{y}} = \frac{1}{nm} \sum_{i=1}^n \sum_{j=1}^m y_{ij}$$

The estimate of the variance of

$$\bar{\bar{y}} \text{ is } \frac{N-n}{nN(n-1)} \sum_{i=1}^n (\bar{y}_i - \bar{\bar{y}})^2 + \frac{M-m}{m(m-1)MnN} \sum_{i=1}^n \sum_{j=1}^m (y_{ij} - \bar{y}_i)^2$$

### Example

In this example, we have a sample of two primary units, with five subunits from each. The population consists of 10 primary units with 15 elements each.

```

INTEGER      IDO, NELPOP, NELSAM, NOBS, NOUT, NUNPOP, NUNSAM
REAL         CONPER, PUMEAN(2), PUVAR(2), STAT(9), Y(10)
EXTERNAL     SMPST, UMACH
C
DATA Y/2.7, 5.1, 4.3, 2.8, 1.9, 6.2, 4.8, 5.1, 7.2, 6.5/
C
NUNSAM = 2
NELSAM = 5
NOBS   = 10
NUNPOP = 10
NELPOP = 15
C
                                All data are input at once.
IDO     = 0
CONPER  = 95.0
CALL SMPST (IDO, NUNSAM, NELSAM, NOBS, Y, NUNPOP, NELPOP,
&          CONPER, PUMEAN, PUVAR, STAT)
C
                                Print results
CALL UMACH (2, NOUT)
WRITE (NOUT,99999) STAT
99999 FORMAT (' Mean estimate = ', F8.3, '          Total estimate = ',
&           F8.1, /, ' Vhat of mean = ', F8.3, '          Vhat of total '
&           ', ' = ', F8.1, /, ' Confidence limits for mean ', F8.3,
&           ', ', F8.3, /, ' Confidence limits for total ', F8.1,
&           ', ', F8.1, /, ' C. V.          = ', F8.1, '%')
END

```

### Output

```

Mean estimate =    4.660          Total estimate =    699.0
Vhat of mean   =    1.370          Vhat of total   =   30823.7
Confidence limits for mean    2.366,    6.954
Confidence limits for total   354.9,   1043.1
C. V.          =    25.1%

```

# Chapter 13: Survival Analysis, Life Testing, and Reliability

---

## Routines

<b>13.1. Survival Analysis</b>		
Kaplan-Meier estimates .....	KAPMR	938
Print Kaplan-Meier estimates .....	KTBLE	942
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Analyze time event data using a proportional hazards model.....	PHGLM	951
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Estimates using various parametric models .....	STBLE	985
<b>13.2. Actuarial Tables</b>		
Current and cohort tables .....	ACTBL	992

---

## Usage Notes

The routines described in this chapter have primary application in the areas of reliability and life testing, but they may find application in any situation in which time is a variable of interest. Kalbfleisch and Prentice (1980), Elandt-Johnson and Johnson (1980), Lee (1980), Gross and Clark (1975), Lawless (1982), and Chiang (1968) are general references for discussing the models and methods used here.

Kaplan-Meier (product-limit) estimates of the survival distribution in a single population is available through routine KAPMR (page 938), and these can be printed using KTBLE (page 942). Routine TRNBL (page 946) computes generalized Kaplan-Meier estimates. Routine PHGLM (page 951) computes the parameter estimates in a proportional hazards model. Routine SVGLM (page 967) fits any of several generalized linear models, and STBLE (page 985) computes estimates of survival probabilities based on the same models. Routine ACTBL (page 992) computes and (optionally) prints an actuarial table based either upon a cohort followed over time or a cross-section of a population.



---

## KAPMR/DKAPMR (Single/Double precision)

Compute Kaplan-Meier estimates of survival probabilities in stratified samples.

### Usage

```
CALL KAPMR (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN, IGRP,  
           ISRT, SPROB, LDSPRO, NRMISS)
```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in  $X$ . (Input)

**X** — NOBS by NCOL matrix containing the data. (Input)

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

**IRT** — Column number in  $X$  containing the response variable. (Input)

For the  $i$ -th right-censored observation,  $X(i, IRT)$  contains the right-censoring time. Otherwise,  $X(i, IRT)$  contains the failure time. (See **ICEN**.)

**IFRQ** — Column number in  $X$  containing the frequency of response for this observation. (Input)

If **IFRQ** = 0, a response frequency of 1 for each observation is assumed.

**ICEN** — Column number in  $X$  containing the censoring code for this observation. (Input)

If **ICEN** = 0, a censoring code of 0 is assumed. Valid censoring codes are:

Code	Meaning
------	---------

0	Exact failure at $X(i, IRT)$ .
---	--------------------------------

1	Right censored. The response is greater than $X(i, IRT)$ .
---	--

If  $X(i, ICEN)$  is not 0 or 1, then the  $i$ -th observation is omitted from the analysis.

**IGRP** — Column number in  $X$  containing the stratum number for this observation. (Input)

If **IGRP** = 0, the data is assumed to be from one stratum. Otherwise, column **IGRP** of  $X$  contains a unique value for each stratum in the data. Kaplan-Meier estimates are computed within each stratum.

**ISRT** — Sorting option. (Input)

If **ISRT** = 1, column **IRT** of  $X$  is assumed to be sorted in ascending order within each stratum. Otherwise, a detached sort will be performed by **KAPMR**. If sorting is performed by **KAPMR**, all censored individuals are assumed to follow tied failures.

**SPROB** — NOBS by 2 matrix. (Output)

**SPROB**( $i, 1$ ) contains the estimated survival probability at time  $X(i, IRT)$  in the  $i$ -th observation's stratum, while **SPROB**( $i, 2$ ) contains Greenwood's estimate of

the standard deviation of this estimated probability. If the  $i$ -th observation contains censor codes out of range or if a variable is missing, then the corresponding elements of `SPROB` are set to missing (NaN, not a number). Similarly, if an element in `SPROB` is not defined, then it is set to missing.

***LDSPRO*** — Leading dimension of `SPROB` exactly as specified in the dimension statement in the calling program. (Input)

***NRMISS*** — Number of rows of data in `X` that contain any missing values. (Output)

## Comments

1. Automatic workspace usage is

`KAPMR`  $4 * \text{NOBS} + \text{NCOL} + 3 * m$ , or

`DKAPMR`  $4 * \text{NOBS} + \text{NCOL} + 5 * m$ , where  $m = \max(\text{NOBS}, \text{NCOL})$ .

Workspace may be explicitly provided, if desired, by use of `K2PMR`/`DK2PMR`. The reference is

```
CALL K2PMR (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN,
           IGRP, ISRT, SPROB, LDSPRO, NRMISS,
           IGP, IPERM, INDDR, IWK, WK, IPER)
```

The additional arguments are as follows:

***IGP*** — Work vector of length `NOBS`.

***IPERM*** — Work vector of length `NOBS + NCOL`.

***INDDR*** — Work vector of length `NOBS`.

***IWK*** — Work vector of length  $\max(\text{NOBS}, \text{NCOL})$ .

***WK*** — Work vector of length  $2 * \max(\text{NOBS}, \text{NCOL})$ .

***IPER*** — Work vector of length `NOBS`.

2. Missing values may occur in any of the columns of `X`. Any row of `X` that contains missing values in the `IRT`, `ICEN`, or `IFRQ` columns (when the `ICEN` and `IFRQ` columns are present) is omitted from the analysis. Missing values in the `IGRP` column, if present, are classified into an additional “missing” group.

## Algorithm

Routine `KAPMR` computes Kaplan-Meier (or product-limit) estimates of survival probabilities for a sample of failure times that possibly contain right censoring. A survival probability  $S(t)$  is defined as  $1 - F(t)$ , where  $F(t)$  is the cumulative distribution function of the failure times ( $t$ ). Greenwood’s estimate of the standard errors of the survival probability estimates are also computed. (See Kalbfleisch and Prentice, 1980, pages 13 and 14.)

Let  $(t_i, \delta_i)$ , for  $i = 1, \dots, n$  denote the failure/censoring times and the censoring codes for the  $n$  observations in a single sample. Here,  $t_i = x(i, \text{IRT})$  is a failure time if  $\delta_i$  is 0, where  $\delta_i = x(i, \text{ICEN})$ . Also,  $t_i$  is a censoring time if  $\delta_i$  is 1. Rows in  $X$  containing values other than 0 or 1 for  $\delta_i$  are ignored. Let the number of observations in the sample that have not failed by time  $s_{(i)}$  be denoted by  $n_{(i)}$ , where  $s_{(i)}$  is an ordered (from smallest to largest) listing of the distinct failure times (censoring times are omitted). Then the Kaplan-Meier estimate of the survival probabilities is a step function, which in the interval from  $s_{(i)}$  to  $s_{(i+1)}$  (including the lower endpoint) is given by

$$\hat{S}(t) = \prod_{j=1}^i \left( \frac{n_{(j)} - d_{(j)}}{n_{(j)}} \right)$$

where  $d_{(j)}$  denotes the number of failures occurring at time  $s_{(j)}$ . Note that one row of  $X$  may correspond to more than one failed (or censored) observation when the frequency option is in effect ( $\text{IFRQ}$  is not zero). The Kaplan-Meier estimate of the survival probability prior to time  $s_{(1)}$  is 1.0, while the Kaplan-Meier estimate of the survival probability after the last failure time is not defined.

Greenwood's estimate of the variance of

$$\hat{S}(t)$$

in the interval from  $s_{(i)}$  to  $s_{(i+1)}$  is given as

$$\text{est. var}(\hat{S}(t)) = \hat{S}^2(t) \sum_{j=1}^i \frac{d_{(j)}}{n_{(j)}(n_{(j)} - d_{(j)})}$$

Routine `KAPMR` computes the single sample estimates of the survival probabilities for all samples of data included in  $X$  during a single call. This is accomplished through the `IGRP` column of  $X$ , which if present, must contain a distinct code for each sample of observations. If `IGRP` = 0, there is no grouping column, and all observations are assumed to be from the same sample.

When failures and right-censored observations are tied and the data are to be sorted by `KAPMR` (`ISRT` is not 1), `KAPMR` assumes that the time of censoring for the tied-censored observations is immediately after the tied failure (within the same sample). When the `ISRT` = 1 option is in effect, the data are assumed to be sorted from smallest to largest according to column `IRT` of  $X$  within each stratum. Furthermore, a small increment of time is assumed (theoretically) to elapse between the failed and censored observations that are tied (in the same sample). Thus, when the `ISRT` = 1 option is in effect, the user must sort all of the data in  $X$  from smallest to largest according to column `IRT` (and column `IGRP`, if present). By appropriate sorting of the observations, the user can handle censored and failed observations that are tied in any manner desired.

## Example

The following example is taken from Kalbfleisch and Prentice (1980, page 1). The first column in  $X$  contains the death/censoring times for rats suffering from vaginal cancer. The second column contains information as to which of two forms of treatment were provided, while the third column contains the censoring code. Finally, the fourth column contains the frequency of each observation. The product-limit estimates of the survival probabilities are computed for both groups with one call to KAPMR. In this example, the output in SPROB has been equivalenced with columns 5 and 6 of  $X$  so that the input and output matrices could be printed together. Routine KAPMR could have been called with the ISRT = 1 option in effect if the censored observations had been sorted with respect to the failure time variable.

```

INTEGER      ICEN, IFRQ, IGRP, IRT, ISRT, LDSPRO, LDX, NCOL, NOBS
PARAMETER   (ICEN=3, IFRQ=4, IGRP=2, IRT=1, ISRT=0, LDSPRO=33,
&           LDX=33, NCOL=6, NOBS=33)
C
INTEGER      NOUT, NRMIS
REAL         SPROB(LDSPRO,2), X(LDX,NCOL)
CHARACTER   XLABEL(7)*6, YLABEL(1)*6
EXTERNAL    KAPMR, UMACH, WRRRL
C
EQUIVALENCE (X(1,5), SPROB)
C
DATA XLABEL/'OBS', 'TIME', 'GROUP', 'CENSOR', 'FREQ', 'S-HAT',
&        'SE'/
DATA YLABEL/'NUMBER'/
DATA X/143, 164, 188, 190, 192, 206, 209, 213, 216, 220, 227,
&      230, 234, 246, 265, 304, 216, 244, 142, 156, 163, 198, 205,
&      232, 233, 239, 240, 261, 280, 296, 323, 204, 344, 18*5,
&      15*7, 16*0, 2*1, 13*0, 4*1, 2, 20*1, 2, 4, 3*1, 2*2, 3*1,
&      66*0/
C
CALL KAPMR (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN, IGRP, ISRT,
&          SPROB, LDSPRO, NRMIS)
C
CALL WRRRL ('X/SPROB', NOBS, 6, X, LDSPRO, 0, '(W10.6)', YLABEL,
&          XLABEL)
CALL UMACH (2, NOUT)
WRITE (NOUT, '(//'' NRMIS = '' , I5)') NRMIS
END

```

## Output

OBS	TIME	GROUP	X/SPROB CENSOR	FREQ	S-HAT	SE
1	143.000	5.000	0.000	1.000	0.947	0.051
2	164.000	5.000	0.000	1.000	0.895	0.070
3	188.000	5.000	0.000	2.000	0.789	0.094
4	190.000	5.000	0.000	1.000	0.737	0.101
5	192.000	5.000	0.000	1.000	0.684	0.107
6	206.000	5.000	0.000	1.000	0.632	0.111
7	209.000	5.000	0.000	1.000	0.579	0.113
8	213.000	5.000	0.000	1.000	0.526	0.115
9	216.000	5.000	0.000	1.000	0.474	0.115
10	220.000	5.000	0.000	1.000	0.414	0.115

11	227.000	5.000	0.000	1.000	0.355	0.112
12	230.000	5.000	0.000	1.000	0.296	0.108
13	234.000	5.000	0.000	1.000	0.237	0.101
14	246.000	5.000	0.000	1.000	0.158	0.093
15	265.000	5.000	0.000	1.000	0.079	0.073
16	304.000	5.000	0.000	1.000	0.000	NaN
17	216.000	5.000	1.000	1.000	0.474	0.115
18	244.000	5.000	1.000	1.000	0.237	0.101
19	142.000	7.000	0.000	1.000	0.952	0.046
20	156.000	7.000	0.000	1.000	0.905	0.064
21	163.000	7.000	0.000	1.000	0.857	0.076
22	198.000	7.000	0.000	1.000	0.810	0.086
23	205.000	7.000	0.000	1.000	0.759	0.094
24	232.000	7.000	0.000	2.000	0.658	0.105
25	233.000	7.000	0.000	4.000	0.455	0.111
26	239.000	7.000	0.000	1.000	0.405	0.110
27	240.000	7.000	0.000	1.000	0.354	0.107
28	261.000	7.000	0.000	1.000	0.304	0.103
29	280.000	7.000	0.000	2.000	0.202	0.090
30	296.000	7.000	0.000	2.000	0.101	0.068
31	323.000	7.000	0.000	1.000	0.051	0.049
32	204.000	7.000	1.000	1.000	0.810	0.086
33	344.000	7.000	1.000	1.000	NaN	NaN

NRMISS = 0

---

## KTBLE/DKTBLE (Single/Double precision)

Print Kaplan-Meier estimates of survival probabilities in stratified samples.

### Usage

```
CALL KTBLE (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN, IGRP,
            ISRT, SPROB, LDSPRO)
```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in  $X$ . (Input)

**$X$**  — NOBS by NCOL matrix containing the data. (Input)

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

**IRT** — Column number of  $X$  containing the response variable. (Input)

For the  $i$ -th right-censored observation,  $X(i, IRT)$  contains the right-censoring time. Otherwise,  $X(i, IRT)$  contains the failure time. See argument ICEN.

**IFRQ** — Frequency option. (Input)

IFRQ = 0 means that all frequencies are 1.0. For positive IFRQ, column number IFRQ of  $X$  contains the frequencies.

**ICEN** — Column number of  $X$  containing the censoring code for this observation. (Input)

If  $ICEN = 0$ , a censoring code of 0 is assumed. Valid censoring codes are:

**Code    Meaning**

0        Exact failure at  $X(i, IRT)$ .

1        Right censored. The response is greater than  $X(i, IRT)$ .

If  $X(i, ICEN)$  is not zero or one, then the  $i$ -th observation is omitted from the analysis.

**IGRP** — Column number of  $X$  containing the stratum number for this observation. (Input)

If  $IGRP = 0$ , the data are assumed to be from one stratum. Otherwise, column  $IGRP$  of  $X$  contains a unique value for each stratum in the data. Kaplan-Meier estimates are computed within each stratum.

**ISRT** — Sorting option. (Input)

If  $ISRT = 1$ , column  $IRT$  of  $X$  is assumed to be sorted in ascending order within each stratum. Otherwise, a detached sort will be performed by  $KTBLE$ . If sorting is performed by  $KTBLE$ , all censored observations are assumed to follow failing observations with the same response time in  $X(i, IRT)$ .

**SPROB** — NOBS by 2 matrix. (Input)

$SPROB(i, 1)$  contains the estimated survival probability at time  $X(i, IRT)$  in the  $i$ -th observation's stratum, while  $SPROB(i, 2)$  contains Greenwood's estimate of the standard deviation of this estimated probability.  $SPROB$  will usually be computed by routine  $KAPMR$  (page 938). It may contain missing values after the last failed observation in each group.

**LDSPRO** — Leading dimension of  $SPROB$  exactly as specified in the dimension statement in the calling program. (Input)

**Comments**

1. Automatic workspace usage is

$KTBLE$   $4 * NOBS + 3 * \max(NOBS, NCOL)$  units, or  
 $DKTBLE$   $6 * NOBS + 5 * \max(NOBS, NCOL)$  units.

Workspace may be explicitly provided, if desired, by use of  $K2BLE/DK2BLE$ . The reference is

```
CALL K2BLE (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN,  
           IGRP, ISRT, SPROB, LDSPRO, ALGL, IPERM,  
           INDDR, WK, WK1, IWK)
```

The additional arguments are as follows:

**ALGL** — Work vector of length  $NOBS$  that contains the log likelihoods of the Kaplan-Meier estimates. If the number of groups is known to be  $m$  or less, then  $ALGL$  can be of length  $m$ .

**IPERM** — Work vector of length  $NOBS$ .

**INDDR** — Work vector of length NOBS.

**WK** — Work vector of length NOBS.

**WKI** — Work vector of length  $2 * \max(\text{NOBS}, \text{NCOL})$ .

**IWK** — Work vector of length  $\max(\text{NOBS}, \text{NCOL})$ .

2. Informational errors

Type	Code	
4	1	An invalid value for <code>SPROB</code> has been detected. The estimated survival probability must be between zero and one, inclusive, and nonincreasing with failure time within each group.
4	2	A negative frequency has been detected.
4	3	A missing value for <code>SPROB</code> has been detected but later failures occur. Missing values are not allowed prior to the last failed observation.

3. Missing values may occur in any of the columns of `X`. Any row of `X` that contains missing values in the `IRT`, `ICEN`, or `IFRQ` columns (when the `ICEN` and `IFRQ` columns are present) is omitted from the analysis. Missing values in the `IGRP` column, if present, are classified into an additional “missing” group.

### Algorithm

Routine `KTBLE` prints life tables based upon the Kaplan-Meier estimates of the survival probabilities (see routine `KAPMR`, page 938). One table for each stratum is printed. In addition to the survival probabilities at each failure point, the following is also printed: the number of individuals remaining at risk, Greenwood’s estimate of the standard errors for the survival probabilities, and the Kaplan-Meier log-likelihood. The Kaplan-Meier log-likelihood is computed as:

$$\ell = \sum_j d_{(j)} \ln d_{(j)} + (n_{(j)} - d_{(j)}) \ln(n_{(j)} - d_{(j)}) - n_{(j)} \ln n_{(j)}$$

where the sum is with respect to the distinct failure times  $s_{(j)}$ ,  $d_{(j)}$  is the number of failures occurring at time  $s_{(j)}$ , and  $n_{(j)}$  is the number of observations that had not yet failed immediately prior to  $s_{(j)}$ . Note that sorting is performed by both `KAPMR` (page 938), and by routine `KTBLE`. The user may sort the data to be increasing in failure time and then use the `ISRT = 1` option to avoid this double sorting.

### Example

This example illustrates the typical use of `KTBLE`. First, routine `KAPMR` (page 938) is used to compute the survival probabilities. This is followed by a call to `KTBLE` that performs the printing. The input data is given as:

143, 164, 188(2), 190, 192, 206, 209, 213, 216, 220, 227, 230, 234, 246, 265, 304, 216*, 244*, 142, 156, 163, 198, 205, 232(2), 233(4), 239, 240, 261, 280(2), 296(2), 323, 204 *, 344*
---

where items marked with an \* are right censored; and the frequency of each failure time, if different from 1, is given in parenthesis.

```

INTEGER      ICEN, IFRQ, IGRP, IRT, ISRT, LDSPRO, LDX, NCOL, NOBS
PARAMETER   (ICEN=3, IFRQ=4, IGRP=2, IRT=1, ISRT=0, LDSPRO=33,
&           LDX=33, NCOL=4, NOBS=33)
C
INTEGER      NRMISS
REAL         SPROB(LDSPRO,2), X(LDX,NCOL)
EXTERNAL    KAPMR, KTBLE
C
DATA X/143, 164, 188, 190, 192, 206, 209, 213, 216, 220, 227,
&          230, 234, 246, 265, 304, 216, 244, 142, 156, 163, 198, 205,
&          232, 233, 239, 240, 261, 280, 296, 323, 204, 344, 18*5,
&          15*7, 16*0, 2*1, 13*0, 4*1, 2, 20*1, 2, 4, 3*1, 2*2, 3*1/
C
CALL KAPMR (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN, IGRP, ISRT,
&          SPROB, LDSPRO, NRMISS)
C
CALL KTBLE (NOBS, NCOL, X, LDX, IRT, IFRQ, ICEN, IGRP, ISRT,
&          SPROB, LDSPRO)
END

```

### Output

Kaplan Meier Survival Probabilities  
For Group Value = 5.00000

Number at risk	Number Failing	Time	Survival Probability	Estimated Std. Error
19	1	143	0.94737	0.05123
18	1	164	0.89474	0.07041
17	2	188	0.78947	0.09353
15	1	190	0.73684	0.10102
14	1	192	0.68421	0.10664
13	1	206	0.63158	0.11066
12	1	209	0.57895	0.11327
11	1	213	0.52632	0.11455
10	1	216	0.47368	0.11455
8	1	220	0.41447	0.11452
7	1	227	0.35526	0.11243
6	1	230	0.29605	0.10816
5	1	234	0.23684	0.10145
3	1	246	0.15789	0.09343
2	1	265	0.07895	0.07279
1	1	304	0.00000	NaN

Total number in group = 19  
Total number failing = 17  
Product Limit Likelihood = -49.1692



Kaplan Meier Survival Probabilities  
For Group Value = 7.00000

Number at risk	Number Failing	Time	Survival Probability	Estimated Std. Error
21	1	142	0.95238	0.04647
20	1	156	0.90476	0.06406
19	1	163	0.85714	0.07636
18	1	198	0.80952	0.08569
16	1	205	0.75893	0.09409
15	2	232	0.65774	0.10529
13	4	233	0.45536	0.11137
9	1	239	0.40476	0.10989
8	1	240	0.35417	0.10717
7	1	261	0.30357	0.10311
6	2	280	0.20238	0.09021
4	2	296	0.10119	0.06778
2	1	323	0.05060	0.04928

Total number in group = 21  
Total number failing = 19  
Product Limit Likelihood = -50.4277

---

## TRNBL/DTRNBL (Single/Double precision)

Compute Turnbull's generalized Kaplan-Meier estimates of survival probabilities in samples with interval censoring.

### Usage

```
CALL TRNBL (NOBS, NCOL, X, LDX, ILT, IRT, IFRQ, ICEN,
            MAXIT, EPS, IPRINT, NINTVL, SPROB, LDSPRO,
            ALGL, NRMIS)
```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in *X*. (Input)

**X** — NOBS by NCOL matrix containing the data. (Input)

**LDX** — Leading dimension of *X* exactly as specified in the dimension statement in the calling program. (Input)

**ILT** — For interval-censored and left-censored observations, the column number in *X* that contains the upper endpoint of the failure interval. (Input)  
See argument ICEN. If ILT = 0, left-censored and interval-censored observations cannot be input.

**IRT** — For interval-censored and right-censored observations, the column number in *X* that contains the lower endpoint of the failure interval. (Input)  
See argument ICEN. IRT must not be zero.

**IFRQ** — Frequency option. (Input)

If  $IFRQ = 0$ , a response frequency of 1 for each observation is assumed. For positive  $IFRQ$ , column number  $IFRQ$  contains the frequency of response for each observation.

**ICEN** — Censoring code option. (Input)

If  $ICEN = 0$ , a censoring code of 0 is assumed. For positive  $ICEN$ , column number  $ICEN$  contains the censoring code for each observation. Valid censoring codes are:

**Code    Meaning**

- 0      Exact failure at  $X(i, IRT)$ .
- 1      Right censored. The response is greater than  $X(i, IRT)$ .
- 2      Left censored. The response is less than or equal to  $X(i, ILT)$ .
- 3      Interval censored. The response is greater than  $X(i, IRT)$ , but less than or equal to  $X(i, ILT)$ .

**MAXIT** — Maximum number of iterations. (Input)

**EPS** — Convergence criterion. (Input)

Convergence is assumed when the relative change in the log-likelihood from one iteration to the next is less than  $EPS$ .  $EPS = 0.00001$  is typical.

**IPRINT** — Printing option. (Input)

$IPRINT = 0$  means that no printing is performed.  $IPRINT = 1$  means that printing is performed.

**NINTVL** — Number of failure intervals found. (Output)

**SPROB** —  $NINTVL$  by 4 matrix. (Output)

**Col.    Description**

- 1      Lower endpoint of the failure interval
- 2      Upper endpoint of the failure interval
- 3      Estimated change in the survival probability density within the failure interval
- 4      Estimate of the survival probability for the interval

The estimated survival probability is a constant equal to  $SPROB(i, 4)$  from  $SPROB(i, 2)$  to  $SPROB(i + 1, 1)$ . The estimated survival probability is 1 prior to  $SPROB(1, 1)$ . The estimated survival probability is undefined in the interval  $SPROB(i, 1)$  to  $SPROB(i, 2)$ . If the  $NINTVL$ -th interval is from  $SPROB(NINTVL, 1)$  to infinity, then  $SPROB(NINTVL, 2)$  is set to positive machine infinity.

**LDSPRO** — Leading dimension of  $SPROB$  exactly as specified in the dimension statement in the calling program. (Input)

If  $LDSPRO$  is less than  $NINTVL$ , only the first  $LDSPRO$  intervals are returned in  $SPROB$ .

**ALGL** — Optimized log-likelihood for the input data. (Output)

**NRMISS** — Number of rows of data in  $x$  that contain missing values. (Output)  
 Any row of  $x$  that contains missing values in the  $ILT$ ,  $IRT$ ,  $ICEN$ , or  $IFRQ$  columns (when the  $ILT$ ,  $ICEN$  or  $IFRQ$  is positive) is omitted from the analysis.

### Comments

- Automatic workspace usage is

$TRNBL$   $9 * NOBS + 3 * \max(NOBS, 7)$  units, or  
 $DTRNBL$   $16 * NOBS + 5 * \max(NOBS, 7)$  units.

Workspace may be explicitly provided, if desired, by use of  $T2NBL/DT2NBL$ . The reference is

```
CALL T2NBL (NOBS, NCOL, X, LDX, ILT, IRT, IFRQ,
           ICEN, MAXIT, EPS, IPRINT, NINTVL, SPROB,
           LDSPRO, ALGL, NRMISS, WK, IPERM, INDDR,
           WWK, IWK)
```

The additional arguments are as follows:

**WK** — Work vector of length  $7 * NOBS$ .

**IPERM** — Work vector of length  $NOBS$ .

**INDDR** — Work vector of length  $NOBS$ .

**WWK** — Work vector of length  $2 * \max(NOBS, 7)$ .

**IWK** — Work vector of length  $\max(NOBS, 7)$ .

- Informational errors

Type	Code	
3	3	The maximum number of iterations was exceeded. Convergence is assumed.
4	1	There are no valid observations.
4	2	There are no finite failure intervals present in the data.

### Algorithm

Routine  $TRNBL$  computes nonparametric maximum likelihood estimates of a survival distribution based upon a random sample of data containing exact failure, right-censored, leftcensored (interval censored with a left endpoint of zero), or interval-censored observations. The computational method of Turnbull (1976) is used in computing the probability estimates. The model used is also discussed by Peto (1973).

Routine  $TRNBL$  begins by finding a set of regions or “failure intervals” (to distinguish them from “observation failure intervals”) on the positive real axis in which a change in the survival probability occurs. The survival probability is constant outside of these regions, and undefined within them. Each region (failure interval) is composed of a single left and a single right endpoint obtained from the left and right endpoints of the observation failure intervals

(for exact failure times, the left and right endpoints are equal). The regions are defined by the fact that no observation interval endpoints are allowed within a region, except at its endpoints. Note that the endpoints of the intervals need not correspond to a single observation. Regions defined by endpoints from two distinct observations are often obtained.

Let  $p_i$ ,  $i = 1, \dots, \text{NINTVL}$  denote the change in the survival probability within the  $i$ -th region, and let the region be denoted by  $c_i$ . Let  $n = \text{NOBS}$  and suppose that the observation failure interval for observation  $j$  is denoted by  $I_j$ . The EM (expectation, maximization) algorithm of Dempster, Laird and Rubin (1977) is used to find the optimal

$$\hat{p}_i \text{'s}$$

The algorithm is defined as follows:

For given

$$\hat{p}_i$$

compute the expected contribution of the  $j$ -th observation to the  $i$ -th change interval as

$$\hat{\mu}_{ij} = \frac{f_j \hat{p}_i \delta_{ij}}{\sum_j f_j \hat{p}_i \delta_{ij}}$$

where  $\delta_{ij} = 1$  if  $c_i \subseteq I_j$  and  $\delta_{ij} = 0$  otherwise, and  $f_j$  is the observation frequency.

For given expectations

$$\hat{\mu}_{ij}$$

compute the new probability estimate as

$$\hat{p}_i = \frac{\sum_j \hat{\mu}_{ij}}{\sum_j f_j}$$

Iterate in this manner until convergence. Convergence is assumed when the relative change in the log-likelihood

$$(\ell = \sum_j f_j \ln(\sum_i \delta_{ij} \hat{p}_i))$$

is small (less than `EPS`). Because the algorithm is slow to converge, 5 expectation-maximization cycles are considered to be one iteration of the algorithm. The initial estimate for all the

$$\hat{p}_i \text{'s}$$

is taken to be one divided by the number of regions (failure intervals).

### Example

The following example contains exact failure, right-, left-, and interval-censored observations. The 20 observations yield 15 change intervals. The last interval is from 192 to  $\infty$ , and corresponds to a right-censored observation. When the last interval is infinite, as is the case here, the second column of *SPROB* contains  $+\infty$  in the *NINTVL*-th position. Left- or right-censored observations input in *X* are arbitrarily assigned the value 0.0 for the non-specified endpoint.

```

INTEGER      ICEN, IFRQ, ILT, IPRINT, IRT, LDSPRO, LDX, MAXIT,
&           NCOL, NOBS
REAL        EPS
PARAMETER   (EPS=0.00001, ICEN=4, IFRQ=3, ILT=1, IPRINT=1, IRT=2,
&           LDSPRO=20, LDX=20, MAXIT=30, NCOL=4, NOBS=20)
C
INTEGER      NINTVL, NRMISS
REAL        ALGL, SPROB(LDSPRO,4), X(LDX,NCOL)
EXTERNAL    TRNBL, WRRRN
C
DATA X/0.9, 1.9, 2.5, 3.5, 6.3, 7.1, 18., 25.1, 25.3, 30.3, 45.9,
&      63.5, 70.1, 73.0, 93.0, 94.4, 96.0, 0.0, 191.4, 0.0, 0.9,
&      0.0, 0.0, 0.0, 6.3, 1.9, 1.8, 25.1, 9.5, 30.3, 45.9,
&      60.7, 70.1, 71.0, 74.0, 94.4, 96.0, 96.0, 191.4, 192.0,
&      17*1.0, 5.0, 1.0, 1.0, 0.0, 2.0, 2.0, 2.0, 0.0, 3.0, 3.0,
&      0.0, 3.0, 0.0, 0.0, 3.0, 0.0, 3.0, 3.0, 0.0, 0.0, 1.0, 0.0,
&      1.0/
C
CALL WRRRN ('X', NOBS, NCOL, X, LDX, 0)
C
CALL TRNBL (NOBS, NCOL, X, LDX, ILT, IRT, IFRQ, ICEN, MAXIT,
&          EPS, IPRINT, NINTVL, SPROB, LDSPRO, ALGL, NRMISS)
C
END

```

### Output

	X			
	1	2	3	4
1	0.9	0.9	1.0	0.0
2	1.9	0.0	1.0	2.0
3	2.5	0.0	1.0	2.0
4	3.5	0.0	1.0	2.0
5	6.3	6.3	1.0	0.0
6	7.1	1.9	1.0	3.0
7	18.0	1.8	1.0	3.0
8	25.1	25.1	1.0	0.0
9	25.3	9.5	1.0	3.0
10	30.3	30.3	1.0	0.0
11	45.9	45.9	1.0	0.0
12	63.5	60.7	1.0	3.0
13	70.1	70.1	1.0	0.0
14	73.0	71.0	1.0	3.0
15	93.0	74.0	1.0	3.0
16	94.4	94.4	1.0	0.0
17	96.0	96.0	1.0	0.0
18	0.0	96.0	5.0	1.0
19	191.4	191.4	1.0	0.0
20	0.0	192.0	1.0	1.0

Iteration	Log-Likelihood	Relative convergence
0	-54.94	.....
1	-52.14	0.5367E-01
2	-52.09	0.8407E-03
3	-52.09	0.1372E-03
4	-52.09	0.2476E-04
5	-52.08	0.4614E-05

SPROB

Interval	Lower Endpoint	Upper Endpoint	Interval Probability	Survival Probability
1	0.9000	0.9000	0.0972	0.9028
2	1.9000	1.9000	0.1215	0.7813
3	6.3000	6.3000	0.0729	0.7083
4	9.5000	18.0000	0.0000	0.7083
5	25.1000	25.1000	0.0833	0.6250
6	30.3000	30.3000	0.0417	0.5833
7	45.9000	45.9000	0.0417	0.5417
8	60.7000	63.5000	0.0417	0.5000
9	70.1000	70.1000	0.0417	0.4583
10	71.0000	73.0000	0.0417	0.4167
11	74.0000	93.0000	0.0417	0.3750
12	94.4000	94.4000	0.0417	0.3333
13	96.0000	96.0000	0.1111	0.2222
14	191.4000	191.4000	0.1111	0.1111
15	192.0000	Inf	0.1111	0.0000

---

## PHGLM/DPHGLM (Single/Double precision)

Analyze time event data via the proportional hazards model.

### Usage

```
CALL PHGLM (NOBS, NCOL, X, LDX, IRT, IFRQ, IFIX, ICEN,
            ISTRAT, MAXIT, EPS, RATIO, NCLVAR, INDCL, NEF,
            NVEF, INDEF, INIT, ITIE, IPRINT, MAXCL, NCLVAL,
            CLVAL, NCOEF, COEF, LDcoef, ALGL, COV, LDcov,
            XMEAN, CASE, LDCASE, GR, IGRP, NRMISS)
```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in *X*. (Input)

**X** — NOBS by NCOL matrix containing the data. (Input)

When **ITIE** = 1, the observations in *X* must be grouped by stratum and sorted from largest to smallest failure time within each stratum, with the strata separated.

**LDX** — Leading dimension of *X* exactly as specified in the dimension statement in the calling program. (Input)

**IRT** — Column number in *X* containing the response variable. (Input)

For point observations, *X*(*i*, **IRT**) contains the time of the *i*-th event. For right-

censored observations,  $x(i, \text{IRT})$  contains the right-censoring time. Note that because PHGLM only uses the order of the events, negative “times” are allowed.

**IFRQ** — Column number in  $x$  containing the frequency of response for each observation. (Input)

If  $\text{IFRQ} = 0$ , a response frequency of 1 for each observation is assumed.

**IFIX** — Column number in  $x$  containing a constant to be added to the linear response. (Input)

The linear response is taken to be

$$w_i + z_i \hat{\beta}$$

where  $w_i$  is the observation constant,  $z_i$  is the observation design row vector, and

$$\hat{\beta}$$

is the vector of estimated parameters. The “fixed” constant allows one to test hypotheses about parameters via the log-likelihoods. If  $\text{IFIX} = 0$ , the fixed parameter is assumed to be 0.

**ICEN** — Column number in  $x$  containing the censoring code for each observation. (Input)

If  $\text{ICEN} = 0$  a censoring code of 0 is assumed for all observations.

$x(i, \text{ICEN})$	Censoring
0	Point observation at $x(i, \text{IRT})$ .
1	Right censored. The response is greater than $x(i, \text{IRT})$ .

**ISTRAT** — Column number in  $x$  containing the stratification variable. (Input)

If  $\text{ISTRAT} = 0$ , all observations are considered to be in one stratum. Otherwise, column  $\text{ISTRAT}$  in  $x$  contains a unique number for each stratum. The risk set for an observation is determined by the its stratum.

**MAXIT** — Maximum number of iterations. (Input)

$\text{MAXIT} = 30$  will usually be sufficient. Use  $\text{MAXIT} = 0$  to compute the Hessian and gradient, stored in  $\text{COV}$  and  $\text{GR}$ , at the initial estimates. When  $\text{MAXIT} = 0$ ,  $\text{INIT}$  must be 1.

**EPS** — Convergence criterion. (Input)

Convergence is assumed when the relative change in  $\text{ALGL}$  from one iteration to the next is less than  $\text{EPS}$ . If  $\text{EPS}$  is zero,  $\text{EPS} = 0.0001$  is assumed.

**RATIO** — Ratio at which a stratum is split into two strata. (Input)

Let

$$r_k = \exp(z_k \hat{\beta} + w_k)$$

be the observation proportionality constant, where  $z_k$  is the design row vector for the  $k$ -th observation and  $w_k$  is the optional fixed parameter specified by  $x(k, \text{IFIX})$ . Let  $r_{\min}$  be the minimum value  $r_k$  in a stratum, where, for failed observations, the minimum is over all times less than or equal to the time of

occurrence of the  $k$ -th observation. Let  $r_{\max}$  be the maximum value of  $r_k$  for the remaining observations in the group. Then, if  $r_{\min} > \text{RATIO } r_{\max}$ , the observations in the group are divided into two groups at  $k$ .  $\text{RATIO} = 1000$  is usually a good value. Set  $\text{RATIO} = -1.0$  if no division into strata is to be made.

**NCLVAR** — Number of classification variables. (Input)

Dummy variables are generated for classification variables using the  $\text{IDUMMY} = 2$  option of IMSL routine  $\text{GRGLM}$  (page 210). See Comment 3.

**INDCL** — Index vector of length  $\text{NCLVAR}$  containing the column numbers of  $\mathbf{X}$  that are the classification variables. (Input, if  $\text{NCLVAR}$  is positive, not used otherwise)

If  $\text{NCLVAR}$  is 0,  $\text{INDCL}$  is not referenced and can be dimensioned of length 1 in the calling program.

**NEF** — Number of effects in the model. (Input)

In addition to effects involving classification variables, simple covariates and the product of simple covariates are also considered effects.

**NVEF** — Vector of length  $\text{NEF}$  containing the number of variables associated with each effect in the model. (Input)

**INDEF** — Index vector of length  $\text{NVEF}(1) + \dots + \text{NVEF}(\text{NEF})$  containing the column numbers of  $\mathbf{X}$  associated with each effect. (Input)

The first  $\text{NVEF}(1)$  elements of  $\text{INDEF}$  contain the column numbers of  $\mathbf{X}$  for the variables in the first effect. The next  $\text{NVEF}(2)$  elements in  $\text{INDEF}$  contain the column numbers for the second effect, etc.

**INIT** — Initialization option. (Input)

If  $\text{INIT} = 1$ , then the  $\text{NCOEF}$  elements of column 1 of  $\text{COEF}$  contain the initial estimates on input to  $\text{PHGLM}$ . For  $\text{INIT} = 0$ , all initial estimates are taken to be 0.

**ITIE** — Option parameter containing the method to be used for handling ties. (Input)

**ITIE Method**

- |   |  |
|---|--|
| 0 | Breslow's approximate method   |
| 1 | Failures are assumed to occur in the same order as the observations input in $\mathbf{x}$ . The observations in $\mathbf{x}$ must be sorted from largest to smallest failure time within each stratum, and grouped by stratum. All observations are treated as if their failure/censoring times were distinct when computing the log-likelihood. |

**IPRINT** — Printing option. (Input)

**IPRINT Action**

- |   |  |
|---|--|
| 0 | No printing is performed.  |
| 1 | Printing is performed, but observational statistics are not printed. |
| 2 | All output statistics are printed.                                   |



**MAXCL** — An upper bound on the sum of the number distinct values taken by the classification variables. (Input)

**NCLVAL** — Vector of length NCLVAR containing the number of values taken by each classification variable. (Output, if NCLVAR is positive, not used otherwise) NCLVAL(*i*) is the number of distinct values for the *i*-th classification variable. If NCLVAR is zero, NCLVAL is not used and can be dimensioned of length 1 in the calling program.

**CLVAL** — Vector of length NCLVAL(1) + NCLVAL(2) + ... + NCLVAL(NCLVAR) containing the distinct values of the classification variables. (Output, if NCLVAR is positive, not used otherwise)

The first NCLVAL(1) elements of CLVAL contain the values for the first classification variable, the next NCLVAL(2) elements contain the values for the second classification variable, etc. If NCLVAR is zero, then NCLVAL is not referenced and can be dimensioned of length 1 in the calling program.

**NCOEF** — Number of estimated coefficients in the model. (Output)

**COEF** — NCOEF by 4 matrix containing the parameter estimates and associated statistics. (Output, if INIT = 0; input, if INIT = 1 and MAXIT = 0, input/output, if INIT = 1 and MAXIT > 0)

**Col.     Statistic**

1         Coefficient estimate

$$\hat{\beta}$$

2         Estimated standard deviation of the estimated coefficient.

3         Asymptotic normal score for testing that the coefficient is zero against the two-sided alternative.

4         

-value associated with the normal score in column 3.

When COEF is input, only column 1 needs to be given.

**LDCOEF** — Leading dimension of COEF exactly as specified in the dimension statement in the calling program. (Input)

**ALGL** — The maximized log-likelihood. (Output)

**COV** — NCOEF by NCOEF matrix containing the estimated asymptotic variance-covariance matrix of the parameters. (Output)

For MAXIT = 0, COV is the inverse of the Hessian of the negative of the log-likelihood, computed at the estimates input in COEF.

**LDCOV** — Leading dimension of COV exactly as specified in the dimension statement in the calling program. (Input)

**XMEAN** — Vector of length NCOEF containing the means of the design variables. (Output)

**CASE** — NOBS by 5 matrix containing the case statistics for each observation. (Output if MAXIT > 0; used as working storage otherwise)

**Col.   Statistic**

- 1     Estimated survival probability at the observation time.
- 2     Estimated observation influence or leverage.
- 3     A residual estimate.
- 4     Estimated cumulative baseline hazard rate.
- 5     Observation proportionality constant.

**LDCASE** — Leading dimension of CASE exactly as specified in the dimension statement in the calling program. (Input)

**GR** — Vector of length NCOEF containing the last parameter updates (excluding step halvings). (Output)

For MAXIT = 0, GR contains the inverse of the Hessian times the gradient vector computed at the estimates input in COEF.

**IGRP** — Vector of length NOBS giving the stratum number used for each observation. (Output)

If RATIO is not -1.0, additional "strata" (other than those specified by column ISTRAT of X) may be generated. IGRP also contains a record of the generated strata. See the algorithm section for more detail.

**NRMIS** — Number of rows of data in X that contain missing values in one or more columns IRT, IFRQ, IFIX, ICEN, ISTRAT, INDCL, or INDEF of X. (Output)

**Comments**

1.     Automatic workspace usage is

PHGLM   2 \* NOBS + 3 \* NCOEF + max(NCOEF \* NCOEF, 2) +  
          3 \* MAX(NOBS, NCOL) + 1 units, or

DPHGLM  2 \* NOBS + 5 \* NCOEF + 2 \* max(NCOEF \* NCOEF,  
          2) + 3 \* max(NOB, NCOL) + 2 units.

Workspace may be explicitly provided, if desired, by use of P2GLM/DP2GLM. The reference is

CALL P2GLM (NOBS, NCOL, X, LDX, IRT, IFRQ, IFIX,  
            ICEN, ISTRAT, MAXIT, EPS, RATIO, NCLVAR,  
            INDCL, NEF, NVEF, INDEF, INIT, ITIE,  
            IPRINT, MAXCL, NCLVAL, CLVAL, NCOEF,  
            COEF, LDCEOF, ALGL, COV, LDCEOF, XMEAN,  
            CASE, LDCASE, GR, IGRP, NRMIS, OBS,  
            SMG, SMH, IPTR, IDT, IWK)

The additional arguments are as follows:

**OBS** — Work vector of length NCOEF + 1.

**SMG** — Work vector of length NCOEF.

**SMH** — Work vector of length max(NCOEF \* NCOEF, 2).

**IPTR** — Work vector of length NOBS + NCOEF.

**IDT** — Work vector of length NOBS.

**IWK** — Work vector of length  $3 * \max(\text{NOBS}, \text{NCOL})$

2. Informational errors

Type	Code	
3	1	Too many iterations required. Convergence assumed.
3	2	Too many step halvings. Convergence assumed.
3	3	Additional strata were formed as required because of the detection of infinite parameter estimates.
4	4	The number of distinct values of the classification variables exceeds MAXCL.
4	5	The model specified by NEF, NVEF, and INDEF yields no covariates.
4	6	After eliminating observations with missing values, no valid observations remain.
4	7	After eliminating observations with missing values, only one covariate vector remains.
4	8	The number of distinct values for each classification variable must be greater than one.
4	9	LDCOEFF or LDCOV must be greater or equal to NCOEFF.

3. Dummy variables are generated for the classification variables as follows: An ascending list of all distinct values of the classification variable is obtained and stored in CLVAL. Dummy variables are then generated for each but the last of these distinct values. Each dummy variable is zero unless the classification variable equals the list value corresponding to the dummy variable, in which case, the dummy variable is one. See argument IDUMMY for IDUMMY = 2 in routine GRGLM (page 210) in Chapter 2.
4. The “product” of a classification variable with a covariate yields dummy variables equal to the product of the covariate with each of the dummy variables associated with the classification variable.
5. The “product” of two classification variables yields dummy variables in the usual manner. Each dummy variable associated with the first classification variable multiplies each dummy variable associated with the second classification variable. The resulting dummy variables are such that the index of the second classification variable varies fastest.

### Algorithm

Routine PHGLM computes parameter estimates and other statistics in Proportional Hazards Generalized Linear Models. These models were first proposed by Cox (1972). Two methods for handling ties are allowed in PHGLM. Time-dependent covariates are not allowed. The user is referred to Cox and Oakes (1984), Kalbfleisch and Prentice (1980), Elandt-Johnson and Johnson (1980), Lee (1980), or Lawless (1982), among other texts, for a thorough discussion of the Cox proportional hazards model.

Let  $\lambda(t, z_i)$  represent the hazard rate at time  $t$  for observation number  $i$  with covariables contained as elements of row vector  $z_i$ . The basic assumption in the proportional hazards model (the proportionality assumption) is that the hazard rate can be written as a product of a time varying function  $\lambda_0(t)$ , which depends only on time, and a function  $f(z_i)$ , which depends only on the covariable values. The function  $f(z_i)$  used in PHGLM is given as  $f(z_i) = \exp(w_i + \beta z_i)$  where  $w_i$  is a fixed constant assigned to the observation, and  $\beta$  is a vector of coefficients to be estimated. With this function one obtains a hazard rate  $\lambda(t, z_i) = \lambda_0(t) \exp(w_i + \beta z_i)$ . The form of  $\lambda_0(t)$  is not important in proportional hazards models.

The constants  $w_i$  may be known theoretically. For example, the hazard rate may be proportional to a known length or area, and the  $w_i$  can then be determined from this known length or area. Alternatively, the  $w_i$  may be used to fix a subset of the coefficients  $\beta$  (say,  $\beta_1$ ) at specified values. When  $w_i$  is used in this way, constants  $w_i = \beta_1 z_{1i}$  are used, while the remaining coefficients in  $\beta$  are free to vary in the optimization algorithm. If user-specified constants are not desired, the user should set `IFIX` to 0 so that  $w_i = 0$  will be used.

With this definition of  $\lambda(t, z_i)$ , the usual partial (or marginal, see Kalbfleisch and Prentice (1980)) likelihood becomes

$$L = \prod_{i=1}^{n_d} \frac{\exp(w_i + \beta z_i)}{\sum_{j \in R(t_i)} \exp(w_j + \beta z_j)}$$

where  $R(t_i)$  denotes the set of indices of observations that have not yet failed at time  $t_i$  (the risk set),  $t_i$  denotes the time of failure for the  $i$ -th observation,  $n_d$  is the total number of observations that fail. Right-censored observations (i.e., observations that are known to have survived to time  $t_i$ , but for which no time of failure is known) are incorporated into the likelihood through the risk set  $R(t_i)$ . Such observations never appear in the numerator of the likelihood. When `ITIE` = 0, all observations that are censored at time  $t_i$  are not included in  $R(t_i)$ , while all observations that fail at time  $t_i$  are included in  $R(t_i)$ .

If it can be assumed that the dependence of the hazard rate upon the covariate values remains the same from stratum to stratum, while the time-dependent term,  $\lambda_0(t)$ , may be different in different strata, then PHGLM allows the incorporation of strata into the likelihood as follows. Let  $k$  index the  $m = \text{NSTRAT}$  strata. Then, the likelihood is given by

$$L_s = \prod_{k=1}^m \left[ \prod_{i=1}^{n_k} \frac{\exp(w_{ki} + \beta z_{ki})}{\sum_{j \in R(t_{ki})} \exp(w_{kj} + \beta z_{kj})} \right]$$

In PHGLM, the log of the likelihood is maximized with respect to the coefficients  $\beta$ . A quasi-Newton algorithm approximating the Hessian via the matrix of sums of squares and cross products of the first partial derivatives is used in the initial

iterations (the “Q-N” method in the output). When the change in the log-likelihood from one iteration to the next is less than  $100 * \text{EPS}$ , Newton-Raphson iteration is used (the “N-R” method). If, during any iteration, the initial step does not lead to an increase in the log-likelihood, then step halving is employed to find a step that will increase the log-likelihood.

Once the maximum likelihood estimates have been computed, PHGLM computes estimates of a probability associated with each failure. Within stratum  $k$ , an estimate of the probability that the  $i$ -th observation fails at time  $t_i$  given the risk set  $R(t_{ki})$  is given by

$$p_{ki} = \frac{\exp(w_{ki} + z_{ki}\beta)}{\sum_{j \in R(t_{ki})} \exp(w_{kj} + z_{kj}\beta)}$$

A diagnostic “influence” or “leverage” statistic is computed for each noncensored observation as:

$$l_{ki} = -g'_{ki} H_s^{-1} g'_{ki}$$

where  $H_s$  is the matrix of second partial derivatives of the log-likelihood, and

$$g'_{ki}$$

is computed as:

$$g'_{ki} = z_{ki} - \frac{z_{ki} \exp(w_{ki} + z_{ki}\beta)}{\sum_{j \in R(t_{ki})} \exp(w_{kj} + z_{kj}\beta)}$$

Influence statistics are not computed for censored observations.

A “residual” is computed for each of the input observations according to methods given in Cox and Oakes (1984, page 108). Residuals are computed as

$$r_{ki} = \exp(w_{ki} + z_{ki}\hat{\beta}) \sum_{j \in R(t_{ki})} \frac{d_{kj}}{\sum_{l \in R(t_{kj})} \exp(w_{kl} + z_{kl}\hat{\beta})}$$

where  $d_{kj}$  is the number of tied failures in group  $k$  at time  $t_{kj}$ . Assuming that the proportional hazards assumption holds, the residuals should approximate a random sample (with censoring) from the unit exponential distribution. By subtracting the expected values, centered residuals can be obtained. (The  $j$ -th expected order statistic from the unit exponential with censoring is given as

$$e_j = \sum_{l \leq j} \frac{1}{h-l+1}$$

where  $h$  is the sample size, and censored observations are not included in the summation.)

An estimate of the cumulative baseline hazard within group  $k$  is given as

$$\hat{H}_{k0}(t_{ik}) = \sum_{t_{kj} \leq t_{ki}} \frac{d_{kj}}{\sum_{l \in R(t_{kj})} \exp(w_{kl} + z_{kl}\hat{\beta})}$$

The observation proportionality constant is computed as

$$\exp(w_{ki} + z_{ki}\hat{\beta})$$

### Programming Notes

1. The covariate vectors  $z_{ki}$  are computed from each row of the input matrix  $\mathbf{X}$  via routine GRGLM (page 210). Thus, class variables are easily incorporated into the  $z_{ki}$ . The reader is referred to the document for GRGLM in the regression chapter for a more detailed discussion. Note that PHGLM calls GRGLM with the option IDUMMY = 2.
2. The average of each of the explanatory variables is subtracted from the variable prior to computing the product  $z_{ki}\hat{\beta}$ . Subtraction of the mean values has no effect on the computed log-likelihood or the estimates since the constant term occurs in both the numerator and denominator of the likelihood. Subtracting the mean values does help to avoid invalid exponentiation in the algorithm and may also speed convergence.
3. Routine PHGLM allows for two methods of handling ties. In the first method (ITIE = 1), the user is allowed to break ties in any manner desired. When this method is used, it is assumed that the user has sorted the rows in  $\mathbf{X}$  from largest to smallest with respect to the failure/censoring times  $\mathbf{X}(i, \text{IRT})$  within each stratum (and across strata), with tied observations (failures or censored) broken in the manner desired. The same effect can be obtained with ITIE = 0 by adding (or subtracting) a small amount from each of the tied observations failure/censoring times  $t_i = \mathbf{X}(i, \text{IRT})$  so as to break the ties in the desired manner.

The second method for handling ties (ITIE = 0) uses an approximation for the tied likelihood proposed by Breslow (1974). The likelihood in Breslow's method is as specified above, with the risk set at time  $t_i$  including all observations that fail at time  $t_i$ , while all observations that are censored at time  $t_i$  are not included. (Tied censored observations are assumed to be censored immediately prior to the time  $t_i$ ).

4. If INIT = 1, then it is assumed that the user has provided initial estimates for the model coefficients  $\beta$  in the first column of the matrix COEF. When initial estimates are provided by the user, care should be taken to ensure that the estimates correspond to the generated covariate vector  $z_{ki}$ . If INIT = 0, then initial estimates of zero are used for all of the coefficients. This corresponds to no effect from any of the covariate values.
5. If a linear combination of covariates is monotonically increasing or decreasing with increasing failure times, then one or more of the

estimated coefficients is infinite and extended maximum likelihood estimates must be computed. Such estimates may be written as

$$\hat{\beta} = \hat{\beta}_f + \rho \hat{\gamma}$$

where  $\rho = \infty$  at the supremum of the likelihood so that

$$\hat{\beta}_f$$

is the finite part of the solution. In PHGLM, it is assumed that extended maximum likelihood estimates must be computed if, within any group  $k$ , for any time  $t$ ,

$$\min_{t_{ki} < t} \exp(w_{ki} + z_{ki} \hat{\beta}) > \rho \max_{t_{ki} < t} \exp(w_{ki} + z_{ki} \hat{\beta})$$

where  $\rho = \text{RATIO}$  is specified by the user. Thus, for example, if  $\rho = 10000$ , then PHGLM does not compute extended maximum likelihood estimates until the estimated proportionality constant

$$\exp(w_{ki} + z_{ki} \hat{\beta})$$

is 10000 times larger for all observations prior to  $t$  than for all observations after  $t$ . When this occurs, PHGLM computes estimates for

$$\hat{\beta}_f$$

by splitting the failures in stratum  $k$  into two strata at  $t$  (see Bryson and Johnson 1981). Censored observations in stratum  $k$  are placed into a stratum based upon the associated value for

$$\exp(w_{ki} + z_{ki} \hat{\beta})$$

The results of the splitting are returned in IGRP.

The estimates

$$\hat{\beta}_f$$

based upon the stratified likelihood represent the finite part of the extended maximum likelihood solution. Routine PHGLM does not compute

$$\hat{\gamma}$$

explicitly, but an estimate for

$$\hat{\gamma}$$

may be obtained in some circumstances by setting  $\text{RATIO} = -1$  and optimizing the log-likelihood without forming additional strata. The solution

$$\hat{\beta}$$

obtained will be such that

$$\hat{\beta} = \hat{\beta}_f + \rho \hat{\gamma}$$

for some finite value of  $\rho > 0$ . At this solution, the Newton-Raphson algorithm will not have “converged” because the Newton-Raphson step sizes returned in GR will be large, at least for some variables.

Convergence will be declared, however, because the relative change in the log-likelihood during the final iterations will be small.

### Example 1

The following data are taken from Lawless (1982, page 287) and involve the survival of lung cancer patients based upon their initial tumor types and treatment type. In the first example, the likelihood is maximized with no strata present in the data. This corresponds to Example 7.2.3 in Lawless (1982, page 367). The input data is printed in the output. The model is given as:

$$\ln(\lambda) = \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_3 + \alpha_i + \gamma_j$$

where  $\alpha_i$  and  $\gamma_j$  correspond to dummy variables generated from columns 6 and 7 of  $x$ , respectively,  $x_1$  corresponds to column 3 of  $x$ ,  $x_2$  corresponds to column 4 of  $x$ , and  $x_3$  corresponds to column 5 of  $x$ .

```

INTEGER      ICEN, IFIX, IFRQ, INIT, IPRINT, IRT, ISTRAT, ITIE,
&            LDCASE, LDcoef, LDCOV, LDX, MAXCL, MAXIT, NCLVAR,
&            NCOL, NEF, NOBS
REAL         EPS, RATIO
PARAMETER   (EPS=0.0001, ICEN=2, IFIX=0, IFRQ=0, INIT=0,
&            IPRINT=2, IRT=1, ISTRAT=0, ITIE=0, LDcoef=7, LDX=40,
&            MAXCL=10, MAXIT=30, NCLVAR=2, NCOL=7, NEF=5,
&            RATIO=10000.0, LDCASE=LDX, LDCOV=LDcoef, NOBS=LDX)
C
INTEGER      IGRP(NOBS), INDCL(NCLVAR), INDEF(5), NCLVAL(NCLVAR),
&            NCOEF, NRMISS, NVEF(NEF)
REAL         ALGL, CASE(LDCASE,5), CLVAL(6), COEF(LDcoef,4),
&            COV(LDCOV,LDCOV), GR(LDCOV), X(LDX,NCOL), XMEAN(LDCOV)
EXTERNAL     PHGLM, WRRRL
C
DATA X/411, 126, 118, 92, 8, 25, 11, 54, 153, 16, 56, 21, 287,
&      10, 8, 12, 177, 12, 200, 250, 100, 999, 231, 991, 1, 201,
&      44, 15, 103, 2, 20, 51, 18, 90, 84, 164, 19, 43, 340, 231,
&      5*0, 1, 16*0, 1, 5*0, 1, 11*0, 7, 6, 7, 4, 4, 7, 7, 8, 6,
&      3, 8, 4, 6, 4, 2, 5, 5, 4, 8, 7, 6, 9, 5, 7, 2, 8, 6, 5, 7,
&      4, 3, 3, 4, 6, 8, 7, 3, 6, 8, 7, 64, 63, 65, 69, 63, 48,
&      48, 63, 63, 53, 43, 55, 66, 67, 61, 63, 66, 68, 41, 53, 37,
&      54, 52, 50, 65, 52, 70, 40, 36, 44, 54, 59, 69, 50, 62, 68,
&      39, 49, 64, 67, 5, 9, 11, 10, 58, 9, 11, 4, 14, 4, 12, 2,
&      25, 23, 19, 4, 16, 12, 12, 8, 13, 12, 8, 7, 21, 28, 13, 13,
&      22, 36, 9, 87, 5, 22, 4, 15, 4, 11, 10, 18, 7*1, 7*2, 2*3,
&      5*4, 7*1, 4*2, 3*3, 5*4, 21*0, 19*1/
DATA NVEF/1, 1, 1, 1, 1/, INDEF/3, 4, 5, 6, 7/, INDCL/6, 7/
C
CALL WRRRL ('The First 10 Rows of the Input Data', 10, NCOL,
&          X, LDX, 0, '(I7)', 'Number', 'Number')

```



```

C      CALL PHGLM (NOBS, NCOL, X, LDX, IRT, IFRQ, IFIX, ICEN, ISTRAT,
&              MAXIT, EPS, RATIO, NCLVAR, INDCL, NEF, NVEF, INDEF,
&              INIT, ITIE, IPRINT, MAXCL, NCLVAL, CLVAL, NCOEF,
&              COEF, LDCOEF, ALGL, COV, LDICOV, XMEAN, CASE, LDCASE,
&              GR, IGRP, NRMIS)
C
      END

```

**Output**

The First 10 Rows of the Input Data

	1	2	3	4	5	6	7
1	411	0	7	64	5	1	0
2	126	0	6	63	9	1	0
3	118	0	7	65	11	1	0
4	92	0	4	69	10	1	0
5	8	0	4	63	58	1	0
6	25	1	7	48	9	1	0
7	11	0	7	48	11	1	0
8	54	0	8	63	4	2	0
9	153	0	6	63	14	2	0
10	16	0	3	53	4	2	0

Initial Estimates

1	2	3	4	5	6	7
0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000

Method	Iteration	Step size	Maximum scaled coef. update	Log likelihood
Q-N	0			-102.4
Q-N	1	1.0000	0.5034	-91.04
Q-N	2	1.0000	0.5782	-88.07
N-R	3	1.0000	0.1131	-87.92
N-R	4	1.0000	0.6958E-01	-87.89
N-R	5	1.0000	0.8144E-03	-87.89

Log-likelihood -87.88779

Coefficient Statistics

	Coefficient	Standard error	Asymptotic z-statistic	Asymptotic p-value
1	-0.585	0.137	-4.272	0.000
2	-0.013	0.021	-0.634	0.526
3	0.001	0.012	0.064	0.949
4	-0.367	0.485	-0.757	0.449
5	-0.008	0.507	-0.015	0.988
6	1.113	0.633	1.758	0.079
7	0.380	0.406	0.936	0.349

Asymptotic Coefficient Covariance

	1	2	3	4	5
1	0.1873E-01	0.2530E-03	0.3345E-03	0.5745E-02	0.9750E-02
2		0.4235E-03	-0.4120E-04	-0.1663E-02	-0.7954E-03
3			0.1397E-03	0.8111E-03	-0.1831E-02
4				0.2350	0.9799E-01
5					0.2568

		6	7
1	0.4264E-02	0.2082E-02	
2	-0.3079E-02	-0.2898E-02	
3	0.5995E-03	0.1684E-02	
4	0.1184	0.3735E-01	
5	0.1253	-0.1944E-01	
6	0.4008	0.6289E-01	
7		0.1647	

Case Analysis

	Survival Probability	Influence	Residual	Cumulative hazard	Proportionality constant
1	0.00	0.04	2.05	6.10	0.3
2	0.30	0.11	0.74	1.21	0.61
3	0.34	0.12	0.36	1.07	0.33
4	0.43	0.16	1.53	0.84	1.83
5	0.96	0.56	0.09	0.05	2.05
6	0.74	NaN	0.13	0.31	0.42
7	0.92	0.37	0.03	0.08	0.42
8	0.59	0.26	0.14	0.53	0.27
9	0.26	0.12	1.20	1.36	0.88
10	0.85	0.15	0.97	0.17	5.76
11	0.55	0.31	0.21	0.60	0.36
12	0.74	0.21	0.96	0.31	3.12
13	0.03	0.06	3.02	3.53	0.86
14	0.94	0.09	0.17	0.06	2.71
15	0.96	0.16	1.31	0.05	28.89
16	0.89	0.23	0.59	0.12	4.82
17	0.18	0.09	2.62	1.71	1.54
18	0.89	0.19	0.33	0.12	2.68
19	0.14	0.23	0.72	1.96	0.37
20	0.05	0.09	1.66	2.95	0.56
21	0.39	0.22	1.17	0.94	1.25
22	0.00	0.00	1.73	21.11	0.08
23	0.08	NaN	2.19	2.52	0.87
24	0.00	0.00	2.46	8.89	0.28
25	0.99	0.31	0.05	0.01	4.28
26	0.11	0.17	0.34	2.23	0.15
27	0.66	0.25	0.16	0.41	0.38
28	0.87	0.22	0.15	0.14	1.02
29	0.39	NaN	0.45	0.94	0.48
30	0.98	0.25	0.06	0.02	2.53
31	0.77	0.26	1.03	0.26	3.90
32	0.63	0.35	1.80	0.46	3.88
33	0.82	0.26	1.06	0.19	5.47
34	0.47	0.26	1.65	0.75	2.21
35	0.51	0.32	0.39	0.67	0.58
36	0.22	0.18	0.49	1.53	0.32
37	0.80	0.26	1.08	0.23	4.77
38	0.70	0.16	0.26	0.36	0.73
39	0.01	0.23	0.87	4.66	0.19
40	0.08	0.20	0.81	2.52	0.32

Last Coefficient Update

	1	2	3	4	5	6
	-1.016E-07	1.918E-09	-1.305E-08	-7.190E-07	-2.854E-07	2.108E-08

7  
-6.947E-08

Covariate Means

1	2	3	4	5	6	7
5.65	56.58	15.65	0.35	0.28	0.12	0.53

Distinct Values For Each Class Variable

Variable 1:	1.0	2.0	3.0	4.0
Variable 2:	0.	1.0		

Stratum Numbers For Each Observation

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	
1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	
1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1

Number of Missing Values 0

## Example 2

This example illustrates the use of PHGLM when there are strata present in the data. The observations from Example 1 are arbitrarily grouped into four strata (the first ten observations form stratum 1, the next 10 for stratum 2, etc.). Otherwise, the problem is unchanged. The resulting coefficients are very similar to those obtained when there is no stratification variable. The model is the same as in Example 1.

```

C      INTEGER      LDCASE, LDCCOEF, LDCOV, LDX, MAXCL, NCLVAR, NCOL, NEF,
      &              NOBS
      REAL          RATIO
      PARAMETER    (LDCCOEF=7, LDX=40, MAXCL=10, NCLVAR=2, NCOL=8, NEF=5,
      &              LDCCASE=LDX, LDCOV=LDCCOEF, NOBS=LDX, RATIO=10000.0)
C      SPECIFICATIONS FOR PARAMETERS
      INTEGER      ICEN, IFRQ, INIT, IFIX, IPRINT, IRT, ISTRAT, ITIE,
      &              MAXIT
      REAL          EPS
      PARAMETER    (EPS=0.0001, ICEN=2, IFRQ=0, INIT=0, IFIX=0,
      &              IPRINT=2, IRT=1, ISTRAT=8, ITIE=0, MAXIT=30)
C      SPECIFICATIONS FOR LOCAL VARIABLES
      INTEGER      IGRP(NOBS), NCLVAL(NCLVAR), NCCOEF, NRMIS
      REAL          ALGL, CASE(LDCASE,5), CLVAL(6), COEF(LDCCOEF,4),
      &              COV(LDCOV,LDCOV), GR(LDCOV), XMEAN(LDCOV)
C      SPECIFICATIONS FOR SAVE VARIABLES
      INTEGER      INDCL(NCLVAR), INDEF(NEF), NVEF(NEF)
      REAL          X(LDX,NCOL)
      SAVE         INDCL, INDEF, NVEF, X
C      SPECIFICATIONS FOR SUBROUTINES
      EXTERNAL    PHGLM
C
      DATA X/411, 126, 118, 92, 8, 25, 11, 54, 153, 16, 56, 21, 287,
      &          10, 8, 12, 177, 12, 200, 250, 100, 999, 231, 991, 1, 201,
      &          44, 15, 103, 2, 20, 51, 18, 90, 84, 164, 19, 43, 340, 231,
      &          5*0, 1, 16*0, 1, 5*0, 1, 11*0, 7, 6, 7, 4, 4, 7, 7, 8, 6,
      &          3, 8, 4, 6, 4, 2, 5, 5, 4, 8, 7, 6, 9, 5, 7, 2, 8, 6, 5, 7,
      &          4, 3, 3, 4, 6, 8, 7, 3, 6, 8, 7, 64, 63, 65, 69, 63, 48,

```

```

&      48, 63, 63, 53, 43, 55, 66, 67, 61, 63, 66, 68, 41, 53, 37,
&      54, 52, 50, 65, 52, 70, 40, 36, 44, 54, 59, 69, 50, 62, 68,
&      39, 49, 64, 67, 5, 9, 11, 10, 58, 9, 11, 4, 14, 4, 12, 2,
&      25, 23, 19, 4, 16, 12, 12, 8, 13, 12, 8, 7, 21, 28, 13, 13,
&      22, 36, 9, 87, 5, 22, 4, 15, 4, 11, 10, 18, 7*1, 7*2, 2*3,
&      5*4, 7*1, 4*2, 3*3, 5*4, 21*0, 19*1, 10*1, 10*2, 10*3, 10*4/
DATA NVEF/1, 1, 1, 1, 1/, INDEF/3, 4, 5, 6, 7/, INDCL/6, 7/
C
CALL PHGLM (NOBS, NCOL, X, LDX, IRT, IFRQ, IFIX, ICEN, ISTRAT,
&          MAXIT, EPS, RATIO, NCLVAR, INDCL, NEF, NVEF, INDEF,
&          INIT, ITIE, IPRINT, MAXCL, NCLVAL, CLVAL, NCOEF,
&          COEF, LDcoef, ALGL, COV, LDcov, XMEAN, CASE, LDCASE,
&          GR, IGRP, NRMISS)
C
END

```

### Output

```

Initial Estimates
      1      2      3      4      5      6      7
0.0000  0.0000  0.0000  0.0000  0.0000  0.0000  0.0000

```

```

Method  Iteration  Step size  Maximum scaled  Log
                    coef. update  likelihood
Q-N      0
Q-N      1      1.0000      0.6748      -45.79
Q-N      2      1.0000      0.7105      -42.85
N-R      3      1.0000      0.2315      -42.59
N-R      4      1.0000      0.1674      -42.55
N-R      5      1.0000      0.3372E-02  -42.55

```

Log-likelihood -42.54570

```

Coefficient Statistics
      Coefficient      Standard      Asymptotic      Asymptotic
      error      z-statistic      p-value
1      -0.716      0.170      -4.222      0.000
2      -0.033      0.030      -1.122      0.262
3      0.001      0.015      0.048      0.961
4      -0.100      0.999      -0.100      0.921
5      -0.405      0.729      -0.555      0.579
6      1.136      0.769      1.478      0.139
7      -0.087      1.454      -0.060      0.952

```

```

Asymptotic Coefficient Covariance
      1      2      3      4      5
1      0.2877E-01      0.8662E-03      0.3119E-03      0.5057E-02      0.2480E-01
2      0.8842E-03      -0.8137E-04      -0.7623E-02      -0.6925E-03
3      0.2158E-03      -0.2567E-02      -0.3738E-02
4      0.9975      0.5109
5      0.5319
      6      7
1      -0.7669E-02      0.6405E-02
2      -0.8800E-03      0.4120E-02
3      0.1170E-02      -0.3699E-02
4      0.1944      0.8056
5      0.1802      0.4905
6      0.5909      0.1858

```

		Case Analysis				
	Survival Probability	Influence	Residual	Cumulative hazard	Proportionality constant	
1	0.00	0.00	2.01	7.83	0.26	
2	0.09	0.06	1.32	2.42	0.55	
3	0.20	0.04	0.40	1.59	0.25	
4	0.40	0.04	1.69	0.91	1.87	
5	0.92	0.47	0.21	0.09	2.36	
6	0.73	NaN	0.14	0.31	0.44	
7	0.82	0.47	0.09	0.20	0.44	
8	0.55	0.67	0.06	0.61	0.10	
9	0.02	0.07	1.59	3.94	0.40	
10	0.73	0.10	1.50	0.31	4.79	
11	0.39	0.68	0.17	0.93	0.19	
12	0.60	0.14	1.12	0.51	2.19	
13	0.00	0.00	2.32	6.32	0.37	
14	0.90	0.16	0.15	0.10	1.49	
15	0.98	0.04	0.75	0.02	35.42	
16	0.75	0.21	1.12	0.29	3.83	
17	0.25	0.07	1.55	1.39	1.12	
18	0.75	0.21	0.63	0.29	2.14	
19	0.10	0.18	0.69	2.31	0.30	
20	0.03	0.11	1.48	3.60	0.41	
21	0.50	0.61	1.00	0.70	1.44	
22	0.00	0.00	1.28	13.59	0.09	
23	0.33	NaN	1.92	1.09	1.76	
24	0.05	0.00	1.32	2.94	0.45	
25	0.95	0.15	0.47	0.05	9.84	
26	0.33	0.24	0.23	1.09	0.21	
27	0.62	0.40	0.22	0.47	0.47	
28	0.76	0.13	0.71	0.27	2.63	
29	0.50	NaN	0.37	0.70	0.53	
30	0.87	0.23	0.49	0.14	3.53	
31	0.88	0.35	0.67	0.13	5.07	
32	0.71	0.22	1.56	0.34	4.54	
33	0.97	0.52	0.20	0.03	7.00	
34	0.44	0.03	2.64	0.83	3.19	
35	0.56	0.20	0.29	0.57	0.50	
36	0.11	0.00	0.61	2.24	0.27	
37	0.94	0.19	0.82	0.07	12.50	
38	0.79	0.43	0.24	0.23	1.05	
39	0.00	0.00	1.69	11.13	0.15	
40	0.01	0.00	1.28	4.54	0.28	

Last Coefficient Update						
1	2	3	4	5	6	7
-7.363E-07	8.762E-09	1.252E-08	-1.697E-06	-1.642E-06	1.075E-06	
						-1.772E-06

Covariate Means						
1	2	3	4	5	6	7
5.65	56.58	15.65	0.35	0.28	0.12	0.53

Distinct Values For Each Class Variable  
 Variable 1: 1.0 2.0 3.0 4.0

```

Variable 2:      0.      1.0

                Stratum Numbers For Each Observation
 1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20
 1  1  1  1  1  1  1  1  1  1  2  2  2  2  2  2  2  2  2  2
21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40
 3  3  3  3  3  3  3  3  3  3  4  4  4  4  4  4  4  4  4  4
Number of Missing Values      0

```

---

## SVGLM/DSVGLM (Single/Double precision)

Analyze censored survival data using a generalized linear model.

### Usage

```

CALL SVGLM (NOBS, NCOL, X, LDX, MODEL, ILT, IRT, IFRQ,
            IFIX, ICEN, INFIN, MAXIT, EPS, INTCEP, NCLVAR,
            INDCL, NEF, NVEF, INDEF, INIT, IPRINT, MAXCL,
            NCLVAL, CLVAL, NCOEF, COEF, LDCEOF, ALGL, COV,
            LDCOV, XMEAN, CASE, LDCASE, GR, IADD, NRMISS)

```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in *x*. (Input)

**X** — NOBS by NCOL matrix containing the data. (Input)

**LDX** — Leading dimension of *x* exactly as specified in the dimension statement in the calling program. (Input)

**MODEL** — Model option parameter. (Input)

MODEL specifies the distribution of the response variable and the relationship of the linear model to a distribution parameter.

#### MODEL Distribution

0	Exponential
1	Linear hazard
2	Log-normal
3	Normal
4	Log-logistic
5	Logistic
6	Log least extreme value
7	Least extreme value
8	Log extreme value
9	Extreme value
10	Weibull

For further discussion of the models and parameterizations used, see the Algorithm section.

**ILT** — For interval-censored and left-censored observations, the column number in  $X$  that contains the upper endpoint of the failure interval. (Input)  
See argument **ICEN**. If **ILT** = 0, left-censored and interval-censored observations cannot be input.

**IRT** — For interval-censored and right-censored observations, the column number in  $X$  that contains the lower endpoint of the failure interval. (Input)  
For exact-failure observations,  $X(i, IRT)$  contains the exact-failure time. **IRT** must not be zero. See argument **ICEN**.

**IFRQ** — Column number in  $X$  containing the frequency of response for each observation. (Input)  
If **IFRQ** = 0, a response frequency of 1 for each observation is assumed.

**IFIX** — Column number in  $X$  containing a constant to be added to the linear response. (Input)  
The estimated linear response is taken to be

$$w_i + z_i \hat{\beta}$$

where  $w_i$  is the observation constant,  $z_i$  is the observation design vector,

$$\hat{\beta}$$

is the vector of estimated parameters output in the first column of **COEF**, and  $i$  indexes the observations. The “fixed” constant allows one to test hypotheses about parameters via the log-likelihoods. If **IFIX** = 0, the fixed parameter is assumed to be 0.

**ICEN** — Column number in  $X$  containing the censoring code for each observation. (Input)  
If **ICEN** = 0, a censoring code of 0 is assumed. Valid censoring codes are:

<b>x(i, ICEN)</b>	<b>Censoring</b>
0	Exact failure at $X(i, IRT)$ .
1	Right censored. The response is greater than $X(i, IRT)$ .
2	Left censored. The response is less than or equal to $X(i, ILT)$ .
3	Interval censored. The response is greater than $X(i, IRT)$ , but less than or equal to $X(i, ILT)$ .

**INFIN** — Method to be used for handling infinite estimates. (Input)

<b>INFIN</b>	<b>Method</b>
0	Remove a right or left-censored observation from the loglikelihood whenever the probability of the observation exceeds 0.995. At convergence, use linear programming to check that all removed observations actually have infinite linear response

$$z_i \hat{\beta}$$

Set  $IADD(i)$  for observation  $i$  to 2 if the linear response is infinite. If not all removed observations have infinite linear response, recompute the estimates based upon the observations with finite

$$z_i \hat{\beta}$$

1 Iterate without checking for infinite estimates.

See the algorithm section for more discussion.

**MAXIT** — Maximum number of iterations. (Input)

MAXIT = 30 will usually be sufficient. Use MAXIT = 0 to compute the Hessian and score vector at the initial estimates.

**EPS** — Convergence criterion. (Input)

Convergence is assumed when the maximum relative change in any coefficient estimate is less than EPS from one iteration to the next, or when the relative change in the log-likelihood, ALGL, from one iteration to the next is less than EPS/100. If EPS is negative, EPS = 0.001 is assumed.

**INTCEP** — Intercept option. (Input)

**INTCEP Action**

0 No intercept is in the model (unless otherwise provided for by the user).

1 An intercept is automatically included in the model.

**NCLVAR** — Number of classification variables. (Input)

Dummy or indicator variables are generated for classification variables using the IDUMMY = 2 option of routine GRGLM (page 210). See Comment 3.

**INDCL** — Index vector of length NCLVAR containing the column numbers of X that are classification variables. (Input, if NCLVAR is positive, not used otherwise)

If NCLVAR is 0, INDCL is not referenced and can be dimensioned of length 1 in the calling program.

**NEF** — Number of effects in the model. (Input)

In addition to effects involving classification variables, simple covariates and the product of simple covariates are also considered effects.

**NVEF** — Vector of length NEF containing the number of variables associated with each effect in the model. (Input, if NEF is positive; not used otherwise)

If NEF is zero, NVEF is not used and can be dimensioned of length 1 in the calling program.

**INDEF** — Index vector of length NVEF(1) + NVEF(2) + ... + NVEF(NEF) containing the column numbers in X associated with each effect. (Input, if NEF is positive; not used otherwise)

The first NVEF(1) elements of INDEF give the column numbers in X of the variables in the first effect. The next NVEF(2) elements of INDEF give the column numbers for the second effect, etc. If NEF is zero, INDEF is not used and can be dimensioned of length one in the calling program.

**INIT** — Initialization option. (Input)



**INIT Action**  
 0 Unweighted linear regression is used to obtain initial estimates.  
 1 The NCOEF elements in the first column of COEF contain initial estimates of the parameters on input to SVGLM (requiring that the user know NCOEF prior to calling SVGLM).

**IPRINT** — Printing option. (Input)

**IPRINT Action**  
 0 No printing is performed.  
 1 Printing is performed, but observational statistics are not printed.  
 2 All output statistics are printed.

**MAXCL** — An upper bound on the sum of the number of distinct values taken by the classification variables. (Input)

**NCLVAL** — Vector of length NCLVAR containing the number of values taken by each classification variable. (Output, if NCLVAR is positive; not used otherwise) NCLVAL(*i*) is the number of distinct values for the *i*-th classification variable. If NCLVAR is zero, NCLVAL is not used and can be dimensioned of length 1 in the calling program.

**CLVAL** — Vector of length NCLVAL(1) + NCLVAL(2) + ... + NCLVAL(NCLVAR) containing the distinct values of the classification variables in ascending order. (Output, if NCLVAR is positive, not used otherwise)

The first NCLVAL(1) elements contain the values for the first classification variables, the next NCLVAL(2) elements contain the values for the second classification variable, etc. If NCLVAR is zero, then CLVAL is not referenced and can be dimensioned of length 1 in the calling program.

**NCOEF** — Number of estimated coefficients in the model. (Output, if INIT = 0; input, if INIT = 1)

**COEF** — NCOEF by 4 matrix containing parameter estimates and associated statistics. (Output, if INIT = 0; input/output, if INIT = 1; input, if MAXIT = 0)

<b>Col.</b>	<b>Statistic</b>
1	Coefficient estimate.
2	Estimated standard deviation of the estimated coefficient.
3	Asymptotic normal score for testing that the coefficient is zero.
4	<i>p</i> -value associated with the normal score in column 3.

When COEF is input, only column 1 is referenced as input data, and columns 2 to 4 need not be set. When present in the model, the initial coefficient in COEF estimates a “nuisance” parameter, and the remaining coefficients estimate parameters associated with the “linear” model, beginning with the intercept, if present. Nuisance parameters are as follows:

<b>Model</b>	<b>Nuisance Parameter</b>
1	Coefficient of the quadratic term in time, $\theta$
2 – 9	Scale parameter, $\sigma$
10	Shape parameter, $\theta$

**LDCOEF** — Leading dimension of **COEF** exactly as specified in the dimension statement in the calling program. (Input)

**ALGL** — Maximized log-likelihood. (Output)

**COV** — **NCOEF** by **NCOEF** matrix containing the estimated asymptotic covariance matrix of the coefficients. (Output)

**COV** is computed as the inverse of the matrix of second partial derivatives of negative one times the log-likelihood. When **MAXIT** = 0, **COV** is computed at the initial estimates.

**LDCOV** — Leading dimension of **COV** exactly as specified in the dimension statement in the calling program. (Input)

**XMEAN** — Vector of length **NCOEF** containing the means of the design variables. (Output)

**CASE** — **NOBS** by 5 vector containing the case analysis. (Output)

**Col.     Statistic**

- 1     Estimated predicted value
- 2     Estimated influence or leverage
- 3     Residual estimate
- 4     Estimated cumulative hazard
- 5     For non-censored observations, the estimated density at the observation failure time and covariate values. For censored observations, the corresponding estimated probability.

If **MAXIT** = 0, **CASE** is a **NOBS** by 1 vector containing the estimated probability (for censored observations) or the estimated density (for non censored observations).

**LDCASE** — Leading dimension of **CASE** exactly as specified in the dimension statement in the calling program. (Input)

**GR** — Vector of length **NCOEF** containing the last parameter updates, excluding step halvings. (Output)

**GR** is computed as the inverse of the matrix of second partial derivatives times the vector of first partial derivatives of the log-likelihood. When **MAXIT** = 0, the derivatives are computed at the initial estimates.

**IADD** — Vector of length **NOBS** indicating which observations have and have not been included in the model. (Output, if **MAXIT** > 0; input/output, if **MAXIT** = 0)

**Value    Status of Observation**

- 0     Observation *i* has been included in the model.
- 1     Observation *i* has not been included in the model due to missing values in the **X** matrix.
- 2     Observation *i* has not been included in the model because of infinite estimates in extended maximum likelihood estimation. If **MAXIT** = 0, then the **IADD** array must be initialized prior to calling **SVGLM**.

***NRMIS*** — Number of rows of data in *x* that contain missing values in one or more columns *ILT*, *IRT*, *IFRQ*, *ICOEF*, *ICEN*, *INDCL* or *INDEF* of *x*. (Output)

### Comments

1. Automatic workspace usage is

**SVGLM**  $6 * NMAX + 5 * NCOEF + 7 * NOBS + NMAX * NCOEF + 8$  units if *INFIN* = 0 and  $5 * NCOEF + 7 * NOBS + 8$  units if *INFIN* = 1, or

**DSVGLM**  $10 * NMAX + 10 * NCOEF + 9 * NOBS + 2 * NMAX * NCOEF + 10$  units if *INFIN* = 0 and  $10 * NCOEF + 9 * NOBS + 10$  units if *INFIN* = 1. *NMAX* is defined in the list below.

Workspace may be explicitly provided, if desired, by use of **S2GLM/DS2GLM**. The reference is

```
CALL S2GLM (NOBS, NCOL, X, LDX, MODEL, ILT,
            IRT, IFRQ, IFIX, ICEN, INFIN, MAXIT,
            EPS, INTCEP, NCLVAR, INDCL, NEF, NVEF,
            INDEF, INIT, IPRINT, MAXCL, NCLVAL, CLVAL,
            NCOEF, COEF, LDCEP, ALGL, COV, LDCOV,
            XMEAN, CASE, LDCASE, GR, IADD, NRMIS,
            NMAX, OBS, ADD, XD, WK, KBASIS, RWK,
            IWK)
```

The additional arguments are as follows:

***NMAX*** — Maximum number of observations that can be handled in the linear programming. (Input)

If workspace is not explicitly provided, *NMAX* is set to

$NMAX = (n - 6) / (6 + NCOEF)$  in **SVGLM** and  $NMAX = (n - 6) / (10 + 2 * NCOEF)$  in **DSVGLM**, where *n* is the number of units of workspace remaining after allocating workspace for *OBS*, *RWK*, and *IWK*. If *INFIN* .EQ. 1, then set *NMAX* to 0.

***OBS*** — Work array of length  $2 * NCOEF + 2$ .

***ADD*** — Logical work array of length *NMAX*. If *ADD*(*I*) = .TRUE., the *I*-th observation deleted from the model was returned to the model. *ADD* is not needed and can be a array of length 1 in the calling program if *NMAX* = 0.

***XD*** — Work array of length *NMAX* \* *NCOEF*. *XD* is not needed and can be a array of length 1 in the calling program if *NMAX* = 0.

***WK*** — Work array of length  $4 * NMAX$ . *WK* is not needed and can be a array of length 1 in the calling program if *NMAX* = 0.

***KBASIS*** — Work array of length *NMAX*. *KBASIS* is not needed and can be a array of length 1 in the calling program if *NMAX* = 0.

***RWK*** — Work array of length  $2 * NOBS + 3 * NCOEF$ .

***IWK*** — Work array of length  $5 * NOBS + 6$ .

2. Informational errors
- | Type | Code |  |
|------|------|--|
| 3    | 1    | There were too many iterations required. Convergence is assumed.   |
| 3    | 2    | There were too many step halvings. Convergence is assumed.   |
| 3    | 4    | The censoring interval has a length of 0. The censoring code for this observation is set to 0.   |
| 3    | 5    | COEF(1, 1) > 1.0. The expected value for the log logistic distribution, MODEL = 4, does not exist. Predicted values are not calculated.      |
| 3    | 6    | COEF(1, 1) > 1.0. The expected value for the log extreme value distribution, MODEL = 8, does not exist. Predicted values are not calculated. |
| 4    | 7    | The number of distinct values of the classification variables exceeds MAXCL. MAXCL must be increased for the computations to proceed.        |
| 4    | 8    | The number of distinct values for each classification variable must be greater than one.   |
| 4    | 9    | INIT = 1 and the number of coefficients input in NCOEF does not equal the number of coefficients required by the specified model.            |
| 4    | 10   | For the exponential model, NCOEF has been determined to equal 0. With no coefficients in the model, processing cannot continue.              |
| 4    | 11   | LDCOEFF or LDICOV is less than NCOEF.  |
| 4    | 12   | NOBS - NRMISS must be greater than or equal to 2 in order to estimate the coefficients.  |
| 4    | 13   | The number of observations to be deleted has exceeded NMAX. Rerun with a different model or increase the workspace.                          |
3. Dummy variables are generated for the classification variables as follows: An ascending list of all distinct values of each classification variable is obtained and stored in CLVAL. Dummy variables are then generated for each but the last of these distinct values. Each dummy variable is zero unless the classification variable equals the list value corresponding to the dummy variable, in which case the dummy variable is one. See Argument IDUMMY for IDUMMY = 2 in routine GRGLM (page 210).
4. The “product” of a classification variable with a covariate yields dummy variables equal to the product of the covariate with each of the dummy variables associated with the classification variable.
5. The “product” of two classification variables yields dummy variables in the usual manner. Each dummy variable associated with the first classification variable multiplies each dummy variable associated with

the second classification variable. The resulting dummy variables are such that the index of the second classification variable varies fastest.

### Algorithm

Routine *SVGLM* computes maximum likelihood estimates of parameters and associated statistics in generalized linear models commonly found in survival (reliability) analysis. Although the terminology used will be from the survival area, the methods discussed have application in many areas of data analysis, including reliability analysis and event history analysis. Indeed, these methods may be used anywhere a random variable from one of the discussed distributions is parameterized via one of the models available in *SVGLM*. Thus, while it is not advisable to do so, standard multiple linear regression may be performed by routine *SVGLM*. Estimates for any of ten standard models can be computed. Exact, left-censored, right-censored, or interval-censored observations are allowed. (Note that left censoring is the same as interval censoring with left endpoint equal to the left endpoint of the support of the distribution.)

Let  $\eta = x^T \beta$  be the linear parameterization, where  $x$  is a design vector obtained in *SVGLM* via routine *GRGLM* (page 210) from a row of  $x$ , and  $\beta$  is a vector of parameters associated with the linear model. Let  $T$  denote the random response variable and  $S(t)$  denote the probability that  $T > t$ . All models considered also allow a fixed parameter  $w_i$  for observation  $i$  (input in column *IFIX* of  $x$ ). Use of this parameter is discussed below. There may also be nuisance parameters  $\theta > 0$ , or  $\sigma > 0$  to be estimated (along with  $\beta$ ) in the various models. Let  $\Phi$  denote the cumulative normal distribution. The survival models available in *SVGLM* are:

Model	Name	$S(t)$
0	Exponential	$\exp\{-\exp(w_i + \eta)\}$
1	Linear hazard	$\exp\{-(t + \frac{\theta t^2}{2})\exp[(w_i + \eta)]\}$
2	Log-normal	$1 - \Phi(\frac{\ln(t) - \eta - w_i}{\sigma})$
3	Normal	$1 - \Phi(\frac{t - \eta - w_i}{\sigma})$
4	Log-logistic	$\{1 + \exp(\frac{\ln(t) - \eta - w_i}{\sigma})\}^{-1}$
5	Logistic	$\{1 + \exp(\frac{t - \eta - w_i}{\sigma})\}^{-1}$
6	Log least extreme value	$\exp\{-\exp(\frac{\ln(t) - \eta - w_i}{\sigma})\}$
7	Least extreme value	$\exp\{-\exp(\frac{t - \eta - w_i}{\sigma})\}$
8	Log extreme value	$1 - \exp\{-\exp[-\frac{\ln(t) - \eta - w_i}{\sigma}]\}$
9	Extreme value	$1 - \exp\{-\exp[-(\frac{t - \eta - w_i}{\sigma})]\}$
10	Weibull	$\exp\{-(\frac{t}{\exp(w_i + \eta)})^\theta\}$

Note that the log-least-extreme-value model is a reparameterization of the Weibull model. Moreover, models 0, 1, 2, 4, 6, 8, and 10 require that  $T > 0$ , while all of the remaining models allow any value for  $T, -\infty < T < \infty$ .

Each row in the data matrix can represent a single observation, or, through the use of column `IFRQ`, it can represent several observations. Classification variables and their products are easily incorporated into the models via the usual GLM type specifications through the use of variables `NCLVAR` and `INDCL`, and the model variables `NEF`, `NVEF`, and `INDEF`.

The constant parameter  $w_i$  is input in  $X$  and may be used for a number of purposes. For example, if the parameter in an exponential model is known to depend upon the size of the area tested, volume of a radioactive mass, or population density, etc., then a multiplicative factor of the exponential parameter  $\lambda = \exp(x\beta)$  may be known a priori. This factor can be input in  $w_i$  ( $w_i$  is the log of the factor). An alternate use of  $w_i$  is as follows: It may be that  $\lambda = \exp(x_1\beta_1 + x_2\beta_2)$ , where  $\beta_2$  is known. Letting  $w_i = x_2\beta_2$ , estimates for  $\beta_1$  can be obtained via `SVGLM` with the known fixed values for  $\beta_2$ . Standard methods can then be used to test hypotheses about  $\beta_2$  via computed log-likelihoods.

### Computational details

The computations proceed as follows:

1. The input arguments are checked for consistency and validity.
2. Estimates for the means of the explanatory variables  $x$  (as generated from the model specification via `GRGLM`, page 210) are computed. Let  $f_i$  denote the frequency of the observation. Means are computed as

$$\bar{x} = \frac{\sum_i f_i x_i}{\sum_i f_i}$$

3. If `INIT = 0`, initial estimates of the parameters for all but the exponential models (`MODEL = 0, 1`) are obtained as follows:
  - A. Routine `KAPMR` (page 938) is used to compute a nonparametric estimate of the survival probability at the upper limit of each failure interval. (Because upper limits are used, interval and left-censored data are taken to be exact failures at the upper endpoint of the failure interval.) The Kaplan-Meier estimate is computed under the assumption that all failure distributions are identical (i.e., all  $\beta$ 's but the intercept, if present, are assumed to be zero).
  - B. If `INTCEP = 0`, a simple linear regression is performed predicting

$$S^{-1}(\hat{S}(t)) - w_i = \alpha + \phi t^*$$

where  $t^*$  is computed at the upper endpoint of each failure interval,  $t^* = t$  in models 3, 5, 7, and 9, and  $t^* = \ln(t)$  in models 2, 4, 6, 8, and 10, and  $w_i$  is the fixed constant, if present. If INTCEP is zero,  $\alpha$  is fixed at zero, and the model

$$S^{-1}(\hat{S}(t)) - \hat{\phi}t^* - w_i = x^T\beta$$

is fit instead of the model above. In this model, the coefficients  $\beta$  are used in place of the location estimate  $\alpha$  above. Here,

$$\hat{\phi}$$

is estimated from the simple linear regression with  $\alpha = 0$ .

- C. If the intercept is in the model, then in log-location-scale models (models 1–8),

$$\hat{\sigma} = \hat{\phi}$$

and the initial estimate of the intercept, if present, is taken to be

$$\hat{\alpha}$$

In the Weibull model,

$$\hat{\theta} = 1 / \hat{\phi}$$

and the intercept, if present, is taken to be

$$\hat{\alpha}$$

Initial estimates of all parameters  $\beta$ , other than the intercept, are taken to be zero.

If no intercept is in the model, the scale parameter is estimated as above, and the estimates

$$\hat{\beta}$$

from Step B are used as initial estimates for the  $\beta$ 's.

For exponential models (MODEL = 0, 1), the average total time on test statistic is used to obtain an estimate for the intercept. Specifically, let  $T_i$  denote the total number of failures divided by the total time on test. The initial estimate for the intercept is then  $\ln(T_i)$ . Initial estimates for the remaining parameters  $\beta$  are taken as zero, and, if MODEL = 1, the initial estimate for the linear hazard parameter  $\theta$  is taken to be a small positive number. When the intercept is not in the model, the initial estimate for the parameter  $\theta$  is taken as a small positive number, and initial estimates of the parameters  $\beta$  are computed via multiple linear regression as above.

4. A quasi-Newton algorithm is used in the initial iterations based upon a Hessian estimate

$$\hat{H}_{\kappa_j \kappa_l} = \sum_i \ell'_{i\alpha_j} \ell'_{i\alpha_l}$$

where

$$\ell'_{i\alpha_j}$$

is the partial derivative of the  $i$ -th term in the log-likelihood with respect to the parameter  $\alpha_j$ , and  $\alpha_j$  denotes one of the parameters to be estimated.

When the relative change in the log-likelihood from one iteration to the next is 0.1 or less, exact second partial derivatives are used for the Hessian so that Newton-Raphson iteration is used.

If the initial step size results in an increase in the log-likelihood, the full step is used. If the log-likelihood decreases for the initial step size, the step size is halved, and a check for an increase in the log-likelihood performed. Step-halving is performed (as a simple line search) until an increase in the log-likelihood is detected, or until the step size is less than 0.0001 (where the initial step size is 1).

5. Convergence is assumed when the maximum relative change in any coefficient update from one iteration to the next is less than `EPS`, or when the relative change in the loglikelihood from one iteration to the next is less than `EPS/100`. Convergence is also assumed after `MAXIT` iterations, or when step halving leads to a step size of less than `.0001`, with no increase in the log-likelihood.
6. If requested (`INFIN = 0`), then the methods of Clarkson and Jennrich (1988) are used to check for the existence of infinite estimates in

$$\eta_i = x_i^T \beta$$

As an example of a situation in which infinite estimates can occur, suppose that observation  $j$  is right censored with  $t_j > 15$  in a normal distribution model in which we fit the mean as

$$\mu_j = x_j^T \beta = \eta_j$$

where  $x_j$  is the observation design vector. If design vector  $x_j$  for parameter  $\beta_m$  is such that  $x_{jm} = 1$  and  $x_{im} = 0$  for all  $i \neq j$ , then the optimal estimate of  $\beta_m$  occurs at

$$\hat{\beta}_m = \infty$$

leading to an infinite estimate of both  $\beta_m$  and  $\eta_j$ . In `SVGLM`, such estimates may be “computed.”



In all models fit by SVGLM, infinite estimates can only occur when the optimal estimated probability associated with the left or right-censored observation is 1. If  $\text{INFIN} = 0$ , left- or right-censored observations that have estimated probability greater than 0.995 at some point during the iterations are excluded from the log-likelihood, and the iterations proceed with a log-likelihood based upon the remaining observations. This allows convergence of the algorithm when the maximum relative change in the estimated coefficients is small and also allows for a more precise determination of observations with infinite

$$\eta_i = x_i^T \beta$$

At convergence, linear programming is used to ensure that the eliminated observations have infinite  $\eta_i$ . If some (or all) of the removed observations should not have been removed (because their estimated  $\eta_i$ 's must be finite), then the iterations are restarted with a log-likelihood based upon the finite  $\eta_i$  observations. See Clarkson and Jennrich (1988) for more details.

When  $\text{INFIN} = 1$ , no observations are eliminated during the iterations. In this case, when infinite estimates occur, some (or all) of the coefficient estimates

$$\hat{\beta}$$

will become large, and it is likely that the Hessian will become (numerically) singular prior to convergence.

7. The case statistics are computed as follows:

Let

$$l_i(\theta_i)$$

denote the log-likelihood of the  $i$ -th observation evaluated at  $\theta_i$ , let

$$l'_i$$

denote the vector of derivatives of

$$l_i$$

with respect to all parameters,

$$l'_{\eta,i}$$

denote the derivative of

$$l_i$$

with respect to  $\eta = x^T \beta$ ,  $H$  denote the Hessian, and  $E$  denote expectation. Then, the columns of CASE are:

- A. Predicted values are computed as  $E(T|x)$  according to standard formulas. If MODEL is 4 or 8, and if  $\sigma \geq 1$ , then the expected values cannot be computed because they are infinite.

- B. Following Cook and Weisberg (1982), we take the influence (or leverage) of the  $i$ -th observation to be

$$(\ell'_i)^T H^{-1} \ell'_i$$

This quantity is a one-step approximation to the change in the estimates when the  $i$ -th observation is deleted (ignoring the nuisance parameters).

- C. The “residual” is computed as

$$\ell'_{\hat{\eta},i}$$

- D. The cumulative hazard is computed at the observation covariate values and, for interval observations, the upper endpoint of the failure interval. The cumulative hazard can also be used as a “residual” estimate. If the model is correct, the cumulative hazards should follow a standard exponential distribution. See Cox and Oakes (1984).

- E. The density (for exact failures) or the interval probability (for censored observations) is computed for given  $x$ .

### Programming Notes

Classification variables are specified by parameters NCLVAR and INDCL. Indicator variables are created for the classification variables using routine GRGLM (page 210) with IDUMMY = 2.

### Example 1

This example is from Lawless (1982, page 287) and involves the mortality of patients suffering from lung cancer. (The first ten rows of the input data are printed in the output.) An exponential distribution is fit for model

$$\eta = \mu + \beta_1 x_3 + \beta_2 x_4 + \beta_3 x_5 + \alpha_i + \gamma_k$$

where  $\alpha_i$  is associated with a classification variable with 4 levels, and  $\gamma_k$  is associated with a classification variable with 2 levels. Note that because the computations are performed in single precision, there will be some small variation in the estimated coefficients across different machine environments.

```

C
  INTEGER      ICEN, IFIX, IFRQ, ILT, INFIN, INIT, INTCEP, IPAR,
&             IPRINT, IRT, LDCASE, LDCEOF, LDCEOV, LDX, MAXCL,
&             MAXIT, MODEL, NCLVAR, NCOL, NEF, NOBS
  REAL        EPS
  PARAMETER   (EPS=0.001, ICEN=2, IFIX=0, IFRQ=0, ILT=0, INFIN=0,
&             INIT=0, INTCEP=1, IPAR=0, IPRINT=2, IRT=1, LDCASE=40,
&             LDCEOF=8, LDCEOV=8, LDX=40, MAXCL=6, MAXIT=15,
&             MODEL=0, NCLVAR=2, NCOL=7, NEF=5, NOBS=40)
  INTEGER      IADD(NOBS), INDCL(NCLVAR), INDEF(5), NCLVAL(NCLVAR),
&             NCOEF, NRMIS, NVEF(NEF)
  REAL        ALGL, CASE(LDCASE,5), CLVAL(MAXCL), COEF(LDCEOF,4),

```

```

&          COV(LDCOV,LDCOV), GR(LDCOV), X(LDX,NCOL), XMEAN(LDCOV)
EXTERNAL   SVGLM, WRRRL
C
  DATA X/411, 126, 118, 92, 8, 25, 11, 54, 153, 16, 56, 21, 287,
&        10, 8, 12, 177, 12, 200, 250, 100, 999, 231, 991, 1, 201,
&        44, 15, 103, 2, 20, 51, 18, 90, 84, 164, 19, 43, 340, 231,
&        5*0, 1, 16*0, 1, 5*0, 1, 11*0, 7, 6, 7, 4, 4, 7, 7, 8, 6,
&        3, 8, 4, 6, 4, 2, 5, 5, 4, 8, 7, 6, 9, 5, 7, 2, 8, 6, 5, 7,
&        4, 3, 3, 4, 6, 8, 7, 3, 6, 8, 7, 64, 63, 65, 69, 63, 48,
&        48, 63, 63, 53, 43, 55, 66, 67, 61, 63, 66, 68, 41, 53, 37,
&        54, 52, 50, 65, 52, 70, 40, 36, 44, 54, 59, 69, 50, 62, 68,
&        39, 49, 64, 67, 5, 9, 11, 10, 58, 9, 11, 4, 14, 4, 12, 2,
&        25, 23, 19, 4, 16, 12, 12, 8, 13, 12, 8, 7, 21, 28, 13, 13,
&        22, 36, 9, 87, 5, 22, 4, 15, 4, 11, 10, 18, 7*1, 7*2, 2*3,
&        5*4, 7*1, 4*2, 3*3, 5*4, 21*0, 19*1/
  DATA NVEF/1, 1, 1, 1, 1/, INDEF/3, 4, 5, 6, 7/, INDCL/6, 7/
C
  CALL WRRRL ('First 10 rows of the input data.', 10, NCOL, X,
&           LDX, 0, '(F5.1)', 'NUMBER', 'NUMBER')
C
  CALL SVGLM (NOBS, NCOL, X, LDX, MODEL, ILT, IRT, IFRQ, IFIX,
&           ICEN, INFIN, MAXIT, EPS, INTCEP, NCLVAR, INDCL, NEF,
&           NVEF, INDEF, INIT, IPRINT, MAXCL, NCLVAL, CLVAL,
&           NCOEF, COEF, LDCEOF, ALGL, COV, LDCOV, XMEAN, CASE,
&           LDCASE, GR, IADD, NRMISS)
C
  END

```

### Output

First 10 rows of the input data.

	1	2	3	4	5	6	7
1	411.0	0.0	7.0	64.0	5.0	1.0	0.0
2	126.0	0.0	6.0	63.0	9.0	1.0	0.0
3	118.0	0.0	7.0	65.0	11.0	1.0	0.0
4	92.0	0.0	4.0	69.0	10.0	1.0	0.0
5	8.0	0.0	4.0	63.0	58.0	1.0	0.0
6	25.0	1.0	7.0	48.0	9.0	1.0	0.0
7	11.0	0.0	7.0	48.0	11.0	1.0	0.0
8	54.0	0.0	8.0	63.0	4.0	2.0	0.0
9	153.0	0.0	6.0	63.0	14.0	2.0	0.0
10	16.0	0.0	3.0	53.0	4.0	2.0	0.0

### Initial Estimates

	1	2	3	4	5	6	7	8
	-5.054	0.000	0.000	0.000	0.000	0.000	0.000	0.000

Method	Iteration	Step size	Maximum scaled coef. update	Log likelihood
Q-N	0			-224.0
Q-N	1	1.0000	0.9839	-213.4
N-R	2	1.0000	3.603	-207.3
N-R	3	1.0000	10.12	-204.3
N-R	4	1.0000	0.1430	-204.1
N-R	5	1.0000	0.1174E-01	-204.1

Log-likelihood            -204.1392

Coefficient Statistics				
	Coefficient	Standard error	Asymptotic z-statistic	Asymptotic p-value
1	-1.103	1.309	-0.842	0.3998
2	-0.540	0.108	-4.995	0.0000
3	-0.009	0.020	-0.459	0.6460
4	-0.003	0.012	-0.291	0.7710
5	-0.363	0.445	-0.816	0.4149
6	0.127	0.486	0.261	0.7939
7	0.869	0.586	1.483	0.1385
8	0.270	0.388	0.695	0.4873

Asymptotic Coefficient Covariance					
	1	2	3	4	5
1	1.714	-8.1873E-02	-1.9753E-02	-2.2481E-03	-6.5707E-02
2		1.1690E-02	6.4506E-05	2.8955E-04	-3.8734E-04
3			3.8676E-04	-3.9067E-05	-1.2359E-03
4				1.3630E-04	7.5656E-04
5					0.1976

	6	7	8
1	-0.1038	-0.1554	-4.2370E-05
2	8.5772E-03	1.8120E-02	6.5272E-03
3	-3.2789E-04	-1.6986E-03	-2.7794E-03
4	-1.6742E-03	6.2668E-04	1.5432E-03
5	9.0035E-02	0.1122	4.3157E-02
6	0.2365	0.1142	-1.3527E-02
7		0.3436	5.1948E-02
8			0.1507

Case Analysis					
	Predicted	Influence	Residual	Cumulative Hazard	Density or Probability
1	262.7	0.0450	-0.565	1.565	0.0008
2	153.8	0.0042	0.181	0.819	0.0029
3	270.5	0.0482	0.564	0.436	0.0024
4	55.3	0.0844	-0.663	1.663	0.0034
5	61.7	0.3765	0.870	0.130	0.0142
6	230.4	0.0025	-0.108	0.108	0.8972
7	232.0	0.1960	0.953	0.047	0.0041
8	272.8	0.1677	0.802	0.198	0.0030
9	95.9	0.0505	-0.596	1.596	0.0021
10	16.8	0.0005	0.045	0.955	0.0230
11	234.0	0.1911	0.761	0.239	0.0034
12	29.1	0.0156	0.278	0.722	0.0167
13	102.2	0.4609	-1.807	2.807	0.0006
14	34.8	0.0686	0.713	0.287	0.0216
15	5.3	0.0838	-0.521	1.521	0.0415
16	25.7	0.0711	0.533	0.467	0.0244
17	65.6	0.4185	-1.698	2.698	0.0010
18	38.4	0.0886	0.688	0.312	0.0191
19	261.0	0.0155	0.234	0.766	0.0018
20	167.2	0.0338	-0.495	1.495	0.0013
21	85.8	0.0082	-0.166	1.166	0.0036
22	947.8	0.0005	-0.054	1.054	0.0004
23	105.9	0.6402	-2.181	2.181	0.1129
24	305.2	0.5757	-2.247	3.247	0.0001
25	24.6	0.3203	0.959	0.041	0.0390

26	572.8	0.0762	0.649	0.351	0.0012
27	217.5	0.1246	0.798	0.202	0.0038
28	96.6	0.1494	0.845	0.155	0.0089
29	173.4	0.1096	-0.594	0.594	0.5522
30	38.7	0.1928	0.948	0.052	0.0245
31	22.5	0.0040	0.112	0.888	0.0183
32	30.7	0.2270	-0.661	1.661	0.0062
33	20.8	0.0058	0.134	0.866	0.0202
34	54.6	0.1094	-0.648	1.648	0.0035
35	168.6	0.0923	0.502	0.498	0.0036
36	256.8	0.0341	0.361	0.639	0.0021
37	21.9	0.0069	0.134	0.866	0.0192
38	124.3	0.0680	0.654	0.346	0.0057
39	417.9	0.0087	0.186	0.814	0.0011
40	257.1	0.0025	0.101	0.899	0.0016

Last Coefficient Update

	1	2	3	4	5	6
-1.031E-05	-1.437E-06	3.098E-07	4.722E-08	-1.844E-05	-1.671E-06	
	7	8				
-2.520E-06	8.139E-06					

Covariate Means

	1	2	3	4	5	6	7
5.65	56.58	15.65	0.35	0.28	0.12	0.53	

Distinct Values For Each Class Variable

Variable 1:	1.0	2.0	3.0	4.0
Variable 2:	0.	1.0		

Observation Codes

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

Number of Missing Values 0

**Example 2**

As a second example, the MAXIT = 0 option is used for the model in Example 1 with the coefficients restricted such that  $\mu = -1.25$ ,  $\beta_1 = -.6$ , and the remaining 6 coefficients are zero. A chi-squared statistic with 8 degrees of freedom for testing that the coefficients are specified as above (versus the alternative that they are not as specified) may be computed from the output as

$$\chi^2 = g^T \hat{\Sigma}^{-1} g$$

where

$$\hat{\Sigma}$$

is output in COV, and g is output in GR. The resulting test statistic (6.107), based upon no iterations, is comparable to the likelihood ratio test statistic that may be

computed from the log-likelihood output in Example 2 (-206.6835) and the log-likelihood output in Example 1 (-204.1392).

$$\chi^2_{LR} = 2(206.6835 - 204.1392) = 5.0886$$

Neither test statistic is significant at the  $\alpha = 0.05$  level.

```

INTEGER      ICEN, IFIX, IFRQ, ILT, INFIN, INIT, INTCEP, IPAR,
&            IPRINT, IRT, LDCASE, LDCOEF, LDCEV, LDX, MAXCL,
&            MAXIT, MODEL, NCLVAR, NCOL, NEF, NOBS
REAL         EPS
PARAMETER    (EPS=0.001, ICEN=2, IFIX=0, IFRQ=0, ILT=0, INFIN=0,
&            INIT=1, INTCEP=1, IPAR=0, IPRINT=2, IRT=1, LDCASE=40,
&            LDCOEF=8, LDCEV=8, LDX=40, MAXCL=6, MAXIT=0, MODEL=0,
&            NCLVAR=2, NCOL=7, NEF=5, NOBS=40)
C
INTEGER      IADD(NOBS), INDCL(NCLVAR), INDEF(5), IRANK,
&            NCLVAL(NCLVAR), NCOEF, NRMISS, NVEF(NEF)
REAL         ALGL, CASE(LDCASE,5), CHI, CLVAL(MAXCL),
&            COEF(LDCOEF,4), COV(LDCEV,LDCEV), GR(LDCEV),
&            GRD(LDCEV), SDOT, X(LDX,NCOL), XMEAN(LDCEV)
EXTERNAL     CHFAC, GIRTS, SDOT, SSET, SVGLM
C
DATA X/411, 126, 118, 92, 8, 25, 11, 54, 153, 16, 56, 21, 287,
&    10, 8, 12, 177, 12, 200, 250, 100, 999, 231, 991, 1, 201,
&    44, 15, 103, 2, 20, 51, 18, 90, 84, 164, 19, 43, 340, 231,
&    5*0, 1, 16*0, 1, 5*0, 1, 11*0, 7, 6, 7, 4, 4, 7, 7, 8, 6,
&    3, 8, 4, 6, 4, 2, 5, 5, 4, 8, 7, 6, 9, 5, 7, 2, 8, 6, 5, 7,
&    4, 3, 3, 4, 6, 8, 7, 3, 6, 8, 7, 64, 63, 65, 69, 63, 48,
&    48, 63, 63, 53, 43, 55, 66, 67, 61, 63, 66, 68, 41, 53, 37,
&    54, 52, 50, 65, 52, 70, 40, 36, 44, 54, 59, 69, 50, 62, 68,
&    39, 49, 64, 67, 5, 9, 11, 10, 58, 9, 11, 4, 14, 4, 12, 2,
&    25, 23, 19, 4, 16, 12, 12, 8, 13, 12, 8, 7, 21, 28, 13, 13,
&    22, 36, 9, 87, 5, 22, 4, 15, 4, 11, 10, 18, 7*1, 7*2, 2*3,
&    5*4, 7*1, 4*2, 3*3, 5*4, 21*0, 19*1/
DATA NVEF/1, 1, 1, 1, 1/, INDEF/3, 4, 5, 6, 7/, INDCL/6, 7/
C
NCOEF = 8
CALL SSET (NCOEF, 0.0, COEF(3,1), 1)
CALL SSET (NOBS, 0, IADD, 1)
COEF(1,1) = -1.25
COEF(2,1) = -0.60
CALL SVGLM (NOBS, NCOL, X, LDX, MODEL, ILT, IRT, IFRQ, IFIX,
&           ICEN, INFIN, MAXIT, EPS, INTCEP, NCLVAR, INDCL, NEF,
&           NVEF, INDEF, INIT, IPRINT, MAXCL, NCLVAL, CLVAL,
&           NCOEF, COEF, LDCOEF, ALGL, COV, LDCEV, XMEAN, CASE,
&           LDCASE, GR, IADD, NRMISS)
C
                                Compute Chi-squared
CALL CHFAC (NCOEF, COV, LDCEV, EPS, IRANK, COV, LDCEV)
CALL GIRTS (NCOEF, COV, LDCEV, 1, GR, NCOEF, 2, IRANK, GRD,
&           NCOEF, COV, LDCEV)
C
CHI = SDOT(NCOEF,GRD,1,GRD,1)
WRITE (6,99999) ' Chi-squared statistic with 8 degrees of '//
&              'freedom ', CHI
C
99999 FORMAT (/, A, G12.4)
END

```

## Output

Log-likelihood            -206.6835

Coefficient Statistics				
	Coefficient	Standard error	Asymptotic z-statistic	Asymptotic p-value
1	-1.25	1.377	-0.908	0.364
2	-0.60	0.112	-5.365	0.000
3	0.00	0.021	0.000	1.000
4	0.00	0.011	0.000	1.000
5	0.00	0.429	0.000	1.000
6	0.00	0.530	0.000	1.000
7	0.00	0.775	0.000	1.000
8	0.00	0.405	0.000	1.000

Hessian					
	1	2	3	4	5
1	1.897	-8.1835E-02	-2.3464E-02	-1.1634E-03	-9.0646E-02
2		1.2507E-02	2.0883E-06	3.1320E-04	-5.3147E-04
3			4.6174E-04	-5.5344E-05	-8.1929E-04
4				1.1797E-04	6.0699E-04
5					0.1839
	6	7	8		
1	-0.1641	-0.1681	7.7768E-02		
2	1.0372E-02	1.9269E-02	5.9762E-03		
3	5.1246E-04	-1.6419E-03	-4.0106E-03		
4	-2.0693E-03	6.9029E-04	1.7020E-03		
5	9.9640E-02	0.1191	3.5786E-02		
6	0.2808	0.1264	-2.2602E-02		
7		0.6003	4.6015E-02		
8			0.1641		

Estimated Probability (censored) or Estimated Density (non-censored)								
	1	2	3	4	5	6	7	8
0.0007	0.0029	0.0026	0.0024	0.0211	0.8982	0.0041	0.0021	
0.00240	.0222	0.0021	0.0151	0.0008	0.0200	0.0433	0.0120	
0.0011	0.0190	0.0015	0.0015	0.0036	0.0004	0.0371	0.0001	
0.0792	0.0015	0.0055	0.0115	0.6424	0.0247	0.0184	0.0042	
0.0163	0.0039	0.0019	0.0021	0.0193	0.0056	0.0011	0.0016	

Newton-Raphson Step								
	1	2	3	4	5	6	7	8
0.171	0.062	-0.011	-0.003	-0.336	0.133	1.297	0.298	

Covariate Means							
	1	2	3	4	5	6	7
5.65	56.58	15.65	0.35	0.28	0.12	0.53	

Distinct Values For Each Class Variable

```

Variable 1:      1.0          2.0          3.0          4.0
Variable 2:      0.          1.0

                Observation Codes
1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20
0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0

21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40
0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0  0

Number of Missing Values          0

Chi-squared statistic with 8 degrees of freedom    6.335

```

---

## STBLE/DSTBLE (Single/Double precision)

Estimate survival probabilities and hazard rates for various parametric models.

### Usage

```
CALL STBLE (NOBS, NCOL, XPT, LDXPT, MODEL, TIME, NPT,
            DELTA, IFIX, INTCEP, NCLVAR, INDCL, NCLVAL,
            CLVAL, NEF, NVEF, INDEF, NCOEF, COEF, IPRINT,
            SPROB, LDSPRO, XBETA)
```

### Arguments

**NOBS** — Number of observations. (Input)

**NCOL** — Number of columns in XPT. (Input)

**XPT** — NOBS by NCOL matrix, each row of which contains the covariates for a group for which survival estimates are desired. (Input)

**LDXPT** — Leading dimension of XPT exactly as specified in the dimension statement of the calling program. (Input)

**MODEL** — Model option parameter. (Input)

MODEL specifies the distribution of the response variable and the relationship of the linear model to a distribution parameter.

#### MODEL Distribution

0	Exponential
1	Linear hazard
2	Log-normal
3	Normal
4	Log-logistic
5	Logistic
6	Log least extreme value
7	Least extreme value
8	Log extreme value
9	Extreme value
10	Weibull



For further discussion of the models, see the Algorithm section.

**TIME** — Beginning of the time grid for which the survival estimates are desired. (Input)

Survival probabilities and hazard rates are computed for each covariate vector over the grid of time points  $\text{TIME} + i * \text{DELTA}$  for  $i = 0, 1, \dots, \text{NPT} - 1$ .

**NPT** — Number of points on the time grid for which survival probabilities are desired. (Input)

**DELTA** — Increment between time points on the time grid. (Input)

**IFIX** — Column number in XPT containing a constant to be added to the linear response. (Input)

The estimated linear response is  $w + \text{COEF}(1) * z(1) + \text{COEF}(2) * z(2) + \dots + \text{COEF}(\text{NCOEF}) * z(\text{NCOEF})$ , where  $z$  is the design vector for the  $I$ -th observation obtained from a row of XPT.  $w = \text{XPT}(I, \text{IFIX})$  if  $\text{IFIX}$  is positive, and  $w = 0$  otherwise.

**INTCEP** — Intercept option. (Input)

**INTCEP Action**

- 0 No intercept is in the model (unless otherwise provided for by the user).
- 1 An intercept is automatically included in the model.

**NCLVAR** — Number of classification variables. (Input)

Dummy or indicator variables are generated for classification variables using the  $\text{IDUMMY} = 2$  option of routine GRGLM (page 210). See Comment 2.

**INDCL** — Index vector of length NCLVAR containing the column numbers of X that are classification variables. (Input, if NCLVAR is positive, not used otherwise)

If NCLVAR is 0, INDCL is not referenced and can be dimensioned of length 1 in the calling program.

**NCLVAL** — Vector of length NCLVAR containing the number of values taken on by each classification variable. (Input, if NCLVAR is positive, not referenced otherwise)

$\text{NCLVAL}(I)$  is the number of distinct values for the  $I$ -th classification variable. NCLVAL is not referenced and can be dimensioned of length 1 in the calling program if NCLVAR is zero.

**CLVAL** — Vector of length  $\text{NCLVAL}(1) + \text{NCLVAL}(2) + \dots + \text{NCLVAL}(\text{NCLVAR})$  containing the distinct values of the classification variables. (Input, if NCLVAR is positive; not used otherwise)

The first  $\text{NCLVAL}(1)$  elements contain the values for the first classification variables, the next  $\text{NCLVAL}(2)$  elements contain the values for the second classification variable, etc. If NCLVAR is zero, then CLVAL is not referenced and can be dimensioned of length 1 in the calling program.

**NEF** — Number of effects in the model. (Input)

In addition to effects involving classification variables, simple covariates and the product of simple covariates are also considered effects.

**NVEF** — Vector of length **NVEF** that contains the number of variables associated with each effect. (Input, if **NEF** is greater than 0; not referenced otherwise)  
**NVEF** is not referenced and can be dimensioned of length 1 in the calling program if **NEF** is zero.

**INDEF** — Vector of length  $\text{NVEF}(1) + \dots + \text{NVEF}(\text{NEF})$  that contains the column numbers in **X** associated with each effect. (Input, if **NEF** is greater than 0; not used otherwise)

The first **NVEF**(1) elements of **INDEF** contain the column numbers in **XPT** for the variables in the first effect. The next **NVEF**(2) elements in **INDEF** contain the column numbers for the second effect, etc.. If **NCLVAR** is zero, **INDEF** is not referenced and can be dimensioned of length 1 in the calling program.

**NCOEF** — Number of coefficients in the model. (Input)

**COEF** — Vector of length **NCOEF** containing the model parameter estimates. (Input)

Usually routine **SVGLM** (page 967) is first called to estimate **COEF** as the first column of matrix **COEF** in **SVGLM**. When present in the model, the initial coefficient in **COEF** is a “nuisance” parameter, and the remaining coefficients are parameters associated with the “linear” model, beginning with the intercept, if present. Nuisance parameters are as follows:

**Model Nuisance Parameter**

- |     |   |
|-----|---|
| 1   | Coefficient of the quadratic term in time, $\theta$ |
| 2–9 | Scale parameter, $\sigma$                           |
| 10  | Shape parameter, $\theta$                           |

There is no nuisance parameter for model 0.

**IPRINT** — Printing option. (Input)

**IPRINT Action**

- |   |                           |
|---|---------------------------|
| 0 | No printing is performed. |
| 1 | Printing is performed.    |

**SPROB** — **NPT** by  $2 * \text{NOBS} + 1$  matrix. (Output)

**SPROB**(*i*, 2) contains the estimated survival probability at time

**SPROB**(*i*, 1) = **TIME** + (*i* – 1) \* **DELTA** for observations with covariates given in row 1 of **XPT**. **SPROB**(*i*, 3) contains the estimate for the hazard rate at this time point. Columns 4 and 5 contain the estimated survival probabilities and hazard rates for observations with covariates given in the second row in **XPT**, etc., up to columns  $2 * \text{NOBS}$  and  $2 * \text{NOBS} + 1$ , which contain these statistics for observations with covariates in the last row of **XPT**.

**LDSPRO** — Leading dimension of **SPROB** exactly as specified in the dimension statement in the calling program. (Input)

***XBETA*** — Vector of length NOBS containing the estimated linear response  $w + \text{COEF}(1) * z(1) + \dots + \text{COEF}(\text{NCOEF}) * z(\text{NCOEF})$  for each row of XPT. (Output)

### Comments

- Automatic workspace usage is

STBLE NCOEF numeric units if IPRINT is zero, or NCOEF + max(7, NCOL) numeric units + NCOL \* 10 character units if IPRINT is 1, or

DSTBLE 2 \* NCOEF numeric units if IPRINT is zero, or 2 \* NCOEF + max(7, NCOL) numeric units + NCOL \* 10 character units if IPRINT is 1.

Workspace may be explicitly provided, if desired, by use of S2BLE/DS2BLE. The reference is

```
CALL S2BLE (NOBS, NCOL, XPT, LDXPT, MODEL, TIME,
           NPT, DELTA, IFIX, INTCEP, NCLVAR, INDCL,
           NCLVAL, CLVAL, NEF, NVEF, INDEF, NCOEF,
           COEF, IPRINT, SPROB, LDSPRO, XBETA,
           CHWK, Z, RWK)
```

The additional arguments are as follows:

***CHWK*** — CHARACTER \* 10 work vector of length NCOL.

***Z*** — Work vector of length NCOEF.

***RWK*** — Work vector of length MAX(7, NCOL) if IPRINT = 1, or of length 1 if IPRINT = 0.

- Dummy variables are generated for the classification variables as follows: The list of all distinct values of each classification variable is as stored in CLVAL. Dummy variables are generated for each but the last of these distinct values. Each dummy variable is zero unless the classification variable equals the list value corresponding to the dummy variable, in which case the dummy variable is one. See argument IDUMMY for IDUMMY = 2 in routine GRGLM (page 210).

- Informational errors

Type	Code	Description
3	1	Some survival probabilities are less than or equal to zero. The corresponding hazard values cannot be computed.
4	2	The specified number of coefficients, NCOEF, is incorrect.
4	3	The model specified is not defined for negative time.

### Algorithm

Routine STBLE computes estimates of survival probabilities and hazard rates for the parametric survival/reliability models fit by routine SVGLM (page 967) for one or more vectors of covariate values. Because estimates for the parameters of the model must be given, routine SVGLM is usually invoked to obtain these estimates prior to invoking STBLE.

Let  $\eta = x^T \beta$  be the linear parameterization, where  $x$  is a design vector obtained in STBLE via routine GRGLM (page 210) from a row of XPT, and  $\beta$  is a vector of parameters associated with the linear model. Let  $T$  denote the random response variable and  $S(t)$  denote the probability that  $T > t$ . All models considered also allow a fixed parameter  $w$  (input in column IFIX of XPT). Use of this parameter is discussed in the document for routine SVGLM. There may also be nuisance parameters  $\theta > 0$ , or  $\sigma > 0$ . Let  $\Phi$  denote the cumulative normal distribution. The survival models available in STBLE are

Model	Name	$S(t)$
0	Exponential	$\exp\{-t \exp(w + \eta)\}$
1	Linear hazard	$\exp\{-(t + \frac{\theta t^2}{2}) \exp[(w + \eta)]\}$
2	Log-normal	$1 - \Phi(\frac{\ln(t) - \eta - w}{\sigma})$
3	Normal	$1 - \Phi(\frac{t - \eta - w}{\sigma})$
4	Log-logistic	$\{1 + \exp(\frac{\ln(t) - \eta - w}{\sigma})\}^{-1}$
5	Logistic	$\{1 + \exp(\frac{t - \eta - w}{\sigma})\}^{-1}$
6	Log least extreme value	$\exp\{-\exp(\frac{\ln(t) - \eta - w}{\sigma})\}$
7	Least extreme value	$\exp\{-\exp(\frac{t - \eta - w}{\sigma})\}$
8	Log extreme value	$1 - \exp\{-\exp[-(\frac{\ln(t) - \eta - w}{\sigma})]\}$
9	Extreme value	$1 - \exp\{-\exp[-(\frac{t - \eta - w}{\sigma})]\}$
10	Weibull	$\exp\{-\left(\frac{t}{\exp(w + \eta)}\right)^\theta\}$

Let  $\lambda(t)$  denote the hazard rate at time  $t$ . Then  $\lambda(t)$  and  $S(t)$  are related as

$$S(t) = \exp\left\{-\int_{-\infty}^t \lambda(s) ds\right\}$$

Models 0, 1, 2, 4, 6, 8, and 10 require that  $T > 0$  (in which case, we assume  $\lambda(s) = 0$  for  $s < 0$ ), while the remaining models allow arbitrary values for  $T$ ,  $-\infty < T < \infty$ . The computations proceed in routine STBLE as follows:

1. The input arguments are checked for consistency and validity.

- For each row of XPT, the explanatory variables are generated from the classification and variables and the covariates using routine GRGLM with the IDUMMY = 2 option. (When IDUMMY is two, GRGLM assigns an indicator variable the value 1.0 when the observation is in the class, assigns the value 0.0 otherwise, and omits the last indicator variable from the design vector. See the manual documentation for GRGLM.)

Given the explanatory variables  $x$ ,  $\eta$  is computed as  $\eta = x^T \beta$ , where  $\beta$  in input in COEF.

- For each time point requested in the time grid, the survival probabilities and hazard rates are computed.

### Example

The example is a continuation of the first example given for routine SVGLM (page 967). Prior to calling STBLE, SVGLM is invoked to compute the parameter estimates. The example is taken from Lawless (1982, page 287) and involves the mortality of patients suffering from lung cancer.

```

C
INTEGER      ICEN, IFIX, IFRQ, ILT, INFIN, INIT, INTCEP, IPRINT,
&            IRT, ITIE, LDCASE, LDCCOEF, LDCCOV, LDSPRO, LDX, LDXPT,
&            MAXCL, MAXIT, MODEL, NCLVAR, NCOL, NEF, NOBS, NPT,
&            NOBS
REAL         DELTA, EPS, TIME, XPWR
PARAMETER   (DELTA=20.0, EPS=0.001, ICEN=2, IFIX=0, IFRQ=0,
&            ILT=0, INFIN=0, INIT=0, INTCEP=1, IPRINT=1, IRT=1,
&            ITIE=0, LDCASE=40, LDCCOEF=9, LDCCOV=9, LDX=40,
&            LDXPT=2, MAXCL=6, MAXIT=15, MODEL=0, NCLVAR=2,
&            NCOL=7, NEF=5, NOBS=40, NPT=10, NOBS=2, TIME=10.0,
&            XPWR=0.0, LDSPRO=NPT)

C
INTEGER      IADD(NOBS), INDCL(NCLVAR), INDEF(5), NCLVAL(NCLVAR),
&            NCOEF, NRMISS, NVEF(NEF)
REAL         ALGL, CASE(LDCASE,5), CLVAL(MAXCL), COEF(LDCCOEF,4),
&            COV(LDCCOV,LDCCOV), GR(LDCCOV), SPROB(LDSPRO,2*NOBS+1),
&            X(LDX,NCOL), XBETA(NOBS), XMEAN(LDCCOV),
&            XPT(LDXPT,NCOL)
EXTERNAL    SCOPY, STBLE, SVGLM

C
DATA X/411, 126, 118, 92, 8, 25, 11, 54, 153, 16, 56, 21, 287,
&      10, 8, 12, 177, 12, 200, 250, 100, 999, 231, 991, 1, 201,
&      44, 15, 103, 2, 20, 51, 18, 90, 84, 164, 19, 43, 340, 231,
&      5*0, 1, 16*0, 1, 5*0, 1, 11*0, 7, 6, 7, 4, 4, 7, 7, 8, 6,
&      3, 8, 4, 6, 4, 2, 5, 5, 4, 8, 7, 6, 9, 5, 7, 2, 8, 6, 5, 7,
&      4, 3, 3, 4, 6, 8, 7, 3, 6, 8, 7, 64, 63, 65, 69, 63, 48,
&      48, 63, 63, 53, 43, 55, 66, 67, 61, 63, 66, 68, 41, 53, 37,
&      54, 52, 50, 65, 52, 70, 40, 36, 44, 54, 59, 69, 50, 62, 68,
&      39, 49, 64, 67, 5, 9, 11, 10, 58, 9, 11, 4, 14, 4, 12, 2,
&      25, 23, 19, 4, 16, 12, 12, 8, 13, 12, 8, 7, 21, 28, 13, 13,
&      22, 36, 9, 87, 5, 22, 4, 15, 4, 11, 10, 18, 7*1, 7*2, 2*3,
&      5*4, 7*1, 4*2, 3*3, 5*4, 21*0, 19*1/
DATA NVEF/1, 1, 1, 1, 1/, INDEF/3, 4, 5, 6, 7/, INDCL/6, 7/

C
CALL SVGLM (NOBS, NCOL, X, LDX, MODEL, ILT, IRT, IFRQ, IFIX,
&          ICEN, INFIN, MAXIT, EPS, INTCEP, NCLVAR, INDCL, NEF,

```

```

&          NVEF, INDEF, INIT, 0, MAXCL, NCLVAL, CLVAL, NCOEF,
&          COEF, LDcoef, ALGL, COV, LDCOV, XMEAN, CASE, LDCASE,
&          GR, IADD, NRMISS)
C
CALL SCOPY (NCOL, X(1,1), LDX, XPT(1,1), LDXPT)
CALL SCOPY (NCOL, X(2,1), LDX, XPT(2,1), LDXPT)
C
CALL STBLE (NOBS, NCOL, XPT, LDXPT, MODEL, TIME, NPT, DELTA,
&          IFIX, INTCEP, NCLVAR, INDCL, NCLVAL, CLVAL, NEF,
&          NVEF, INDEF, NCOEF, COEF, IPRINT, SPROB, LDSPRO,
&          XBETA)
C
END

```

### Output

```

              group  1
              xpt
  1          2          3          4          5          6
411         0          7         64          5          1

7
0

```

```

              design vector
  1          2          3          4          5          6
  1          7          64          5          1          0

7          8
0          1

```

xbeta = -5.57097

```

              group  2
              xpt
  1          2          3          4          5          6
126         0          6          63          9          1

7
0

```

```

              design vector
  1          2          3          4          5          6
  1          6          63          9          1          0

7          8
0          1

```

xbeta = -5.03551

### survival and hazard estimates (sprob)

time	s1	h1	s2	h2
10.00	0.9626	0.003807	0.9370	0.006503
30.00	0.8921	0.003807	0.8228	0.006503
50.00	0.8267	0.003807	0.7224	0.006503
70.00	0.7661	0.003807	0.6343	0.006503
90.00	0.7099	0.003807	0.5570	0.006503
110.00	0.6579	0.003807	0.4890	0.006503

130.00	0.6096	0.003807	0.4294	0.006503
150.00	0.5649	0.003807	0.3770	0.006503
170.00	0.5235	0.003807	0.3310	0.006503
190.00	0.4852	0.003807	0.2907	0.006503

Note that in simple exponential models the hazard rate is constant over time.

---

## ACTBL/DACTBL (Single/Double precision)

Produce population and cohort life tables.

### Usage

```
CALL ACTBL (IMTH, N, NPOP, AGE, A, IPOP, IDTH, IPRINT,
           TABLE, LDTABL)
```

### Arguments

**IMTH** — Type of life table. (Input)

IMTH = 0 indicates a population (current) table. IMTH = 1 indicates a cohort table.

**N** — Number of age classes. (Input)

**NPOP** — Population size. (Input, if IMTH = 0; not used otherwise)

For IMTH = 0, the population size at the beginning of the first age interval. The value is somewhat arbitrary. NPOP = 10000 is reasonable. Not used if IMTH = 1.

**AGE** — Vector of length N + 1 containing the lowest age in each age interval, and in AGE(N + 1), the endpoint of the last age interval. (Input)

Negative AGE(1) indicates that the age intervals are all of length |AGE(1)| and that the initial age interval is from 0.0 to |AGE(1)|. In this case, all other elements of AGE need not be specified. AGE(N + 1) need not be specified when IMTH = 1.

**A** — Vector of length N containing the fraction of those dying within each interval who die before the interval midpoint. (Input)

A common choice for all A(I) is 0.5. This choice may also be specified by setting A(1) to any negative value. In this case, the remaining values of A need not be specified.

**IPOP** — Vector of length N containing the cohort sizes during each interval. (Input)

If IMTH = 0, then IPOP(I) contains the size of the population at the midpoint of interval I. If IMTH = 1, then IPOP(I) contains the size of the cohort at the beginning of interval I. When IMTH = 0, the population sizes in IPOP may need to be adjusted to correspond to the number of deaths in IDTH. See the algorithm section of the document for more information.

**IDTH** — Vector of length N containing the number of deaths in each age interval. (Input, if IMTH = 0; not used otherwise)

If IMTH = 1, IDTH is not used and may be dimensioned of length 1.

**IPRINT** — Printing option. (Input)

If `IPRINT = 1`, the life table is printed. Otherwise, no printing is done.

**TABLE** —  $N$  by 12 matrix containing the life table. (Output)

The rows of `TABLE` correspond to the age intervals.

**Col. Description**

- 1 Lowest age in the age interval.
- 2 Fraction of those dying within the interval who die before the interval midpoint.
- 3 Number surviving to the beginning of the interval.
- 4 Number of deaths in the interval.
- 5 Death rate in the interval. If `IMTH = 1`, this column is set to NaN (not a number).
- 6 Proportion dying in the interval.
- 7 Standard error of the proportion dying in the interval.
- 8 Proportion of survivors at the beginning of the interval.
- 9 Standard error of the proportion of survivors at the beginning of the interval.
- 10 Expected lifetime at the beginning of the interval.
- 11 Standard error of the expected life at the beginning of the interval.
- 12 Total number of time units lived by all of the population in the interval.

**LDTABLE** — Leading dimension of `TABLE` exactly as specified in the dimension statement in the calling program. (Input)

**Algorithm**

Routine `ACTBL` computes population (current) or cohort life tables based upon the observed population sizes at the middle (`IMTH = 0`) or the beginning (`IMTH = 1`) of some userspecified age intervals. The number of deaths in each of these intervals must also be observed.

The probability of dying prior to the middle of the interval, given that death occurs somewhere in the interval, may also be specified. Often, however, this probability is taken to be 0.5. For a discussion of the probability models underlying the life table here, see the references.

Let  $t_i$ , for  $i = 0, 1, \dots, t_n$  denote the time grid defining the  $n$  age intervals, and note that the length of the age intervals may vary. Following Gross and Clark (1975, page 24), let  $d_i$  denote the number of individuals dying in age interval  $i$ , where age interval  $i$  ends at time  $t_i$ . If `IMTH = 0`, the death rate at the middle of the interval is given by  $r_i = d_i / (M_i h_i)$ , where  $M_i$  is the number of individuals alive at the middle of the interval, and  $h_i = t_i - t_{i-1}$ ,  $t_0 = 0$ . The number of individuals alive at the beginning of the interval may be estimated by  $P_i = M_i + (1 - a_i)d_i$  where  $a_i$  is the probability that an individual dying in the interval dies prior to the interval midpoint. When `IMTH = 1`,  $P_i$  is input directly while the death rate in the interval,  $r_i$ , is not needed.



The probability that an individual dies during the age interval from  $t_{i-1}$  to  $t_i$  is given by  $q_i = d_i/P_i$ . It is assumed that all individuals alive at the beginning of the last interval die during the last interval. Thus,  $q_n = 1.0$ . The asymptotic variance of  $q_i$  can be estimated by

$$\sigma_i^2 = q_i(1 - q_i) / P_i$$

When  $\text{IMTH} = 0$ , the number of individuals alive in the middle of the time interval (input in  $\text{IPOP}(\text{I})$ ) must be adjusted to correspond to the number of deaths observed in the interval. Routine  $\text{ACTBL}$  assumes that the number of deaths observed in interval  $h_i$  occur over a time period equal to  $h_i$ . If  $d_i$  is measured over a period  $u_i$ , where  $u_i \neq h_i$ , then  $\text{IPOP}(\text{I})$  must be adjusted to correspond to  $d_i$  by multiplication by  $u_i/h_i$ , i.e., the value  $M_i$  input into  $\text{ACTBL}$  as  $\text{IPOP}(\text{I})$  is computed as

$$M_i^* = M_i u_i / h_i$$

Let  $S_i$  denote the number of survivors at time  $t_i$  from a hypothetical ( $\text{IMTH} = 0$ ) or observed ( $\text{IMTH} = 1$ ) population. Then,  $S_0 = \text{NPOP}$  when  $\text{IMTH} = 0$ , and  $S_0 = \text{IPOP}(1)$  for  $\text{IMTH} = 1$ , and  $S_i$  is given by  $S_i = S_{i-1} - \delta_{i-1}$  where  $\delta_i = S_i q_i$  is the number of individuals who die in the  $i$ -th interval. The proportion of survivors in the interval is given by  $V_i = S_i/S_0$  while the asymptotic variance of  $V_i$  can be estimated as follows.

$$\text{var}(V_i) = V_i^2 \sum_{j=1}^{i-1} \frac{\sigma_j^2}{(1 - q_j)^2}$$

The expected lifetime at the beginning of the interval is calculated as the total lifetime remaining for all survivors alive at the beginning of the interval divided by the number of survivors at the beginning of the interval. If  $e_i$  denotes this average expected lifetime, then the variance of  $e_i$  can be estimated as (see Chiang 1968)

$$\text{var}(e_i) = \frac{\sum_{j=i}^{n-1} P_j^2 \sigma_j^2 [e_{j+1} + h_{j+1}(1 - a_j)]^2}{P_i^2}$$

where  $\text{var}(e_n) = 0.0$ .

Finally, the total number of time units lived by all survivors in the time interval can be estimated as:

$$U_i = h_i [S_i - \delta_i(1 - a_i)]$$

### Example

The following example is taken from Chiang (1968). The cohort life table has thirteen equally spaced intervals, so  $\text{AGE}(1)$  is set to  $-5.0$ . Similarly, the

probabilities of death prior to the middle of the interval are all taken to be 0.5, so A(1) is set to -1.0. Since IPRINT = 1, the life table is printed by ACTBL.

```

C                               Specifications
INTEGER      IMTH, IPRINT, LDTABL, N, NPOP
PARAMETER    (IMTH=1, IPRINT=1, N=13, NPOP=10000, LDTABL=N)

C
INTEGER      IDTH(13), IPOP(13)
REAL         A(1), AGE(1), TABLE(13,12)
EXTERNAL     ACTBL

C
DATA AGE/-5.0/, A/-1.0/
DATA IPOP/270, 268, 264, 261, 254, 251, 248, 232, 166, 130, 76,
&      34, 13/

C
CALL ACTBL (IMTH, N, NPOP, AGE, A, IPOP, IDTH, IPRINT, TABLE,
&          LDTABL)

C
END

```

### Output

Life Table					
Age Class	Age	PDHALF	Alive	Deaths	Death Rate
1	0	0.5	270	2	NaN
2	5	0.5	268	4	NaN
3	10	0.5	264	3	NaN
4	15	0.5	261	7	NaN
5	20	0.5	254	3	NaN
6	25	0.5	251	3	NaN
7	30	0.5	248	16	NaN
8	35	0.5	232	66	NaN
9	40	0.5	166	36	NaN
10	45	0.5	130	54	NaN
11	50	0.5	76	42	NaN
12	55	0.5	34	21	NaN
13	60	0.5	13	13	NaN

Age Class	P(D)	Std(P(D))	P(S)	Std(P(S))	Lifetime
1	0.007	0.00522	1.000	0.00000	43.19
2	0.015	0.00741	0.993	0.00522	38.49
3	0.011	0.00652	0.978	0.00897	34.03
4	0.027	0.01000	0.967	0.01092	29.40
5	0.012	0.00678	0.941	0.01437	25.14
6	0.012	0.00686	0.930	0.01557	20.41
7	0.065	0.01560	0.919	0.01665	15.62
8	0.284	0.02962	0.859	0.02116	11.53
9	0.217	0.03199	0.615	0.02962	10.12
10	0.415	0.04322	0.481	0.03041	7.23
11	0.553	0.05704	0.281	0.02737	5.59
12	0.618	0.08334	0.126	0.02019	4.41
13	1.000	0.00000	0.048	0.01303	2.50

Age Class	Std(Life)	Time Units
1	0.6993	1345.0
2	0.6707	1330.0
3	0.6230	1312.5
4	0.5940	1287.5
5	0.5403	1262.5

6	0.5237	1247.5
7	0.5149	1200.0
8	0.4982	995.0
9	0.4602	740.0
10	0.4328	515.0
11	0.4361	275.0
12	0.4167	117.5
13	0.0000	32.5

# Chapter 14: Multidimensional Scaling

---

## Routines

<b>14.1. Multidimensional Scaling Routines</b>		
Individual differences model .....	MSIDV	1003
<b>14.2. Utility Routines</b>		
Compute distance matrices based upon a model.....	MSDST	1017
Standardize the input data .....	MSSTN	1020
Double center a dissimilarity matrix .....	MSDBL	1024
Compute initial estimates.....	MSINI	1028
Compute stress given disparities and distances.....	MSTRS	1035

---

## Usage Notes

The routines described in this chapter all involve multidimensional scaling. Routine `MSIDV` (page 1003) performs computations for the individual differences metric scaling models. The utility routines are useful for associated computations as well as for programming other methods of multidimensional scaling.

The following is a brief introduction to multidimensional scaling meant to acquaint the user with the purposes of the routines described in this chapter. Also of interest is the table at the end of this section giving the notation used. A more complete description of procedures in multidimensional scaling may be found in the references, as well as in the algorithm sections for the routines.

### Multidimensional Scaling Data Types

A “dissimilarity” is a subject’s measure of the “distance” between two objects. For example, a subject’s estimate of the distance between two cities is a dissimilarity measure that may, or may not, be the actual distance between the cities (depending upon the subjects familiarity with the two cities). Dissimilarities usually have less relationship to distance. For example, the

subject may estimate, on a given scale, the difference between two smells, two tastes, two colors, two shapes, etc. As a concrete example, the subject is asked to compare two wines and indicate whether they have very similar tastes (scale value 0), or very different tastes (scale value 10), or are somewhere in between. In this case, no objective measure of “distance” is available, yet the dissimilarity may be measured. In all cases, however, the larger the difference between the objects, the larger the dissimilarity measure.

If instead the measure increases as the objects become more similar, then a “similarity” measure rather than a “dissimilarity” measure is obtained. Most routines in this chapter require dissimilarities as input so that similarities must be converted to dissimilarities before most routines in this chapter can be used. Routine `MSSTN` (page 1020) provides two common methods for performing these conversions.

In general, dissimilarities between all objects in a set are measured (yielding a matrix of dissimilarities), and the multidimensional scaling problem is to locate the objects in a Euclidean (or other) space of known dimension given the matrix of dissimilarities. The estimates of object locations should yield predicted distances between the objects that “closely approximate” the observed dissimilarities. In many multidimensional scaling methods, “closely approximates” means that a predefined measure of the discrepancy (the “stress”) is minimized. The simplest stress measure is the sum of the squared differences between the observed dissimilarities and the distances predicted by the estimated object locations. This stress measure, as well as all other stress measures used in this chapter, is discussed more fully in the manual document for routine `MSTRS` (page 1035).

Note that the predicted distances between objects may not be Euclidean distance. Indeed, in one of the more popular multidimensional scaling models, the individual differences model, weighted Euclidean distance is used. Let  $\lambda_{1k}$  and  $\lambda_{2k}$ ,  $k = 1, \dots, d$ , be the location estimates of two objects (stimuli) in a  $d$  dimensional space. Then, the weighted Euclidean distance used in the individual difference model is given by

$$\delta_{12} = \sqrt{\sum_{k=1}^d w_k (\lambda_{1k} - \lambda_{2k})^2}$$

Many other distance models are possible. The models used in this chapter are discussed in the manual document for routine `MSDST` (page 1017).

A dissimilarity is a subject’s estimate of the difference (“distance”) between two objects. From the observed dissimilarities, a predicted distance between the objects is obtained by estimating the location of the objects in a Euclidean space of given dimension. In metric scaling, the dissimilarity may be a ratio measure (in which case a dissimilarity of zero means that the objects are in the same location) or an interval measure (in which case “distance” plus a constant is observed). When an interval measure is observed, the interval constant,  $c$ , must also be estimated in order to relate the dissimilarity to the predicted distance.

For ratio measures,  $c$  is not required. A couple of methods for estimating  $c$  are used by the routines in this chapter. These methods are explained in the routines that use them.

In nonmetric scaling, the dissimilarity is an ordinal (rank) or categorical measure. In this case, the stress function need only assure that the predicted distances satisfy, as closely as possible, the ordinal or categorical relationships observed in the data. Thus, the stress should be zero if the predicted distances maintain the observed rankings in the dissimilarities in ordinal data. The meaning of a stress in categorical data is more obtuse and is discussed further below.

In ordinal data, the stress function is computed as follows: First, the dissimilarities are transformed so that they correspond as closely as possible to the predicted distances, but such that the observed ordinal relationships are maintained. The transformed dissimilarities are called “disparities”, and the stress function is computed from the disparities and the predicted distances. (In ratio and interval data, disparities may be taken as the dissimilarities.) Thus, if the predicted distances preserve the observed ordinal relationships, a stress of zero will be computed. If the predicted distances do not preserve these relationships, then new estimates for the distances based upon the disparities can be computed. These can be followed by new estimates of the disparities. When the new estimates do not lead to a lower stress, convergence of the algorithm is assumed.

In categorical data, all that is observed is a category for the “distance” between the objects, and there are no known relationships between the categories. In categorical data, the disparities are such that the categories are preserved. A score minimizing the stress is found for each category. As with ordinal data, new distances are computed from this score, followed by new scores for the categories, etc., with convergence occurring when the stress cannot be lowered further. In categorical data, a stress of zero should be relatively uncommon.

The individual differences model assumes that the squared distance between stimuli  $i$  and  $j$  for subject  $l$ ,

$$\delta_{ijl}^2$$

is given as

$$\delta_{ijl}^2 = \sum_{k=1}^d w_{lk} (\lambda_{ik} - \lambda_{jk})^2$$

where  $d$  is the number of dimensions (always assumed to be known),  $\lambda_{ik}$  is the location of the  $i$ -th stimulus in the  $k$ -th dimension, and  $w_{lk}$  is the weight given by subject  $l$  to the  $k$ -th dimension. Let

$$\bar{\delta}_{i \bullet l}^2$$

denote the average of the squared distances in the  $i$ -th row of the dissimilarity matrix for the  $l$ -th subject, let

$$\bar{\delta}_{\bullet j l}^2$$

be similarly defined for the  $j$ -th column, and let

$$\bar{\delta}_{\bullet \bullet l}^2$$

denote the average of all squared distances for the  $l$ -th subject. Then, the product moment (double centering) transformation is given by

$$p_{ijl} = -(\delta_{ijl}^2 - \bar{\delta}_{i \bullet l}^2 - \bar{\delta}_{\bullet j l}^2 + \bar{\delta}_{\bullet \bullet l}^2) / 2.0$$

The advantage of the product-moment transformations is that the “product-moment” (double centered) matrices  $P_l = (p_{ijl})$  can be expressed as

$$P_l = \Lambda[\text{diag}(W_l)]\Lambda^T$$

where  $\Lambda = (\lambda_{ik})$  is the configuration matrix, and where  $\text{diag}(W_l)$  is a diagonal matrix with the subject weights for subject  $l$ ,  $w_{lk}$ , along the diagonal. If one assumes that the dissimilarities are measured without error, then the dissimilarities can be used in place of the distances, and the above relationship allows one to compute both  $\text{diag}(W_l)$  and  $\Lambda$  directly from the product-moment matrices so obtained. If error is present but small, then very good estimates of  $\Lambda$  and  $\text{diag}(W_l)$  can still be obtained (see De Leeuw and Pruzansky 1978). Routine `MSDBL` (page 1024) computes the product-moment matrices while `MSINI` (page 1028) computes the above estimates for  $X$  and  $\text{diag}(W_l)$ .

## Data Structures

The data input to a multidimensional scaling routine is, conceptually, one or more dissimilarity (or similarity) matrices where a dissimilarity matrix contains the dissimilarity measure between the  $i$ -th and  $j$ -th stimuli (objects) in position  $(i, j)$  of the matrix. In multidimensional scaling, the dissimilarity matrix need not be symmetric (asymmetric distances can also be modelled, see routine `MSDST`, page 1017) but if it is, only elements above the diagonal need to be observed. Moreover, in the multidimensional “unfolding” models, the distances between all pairs of objects are not observed. Rather, all (or at least many) of the dissimilarities between one set of objects and a second set are measured. When these types of input are combined with the fact that missing values are also allowed in many multidimensional scaling routines, it is easy to see that data structures required in multidimensional scaling can be quite complicated. Three types of structures are allowed for the routines described in this chapter. These are discussed below.

Let  $x$  denote a matrix containing the input dissimilarities. The columns of  $x$  correspond to the different subjects, and a subjects dissimilarity matrix is contained within the column. Thus,  $x$  is a matrix containing a set of

dissimilarity matrices, one dissimilarity matrix within each column. For any one problem, the form (structure) of all dissimilarity matrices input in  $x$  must be consistent over all subjects. The form can vary from problem to problem, however. In the following,  $x$  contains only one column and the index for subject is ignored to simplify the notation. The three storage forms used by the routines described in this chapter are

1. **Square symmetric:** For this form, each column of  $x$  contains the upper triangular part of the dissimilarity matrix, excluding the diagonal elements (which should be zero anyway). Specifically,  $x(1)$  contains the (1, 2) element of the dissimilarity matrix,  $x(2)$  contains the (1, 3) element,  $x(3)$  contains the (2, 3) element, etc. Let  $q$  denote the number of stimuli in the matrix. All  $q(q - 1)/2$  off-diagonal elements are stored.
2. **Square asymmetric:**  $x$  contains all elements of each square matrix, including the diagonal elements, which are not used. The dissimilarities are stored in  $x$  as if  $x$  were dimensioned  $q \times q$ . The diagonal elements are ignored.
3. **Rectangular:** This corresponds to the “unfolding models” in which not all of the dissimilarities in each matrix are observed. In this storage mode, the row stimuli do not correspond to the column stimuli. Because of the form of the data, no diagonal elements are present, and the data are stored in  $x$  as if  $x$  were dimensioned  $r \times s$  where  $r$  is the number of row stimuli and  $s$  is the number of column stimuli.

Missing values are also allowed. They are indicated in  $x$  in either of two ways: 1) The standard IMSL missing value indicator NaN (not a number) may be used to indicate missing values, or 2) negative elements of  $x$  are taken to be missing dissimilarities.

Table 14.1 gives some notation commonly used in this chapter. In general, an element of a matrix is denoted by the lowercase matrix name with subscripts. The notation is generally consistent, but there are some variations when variation seems appropriate.



**Table 14.1: Commonly Used Notation**

Symbol	Fortran	Meaning
$\delta_{ijl}$	DIST	Distance between objects $i$ and $j$ for subject $l$ .
$\delta_{ijl}^*$	DISP	Disparity for objects $i$ and $j$ for subject $l$ .
$X$	X	The input array of dissimilarities.
$d$	NDIM	The number of dimensions in the solution.
$W$	W	The matrix of subject weights.
$\text{diag}(W_l)$		The diagonal matrix of subject weights for subject $l$ .
$\pi$	WS	The matrix of stimulus weights.
$\Lambda$	CFL	The configuration matrix.
$\alpha_h$	A	The intercept for strata $h$ .
$\beta_h$	B	The slope for strata $h$ .
$v_h$	WT	The stratum weight for stratum $h$ .
$n_h$	NCOM	The number nonmissing dissimilarities in stratum $h$ .
$P_l$	P	The product-moment matrix for subject $l$ .
$\phi$	STRSS	The stress criterion (over all strata).
$\phi_l$	STRS	The stress within stratum $l$ .
$p$	POWER	The power to use in the stress criterion.
$q$	NSTIM	The total number of stimuli.
$\eta$	NSUB	The number of matrices input.
$\Gamma$		Normalized eigenvectors.
	IFORM	Option giving the form of the dissimilarity input.
	ICNVT	Option giving the method for converting to dissimilarities.
	MODEL	Vector giving the parameters in the distance model.
	ISTRS	Option giving the stress formula to use.
	ITRANS	Option giving the transformation to use.
	IDISP	The method to be used in estimating disparities.
	EPS	Convergence tolerance.

---

## MSIDV/DMSIDV (Single/Double precision)

Perform individual-differences multidimensional scaling for metric data using alternating least squares.

### Usage

```
CALL MSIDV (NSTIM, NSUB, X, ICNVT, MODEL, ISTRS, ITRANS,  
           NDIM, IPRINT, DIST, CFL, LDCFL, W, LDW, A, B,  
           WT, STRS, STRSS, RESID)
```

### Arguments

**NSTIM** — Number of stimuli in each similarity/dissimilarity matrix. (Input)

**NSUB** — Number of matrices to be used in the analysis. (Input)

**X** — NSUB similarity or dissimilarity matrices in symmetric storage mode. (Input)

Each matrix must occupy consecutive memory positions, and must be stored as a column in X. X must be dimensioned as

```
DIMENSION X(NC2, NSUB)
```

where  $NC2 = NSTIM * (NSTIM - 1)/2$ . Each matrix is stored without the diagonal elements by column as upper triangular matrices. For example, a 3 by 3 matrix would be stored with the (1, 2), (1, 3), (2, 3) elements as the first three elements of the first column of X.

**ICNVT** — Option for converting from similarity to dissimilarity data. (Input)

If **ICNVT** = 0, the input data contains dissimilarities and no conversion is performed. If **ICNVT** = 1, the data are converted from similarity to dissimilarity data by subtracting each similarity from the largest similarity for the subject. If **ICNVT** = 2, the data are converted to dissimilarities by reciprocating each similarity.

**MODEL** — Model option parameter. (Input)

**MODEL** = 0 means the Euclidean model is used, otherwise, the individual differences model is used.

**ISTRS** — Option giving the stress formula to be used. (Input)

Stress formulas differ in the weighting given to each subject. The valid values of **ISTRS** are:

#### **ISTRS** Weighting used

- |   |  |
|---|--|
| 0 | Inverse of within-subject variance of observed dissimilarities about the predicted distances |
| 1 | Inverse of within-subject sum of squared dissimilarities                                     |
| 2 | Inverse of within-subject variance of dissimilarities about the subject mean                 |

See the Algorithm section for further discussion of the stress formula weights.

**ITRANS** — Option giving the transformation to be used on the observed and predicted dissimilarities when computing the criterion function. (Input)

**ITRANS Transformation**

- 0 Squared distances
- 1 Distances (that is, no transformation is performed)
- 2 Log of the distances

See the Algorithm section for further discussion of stress formula transformations.

**NDIM** — Number of dimensions desired in the solution. (Input)

**IPRINT** — Printing option. (Input)

**IPRINT Action**

- 0 No printing is performed.
- 1 Printing is performed, but the output is abbreviated.
- 2 All printing is performed.

**DIST** — Vector of length  $NSUB * NC2$ , where  $NC2 = NSTIM * (NSTIM - 1)/2$ , containing the predicted distances. (Output)

**DIST** contains the distances as predicted by the estimated parameters in the model. **DIST** has the same storage mode as **X** and may be treated as a series of **NSUB** matrices in symmetric storage mode but without the diagonal elements.

**CFL** — Matrix of size **NSTIM** by **NDIM** containing the configuration of points obtained from the multidimensional scaling. (Output)

**LDCFL** — Leading dimension of **CFL** exactly as specified in the dimension statement in the calling program. (Input)

**W** — **NSUB** by **NDIM** matrix containing the subject weights. (Output when **MODEL** is not zero, not referenced otherwise)

**w** is not used and may be dimensioned of length 1 if **MODEL** = 0.

**LDW** — Leading dimension of **w** exactly as specified in the dimension statement in the calling program. (Input)

**A** — Vector of length **NSUB** containing the intercepts for each subject. (Output)

**B** — Vector of length **NSUB** containing the slopes for each subject. (Output)

**WT** — Vector of length **NSUB** containing the criterion function weights for each subject. (Output)

**STRS** — Vector of length **NSUB** containing the value of the weighted optimized criterion within each subject. (Output)

**STRSS** — Value of the weighted optimized criterion function (summed over subjects). (Output)

**RESID** —  $NSUB * NC2$  vector containing the observation residuals. (Output)  
Here,  $NC2 = NSTIM * (NSTIM - 1)/2$ .

## Comments

### 1. Automatic workspace usage is

**MSIDV**  $\max(\text{NSUB}, \text{NDIM} * \text{NSTIM}, \text{ND} + 1) + \text{NDIM} * (\text{NSTIM} + 4) +$   
 $\text{NDIM} * \text{NDIM} * (2 + \max(\text{NSUB}, \text{NSTIM})) + \text{NDSS} * (\text{NDSS} + 9)$   
 $+ \text{NSUB} * (\text{NSUB} + \text{NC2} + 1) + 2 * \text{NSTIM} * \text{NSTIM} + \text{NC2} + 2 * \text{NDIM} * \text{NDIM} * \text{NSTIM} * \text{NSTIM} + 3 * \text{NPAR} + 2$

**DMSIDV**  $2 * \max(\text{NSUB}, \text{NDIM} * \text{NSTIM}, \text{ND} + 1) + 2 * \text{NDIM} * (\text{NSTIM} +$   
 $2 * \text{NDIM} * \text{NDIM} * (2 + \max(\text{NSUB}, \text{NSTIM})) + \text{NDSS} * (2 * \text{NDS} + 17) + 2 * \text{NSUB} * (\text{NSUB} + \text{NC2} + 1) + 4 * \text{NSTIM} * \text{NSTIM} + 2 * \text{NC2} + 2 * \text{NDIM} * \text{NDIM} * \text{NSTIM} * \text{NSTIM} + 4 * \text{NPAR} + 2$

Workspace may be explicitly provided, if desired, by use of M2IDV/DM2IDV. The reference is

```
CALL M2IDV (NSTIM, NSUB, X, ICNVT, MODEL, ISTRS,
            ITRANS, NDIM, IPRINT, DIST, CFL, LDCFL,
            W, LDW, A, B, WT, STRS, STRSS, RESID,
            WK1, WK2, WK3, WK4, WK5, WK6, WK7, WK8,
            WK10, WK11, WK12, WK13, ID, WKDER,
            DWKHES, DWKGRA, WKDDP, NCOM, DISP)
```

The additional arguments are as follows:

**WK1** — Work vector of length equal to  $\max(\text{NSUB}, \text{NDIM} * \text{NSTIM}, \text{ND} + 1)$

**WK2** — Work vector of length equal to  $\text{NDIM} * \text{NDIM}$

**WK3** — Work vector of length equal to  $\text{NSTIM} * \text{NSTIM}$

**WK4** — Work vector of length equal to  $\text{NSTIM} * \text{NSTIM}$

**WK5** — Work vector of length equal to  $\text{NDSS} * \text{NDSS}$

**WK6** — Work vector of length equal to  $3 * \text{NDSS}$

**WK7** — Work vector of length equal to  $5 * \text{NDSS}$

**IWK8** — Integer work vector of length equal to  $\text{NDSS}$

**WK10** — Work vector of length equal to  $\text{NDIM} * \text{NDIM}$

**WK11** — Work vector of length equal to  $\text{NSUB} * \text{NSUB}$

**WK12** — Work vector of length equal to  $\text{NDIM} * \text{NDIM} * \max(\text{NSUB}, \text{NSTIM})$

**WK13** — Work vector of length equal to  $\text{NSTIM} * \text{NDIM}$

**ID** — Integer work vector of length equal to  $4 * \text{NDIM} + 2$

**WKDER** — Work vector of length equal to  $\text{NPAR}$

**DWKHES** — Double precision work vector of length equal to  $\text{NDIM} * \text{NDIM} * \text{NSTIM} * \text{NSTIM}$

**DWKGRA** — Double precision work vector of length equal to NPAR

**WKDDP** — Work vector of length equal to NC2

**NCOM** — Work vector of length equal to NSUB

**DISP** — Work vector of length equal to NSUB \* NC2

where  $ND = NDIM * (NDIM + 1)/2$ ,  $NC2 = NSTIM * (NSTIM - 1)/2$ ,  
 $NDSS = \max(NDIM, NSTIM, NSUB)$ , and where  $NPAR = NDIM * NSTIM + 2 * NSUB$  when  $MODEL = 0$ ; otherwise  $NPAR = NDIM * NSTIM + (NDIM + 2) * NSUB$ .

## 2. Informational errors

Type	Code	
3	1	At some point during the iterations there were too many step halvings. This is usually not serious.
4	1	The program cannot continue because a Hessian matrix is ill defined. A different model may be required. This error should only occur when there is serious numerical imprecision.
4	2	A dissimilarity matrix has every element missing.

## Algorithm

Routine `MSIDV` performs multidimensional scaling analysis according to an alternating optimization algorithm. Input to `MSIDV` consists of symmetric dissimilarity matrices measuring distances between the row and column objects. Optionally, similarities can be input, and these can be converted to dissimilarities by use of the `ICNVT` option. In `MSIDV`, the row and column objects (stimuli) must be identical. Dissimilarities in multidimensional scaling are used to position the objects within a  $d = NDIM$  dimensional space, where  $d$  is specified by the user. Optionally, in the individual differences scaling model ( $MODEL \neq 0$ ), the weight assigned to each dimension for each subject may be changed.

## The Input Data

The data input in `X` must be in a special symmetric storage form. For this storage mode, the input array `X` contains only the upper triangular part of each dissimilarity matrix and does not contain the diagonal elements (which should all be zero anyway). Storage of symmetric data in `X` is as follows: `X(1)` corresponds to the (1, 2) element in the first matrix (which is a measure of the distance between objects 1 and 2), `X(2)` corresponds to the (1, 3) element, `X(3)` corresponds to the (2, 3) element, etc., until all  $t = q(q - 1)/2$  off-diagonal elements in the first matrix are stored, where  $q = NSTIM$ . The  $t + 1$  element in `X` contains the (1, 2) element in the second matrix, and so on.

Missing values are indicated in either of two ways: 1) The standard missing value indicator NaN (not a number), specified via routine `AMACH(6)` (Reference Material) may be used to indicate missing values, or 2) Negative elements of `X`

may be used to indicate missing observations. In either case, missing values are estimated as the mean dissimilarity for the subject and used as such when computing initial estimates, and they are omitted from the criterion function when optimal estimates are computed.

Routine MSIDV assumes a metric scaling model. When no transformation is specified ( $\text{ITRANS} = 1$ ), then each datum (after transforming to dissimilarities) is a measure of distance plus a constant,  $\alpha_m$ . In this case, the constant (which is always called the “intercept”) is assumed to vary with subject and must first be added to the observed dissimilarities in order to obtain a metric. When a transformation is specified ( $\text{ITRANS} \neq 1$ ), the meaning of  $\alpha_m$  changes (with respect to metrics). Thus, when  $\text{ITRANS} = 1$ , the data is assumed to be interval (see the chapter introduction) while when  $\text{ITRANS} \neq 1$  ratio data is assumed. A scaling factor, the “slope”, is also always estimated for each subject.

### The Criterion Function

When  $\text{ISTRS} = 1$  or  $2$ , the criterion function in MSIDV is given as

$$\phi = \sum_m v_m \sum_{i,j} \left( f(\delta_{ijm}^*) - \alpha_m - \beta_m f(\delta_{ijm}) \right)^2$$

where  $\delta_{ijm}$  denotes the predicted distance between objects  $i$  and  $j$  on subject  $m$ ,

$$\delta_{ijm}^*$$

denotes the corresponding dissimilarity (the observed distance),  $v_m$  is the subject weight,  $f$  is one of the transformations  $f(x) = x^2$ ,  $f(x) = x$ , or  $f(x) = \ln(x)$  specified by parameter  $\text{ITRANS}$ ,  $\alpha_m$  is the intercept added to the transformed observation within each subject, and  $\beta_m$  is the slope for the subject. For  $\text{ISTRS} = 0$ , the criterion function is given as

$$\phi = \sum_m n_m \ln \left( \sum_{i,j} \left( f(\delta_{ijm}^*) - \alpha_m - \beta_m f(\delta_{ijm}) \right)^2 \right)$$

where  $n_m$  is the number of nonmissing observations on the  $m$ -th subject.

Assuming fixed weights, the first derivatives of the criterion for  $\text{ISTRS} = 0$  are identical to the first derivatives of the criterion when  $\text{ISTRS} = 1$  or  $2$ , but with weights

$$v_m^{-1} = \sum_{i,j} \left( f(\delta_{ijm}^*) - \alpha_m - \beta_m f(\delta_{ijm}) \right)^2 / n_m$$

$\text{ISTRS}$  can, thus, be thought of as changing the weighting to be used in the criterion function.

The transformation  $f(x)$  specified by parameter  $\text{ITRANS}$  is used to obtain constant within-subject variance of the subject dissimilarities. If the variance of

the log of the observed dissimilarities (about the predicted dissimilarities) is constant within subject, then the log transformation should be used. In this case, the variance of a dissimilarity should be proportional to its magnitude. Alternatively, the within-subject variance may be constant when distances (or squared distances) are used.

### The Distance Models

The distance models for  $\delta_{ijm}$  available in MSIDV are given by:

1. The Euclidean model:

$$\delta_{ijm}^2 = \sum_{k=1}^d (\lambda_{ik} - \lambda_{jk})^2$$

2. The individual-differences model:

$$\delta_{ijm}^2 = \sum_{k=1}^d w_{mk} (\lambda_{ik} - \lambda_{jk})^2$$

where  $\Lambda$  denotes the configuration (CFL) so that  $\lambda_{ik}$  is the location of the  $i$ -th stimulus in the  $k$ -th dimension, where  $d$  is the number of dimensions, and where  $w_{mk}$  is the weight assigned by the  $m$ -th subject to the  $k$ -th dimension ( $w$ ).

### The Subject Weights

Weights that are inversely proportional to the estimated variance of the dissimilarities (about their predicted values) within each subject may be preferred because such weights lead to normal distribution theory maximum likelihood estimates (when it is assumed that the dissimilarities are independently normally distributed with constant residual variance). The estimated (conditional) variance used as the inverse of the weight  $v_m$  for the  $m$ -th subject in MSIDV (when ISTRS = 0) is computed as

$$v_m^{-1} = \sum_{i,j} \frac{(f(\delta_{ijm}^*) - \alpha_m - \beta_m f(\delta_{ijm}))^2}{n_m}$$

where the sum is over the observations for the subject, and where  $n_m$  is the number of observed nonmissing dissimilarities for the subject. These weights are used in the first derivatives of the criterion function.

When ISTRS = 1, the within-subject average sum of squared dissimilarities are used for the weights. They are computed as

$$v_m^{-1} = \frac{\sum_{i,j} f(\delta_{ijm}^*)^2}{n_m}$$

Finally, when  $\text{ISTRS} = 2$ , the within-subject variance of the dissimilarities is used for the weights. These are computed as follows

$$v_m^{-1} = \frac{\sum_{i,j} (f(\delta_{ijm}^*) - \overline{f(\delta_{ijk}^*)})^2}{n_m}$$

where

$$\overline{f(\delta_{ijk}^*)}$$

denotes the average of the transformed dissimilarities in the stratum.

### The Optimization Procedure

Initial estimates of all parameters are obtained through methods discussed in routine `MSINI` (page 1028). After obtaining initial estimates, a modified Gauss-Newton algorithm is used to obtain estimates for the parameters that optimize the criterion function. The parameters are optimized sequentially as follows:

1. Optimize the configuration estimates,  $\Lambda = \text{CFL}$ .
2. If required, estimate the optimal subject weights,  $w_{mk} = \overline{w(m, k)}$ , one subject at a time.
3. Optimize the parameters  $\alpha_m = A(m)$  and  $\beta_m = B(m)$ , one subject at a time.
4. If convergence has not been reached, continue at Step 1.

An iteration is defined to be all of the Steps 1, 2, and 3. Convergence is assumed when the maximum absolute change in any parameter during an iteration is less than  $10^{-4}$  or if there is no change in the criterion function during an iteration.

### The $L_p$ Gauss-Newton Algorithm

A modified Gauss-Newton algorithm is used in the estimation of all parameters. This algorithm, which is discussed in detail by Merle and Spath (1974), uses iteratively reweighted least squares on a Taylor series linearization of the parameters in  $\delta_{ijm}$ . During each iteration, the subject weights, which may depend upon the parameters in the model, are assumed to be fixed.

### Standardization

All models available are overparameterized so the resulting parameter estimates are not uniquely defined. For example, in the Euclidean model, the columns of  $X$  can be translated or “rotated” (multiplied by an orthonormal matrix), and the resulting stress will not be changed. To eliminate lack of uniqueness due to translation, model estimates for the configuration are centered in all models. No attempt at eliminating the rotation problem is made, but note that rotation invariance is not a problem in many of the models given. With more general



models than the Euclidean model, other kinds of overparameterization occur. Further restrictions on the parameters to eliminate this overparameterization are given below by the model transformation type specified by ITRANS. In the following,  $w_{ik} \in W$ , where  $W$  is the matrix of subject weights. The restrictions to be applied by model transformation type are

1. For all models:

$$(a) \quad \sum_{i=1}^q x_{ik} = 0$$

where  $q = \text{NSTIM}$ . i.e., center the columns of  $X$ .

(b) If  $W$  is in the model, scale the columns of  $W$  so that

$$\sum_{i=1}^q x_{ik}^2 = 1$$

2. For  $f(x) = x$  and  $f(x) = x^2$ :

(a) Set  $b_h = 1$  if the data are matrix conditional and  $W$  is in the model or if the data are unconditional. (Matrix conditional with one matrix is considered to be unconditional data.)

(b) If  $W$  is not in the model, scale all elements in  $X$  so that

$$\sum_{h=1}^{\eta} b_h^2 = \eta$$

where  $\eta = \text{NSUB}$  is the number of matrices observed.

3. For  $f(x) = \ln(x)$ , substitute  $a_h$  for  $b_h$  (but set  $a_h$  to 0 instead of 1) in all restrictions in Item 2.

### Example 1

The following example concerns some intercity distance rankings. The data are described by Young and Lewyckyj (1979, page 83). The driving mileages between various cities in the United States are ranked, yielding a symmetric ordinal dissimilarity matrix. These rankings are used as input to MSIDV. A Euclidean model is fit. The resulting two-dimensional scaling yields results closely resembling the locations of the major cities in the U.S. Note that MSIDV assumes continuous, not ranked, data.

The original rankings are given as:

$$\begin{pmatrix} - & 4 & 22 & 8 & 34 & 6 & 10 & 35 & 36 & 3 \\ & - & 13 & 15 & 31 & 21 & 9 & 32 & 30 & 5 \\ & & - & 12 & 11 & 29 & 27 & 16 & 19 & 26 \\ & & & - & 24 & 18 & 25 & 28 & 33 & 23 \\ & & & & - & 39 & 42 & 2 & 17 & 37 \\ & & & & & - & 20 & 44 & 45 & 14 \\ & & & & & & - & 43 & 40 & 1 \\ & & & & & & & - & 7 & 41 \\ & & & & & & & & - & 38 \\ & & & & & & & & & - \end{pmatrix}$$

```

INTEGER      ICNVT, IPRINT, ISTRS, ITRANS, LDCFL, LDW, LNX, MODEL,
&            NDIM, NSTIM, NSUB
PARAMETER    (ICNVT=0, IPRINT=2, ISTRS=1, ITRANS=0, LDCFL=10,
&            LDW=1, LNX=45, MODEL=0, NDIM=2, NSTIM=10, NSUB=1)
C
REAL         A(1), B(1), CFL(LDCFL,NDIM), DIST(45), RESID(LNX),
&            STRS(1), STRSS, W(LDW), WT(1), X(LNX)
EXTERNAL     MSIDV, PGOPT
C
DATA X/4, 22, 13, 8, 15, 12, 34, 31, 11, 24, 6, 21, 29, 18, 39,
&      10, 9, 27, 25, 42, 20, 35, 32, 16, 28, 2, 44, 43, 36, 30,
&      19, 33, 17, 45, 40, 7, 3, 5, 26, 23, 37, 14, 1, 41, 38/
C
C            Call PGOPT to set page length for
C            the plotting
CALL PGOPT (-2, 50)
C
CALL MSIDV (NSTIM, NSUB, X, ICNVT, MODEL, ISTRS, ITRANS, NDIM,
&          IPRINT, DIST, CFL, LDCFL, W, LDW, A, B, WT, STRS,
&          STRSS, RESID)
C
END

```

### Output

Initial parameter estimates.

```

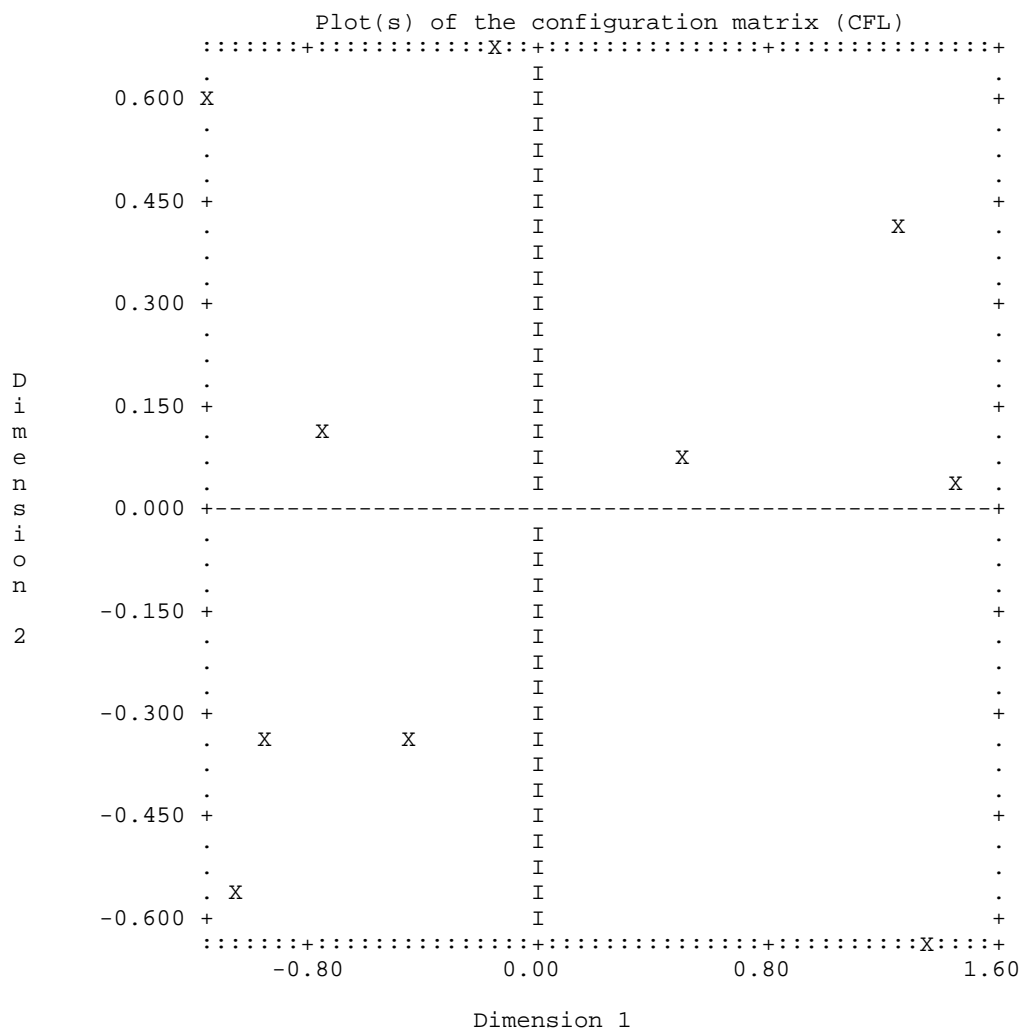
      CFL
      1      2
1  -0.762   0.124
2  -0.451  -0.349
3   0.496   0.073
4  -0.151   0.651
5   1.237   0.392
6  -1.114   0.588
7  -1.077  -0.566
8   1.461   0.034
9   1.321  -0.614

```

10 -0.961 -0.333

Iteration history

Iter	Source	Stress	Stress change	Maximum Change
1	INIT STRSS	0.3755E-02		
1	CONFIG STRSS	0.3399E-02	0.3559E-03	0.8062E-03
1	LINES STRSS	0.3142E-02	0.2564E-03	0.8062E-03
2	CONFIG STRSS	0.3068E-02	0.7382E-04	0.1022E-04
2	LINES STRSS	0.3047E-02	0.2156E-04	0.1022E-04



Final parameter estimates.

NCOM  
45

CFL

	1	2
1	-0.738	0.095

```

2  -0.447  -0.337
3   0.497   0.077
4  -0.153   0.661
5   1.237   0.399
6  -1.132   0.609
7  -1.074  -0.571
8   1.445   0.035
9   1.325  -0.624
10  -0.960  -0.343

```

```

A
-0.04255

```

```

B
0.4019

```

```

WT
0.01248

```

```

STRS
0.003047

```

```

STRESS =      3.04681E-03

```

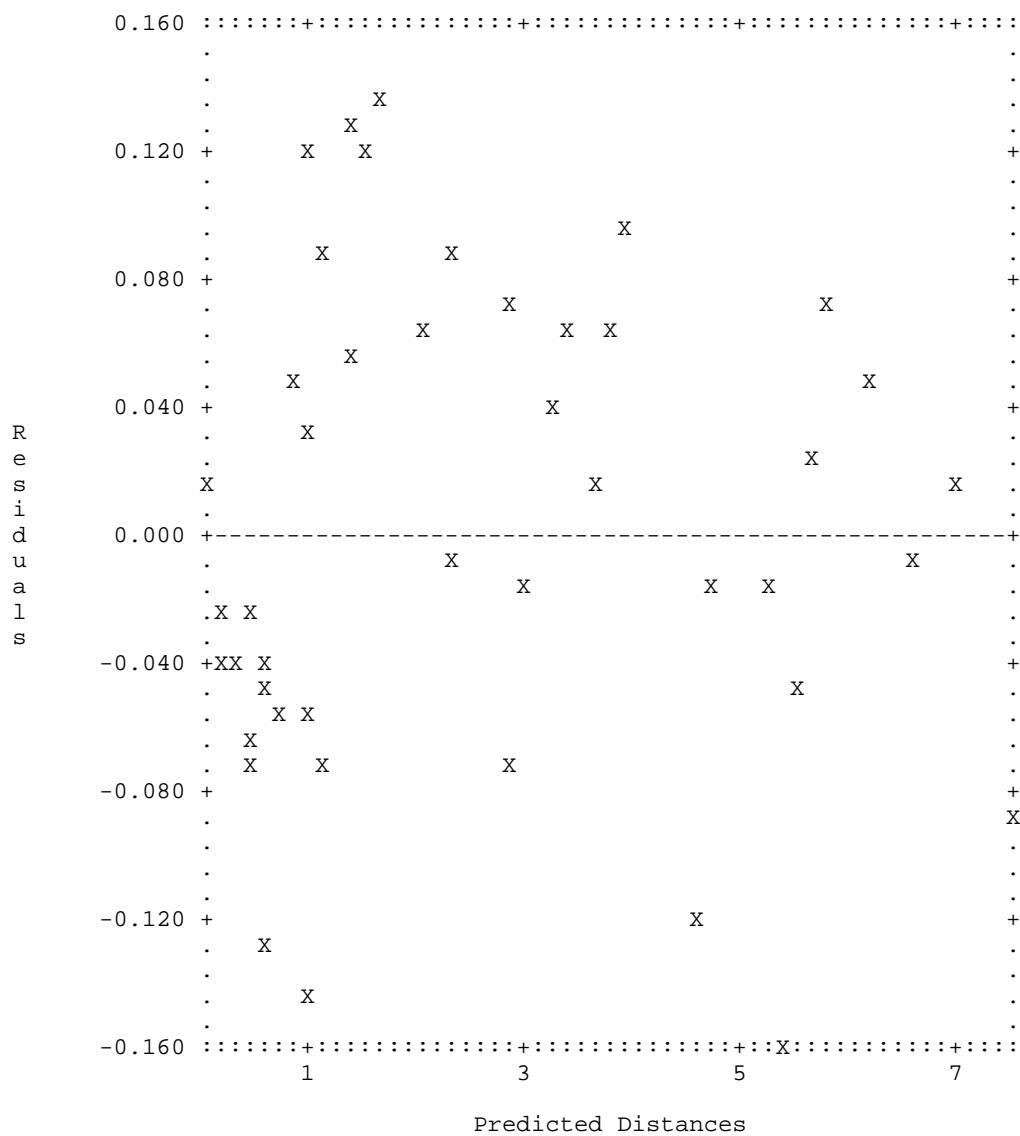
```

                Residuals
Subject  Row Stimulus  Column Stimulus Residual
  1         2           1         -0.0436
  1         3           1          0.1230
  1         3           2        -0.1422
  1         4           1        -0.1318
  1         4           2        -0.0697
  1         4           3       -0.0581
  1         5           1          0.0950
  1         5           2          0.0631
  1         5           3       -0.0456
  1         5           4          0.0639
  1         6           1       -0.0742
  1         6           2          0.1268
  1         6           3          0.0681
  1         6           4          0.1212
  1         6           5       -0.0495
  1         7           1       -0.0376
  1         7           2       -0.0216
  1         7           3       -0.0736
  1         7           4       -0.0119
  1         7           5          0.0464
  1         7           6          0.0558
  1         8           1       -0.1177
  1         8           2          0.0169
  1         8           3          0.0480
  1         8           4       -0.0173
  1         8           5       -0.0223
  1         8           6          0.0178
  1         8           7       -0.0047
  1         9           1       -0.0185
  1         9           2          0.0373
  1         9           3          0.0872
  1         9           4          0.0618
  1         9           5          0.0335

```

1	9	6	-0.0913
1	9	7	0.0202
1	9	8	-0.0671
1	10	1	-0.0415
1	10	2	-0.0276
1	10	3	0.0869
1	10	4	0.1342
1	10	5	-0.1565
1	10	6	-0.0522
1	10	7	0.0179
1	10	8	0.0701
1	10	9	-0.0191

Residual Plot



### Example 2

The second example involves three subjects' assessment of the dissimilarity between rectangles that vary in height and width. An analysis is performed in  $k = 2$  dimensions using the individual-differences scaling model. The estimated subject weights,  $w_{mk}$ , indicate how each subject weight the dimensions. The raw data are given as follows:

$$\begin{pmatrix} - & 1.00 & 1.41 & 2.24 & 2.00 & 2.24 & 1.41 & 1.00 & 1.00 \\ & - & 1.00 & 2.00 & 2.24 & 2.83 & 2.24 & 2.00 & 1.41 \\ & & - & 1.00 & 1.41 & 2.24 & 2.00 & 2.24 & 1.00 \\ & & & - & 1.00 & 2.00 & 2.24 & 2.83 & 1.41 \\ & & & & - & 1.00 & 1.41 & 2.24 & 1.00 \\ & & & & & - & 1.00 & 2.00 & 1.41 \\ & & & & & & - & 1.00 & 1.00 \\ & & & & & & & - & 1.41 \\ & & & & & & & & - \end{pmatrix}$$
$$\begin{pmatrix} - & 1.50 & 1.68 & 2.12 & 1.50 & 2.12 & 1.68 & 1.50 & 0.75 \\ & - & 0.75 & 1.50 & 2.12 & 3.35 & 3.09 & 3.00 & 1.68 \\ & & - & 0.75 & 1.68 & 3.09 & 3.00 & 3.09 & 1.50 \\ & & & - & 1.50 & 3.00 & 3.09 & 3.35 & 1.68 \\ & & & & - & 1.50 & 1.68 & 2.12 & 0.75 \\ & & & & & - & 0.75 & 1.50 & 1.68 \\ & & & & & & - & 0.75 & 1.50 \\ & & & & & & & - & 1.68 \\ & & & & & & & & - \end{pmatrix}$$

$$\begin{pmatrix} - & 0.50 & 2.06 & 4.03 & 4.00 & 4.03 & 2.06 & 0.50 & 2.00 \\ & - & 2.00 & 4.00 & 4.03 & 4.12 & 2.24 & 1.00 & 2.06 \\ & & - & 2.00 & 2.06 & 2.24 & 1.00 & 2.24 & 0.50 \\ & & & - & 0.50 & 1.00 & 2.24 & 4.12 & 2.06 \\ & & & & - & 0.50 & 2.06 & 4.03 & 2.00 \\ & & & & & - & 2.00 & 4.00 & 2.06 \\ & & & & & & - & 2.00 & 0.50 \\ & & & & & & & - & 2.06 \\ & & & & & & & & - \end{pmatrix}$$

```

INTEGER      ICNVT, IPRINT, ISTRS, ITRANS, LDCFL, LDW, LNX, MODEL,
&            NDIM, NSTIM, NSUB
PARAMETER    (ICNVT=0, IPRINT=1, ISTRS=0, ITRANS=0, LDCFL=9,
&            LDW=3, LNX=108, MODEL=1, NDIM=2, NSTIM=9, NSUB=3)
C
REAL         A(NSUB), B(NSUB), CFL(LDCFL,NDIM), DIST(LNX),
&            RESID(LNX), STRS(NSUB), STRSS, W(LDW,NDIM), WT(NSUB),
&            X(LNX)
EXTERNAL     MSIDV
C
DATA X/1.00, 1.41, 1.00, 2.24, 2.00, 1.00, 2.00, 2.24, 1.41,
&     1.00, 2.24, 2.83, 2.24, 2.00, 1.00, 1.41, 2.24, 2.00, 2.24,
&     1.41, 1.00, 1.00, 2.00, 2.24, 2.83, 2.24, 2.00, 1.00, 1.00,
&     1.41, 1.00, 1.41, 1.00, 1.41, 1.00, 1.41, 1.50, 1.68, 0.75,
&     2.12, 1.50, 0.75, 1.50, 2.12, 1.68, 1.50, 2.12, 3.35, 3.09,
&     3.00, 1.50, 1.68, 3.09, 3.00, 3.09, 1.68, 0.75, 1.50, 3.00,
&     3.09, 3.35, 2.12, 1.50, 0.75, 0.75, 1.68, 1.50, 1.68, 0.75,
&     1.68, 1.50, 1.68, 0.50, 2.06, 2.00, 4.03, 4.00, 2.00, 4.00,
&     4.03, 2.06, 0.50, 4.03, 4.12, 2.24, 1.00, 0.50, 2.06, 2.24,
&     1.00, 2.24, 2.06, 2.00, 0.50, 1.00, 2.24, 4.12, 4.03, 4.00,
&     2.00, 2.00, 2.06, 0.50, 2.06, 2.00, 2.06, 0.50, 2.06/
C
CALL MSIDV (NSTIM, NSUB, X, ICNVT, MODEL, ISTRS, ITRANS, NDIM,
&          IPRINT, DIST, CFL, LDCFL, W, LDW, A, B, WT, STRS,
&          STRSS, RESID)
C
END

```

### Output

Iteration history					
Iter	Source	Stress	Stress change	Maximum Change	
1	INIT	STRSS	-0.3590E+03		
1	CONFIG	STRSS	-0.3590E+03	0.0000E+00	0.5708E-03
1	SUB WT	STRSS	-0.3590E+03	0.0000E+00	0.1581E-02
1	LINES	STRSS	-0.3590E+03	0.0000E+00	0.2727E-02
2	CONFIG	STRSS	-0.3590E+03	0.0000E+00	0.1442E-06

```

2      SUB WT STRSS -0.3590E+03    0.0000E+00    0.7165E-04
2      LINES STRSS -0.3590E+03    0.0000E+00    0.7165E-04
Final parameter estimates.

```

```

      NCOM
1      2      3
36     36     36

```

```

      CFL
1      2
1      1.225    0.000
2      1.225   -1.225
3      0.000   -1.225
4     -1.225   -1.225
5     -1.225    0.000
6     -1.225    1.225
7      0.000    1.225
8      1.225    1.225
9      0.000    0.000

```

```

      W
1      2
1      1.000    1.000
2      0.342    1.372
3      1.411    0.089

```

```

      A
1      2      3
-0.002773    0.001941    0.000055

```

```

      B
1      2      3
0.2229    0.2587    0.2963

```

```

      WT
1      2      3
1000.0    1000.0    1000.0

```

```

      STRS
1      2      3
-119.7   -119.7   -119.7

```

```

STRESS =      -359.018

```

---

## MSDST/DMSDST (Single/Double precision)

Compute distances in a multidimensional scaling model.

### Usage

```
CALL MSDST (NSTIM, NDIM, CFL, LDCFL, NSUB, IMOD, IFORM,
            ITRANS, W, LDW, WS, LDWS, DIST)
```

### Arguments

*NSTIM* — Number of stimuli. (Input)



**NDIM** — Number of dimensions in the model. (Input)

**CFL** — NSTIM by NDIM matrix containing the stimulus configuration. (Input)

**LDCFL** — Leading dimension of CFL exactly as specified in the dimension statement in the calling program. (Input)

**NSUB** — Number of subjects. (Input)

**IMOD** — Vector of length 3 describing the weighting to be used. (Input)

<b>I</b>	<b>WEIGHT</b>
1	Not used. Reserved for other scaling subroutines.
2	Subject weights (in <i>w</i> ).
3	Stimulus weights (in <i>ws</i> ).

If **IMOD**(*i*) is zero, then the *i*-th set of weights is not used. Otherwise, the weights are used. For the Euclidean model, set **IMOD**(2) = **IMOD**(3) = 0. For the individual differences model, **IMOD**(2) should not be zero. For the stimulus weighted individual differences model, both **IMOD**(2) and **IMOD**(3) are not zero.

**IFORM** — Form option. (Input)

If **IFORM** = 0, the computed distances are stored as the upper triangle of square matrices stored columnwise without the diagonal elements. Otherwise, the distances are stored as square matrices and include the diagonal elements. See argument **DIST**.

**ITRANS** — Transformation option. (Input)

**ITRANS** determines the output returned in **DIST**.

**ITRANS Output in DIST**

0	Squared distances
1	Distances
2	Log of the distances

**W** — NSUB by NDIM matrix of individual weights. (Input)

If **IMOD**(2) is zero, then **W** is not referenced and can be an array of length 1.

**LDW** — Leading dimension of **W** exactly as specified in the dimension statement in the calling program. (Input)

**WS** — NSTIM by NDIM matrix of stimulus weights. (Input)

If **IMOD**(3) is zero, then **W** is not referenced and can be an array of length 1.

**LDWS** — Leading dimension of **WS** exactly as specified in the dimension statement in the calling program. (Input)

**DIST** — Vector of length  $n_v * NSUB$ , where  $n_v = NSTIM * (NSTIM - 1)/2$  if **IFORM** = 0, and  $n_v = NSTIM * NSTIM$  otherwise. (Output)

**DIST** may be treated as NSUB distance matrices. Storage in **DIST** is such that the elements of each column of a subject's distance matrix are adjacent. Each column in the matrix is immediately followed by the elements in the next column. If **IFORM** = 0, then only the elements in each column above the diagonal are stored. Otherwise, all elements are stored.

## Algorithm

Routine MSDST computes squared distances, distances, or log distances for various metrics in multidimensional scaling. The “distances” are computed and stored as either square matrices or as upper triangular symmetric matrices stored columnwise without the diagonal. In both cases, the distances are output in a vector of the required length. The terminology and metrics used here are the same as those used in the ALSICAL program of Takane, Young, De Leeuw (1977).

Suppose that there are  $q$  stimuli,  $M$  subjects, and  $d$  dimensions. Let  $\lambda_{ik}$  denote the location of the  $i$ -th stimulus in the  $k$ -th dimension. If  $w_{ik}$  denotes the weight of the  $i$ -th subject on the  $k$ -th dimension (matrix  $W$ ) and  $\pi_{ik}$  denotes the weight for the  $i$ -th stimulus on the  $k$ -th dimension (matrix  $\Pi$ ), then the distance models computed are the same as the distance models in MSIDV. They are given by:

Euclidean Model

$$\delta_{ijm}^2 = \sum_{k=1}^d (\lambda_{ik} - \lambda_{jk})^2$$

Individual Differences Model

$$\delta_{ijm}^2 = \sum_{k=1}^d w_{mk} (\lambda_{ik} - \lambda_{jk})^2$$

Stimulus-Weighted Model

$$\delta_{ijm}^2 = \sum_{k=1}^d \pi_{ik} (\lambda_{ik} - \lambda_{jk})^2$$

Stimulus-Weighted Individual Differences Model

$$\delta_{ijm}^2 = \sum_{k=1}^d \pi_{ik} w_{mk} (\lambda_{ik} - \lambda_{jk})^2$$

where  $\delta_{ijm}$  is the distance between the  $i$ -th and  $j$ -th stimuli on the  $m$ -th subject.

## Example

The following small example illustrates the distance computations in symmetric matrices. The data are fictional.

```
C
INTEGER      IFORM, ITRANS, LDCFL, LDW, LDWS, NDIM, NSTIM, NSUB
PARAMETER   (IFORM=0, ITRANS=0, LDCFL=4, LDW=2, LDWS=4, NDIM=2,
&           NSTIM=4, NSUB=2)

INTEGER      IMOD(3), NOUT
REAL        CFL(NSTIM,NDIM), DIST(12), W(NSUB,NDIM), WS(1)
```

```

EXTERNAL  MSDST, UMACH
C
DATA IMOD/0, 1, 0/
C
DATA CFL/1.0, -1.0, 1.0, -1.0,
&      1.0, 1.0, -1.0, -1.0/
C
DATA W/1.0, 2.0, 1.0, 2.0/
C
CALL MSDST (NSTIM, NDIM, CFL, LDCFL, NSUB, IMOD, IFORM, ITRANS,
&          W, LDW, WS, LDWS, DIST)
C
CALL UMACH (2, NOUT)
WRITE (NOUT,*) DIST
END

```

### Output

```

4.00000  4.00000  8.00000  8.00000  4.00000  4.00000  8.00000
8.00000  16.0000  16.0000  8.00000  8.00000

```

---

## MSSTN/DMSSTN (Single/Double precision)

Transform dissimilarity/similarity matrices and replace missing values by estimates to obtain standardized dissimilarity matrices.

### Usage

```
CALL MSSTN (NROW, NCOL, NSUB, IFORM, X, LDX, ICNVT, ISTRAT,
           ISCALE, NCOM, XOUT)
```

### Arguments

**NROW** — Number of row stimuli in each dissimilarity/similarity matrix. (Input)

**NCOL** — Number of column stimuli in each dissimilarity/similarity matrix. (Input)

If **IFORM** = 0 or 1, **NCOL** must equal **NROW**, and the stimuli in the rows and columns must correspond to one another.

**NSUB** — Number of dissimilarity/similarity matrices. (Input)

**IFORM** — Storage option indicating the storage mode for the input data in each column of **X**. (Input)

Array **X** contains **NSUB** columns, and each column of **X** contains a dissimilarity/similarity matrix stored as specified by option **IFORM**.

#### **IFORM** Data Storage Mode

0 Symmetric storage mode without the diagonal elements. (Upper triangular matrix stored columnwise.) In this storage mode, consecutive elements of each column of **X** contain the (1, 2), (1, 3), (2, 3), (1, 4), (2, 4), (3, 4), ..., (**NROW** - 1, **NROW**) elements of the corresponding dissimilarity/similarity matrix.

- 1 Square matrix in full storage mode. Consecutive elements of each column of  $X$  contain the  $(1, 1), (2, 1), (3, 1), \dots, (\text{NROW}, 1), (1, 2), (2, 2), \dots, (\text{NROW}, \text{NROW})$  elements of the corresponding dissimilarity/similarity matrix.
- 2 Rectangular matrix in full storage mode. In this storage mode, the row and column stimuli input in  $X$  do not correspond to each other. Let  $m = \text{NROW}$ . Consecutive elements of each column of  $X$  contain the  $(1, m + 1), (2, m + 1), \dots, (\text{NROW}, m + 1), (1, m + 2), \dots, (\text{NROW}, m + 2), \dots, (\text{NROW}, m + \text{NCOL})$  elements of the corresponding dissimilarity/similarity matrix.

$X$  —  $\text{NSUB}$  similarity or dissimilarity matrices in storage mode as determined by  $\text{IFORM}$ . (Input)

$X$  must be dimensioned as:

```
DIMENSION X(LDX, NSUB)
```

where  $\text{LDX} \geq \text{NROW} * \text{NCOL}$  in full storage mode and  $\text{LDX} \geq \text{NROW} * (\text{NROW} - 1)/2$  in symmetric storage mode. See argument  $\text{IFORM}$  for the method of storage used for each storage mode. Negative elements of  $X$ , or elements equal to NaN (“not a number”) are presumed to be missing values and will be estimated as an appropriate average in  $\text{MSSTN}$ .

$\text{LDX}$  — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

$\text{ICNVT}$  — Option for converting from similarity to dissimilarity matrices. (Input)

**ICNVT Conversion**

- 0 No conversion performed.
- 1 Subtracting each similarity from the largest similarity in the strata (see  $\text{ISTRAT}$ ).
- 2 Take the reciprocal of each similarity (elements of  $X$  equal to zero are assumed to be missing).

$\text{ISTRAT}$  — Option giving the level of stratification to be used. (Input)

If  $\text{ISTRAT} = 1$ , each dissimilarity/similarity matrix in  $X$  is considered to be in a different stratum. The data are said to be matrix conditional. If  $\text{ISTRAT} = 2$ , each column of each dissimilarity matrix is considered to be in a different stratum. (Thus, each column of array  $X$  contains  $\text{NCOL}$  strata.) For  $\text{ISTRAT}$  to be 2,  $\text{IFORM}$  must be 1 or 2. The data are said to be column conditional. If  $\text{ISTRAT} = 3$ , all of the dissimilarity/similarity matrices in  $X$  are considered to be in the same stratum. The data are said to be unconditional.

$\text{ISCALE}$  — Scaling option. (Input)

**ISCALE Scaling**

- 0 No scaling is performed.
- 1 The data in each stratum are scaled such that the sum of the squared dissimilarities equals the number of elements in the stratum.

**NCOM** — Vector containing the number of nonmissing observations in each stratum. (Output)

The diagonal elements of each dissimilarity/similarity matrix are not counted.

**ISTRAT Length of NCOM**

- 1 NSUB
- 2 NSUB \* NSTIM, where NSTIM = NROW when IFORM = 0 or 1, and NSTIM = NROW + NCOL when IFORM = 2
- 3 1

**XOUT** — Vector of length NV \* NSUB containing the standardized dissimilarity matrices where NV = NROW \* (NROW - 1)/2 if IFORM = 0 and NV = NSTIM \* NSTIM otherwise. (Output)

The value of NSTIM is as described in parameter NCOM. XOUT contains the standardized dissimilarity matrices in the same storage mode as X if IFORM = 0 or 1 and stored as square matrices when IFORM = 2. Missing values are replaced by an appropriate average dissimilarity and changed in sign. Scaling is performed as requested.

**Comments**

- 1. Automatic workspace usage is

MSSTN 2 \* NSTIM \* NSTIM units, or  
DMSSTN 4 \* NSTIM \* NSTIM units.

Workspace may be explicitly provided, if desired, by use of M2STN/DM2STN. The reference is

```
CALL M2STN (NROW, NCOL, IFORM, NSUB, X, LDX, ICNVT,  
           ISTRAT, ISCALE, NCOM, XOUT, NSTIM, XX,  
           XMIS)
```

The additional arguments are as follows:

**NSTIM** — Integer scalar. NSTIM = NROW when IFORM = 0 or 1, and NSTIM = NROW + NCOL when IFORM = 2.

**XX** — Work vector of length NSTIM \* NSTIM.

**XMIS** — Work vector of length NSTIM \* NSTIM.

- 2. Informational errors

Type	Code	
3	1	At least one column in column conditional data has all elements missing.
4	2	A dissimilarity matrix has every element missing.

**Algorithm**

Routine MSSTN standardizes dissimilarity/similarity data to be usable by other routines in the multidimensional scaling chapter. Routine MSSTN converts similarity to dissimilarity data, estimates missing values within specified strata

("conditionality groups"), scales the data, computes the number of nonmissing data elements within each stratum, and stores the data in a standard form.

The computations proceed as follows:

1. Routine MSSTN begins by expanding rectangular or symmetric storage-form data into square storage mode (the form when IFORM = 1).
2. Missing values are replaced by the average nonmissing value within the stratum, or when there is only one stratum, the average within each matrix is used. If all elements in a stratum are missing and the stratum is a column of the dissimilarity/similarity matrix, then the average of the nonmissing elements in the matrix is used as the missing value estimate. (Missing values are estimated primarily for use in routines computing estimates via "double-centering", routines MSINI, page 1028, and MSDBL, page 1024.) Missing values are denoted in the output by changing the signs of the estimated missing elements to be negative.
3. The data are converted to dissimilarity data from similarity data according to the method specified by the parameter ICNVT.
4. The data are scaled according to the method specified by the ISCALE parameter.

### Example

The following example illustrates the use of MSSTN on similarity data that are converted to dissimilarity data with the ICNVT = 1 option. Standardization within each matrix is used. The input data is such that IFORM = 0. Since ICNVT = 1 and all elements of the input data are nonnegative, no missing values are estimated. The input data is given by the following two similarity matrices:

$$\begin{pmatrix} - & 4 & 0 & 3 & 1 \\ & - & 1 & 1 & 3 \\ & & - & 0 & 2 \\ & & & - & 4 \\ & & & & - \end{pmatrix} \begin{pmatrix} - & 1 & 2 & 3 & 1 \\ & - & 1 & 2 & 0 \\ & & - & 1 & 3 \\ & & & - & 4 \\ & & & & - \end{pmatrix}$$

```
C  
INTEGER      ICNVT, IFORM, ISCALE, ISTRAT, LDX, NCOL, NROW, NSUB  
PARAMETER   (ICNVT=1, IFORM=0, ISCALE=1, ISTRAT=1, LDX=10,  
&           NCOL=5, NROW=5, NSUB=2)  
C  
INTEGER      I, J, K, N, NCOM(NSUB), NOUT  
REAL        X(LDX,NSUB), XOUT(NROW*(NROW-1))  
EXTERNAL    MSSTN, UMACH, WRIRN  
C  
DATA X/4.0, 0.0, 1.0, 3.0, 1.0, 0.0, 1.0, 3.0, 2.0, 4.0, 1.0,  
&      2.0, 1.0, 3.0, 2.0, 1.0, 1.0, 0.0, 3.0, 4.0/  
C  
CALL MSSTN (NROW, NCOL, NSUB, IFORM, X, LDX, ICNVT, ISTRAT,  
&          ISCALE, NCOM, XOUT)  
C
```

```

CALL WRIRN ('NCOM', 1, NSUB, NCOM, 1, 0)
CALL UMACH (2, NOUT)
C
N = 1
DO 20 I=1, 2
  WRITE (NOUT,99998) I
  DO 10 J=1, 4
    WRITE (NOUT,99999) (XOUT(K),K=N,N+J-1)
    N = N + J
  10 CONTINUE
  20 CONTINUE
C
99998 FORMAT (///' Output matrix (in XOUT)', I2)
99999 FORMAT (1X, 4F8.3)
C
END

```

### Output

```

NCOM
 1   2
10  10

Output matrix (in XOUT) 1
 0.000
 1.569  1.177
 0.392  1.177  1.569
 1.177  0.392  0.784  0.000

Output matrix (in XOUT) 2
 1.205
 0.803  1.205
 0.402  0.803  1.205
 1.205  1.606  0.402  0.000

```

---

## MSDBL/DMSDBL (Single/Double precision)

Obtain normalized product-moment (double centered) matrices from dissimilarity matrices.

### Usage

```

CALL MSDBL (NSTIM, NSUB, IFORM, X, LDX, ISCALE, DISP,
           LDDISP, P, LDP, DS, LDDS)

```

### Arguments

**NSTIM** — Number of stimuli in each dissimilarity matrix. (Input)

**NSUB** — Number of dissimilarity matrices. (Input)

**IFORM** — Storage option for the data in each dissimilarity matrix. (Input) Each column of X contains one of the NSUB dissimilarity matrices in the storage mode specified by IFORM.

**IFORM Data Storage Mode**

- 0 Symmetric storage mode without the diagonal elements. (Upper triangular matrix stored columnwise.) In this storage mode, consecutive elements of each column of  $X$  contain the (1, 2), (1, 3), (2, 3), (1, 4), (2, 4), (3, 4), ..., (NSTIM - 1, NSTIM) elements of the corresponding dissimilarity matrix.
- 1 Square matrix in full storage mode. Consecutive elements of each column of  $X$  contain the (1, 1), (2, 1), (3, 1), ..., (NROW, 1), (1, 2), (2, 2), ..., (NSTIM, NSTIM) elements of the corresponding dissimilarity matrix.

$X$  —  $NV$  by  $NSUB$  matrix containing the  $NSUB$  dissimilarity matrices, where  $NV = NSTIM * (NSTIM - 1)/2$  if  $IFORM = 0$ , and  $NV = NSTIM * NSTIM$  if  $IFORM = 1$ . (Input)

Missing values (NaN, “not a number”) are not allowed in  $X$ , but the position of a missing element may be indicated as a negative dissimilarity. Since  $MSDBL$  uses the absolute value of each element in  $X$  in the estimation procedure, the signs of elements in  $X$  have no effect. See Comments.

$LDX$  — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

$ISCALE$  — Scaling option. (Input)

**ISCALE Type of Scaling**

- 0 No scaling
- 1 Scaling within each matrix
- 2 Scaling over all matrices

Scaling is such that the Euclidean norm of the vector of scaled data is equal to the number of elements in vector.

$DISP$  —  $NSTIM$  by  $NSTIM$  by  $NSUB$  array containing the  $NSUB$  dissimilarity matrices in full storage mode. (Output)

In  $DISP$ , missing value estimates are positive, and all elements represent the square of distances.

$LDDISP$  — Leading and second dimension of  $DISP$  exactly as specified in the dimension statement in the calling program. (Input)

$P$  —  $NSTIM$  by  $NSTIM$  by  $NSUB$  array containing the standardized product-moment matrices in full storage mode. (Output)

$P$  contains  $NSUB$  matrices, each of size  $NSTIM$  by  $NSTIM$ . If  $DISP$  is not needed,  $DISP$  and  $P$  can occupy the same storage locations.

$LDP$  — Leading and second dimension of  $P$  exactly as specified in the dimension statement in the calling program. (Input)

$DS$  —  $NSTIM$  by  $NSTIM$  array containing the sum of the  $NSUB$  matrices in  $P$ . (Output)

$LD DS$  — Leading dimension of  $DS$  exactly as specified in the dimension statement in the calling program. (Input)



## Comments

Routine `MSSTN` (page 1020) may be used to obtain the matrix  $X$  with missing values estimated and changed in sign so that all estimates of missing values are negative. Routine `MSSTN` will also convert similarities to dissimilarities. Unless a ratio distance measure is observed, the user will usually call `MSSTN` prior to calling `MSDBL`.

## Algorithm

Routine `MSDBL` computes product-moment (double-centered) matrices from input dissimilarity matrices. The product-moment matrices output from `MSDBL` may be scaled either within each matrix, over all matrices input, or not at all.

The interest in product-moment matrices can be explained as follows: Let  $\Lambda$  denote a configuration of points in an  $d$ -dimensional Euclidean space with center at the origin. When the data is measured without error, the matrix

$P = \Lambda\Lambda^T$  can also be written as the “double-centered” matrix (defined below) obtained from the matrix of squares of distances between the rows of

$$\Lambda \left( \delta_{ij}^2 = \sum_k (\lambda_{ik} - \lambda_{jk})^2 \right)$$

These distances are input, approximately, in the dissimilarities. Thus, an estimate for  $\Lambda$  can be obtained, approximately, by computing the double-centered matrix  $P$  from the squared dissimilarities and then computing  $\Lambda$  from the scaled eigenvectors of  $P$  (such that  $P = \Lambda\Lambda^T$ ).

The computation in `MSDBL` proceeds as follows:

1. Each input dissimilarity matrix is transformed into a square symmetric matrix of distances. Asymmetric matrices are made symmetric by averaging the matrix of dissimilarities with its transpose.
2. Estimates for the square of the distances,

$$\overline{\delta^2}$$

are computed as the square of the estimated distances.

3. Let

$$\overline{\delta^2}_{m\bullet\bullet}$$

denote the average squared distance in a matrix  $m$  of squared distances, let

$$\overline{\delta^2}_{mi\bullet}$$

denote the average of the  $i$ -th row of estimated squared distances in matrix  $m$  and let

$$\overline{\delta^2}_{m\bullet j}$$

denote the average of the  $j$ -th column. The  $m$ -th product-moment matrix is computed from the  $m$ -th estimated squared distance matrix as

$$P_{mij} = -\left(\bar{\delta}_{mij}^2 - \bar{\delta}_{mi\bullet}^2 - \bar{\delta}_{m\bullet j}^2 + \bar{\delta}_{m\bullet\bullet}^2\right) / 2$$

The resulting matrix is said to be double-centered.

4. If the elements of  $P_m$  are to be scaled within matrix  $m$ , then the elements of  $P_m$  are divided by

$$\sqrt{\sum_{i,j} P_{mij}^2 / q^2}$$

where  $q = \text{NSTIM}$  so that  $q^2$  is the total number of elements in the matrix. If the elements of  $P$  are to be scaled over all matrices, then the elements of each matrix are divided by

$$\sqrt{\sum_{m,i,j} P_{mij}^2 / (sq^2)}$$

where  $s = \text{NSUB}$ .

5. The matrix DS is computed as the sum over all subjects of the product-moment matrices,  $P_m$ .

### Example

The following example illustrates the use of MSDBL in computing product-moment matrices for two input dissimilarity matrices. The input matrices are given as:

$$\begin{pmatrix} - & 4 & 1 & 3 & 1 \\ & - & 1 & 1 & 3 \\ & & - & 2 & 2 \\ & & & - & 4 \\ & & & & - \end{pmatrix} \quad \begin{pmatrix} - & 1 & 2 & 3 & 1 \\ & - & 1 & 2 & 2 \\ & & - & 1 & 3 \\ & & & - & 4 \\ & & & & - \end{pmatrix}$$

```

C      INTEGER      IFORM, ISCALE, LDDISP, LDDS, LDP, LDX, NSTIM, NSUB
PARAMETER      (IFORM=0, ISCALE=1, LDDISP=5, LDDS=5, LDP=5, LDX=10,
&              NSTIM=5, NSUB=2)
C
C      REAL          DISP(LDDISP,LDDISP,NSUB), DS(LDDS,NSTIM),
&              P(LDP,LDP,NSUB), X(LDX,NSUB)
EXTERNAL      MSDBL, WRRRN
C
C      DATA X/4.0, 1.0, 1.0, 3.0, 1.0, 2.0, 1.0, 3.0, 2.0, 4.0, 1.0,
&          2.0, 1.0, 3.0, 2.0, 1.0, 1.0, 2.0, 3.0, 4.0/
C
C      CALL MSDBL (NSTIM, NSUB, IFORM, X, LDX, ISCALE, DISP, LDDISP, P,
&              LDP, DS, LDDS)
C
C      CALL WRRRN ('The first matrix in DISP', NSTIM, NSTIM, DISP,
```

```

&          LDDISP, 0)
CALL WRRRN ('The second matrix in DISP', NSTIM, NSTIM,
&          DISP(1,1,2), LDDISP, 0)
CALL WRRRN ('The first matrix in P', NSTIM, NSTIM, P, LDP, 0)
CALL WRRRN ('The second matrix in P', NSTIM, NSTIM, P(1,1,2),
&          LDP, 0)
CALL WRRRN ('DS', NSTIM, NSTIM, DS, LDDS, 0)
C
END

```

### Output

```

The first matrix in DISP
  1      2      3      4      5
1  0.00  16.00   1.00   9.00   1.00
2  16.00   0.00   1.00   1.00   9.00
3   1.00   1.00   0.00   4.00   4.00
4   9.00   1.00   4.00   0.00  16.00
5   1.00   9.00   4.00  16.00   0.00

The second matrix in DISP
  1      2      3      4      5
1  0.00   1.00   4.00   9.00   1.00
2   1.00   0.00   1.00   4.00   4.00
3   4.00   1.00   0.00   1.00   9.00
4   9.00   4.00   1.00   0.00  16.00
5   1.00   4.00   9.00  16.00   0.00

The first matrix in P
  1      2      3      4      5
1  1.110 -1.931  0.274 -0.487  1.034
2 -1.931  1.110  0.274  1.034 -0.487
3  0.274  0.274 -0.182 -0.182 -0.182
4 -0.487  1.034 -0.182  1.338 -1.703
5  1.034 -0.487 -0.182 -1.703  1.338

The second matrix in P
  1      2      3      4      5
1  0.500  0.000 -0.500 -1.000  1.000
2  0.000  0.000  0.000  0.000  0.000
3 -0.500  0.000  0.500  1.000 -1.000
4 -1.000  0.000  1.000  2.000 -2.000
5  1.000  0.000 -1.000 -2.000  2.000

DS
  1      2      3      4      5
1  0.805 -0.966 -0.113 -0.743  1.017
2 -0.966  0.555  0.137  0.517 -0.243
3 -0.113  0.137  0.159  0.409 -0.591
4 -0.743  0.517  0.409  1.669 -1.852
5  1.017 -0.243 -0.591 -1.852  1.669

```

---

## MSINI/DMSINI (Single/Double precision)

Compute initial estimates in multidimensional scaling models.

### Usage

```

CALL MSINI (NSTIM, NSUB, IFORM, X, LDX, IMOD, NDIM, CFL,
           LDCFL, W, LDW, WS, LDWS, WMIN, WSMIN)

```

## Arguments

***NSTIM*** — Number of stimuli in each dissimilarity matrix. (Input)

***NSUB*** — Number of dissimilarity matrices to be used in the analysis. (Input)

***IFORM*** — Storage option for the data in each dissimilarity matrix. (Input)  
Each column of ***X*** contains one of the ***NSUB*** dissimilarity matrices in the storage mode specified by ***IFORM***.

### ***IFORM*** Data Storage Mode

- 0 Symmetric storage mode without the diagonal elements. (Upper triangular matrix stored columnwise.) Consecutive elements of each column of ***X*** contain the (1, 2), (1, 3), (2, 3), (1, 4), (2, 4), (3, 4), ..., (***NSTIM*** - 1, ***NSTIM***) elements of the dissimilarity matrix.
- 1 Square matrix in full storage mode. Consecutive elements of each column of ***X*** contain the (1, 1), (2, 1), (3, 1), ..., (***NSTIM***, 1), (1, 2), (2, 2), ..., (***NSTIM***, ***NSTIM***) elements of the dissimilarity matrix.

***X*** — ***NV*** by ***NSUB*** matrix containing the ***NSUB*** dissimilarity matrices, where  $NV = NSTIM * (NSTIM - 1)/2$  if ***IFORM*** = 0, and  $NV = NSTIM * NSTIM$  if ***IFORM*** = 1. (Input)

If ***IFORM*** = 0, then the input data is assumed to be symmetric, and the elements below and on the diagonal are not input. If ***IFORM*** = 1, all elements of each column of ***X*** are input, and the data for the column need not form a symmetric matrix. Missing values (NaN, “not a number”) are not allowed in ***X***, but the position of a missing element may be indicated as a negative dissimilarity. Since ***MSINI*** uses the absolute value of each element in ***X*** as the dissimilarity to be used in the estimation procedure, the sign of an element in ***X*** has no effect. See Comment 3.

***LDX*** — Leading dimension of ***X*** exactly as specified in the dimension statement in the calling program. (Input)

***IMOD*** — Vector of length 3 giving the model parameters to be estimated. (Input)

***IMOD*** also gives the method of initialization to be used for each set of parameters. Each element of ***IMOD*** corresponds to a different parameter matrix. The correspondence is given as:

### **ElementParameter Matrix**

- |   |                         |
|---|-------------------------|
| 1 | CFL—The configuration   |
| 2 | w—The subject weights   |
| 3 | WS—The stimulus weights |

The value used for each element of ***IMOD*** tells how the parameter matrix is to be initialized.

### **Value Effect on Parameter Matrix**

- |   |   |
|---|---|
| 0 | The parameter matrix is not used.   |
| 1 | The parameter matrix is input and its values are fixed. The parameter matrix may be standardized. |

- 2 Initial estimates are input, but they may be changed by *MSINI*.  
 3 *MSINI* calculates the initial estimates.

*IMOD(1)* must be nonzero. *IFORM* must not be 0 if *IMOD(3)* is not zero. If *IMOD(2)* or *IMOD(3)* is 1, *IMOD(1)* must be 1. If *IMOD(3)* is 1, *IMOD(2)* must not be 2 or 3.

***NDIM*** — Number of dimensions in the solution. (Input)

***CFL*** — *NSTIM* by *NDIM* matrix containing the estimated stimulus coordinates. (Input/Output, if *IMOD(1)* = 1 or 2; output, otherwise)

***LDCFL*** — Leading dimension of *CFL* exactly as specified in the dimension statement in the calling program. (Input)

***W*** — *NSUB* by *NDIM* matrix of subject weights. (Input/Output, if *IMOD(2)* = 1 or 2, output, if *IMOD(2)* = 3, not referenced if *IMOD(2)* = 0)

*W* is not referenced and can be dimensioned as a 1 by 1 matrix if *IMOD(2)* = 0.

***LDW*** — Leading dimension of *W* exactly as specified in the dimension statement in the calling program. (Input)

***WS*** — *NSTIM* by *NDIM* matrix of stimulus weights. (Input/Output, if *IMOD(3)* = 1 or 2; output, if *IMOD(3)* = 3, not referenced if *IMOD(3)* = 0)

*WS* is not referenced and can be dimensioned as a 1 by 1 matrix if *IMOD(3)* = 0.

***LDWS*** — Leading dimension of *WS* exactly as specified in the dimension statement in the calling program. (Input)

***WMIN*** — Minimum weight in *W* prior to adjustment. (Output, if *IMOD(2)* = 2 or 3; not referenced if *IMOD(2)* = 0 or 1)

If *WMIN* is negative, the weights in *W* are adjusted such that all weights are positive by subtracting *WMIN* from each element in *W*.

***WSMIN*** — Minimum weight in *WS* prior to adjustment. (Output, if *IMOD(3)* = 2 or 3; not referenced if *IMOD(3)* = 0 or 1)

If *WSMIN* is negative, the weights in *WS* are adjusted such that all weights are positive by subtracting *WSMIN* from each element in *WS*.

### Comments

- Automatic workspace usage is

*MSINI*  $\max(\text{NDIM} + 1, \text{NSUB}, \text{NSTIM}) + 2 * \text{NSTIM} * \text{NSTIM} + \text{NSTIM} * \text{NSTIM} * \text{NSUB} + 4 * \text{NSTIM} + \max(5 * \text{NSTIM}, 4 * \text{NSUB}) + \text{NDIM} * \text{NDIM} * \text{NSUB}$  units, or

*DMSINI*  $2 * \max(\text{NDIM} + 1, \text{NSUB}, \text{NSTIM}) + 4 * \text{NSTIM} * \text{NSTIM} + 2 * \text{NSTIM} * \text{NSTIM} * \text{NSUB} + 7 * \text{NSTIM} + 2 * \max(5 * \text{NSTIM}, 4 * \text{NSUB}) + 2 * \text{NDIM} * \text{NDIM} * \text{NSUB}$  units

Workspace may be explicitly provided, if desired, by use of *M2INI*/*DM2INI*. The reference is

```
CALL M2INI (NSTIM, NSUB, IFORM, X, LDX, IMOD, NDIM, CFL,
           LDCFL, W, LDW, WS, LDWS, WMIN, WSMIN, TR, XX,
           DISP, DS, EWK1, EWK2, IEWK, C)
```

The additional arguments are as follows:

**TR** — Real work vector of length  $\max(\text{NDIM} + 1, \text{NSUB}, \text{NSTIM})$ .

**XX** — Real work vector of length  $\text{NSTIM} * \text{NSTIM}$ .

**DISP** — Real work vector of length  $\text{NSTIM} * \text{NSTIM} * \text{NSUB}$ .

**DS** — Real work vector of length  $\text{NSTIM} * \text{NSTIM}$ .

**EWK1** — Real work vector of length  $3 * \text{NSTIM}$ .

**EWK2** — Real work vector of length  $\max(5 * \text{NSTIM}, 4 * \text{NSUB})$ .

**IEWK** — Integer work vector of length  $\text{NSTIM}$ .

**C** — Real work vector of length  $\text{NDIM} * \text{NDIM} * \text{NSUB}$ .

2. Informational error

Type	Code	
4	1	The sum of the product moment matrices for the data input in X has less than $\text{NDIM}$ positive eigenvalues. Rerun with $\text{NDIM} =$ number of positive eigenvalues or less or provide initial estimates for the configuration matrix CFL.

3. Routine MSSTN (page 1020) may be used to obtain the matrix X with missing values estimated and changed in sign so that all estimates of missing values are negative. Routine MSSTN will also convert similarities to dissimilarities. Unless a ratio distance measure is observed, the user will usually call MSSTN prior to calling MSINI.

### Algorithm

Routine MSINI computes initial estimates for the stimulus configuration ( $\Lambda = \text{CFL}$ ), subject weights ( $W = w$ ), and stimulus weights ( $\Pi = \text{WS}$ ) in multidimensional scaling models. The number of dimensions in the solution must also be input. Routine MSINI requires complete (i.e., no missing values) dissimilarity matrices as input. Consequently, missing data must be replaced by an estimate (often an average of other dissimilarities). Because the absolute values of dissimilarities are used, missing dissimilarities may be denoted by changing their sign to be negative. Estimation of missing values, and further standardization, can be performed through the use of routine MSSTN (page 1020).

In some cases, MSINI can use values input in parameter matrices CFL, W, or WS in order to compute initial estimates for other parameter matrices. For example, values input in matrix CFL may be used in the estimation of initial estimates for W or WS. Because of the method of estimation, values input for some parameter matrices will not effect the estimate computed for other matrices. In particular,

values input in  $w$  will not effect the estimation of  $CFL$ , and values input in  $ws$  will not effect the estimation of either  $CFL$  or  $w$ . Note that some combinations of input and estimated matrices are not even allowed (see the option parameter  $IMOD$ ). Also, note that when the configuration matrix  $CFL$  is input and fixed (except for standardization), computed estimates for all weights  $w$  and  $ws$  are arbitrarily taken as 1.

Let

$$\tilde{\delta}_{ijl}^2$$

denote the squared distance between stimulus  $i$  and stimulus  $j$  for matrix (subject)  $l$ , let

$$\bar{\delta}_{i\bullet l}^2$$

denote the average of the squared distances in the  $i$ -th row for the  $l$ -th subject, let

$$\bar{\delta}_{\bullet jl}^2$$

be similarly defined, and let

$$\bar{\delta}_{\bullet\bullet l}^2$$

denote the average of all squared distances for the  $l$ -th subject. If each dissimilarity input in  $x$  is measured without error, then the dissimilarities and the distances are identical. In  $MSINI$ , the errors observed in the dissimilarities,

$$\tilde{\delta}_{ijl}^2$$

are assumed to be small so that good estimates for the squared distances may be computed by squaring each dissimilarity (after first subtracting the constant obtained in Step 1 below). The computations proceed as follows:

1. The squared distance matrices are double-centered using the product moment transformation

$$p_{ijl} = -\left(\tilde{\delta}_{ijl}^2 - \bar{\delta}_{i\bullet l}^2 - \bar{\delta}_{\bullet jl}^2 + \bar{\delta}_{\bullet\bullet l}^2\right) / 2$$

The matrix formed by averaging the product moment matrices  $P_l$  (over subjects) is computed as  $\bar{P}$

2. If the configuration has been input and cannot be modified (i.e., if  $IMOD(1)$  is 1), then all weights to be estimated are taken as 1, and the computations continue in Step 8 below.
3. If the configuration matrix has not been input, then a preliminary estimate is obtained by first computing the eigenvectors ( $\Gamma$ ) corresponding to the  $d$ -largest eigenvalues of  $\bar{P}$ .

The configuration is then estimated as  $\Gamma\Delta^{1/2}$ , where  $\Delta$  is the square matrix containing the eigenvalues along the diagonal and zeros off the diagonal.

4. If the subject weights  $w$  are to be estimated, or if they can be modified (i.e., if `IMOD(2)` is 2 or greater), then a `SUMSCALE` procedure (De Leeuw and Pruzansky, 1978) is used to estimate the weights (regardless of the values input) and to “rotate” the configuration estimates. This is done as follows:

- A. The matrices

$$C_l = \Phi^{-1} \Lambda^T P_l \Lambda \Phi^{-1}$$

are computed, where  $\Phi = \Delta$  if  $\Delta$  has been computed, and where the diagonal elements of  $\Phi$  are the diagonal elements of  $\Lambda^T \Lambda$  otherwise (the off-diagonal elements of  $\Phi$  are always zero).

- B. An orthogonal matrix  $Q$  is found such that the sum of the squared off-diagonal elements of  $Q^T C_l Q$  is minimized over all matrices  $Q$ . (See IMSL routine `KPRIN`, page 797.)
  - C. A new configuration estimate is obtained by “rotating” the current estimate, i.e.,  $\Lambda_n = \Lambda$ .
  - D. The subject weights for subject  $l$  are taken as the diagonal elements of  $Q^T C_l Q$ .
5. If the subject weights have been computed and the minimum weight in  $w$  is negative, add its absolute value to all elements in  $w$  to ensure that all estimated stimulus weights are nonnegative.
  6. If the stimulus weights are to be estimated (i.e., if `(IMOD(3))` is 2 or 3), then least-squares estimates are used. The least-squares model is obtained by substituting predicted distance for actual distance in the multidimensional scaling model specified by `IMOD` (see the chapter introduction for a discussion of the models available). Least-squares fitting is then performed over the `NSUB` subjects.
  7. If the stimulus weights have been computed and the minimum weight in  $WS$  is negative, its absolute value is added to all elements in  $WS$  to ensure that all estimated stimulus weights are nonnegative.
  8. The estimates are standardized (even when `IMOD(i) = 2`) as follows:

- A. If `IMOD(2)` is not zero, then let

$$r_i = \lambda_i^T \lambda_i$$

where  $i$ -th is the  $i$ th column of the configuration matrix. Let  $w_i$  denote the  $i$ -th column of the subject weight matrix.



Standardize  $\Lambda$  such that the diagonal elements of  $\Lambda^T \Lambda$  are 1.  
Multiply  $w_i$  by  $r_i$ .

**B.** If  $\text{IMOD}(2) = 0$  but  $\text{IMOD}(3)$  is not zero, then compute  $r_i$  and standardize the configuration matrix as above. Multiply the  $i$ -th column of  $WS$  by  $r_i$ .

**C.** If both  $\text{IMOD}(2)$  and  $\text{IMOD}(3)$  are nonzero, then compute

$$s_i = \sqrt{w_i^T w_i}$$

and standardize  $w$  such that  $W^T W$  is an identity matrix. Multiply the  $i$ -th column of  $WS$  by  $c_i$ .

### Example

The following example illustrates the use of `MSINI` to obtain initial estimates for an individual differences model when symmetric dissimilarities matrices obtained from two subjects are input. The input matrices are given as:

$$\begin{pmatrix} - & 4 & 1 & 3 & 1 \\ & - & 1 & 1 & 3 \\ & & - & 2 & 2 \\ & & & - & 4 \\ & & & & - \end{pmatrix} \begin{pmatrix} - & 1 & 2 & 3 & 1 \\ & - & 1 & 2 & 2 \\ & & - & 1 & 3 \\ & & & - & 4 \\ & & & & - \end{pmatrix}$$

Estimates obtained from `MSINI` are not optimal. Usually an optimizing multidimensional scaling routine will be called with the initial estimates computed in `MSINI`.

```

C      INTEGER      IFORM, LDCFL, LDW, LDWS, LDX, NDIM, NSTIM, NSUB
PARAMETER      (IFORM=0, LDCFL=5, LDW=2, LDWS=5, LDX=10, NDIM=2,
&              NSTIM=5, NSUB=2)
C
C      INTEGER      IMOD(3), NOUT
REAL           CFL(LDCFL,NDIM), W(LDW,NDIM), WMIN, WS(LDWS,NDIM),
&              WSMIN, X(LDX,2)
EXTERNAL      MSINI, UMACH, WRRRN
C
C      DATA X/4.0, 1.0, 1.0, 3.0, 1.0, 2.0, 1.0, 3.0, 2.0, 4.0, 1.0,
&          2.0, 1.0, 3.0, 2.0, 1.0, 1.0, 2.0, 3.0, 4.0/
C      DATA IMOD/3, 3, 0/
C
C      CALL UMACH (2, NOUT)
C
C      CALL MSINI (NSTIM, NSUB, IFORM, X, LDX, IMOD, NDIM, CFL, LDCFL,
&              W, LDW, WS, LDWS, WMIN, WSMIN)
C
C      CALL WRRRN ('The Configuration', NSTIM, NDIM, CFL, LDCFL, 0)
CALL WRRRN ('Subject weights', NSUB, NDIM, W, LDW, 0)
WRITE (NOUT,99999) WMIN

```

```

C
99999 FORMAT (/, ' WMIN = ', F12.4)
C
      END

```

### Output

```

The Configuration
      1      2
1  0.2279  0.6854
2 -0.0808 -0.6584
3 -0.1728 -0.0090
4 -0.6621 -0.2287
5  0.6879  0.2107
Subject weights
      1      2
1  7.078  8.533
2  9.615  0.000
WMIN =      0.0000

```

---

## MSTRS/DMSTRS (Single/Double precision)

Compute various stress criteria in multidimensional scaling.

### Usage

```

CALL MSTRS (N, DIST, DISP, INTCEP, A, ISLOPE, B, POWER,
           ISTRS, STRSS, WT)

```

### Arguments

**N** — Number of distances and disparities. (Input)

**DIST** — Vector of length **N** containing the distances. (Input)  
Missing values are not allowed in **DIST**.

**DISP** — Vector of length **N** containing the disparities. (Input)

**INTCEP** — Intercept option parameter. (Input)

If **INTCEP** = 0, the intercept is not used in the model. If **INTCEP** = 1, the intercept is used in the model.

**A** — The intercept. (Input)

If **INTCEP** = 0, **A** is not used.

**ISLOPE** — Slope option parameter. (Input)

If **ISLOPE** = 0, the slope **B** is not used. If **ISLOPE** = 1, the slope is used.

**B** — The slope. (Input)

If **ISLOPE** = 0, **B** is not used.

**POWER** — Power to use in the stress function. (Input)

**POWER** must be greater than or equal to 1.

**ISTRS** — Stress option parameter. (Input)

**ISTRSS Stress Criterion Used**

- 0 Log stress
- 1 Stress weighted by the inverse of the sum of the squared disparities
- 2 Stress weighted by the inverse of the sum of the centered squared disparities

**STRSS** — The computed stress criterion. (Output)

**WT** — The weight used in computing the stress. (Output)

If the weight is too large, a maximum weight is used. See the algorithm section of the manual document.

**Algorithm**

Routine **MSTRS** computes the value of stress criteria commonly used in multidimensional scaling. Routine **MSTRS** allows transformed values of the disparities and distances to be input and will compute the stress on the transformed values. Additionally, the user can input a slope and/or an intercept to be used in the stress computations, and the stress can be computed using an arbitrary  $L_p$  norm as well as the squared error norm in which  $p = 2$ .

Let

$$\delta_i^*$$

denote a disparity,  $\delta_i$  denote the corresponding distance,  $\alpha$  denote the intercept, and let  $\beta$  denote the slope. If **INTCEP** = 0, then set  $\alpha = 0$ . If **ISLOPE** = 0, then set  $\beta = 1$ .

Set  $\epsilon = 0.001$ , and let

$$\tau = \sum_{i=1}^n |\delta_i^* - \alpha - \beta\delta_i|^p$$

When **ISTRSS** = 0, the stress is computed as

$$\phi_0 = n \ln[\max(n\epsilon, \tau)]$$

where  $n$  is the number of nonmissing disparities, and  $p = \text{POWER}$  is the power to be used. This stress formula, when optimized, can lead to to normal distribution theory maximum likelihood estimation. It can not be used in nonmetric scaling. The weight is computed as  $n/\max(n\epsilon, \tau)$ .

When **ISTRSS** is 1, the stress is computed as

$$\phi_1 = \frac{\tau}{\max(\epsilon\tau, \sum_{i=1}^n |\delta_i^*|^p)}$$

and the weight returned is given as

$$1 / \max\left(\epsilon\tau, \sum_{i=1}^n |\delta_i^*|^p\right)$$

Takane, Young, and de Leeuw (1977) recommend using this formula when the data is not column conditional (i.e., whenever the stress is computed over one or more dissimilarity matrices rather than over one column in a single matrix). When ISTRS = 2, the stress is given by

$$\phi_2 = \frac{\tau}{\max\left(\epsilon\tau, \sum_{i=1}^n |\delta_i^* - \bar{\delta}_\bullet^*|^p\right)}$$

where

$$\bar{\delta}_\bullet^* = \sum_{i=1}^n \delta_i^*$$

is the average of the nonmissing disparities. The weight is computed as

$$1 / \max\left(\epsilon\tau, \sum_{i=1}^n |\delta_i^* - \bar{\delta}_\bullet^*|^p\right)$$

Takane, Young, and de Leeuw (1977) recommend this stress for column conditional data.

Missing values (NaN) are not allowed in DIST while missing disparities in DISP are not used in the computations. If all disparities are missing, the stress criteria is set to 0, and the weight (WT) is set to missing (NaN).

In general, a single call to MSTRS would be made for each strata (“conditionality group”) in the data.

### Example

The following example illustrates the computation of stress when the log of the distances and disparities are input. For this example, ISTRS is 1 and POWER is 2.

```

C      INTEGER      INTCEP, ISLOPE, ISTRS, N
      REAL          A, POWER
      PARAMETER     (A=0.0, INTCEP=0, ISLOPE=1, ISTRS=1, N=10, POWER=2.0)
C
      INTEGER      I, NOUT
      REAL          ALOG, B, DISP(N), DIST(N), SDOT, STRSS, WT
      INTRINSIC     ALOG
      EXTERNAL      MSTRS, SDOT, UMACH
C
      DATA DIST/4.0, 1.5, 1.25, 3.0, 1.75, 2.0, 1.0, 3.5, 2.5, 3.75/
      DATA DISP/4.0, 1.0, 1.0, 3.0, 1.0, 2.0, 1.0, 3.0, 2.0, 4.0/
C
C          Transform the data
      DO 10 I=1, N
          DIST(I) = ALOG(DIST(I))
          DISP(I) = ALOG(DISP(I))
10 CONTINUE
C
C          Compute a slope
      B = SDOT(N,DISP,1,DIST,1)/SDOT(N,DIST,1,DIST,1)

```

```
C          Compute the stress
  CALL MSTRS (N, DIST, DISP, INTCEP, A, ISLOPE, B, POWER, ISTRS,
&          STRSS, WT)
C          Print results
  CALL UMACH (2, NOUT)
  WRITE (NOUT,99999) STRSS, WT
C
99999 FORMAT (' STRSS = ', F12.4, '      WT = ', F12.4)
      END
```

**Output**

```
STRSS =      0.0720      WT =      0.1385
```

# Chapter 15: Density and Hazard Estimation

---

## Routines

<b>15.1. Estimates for a Density</b>		
Penalized likelihood estimates .....	DESPL	1040
Kernel estimates .....	DESKN	1044
Gaussian kernel estimates via fast Fourier transform .....	DNFFT	1047
Point estimates .....	DESPT	1052
<b>15.2. Modified Likelihood Estimates for Hazards</b>		
Estimates of the smoothing parameters, general case .....	HAZRD	1054
Estimates of the smoothing parameters, easy-to-use version .....	HAZEZ	1061
Estimation of the hazard function .....	HAZST	1069

---

## Usage Notes

The routines described in this chapter compute estimates for smoothing parameters and estimates in models for estimating density and hazard functions. For density estimation, the penalized likelihood method of Scott (1976) may be used to obtain smooth estimates for arbitrary (smooth) densities. Alternatively, the routines `DESKN` (page 1044) and `DNFFT` (page 1047) obtain density estimates by the kernel method for a given window width and kernel function. Routine `DNFFT` uses a Gaussian kernel, while for routine `DESKN`, the kernel is provided by the user. Finally, routine `DESPT` (page 1052) finds linear or quasi-cubic interpolated estimates of a density. Tapia and Thompson (1978) discuss all of these methods.

For hazard estimation, the methods of Tanner and Wong (1984) are used to obtain estimates of the smoothing parameters in a modified likelihood. These methods are implemented in routines `HAZRD` (page 1054) and `HAZEZ` (page 1061), the difference between the routines is in the ease of use and the options offered. For given smoothing parameters, the routine `HAZST` (page 1069) may be used to obtain estimates for the hazard function.

---

## DESPL/DDESPL (Single/Double precision)

Perform nonparametric probability density function estimation by the penalized likelihood method.

### Usage

```
CALL DESPL (NOBS, X, NODE, BNDS, INIT, ALPHA, MAXIT, EPS,  
           DENS, STAT, NMISS)
```

### Arguments

**NOBS** — Number of observations. (Input)

**X** — Vector of length NOBS containing the random sample of responses. (Input)

**NODE** — Number of mesh nodes for the discrete probability density estimate. (Input)

NODE must be an odd integer greater than 4.

**BNDS** — Vector of length 2 containing the upper and lower endpoints for the interval of support of the density. (Input)

The node values are taken as  $BNDS(1)$ ,  $BNDS(1) + h$ , ...,  $BNDS(2)$ , where  $h = (BNDS(2) - BNDS(1))/(NODE - 1)$ . All observations in vector **x** should be in the support interval.

**INIT** — Initialization option. (Input)

INIT = 0 means that a bootstrap procedure is used to obtain initial estimates for the density. Otherwise, user-supplied initial estimates are contained in **DENS** on entry into DESPL.

**ALPHA** — Penalty-weighting factor that controls the smoothness of the estimate. (Input)

For standard normal data, ALPHA = 10.0 works well. Other values that might be tried are 1.0 and 100.0. ALPHA must be greater than 0.0.

**MAXIT** — Maximum number of iterations allowed in the iterative procedure. (Input)

MAXIT = 30 is typical.

**EPS** — Convergence criterion. (Input)

When the Euclidean norm of the changes to **DENS** is less than EPS, convergence is assumed. EPS = 0.0001 is typical.

**DENS** — Vector of length NODE containing the estimated values of the discrete pdf at the NODE equally spaced mesh nodes. (Input/Output, if INIT ≠ 0; output, otherwise)

If INIT is not zero, then DENS(1) through DENS(NODE) contain the (positive) initial estimates on input. The sum of these estimates times the window width  $h$  (see BNDS) must equal 1.0, i.e., the integral of the density must be 1.

**STAT** — Vector of length 4 containing output statistics. (Output)  
 STAT(1) and STAT(2) contain the log-likelihood and the log-penalty terms, respectively. STAT(3) and STAT(4) contain the estimated mean and variance for the estimated density.

**NMISS** — Number of missing values in  $x$ . (Output)

### Comments

1. Automatic workspace usage is

DESPL NOBS + 6 \* NODE + 9 \* (NODE - 2) units, or  
 DDESPL 2 \* NOBS + 12 \* NODE + 17 \* (NODE - 2) units.

Workspace may be explicitly provided, if desired, by use of  
 D2SPL/DD2SPL. The reference is

```
CALL D2SPL (NOBS, X, NODE, BNDS, INIT, ALPHA, MAXIT,
           EPS, DENS, STAT, NMISS, HESS, LDHESS,
           ILOHI, DENEST, B, IPVT, WK2, XWK)
```

The additional arguments are as follows.

**HESS** — Work vector of length  $7 * (NODE - 2)$ .

**LDHESS** — Leading dimension of HESS exactly as specified in the  
 dimension statement in the calling program. (Input)  
 The leading dimension must be set to 7.

**ILOHI** — Integer work vector of length  $2 * NODE$ .

**DENEST** — Work vector of length  $3 * NODE$ .

**B** — Work vector of length NODE.

**IPVT** — Integer work vector of length  $NODE - 2$ .

**WK2** — Work vector of length  $NODE - 2$ .

**XWK** — Work vector of length NOBS. If  $x$  is sorted with all missing  
 (NaN, not a number) values at the end, then XWK is not needed. If  $x$  is  
 not needed,  $x$  and XWK can share the same storage location.

2. Informational error

Type	Code	
3	1	The maximum number of iterations is exceeded.

- 3 Routine DESPT (page 1052) may be used after the estimates DENS have  
 been obtained in order to obtain an interpolated estimate of the density at  
 new points. Use AMESH = BNDS in calling DESPT.

### Algorithm

Routine DESPL computes piecewise linear estimates of a one-dimensional density  
 function for a given random sample of observations. These estimates are  
 discussed in detail in Scott et al. (1980), and in Tapia and Thompson (1978,



Chapter 5). The estimator of the density function is piecewise linear over the finite interval (BNDS(1) to BNDS(2)), is nonnegative, and integrates to one. A penalty method is used to ensure “smooth” behavior of the estimator. The criterion function to be maximized is a discrete approximation to

$$\Phi = \prod_{i=1}^n f(x_i) \exp \left( -\alpha \int \left| \frac{d^2 f(t)}{dt^2} \right|^2 dt \right)$$

where  $n = \text{NOBS}$  and  $f(t)$  is a density function. Let  $m = \text{NODE}$ . The discrete approximation is as follows: The density  $f$  is estimated at each of the equally spaced grid points  $t_j$ , for  $j = 1, \dots, m$ , with restriction  $f(t_1) = f(t_m) = 0.0$ ; the density at each data point  $x_i$  is then estimated using linear interpolation. The integral of the second derivative of the square of  $f$  is approximated using the piecewise linear function defined by the estimates of  $f$  at the grid points  $t_j$ .

Because  $\ln \Phi$  is actually maximized, the criterion can be separated into a likelihood term (returned in STAT(1)) and a penalty term (returned in STAT(2)).

The parameter  $\alpha$  (= ALPHA) determines the amount of “smoothness” in the estimate. The larger the value of  $\alpha$ , the smoother the resulting estimator for  $f$ . In practice, the user should pick  $\alpha$  as small as possible such that there is not excessive bumpiness in the estimator. One way of doing this is to try several values of  $\alpha$  that differ by factors of 10. The resulting estimators can then be graphically displayed and examined for bumpiness.  $\alpha$  could then be chosen from the displayed density estimates. IMSL routines can be used to produce line printer plots (PLOT, page 1096) of the estimated density. For a random sample from the standard normal distribution,  $\alpha = 10.0$  works well. Note that  $\alpha$  changes with scale. If  $x$  is multiplied by a factor  $\beta$ ,  $\alpha$  should be multiplied by a factor  $\beta^5$ .

The second choice to be made in using DESPL is the mesh for the estimator. The mesh interval (BNDS(1), BNDS(2)) should be picked as narrow as possible since a narrow mesh will speed algorithm convergence. Note, however, that points outside the interval (BNDS(1), BNDS(2)) are not included in the likelihood. Because of this fact, DESPL actually estimates a density that is conditional on the mesh interval (BNDS(1), BNDS(2)). The number of mesh nodes, NODE, should be as small as possible, but large enough to exhibit the “fine” structure of the density. One possible method for determining NODE is to use  $\text{NODE} = 21$  initially. With  $\text{NODE} = 21$ , find an acceptable value for  $\alpha$ . When an acceptable value for  $\alpha$  has been found, increase or decrease NODE as required.

STAT(3) and STAT(4) contain “exact” estimates of the mean and variance when the estimated piecewise linear density is used in the required integrals. Routine DESPT (page 1052) may be used to find interpolated estimates for the density at any point  $x$  given the NODE estimates of the density returned in DENS.

### Example

An estimate for a density function of unknown form using a random sample of size 10 and 13 mesh points with  $\alpha = 10$  is estimated as follows:

```
INTEGER      INIT, MAXIT, NOBS, NODE
REAL         ALPHA, EPS
PARAMETER    (ALPHA=10.0, EPS=0.0001, INIT=0, MAXIT=25, NOBS=10,
&           NODE=13)
C
INTEGER      NMISS, NOUT
REAL         BNDS(2), DENS(NODE), STAT(4), X(NOBS)
EXTERNAL     DESPL, UMACH
C
DATA BNDS/-3., 3./
DATA X/-.9471, -.7065, -.2933, -.1169, .2217, .4425, .4919,
&       .5752, 1.1439, 1.3589/
C
CALL DESPL (NOBS, X, NODE, BNDS, INIT, ALPHA, MAXIT, EPS, DENS,
&          STAT, NMISS)
C
CALL UMACH (2, NOUT)
WRITE (NOUT, '( ' DENS = ' ', 9F7.4, /, 9X, 4F7.4)') DENS
WRITE (NOUT, '( ' Log-likelihood term = ' ', F7.3, /,
&         ' ' Log-penalty term      = ' ', F7.3, /,
&         ' ' Mean                  = ' ', F7.3, /,
&         ' ' Variance               = ' ', F7.3)') STAT
END
```

### Output

```
DENS =  0.0000 0.0014 0.0356 0.1111 0.2132 0.3040 0.3575 0.3565 0.2947
        0.1986 0.0986 0.0288 0.0000
Log-likelihood term = -11.968
Log-penalty term   = -1.303
Mean               =  0.217
Variance           =  1.042
```

The following figure shows the affect of various choices of  $\alpha$ . For larger  $\alpha$ , the density estimate is smoother.

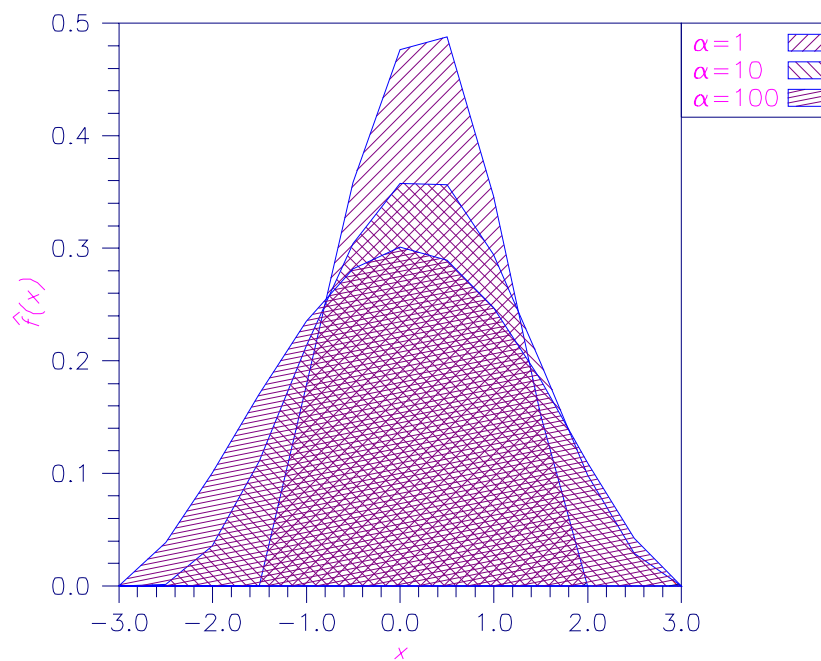


Figure 15-1 Density Estimates Using  $\alpha = 1, 10, 100$

---

## DESKN/DDESKN (Single/Double precision)

Perform nonparametric probability density function estimation by the kernel method.

### Usage

```
CALL DESKN (XKER, NOBS, X, WINDOW, XMAX, NXPT, XPT, DENS,
           NMISS)
```

### Arguments

**XKER** — User-supplied FUNCTION to compute the kernel at any point on the real line. The form is XKER(Y), where

**Y** — Point at which the kernel is to be evaluated.

**XKER** — Value of the kernel at point Y.

**XKER** — must be declared EXTERNAL in the calling program.

**NOBS** — Number of observations. (Input)

**X** — Vector of length NOBS containing the random sample of observations. (Input)

**WINDOW** — Window width for the kernel function. (Input)  
Generally, several different values of **WINDOW** should be tried.

**XMAX** — Cutoff value such that  $x_{KER}(y) = 0.0$  for all  $|y|$  greater than **XMAX**. (Input)

If **XMAX** exists, then the kernel function is 0.0 for all  $y$  greater in absolute value than **XMAX**, and the efficiency of the computations is enhanced. If no such **XMAX** exists or the user does not wish to make use of **XMAX**, then **XMAX** should be assigned any nonpositive value.

**NXPT** — Number of points at which a density estimate is desired. (Input)

**XPT** — Vector of length **NXPT** containing the values at which a density estimate is desired. (Input)

If **XMAX** is greater than zero, then **XPT** must be sorted from smallest to largest.

**DENS** — Vector of length **NXPT** containing the density estimates at the points specified in **XPT**. (Output)

**NMISS** — Number of missing (NaN, not a number) values in **x**. (Output)

### Comments

1. Informational error  
Type Code  
4 7 Negative kernel functions are not allowed.
2. Routine **DESPT** (page 1052) may be used to obtain interpolated density estimates from the **NXPT** density estimates returned in **DENS**. Array **AMESH** in **DESPT** corresponds to array **XPT** in **DESKN**.

### Algorithm

Routine **DESKN** computes kernel estimates of the density function for a random sample of (scalar-valued) observations. The kernel estimate of the density at the point  $y$  is given by.

$$\hat{f}(y) = \frac{1}{nh} \sum_{i=1}^n K[(y - x_i) / h]$$

where

$$\hat{f}(y)$$

is the estimated density at  $y$ ,  $K$  is the kernel function,  $x_i$  denotes the  $i$ -th observation,  $n$  is the number of observations, and  $h$  is a fixed constant (called the “window width”) supplied by the user.

One is usually interested in computing the density estimates using several values of the window width  $h$ . Tapia and Thompson (1978), Chapter 2, give some considerations relevant to the choice of  $h$ . Some common kernel functions (see Tapia and Thompson 1978, page 60) are given as follows.

Name	Function
Uniform	$K(y) = \begin{cases} 0.5 & \text{for }  y  < 1 \\ 0 & \text{elsewhere} \end{cases}$
Triangular	$K(y) = \begin{cases} 1 -  y  & \text{for }  y  < 1 \\ 0 & \text{elsewhere} \end{cases}$
Biweight	$K(y) = \begin{cases} 15(1 - y^2)^2 / 16 & \text{for }  y  < 1 \\ 0 & \text{elsewhere} \end{cases}$
Normal	$K(y) = \frac{1}{\sqrt{2\pi}} e^{-y^2/2} \quad -\infty < y < \infty$

The computation can be made much more efficient when the kernel is nonzero over a finite range since observations outside this range can be ignored in the computation of the density. In this case, the array XPT is assumed to be sorted.

### Example

In this example, the standard normal density function is estimated at 13 points using a random sample of 10 points from a standard normal distribution. The biweight kernel function is used. The actual density for the standard normal density is also reported in the output for comparison. The random sample is generated using routines RNSET (page 1166) and RNNOR (page 1208).

```

INTEGER      NOBS, NXPT
REAL         C1, WINDOW, XMAX
PARAMETER   (C1=0.3989423, NOBS=10, NXPT=13, WINDOW=2.0, XMAX=1.0)
C
INTEGER      I, NMISS, NOUT
REAL         DENS(NXPT), EXP, X(NOBS), XKER, XPT(NXPT)
INTRINSIC    EXP
EXTERNAL     DESKN, RNNOR, RNSET, UMACH, XKER
C
DATA XPT/-3.0, -2.5, -2.0, -1.5, -1.0, -0.5, 0.0, 0.5, 1.0, 1.5,
&        2.0, 2.5, 3.0/
C
CALL RNSET (1234457)
CALL RNNOR (NOBS, X)
C
CALL DESKN (XKER, NOBS, X, WINDOW, XMAX, NXPT, XPT, DENS, NMISS)
C
CALL UMACH (2, NOUT)
WRITE (NOUT, '( " NMISS = ', I1)') NMISS
WRITE (NOUT, '( " DENS Estimate = ', 10F6.4, ', , 8X, 3F6.4)') DENS
WRITE (NOUT, '( " DENS Exact   = ', 10F6.4, ', , 8X, 3F6.4)')
&                                     (C1*EXP(-XPT(I)*XPT(I)/2.0), I=1, NXPT)
END
REAL FUNCTION XKER (Y)
REAL      Y
C
REAL      ABS
```

```

C      INTRINSIC  ABS
      IF (ABS(Y) .LT. 1.0) THEN
          XKER = 15.0*(1.0-Y*Y)*(1.0-Y*Y)/16.0
      ELSE
          XKER = 0.0
      END IF
      RETURN
      END
  
```

### Output

```

NMISS = 0
DENS Estimate = 0.00000.01180.07900.16980.26780.34670.36870.31840.22340.1391
                0.06120.01350.0005
DENS Exact    = 0.00440.01750.05400.12950.24200.35210.39890.35210.24200.1295
                0.05400.01750.0044
  
```

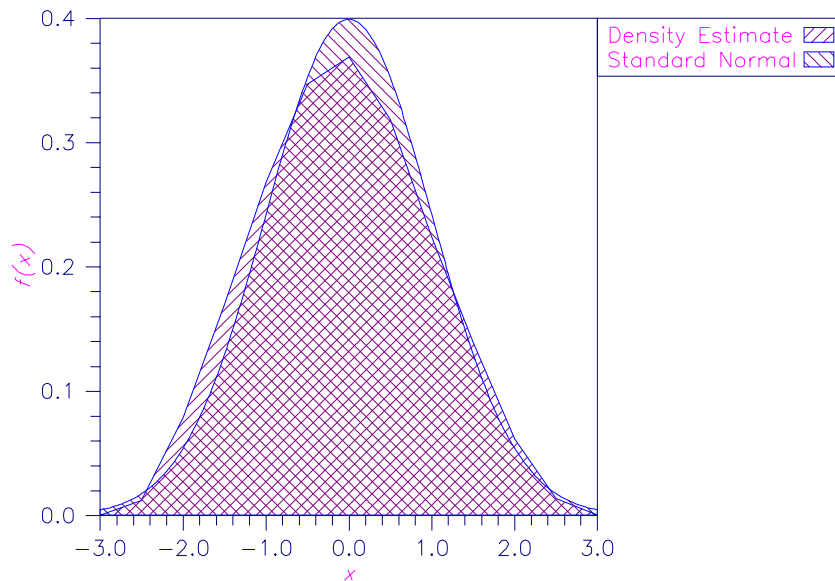


Figure 15-2 Density Estimate and Standard Normal Density

---

## DNFFT/DDNFFT (Single/Double precision)

Compute Gaussian kernel estimates of a univariate density via the fast Fourier transform over a fixed interval.

### Usage

```

CALL DNFFT (NOBS, X, FREQ, BNDS, WINDOW, IFFT, NXPT, COEF,
            DENS, NRMIS)
  
```

## Arguments

**NOBS** — Number of observations. (Input)

**X** — Vector of length NOBS containing the data for which a univariate density estimate is desired. (Input)

X is not referenced and may be dimensioned of length 1 in the calling program if IFFT = 1.

**FREQ** — Vector of length NOBS containing the frequency of the corresponding element of X. (Input)

If FREQ(1) is -1.0, then the vector FREQ is not used and all frequencies are taken to be one. FREQ is also not used if IFFT = 1. In either case, FREQ may be dimensioned of length 1 in the calling program.

**BNDS** — Vector of length 2 containing the minimum and maximum values of X at which the density is to be estimated. (Input)

Observations less than BNDS(1) or greater than BNDS(2) are ignored. If either range of the hypothesized density is infinite, a value equal to the smallest observation minus 3 \* WINDOW is a good choice for BNDS(1), and a value equal to the largest observation plus 3 \* WINDOW is a good choice for BNDS(2). Let STEP = (BNDS(2) - BNDS(1))/(NXPT - 1), and note that the density is estimated at the points BNDS(1) + i \* STEP where  $i = 0, 1, \dots, NXPT - 1$ . The density is assumed constant over the interval from BNDS(1) + i \* STEP to BNDS(1) + (i + 1) \* STEP.

**WINDOW** — Window width for the kernel function. (Input)

Generally, several different values for WINDOW should be tried. When several different values are tried, use the IFFT option.

**IFFT** — Fourier transform option parameter. (Input)

If IFFT = 1, then COEF contains the Fourier coefficients on input, and the coefficients are not computed. Otherwise, the coefficients are computed. This option is used when several different values for WINDOW are to be tried. On the first call to DNFFT, IFFT = 0 and the coefficients COEF are computed. On subsequent calls, these coefficients do not need to be recomputed (but only if NXPT also remains fixed).

**NXPT** — Number of equally-spaced points at which the density is to be estimated. (Input)

Routine DNFFT is most efficient when NXPT is a power of 2. Little efficiency is lost if NXPT is a product of small primes. Because of the method of estimation, NXPT should be large, say greater than 64.

**COEF** — Vector of length NXPT containing the Fourier coefficients. (Input, if IFFT= 1; output, otherwise)

**DENS** — Vector of length NXPT containing the density estimates. (Output)

The density is estimated at the points BNDS(1) + i \* STEP,  $i = 0, 1, \dots, NXPT - 1$ , where STEP = (BNDS (2) - BNDS(1))/(NXPT - 1).

***NRMISS*** — Number of rows of data that contain missing values in *X* or *FREQ*.  
 (Output) *NRMISS* is not referenced if *IFFT* = 1.

### Comments

- Automatic workspace usage is

*DNFFT* 2 \* *NXPT* + 15 units, or  
*DDNFFT* 4 \* *NXPT* + 30 units.

Workspace may be explicitly provided, if desired, by use of  
*D2FFT/DD2FFT*. The reference is

CALL *D2FFT* (*NOBS*, *X*, *BNDS*, *WINDOW*, *IFFT*, *NXPT*,  
*COEF*, *DENS*, *NRMISS*, *WFFTR*)

The additional argument is

***WFFTR*** – Work vector of length 2 \* *NXPT* + 15. See Comment 3.  
 (Input)

- Informational errors

Type	Code	
4	1	The sum of the frequencies must be positive.
4	2	Each frequency must be nonnegative.
4	3	There are no valid observations remaining after all missing values are eliminated.

- WFFTR* is computed in *DNFFT*. If *D2FFT* is to be called, *WFFTR* must first be computed via the following FORTRAN statement:

CALL *FFTRI* (*NXPT*, *WFFTR*)

If *DD2FFT* is used, call *DDFFTRI* instead of *FFTRI*. *WFFTR* need not be recomputed between successive calls to *D2FFT* if *NXPT* does not change.

### Algorithm

Routine *DNFFT* computes Gaussian kernel estimates of the density function for a random sample of (scalar-valued) observations using a Gaussian kernel (normal density). The computations are comparatively fast because they are performed through the use of the fast Fourier transform. Routine *DESKN* (page 1044) should be used in place of *DNFFT* if a kernel other than the Gaussian kernel is to be used, if a irregular grid is desired, or if the approximations in *DNFFT* are not acceptable. Because of its speed, *DNFFT* will usually be preferred to *DESKN*.

A Gaussian kernel estimate of the density at the point *y* is given by:

$$\hat{f}(y) = \frac{1}{nh} \sum_{i=1}^n \frac{1}{\sqrt{2\pi}} \exp \left[ -\frac{1}{2} \left( \frac{y - x_i}{h} \right)^2 \right]$$

where



$$\hat{f}(y)$$

is the estimated density at  $y$ ,  $x_i$  denotes the  $i$ -th observation,  $n$  is the number of observations, and  $h$  is a fixed constant called the *window width* supplied by the user. If density estimates for several different window sizes are to be computed, then `DNFFT` performs a fast Fourier transform on the data only during the first call (when `IFFT` is zero). On subsequent calls (with `IFFT` set at 1), the Fourier transform of the data need not be recomputed.

If the same value of `NXPT` is to be used with several different input vectors  $x$ , then the computations can be made faster by the use of `D2FFT`. In `D2FFT`, it is assumed that some constants required by the Fourier transform and its inverse have already been computed via routine `FFTRI` (IMSL MATH/LIBRARY) in work array `WFFTR`. If `D2FFT` is called repeatedly with the same value of `NXPT`, `WFFTR` need only be computed once.

Routine `DNFFT` is an implementation of *Applied Statistics* algorithm AS 176 (Silverman 1982) using IMSL routines for the fast Fourier transforms. Modification to algorithm AS 176, as discussed in Silverman (1986, pages 61–66), gives the details of the computational method. The basic idea is to partition the support of the density into `NXPT` equally-sized nonoverlapping intervals. The frequency of the observations within each interval is then computed, and the Fourier transform of the frequencies obtained. Since the kernel density estimate is the convolution of the frequencies with the Gaussian kernel (for given window size), the Fourier coefficients for the Gaussian kernel density estimates are computed as the product of the coefficients obtained for the frequencies, times the Fourier coefficients for the Gaussian kernel function. The discrete Fourier coefficients for the Gaussian kernel may be estimated from the continuous transform. The inverse transform is then used to obtain the density estimates.

Because the fast Fourier transform is used in computing

$$\hat{f}(y)$$

the computations are relatively fast (providing that `NXPT` is a product of small primes). To maintain precision, a large number of intervals, say 256, is usually recommended. Tapia and Thompson (1978), Chapter 2, give some considerations relevant to the choice of the window size parameter `WINDOW`. Generally, several different window sizes should be tried in order to obtain the best value for this parameter.

### Example

In this example, the density function is estimated at 64 points using a random sample of 150 points from a standard normal distribution. The actual density for the standard normal density is also reported in the output for comparison. The random sample is generated using routines `RNSET` (page 1166) and `RNNOR` (page 1208).

```
INTEGER      IFFT, NOBS, NXPT
REAL         CONS, WINDOW
```

```

      PARAMETER (CONS=0.39894228, IFFT=0, NOBS=150, NXPT=64,
&              WINDOW=0.25)
C
      INTEGER    I, NOUT, NRMISS
      REAL       BNDS(2), COEF(NXPT), DENS(NXPT), EXP, FREQ(1), STEP,
&              X(NOBS), XX
      INTRINSIC  EXP
      EXTERNAL   DNFFT, RNNOR, RNSET, UMACH
C
      DATA BNDS/-4.0, 3.875/
C
      CALL RNSET (123457)
      CALL RNNOR (NOBS, X)
C
      FREQ(1) = -1.0
      CALL DNFFT (NOBS, X, FREQ, BNDS, WINDOW, IFFT, NXPT, COEF,
&              DENS, NRMISS)
C
      CALL UMACH (2, NOUT)
      WRITE (NOUT,99998)
99998 FORMAT (' X DENSITY POPULATION')
      STEP = (BNDS(2)-BNDS(1))/(NXPT-1)
      XX = BNDS(1)
      DO 10 I=1, NXPT, 2
          WRITE (NOUT,99999) XX, DENS(I), CONS*EXP(-XX*XX/2.0)
99999  FORMAT (F6.2, 2F8.4)
          XX = XX + STEP*2.0
      10 CONTINUE
C
      END

```

### Output

```

X DENSITY POPULATION
-4.00 0.0000 0.0001
-3.75 0.0000 0.0004
-3.50 0.0000 0.0009
-3.25 0.0000 0.0020
-3.00 0.0001 0.0044
-2.75 0.0011 0.0091
-2.50 0.0089 0.0175
-2.25 0.0345 0.0317
-2.00 0.0772 0.0540
-1.75 0.1204 0.0863
-1.50 0.1573 0.1295
-1.25 0.2076 0.1826
-1.00 0.2682 0.2420
-0.75 0.2987 0.3011
-0.50 0.2976 0.3521
-0.25 0.3072 0.3867
 0.00 0.3336 0.3989
 0.25 0.3458 0.3867
 0.50 0.3169 0.3521
 0.75 0.2834 0.3011
 1.00 0.2683 0.2420
 1.25 0.2242 0.1826
 1.50 0.1557 0.1295
 1.75 0.1182 0.0863

```

2.00	0.0946	0.0540
2.25	0.0569	0.0317
2.50	0.0199	0.0175
2.75	0.0033	0.0091
3.00	0.0002	0.0044
3.25	0.0000	0.0020
3.50	0.0000	0.0009
3.75	0.0000	0.0004

---

## DESPT/DDESPT (Single/Double precision)

Estimate a probability density function at specified points using linear or cubic interpolation.

### Usage

CALL DESPT (NODE, XPT, IOPT, NORD, AMESH, DENS, DENEST)

### Arguments

**NODE** — Number of points at which the density is desired. (Input)

**XPT** — Vector of length **NODE** containing the points at which an estimate of the probability density is desired. (Input)

**IOPT** — Interpolation option parameter. (Input)

#### **IOPT** Method of interpolation

- |   |                                 |
|---|---------------------------------|
| 1 | Linear on equally spaced points |
| 2 | Linear with unequal spacing     |
| 3 | Cubic on equally spaced points  |
| 4 | Cubic with unequal spacing      |

**NORD** — Number of ordinates supplied. (Input)

**NORD** must be greater than one for linear interpolation, and greater than three for cubic interpolation.

**AMESH** — Vector of length **NORD** for **IOPT** = 2 or 4, and of length 2 for **IOPT** = 1 or 3. (Input)

If **IOPT** = 2 or 4, **AMESH(I)** contains the abscissas corresponding to each density estimate in **DENS(I)**. In this case, the abscissas must be specified in increasing order. If **IOPT** = 1 or 3 (i.e., for an equally spaced mesh), then the lower and upper ends of the mesh are specified by **AMESH(1)** and **AMESH(2)**, respectively, with the increment between mesh points given by  $(\text{AMESH}(2) - \text{AMESH}(1))/(\text{NORD} - 1)$ .

**DENS** — Vector of length **NORD** containing the density function values corresponding to each of the **NORD** abscissa values. (Input)

**DENEST** — Vector of length **NODE** containing the density function estimates for the points in **XPT**. (Output)

## Comments

- Automatic workspace usage is  
DESPT 6 \* NORD units if IOPT = 3, 5 \* NORD + 1 units if IOPT = 4, and  
1 unit otherwise, or  
DDESPT 12 \* NORD units if IOPT = 3, 10 \* NORD + 2 units if IOPT = 4,  
and 2 units otherwise.  
  
Workspace may be explicitly provided, if desired, by use of  
D2SPT/DD2SPT . The reference is  
CALL D2SPT (NODE, XPT, IOPT, NORD, AMESH, DENS,  
DENEST, CF, X, BREAK)  
  
The additional arguments are as follows:  
  
**CF** — Work vector of length 4 \* NORD for IOPT = 3 or 4. CF is not used  
for other values of IOPT and may be dimensioned of length 1.  
  
**X** — Work vector of length NORD for IOPT = 3 or 4. X is not used for  
other values of IOPT and may be dimensioned of length 1.  
  
**BREAK** — Work vector of length NORD for IOPT = 3 or 4. BREAK is not  
used for other values of IOPT and may be dimensioned of length 1.
- Array AMESH is the same as array BNDS in DESPL (page 1040) when  
IOPT is 1 or 3, and the same as array XPT in DESKN (page 1044) when  
IOPT is 2 or 4.

## Algorithm

Routine DESPT computes an estimate of a density function using either linear or cubic spline interpolation on a set  $\{(X_i, F_i), \text{ for } i = 1, \dots, N\}$ , where  $F_i = \text{DENS}(i)$ ,  $N = \text{NODE}$ , and where the values of the the grid points  $X_i$  can be obtained from the vector AMESH. The value of IOPT indicates the type of interpolation (linear or cubic) to be performed and whether the mesh values are equally spaced. When IOPT is 1 or 3, then an equally spaced mesh is used with mesh values given by

$$\text{AMESH}(1) + i * \text{DELTA}$$

for  $i = 0, 1, \dots, N - 1$ , where

$$\text{DELTA} = (\text{AMESH}(2) - \text{AMESH}(1)) / (\text{NORD} - 1)$$

IOPT = 2 or 4 yields an unequally spaced mesh with all mesh values contained in the vector AMESH.

The Akima cubic spline method of interpolation (Akima 1970) is used for the cubic interpolation.

### Example

The standard normal density is to be estimated via a grid of points over which the density is provided. Grid points are given by (0.0, 0.5, 1.0, 1.5, 2.0) while the density is to be estimated (via linear interpolation) at the four points (0.25, 0.75, 1.25, 1.75). For comparison, both the exact and the estimated density values at each of the four points are printed.

```
INTEGER      IOPT, NODE, NORD
PARAMETER   (IOPT=1, NODE=4, NORD=5)
C
INTEGER      I, NOUT
REAL         AMESH(2), DENEST(NODE), DENS(NORD), EXP, F, H, X, X0,
&           XPT(NODE)
INTRINSIC    EXP
EXTERNAL     DESPT, UMACH
C
DATA XPT/0.25, 0.75, 1.25, 1.75/
DATA AMESH/0, 2/
C
F(X) = 0.3989423*EXP(-X*X/2.0)
C                               Get the grid values
H = (AMESH(2)-AMESH(1))/(NORD-1)
X0 = AMESH(1)
DO 10 I=1, NORD
    DENS(I) = F(X0)
    X0 = X0 + H
10 CONTINUE
C                               Get the density estimates
CALL DESPT (NODE, XPT, IOPT, NORD, AMESH, DENS, DENEST)
C                               Print the results
CALL UMACH (2, NOUT)
WRITE (NOUT, '( ' X DENEST EXACT ' )' )
DO 20 I=1, NODE
    WRITE (NOUT, '(F5.2, 2F12.5)' ) XPT(I), DENEST(I), F(XPT(I))
20 CONTINUE
END
```

### Output

X	DENEST	EXACT
0.25	0.37550	0.38667
0.75	0.29702	0.30114
1.25	0.18574	0.18265
1.75	0.09175	0.08628

---

## HAZRD/DHAZRD (Single/Double precision)

Perform nonparametric hazard rate estimation using kernel functions and quasi-likelihoods.

### Usage

```
CALL HAZRD (NOBS, X, LDX, IRT, ICEN, IWTO, NGRID, BSTART,
            GINC, KMIN, INK, NK, IPRINT, ISORT, ST, JCEN,
            ALPHA, BETA, K, VML, H, NMISS)
```

## Arguments

**NOBS** — Number of observations. (Input)

**X** — NOBS by  $m$  matrix containing the raw data, where  $m = 1$  if ICEN = 0, and  $m = 2$  otherwise. (Input)

**LDX** — Leading dimension of  $x$  exactly as specified in the dimension statement in the calling program. (Input)

**IRT** — Column number in  $x$  of the event times. (Input)

**ICEN** — Censoring option. (Input)

If ICEN = 0, then all of the data is treated as exact data with no censoring. For ICEN > 0, column ICEN of  $x$  contains the censoring codes. A censoring code of 0 means an exact event (failure). A censoring code of 1 means that the observation was right censored at the event time.

**IWTO** — Weight option. (Input)

If IWTO = 1, then weight  $\ln(1 + 1/(\text{NOBS} - i + 1))$  is used for the  $i$ -th smallest observation. Otherwise, weight  $1/(\text{NOBS} - i + 1)$  is used.

**NGRID** — Grid option. (Input)

If NGRID = 0, a default grid is used to locate an initial starting value for parameter BETA. For NGRID > 0, a user-defined grid is used. This grid is defined as  $\text{BSTART} + (j - 1) * \text{GINC}$ , for  $j = 1, \dots, \text{NGRID}$ , where BSTART, GINC, and NGRID are input.

**BSTART** — First value to be used in the user-defined grid. (Input)

Not used if NGRID = 0.

**GINC** — For a user-defined grid, the increment between successive grid values of BETA. (Input)

Not used if NGRID = 0.

**KMIN** — Minimum number for parameter  $k$ . (Input)

Parameter  $k$  is the number of nearest neighbors to be used in computing the  $k$ -th nearest neighbor distance.

**INK** — Increment between successive values of parameter  $k$ . (Input)

**NK** — Number of values of  $k$  to be considered. (Input)

HAZRD finds the optimal value of  $k$  over the grid given by:  $\text{KMIN} + (j - 1) * \text{INK}$ , for  $j = 1, \dots, \text{NK}$ .

**IPRINT** — Printing option. (Input)

If IPRINT = 1, the grid estimates and the optimized estimates are printed for each value of  $k$ . Otherwise, no printing is performed.

**ISORT** — Sorting option. (Input)

If ISORT = 1, then the event times are not automatically sorted by HAZRD. Otherwise, sorting is performed with exact failure times following tied right-censored times.

**ST** — Vector of length `NOBS` containing the times of occurrence of the events, sorted from smallest to largest. (Output)

Vector `ST` is obtained from the matrix `X` and should be used as input to routine `HAZST` (page 1069).

**JCEN** — Vector of length `NOBS` containing the sorted censor codes. (Output)

Censor codes are sorted corresponding to the events `ST(i)`, with censored observations preceding tied failures. Vector `JCEN` is obtained from the censor codes in `X`, if present, and is used as input to routine `HAZST` (page 1069).

**ALPHA** — Optimal estimate for the parameter  $\alpha$ . (Output)

**BETA** — Optimal estimate for the parameter  $\beta$ . (Output)

**K** — Optimal estimate for the parameter  $k$ . (Output)

**VML** — Optimum value of the criterion function. (Output)

**H** — Vector of length `NOBS * 5` containing constants needed to compute the  $k$ -th nearest failure distances, and the observation weights. (Output)

`H` is used as input to routine `HAZST` (page 1069).

**NMISS** — Number of missing (NaN, not a number) values in `X`. (Output)

### Comments

1. Informational Errors  

Type	Code	
4	18	All observations are missing (NaN, not a number) values.
2. In the optimization routines, the parameterization is changed to  $\beta^*$  and  $\alpha^*$ , where  $\beta^* = -\ln(\beta)$  and  $\alpha^* = -\ln(\alpha)$ . The default grid uses  $-8, -4, -3, -2.5, -2, -1.5, -1, -0.5$ , and  $0.5$  for  $\beta^*$ . This corresponds to a grid in  $\beta$  of  $2981, 54.6, 20.08, 12.18, 7.39, 4.48, 2.72, 1.64$ , and  $.61$ . The grid  $\beta$  that maximizes the modified “likelihood” is used as the starting point for the iterations.
3. If the initial estimate of  $\beta$  as determined from the grid or as given by the user is greater than 400 (actually  $e^6$ ), then infinite  $\beta$  is assumed, and an analytic estimate of  $\alpha$  based upon infinite  $\beta$  is used. In the optimization, if it is determined that  $\beta$  must be greater than 1000, then an infinite  $\beta$  is assumed. Infinite  $\beta$  corresponds to a “flat” hazard rate.

### Algorithm

Routine `HAZRD` is an implementation of the methods discussed by Tanner and Wong (1984) for estimating the hazard rate in survival or reliability data with right censoring. It uses the biweight kernel,

$$K(x) = \begin{cases} \frac{15}{16}(1-x^2)^2 & \text{for } |x| < 1 \\ 0 & \text{elsewhere} \end{cases}$$

and a modified likelihood to obtain data-based estimates of the smoothing parameters  $\alpha$ ,  $\beta$ , and  $k$  needed in the estimation of the hazard rate. For kernel  $K(x)$ , define the “smoothed” kernel  $K_s(x - x_{(j)})$  as follows:

$$K_s(x - x_{(j)}) = \frac{1}{\alpha d_{jk}} K\left(\frac{x - x_{(j)}}{\beta d_{jk}}\right)$$

where  $d_{jk}$  is the distance to the  $k$ -th nearest failure from  $x_{(j)}$ , and  $x_{(j)}$  is the  $j$ -th ordered observation (from smallest to largest). For given  $\alpha$  and  $\beta$ , the hazard at point  $x$  is then

$$h(x) = \sum_{i=1}^N \{(1 - \delta_i) w_i K_s(x - x_{(i)})\}$$

where  $N = \text{NOBS}$ ,  $\delta_i$  is the  $i$ -th observation’s censor code (1 = censored, 0 = failed), and  $w_i$  is the  $i$ -th ordered observation’s weight, which may be chosen as either  $1/(N - i + 1)$ , or  $\ln(1 + 1/(N - i + 1))$ . After the smoothing parameters have been obtained, the hazard may be estimated via HAZST (page 1069).

Let

$$H(x) = \int_0^x h(s) ds$$

The likelihood is given by

$$L = \prod_{i=1}^N \{h(x_i)^{(1-\delta_i)} \exp(-H(x_i))\},$$

where  $\Pi$  denotes product. Since the likelihood leads to degenerate estimates, Tanner and Wong (1984) suggest the use of a modified likelihood. The modification consists of deleting observation  $x_i$  in the calculation of  $h(x_i)$  and  $H(x_i)$  when the likelihood term for  $x_i$  is computed using the usual optimization techniques.  $\alpha$  and  $\beta$  for given  $k$  can then be estimated.

Estimates for  $\alpha$  and  $\beta$  are computed as follows: for given  $\beta$ , a closed form solution is available for  $\alpha$ . The problem is thus reduced to the estimation of  $\beta$ . A grid search for  $\beta$  is first performed. Experience indicates that if the initial estimate of  $\beta$  from this grid search is greater than, say,  $e^6$ , then the modified likelihood is degenerate because the hazard rate does not change with time. In this situation,  $\beta$  should be taken to be infinite, and an estimate of  $\alpha$  corresponding to infinite  $\beta$  should be directly computed. When the estimate of  $\beta$  from the grid search is less than  $e^6$ , a secant algorithm is used to optimize the modified likelihood. The secant algorithm iteration stops when the change in  $\beta$  from one iteration to the next is less than  $10^{-5}$ . Alternatively, the iterations may



cease when the value of  $\beta$  becomes greater than  $e^6$ , at which point an infinite  $\beta$  with a degenerate likelihood is assumed.

To find the optimum value of the likelihood with respect to  $k$ , a user-specified grid of  $k$ -values is used. For each grid value, the modified likelihood is optimized with respect to  $\alpha$  and  $\beta$ . That grid point, which leads to the smallest likelihood, is taken to be the optimal  $k$ .

### Programming Notes

1. The routine HAZST (page 1069) may be used to estimate the hazard on a grid of points once the optimal values for  $\alpha$ ,  $\beta$  and  $k$  have been found. The user should also consider using the “easy-to-use” version of HAZRD, routine HAZEZ (page 1061).
2. If sorting of the data is performed by HAZRD, then the sorted array will be such that all censored observations at a given time precede all failures at that time. To specify an arbitrary pattern of censored/failed observations at a given time point, the ISORT = 1 option must be used. In this case, it is assumed that the times have already been sorted from smallest to largest.
3. The smallest value of  $k$  must be greater than the largest number of tied failures since  $d_{jk}$  must be positive for all  $j$ . (Censored observations are not counted.) Similarly, the largest value of  $k$  must be less than the total number of failures. If the grid specified for  $k$  includes values outside the allowable range, then a warning error is issued; but  $k$  is still optimized over the allowable grid values.
4. The secant algorithm iterates on the transformed parameter  $\beta^* = \exp(-\beta)$ . This assures a positive  $\beta$ , and it also seems to lead to a more desirable grid search. All results returned to the user are in the original parameterization, however.
5. Since local minimums have been observed in the modified likelihood, it is recommended that more than one grid of initial values for  $\alpha$  and  $\beta$  be used.

### Example

The following example is taken from Tanner and Wong (1984). The data are from Stablein, Carter, and Novak (1981) and involve the survival times of individuals with nonresectable gastric carcinoma. Individuals treated with radiation and chemotherapy are used. For each value of  $k$  from 18 to 22 with increment of 2, the default grid search for  $\beta$  is performed. Using the optimal value of  $\beta$  in the grid, the optimal parameter estimates of  $\alpha$  and  $\beta$  are computed for each value of  $k$ . The final solution is the parameter estimates for the value of  $k$  which optimizes the modified likelihood (VML). Because the IPRINT = 1 option is in effect, HAZRD prints all of the results in the output.

```

    INTEGER    ICEN, INK, IPRINT, IRT, ISORT, IWTO, KMIN, LDX,
&             NGRID, NK, NOBS
    REAL       BSTART, GINC
    PARAMETER  (BSTART=0.0, GINC=0.0, ICEN=2, INK=2, IPRINT=1,
&             IRT=1, ISORT=1, IWTO=0, KMIN=18, LDX=45, NGRID=0,
&             NK=3, NOBS=45)
C
    INTEGER    JCEN(NOBS), K, NMISS, NOUT
    REAL       ALPHA, BETA, H(5*NOBS), ST(NOBS), VML, X(NOBS,2)
    EXTERNAL   HAZRD, UMACH, WRIRN, WRRRN
C
    DATA X/17, 42, 44, 48, 60, 72, 74, 95, 103, 108, 122, 144, 167,
&         170, 183, 185, 193, 195, 197, 208, 234, 235, 254, 307, 315,
&         401, 445, 464, 484, 528, 542, 567, 577, 580, 795, 855, 882,
&         892, 1031, 1033, 1306, 1335, 1366, 1452, 1472, 36*0, 9*1/
C
    CALL HAZRD (NOBS, X, LDX, IRT, ICEN, IWTO, NGRID, BSTART, GINC,
&             KMIN, INK, NK, IPRINT, ISORT, ST, JCEN, ALPHA, BETA,
&             K, VML, H, NMISS)
C
    CALL UMACH (2, NOUT)
    WRITE (NOUT,99999) NMISS
99999 FORMAT (' NMISS = ', I4/)
    CALL WRRRN ('ST', 1, NOBS, ST, 1, 0)
    CALL WRIRN ('JCEN', 1, NOBS, JCEN, 1, 0)
    CALL WRRRN ('H', NOBS, 5, H, NOBS, 0)
    END

```

### Output

```

*** GRID SEARCH FOR K =      18 ***
ALPHA          BETA          VML
4.578322      2980.958008     -266.804504
4.543117          54.598148     -266.619690
4.336464          20.085537     -265.541168
4.019334          12.182494     -264.001404
3.542742          7.389056      -262.540100
2.990577          4.481689      -262.511810
2.351537          2.718282      -262.633911
1.584173          1.648721      -262.158264
0.966332          1.000000      -262.868408

```

```

*** OPTIMAL PARAMETER ESTIMATES ***
ALPHA          BETA          VML
1.695147          1.769263      -262.118530

```

```

*** GRID SEARCH FOR K =      20 ***
ALPHA          BETA          VML
4.053934      2980.958008     -266.525970
4.032835          54.598148     -266.401428
3.905046          20.085537     -265.648315
3.687815          12.182494     -264.401672
3.304344          7.389056      -262.665924
2.822716          4.481689      -262.080078
2.252759          2.718282      -262.445251
1.555777          1.648721      -261.772278
0.955586          1.000000      -262.617645

```

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*  
 ALPHA BETA VML  
 1.540533 1.631551 -261.771484

\*\*\* GRID SEARCH FOR K = 22 \*\*\*  
 ALPHA BETA VML  
 3.656405 2980.958008 -267.595337  
 3.641593 54.598148 -267.498596  
 3.550560 20.085537 -266.903870  
 3.388752 12.182494 -265.859131  
 3.071474 7.389056 -264.066040  
 2.645036 4.481689 -263.038696  
 2.137399 2.718282 -263.334717  
 1.512606 1.648721 -262.639740  
 0.936368 1.000000 -262.682739

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*  
 ALPHA BETA VML  
 1.342176 1.450016 -262.561188

\*\*\* THE FINAL SOLUTION (K = 20) \*\*\*  
 ALPHA BETA VML  
 1.540533 1.631551 -261.771484

NMISS = 0

ST							
1	2	3	4	5	6	7	8
17.0	42.0	44.0	48.0	60.0	72.0	74.0	95.0
9	10	11	12	13	14	15	16
103.0	108.0	122.0	144.0	167.0	170.0	183.0	185.0
17	18	19	20	21	22	23	24
193.0	195.0	197.0	208.0	234.0	235.0	254.0	307.0
25	26	27	28	29	30	31	32
315.0	401.0	445.0	464.0	484.0	528.0	542.0	567.0
33	34	35	36	37	38	39	40
577.0	580.0	795.0	855.0	882.0	892.0	1031.0	1033.0
41	42	43	44	45			
1306.0	1335.0	1366.0	1452.0	1472.0			

JCEN																			
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	1	1
41	42	43	44	45															
1	1	1	1	1															

H				
1	2	3	4	5
1	217.0	218.0	1.0	21.0
				1.0

2	192.0	193.0	1.0	21.0	0.5
3	190.0	191.0	1.0	21.0	0.3
4	186.0	187.0	1.0	21.0	0.2
5	174.0	175.0	1.0	21.0	0.2
6	162.0	163.0	1.0	21.0	0.2
7	160.0	161.0	1.0	21.0	0.1
8	139.0	140.0	1.0	21.0	0.1
9	131.0	132.0	1.0	21.0	0.1
10	126.0	127.0	1.0	21.0	0.1
11	112.0	113.0	1.0	21.0	0.1
12	102.0	110.0	2.0	22.0	0.1
13	123.0	125.0	3.0	23.0	0.1
14	126.0	128.0	3.0	23.0	0.1
15	132.0	135.0	5.0	25.0	0.1
16	130.0	137.0	5.0	25.0	0.1
17	133.0	145.0	5.0	25.0	0.1
18	135.0	147.0	5.0	25.0	0.1
19	137.0	149.0	5.0	25.0	0.1
20	148.0	160.0	5.0	25.0	0.1
21	167.0	174.0	6.0	26.0	0.0
22	166.0	175.0	6.0	26.0	0.0
23	182.0	191.0	6.0	26.0	0.0
24	204.0	212.0	9.0	29.0	0.0
25	212.0	213.0	9.0	29.0	0.0
26	231.0	234.0	14.0	34.0	0.0
27	275.0	278.0	14.0	34.0	0.0
28	294.0	297.0	14.0	34.0	0.0
29	311.0	314.0	15.0	35.0	0.0
30	343.0	345.0	16.0	36.0	0.0
31	357.0	359.0	16.0	38.0	0.0
32	382.0	384.0	16.0	38.0	0.0
33	392.0	394.0	16.0	38.0	0.0
34	395.0	397.0	16.0	38.0	0.0
35	610.0	612.0	16.0	43.0	0.0
36	670.0	672.0	16.0	45.0	0.0
37	689.0	697.0	17.0	45.0	0.0
38	699.0	707.0	17.0	45.0	0.0
39	838.0	846.0	17.0	45.0	0.0
40	840.0	848.0	17.0	45.0	0.0
41	1113.0	1121.0	17.0	45.0	0.0
42	1142.0	1150.0	17.0	45.0	0.0
43	1173.0	1181.0	17.0	45.0	0.0
44	1259.0	1267.0	17.0	45.0	0.0
45	1279.0	1287.0	17.0	45.0	0.0

---

## HAZEZ/DHAZEZ (Single/Double precision)

Perform nonparametric hazard rate estimation using kernel functions. Easy-to-use version of HAZRD.

### Usage

```
CALL HAZEZ (NOBS, X, LDX, IRT, ICEN, IPRINT, ST, JCEN,
           ALPHA, BETA, K, VML, H, NMISS)
```

## Arguments

**NOBS** — Number of observations. (Input)

**X** — NOBS by  $m$  matrix containing the raw data, where  $m = 1$  if ICEN = 0, and  $m = 2$  otherwise. (Input)

**LDX** — Leading dimension of  $x$  exactly as specified in the dimension statement in the calling program. (Input)

**IRT** — Column number in  $x$  containing the times of occurrence of the events. (Input)

**ICEN** — Censoring option. (Input)

If ICEN = 0, then all of the data is treated as exact data with no censoring. For ICEN > 0, column ICEN of  $x$  contains the censoring codes. A censoring code of 0 means an exact event (failure). A censoring code of 1 means that the observation was right censored at the event time.

**IPRINT** — Printing option. (Input)

If IPRINT = 1, the grid estimates and the optimized estimates are printed for each value of  $k$ . Otherwise, no printing is performed.

**ST** — Vector of length NOBS containing the times of occurrence of the events, sorted from smallest to largest. (Output)

Vector ST is obtained from matrix  $x$  and is used as input to routine HAZST (page 1069).

**JCEN** — Vector of length NOBS containing the sorted censor codes. (Output)

Censor codes are sorted corresponding to the events  $ST(i)$ , with censored observations preceding tied failures. Vector JCEN is obtained from the censor codes in  $x$  and is used as input to routine HAZST (page 1069).

**ALPHA** — Optimal estimate for the parameter  $\alpha$ . (Output)

**BETA** — Optimal estimate for the parameter  $\beta$ . (Output)

**K** — Optimal estimate for the parameter  $k$ . (Output)

**VML** — Optimal value of the criterion function. (Output)

VML is the “modified likelihood”.

**H** — Vector of length  $5 * NOBS$  containing the constants needed to compute the  $k$ -th nearest failure distance and the observation weights. (Output)

H is used as input to routine HAZST (page 1069).

**NMISS** — Number of missing (NaN, not a number) values in  $x$ . (Output)

## Comments

1. Informational errors

Type	Code
------	------

4	6	All observations are missing (NaN, not a number) values.
---	---	--

4 7 There are not enough failing observations in  $x$  to continue.

2. The grid values in the initial grid search are given as follows: Let  $\beta^* = -8, -4, -2, -1, -0.5, 0.5, 1,$  and  $2,$  and

$$\beta = e^{-\beta^*}$$

For each value of  $\beta$ ,  $\text{VML}$  is computed at the optimizing  $\beta$ . The maximizing  $\beta$  is used to initiate the iterations.

3. If the initial  $\beta^*$  is determined from the grid search to be less than  $-6$ , then it is presumed that  $\beta$  is infinite, and an analytic estimate of  $\alpha$  based upon infinite  $\beta$  is used. Infinite  $\beta$  corresponds to a flat hazard rate.

### Algorithm

Routine `HAZEZ` is an implementation of the methods discussed by Tanner and Wong (1984) for estimating the hazard rate in survival or reliability data with right censoring. It uses the biweight kernel,

$$K(x) = \begin{cases} \frac{15}{16}(1-x^2)^2 & \text{for } |x| < 1 \\ 0 & \text{elsewhere} \end{cases}$$

and a modified likelihood to obtain data-based estimates of the smoothing parameters  $\alpha$ ,  $\beta$ , and  $k$  needed in the estimation of the hazard rate. For kernel  $K(x)$ , define the “smoothed” kernel  $K_s(x - x(j))$  as follows:

$$K_s(x - x(j)) = \frac{1}{\alpha d_{jk}} K\left(\frac{x - x(j)}{\beta d_{jk}}\right)$$

where  $d_{jk}$  is the distance to the  $k$ -th nearest failure from  $x(j)$ , and  $x(j)$  is the  $j$ -th ordered observation (from smallest to largest). For given  $\alpha$  and  $\beta$ , the hazard at point  $x$  is given by:

$$h(x) = \sum_{i=1}^N \{(1 - \delta_i) w_i K_s(x - x(i))\}$$

where  $N = \text{NOBS}$ ,  $\delta_i$  is the censor code ( $0 = \text{failed}$ ,  $1 = \text{censored}$ ) for the  $i$ -th ordered observation, and  $w_i$  is the weight of the  $i$ -th ordered observation (given by  $1/(N - i + 1)$ ). The hazard may be estimated via routine `HAZST` (page 1069) after the smoothing parameters have been obtained

Let

$$H(x) = \int_0^x h(s) ds$$

The likelihood is given by:

$$L = \prod_{i=1}^N \{h(x_i)^{(1-\delta_i)} \exp(-H(x_i))\}$$

where  $\Pi$  denotes product. Since the likelihood, as specified, will lead to degenerate estimates, Tanner and Wong (1984) suggest the use of a modified likelihood. The modification consists of deleting the observation  $x_i$  in the calculation of  $h(x_i)$  and  $H(x_i)$  when the likelihood term for  $x_i$  is computed. For a given  $k$ ,  $\alpha$  and  $\beta$  can then be estimated via the usual optimization techniques.

Estimates for  $\alpha$  and  $\beta$  are computed as follows. For a given  $\beta$ , a closed form solution is available for  $\alpha$ . The problem is thus reduced to the estimation of  $\beta$ . To estimate  $\alpha$  and  $\beta$ , a grid search is first performed. Experience indicates that if the initial estimate of  $\beta$  from this grid search is greater than  $\exp(6)$ , then the modified likelihood is degenerate because the hazard rate does not change with time. In this situation,  $\beta$  should be taken to be infinite, and an estimate of  $\alpha$  corresponding to infinite  $\beta$  is computed directly. When the estimate of  $\beta$  from the grid search is less than  $\exp(6)$  (approximately 400), a secant algorithm is used to optimize the modified likelihood. The secant algorithm is said to have converged when the change in  $\beta$  from one iteration to the next is less than 0.00001. Additionally, convergence is assumed when the value of  $\beta$  becomes greater than  $\exp(6)$ . This corresponds to an infinite  $\beta$  with a degenerate likelihood.

A grid of  $k$ -values is used to find the optimum value of the likelihood with respect to  $k$ . The grid is determined by HAZEZ and consists of at most 10 points. The starting value in the grid is the smallest possible value of  $k$ . An increment of 2 is then used to obtain the remaining grid points.

For each grid value, the modified likelihood is optimized with respect to  $\alpha$  and  $\beta$ . That grid point, which leads to the smallest likelihood, is taken to be the optimal  $k$ .

### Programming Notes

1. Routine HAZST (page 1069) may be used to estimate the hazard on a grid of points once the optimal values for  $\alpha$ ,  $\beta$  and  $k$  have been found. (The user should also consider using routine HAZRD (page 1054), which allows for more options than HAZEZ.)
2. Routine HAZEZ assumes that censored observations precede failed observations at tied failure/censoring times.
3. The secant algorithm iterates on the transformed parameter  $\beta^* = \exp(-\beta)$ . This assures a positive  $\beta$ , and it also seems to lead to a more desirable grid search. All results returned to the user are in the original parameterization.

## Example

The following example is illustrated in Tanner and Wong (1984), and the data are taken from Stablein, Carter, and Novak (1981). It involves the survival times of individuals with nonresectable gastric carcinoma. Only those individuals treated with radiation and chemotherapy are used.

```
INTEGER ICEN, IPRINT, IRT, LDX, NOBS
PARAMETER (ICEN=2, IPRINT=1, IRT=1, LDX=45, NOBS=45)
C
INTEGER JCEN(NOBS), K, NMISS, NOUT
REAL ALPHA, BETA, H(5*NOBS), ST(NOBS), VML, X(NOBS,2)
EXTERNAL HAZEZ, UMACH, WRIRN, WRRRN
C
DATA X/17, 42, 44, 48, 60, 72, 74, 95, 103, 108, 122, 144, 167,
& 170, 183, 185, 193, 195, 197, 208, 234, 235, 254, 307, 315,
& 401, 445, 464, 484, 528, 542, 567, 577, 580, 795, 855, 882,
& 892, 1031, 1033, 1306, 1335, 1366, 1452, 1472, 36*0, 9*1/
C
CALL HAZEZ (NOBS, X, LDX, IRT, ICEN, IPRINT, ST, JCEN, ALPHA,
& BETA, K, VML, H, NMISS)
C
CALL UMACH (2, NOUT)
WRITE (NOUT,99999) NMISS
99999 FORMAT (/ ' NMISS = ', I4/)
CALL WRRRN ('ST', 1, NOBS, ST, 1, 0)
CALL WRIRN ('JCEN', 1, NOBS, JCEN, 1, 0)
CALL WRRRN ('H', NOBS, 5, H, NOBS, 0)
END
```

## Output

```
*** GRID SEARCH FOR K =      2 ***
ALPHA      BETA      VML
65.157967  2980.958008  -266.543945
32.434208   54.598148    -262.551147
17.100269   20.085537    -263.100769
11.402525   12.182494    -264.410187
 7.263529    7.389056    -267.502014
 4.452315    4.481689    -270.548523
 2.689497    2.718282    -335.407288
 1.633702    1.648721    -338.566162
 0.995799    1.000000    -519.939514

*** OPTIMAL PARAMETER ESTIMATES ***
ALPHA      BETA      VML
32.219337   53.968315    -262.550781

*** GRID SEARCH FOR K =      4 ***
ALPHA      BETA      VML
25.596716  2980.958008  -266.471558
20.476425   54.598148    -262.893860
13.995192   20.085537    -262.792755
10.109113   12.182494    -262.573212
 6.883837    7.389056    -263.030121
 4.407142    4.481689    -265.238647
 2.690131    2.718282    -265.606293
 1.633339    1.648721    -266.822693
```



0.993371 1.000000 -271.831390

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
8.530729	9.683726	-262.545593

\*\*\* GRID SEARCH FOR K = 6 \*\*\*

ALPHA	BETA	VML
16.828691	2980.958008	-266.729248
14.840095	54.598148	-264.019409
11.215133	20.085537	-262.844360
9.013870	12.182494	-263.663391
6.557755	7.389056	-263.283752
4.330785	4.481689	-263.732697
2.691744	2.718282	-264.613800
1.633932	1.648721	-265.381866
0.990891	1.000000	-266.242767

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
12.553377	28.178671	-262.529877

\*\*\* GRID SEARCH FOR K = 8 \*\*\*

ALPHA	BETA	VML
11.377748	2980.958008	-266.746185
10.773529	54.598148	-265.469299
8.766835	20.085537	-262.476807
7.427887	12.182494	-263.109009
5.916299	7.389056	-264.492432
4.184323	4.481689	-264.289886
2.656351	2.718282	-264.807617
1.623750	1.648721	-265.270691
0.989442	1.000000	-264.738403

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
8.522110	18.281288	-262.438568

\*\*\* GRID SEARCH FOR K = 10 \*\*\*

ALPHA	BETA	VML
8.689023	2980.958008	-267.026093
8.412854	54.598148	-266.250366
7.196551	20.085537	-263.192688
6.207793	12.182494	-262.648376
5.143391	7.389056	-264.274384
3.934601	4.481689	-264.523193
2.630993	2.718282	-264.877869
1.611710	1.648721	-264.332581
0.984530	1.000000	-263.920013

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
6.483376	13.956067	-262.589661

\*\*\* GRID SEARCH FOR K = 12 \*\*\*

ALPHA	BETA	VML
6.669007	2980.958008	-266.778259
6.551508	54.598148	-266.347595

5.933167	20.085537	-264.141174
5.252526	12.182494	-262.516205
4.471936	7.389056	-262.691589
3.598284	4.481689	-263.914032
2.557817	2.718282	-263.390106
1.588307	1.648721	-263.879578
0.973723	1.000000	-263.361908

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
4.923379	9.819798	-262.336670

\*\*\* GRID SEARCH FOR K = 14 \*\*\*

ALPHA	BETA	VML
5.668086	2980.958008	-266.747559
5.595870	54.598148	-266.436584
5.195685	20.085537	-264.737946
4.685275	12.182494	-262.971497
4.044650	7.389056	-262.288147
3.335586	4.481689	-263.126434
2.496436	2.718282	-262.891663
1.585763	1.648721	-263.418976
0.969140	1.000000	-263.164032

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
4.145060	7.966486	-262.260559

\*\*\* GRID SEARCH FOR K = 16 \*\*\*

ALPHA	BETA	VML
4.970138	2980.958008	-266.419281
4.924928	54.598148	-266.199646
4.663393	20.085537	-264.938660
4.280633	12.182494	-263.266602
3.741570	7.389056	-262.020355
3.132969	4.481689	-262.401733
2.421248	2.718282	-262.555817
1.586469	1.648721	-262.426025
0.967658	1.000000	-263.135101

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
3.639074	6.767537	-261.987305

\*\*\* GRID SEARCH FOR K = 18 \*\*\*

ALPHA	BETA	VML
4.578322	2980.958008	-266.804504
4.543117	54.598148	-266.619690
4.336464	20.085537	-265.541168
4.019334	12.182494	-264.001404
3.542742	7.389056	-262.540100
2.990577	4.481689	-262.511810
2.351537	2.718282	-262.633911
1.584173	1.648721	-262.158264
0.966332	1.000000	-262.868408

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
-------	------	-----

1.695147                      1.769263                      -262.118530

\*\*\* GRID SEARCH FOR K =    20 \*\*\*

ALPHA	BETA	VML
4.053934	2980.958008	-266.525970
4.032835	54.598148	-266.401428
3.905046	20.085537	-265.648315
3.687815	12.182494	-264.401672
3.304344	7.389056	-262.665924
2.822716	4.481689	-262.080078
2.252759	2.718282	-262.445251
1.555777	1.648721	-261.772278
0.955586	1.000000	-262.617645

\*\*\* OPTIMAL PARAMETER ESTIMATES \*\*\*

ALPHA	BETA	VML
1.540533	1.631551	-261.771484

\*\*\* THE FINAL SOLUTION    (K =    20) \*\*\*

ALPHA	BETA	VML
1.540533	1.631551	-261.771484

NMISS =    0

ST							
1	2	3	4	5	6	7	8
17.0	42.0	44.0	48.0	60.0	72.0	74.0	95.0
9	10	11	12	13	14	15	16
103.0	108.0	122.0	144.0	167.0	170.0	183.0	185.0
17	18	19	20	21	22	23	24
193.0	195.0	197.0	208.0	234.0	235.0	254.0	307.0
25	26	27	28	29	30	31	32
315.0	401.0	445.0	464.0	484.0	528.0	542.0	567.0
33	34	35	36	37	38	39	40
577.0	580.0	795.0	855.0	882.0	892.0	1031.0	1033.0
41	42	43	44	45			
1306.0	1335.0	1366.0	1452.0	1472.0			

JCEN																			
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	1	1
41	42	43	44	45															
1	1	1	1	1															

H					
	1	2	3	4	5
1	217.0	218.0	1.0	21.0	1.0
2	192.0	193.0	1.0	21.0	0.5
3	190.0	191.0	1.0	21.0	0.3

```

      .
      .
      .
43  1173.0  1181.0  17.0  45.0  0.0
44  1259.0  1267.0  17.0  45.0  0.0
45  1279.0  1287.0  17.0  45.0  0.0

```

---

## HAZST/DHAZST (Single/Double precision)

Perform hazard rate estimation over a grid of points using a kernel function.

### Usage

```
CALL HAZST (NOBS, ST, JCEN, IWTO, NGRID, GSTRT, GINC,
           ALPHA, BETA, K, IHCOMP, H, HAZ)
```

### Arguments

**NOBS** — Number of observations. (Input)

If HAZRD (page 1054) or HAZEZ (page 1061) is called prior to this routine and the original data contained missing values, then NOBS in HAZST must be adjusted for the number of missing values from the value used in HAZRD or HAZEZ. That is, NOBS in HAZST is NOBS minus NMISS from HAZRD or HAZEZ .

**ST** — Vector of length NOBS containing the event times, sorted in ascending order. (Input)

ST may not contain missing values.

**JCEN** — Vector of length NOBS containing the censor codes. (Input)

JCEN(*i*) = 1 means that event *i* was (right) censored at time ST(*i*), *i* = 1, ..., NOBS.  
 JCEN(*i*) = 0 means that event *i* was a failure at time ST(*i*).

**IWTO** — Weighting option. (Input)

IWTO = 1 means use weights  $\ln(1 + 1/(\text{NOBS} - i + 1))$ . IWTO = 0 means use weights  $1/(\text{NOBS} - i + 1)$ . Not used if IHCOMP = 1.

**NGRID** — Number of grid points at which to compute the hazard. (Input)

**GSTRT** — First grid value. (Input)

**GINC** — Increment between grid values. (Input)

**ALPHA** — Value for parameter  $\alpha$ . (Input)

**BETA** — Value for parameter  $\beta$ . (Input)

**K** — Value for parameter *k*. (Input)

**IHCOMP** — Option parameter. (Input)

If IHCOMP = 0, H is computed. If IHCOMP = 1, H has already been computed (generally by HAZRD or HAZEZ).

**H** — Vector of length 5 \* NOBS containing the constants used in computing the *k*-th failure distance. (Input, if IHCOMP = 1; Output, otherwise)

**HAZ** — Vector of length `NGRID` containing the estimated hazard rates. (Output)

### Comments

1. Informational error
 

Type	Code	
4	13	At least one missing (NaN, not a number) value was found in <code>ST</code> . Missing values are not allowed in this routine.
2. The user-defined grid is given by  $\text{GSTRT} + j * \text{GINC}, j = 0, \dots, \text{NGRID} - 1$ .
3. Routine `HAZST` assumes that the grid points are new data points.

### Algorithm

Routine `HAZST` estimates the hazard function by use of the biweight kernel,

$$K(x) = \frac{15}{16}(1 - x^2)^2$$

Because a “smoothed” estimate is computed, one generally would use either routine `HAZRD` (page 1054) or routine `HAZEZ` (page 1061) to obtain maximum (modified) likelihood estimates of the smoothing parameters  $\alpha$ ,  $\beta$ , and  $k$ . Maximum (modified) likelihood estimates of these parameters are not required, however. A user-specified grid of points is generated. For each point, the hazard estimate is computed as

$$h(x) = \sum_{i=1}^n (1 - \delta_i) w_i K_s(x - x_{(i)})$$

where  $n = \text{NOBS}$ ,  $\delta_i$  is the  $i$ -th observation’s censoring code (0 = failed, 1 = censored),  $w_i$  is the  $i$ -th observation’s weight (either  $1/(n - i + 1)$  or  $\ln(1 + 1/(n - i + 1))$  depending upon `IWTO`), and  $K_s(x - x_{(i)})$ , the “smoothed kernel”, is as follows:

$$K_s(x - x_{(i)}) = \frac{1}{\alpha d_{ik}} K\left(\frac{x - x_{(i)}}{\beta d_{ik}}\right)$$

Here,  $d_{ik}$  is the distance to the  $k$ -th nearest failure from the  $i$ -th observation. Because of the  $d_{ik}$ , `HAZST` requires the computation of matrix  $H$ , which contains constants needed to quickly compute  $d_{ik}$ . Often,  $H$  will have been computed in routine `HAZRD` or `HAZEZ`. In this case, the parameter `IHCOMP` should be set to zero and  $H$  should be input to `HAZST`. If  $H$  must be computed by `HAZST`, set `IHCOMP` = 1.

## Example

The following example is a continuation of the example from HAZRD. The data are from Stablein, Carter, and Novak (1981), and involve the survival times of individuals with nonresectable gastric carcinoma. Only those individuals treated with both radiation and chemotherapy are used.

```
INTEGER      IHCOMP, IWTO, K, NGRID, NOBS
REAL         ALPHA, BETA, GINC, GSTRT
PARAMETER   (ALPHA=1.540537, BETA=1.631553, GINC=10, GSTRT=0.0,
&           IHCOMP=0, IWTO=0, K=20, NGRID=100, NOBS=45)
C
INTEGER      JCEN(NOBS), NOUT
REAL         H(5*NOBS), HAZ(NGRID), ST(NOBS)
EXTERNAL    HAZST, WRRRN
C
DATA ST/17, 42, 44, 48, 60, 72, 74, 95, 103, 108, 122, 144, 167,
&      170, 183, 185, 193, 195, 197, 208, 234, 235, 254, 307, 315,
&      401, 445, 464, 484, 528, 542, 567, 577, 580, 795, 855, 882,
&      892, 1031, 1033, 1306, 1335, 1366, 1452, 1472/
DATA JCEN/36*0, 9*1/
C
CALL HAZST (NOBS, ST, JCEN, IWTO, NGRID, GSTRT, GINC, ALPHA,
&          BETA, K, IHCOMP, H, HAZ)
C
CALL WRRRN ('Ten elements of HAZ', 1, 10, HAZ, 1, 0)
CALL WRRRN ('The first 10 rows of H', 10, 5, H, NOBS, 0)
END
```

## Output

```
                Ten elements of HAZ
      1          2          3          4          5          6          7
0.000962  0.001111  0.001276  0.001451  0.001634  0.001819  0.002004
      8          9         10
0.002185  0.002359  0.002523
```

```
                The first 10 rows of H
      1          2          3          4          5
1  217.0  218.0   1.0  21.0   1.0
2  192.0  193.0   1.0  21.0   0.5
3  190.0  191.0   1.0  21.0   0.3
4  186.0  187.0   1.0  21.0   0.2
5  174.0  175.0   1.0  21.0   0.2
6  162.0  163.0   1.0  21.0   0.2
7  160.0  161.0   1.0  21.0   0.1
8  139.0  140.0   1.0  21.0   0.1
9  131.0  132.0   1.0  21.0   0.1
10 126.0  127.0   1.0  21.0   0.1
```

# Chapter 16: Line Printer Graphics

---

## Routines

<b>16.1. Histograms</b>		
Vertical histogram plot .....	VHSTP	1074
Vertical histogram plot with bars subdivided into two parts..	VHS2P	1076
Horizontal histogram plot .....	HHSTP	1078
<b>16.2. Scatter Plots</b>		
Scatter plot.....	SCTP	1081
<b>16.3. Exploratory Data Analysis</b>		
Boxplot .....	BOXP	1083
Stem and leaf plot .....	STMLP	1085
<b>16.4. Empirical Probability Distribution</b>		
Cumulative distribution function (CDF) plot .....	CDFP	1087
Plot of two sample CDFs on the same frame .....	CDF2P	1090
Probability plot.....	PROBP	1092
<b>16.5. Other Graphics Routines</b>		
Plot up to 10 sets of points.....	PLOTP	1096
Binary tree plot.....	TREEP	1098

---

## Usage Notes

The routine names in this chapter end with the letter “P” to indicate line printer plotting and every routine starts printing at the beginning of a new page.

Depending on the nature of plots, some routines allow the user to change page width and/or length. This capability is specified in each routine and, if allowed, can be done by calling the routine PGOPT (page 1263) in advance. To change the page width, the user should make the following call to PGOPT:

```
CALL PGOPT(-1, IPAGEW)
```

where IPAGEW indicates the page width in columns. To change the page length, the user should make the following call to PGOPT:

CALL PGOPT(-2, IPAGEL)

where IPAGEL indicates the page length in rows. See the PGOPT document for more information.

---

## VHSTP/DVHSTP (Single/Double precision)

Print a vertical histogram.

### Usage

CALL VHSTP (NBAR, FRQ, ISP, TITLE)

### Arguments

**NBAR** — Number of bars. (Input)

If NBAR exceeds  $100/(ISP + 1)$ , then  $NBAR = 100/(ISP + 1)$  is used. NBAR must be positive.

**FRQ** — Vector of length NBAR containing the frequencies or counts. (Input)  
Elements of FRQ must be nonnegative.

**ISP** — Spacing between histogram bars. (Input)  
ISP may be 0, 1, or 4.

**TITLE** — CHARACTER string containing main title. (Input)

### Comments

1. Informational errors  

Type	Code	
3	1	ISP is out of range. $ISP = 0$ is used.
3	3	$NBAR * (ISP + 1)$ is less than 1 or greater than 100. The width of the histogram is set to 100, and $100/(ISP + 1)$ bars are printed. The number of class intervals will be printed completely if $ISP \neq 0$ and will always be printed up to and including $100/(ISP + 1)$ even though the histogram body is only 100 spaces wide.
3	5	The maximum value in the vector FRQ is less than 1; therefore, the body of the histogram is blank.
3	6	TITLE is too long. TITLE was truncated from the right side.
2. Output is written to the unit specified by the routine UMACH (page 1334).
3. TITLE is centered and placed at the top of the plot. The plot starts on a new page.



## Algorithm

VHSTP prints a vertical histogram on not more than one printer page using not more than 50 vertical and 100 horizontal print positions. Spacing control is allowed on the horizontal axis.

Given a vector containing positive counts, VHSTP determines the maximum count  $T_{\max}$ . Vertical printing position depends on  $K$  defined by

$K = 1 + (T_{\max} - 1)/50$ : If a frequency is greater than  $K$ , then a character is printed on the corresponding position of the first horizontal line from above. Henceforth,  $K$  is reduced by  $K/50$  for each horizontal line, and frequencies are compared to the new  $K$ .

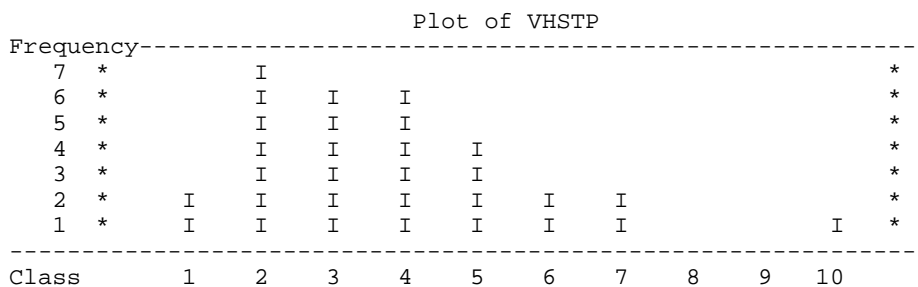
## Example

Consider the data set in Example 1 of the routine OWFRQ (page 3). This data set consists of the measurements (in inches) of precipitation in Minneapolis/St. Paul during the month of March for 30 consecutive years. We use the routine OWFRQ to create a one-way frequency table. A vertical histogram is then generated using VHSTP. A horizontal histogram for the same data set can be found in the document example for the routine HHSTP (page 1078).

```
INTEGER      NBAR, NOBS
PARAMETER   (NBAR=10, NOBS=30)
C
INTEGER      IBEG, IOPT, IREP, ISPACE, LENGTH, NOUT
REAL         CLHW, DIV(NBAR), TABLE(NBAR), X(NOBS), XHI, XLO
EXTERNAL     VHSTP, OWFRQ, UMACH
C
DATA X/0.77, 1.74, 0.81, 1.20, 1.95, 1.20, 0.47, 1.43, 3.37,
&          2.20, 3.00, 3.09, 1.51, 2.10, 0.52, 1.62, 1.31, 0.32, 0.59,
&          0.81, 2.81, 1.87, 1.18, 1.35, 4.75, 2.48, 0.96, 1.89, 0.90,
&          2.05/
C
CALL UMACH (2, NOUT)           Get output unit number
C
CALL OWFRQ (NOBS, X, NBAR, IOPT, XLO, XHI, CLHW, DIV, TABLE)
C
C                               Create a one-way frequency table from
C                               a given data set using intervals of
C                               equal length and user-supplied values
C                               of XLO and XHI
C
IOPT = 1
XLO  = 0.5
XHI  = 4.5
CALL OWFRQ (NOBS, X, NBAR, IOPT, XLO, XHI, CLHW, DIV, TABLE)
WRITE (NOUT,99999) DIV, TABLE
99999 FORMAT (' Midpoints: ', 10F6.2, '/', ' Counts: ', 10F6.0)
C
C                               Create the horizontal histogram
C
ISP = 4
CALL VHSTP (NBAR, TABLE, ISP, 'Plot of VHSTP')
END
```

## Output

```
Midpoints:  0.25  0.75  1.25  1.75  2.25  2.75  3.25  3.75  4.25  4.75
Counts:     2.    7.    6.    6.    4.    2.    2.    0.    0.    1.
```



## VHS2P/DVHS2P (Single/Double precision)

Print a vertical histogram with every bar subdivided into two parts.

### Usage

CALL VHS2P (NBAR, FRQX, FRQY, ISP, TITLE)

### Arguments

**NBAR** — Number of bars. (Input)

NBAR must be positive.

**FRQX** — Vector of length NBAR. (Input)

FRQX contains the frequencies or counts, and the elements of FRQX must be nonnegative.

**FRQY** — Vector of length NBAR. (Input)

FRQY contains the second frequencies or counts, and the elements of FRQY must be nonnegative.

**ISP** — Spacing between histogram bars. (Input)

ISP = 0, 1 or 4 is allowed.

**TITLE** — CHARACTER string containing the title. (Input)

### Comments

- Automatic workspace usage is

VHS2P 2 \* NBAR units, or

DVHS2P 4 \* NBAR units.

Workspace may be explicitly provided, if desired, by use of

V2S2P/DV2S2P. The reference is

CALL V2S2P (NBAR, FRQX, FRQY, ISP, TITLE, WK)

The additional argument is

**WK** — Work vector of length 2 \* NBAR.

- Informational errors

- | Type | Code |   |
|------|------|---|
| 3    | 2    | NBAR * (ISP + 1) is less than 1 or greater than 100. The width of the histogram is set to 100 and 100/(ISP + 1) bars are printed. |
| 3    | 3    | ISP as specified is not valid. The zero option is used.   |
| 3    | 4    | TITLE is too long. TITLE was truncated from the right side.   |
- If NBAR exceeds 100/(ISP + 1), then only 100/(ISP + 1) bars are printed.
  - If the maximum frequency is greater than 9999, the frequency column contains on some lines.
  - Output is written to the unit specified by the routine UMACH (page 1334).
  - TITLE is automatically centered and plot starts on a new page.

### Algorithm

The routine VHS2P prints a vertical histogram on one or more pages, using not more than 50 vertical and 100 horizontal print positions. Spacing control is allowed on the horizontal axis. Given two vectors containing positive counts, VHS2P determines the maximum count of the combined vectors  $T_{\max}$ . Vertical printing position depends on  $K$  defined by  $K = 1 + (T_{\max} - 1)/50$ . If a frequency is greater than  $K$ , then a character is printed on the first line. Henceforth,  $K$  is reduced by  $K/50$  for each position, and frequencies are compared to the new  $K$ .

### Example

Let  $X = \text{FRQX}$  contain 12 months of projected income figures and let  $Y = \text{FRQY}$  contain the actual income figures for the same 12 months. VHS2P produces a histogram that allows projected versus actual figures to be graphically compared.

```

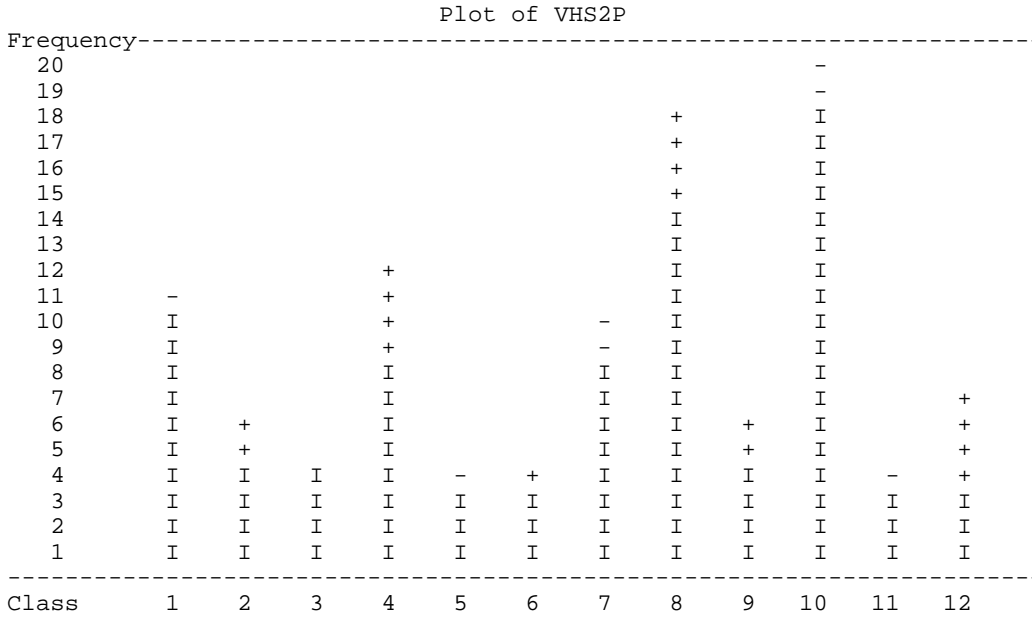
INTEGER      NBAR
PARAMETER   (NBAR=12)
C
INTEGER      ISP, NOUT
REAL         FRQX(NBAR), FRQY(NBAR)
EXTERNAL     UMACH, VHS2P
C
DATA FRQX/11., 4., 4., 8., 4., 3., 10., 14., 4., 20., 4., 3./
DATA FRQY/10., 6., 4., 12., 3., 4., 8., 18., 6., 18., 3., 7./
C
ISP = 4
CALL VHS2P (NBAR, FRQX, FRQY, ISP, 'Plot of VHS2P')
C                                     Get output unit number
CALL UMACH (2, NOUT)
WRITE (NOUT,99999)
99999 FORMAT (/, 3X, 'Twelve months projected sales versus actual ',
&          'sales, in thousands of dollars.', /, 11X, 'A positive ',
&          'sign (+) implies projected exceeded actual.', /, 11X,
&          'A negative sign (-) implies actual exceeded projected.')

```

C

END

### Output



Twelve months projected sales versus actual sales, in thousands of dollars.  
 A positive sign (+) implies projected exceeded actual.  
 A negative sign (-) implies actual exceeded projected.

## HHSTP/DHHSTP (Single/Double precision)

Print a horizontal histogram.

### Usage

```
CALL HHSTP (NBAR, FRQ, IBEG, ISPACE, LENGTH, IREP, IOPT,
           TITLE)
```

### Arguments

**NBAR** — Number of bars. (Input)

NBAR must be positive.

**FRQ** — Vector of length NBAR containing the frequencies or counts. (Input)

Elements of FRQ must be nonnegative.

**IBEG** — Indicates the beginning setting of the plot. (Input)

If IBEG = 0, HHSTP skips to a new page before printing the first line. If IBEG ≠ 0, HHSTP skips two spaces and begins printing on the same page.

**ISPACE** — Indicates spaces between horizontal histogram lines. (Input)  
ISPACE = 0, 1, or 2 is allowed.

**LENGTH** — Indicates the upper limit of the number of lines to print within the histogram per page. (Input)  
After that number of lines is printed, the routine skips to a new page to continue printing. If LENGTH = 0; then the maximum number of lines coincides with the standard printer page, which is 60.

**IREP** — Determines the repeating appearance for the class line (top) and frequency line (bottom) when multiple pages are required. (Input)  
If IREP = 0, the class line and the frequency line are printed on the first and last page of the histogram, respectively. If IREP ≠ 0, both class and frequency line are printed on every page.

**IOPT** — Page width option. (Input)  
IOPT = 0 will cause a full (horizontal) page histogram. IOPT = 1 will limit the width to 80 columns.

**TITLE** — CHARACTER string containing the title of the histogram. (Input)

### Comments

Informational errors

Type	Code	
3	3	ISPACE is not 0, 1, or 2. The zero option is used for ISPACE.
3	6	IOPT is not 0 or 1. The zero option is used for IOPT.
3	7	TITLE is too long and is truncated from the right side.

### Algorithm

The routine HHSTP prints a horizontal histogram on one or more pages. Given a vector containing frequencies or counts, HHSTP determines the maximum count  $T_{\max}$ . Horizontal printing position depends on  $K$  defined by

$$K = 1 + (T_{\max} - 1)/60 \quad \text{for 72 characters}$$

$$K = 1 + (T_{\max} - 1)/120 \quad \text{for 132 characters}$$

If a frequency is greater than  $K$ , then a character is printed in the first position. Henceforth,  $K$  is increased by  $K/60$  or  $K/120$  for each position, and frequencies are compared to the resulting  $K$ .

### Example

Consider the data set in Example 1 of the routine OWFRQ (page 3). We use the routine OWFRQ to create a one-way frequency table. A horizontal histogram is then generated using HHSTP. The user may find a vertical histogram for the same data set in the routine VHSTP (page 1074). Note that classes are listed from left to right in VHSTP.

INTEGER NBAR, NOBS

```

PARAMETER (NBAR=10, NOBS=30)
C
INTEGER  IBEG, IOPT, IREP, ISPACE, LENGTH, NOUT
REAL     CLHW, DIV(NBAR), TABLE(NBAR), X(NOBS), XHI, XLO
EXTERNAL HHSTP, OWFRQ, UMACH
C
DATA X/0.77, 1.74, 0.81, 1.20, 1.95, 1.20, 0.47, 1.43, 3.37,
&     2.20, 3.00, 3.09, 1.51, 2.10, 0.52, 1.62, 1.31, 0.32, 0.59,
&     0.81, 2.81, 1.87, 1.18, 1.35, 4.75, 2.48, 0.96, 1.89, 0.90,
&     2.05/
C
                                Get output unit number
CALL UMACH (2, NOUT)
C
                                Create a one-way frequency table from
C                                a given data set with intervals of
C                                equal length and user-supplied values
C                                of XLO and XHI
C
IOPT = 1
XLO  = 0.5
XHI  = 4.5
CALL OWFRQ (NOBS, X, NBAR, IOPT, XLO, XHI, CLHW, DIV, TABLE)
WRITE (NOUT,99999) DIV, TABLE
99999 FORMAT (' Midpoints: ', 10F6.2, /, ' Counts: ', 10F6.0)
C
                                Create the horizontal histogram
IBEG = 1
ISPACE = 1
LENGTH = 0
IREP = 0
IOPT = 0
CALL HHSTP (NBAR, TABLE, IBEG, ISPACE, LENGTH, IREP, IOPT,
&          'Histogram')
END

```

### Output

```

Midpoints:  .25  .75  1.25  1.75  2.25  2.75  3.25  3.75  4.25  4.75
Counts:     2.   7.   6.   6.   4.   2.   2.   0.   0.   1.

```

```

Histogram
Class -----
10 *I      *
   *      *
 9 *      *
   *      *
 8 *      *
   *      *
 7 *II     *
   *      *
 6 *II     *
   *      *
 5 *IIII   *
   *      *
 4 *IIIIII *
   *      *
 3 *IIIIII *
   *      *
 2 *IIIIII *
   *      *
 1 *II     *
-----

```

---

## SCTP/DSCTP (Single/Double precision)

Print a scatter plot of several groups of data.

### Usage

```
CALL SCTP (NOBS, NVAR, A, LDA, ICOL, RANGE, SYMBOL,
           XTITLE, YTITLE, TITLE)
```

### Arguments

**NOBS** — Number of observations. (Input)

**NVAR** — Number of variables. (Input)

**A** — NOBS by NVAR matrix containing the data. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement of the calling program. (Input)

**ICOL** — Vector of length NVAR representing the nature of each column of matrix A. (Input)

The *I*-th column of A is the independent variable vector if  $ICOL(I) = 1$ . The *I*-th column of A is a dependent variable vector if  $ICOL(I) = 2$ . The *I*-th column of A is ignored otherwise.

**RANGE** — Vector of length four specifying minimum *x*, maximum *x*, minimum *y* and maximum *y*. (Input)

SCTP will calculate the range of the axis if the minimum of that range is greater than or equal to the maximum of that range.

**SYMBOL** — CHARACTER string of length NVAR. (Input)

SYMBOL(*I* : *I*) is the character used to plot the data set represented by column *I*. SYMBOL(*I* : *I*) is ignored if  $ICOL(I) \neq 2$ .

**XTITLE** — CHARACTER string containing the *x*-axis title. (Input)

**YTITLE** — CHARACTER string containing the *y*-axis title. (Input)

**TITLE** — CHARACTER string containing the plot title. (Input)

### Comments

- Informational errors
 

Type	Code	
3	10	XTITLE is too long to fit into the page width determined by the routine PGOPT. XTITLE is truncated from the right side.



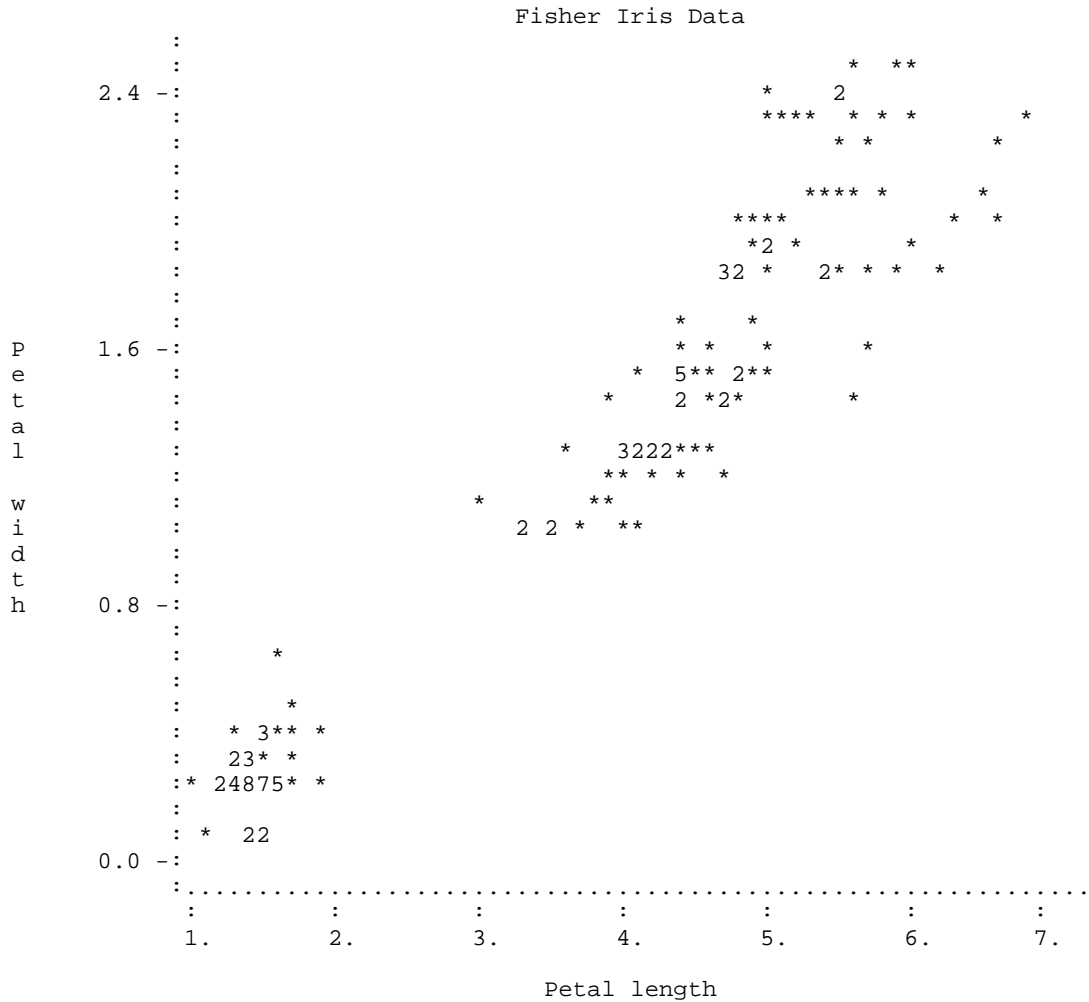


```

C                               Set page width and length
CALL PAGE (-1, 78)
CALL PAGE (-2, 40)
CALL SCTP (NOBS, NVAR, A, LDA, ICOL, RANGE, SYMBOL, 'Petal '//
&          'length', 'Petal width', 'Fisher Iris Data')
C
END

```

### Output



## BOXP/DBOXP (Single/Double precision)

Print boxplots for one or more samples.

### Usage

```
CALL BOXP (NGROUP, NI, X, TITLE)
```

## Arguments

**NGROUP** — The total number of groups of samples. (Input)

**NI** — Vector of length **NGROUP**. (Input)

**NI(I)** is the number of observations in the **I**-th group.

**X** — Vector of length **NI(1) + NI(2) + ... + NI(NGROUP)**. (Input)

The first **NI(1)** positions contain the observations for the first group. The next **NI(2)** positions contain the observations for the second group, and so on.

**TITLE** — CHARACTER string containing the title of the plot. (Input)

## Comments

1. Automatic workspace usage is

**BOXP**  $NI(1) + \dots + NI(NGROUP)$  units, or  
**DBOXP**  $2 * (NI(1) + \dots + NI(NGROUP))$  units.

Workspace may be explicitly provided, if desired, by use of **B2XP/DB2XP**. The reference is

```
CALL B2XP (NGROUP, NI, X, TITLE, WKSP)
```

The additional argument is

**WKSP** — Workspace of length **NI(1) + ... + NI(NGROUP)**. (Input)

The first **NI(1)** positions contain the sorted data from the first **NI(1)** positions of **X**. The next **NI(2)** positions contain sorted data from the next **NI(2)** positions of **X**, and so on.

2. Informational error

Type	Code	
3	5	<b>TITLE</b> is too long to fit into the page width determined by the routine <b>PGOPT</b> . <b>TITLE</b> is truncated from the right side.

3. **TITLE** is centered and placed at the top of the plot. The plot starts on a new page and the default page width is 78. The user may change the width by calling the routine **PGOPT** (page 1263) in advance.

## Algorithm

**BOXP** prints **NGROUP** boxplots. The minimum and maximum of **x** are printed. The median of each data group is marked by “\*” and the upper and lower hinges by “I”. The “H-spread” is the distance between the upper and lower hinges. The observation farthest from the median that still remains within one step (1.5 H-spread) from each hinge also is marked by “+”. The values in the second step (between 1.5 and 3 H-spreads from the hinges) are marked by the letter “o” and the values beyond the second step are marked by “x”. If there are fewer than five data points, each data point is plotted with an “x.” If multiple data points occur at positions marked “x” or “o”, the number of multiple points

is noted. More information on boxplots can be found in Chapter 2 of Chambers et al. (1983).

### Example

This example prints boxplots of three batches of data containing 5, 16 and 7 observations, respectively.

```

INTEGER      NGROUP
PARAMETER    (NGROUP=3)

C
INTEGER      NI(NGROUP)
REAL         X(28)
EXTERNAL     BOXP, PAGE

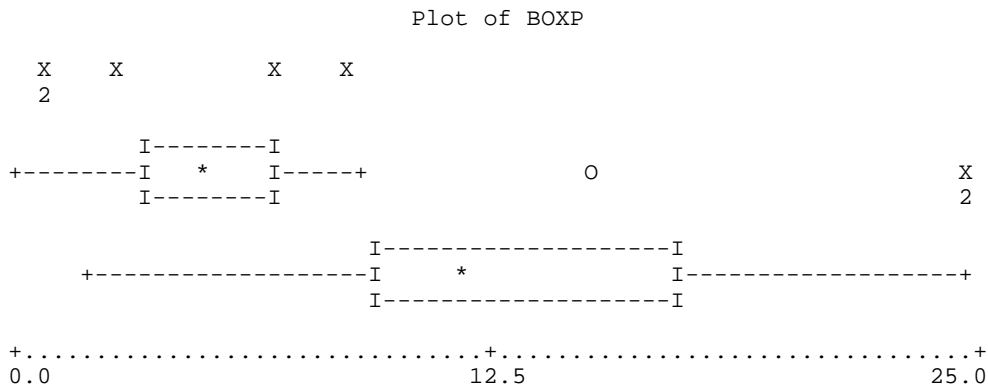
C
DATA (NI(I),I=1,3)/5, 16, 7/
DATA (X(I),I=1,5)/7., 9., 3., 1., 1./
DATA (X(I),I=6,21)/25., 0., 1., 0., 5., 4., 3., 5., 5., 5., 5.,
&    5., 5., 25., 15., 9./
DATA (X(I),I=22,28)/10., 15., 20., 25., 2., 9., 12./

C
                                Set page width.
CALL PAGE (-1, 70)
CALL BOXP (NGROUP, NI, X, 'Plot of BOXP')

C
END

```

### Output




---

## STMLP/DSTMLP (Single/Double precision)

Print a stem-and-leaf plot.

### Usage

```
CALL STMLP (NOBS, X, UNIT, TITLE)
```

### Arguments

**NOBS** — Number of observations. (Input)

**X**—Array of length **NOBS** containing the data. (Input)

**UNIT**— Size of the increment on the stem. (Input)

If **UNIT** is set so small that the length of the stem is more than 60 lines, **STMLP** will use a **UNIT** such that the stem will be no longer than 60 lines. However, if **UNIT** is a negative integer, **STMLP** will use the absolute value of **UNIT**, even if the stem would become very long. A common value for **UNIT** is 10.

**TITLE**— CHARACTER string containing the plot title. (Input)

### Comments

1. Automatic workspace usage is

**STMLP** **NOBS** + **MAXWID** units, or  
**DSTMLP** 2 \* **NOBS** + **MAXWID** units.

Workspace may be explicitly provided, if desired, by use of **S2MLP/DS2MLP**. The reference is

```
CALL S2MLP (NOBS, X, UNIT, TITLE, MAXWID, IWK, WK)
```

The additional arguments are as follows:

**MAXWID**— Page width. (Input)

**MAXWID** = 78 when **STMLP** is called.

**IWK**— Work vector of length **MAXWID**.

**WK**— Vector of length **NOBS**. (Output)

**WK** contains the sorted data from **X**.

2. Informational error

Type	Code	
3	4	<b>TITLE</b> is too long to fit into the page width determined by the routine <b>PGOPT</b> . <b>TITLE</b> is truncated from the right side.

3. Default page width is 78. The user may change it by calling the routine **PGOPT** (page 1263) in advance.

### Algorithm

Routine **STMLP** prints a stem-and-leaf display. The user can specify that the plot be longer than one page, but the default maximum is 60 lines. A plus sign (+) at the end of a line indicates that there are too many data points to fit within the width specifications. A scale marked in units of 10 is printed below the stem-and-leaf display.

### Example

This example prints a stem-and-leaf plot consisting of 27 data points ranging from -21.8 to 106.5.

```
INTEGER    NOBS
```

```

C      PARAMETER (NOBS=27)
C
C      REAL      UNIT, X(NOBS)
C      EXTERNAL  STMLP
C
C      DATA X/6.0, 106.5, 34.0, 88.1, 89.0, 0.3, 0.7, 4.0, 4.0, 5.0,
&      56.0, 62.8, 99.0, 4.0, 15.0, 76.0, 7.6, 101.5, 33.0, 91.0,
&      91.0, -6.3, -21.8, 0.0, 8.99, 5.5, 6.9/
C
C      UNIT = 10.
C      CALL STMLP (NOBS, X, UNIT, 'Stem and leaf plot')
C
C      END

```

### Output

```

                                Stem and leaf plot

Unit =      1.000000
For example: 1 2 represents      12.00000
-2 2
-1
-0 6
0 001444566789
1 5
2
3 34
4
5 6
6 3
7 6
8 89
9 119
10 27

```

---

## CDFP/DCDFP (Single/Double precision)

Print a sample cumulative distribution function (CDF), a theoretical CDF, and confidence band information.

### Usage

```
CALL CDFP (CDF, NOBS, X, N12, N95, IPRINT)
```

### Arguments

**CDF** — User-supplied FUNCTION to compute the cumulative distribution function. The form is CDF(P), where

P — Sample point. (Input)

CDF — Theoretical probability at the point P or integral of the probability density function at the point P. (Output)

CDF must be declared EXTERNAL in the calling program.

**NOBS** — Number of observations. (Input)

**X** —Vector of length **NOBS** containing the sample. (Input)

**N12** —Confidence band option. (Input)

If **N12** = 0, then no confidence bands are printed. If **N12** = 1, then positive or upper one-sided confidence band information is printed. If **N12** = -1, then negative or lower one-sided confidence band information is printed. If **N12** = 2, then two-sided confidence band information is printed.

**N95** —Confidence band option. (Input)

If **N95** = 95, the 95-percent band is desired. Otherwise, the 99-percent band is desired.

**IPRINT** — Print option. (Input)

If **IPRINT** = 1, then **CDFP** prints the sample CDF, the theoretical CDF, and the confidence band on the CDF. If **IPRINT** = 0, then the above information will not be printed.

### Comments

1. Automatic workspace usage is

**CDFP** 5 \* **NOBS** units, or  
**DCDFP** 10 \* **NOBS** units.

Workspace may be explicitly provided, if desired, by use of **C2FP/DC2FP**. The reference is

```
CALL C2FP (CDF, NOBS, X, N12, N95, IPRINT, WKX, WK)
```

The additional arguments are as follows:

**WKX** — Vector of length **NOBS** containing the sorted data **x** in ascending order. (Output)

**WK** — Vector of length 4 \* **NOBS** containing confidence band values. (Output)

**WK** may be dimensioned 3 \* **NOBS** instead of 4 \* **NOBS** for a lower or upper confidence band.

2. Note that sample CDFs are step functions.
3. Confidence bands are plotted around the sample CDF.
4. Output is written to the unit specified by the routine **UMACH** (page 1334).
5. Printing starts on a new page with default page width 78 columns and default page length 60 rows. The user may change these values by calling the routine **PGOPT** (page 1263) in advance.

### Algorithm

When **IPRINT** = 1, **CDFP** prints the sample cumulative distribution function (CDF), the theoretical CDF, and confidence bands on the CDF. The theoretical CDF will be plotted with or without the confidence band information. The sample CDF is calculated. The theoretical CDF is calculated by calling the user

supplied FUNCTION subprogram CDF. Asymptotic critical values are used (from the Smirnov tables) for confidence interval calculations.

### Example

This example prints and plots the sample CDF, the theoretical CDF, and the two-sided 95 percent band information using 70 observations. Routines RNSET (page 1166) and RNUN (page 1171) are called to generate these uniform (0, 1) random numbers.

```
INTEGER      ISEED, N12, N95, NOBS, IPRINT
PARAMETER   (NOBS=70)
REAL        CDF, X(NOBS)
EXTERNAL    CDF, CDFP, PGOPT, RNSET, RNUN

C
ISEED = 123457
IPRINT = 0

C                                Two-sided confidence band option.
N12 = 2

C                                95-percent band option.
N95 = 95

C                                Set page width and length.
CALL PGOPT (-1, 78)
CALL PGOPT (-2, 40)

C                                Initialize the seed.
CALL RNSET (ISEED)

C                                Generate pseudo-random numbers from
C                                a uniform (0,1) distribution.
CALL RNUN (NOBS, X)

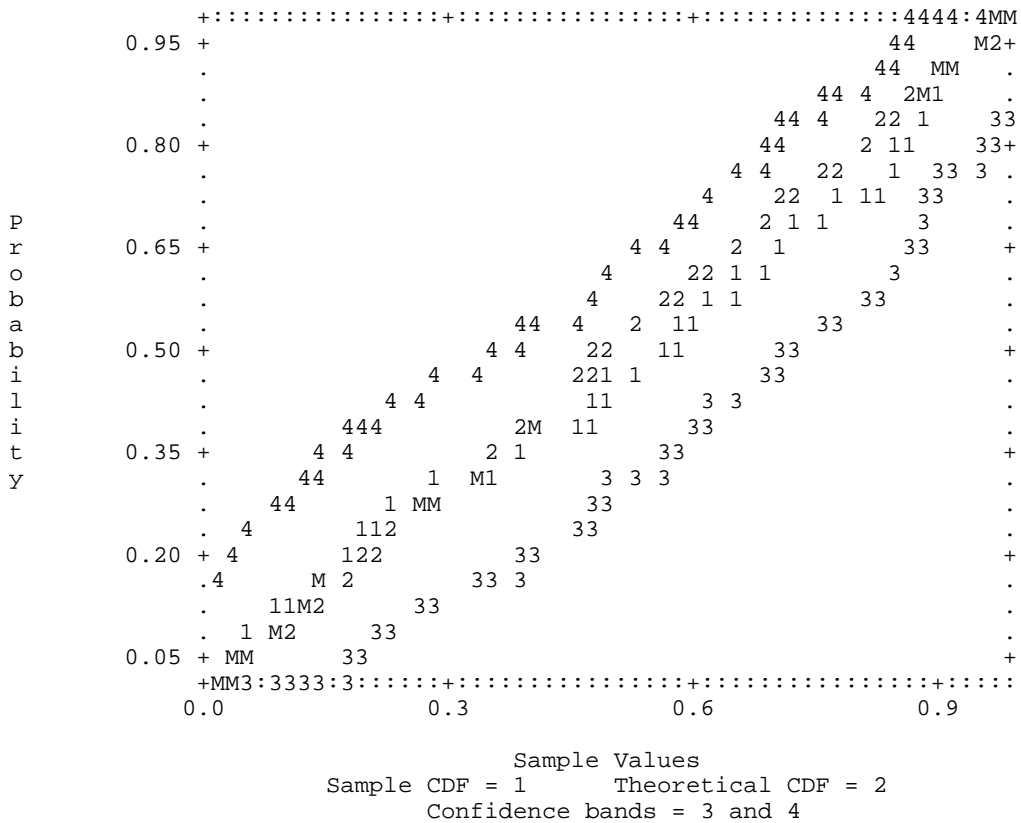
C                                Plot
CALL CDFP (CDF, NOBS, X, N12, N95, IPRINT)
END

C
REAL FUNCTION CDF (X)
REAL      X

C
CDF = X
RETURN
END
```

## Output

Cumulative Sample and Theoretical CDFs



---

## CDF2P/DCDF2P (Single/Double precision)

Print a plot of two sample cumulative distribution functions.

### Usage

CALL CDF2P (NOBS1, NOBS2, X)

### Arguments

**NOBS1** — Size of sample one. (Input)

**NOBS2** — Size of sample two. (Input)

**X** — Vector of length NOBS1 + NOBS2. (Input)  
X contains sample one followed by sample two.



## Comments

1. Automatic workspace usage is  
CDF2P 4 \* (NOBS1 + NOBS2) units, or  
DCDF2P 7 \* (NOBS1 + NOBS2) units.  
  
Workspace may be explicitly provided, if desired, by use of  
C2F2P/DC2F2P. The reference is  
CALL C2F2P (NOBS1, NOBS2, X, WK, IWK)  
  
The additional arguments are as follows:  
**WK** — Work vector of length 3 \* (NOBS1 + NOBS2).  
**IWK** — Work vector of length NOBS1 + NOBS2.
2. Printing starts on a new page with default page width 78 and default  
page length 60. The user may change page width and length by calling  
the routine PGOPT (page 1263) in advance.

## Algorithm

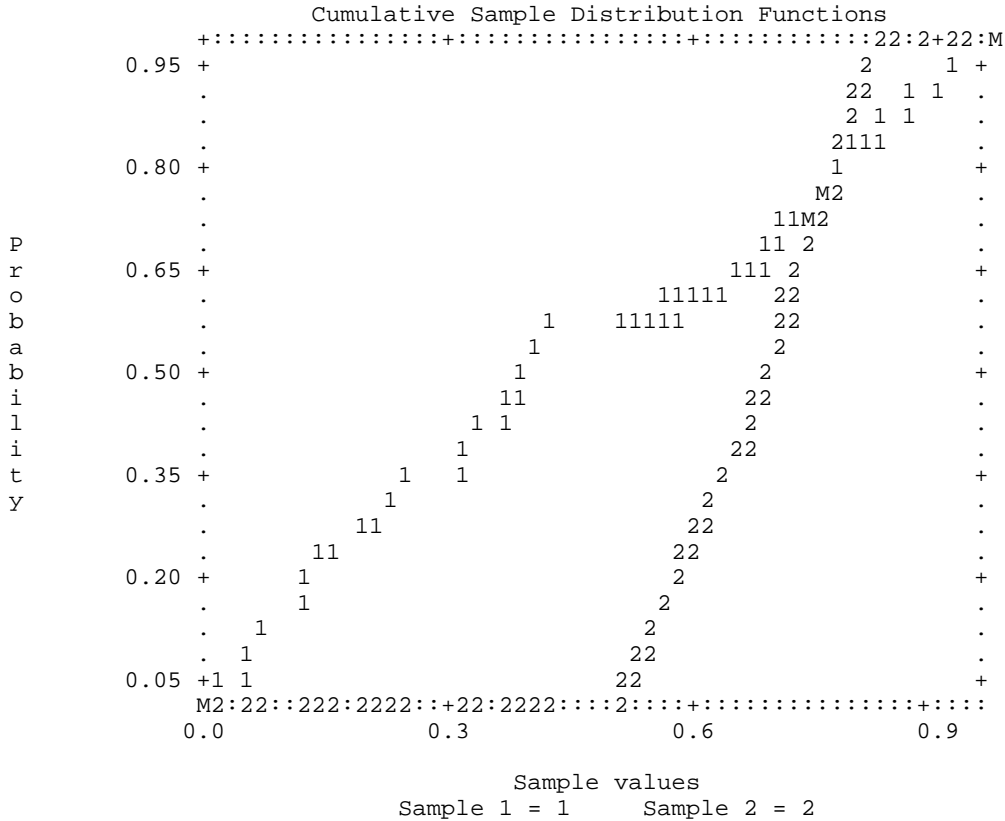
Routine CDF2P plots two sample cumulative probability distribution functions (CDFs). Two samples are first merged and then sorted. The cumulative distribution functions are then calculated. On the plots, the characters “1” and “2” indicate the first and second samples, respectively, and the character “M” indicates multiple points.

## Example

The first sample consists of pseudo-random numbers from a uniform (0, 1) distribution. Routines RNSET (page 1166) and RNUN (page 1171) are used to generate this sample. The second sample consists of points of the standard normal (Gaussian) distribution function generated by the routine ANORDF (page 1122).

```
INTEGER      I
REAL         ANORDF, VAL, X(100)
EXTERNAL     ANORDF, CDF2P, PGOPT, RNSET, RNUN
C           Initialize the seed.
CALL RNSET (1234567)
C           Generate pseudo-random numbers from
C           a uniform (0,1) distribution.
CALL RNUN (50, X)
C           Second sample consists of 50 points of
C           the std normal distribution function.
VAL = 0.
DO 10 I=1, 50
    VAL = VAL + .02
    X(I+50) = ANORDF(VAL)
10 CONTINUE
C           Set page width and length.
CALL PGOPT (-1, 78)
CALL PGOPT (-2, 40)
CALL CDF2P (50, 50, X)
END
```

## Output



---

## PROBP/DPROBP (Single/Double precision)

Print a probability plot.

### Usage

CALL PROBP (NOBS, N1, N2, X, IDIST)

### Arguments

**NOBS** — Total number of observations in uncensored sample. (Input)

**N1** — The rank number of the smallest observation in the sample x, if ranked in the complete sample. (Input)

In other words, the number of observations that have been censored from below is  $N1 - 1$ .

**N2** — The rank number of the largest observation in the sample x, if ranked in the complete sample. (Input)

In other words, the number of observations that have been censored from above is  $\text{NOBS} - \text{N2}$ .

**X** — {Vector of length  $\text{N2} - \text{N1} + 1$ . (Input)}

X contains the data, possibly a censored data set from a complete sample of size NOBS.

**IDIST** — Distribution option. (Input)

IDIST = 1, normal distribution.

IDIST = 2, lognormal distribution.

IDIST = 3, half-normal distribution.

IDIST = 4, exponential distribution.

IDIST = 5, Weibull distribution.

IDIST = 6, extreme value distribution.

### Comments

1. Automatic workspace usage is

PROBP 2 \* NOBS units, or  
DPROBP 4 \* NOBS units.

Workspace may be explicitly provided, if desired, by use of P2OBP/DP2OBP. The reference is

```
CALL P2OBP (NOBS, N1, N2, X, IDIST, M1, M2, WK)
```

The additional arguments are as follows:

**MI** — Rank of the smallest observation actually used. (Output)

**M2** — Rank of the largest observation actually used. (output)

**WK** — Work space of length 2 \* NOBS.

2. Informational error

Type	Code
------	------

3	7	It is necessary to delete some items from the plotting because those items do not satisfy properties of the distribution.
---	---	---

3. NOBS must be greater than or equal to  $\text{N2} - \text{N1} + 1$ . If there is no censoring, then  $\text{N1} = 1$  and  $\text{N2} = \text{NOBS}$ .
4. Output is written to the unit specified by the routine UMACH (page 1334).
5. Printing starts on a new page with default page width 78. The user may change it by calling the routine PGOPT (page 1263) in advance.

### Algorithm

Routine PROBP sorts a data set and plots the observed values along the vertical axis and the ranks along the horizontal axis. In the case of the lognormal and Weibull distributions, the vertical axis has a log scale. The horizontal axis has

the appropriate cumulative distribution function scale. Let  $M = \text{NOBS}$  denote the total number of observations in an uncensored sample. For normal and lognormal distributions, the horizontal plotting distance for the observation with rank  $I$  (out of  $M$ ) is proportional to the inverse normal cumulative distribution function evaluated at  $(3 * I - 1)/(3 * M + 1)$ . For the half-normal plot, the corresponding horizontal distance is proportional to the inverse normal cumulative distribution function evaluated at  $(3 * M + 3 * I - 1)/(6 * M + 1)$ . For other plots, the horizontal distances are proportional to the respective inverse cumulative distribution functions evaluated at  $(I - .5)/M$ .

Let  $N_1 = \text{N1}$  and  $N_2 = \text{N2}$ . In `PROBP` it is assumed that the  $N_1 - 1$  smallest observations and the  $M - N_2$  largest observations have been censored. If there has been no censoring,  $N_1$  should be set to 1 and  $N_2$  set to  $M$ . The smallest observation is plotted against the expected value (or the approximated expected value) of the  $N_1$ -th order statistic from a sample of size  $M$ ; the next smallest observation is plotted as if it were the  $(N_1 + 1)$ -th sample order statistic, and so on.

`PROBP` does not do any shifting of location of the observation in the data set. If any observations fall outside of the range of the distribution (that is, if any observations are nonpositive when the distribution specified is lognormal or Weibull), those observations are censored and  $N_1$  or  $N_2$  is modified to reflect the number censored. In this case an error message of type 3 is generated. A plot which is a straight line provides evidence that the sample is from the distribution specified.

### Example

In this example, a sample of size 250 (artificially generated from a normal distribution by routines `RNSET`, page 1166 and `RNNOR`, page 1208) is plotted by `PROBP` against a normal distribution function. The generally straight line produced is an indication that the sample is from a normal distribution.

```

C      INTEGER      NOBS
C      PARAMETER    (NOBS=250)

C      INTEGER      IDIST, N1, N2
C      REAL          X(NOBS)
C      EXTERNAL     PROBP, RNNOR, RNSET

C      IDIST = 1
C                                     No censoring
C      N1 = 1
C      N2 = 250

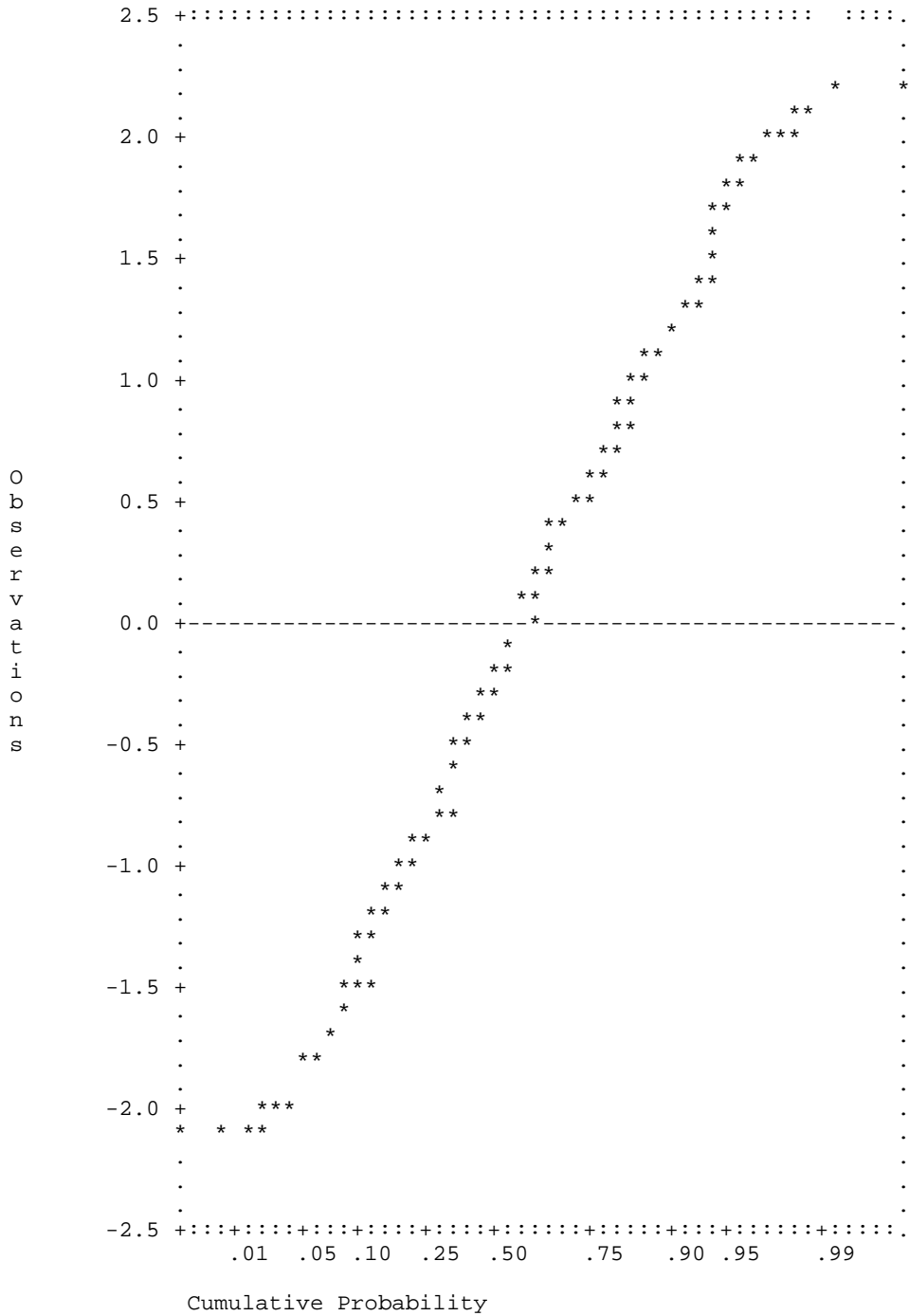
C                                     Initialize the seed
C      CALL RNSET (123457)
C      CALL RNNOR (N2, X)

C      CALL PROBP (NOBS, N1, N2, X, IDIST)
C      END

```

**Output**

Probability plot for normal distribution



---

## PLOTP/DPLOTP (Single/Double precision)

Print a plot of up to 10 sets of points.

### Usage

```
CALL PLOTP (NDATA, NFUN, X, A, LDA, INC, RANGE, SYMBOL,  
           XTITLE, YTITLE, TITLE)
```

### Arguments

**NDATA** — Number of independent variable data points. (Input)

**NFUN** — Number of sets of points. (Input)  
NFUN must be less than or equal to 10.

**X** — Vector of length NDATA containing the values of the independent variable.  
(Input)

**A** — Matrix of dimension NDATA by NFUN containing the NFUN sets of dependent variable values. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement of the calling program. (Input)

**INC** — Increment between elements of the data to be used. (Input)  
PLOTP plots  $X(1 + (I - 1) * INC)$  for  $I = 1, 2, \dots, NDATA$ .

**RANGE** — Vector of length four specifying minimum  $x$ , maximum  $x$ , minimum  $y$  and maximum  $y$ . (Input)  
PLOTP will calculate the range of the axis if the minimum and maximum of that range are equal.

**SYMBOL** — CHARACTER string of length NFUN. (Input)  
SYMBOL ( $I : I$ ) is the symbol used to plot function  $I$ .

**XTITLE** — CHARACTER string used to label the  $x$ -axis. (Input)

**YTITLE** — CHARACTER string used to label the  $y$ -axis. (Input)

**TITLE** — CHARACTER string used to label the plot. (Input)

### Comments

1. Informational errors

Type	Code	
3	7	NFUN is greater than 10. Only the first 10 functions are plotted.
3	8	TITLE is too long. TITLE is truncated from the right side.
3	9	YTITLE is too long. YTITLE is truncated from the right side.

3      10      XTITLE is too long. XTITLE is truncated from the right side. The maximum number of characters allowed depends on the page width and the page length. See Comment 5 below for more information.

2.      YTITLE and TITLE are automatically centered.
3.      For multiple plots, the character M is used if the same print position is shared by two or more data sets.
4.      Output is written to the unit specified by UMACH (page 1334).
5.      Default page width is 78 and default page length is 60. They may be changed by calling PGOPT (page 1263) in advance.

### Algorithm

Routine PLOTP produces a line printer plot of up to ten sets of points superimposed upon the same plot. A character "M" is printed to indicate multiple points. The user may specify the *x* and *y*-axis plot ranges and plotting symbols. Plot width and length may be reset in advance by calling PGOPT (page 1263).

### Example

This example plots the sine and cosine functions from  $-3.5$  to  $+3.5$  and sets page width and length to 78 and 40, respectively, by calling PGOPT (page 1263) in advance.

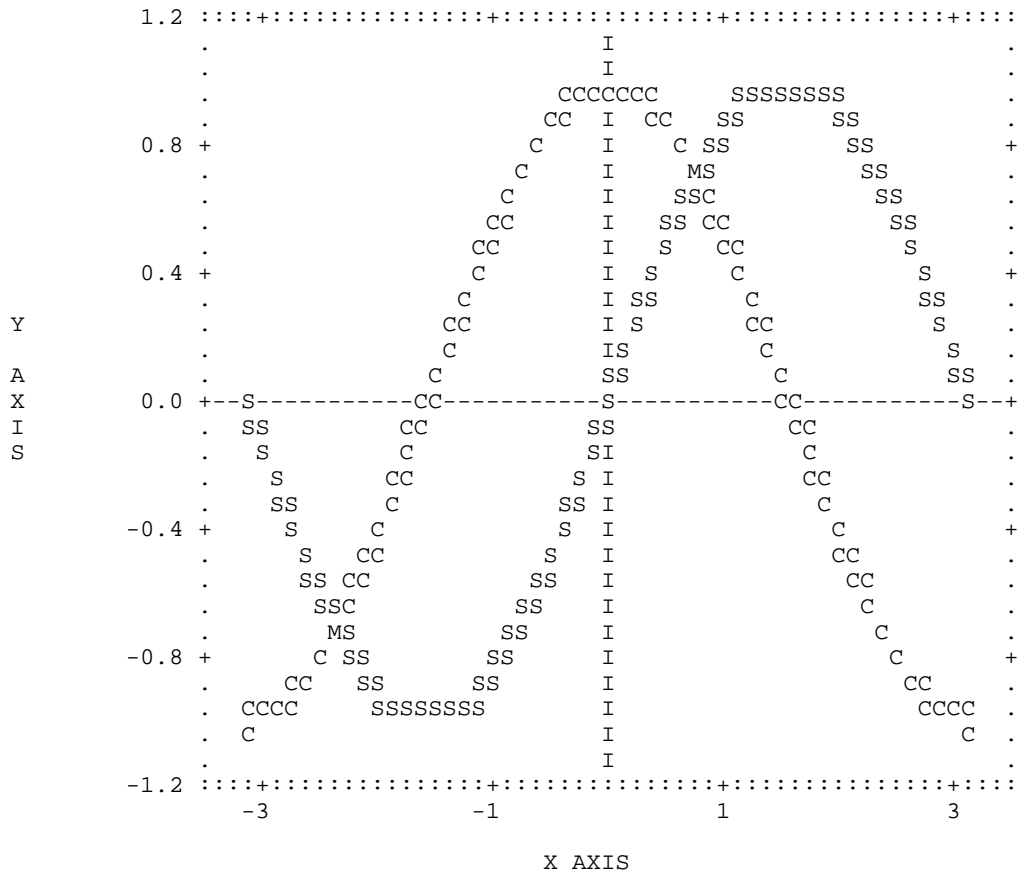
```

INTEGER      I, INC, LDA, NDATA, NFUN
REAL         A(200,2), DELX, PI, RANGE(4), X(200)
CHARACTER    SYMBOL*2
INTRINSIC    COS, SIN
EXTERNAL     CONST, PGOPT, PLOTP
C
DATA SYMBOL/'SC'/
DATA RANGE/-3.5, 3.5, -1.2, 1.2/
C
PI           = 3.14159
NDATA       = 200
NFUN        = 2
LDA         = 200
INC         = 1
DELX        = 2.*PI/199.
DO 10 I= 1, 200
    X(I)     = -PI + FLOAT(I-1) * DELX
    A(I,1)   = SIN(X(I))
    A(I,2)   = COS(X(I))
10 CONTINUE
C
                                Set page width and length
CALL PGOPT (-1, 78)
CALL PGOPT (-2, 40)
CALL PLOTP (NDATA, NFUN, X, A, LDA, INC, RANGE, SYMBOL,
&          'X AXIS', 'Y AXIS', ' C = COS,   S = SIN')
C
END

```

## Output

C = COS, S = SIN




---

## TREEP/DTREEP (Single/Double precision)

Print a binary tree.

### Usage

```
CALL TREEP (NODE, ICLSON, ICRSON, IMETH, CLEVEL, IROOT,
            NSCALE, NFILL, SCALE, NODENM)
```

### Arguments

**NODE** — Initial number of observations or nodes. (Input)  
 NODE must be greater than 2.

**ICLSON** — Vector of length  $NODE - 1$  containing the left son nodes. (Input)  
 Node number  $NODE + K$  has left son given by  $ICLSON(K)$  for  $K = 1, \dots,$   
 $NODE - 1$ .



**ICRSON** — Vector of length  $NODE - 1$  containing the right son nodes. (Input)  
Node number  $NODE + K$  has right son given by  $ICRSON(K)$  for  $K = 1, \dots, NODE - 1$ .

**IMETH** — Method to be used for printing the binary tree. (Input)

<b>IMETH</b>	<b>Method</b>
1	Horizontal tree
2	Horizontal <i>I</i> -tree
3	Vertical tree

**CLEVEL** — Vector of length  $NODE - 1$  containing the level used in merging or splitting the son nodes. (Input)  
 $CLEVEL(K)$  specifies the scale to be used on the vertical ( $IMETH = 1$  or  $2$ ) or horizontal ( $IMETH = 3$ ) axis for node  $NODE + K$ , for  $K = 1, 2, \dots, NODE - 1$ .

**IROOT** — Subtree specification. (Input)  
 $IROOT$  specifies the root node of the subtree to be printed. If  $IROOT = 2 * NODE - 1$  (or zero for the default), the entire tree is printed.  $IROOT$  must be in the range  $NODE + 1$  to  $2 * NODE - 1$ .

**NSCALE** — Number of horizontal slices of tree. (Input)  
 $NSCALE$  must be positive.

**NFILL** — The number of filler lines printed between horizontal or vertical node lines. (Input)  
 $NFILL = 1$  is usually sufficient.  $NFILL$  must be nonnegative.

**SCALE** — Vector of length two giving the interval on the  $CLEVEL$  axis which should be used to plot the tree. (Input)  
 $SCALE(1)$  is the location for printing the terminal nodes. The root node is printed at  $SCALE(2)$ .

**NODENM** —  $CHARACTER(*)$  vector of length  $NODE$  containing the terminal node labels. (Input)  
If terminal node labels are to be  $1, 2, 3, \dots$ , then  $NODENM(1)$  should be "DEFAULT" and the remaining elements of  $NODENM$  are not used. The length of each label is  $M$ , where  $M$  is determined by the user.

### Comments

1. Automatic workspace usage is

$TREEP$   $5 * IROOT + 2$  units, or  
 $DTREEP$   $6 * IROOT + 3$  units.

Workspace may be explicitly provided, if desired, by use of  $T2EEP/DT2EEP$ . The reference is

```
CALL T2EEP (NODE, ICLSON, ICRSON, IMETH, CLEVEL,  
           IROOT, NSCALE, NFILL, SCALE, NODENM,  
           IDTREE, ISTREE, IOTREE, INTREE, TLTREE)
```

The additional arguments are as follows:

**IDTREE** — Work vector of length IROOT. IDTREE is used to store the distance of each node from the vertical axis in vertical tree.

**ISTREE** — Work vector of length IROOT used to store all the nodes. IROOT is the first element of the array.

**IOTREE** — Work vector of length IROOT + 1 used to store the index of each node as TLTREE is sorted.

**INTREE** — Work vector of length IROOT.

**TLTREE** — Work vector of length IROOT + 1 used to store the level of each node in descending order in a vertical tree. It is used to store the distance of each node from the top of the horizontal line in ascending order in a horizontal tree.

2. Printing starts on a new page with default page width 78. The user may change it by calling the routine PGOPT (page 1263) in advance.

### Algorithm

Routine TREEP prints a binary tree which may represent results of hierarchical clustering algorithm such as the routine CLINK.

Let  $M = \text{NODE}$  indicate the number of nodes. A binary tree is composed of  $M$  terminal nodes and  $M - 1$  nonterminal nodes uniquely numbered 1 to  $M$  and  $M + 1$  to  $M + (M - 1)$ , respectively. Each nonterminal node joins together two son nodes which may or may not be terminal. Nonterminal nodes  $M + K$  are printed on the vertical scale interval  $[S_1, S_2]$  at the level given in  $C_K$ , for  $K = 1, 2, \dots, M - 1$ , where  $S_1 = \text{SCALE}(1)$ ,  $S_2 = \text{SCALE}(2)$ , and  $C_K = \text{CLEVEL}(K)$ .

### Example

```

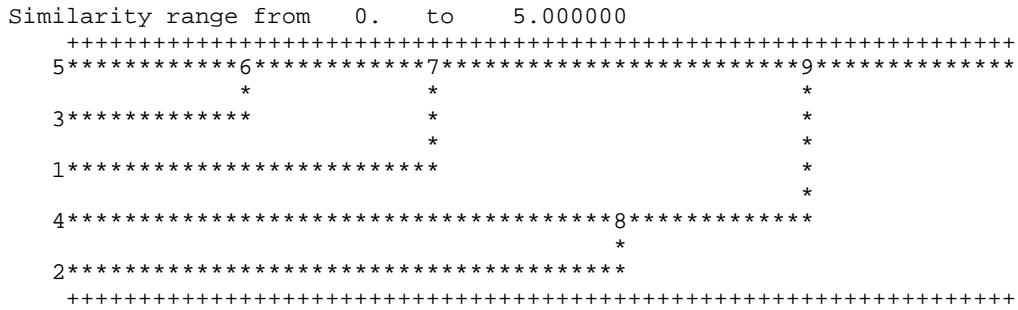
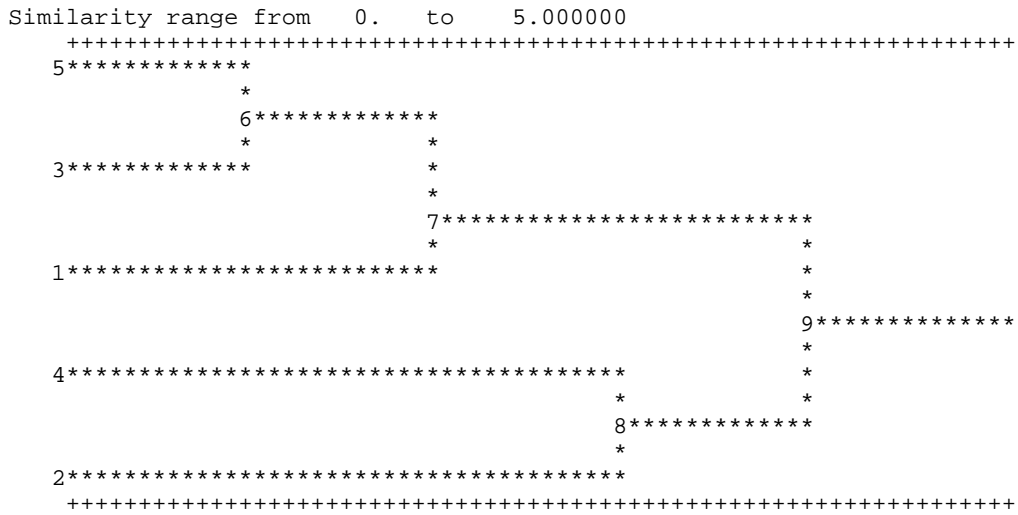
INTEGER      NODE
PARAMETER   (NODE=5)
C
INTEGER      ICLSON(NODE-1), ICRSON(NODE-1), IMETH, IROOT, NFILL,
&            NOUT, NSCALE
REAL         CLEVEL(NODE-1), SCALE(2)
CHARACTER    NODENM(NODE)*7
EXTERNAL     PGOPT, TREEP, UMACH
C
DATA ICLSON/5, 6, 4, 7/
DATA ICRSON/3, 1, 2, 8/
DATA NODENM/'DEFAULT', ' ', ' ', ' ', ' ', ' ', ' '/
DATA CLEVEL/1., 2., 3., 4./
DATA SCALE/0., 5./
C
                                Set page width
CALL PGOPT (-1, 70)
IROOT = 0
NSCALE = 1
NFILL = 1
C
                                Horizontal tree
IMETH = 1
CALL TREEP (NODE, ICLSON, ICRSON, IMETH, CLEVEL, IROOT, NSCALE,
```

```

&          NFFILL, SCALE, NODENM)
CALL UMACH (2, NOUT)
WRITE (NOUT,99999)
99999 FORMAT (1X, //)
C          Horizontal I-tree
      IMETH = 2
      CALL TREEP (NODE, ICLSON, ICRSON, IMETH, CLEVEL, IROOT, NSCALE,
&          NFFILL, SCALE, NODENM)
C
      END

```

**Output**



# Chapter 17: Probability Distribution Functions and Inverses

---

## Routines

<b>17.1. Discrete Random Variables: Distribution Functions and Probability Functions</b>		
Binomial distribution function .....	BINDF	1108
Binomial probability .....	BINPR	1110
Hypergeometric distribution function.....	HYPDF	1111
Hypergeometric probability .....	HYPPR	1113
Poisson distribution function .....	POIDF	1114
Poisson probability .....	POIPR	1115
<b>17.2. Continuous Random Variables: Distribution Functions and Their Inverses</b>		
Kolmogorov-Smirnov one-sided statistic distribution function .....	AKS1DF	1117
Kolmogorov-Smirnov two-sided statistic distribution function .....	AKS2DF	1120
Normal (Gaussian) distribution function .....	ANORDF	1122
Inverse of the normal distribution function .....	ANORIN	1124
Beta distribution function.....	BETDF	1125
Inverse of the beta distribution function .....	BETIN	1127
Bivariate normal distribution function .....	BNRDF	1128
Chi-squared distribution function .....	CHIDF	1129
Inverse of the chi-squared distribution function .....	CHIIN	1132
Noncentral chi-squared distribution function.....	CSNDF	1133
Inverse of the noncentral chi-squared distribution function .....	CSNIN	1136
<i>F</i> distribution function .....	FDf	1137
Inverse of the <i>F</i> distribution function .....	FIN	1139
Gamma distribution function .....	GAMDF	1140
Inverse of the gamma distribution function .....	GAMIN	1142
Student's <i>t</i> distribution function .....	TDF	1143
Inverse of the Student's <i>t</i> distribution function .....	TIN	1145
Noncentral Student's <i>t</i> distribution function.....	TNDF	1146

	Inverse of the noncentral Student's $t$ distribution function .....	TNIN	1149
<b>17.3.</b>	<b>General Continuous Random Variables</b>		
	Distribution function given ordinates of density .....	GCDF	1150
	Inverse of distribution function given ordinates of density .....	GCIN	1152
	Inverse of distribution function given subprogram .....	GFNIN	1155

## Usage Notes

### Comments

Definitions and discussions of the terms basic to this chapter can be found in Johnson and Kotz (1969, 1970a, 1970b). These are also good references for the specific distributions.

In order to keep the calling sequences simple, whenever possible, the subprograms described in this chapter are written for standard forms of statistical distributions. Hence, the number of parameters for any given distribution may be fewer than the number often associated with the distribution. For example, while a gamma distribution is often characterized by two parameters (or even a third, “location”), there is only one parameter that is necessary, the “shape”. The “scale” parameter can be used to scale the variable to the standard gamma distribution. Also, the functions relating to the normal distribution, ANORDF (page 1122) and ANORIN (page 1124), are for a normal distribution with mean equal to zero and variance equal to one. For other means and variances, it is very easy for the user to standardize the variables by subtracting the mean and dividing by the square root of the variance.

The *distribution function* for the (real, single-valued) random variable  $X$  is the function  $F$  defined for all real  $x$  by

$$F(x) = \text{Prob}(X \leq x)$$

where  $\text{Prob}(\cdot)$  denotes the probability of an event. The distribution function is often called the *cumulative distribution function* (CDF).

For distributions with finite ranges, such as the beta distribution, the CDF is 0 for values less than the left endpoint and 1 for values greater than the right endpoint. The subprograms described in this chapter return the correct values for the distribution functions when values outside of the range of the random variable are input, but warning error conditions are set in these cases.

### Discrete Random Variables

For discrete distributions, the function giving the probability that the random variable takes on specific values is called the *probability function*, defined by

$$p(x) = \text{Prob}(X = x)$$

The “PR” routines described in this chapter evaluate probability functions.

The CDF for a discrete random variable is

$$F(x) = \sum_A p(k)$$

where  $A$  is the set such that  $k \leq x$ . The “DF” routines in this chapter evaluate cumulative distribution functions. Since the distribution function is a step function, its inverse does not exist uniquely.

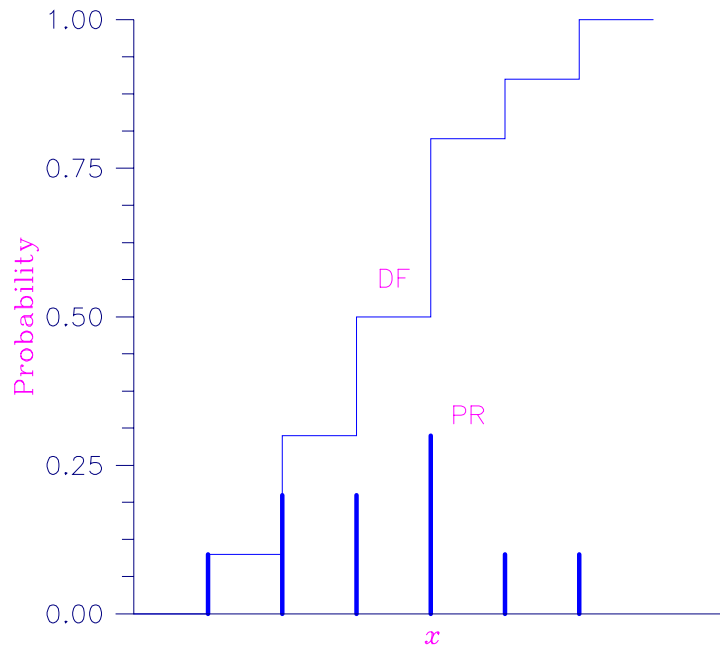


Figure 17-1 Discrete Random Variable

In the plot above, a routine like `BINPR` (page 1110) in this chapter evaluates the individual probability, given  $x$ . A routine like `BINDF` (page 1108) would evaluate the sum of the probabilities up to and including the probability at  $x$ .

## Continuous Distributions

For continuous distributions, a probability function, as defined above, would not be useful because the probability of any given point is 0. For such distributions, the useful analog is the *probability density function* (PDF). The integral of the PDF is the probability over the interval, if the continuous random variable  $X$  has PDF  $f$ , then

$$\text{Prob}(a < X \leq b) = \int_a^b f(x) dx$$

The relationship between the CDF and the PDF is

$$F(x) = \int_{-\infty}^x f(t)dt$$

as shown in Figure 17-2.

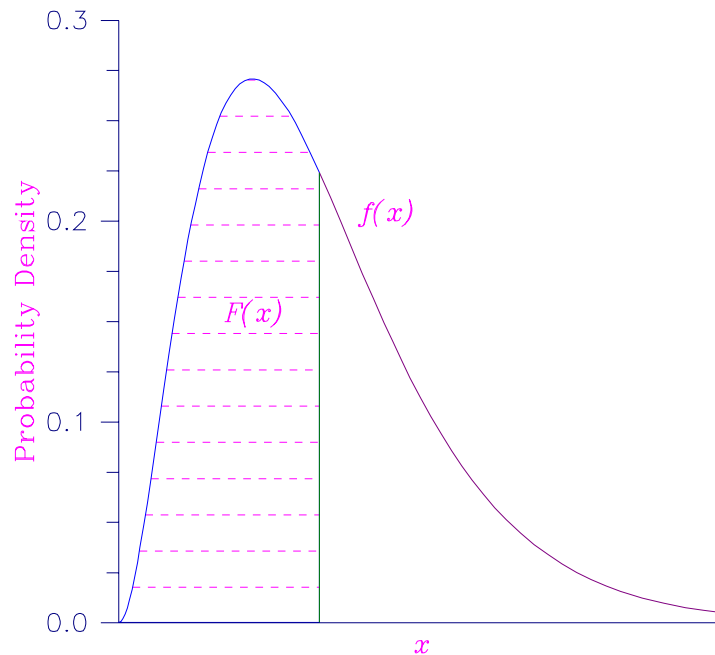


Figure 17-2 Probability Density Function

The “DF” routines described in this chapter evaluate cumulative distribution functions.

For (absolutely) continuous distributions, the value of  $F(x)$  uniquely determines  $x$  within the support of the distribution. The “IN” routines described in this chapter compute the inverses of the distribution functions, that is, given  $F(x)$  (called “P” for “probability”), a routine such as `BETIN` (page 1127) computes  $x$ . The inverses are defined only over the open interval  $(0,1)$ .

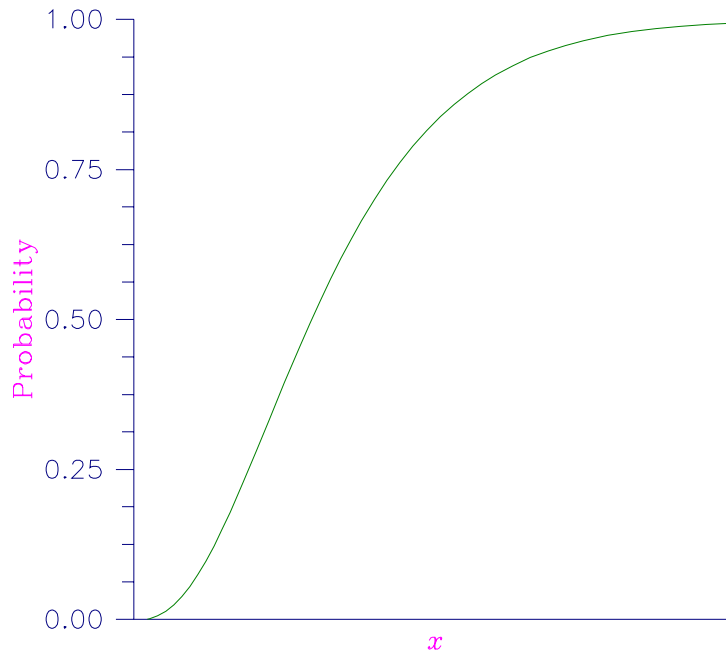


Figure 17-3 Cumulative Probability Distribution Function

There are three routines described in this chapter that deal with general continuous distribution functions. The routine `GCDF` (page 1150) computes a distribution function using values of the density function, and the routine `GCIN` (page 1152) computes the inverse. These two routines may be useful when the user has an estimate of a probability density, as perhaps computed by the routine `DESPL` (page 1040) or `DESKN` (page 1044), or computed from a frequency polygon. The routine `GFNIN` (page 1155) computes the inverse of a distribution function that is specified as a FORTRAN function.

### Additional Comments

Whenever a probability close to 1.0 results from a call to a distribution function or is to be input to an inverse function, it is often impossible to achieve good accuracy because of the nature of the representation of numeric values. In this case, it may be better to work with the complementary distribution function (one minus the distribution function). If the distribution is symmetric about some point (as the normal distribution, for example) or is reflective about some point (as the beta distribution, for example), the complementary distribution function has a simple relationship with the distribution function. For example, to evaluate the standard normal distribution at 4.0, using `ANORIN` (page 1124) directly, the result to six places is 0.999968. Only two of those digits are really useful, however. A more useful result may be 1.000000 minus this value, which can be



obtained to six significant figures as 3.16713E-05 by evaluating ANORIN at -4.0. For the normal distribution, the two values are related by  $\Phi(x) = 1 - \Phi(-x)$ , where  $\Phi(\cdot)$  is the normal distribution function. Another example is the beta distribution with parameters 2 and 10. This distribution is skewed to the right, so evaluating BETDF (page 1125) at 0.7, we obtain 0.999953. A more precise result is obtained by evaluating BETDF with parameters 10 and 2 at 0.3. This yields 4.72392E-5. (In both of these examples, it is wise not to trust the last digit.)

Many of the algorithms used by routines in this chapter are discussed by Abramowitz and Stegun (1964). The algorithms make use of various expansions and recursive relationships and often use different methods in different regions.

Cumulative distribution functions are defined for all real arguments, however, if the input to one of the distribution functions in this chapter is outside the range of the random variable, an error of Type 1 is issued, and the output is set to zero or one, as appropriate. A Type 1 error is of lowest severity, a “note”, and, by default, no printing or stopping of the program occurs. The other common errors that occur in the routines of this chapter are Type 2, “alert”, for a function value being set to zero due to underflow, Type 3, “warning”, for considerable loss of accuracy in the result returned, and Type 5, “terminal”, for incorrect and/or inconsistent input, complete loss of accuracy in the result returned, or inability to represent the result (because of overflow). When a Type 5 error occurs, the result is set to NaN (not a number, also used as a missing value code, obtained by routine AMACH(6), page 1334). (See the section “User Errors” in the Reference Material.)

---

## BINDF/DBINDF (Single/Double precision)

Evaluate the binomial distribution function.

### Usage

BINDF(*K*, *N*, *P*)

### Arguments

*K* — Argument for which the binomial distribution function is to be evaluated. (Input)

*N* — Number of Bernoulli trials. (Input)

*P* — Probability of success on each trial. (Input)

*BINDF* — Function value, the probability that a binomial random variable takes a value less than or equal to *K*. (Output)

BINDF is the probability that *K* or fewer successes occur in *N* independent Bernoulli trials, each of which has a *P* probability of success.

## Comments

Informational errors

Type	Code	
1	3	The input argument, $\kappa$ , is less than zero.
1	4	The input argument, $\kappa$ , is greater than the number of Bernoulli trials, $N$ .

## Algorithm

Function `BINDF` evaluates the distribution function of a binomial random variable with parameters  $n$  and  $p$ . It does this by summing probabilities of the random variable taking on the specific values in its range. These probabilities are computed by the recursive relationship

$$\Pr(X = j) = \frac{(n + 1 - j)p}{j(1 - p)} \Pr(X = j - 1)$$

To avoid the possibility of underflow, the probabilities are computed forward from 0, if  $k$  is not greater than  $n$  times  $p$ , and are computed backward from  $n$ , otherwise. The smallest positive machine number,  $\epsilon$ , is used as the starting value for summing the probabilities, which are rescaled by  $(1 - p)^n \epsilon$  if forward computation is performed and by  $p^n \epsilon$  if backward computation is done. For the special case of  $p = 0$ , `BINDF` is set to 1; and for the case  $p = 1$ , `BINDF` is set to 1 if  $k = n$  and to 0 otherwise.

## Example

Suppose  $X$  is a binomial random variable with  $n = 5$  and  $p = 0.95$ . In this example, we find the probability that  $X$  is less than or equal to 3.

```
INTEGER      K, N, NOUT
REAL         BINDF, P, PR
EXTERNAL     BINDF, UMACH

C
CALL UMACH (2, NOUT)
K = 3
N = 5
P = 0.95
PR = BINDF(K,N,P)
WRITE (NOUT,99999) PR
99999 FORMAT (' The probability that X is less than or equal to 3 is '
&           , F6.4)
END
```

## Output

The probability that  $X$  is less than or equal to 3 is 0.0226

---

## BINPR/DBINPR (Single/Double precision)

Evaluate the binomial probability function.

### Usage

`BINPR(K, N, P)`

### Arguments

**K** — Argument for which the binomial probability function is to be evaluated. (Input)

**N** — Number of Bernoulli trials. (Input)

**P** — Probability of success on each trial. (Input)

**BINPR** — Function value, the probability that a binomial random variable takes a value equal to **K**. (Output)

### Comments

Informational errors

Type	Code	
1	3	The input argument, <b>K</b> , is less than zero.
1	4	The input argument, <b>K</b> , is greater than the number of Bernoulli trials, <b>N</b> .

### Algorithm

The function `BINPR` evaluates the probability that a binomial random variable with parameters  $n$  and  $p$  takes on the value  $k$ . It does this by computing probabilities of the random variable taking on the values in its range less than (or the values greater than)  $k$ . These probabilities are computed by the recursive relationship

$$\Pr(X = j) = \frac{(n + 1 - j)p}{j(1 - p)} \Pr(X = j - 1)$$

To avoid the possibility of underflow, the probabilities are computed forward from 0, if  $k$  is not greater than  $n$  times  $p$ , and are computed backward from  $n$ , otherwise. The smallest positive machine number,  $\epsilon$ , is used as the starting value for computing the probabilities, which are rescaled by  $(1 - p)^n \epsilon$  if forward computation is performed and by  $p^n \epsilon$  if backward computation is done.

For the special case of  $p = 0$ , `BINPR` is set to 0 if  $k$  is greater than 0 and to 1 otherwise; and for the case  $p = 1$ , `BINPR` is set to 0 if  $k$  is less than  $n$  and to 1 otherwise.

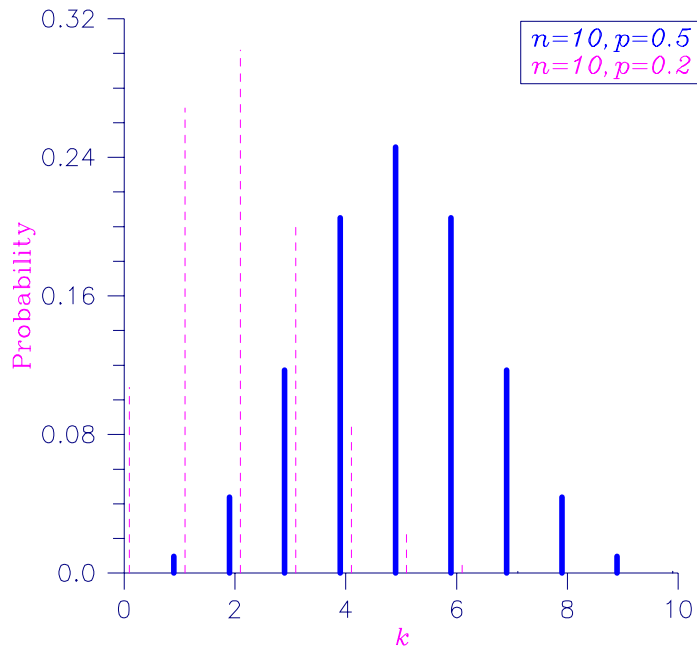


Figure 17-4 Binomial Probability Function

### Example

Suppose  $X$  is a binomial random variable with  $n = 5$  and  $p = 0.95$ . In this example, we find the probability that  $X$  is equal to 3.

```

INTEGER      K, N, NOUT
REAL         BINPR, P, PR
EXTERNAL     BINPR, UMACH

C
CALL UMACH (2, NOUT)
K = 3
N = 5
P = 0.95
PR = BINPR(K,N,P)
WRITE (NOUT,99999) PR
99999 FORMAT (' The probability that X is equal to 3 is ', F6.4)
END

```

### Output

The probability that X is equal to 3 is 0.0214

---

## HYPDF/DHYPDF (Single/Double precision)

Evaluate the hypergeometric distribution function.

## Usage

HYPDF(*K*, *N*, *M*, *L*)

## Arguments

***K*** — Argument for which the hypergeometric distribution function is to be evaluated. (Input)

***N*** — Sample size. (Input)

*N* must be greater than zero and greater than or equal to *K*.

***M*** — Number of defectives in the lot. (Input)

***L*** — Lot size. (Input)

*L* must be greater than or equal to *N* and *M*.

***HYPDF*** — Function value, the probability that a hypergeometric random variable takes a value less than or equal to *K*. (Output)

HYPDF is the probability that *K* or fewer defectives occur in a sample of size *N* drawn from a lot of size *L* that contains *M* defectives.

## Comments

Informational errors

Type	Code	
1	5	The input argument, <i>K</i> , is less than zero.
1	6	The input argument, <i>K</i> , is greater than the sample size.

## Algorithm

The function HYPDF evaluates the distribution function of a hypergeometric random variable with parameters *n*, *l*, and *m*. The hypergeometric random variable *X* can be thought of as the number of items of a given type in a random sample of size *n* that is drawn without replacement from a population of size *l* containing *m* items of this type. The probability function is

$$\Pr(X = j) = \frac{\binom{m}{j} \binom{l-m}{n-j}}{\binom{l}{n}} \quad \text{for } j = i, i+1, i+2, \dots, \min(n, m)$$

where  $i = \max(0, n - l + m)$ .

If *k* is greater than or equal to *i* and less than or equal to  $\min(n, m)$ , HYPDF sums the terms in this expression for *j* going from *i* up to *k*. Otherwise, HYPDF returns 0 or 1, as appropriate. So, as to avoid rounding in the accumulation, HYPDF performs the summation differently depending on whether or not *k* is greater than the mode of the distribution, which is the greatest integer less than or equal to  $(m+1)(n+1)/(l+2)$ .

### Example

Suppose  $X$  is a hypergeometric random variable with  $n = 100$ ,  $l = 1000$ , and  $m = 70$ . In this example, we evaluate the distribution function at 7.

```
INTEGER      K, L, M, N, NOUT
REAL         DF, HYPDF
EXTERNAL     HYPDF, UMACH
C
CALL UMACH (2, NOUT)
K = 7
N = 100
L = 1000
M = 70
DF = HYPDF(K,N,M,L)
WRITE (NOUT,99999) DF
99999 FORMAT (' The probability that X is less than or equal to 7 is '
&           , F6.4)
END
```

### Output

The probability that X is less than or equal to 7 is 0.5995

---

## HYPDR/DHYPDR (Single/Double precision)

Evaluate the hypergeometric probability function.

### Usage

HYPDR(K, N, M, L)

### Arguments

**K** — Argument for which the hypergeometric probability function is to be evaluated. (Input)

**N** — Sample size. (Input)  
N must be greater than zero and greater than or equal to K.

**M** — Number of defectives in the lot. (Input)

**L** — Lot size. (Input)  
L must be greater than or equal to N and M.

**HYPDR** — Function value, the probability that a hypergeometric random variable takes a value equal to K. (Output)

HYPDR is the probability that exactly K defectives occur in a sample of size N drawn from a lot of size L that contains M defectives.

### Comments

Informational errors

Type	Code	
1	5	The input argument, K, is less than zero.

1            6            The input argument,  $k$ , is greater than the sample size.

### Algorithm

The function `HYPBR` evaluates the probability function of a hypergeometric random variable with parameters  $n$ ,  $l$ , and  $m$ . The hypergeometric random variable  $X$  can be thought of as the number of items of a given type in a random sample of size  $n$  that is drawn without replacement from a population of size  $l$  containing  $m$  items of this type. The probability function is

$$\Pr(X = k) = \frac{\binom{m}{k} \binom{l-m}{n-k}}{\binom{l}{n}} \quad \text{for } k = i, i+1, i+2, \dots, \min(n, m)$$

where  $i = \max(0, n - l + m)$ . `HYPBR` evaluates the expression using log gamma functions.

### Example

Suppose  $X$  is a hypergeometric random variable with  $n = 100$ ,  $l = 1000$ , and  $m = 70$ . In this example, we evaluate the probability function at 7.

```
INTEGER      K, L, M, N, NOUT
REAL         HYPBR, PR
EXTERNAL    HYPBR, UMACH

C
CALL UMACH (2, NOUT)
K = 7
N = 100
L = 1000
M = 70
PR = HYPBR(K,N,M,L)
WRITE (NOUT,99999) PR
99999 FORMAT (' The probability that X is equal to 7 is ', F6.4)
END
```

### Output

The probability that X is equal to 7 is 0.1628

---

## POIDF/DPOIDF (Single/Double precision)

Evaluate the Poisson distribution function.

### Usage

```
POIDF(K, THETA)
```

### Arguments

**K** — Argument for which the Poisson distribution function is to be evaluated.  
(Input)

**THETA** — Mean of the Poisson distribution. (Input)  
THETA must be positive.

**POIDF** — Function value, the probability that a Poisson random variable takes a value less than or equal to  $\kappa$ . (Output)

### Comments

Informational error

Type	Code	
1	1	The input argument, $\kappa$ , is less than zero.

### Algorithm

The function POIDF evaluates the distribution function of a Poisson random variable with parameter THETA. THETA, which is the mean of the Poisson random variable, must be positive. The probability function (with  $\theta = \text{THETA}$ ) is

$$f(x) = e^{-\theta} \theta^x / x!, \quad \text{for } x = 0, 1, 2, \dots$$

The individual terms are calculated from the tails of the distribution to the mode of the distribution and summed. POIDF uses the recursive relationship

$$f(x+1) = f(x)\theta/(x+1), \quad \text{for } x = 0, 1, 2, \dots, k-1,$$

with  $f(0) = e^{-\theta}$ .

### Example

Suppose  $X$  is a Poisson random variable with  $\theta = 10$ . In this example, we evaluate the distribution function at 7.

```
INTEGER      K, NOUT
REAL         DF, POIDF, THETA
EXTERNAL     POIDF, UMACH

C
CALL UMACH (2, NOUT)
K          = 7
THETA     = 10.0
DF        = POIDF(K, THETA)
WRITE (NOUT, 99999) DF
99999 FORMAT (' The probability that X is less than or equal to ',
&           '7 is ', F6.4)
END
```

### Output

The probability that  $X$  is less than or equal to 7 is 0.2202

---

## POIPR/DPOIPR (Single/Double precision)

Evaluate the Poisson probability function.



## Usage

`POIPR(K, THETA)`

## Arguments

***K*** — Argument for which the Poisson distribution function is to be evaluated. (Input)

***THETA*** — Mean of the Poisson distribution. (Input)  
*THETA* must be positive.

***POIPR*** — Function value, the probability that a Poisson random variable takes a value equal to *K*. (Output)

## Comments

Informational error

Type	Code
------	------

1	1	The input argument, <i>K</i> , is less than zero.
---	---	---

## Algorithm

The function `POIPR` evaluates the probability function of a Poisson random variable with parameter `THETA`. `THETA`, which is the mean of the Poisson random variable, must be positive. The probability function (with  $\theta = \text{THETA}$ ) is

$$f(x) = e^{-\theta} \theta^k / k!, \quad \text{for } k = 0, 1, 2, \dots$$

`POIPR` evaluates this function directly, taking logarithms and using the log gamma function.

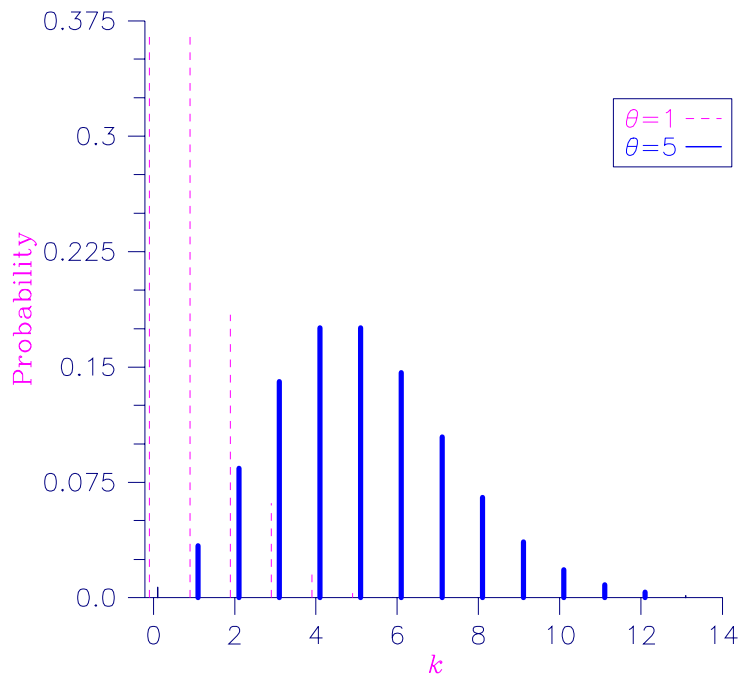


Figure 17-5 Poisson Probability Function

### Example

Suppose  $X$  is a Poisson random variable with  $\theta = 10$ . In this example, we evaluate the probability function at 7.

```

INTEGER      K, NOUT
REAL         POIPR, PR, THETA
EXTERNAL     POIPR, UMACH

C
CALL UMACH (2, NOUT)
K          = 7
THETA     = 10.0
PR        = POIPR(K, THETA)
WRITE (NOUT, 99999) PR
99999 FORMAT (' The probability that X is equal to 7 is ', F6.4)
END

```

### Output

The probability that  $X$  is equal to 7 is 0.0901

---

## AKS1DF/DKS1DF (Single/Double precision)

Evaluate the distribution function of the one-sided Kolmogorov-Smirnov goodness of fit  $D^+$  or  $D^-$  test statistic based on continuous data for one sample.

## Usage

AKS1DF(NOBS, D)

## Arguments

**NOBS** — The total number of observations in the sample. (Input)

**D** — The  $D^+$  or  $D^-$  test statistic. (Input)

D is the maximum positive difference of the empirical cumulative distribution function (CDF) minus the hypothetical CDF or the maximum positive difference of the hypothetical CDF minus the empirical CDF.

**AKSIDF** — The probability of a smaller D. (Output)

## Comments

1. Automatic workspace usage is

AKS1DF 3 \* (NOBS + 1) units, or

DKS1DF 6 \* (NOBS + 1) units.

Workspace may be explicitly provided, if desired, by use of AK21DF/DK21DF. The reference is

AK2DF(NOBS, D, WK)

The additional argument is

**WK** — Work vector of length 3 \* NOBS + 3 if NOBS ≤ 80. WK is not used if NOBS is greater than 80.

2. Informational errors

Type	Code	
------	------	--

1	2	Since the D test statistic is less than zero, the distribution function is zero at D.
---	---	---

1	3	Since the D test statistic is greater than one, the distribution function is one at D.
---	---	--

3. If NOBS ≤ 80, then exact one-sided probabilities are computed. In this case, on the order of NOBS<sup>2</sup> operations are required. For NOBS > 80, approximate one-sided probabilities are computed. These approximate probabilities require very few computations.
4. An approximate two-sided probability for the  $D = \max(D^+, D^-)$  statistic can be computed as twice the AKS1DF probability for D(minus one, if the probability from AKS1DF is greater than 0.5).

## Algorithm

Routine AKS1DF computes the cumulative distribution function (CDF) for the one-sided Kolmogorov-Smirnov one-sample  $D^+$  or  $D^-$  statistic when the theoretical CDF is strictly continuous. Let  $F(x)$  denote the theoretical

distribution function, and let  $S_n(x)$  denote the empirical distribution function obtained from a sample of size NOBS. Then, the  $D^+$  statistic is computed as

$$D^+ = \sup_x [F(x) - S_n(x)]$$

while the one-sided  $D^-$  statistic is computed as

$$D^- = \sup_x [S_n(x) - F(x)]$$

Exact probabilities are computed according to a method given by Conover (1980, page 350) for sample sizes of 80 or less. For sample sizes greater than 80, Smirnov's asymptotic result is used, that is, the value of the CDF is taken as  $1 - \exp(-2nd^2)$ , where  $d$  is  $D^+$  or  $D^-$  (Kendall and Stuart, 1979, page 482). This asymptotic expression is conservative (the value returned by AKS1DF is smaller than the exact value, when the sample size exceeds 80).

### Programming Notes

Routine AKS1DF requires on the order of  $\text{NOBS}^2$  operations to compute the exact probabilities, where an operation consists of taking ten or so logarithms. Because so much computation is occurring within each "operation," AKS1DF is much slower than its two-sample counterpart, function AKS2DF (page 1120).

### Example

In this example, the exact one-sided probabilities for the tabled values of  $D^+$  or  $D^-$ , given, for example, in Conover (1980, page 462), are computed. Tabled values at the 10% level of significance are used as input to AKS1DF for sample sizes of 5 to 50 in increments of 5 (the last two tabled values are obtained using the asymptotic critical values of

$$1.07 / \sqrt{\text{NOBS}}$$

The resulting probabilities should all be close to 0.90.

```

INTEGER      I, NOBS, NOUT
REAL         AKS1DF, D(10)
EXTERNAL     AKS1DF, UMACH

C
DATA D/0.447, 0.323, 0.266, 0.232, 0.208, 0.190, 0.177, 0.165,
&      0.160, 0.151/

C
CALL UMACH (2, NOUT)

C
DO 10 I=1, 10
   NOBS = 5*I

C
   WRITE (NOUT,99999) D(I), NOBS, AKS1DF(NOBS,D(I))

C
99999  FORMAT (' One-sided Probability for D = ', F8.3, ' with NOBS '
```

```

&
10 CONTINUE
END
, '= ', I2, ' is ', F8.4)

```

### Output

```

One-sided Probability for D = 0.447 with NOBS = 5 is 0.9000
One-sided Probability for D = 0.323 with NOBS = 10 is 0.9006
One-sided Probability for D = 0.266 with NOBS = 15 is 0.9002
One-sided Probability for D = 0.232 with NOBS = 20 is 0.9009
One-sided Probability for D = 0.208 with NOBS = 25 is 0.9002
One-sided Probability for D = 0.190 with NOBS = 30 is 0.8992
One-sided Probability for D = 0.177 with NOBS = 35 is 0.9011
One-sided Probability for D = 0.165 with NOBS = 40 is 0.8987
One-sided Probability for D = 0.160 with NOBS = 45 is 0.9105
One-sided Probability for D = 0.151 with NOBS = 50 is 0.9077

```

---

## AKS2DF/DKS2DF (Single/Double precision)

Evaluate the distribution function of the Kolmogorov-Smirnov goodness of fit  $D$  test statistic based on continuous data for two samples.

### Usage

AKS2DF(NOBSX, NOBSY, D)

### Arguments

**NOBSX** — The total number of observations in the first sample. (Input)

**NOBSY** — The total number of observations in the second sample. (Input)

**D** — The  $D$  test statistic. (Input)

$D$  is the maximum absolute difference between empirical cumulative distribution functions (CDFs) of the two samples.

**AKS2DF** — The probability of a smaller  $D$ . (Output)

### Comments

- Automatic workspace usage is

AKS2DF  $\max(\text{NOBSX}, \text{NOBSY}) + 1$  units, or  
 DKS2DF  $2 * \max(\text{NOBSX}, \text{NOBSY}) + 1$  units.

Workspace may be explicitly provided, if desired, by use of  
 AK22DF/DK22DF. The reference is

AK22DF(NOBSX, NOBSY, D, WK)

The additional argument is

**WK** — Work vector of length  $\max(\text{NOBSX}, \text{NOBSY}) + 1$ .

- Informational errors  
 Type Code

- |   |   |  |
|---|---|--|
| 1 | 2 | Since the $D$ test statistic is less than zero, then the distribution function is zero at $D$ .  |
| 1 | 3 | Since the $D$ test statistic is greater than one, then the distribution function is one at $D$ . |

### Algorithm

Function `AKS2DF` computes the cumulative distribution function (CDF) for the two-sided Kolmogorov-Smirnov two-sample  $D$  statistic when the theoretical CDF is strictly continuous. Exact probabilities are computed according to a method given by Kim and Jennrich (1973). Approximate asymptotic probabilities are computed according to methods also given in this reference.

Let  $F_n(x)$  and  $G_m(x)$  denote the empirical distribution functions for the two samples, based on  $n = \text{NOBSX}$  and  $m = \text{NOBSY}$  observations. Then, the  $D$  statistic is computed as

$$D = \sup_x |F_n(x) - G_m(x)|$$

### Programming Notes

Function `AKS2DF` requires on the order of  $\text{NOBSX} * \text{NOBSY}$  operations to compute the exact probabilities, where an operation consists of an addition and a multiplication. For  $\text{NOBSX} * \text{NOBSY}$  less than 10000, the exact probability is computed. If this is not the case, then the Smirnov approximation discussed by Kim and Jennrich (1973) is used if the minimum of  $\text{NOBSX}$  and  $\text{NOBSY}$  is greater than ten percent of the maximum of  $\text{NOBSX}$  and  $\text{NOBSY}$ , or if the minimum is greater than 80. Otherwise, the Kolmogorov approximation discussed by Kim and Jennrich (1973) is used.

### Example

Function `AKS2DF` is used to compute the probability of a smaller  $D$  statistic for a variety of sample sizes using values close to the 0.95 probability value.

```

INTEGER      I, NOBSX(10), NOBSY(10), NOUT
REAL         AKS2DF, D(10)
EXTERNAL     AKS2DF, UMACH
C
DATA NOBSX/5, 20, 40, 70, 110, 200, 200, 200, 100, 100/
DATA NOBSY/10, 10, 10, 10, 10, 20, 40, 60, 80, 100/
DATA D/0.7, 0.55, 0.475, 0.4429, 0.4029, 0.2861, 0.2113, 0.1796,
&      0.18, 0.18/
C
CALL UMACH (2, NOUT)
C
DO 10 I=1, 10
C
      WRITE (NOUT,99999) D(I), NOBSX(I), NOBSY(I),
&      AKS2DF(NOBSX(I),NOBSY(I),D(I))
C
99999      FORMAT (' Probability for D = ', F5.3, ' with NOBSX = ', I3,
```

```

& ' and NOBSY = ', I3, ' is ', F9.6, '.'')
10 CONTINUE
END

```

### Output

```

Probability for D = 0.700 with NOBSX = 5 and NOBSY = 10 is 0.980686.
Probability for D = 0.550 with NOBSX = 20 and NOBSY = 10 is 0.987553.
Probability for D = 0.475 with NOBSX = 40 and NOBSY = 10 is 0.972423.
Probability for D = 0.443 with NOBSX = 70 and NOBSY = 10 is 0.961646.
Probability for D = 0.403 with NOBSX = 110 and NOBSY = 10 is 0.928667.
Probability for D = 0.286 with NOBSX = 200 and NOBSY = 20 is 0.921126.
Probability for D = 0.211 with NOBSX = 200 and NOBSY = 40 is 0.917110.
Probability for D = 0.180 with NOBSX = 200 and NOBSY = 60 is 0.914520.
Probability for D = 0.180 with NOBSX = 100 and NOBSY = 80 is 0.908185.
Probability for D = 0.180 with NOBSX = 100 and NOBSY = 100 is 0.946098.

```

---

## ANORDF/DNORDF (Single/Double precision)

Evaluate the standard normal (Gaussian) distribution function.

### Usage

ANORDF (X)

### Arguments

*X* — Argument for which the normal distribution function is to be evaluated.  
(Input)

*ANORDF* — Function value, the probability that a normal random variable takes a value less than or equal to *x*. (Output)

### Algorithm

Function ANORDF evaluates the distribution function,  $\Phi$ , of a standard normal (Gaussian) random variable, that is,

$$\Phi(x) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^x e^{-t^2/2} dt$$

The value of the distribution function at the point *x* is the probability that the random variable takes a value less than or equal to *x*.

The standard normal distribution (for which ANORDF is the distribution function) has mean of 0 and variance of 1. The probability that a normal random variable with mean and variance  $\sigma^2$  is less than *y* is given by ANORDF evaluated at  $(y - \mu)/\sigma$ .

$\Phi(x)$  is evaluated by use of the complementary error function, erfc. (See ERFC, IMSL MATH/LIBRARY Special Functions). The relationship is:

$$\Phi(x) = \text{erfc}(-x / \sqrt{2.0}) / 2$$

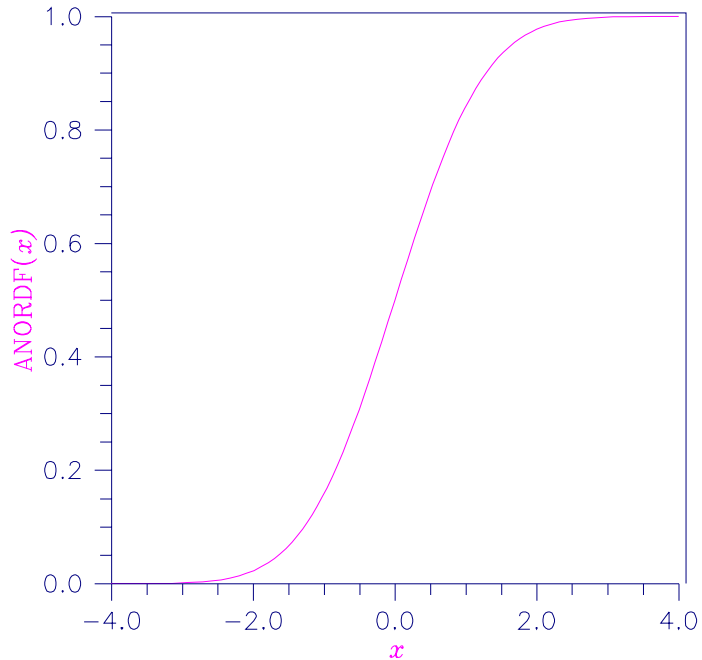


Figure 17-6 Standard Normal Distribution Function

### Example

Suppose  $X$  is a normal random variable with mean 100 and variance 225. In this example, we find the probability that  $X$  is less than 90, and the probability that  $X$  is between 105 and 110.

```

INTEGER      NOUT
REAL         ANORDF, P, X1, X2
EXTERNAL     ANORDF, UMACH

C
CALL UMACH (2, NOUT)
X1 = (90.0-100.0)/15.0
P = ANORDF(X1)
WRITE (NOUT,99998) P
99998 FORMAT (' The probability that X is less than 90 is ', F6.4)
X1 = (105.0-100.0)/15.0
X2 = (110.0-100.0)/15.0
P = ANORDF(X2) - ANORDF(X1)
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that X is between 105 and 110 is ',
&           F6.4)
END

```

### Output

```

The probability that X is less than 90 is 0.2525
The probability that X is between 105 and 110 is 0.1169

```



---

## ANORIN/DNORIN (Single/Double precision)

Evaluate the inverse of the standard normal (Gaussian) distribution function.

### Usage

ANORIN(P)

### Arguments

*P* — Probability for which the inverse of the normal distribution function is to be evaluated. (Input)

*P* must be in the open interval (0.0, 1.0).

*ANORIN* — Function value. (Output)

The probability that a standard normal random variable takes a value less than or equal to *ANORIN* is *P*.

### Algorithm

Function ANORIN evaluates the inverse of the distribution function,  $\Phi$ , of a standard normal (Gaussian) random variable, that is,  $\text{ANORIN}(P) = \Phi^{-1}(p)$ , where

$$\Phi(x) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^x e^{-t^2/2} dt$$

The value of the distribution function at the point  $x$  is the probability that the random variable takes a value less than or equal to  $x$ . The standard normal distribution has a mean of 0 and a variance of 1.

### Example

In this example, we compute the point such that the probability is 0.9 that a standard normal random variable is less than or equal to this point.

```
INTEGER      NOUT
REAL         ANORIN, P, X
EXTERNAL     ANORIN, UMACH
C
CALL UMACH (2, NOUT)
P = 0.9
X = ANORIN(P)
WRITE (NOUT,99999) X
99999 FORMAT (' The 90th percentile of a standard normal is ', F6.4)
END
```

### Output

The 90th percentile of a standard normal is 1.2816

---

## BETDF/DBETDF (Single/Double precision)

Evaluate the beta probability distribution function.

### Usage

BETDF(*X*, *PIN*, *QIN*)

### Arguments

*X* — Argument for which the beta distribution function is to be evaluated. (Input)

*PIN* — First beta distribution parameter. (Input)  
*PIN* must be positive.

*QIN* — Second beta distribution parameter. (Input)  
*QIN* must be positive.

*BETDF* — Probability that a random variable from a beta distribution having parameters *PIN* and *QIN* will be less than or equal to *X*. (Output)

### Comments

Informational errors

Type	Code	
1	1	Since the input argument <i>x</i> is less than or equal to zero, the distribution function is equal to zero at <i>x</i> .
1	2	Since the input argument <i>x</i> is greater than or equal to one, the distribution function is equal to one at <i>x</i> .

### Algorithm

Function BETDF evaluates the distribution function of a beta random variable with parameters *PIN* and *QIN*. This function is sometimes called the *incomplete beta ratio* and, with  $p = \text{PIN}$  and  $q = \text{QIN}$ , is denoted by  $I_x(p, q)$ . It is given by

$$I_x(p, q) = \frac{\Gamma(p)\Gamma(q)}{\Gamma(p+q)} \int_0^x t^{p-1} (1-t)^{q-1} dt$$

where  $\Gamma(\cdot)$  is the gamma function. The value of the distribution function  $I_x(p, q)$  is the probability that the random variable takes a value less than or equal to *x*.

The integral in the expression above is called the *incomplete beta function* and is denoted by  $\beta_x(p, q)$ . The constant in the expression is the reciprocal of the *beta function* (the incomplete function evaluated at one) and is denoted by  $\beta(p, q)$ .

Function BETDF uses the method of Bosten and Battiste (1974).

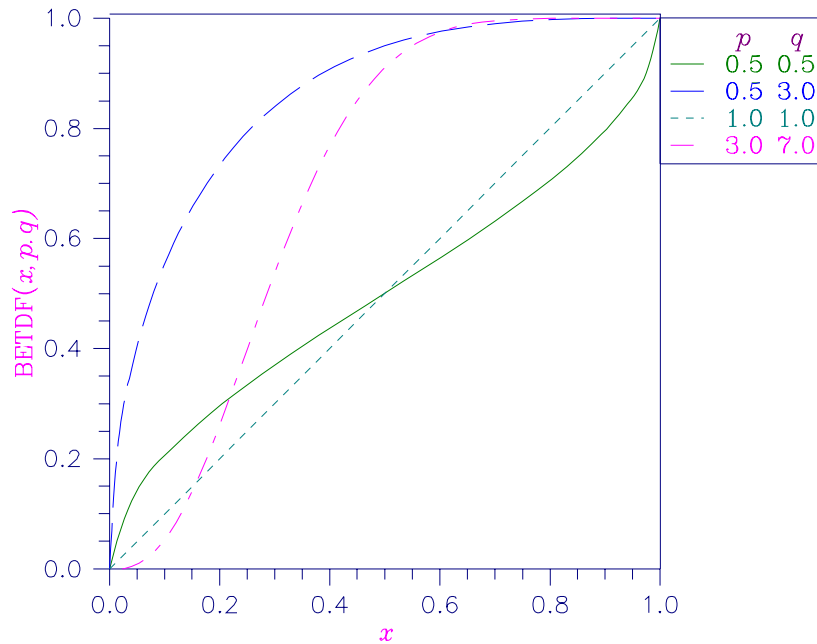


Figure 17-7 Beta Distribution Function

### Example

Suppose  $X$  is a beta random variable with parameters 12 and 12. ( $X$  has a symmetric distribution.) In this example, we find the probability that  $X$  is less than 0.6 and the probability that  $X$  is between 0.5 and 0.6. (Since  $X$  is a symmetric beta random variable, the probability that it is less than 0.5 is 0.5.)

```

INTEGER      NOUT
REAL         BETDF, P, PIN, QIN, X
EXTERNAL    BETDF, UMACH
C
CALL UMACH (2, NOUT)
PIN = 12.0
QIN = 12.0
X = 0.6
P = BETDF(X,PIN,QIN)
WRITE (NOUT,99998) P
99998 FORMAT (' The probability that X is less than 0.6 is ', F6.4)
X = 0.5
P = P - BETDF(X,PIN,QIN)
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that X is between 0.5 and 0.6 is ',
&           F6.4)
END

```

### Output

```

The probability that X is less than 0.6 is 0.8364
The probability that X is between 0.5 and 0.6 is 0.3364

```

---

## BETIN/DBETIN (Single/Double precision)

Evaluate the inverse of the beta distribution function.

### Usage

BETIN(*P*, *PIN*, *QIN*)

### Arguments

*P* — Probability for which the inverse of the beta distribution function is to be evaluated. (Input)

*P* must be in the open interval (0.0, 1.0).

*PIN* — First beta distribution parameter. (Input)

*PIN* must be positive.

*QIN* — Second beta distribution parameter. (Input)

*QIN* must be positive.

*BETIN* — Function value. (Output)

The probability that a beta random variable takes a value less than or equal to *BETIN* is *P*.

### Comments

Informational error

Type	Code
------	------

3	1	The value for the inverse Beta distribution could not be found in 100 iterations. The best approximation is used.
---	---	---

### Algorithm

The function `BETIN` evaluates the inverse distribution function of a beta random variable with parameters `PIN` and `QIN`, that is, with  $P = P$ ,  $p = PIN$ , and  $q = QIN$ , it determines  $x$  (equal to `BETIN(P, PIN, QIN)`), such that

$$P = \frac{\Gamma(p)\Gamma(q)}{\Gamma(p+q)} \int_0^x t^{p-1} (1-t)^{q-1} dt$$

where  $\Gamma(\cdot)$  is the gamma function. The probability that the random variable takes a value less than or equal to  $x$  is  $P$ .

### Example

Suppose  $X$  is a beta random variable with parameters 12 and 12. ( $X$  has a symmetric distribution.) In this example, we find the value  $x_0$  such that the probability that  $X \leq x_0$  is 0.9.

INTEGER	NOUT
REAL	BETIN, P, PIN, QIN, X
EXTERNAL	BETIN, UMACH

```

C
CALL UMACH (2, NOUT)
PIN = 12.0
QIN = 12.0
P   = 0.9
X   = BETIN(P,PIN,QIN)
WRITE (NOUT,99999) X
99999 FORMAT (' X is less than ', F6.4, ' with probability 0.9.')
END

```

### Output

X is less than 0.6299 with probability 0.9.

---

## BNRDF/DBNRDF (Single/Double precision)

Evaluate the bivariate normal distribution function.

### Usage

BNRDF(*X*, *Y*, *RHO*)

### Arguments

*X* — One argument for which the bivariate normal distribution function is to be evaluated. (Input)

*Y* — The other argument for which the bivariate normal distribution function is to be evaluated. (Input)

*RHO* — Correlation coefficient. (Input)

*BNRDF* — Function value, the probability that a bivariate normal random variable with correlation *RHO* takes a value less than or equal to *X* and less than or equal to *Y*. (Output)

### Algorithm

Function *BNRDF* evaluates the distribution function *F* of a bivariate normal distribution with means of zero, variances of one, and correlation of *RHO*; that is, with  $\rho = \text{RHO}$ , and  $|\rho| < 1$ ,

$$F(x, y) = \frac{1}{2\pi\sqrt{1-\rho^2}} \int_{-\infty}^x \int_{-\infty}^y \exp\left(-\frac{u^2 - 2\rho uv + v^2}{2(1-\rho^2)}\right) du dv$$

To determine the probability that  $U \leq u_0$  and  $V \leq v_0$ , where  $(U, V)^T$  is a bivariate normal random variable with mean  $\mu = (\mu_U, \mu_V)^T$  and variance-covariance matrix

$$\Sigma = \begin{pmatrix} \sigma_U^2 & \sigma_{UV} \\ \sigma_{UV} & \sigma_V^2 \end{pmatrix}$$

transform  $(U, V)^T$  to a vector with zero means and unit variances. The input to BNRDF would be  $X = (u_0 - \mu_U)/\sigma_U$ ,  $Y = (v_0 - \mu_V)/\sigma_V$ , and  $\rho = \sigma_{UV}/(\sigma_U\sigma_V)$ .

Function BNRDF uses the method of Owen (1962, 1965). For  $|\rho| = 1$ , the distribution function is computed based on the univariate statistic,  $Z = \min(x, y)$ , and on the normal distribution function ANORDF (page 1122).

### Example

Suppose  $(X, Y)$  is a bivariate normal random variable with mean  $(0, 0)$  and variance-covariance matrix

$$\begin{pmatrix} 1.0 & 0.9 \\ 0.9 & 1.0 \end{pmatrix}$$

In this example, we find the probability that  $X$  is less than  $-2.0$  and  $Y$  is less than  $0.0$ .

```

C
INTEGER      NOUT
REAL         BNRDF, P, RHO, X, Y
EXTERNAL     BNRDF, UMACH

CALL UMACH (2, NOUT)
X   = -2.0
Y   = 0.0
RHO = 0.9
P   = BNRDF(X,Y,RHO)
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that X is less than -2.0  and Y ',
&           'is less than 0.0 is ', F6.4)
END
```

### Output

The probability that  $X$  is less than  $-2.0$  and  $Y$  is less than  $0.0$  is  $0.0228$

---

## CHIDF/DCHIDF (Single/Double precision)

Evaluate the chi-squared distribution function.

### Usage

CHIDF(CHSQ, DF)

### Arguments

**CHSQ** — Argument for which the chi-squared distribution function is to be evaluated. (Input)

**DF** — Number of degrees of freedom of the chi-squared distribution. (Input)  
DF must be greater than or equal to 0.5.

**CHIDF** — Function value, the probability that a chi-squared random variable takes a value less than or equal to CHSQ. (Output)

### Comments

Informational errors

Type	Code	
1	1	Since the input argument, CHSQ, is less than zero, the distribution function is zero at CHSQ.
2	3	The normal distribution is used for large degrees of freedom. However, it has produced underflow. Therefore, the probability, CHIDF, is set to zero.

### Algorithm

Function CHIDF evaluates the distribution function,  $F$ , of a chi-squared random variable with DF degrees of freedom, that is, with  $\nu = DF$ , and  $x = CHSQ$ ,

$$F(x) = \frac{1}{2^{\nu/2} \Gamma(\nu/2)} \int_0^x e^{-t/2} t^{\nu/2-1} dt$$

where  $\Gamma(\cdot)$  is the gamma function. The value of the distribution function at the point  $x$  is the probability that the random variable takes a value less than or equal to  $x$ .

For  $\nu > 65$ , CHIDF uses the Wilson-Hilferty approximation (Abramowitz and Stegun 1964, equation 26.4.17) to the normal distribution, and routine ANORDF (page 1122) is used to evaluate the normal distribution function.

For  $\nu \leq 65$ , CHIDF uses series expansions to evaluate the distribution function. If  $x < \max(\nu/2, 26)$ , CHIDF uses the series 6.5.29 in Abramowitz and Stegun (1964), otherwise, it uses the asymptotic expansion 6.5.32 in Abramowitz and Stegun.

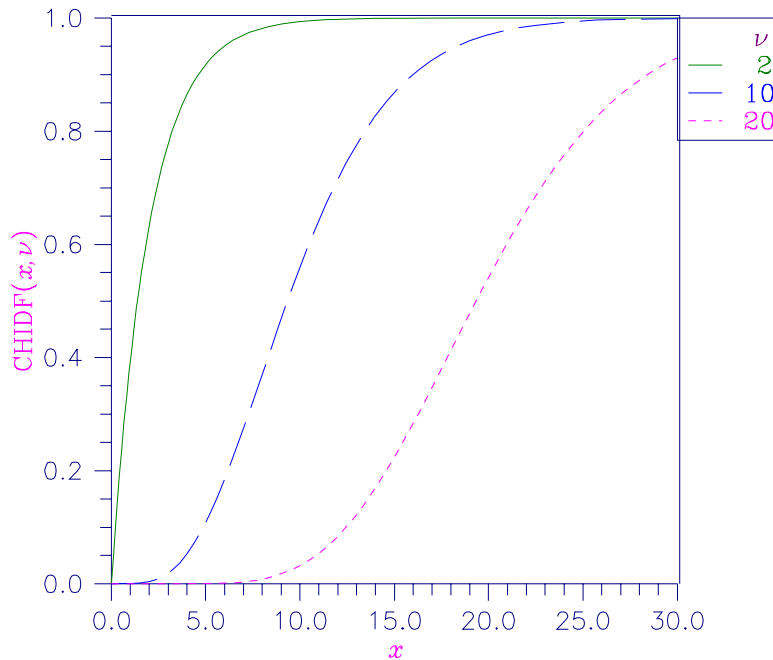


Figure 17-8 Chi-Squared Distribution Function

### Example

Suppose  $X$  is a chi-squared random variable with 2 degrees of freedom. In this example, we find the probability that  $X$  is less than 0.15 and the probability that  $X$  is greater than 3.0.

```

INTEGER      NOUT
REAL         CHIDF, CHSQ, DF, P
EXTERNAL     CHIDF, UMACH

C
CALL UMACH (2, NOUT)
DF   = 2.0
CHSQ = 0.15
P    = CHIDF(CHSQ,DF)
WRITE (NOUT,99998) P
99998 FORMAT (' The probability that chi-squared with 2 df is less ',
&            ' than 0.15 is ', F6.4)
CHSQ = 3.0
P    = 1.0 - CHIDF(CHSQ,DF)
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that chi-squared with 2 df is greater ',
&            ' than 3.0 is ', F6.4)
END

```

### Output

The probability that chi-squared with 2 df is less than 0.15 is 0.0723  
The probability that chi-squared with 2 df is greater than 3.0 is 0.2231



---

## CHIIN/DCHIIN (Single/Double precision)

Evaluate the inverse of the chi-squared distribution function.

### Usage

CHIIN(*P*, *DF*)

### Arguments

*P* — Probability for which the inverse of the chi-squared distribution function is to be evaluated. (Input)

*P* must be in the open interval (0.0, 1.0).

*DF* — Number of degrees of freedom of the chi-squared distribution. (Input)

*DF* must be greater than or equal to 0.5.

*CHIIN* — Function value. (Output)

The probability that a chi-squared random variable takes a value less than or equal to *CHIIN* is *P*.

### Comments

Informational errors

Type	Code	
4	1	Over 100 iterations have occurred without convergence. Convergence is assumed.

### Algorithm

Function *CHIIN* evaluates the inverse distribution function of a chi-squared random variable with *DF* degrees of freedom, that is, with  $P = P$  and  $\nu = DF$ , it determines  $x$  (equal to *CHIIN*(*P*, *DF*)), such that

$$P = \frac{1}{2^{\nu/2} \Gamma(\nu/2)} \int_0^x e^{-t/2} t^{\nu/2-1} dt$$

where  $\Gamma(\cdot)$  is the gamma function. The probability that the random variable takes a value less than or equal to  $x$  is  $P$ .

For  $\nu < 40$ , *CHIIN* uses bisection (if  $\nu \leq 2$  or  $P > 0.98$ ) or regula falsi to find the point at which the chi-squared distribution function is equal to  $P$ . The distribution function is evaluated using routine *CHIDF* (page 1129).

For  $40 \leq \nu < 100$ , a modified Wilson-Hilferty approximation (Abramowitz and Stegun 1964, equation 26.4.18) to the normal distribution is used, and routine *ANORIN* (page 1124) is used to evaluate the inverse of the normal distribution function. For  $\nu \geq 100$ , the ordinary Wilson-Hilferty approximation (Abramowitz and Stegun 1964, equation 26.4.17) is used.

### Example

In this example, we find the 99-th percentage points of a chi-squared random variable with 2 degrees of freedom and of one with 64 degrees of freedom.

```
INTEGER      NOUT
REAL         CHIIN, DF, P, X
EXTERNAL     CHIIN, UMACH

C
CALL UMACH (2, NOUT)
P = 0.99
DF = 2.0
X = CHIIN(P,DF)
WRITE (NOUT,99998) X
99998 FORMAT (' The 99-th percentage point of chi-squared with 2 df '
&           , 'is ', F7.3)
DF = 64.0
X = CHIIN(P,DF)
WRITE (NOUT,99999) X
99999 FORMAT (' The 99-th percentage point of chi-squared with 64 df '
&           , 'is ', F7.3)
END
```

### Output

```
The 99-th percentage point of chi-squared with 2 df is 9.210
The 99-th percentage point of chi-squared with 64 df is 93.217
```

---

## CSNDF/DCSNDF (Single/Double precision)

Evaluate the noncentral chi-squared distribution function.

### Usage

```
CSNDF(CHSQ, DF, ALAM)
```

### Arguments

**CHSQ** — Argument for which the noncentral chi-squared distribution function is to be evaluated. (Input)

**DF** — Number of degrees of freedom of the noncentral chi-squared distribution. (Input)

DF must be greater than or equal to 0.5 and less than or equal to 200,000.

**ALAM** — The noncentrality parameter. (Input)

ALAM must be nonnegative, and ALAM + DF must be less than or equal to 200,000.

**CSNDF** — Function value, the probability that a noncentral chi-squared random variable takes a value less than or equal to CHSQ. (Output)

### Comments

1. Informational errors

Type	Code	
1	1	Since the input argument, CHSQ, is less than or equal to zero, the distribution function is zero at CHSQ.
3	2	Convergence was not obtained. The best approximation to the probability is returned.

2. This subroutine sums terms of an infinite series of central chi-squared distribution functions weighted by Poisson terms. Summing terminates when either the current term is less than  $10 * AMACH(4)$  times the current sum or when 1000 terms have been accumulated. In the latter case, a warning error is issued.

### Algorithm

Function CSNDF evaluates the distribution function of a noncentral chi-squared random variable with DF degrees of freedom and noncentrality parameter ALAM, that is, with  $\nu = DF$ ,  $\lambda = ALAM$ , and  $x = CHSQ$ ,

$$CSNDF(x) = \sum_{i=0}^{\infty} \frac{e^{-\lambda/2} (\lambda/2)^i}{i!} \int_0^x \frac{t^{(\nu+2i)/2-1} e^{-t/2}}{2^{(\nu+2i)/2} \Gamma(\frac{\nu+2i}{2})} dt$$

where  $\Gamma(\cdot)$  is the gamma function. This is a series of central chi-squared distribution functions with Poisson weights. The value of the distribution function at the point  $x$  is the probability that the random variable takes a value less than or equal to  $x$ .

The noncentral chi-squared random variable can be defined by the distribution function above, or alternatively and equivalently, as the sum of squares of independent normal random variables. If  $Y_i$  have independent normal distributions with means  $\mu_i$  and variances equal to one and

$$X = \sum_{i=1}^n Y_i^2$$

then  $X$  has a noncentral chi-squared distribution with  $n$  degrees of freedom and noncentrality parameter equal to

$$\sum_{i=1}^n \mu_i^2$$

With a noncentrality parameter of zero, the noncentral chi-squared distribution is the same as the chi-squared distribution.

Function CSNDF determines the point at which the Poisson weight is greatest, and then sums forward and backward from that point, terminating when the additional terms are sufficiently small or when a maximum of 1000 terms have been accumulated. The recurrence relation 26.4.8 of Abramowitz and Stegun (1964) is used to speed the evaluation of the central chi-squared distribution functions.

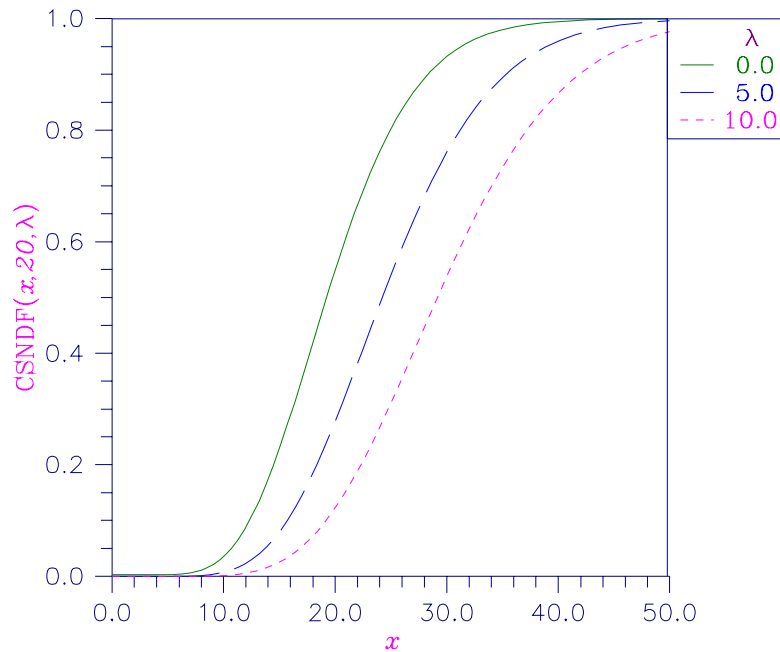


Figure 17-9 Noncentral Chi-squared Distribution Function

### Example

In this example, CSNDF is used to compute the probability that a random variable that follows the noncentral chi-squared distribution with noncentrality parameter of 1 and with 2 degrees of freedom is less than or equal to 8.642.

```

INTEGER      NOUT
REAL         ALAM, CHSQ, CSNDF, DF, P
EXTERNAL    CSNDF, UMACH

C
CALL UMACH (2, NOUT)
DF   = 2.0
ALAM = 1.0
CHSQ = 8.642
P     = CSNDF(CHSQ,DF,ALAM)
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that a noncentral chi-squared random',
&           //, ' variable with 2 df and noncentrality 1.0 is less',
&           //, ' than 8.642 is ', F5.3)
END

```

### Output

```

The probability that a noncentral chi-squared random
variable with 2 df and noncentrality 1.0 is less
than 8.642 is 0.950

```

---

## CSNIN/DCSNIN (Single/Double precision)

Evaluate the inverse of the noncentral chi-squared function.

### Usage

CSNIN(*P*, *DF*, *ALAM*)

### Arguments

*P* — Probability for which the inverse of the noncentral chi-squared distribution function is to be evaluated. (Input)

*P* must be in the open interval (0.0, 1.0).

*DF* — Number of degrees of freedom of the noncentral chi-squared distribution. (Input)

*DF* must be greater than or equal to 0.5 and less than or equal to 200,000.

*ALAM* — The noncentrality parameter. (Input)

*ALAM* must be nonnegative, and *ALAM* + *DF* must be less than or equal to 200,000.

*CSNIN* — Function value. (Output)

The probability that a noncentral chi-squared random variable takes a value less than or equal to *CSNIN* is *P*.

### Comments

Informational error

Type	Code
------	------

4	1	Over 100 iterations have occurred without convergence. Convergence is assumed.
---	---	---

### Algorithm

Function CSNIN evaluates the inverse distribution function of a noncentral chi-squared random variable with *DF* degrees of freedom and noncentrality parameter *ALAM*; that is, with  $P = P$ ,  $\nu = DF$ , and  $\lambda = ALAM$ , it determines  $c_0 (= CSNIN(P, DF, ALAM))$ , such that

$$P = \sum_{i=0}^{\infty} \frac{e^{-\lambda/2} (\lambda/2)^i}{i!} \int_0^{c_0} \frac{x^{(\nu+2i)/2-1} e^{-x/2}}{2^{(\nu+2i)/2} \Gamma(\frac{\nu+2i}{2})} dx$$

where  $\Gamma(\cdot)$  is the gamma function. The probability that the random variable takes a value less than or equal to  $c_0$  is  $P$ .

Function CSNIN uses bisection and modified regula falsi to invert the distribution function, which is evaluated using routine CSNDF (page 1133). See CSNDF for an alternative definition of the noncentral chi-squared random variable in terms of normal random variables.

### Example

In this example, we find the 95-th percentage point for a noncentral chi-squared random variable with 2 degrees of freedom and noncentrality parameter 1.

```
INTEGER      NOUT
REAL         ALAM, CHSQ, CSNIN, DF, P
EXTERNAL     CSNIN, UMACH

C
CALL UMACH (2, NOUT)
DF      = 2.0
ALAM    = 1.0
P       = 0.95
CHSQ    = CSNIN(P,DF,ALAM)
WRITE (NOUT,99999) CHSQ

C
99999 FORMAT (' The 0.05 noncentral chi-squared critical value is ',
&           F6.3, '.')

C
END
```

### Output

The 0.05 noncentral chi-squared critical value is 8.642.

---

## FDF/DFDF (Single/Double precision)

Evaluate the  $F$  distribution function.

### Usage

FDF( $F$ ,  $DFN$ ,  $DFD$ )

### Arguments

$F$  — Argument for which the  $F$  distribution function is to be evaluated. (Input)

$DFN$  — Numerator degrees of freedom. (Input)

$DFN$  must be positive.

$DFD$  — Denominator degrees of freedom. (Input)

$DFD$  must be positive.

$FDF$  — Function value, the probability that an  $F$  random variable takes a value less than or equal to the input  $F$ . (Output)

### Comments

Informational error

Type	Code
------	------

1	3	Since the input argument $F$ is not positive, the distribution function is zero at $F$ .
---	---	--

## Algorithm

Function `FDF` evaluates the distribution function of a Snedecor's  $F$  random variable with `DFN` numerator degrees of freedom and `DFD` denominator degrees of freedom. The function is evaluated by making a transformation to a beta random variable and then using the routine `BETDF` (page 1125). If  $X$  is an  $F$  variate with  $\nu_1$  and  $\nu_2$  degrees of freedom and  $Y = \nu_1 X / (\nu_2 + \nu_1 X)$ , then  $Y$  is a beta variate with parameters  $p = \nu_1/2$  and  $q = \nu_2/2$ . The function `FDF` also uses a relationship between  $F$  random variables that can be expressed as follows.

$$\text{FDF}(X, \text{DFN}, \text{DFD}) = 1.0 - \text{FDF}(1.0/X, \text{DFD}, \text{DFN})$$

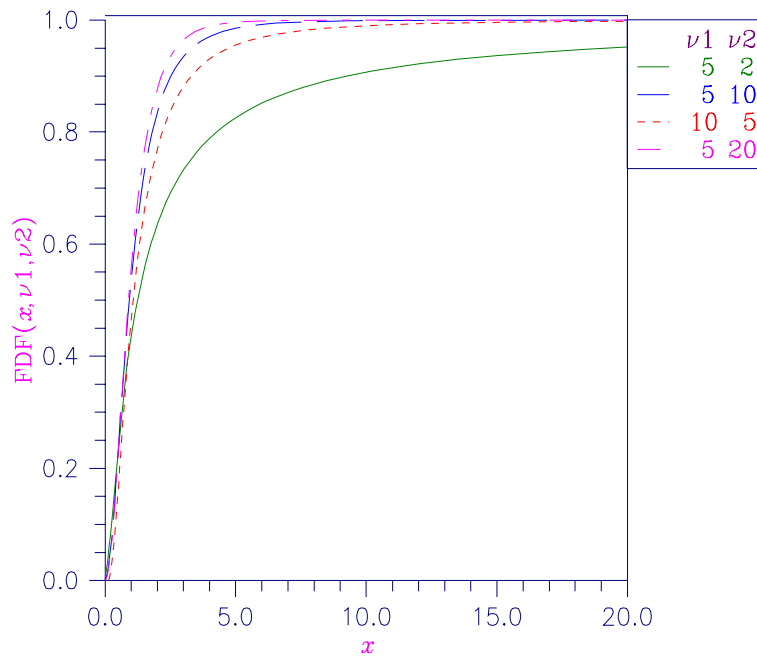


Figure 17-10  $F$  Distribution Function

## Example

In this example, we find the probability that an  $F$  random variable with one numerator and one denominator degree of freedom is greater than 648.

```
C
INTEGER      NOUT
REAL         DFD, DFN, F, FDF, P
EXTERNAL     FDF, UMACH

CALL UMACH (2, NOUT)
F = 648.0
DFN = 1.0
DFD = 1.0
P = 1.0 - FDF(F, DFN, DFD)
WRITE (NOUT, 99999) P
```

```

99999 FORMAT (' The probability that an F(1,1) variate is greater ',
&           ' than 648 is ', F6.4)
END

```

### Output

The probability that an F(1, 1) variate is greater than 648 is 0.0250

## FIN/DFIN (Single/Double precision)

Evaluate the inverse of the  $F$  distribution function.

### Usage

FIN( $P$ ,  $DFN$ ,  $DFD$ )

### Arguments

$P$  — Probability for which the inverse of the  $F$  distribution function is to be evaluated. (Input)

$P$  must be in the open interval (0.0, 1.0).

$DFN$  — Numerator degrees of freedom. (Input)

$DFN$  must be positive.

$DFD$  — Denominator degrees of freedom. (Input)

$DFD$  must be positive.

$FIN$  — Function value. (Output)

The probability that an  $F$  random variable takes a value less than or equal to  $FIN$  is  $P$ .

### Comments

Informational error

Type	Code
4	4

$FIN$  is set to machine infinity since overflow would occur upon modifying the inverse value for the  $F$  distribution with the result obtained from the inverse beta distribution.

### Algorithm

Function  $FIN$  evaluates the inverse distribution function of a Snedecor's  $F$  random variable with  $DFN$  numerator degrees of freedom and  $DFD$  denominator degrees of freedom. The function is evaluated by making a transformation to a beta random variable and then using the routine  $BETIN$  (page 1127). If  $X$  is an  $F$  variate with  $\nu_1$  and  $\nu_2$  degrees of freedom and  $Y = \nu_1 X / (\nu_2 + \nu_1 X)$ , then  $Y$  is a beta variate with parameters  $p = \nu_1/2$  and  $q = \nu_2/2$ . If  $P \leq 0.5$ ,  $FIN$  uses this relationship directly, otherwise, it also uses a relationship between  $F$  random variables that can be expressed as follows, using routine  $FDF$  (page 1137), which is the  $F$  cumulative distribution function:



$FDF(F, DFN, DFD) = 1.0 - FDF(1.0/F, DFD, DFN).$

### Example

In this example, we find the 99-th percentage point for an  $F$  random variable with 1 and 7 degrees of freedom.

```
INTEGER      NOUT
REAL         DFD, DFN, F, FIN, P
EXTERNAL    FIN, UMACH
C
CALL UMACH (2, NOUT)
P = 0.99
DFN = 1.0
DFD = 7.0
F = FIN(P, DFN, DFD)
WRITE (NOUT, 99999) F
99999 FORMAT (' The F(1,7) 0.01 critical value is ', F6.3)
END
```

### Output

The F(1, 7) 0.01 critical value is 12.246

---

## GAMDF/DGAMDF (Single/Double precision)

Evaluate the gamma distribution function.

### Usage

$GAMDF(X, A)$

### Arguments

$X$  — Argument for which the gamma distribution function is to be evaluated. (Input)

$A$  — The shape parameter of the gamma distribution. (Input)  
This parameter must be positive.

$GAMDF$  — Function value, the probability that a gamma random variable takes a value less than or equal to  $x$ . (Output)

### Comments

Informational error

Type Code

1	2	Since the input argument $x$ is less than zero, the distribution function is set to zero.
---	---	---

### Algorithm

Function  $GAMDF$  evaluates the distribution function,  $F$ , of a gamma random variable with shape parameter  $a$ ; that is,

$$F(x) = \frac{1}{\Gamma(a)} \int_0^x e^{-t} t^{a-1} dt$$

where  $\Gamma(\cdot)$  is the gamma function. (The gamma function is the integral from 0 to  $\infty$  of the same integrand as above). The value of the distribution function at the point  $x$  is the probability that the random variable takes a value less than or equal to  $x$ .

The gamma distribution is often defined as a two-parameter distribution with a scale parameter  $b$  (which must be positive), or even as a three-parameter distribution in which the third parameter  $c$  is a location parameter. In the most general case, the probability density function over  $(c, \infty)$  is

$$f(t) = \frac{1}{b^a \Gamma(a)} e^{-(t-c)/b} (x-c)^{a-1}$$

If  $T$  is such a random variable with parameters  $a$ ,  $b$ , and  $c$ , the probability that  $T \leq t_0$  can be obtained from GAMDF by setting  $x = (t_0 - c)/b$ .

If  $x$  is less than  $a$  or if  $x$  is less than or equal to 1.0, GAMDF uses a series expansion. Otherwise, a continued fraction expansion is used. (See Abramowitz and Stegun, 1964.)

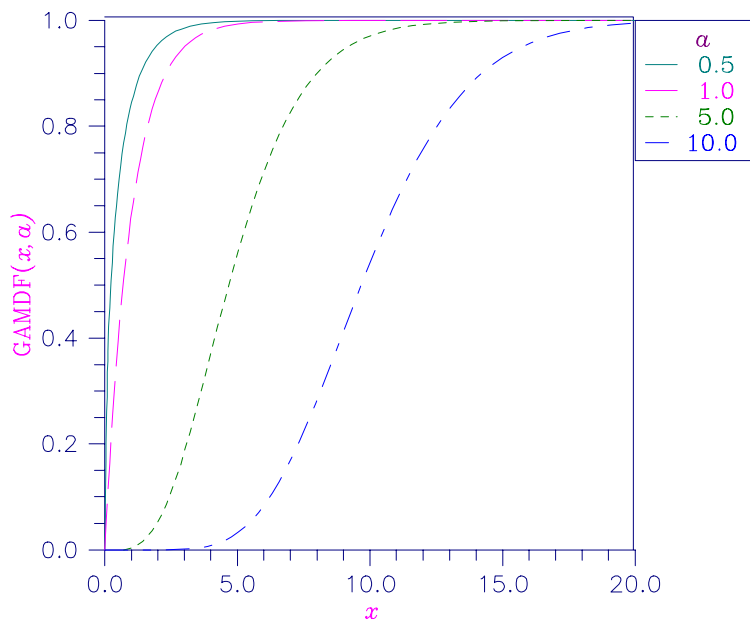


Figure 17-11 Gamma Distribution Function

### Example

Suppose  $X$  is a gamma random variable with a shape parameter of 4. (In this case, it has an *Erlang distribution* since the shape parameter is an integer.) In this example, we find the probability that  $X$  is less than 0.5 and the probability that  $X$  is between 0.5 and 1.0.

```
INTEGER      NOUT
REAL         A, GAMDF, P, X
EXTERNAL     GAMDF, UMACH

C
CALL UMACH (2, NOUT)
A = 4.0
X = 0.5
P = GAMDF(X,A)
WRITE (NOUT,99998) P
99998 FORMAT (' The probability that X is less than 0.5 is ', F6.4)
X = 1.0
P = GAMDF(X,A) - P
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that X is between 0.5 and 1.0 is ',
&           F6.4)
END
```

### Output

```
The probability that X is less than 0.5 is 0.0018
The probability that X is between 0.5 and 1.0 is 0.0172
```

---

## GAMIN/DGAMIN (Single/Double precision)

Evaluate the inverse of the gamma distribution function.

### Usage

GAMIN(P, A)

### Arguments

**P** — Probability for which the inverse of the gamma distribution function is to be evaluated. (Input)

**P** must be in the open interval (0.0, 1.0).

**A** — The shape parameter of the gamma distribution. (Input)

This parameter must be positive.

**GAMIN** — Function value. (Output)

The probability that a gamma random variable takes a value less than or equal to GAMIN is P.

### Comments

Informational error

Type Code

4            1            Over 100 iterations have occurred without convergence.  
Convergence is assumed.

### Algorithm

Function `GAMIN` evaluates the inverse distribution function of a gamma random variable with shape parameter  $a$ , that is, it determines  $x$  ( $= \text{GAMIN}(P, A)$ ), such that

$$P = \frac{1}{\Gamma(a)} \int_0^x e^{-t} t^{a-1} dt$$

where  $\Gamma(\cdot)$  is the gamma function. The probability that the random variable takes a value less than or equal to  $x$  is  $P$ . See the documentation for routine `GAMDF` (page 1140) for further discussion of the gamma distribution.

Function `GAMIN` uses bisection and modified regula falsi to invert the distribution function, which is evaluated using routine `GAMDF`.

### Example

In this example, we find the 95-th percentage point for a gamma random variable with shape parameter of 4.

```
INTEGER    NOUT
REAL       A, GAMIN, P, X
EXTERNAL   GAMIN, UMACH

C
CALL UMACH (2, NOUT)
A = 4.0
P = 0.95
X = GAMIN(P,A)
WRITE (NOUT,99999) X

C
99999 FORMAT (' The 0.05 gamma(4) critical value is ', F6.3,
&            '.')
```

### Output

The 0.05 gamma(4) critical value is 7.754.

---

## TDF/DTDF (Single/Double precision)

Evaluate the Student's  $t$  distribution function.

### Usage

`TDF(T, DF)`

## Arguments

**$T$**  — Argument for which the Student's  $t$  distribution function is to be evaluated. (Input)

**$DF$**  — Degrees of freedom. (Input)  
 $DF$  must be greater than or equal to 1.0.

**$TDF$**  — Function value, the probability that a Student's  $t$  random variable takes a value less than or equal to the input  $T$ . (Output)

## Algorithm

Function  $TDF$  evaluates the distribution function of a Student's  $t$  random variable with  $DF$  degrees of freedom. If the square of  $T$  is greater than or equal to  $DF$ , the relationship of a  $t$  to an  $F$  random variable (and subsequently, to a beta random variable) is exploited, and routine  $BETDF$  (page 1125) is used. Otherwise, the method described by Hill (1970) is used. If  $DF$  is not an integer, if  $DF$  is greater than 19, or if  $DF$  is greater than 200, a Cornish-Fisher expansion is used to evaluate the distribution function. If  $DF$  is less than 20 and  $ABS(T)$  is less than 2.0, a trigonometric series (see Abramowitz and Stegun 1964, equations 26.7.3 and 26.7.4, with some rearrangement) is used. For the remaining cases, a series given by Hill (1970) that converges well for large values of  $T$  is used.

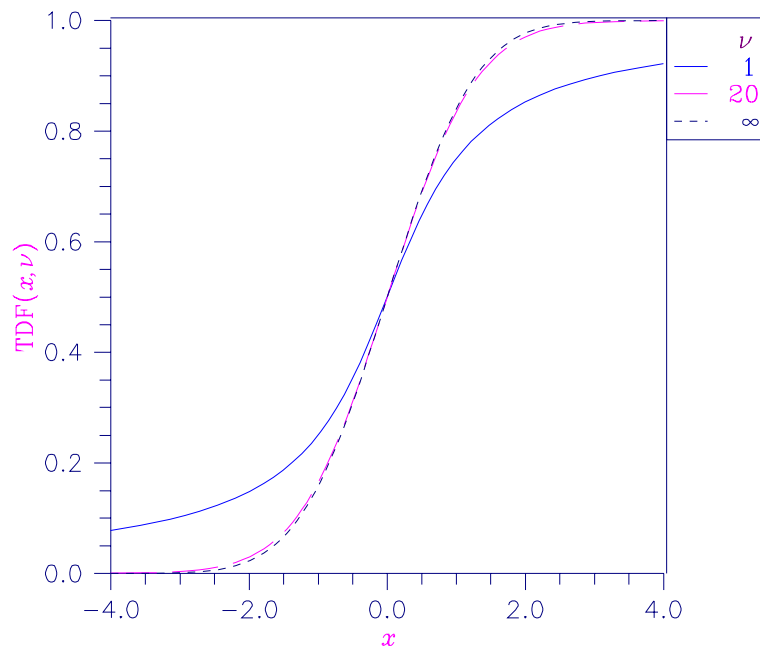


Figure 17-12 Student's  $t$  Distribution Function

### Example

In this example, we find the probability that a  $t$  random variable with 6 degrees of freedom is greater in absolute value than 2.447. We use the fact that  $t$  is symmetric about 0.

```
INTEGER      NOUT
REAL         DF, P, T, TDF
EXTERNAL     TDF, UMACH

C
CALL UMACH (2, NOUT)
T = 2.447
DF = 6.0
P = 2.0*TDF(-T,DF)
WRITE (NOUT,99999) P
99999 FORMAT (' The probability that a t(6) variate is greater ',
&           'than 2.447 in', '/', ' absolute value is ', F6.4)
END
```

### Output

The probability that a  $t(6)$  variate is greater than 2.447 in absolute value is 0.0500

---

## TIN/DTIN (Single/Double precision)

Evaluate the inverse of the Student's  $t$  distribution function.

### Usage

TIN(P, DF)

### Arguments

**P** — Probability for which the inverse of the Student's  $t$  distribution function is to be evaluated. (Input)

P must be in the open interval (0.0, 1.0).

**DF** — Degrees of freedom. (Input)

DF must be greater than or equal to 1.0.

**TIN** — Function value. (Output)

The probability that a Student's  $t$  random variable takes a value less than or equal to TIN is P.

### Comments

Informational error

Type	Code
------	------

4	3
---	---

TIN is set to machine infinity since overflow would occur upon modifying the inverse value for the  $F$  distribution with the result obtained from the inverse  $\beta$  distribution.

## Algorithm

Function `TIN` evaluates the inverse distribution function of a Student's  $t$  random variable with `DF` degrees of freedom. Let  $\nu = DF$ . If  $\nu$  equals 1 or 2, the inverse can be obtained in closed form, if  $\nu$  is between 1 and 2, the relationship of a  $t$  to a beta random variable is exploited and routine `BETIN` (page 1127) is used to evaluate the inverse; otherwise the algorithm of Hill (1970) is used. For small values of  $\nu$  greater than 2, Hill's algorithm inverts an integrated expansion in  $1/(1 + t^2/\nu)$  of the  $t$  density. For larger values, an asymptotic inverse Cornish-Fisher type expansion about normal deviates is used.

## Example

In this example, we find the 0.05 critical value for a two-sided  $t$  test with 6 degrees of freedom.

```
C
INTEGER      NOUT
REAL         DF, P, T, TIN
EXTERNAL     TIN, UMACH

CALL UMACH (2, NOUT)
P = 0.975
DF = 6.0
T = TIN(P,DF)
WRITE (NOUT,99999) T
99999 FORMAT (' The two-sided t(6) 0.05 critical value is ', F6.3)
END
```

## Output

```
The two-sided t(6) 0.05 critical value is  2.447
```

---

# TNDF/DTNDF (Single/Double precision)

Evaluate the noncentral Student's  $t$  distribution function.

## Usage

```
TNDF(T, IDF, DELTA)
```

## Arguments

***T*** — Argument for which the noncentral Student's  $t$  distribution function is to be evaluated. (Input)

***IDF*** — Number of degrees of freedom of the noncentral Student's  $t$  distribution. (Input)

*IDF* must be positive.

***DELTA*** — The noncentrality parameter. (Input)

***TNDF*** — Function value, the probability that a noncentral Student's  $t$  random variable takes a value less than or equal to  $T$ . (Output)

### Algorithm

Function `TNDF` evaluates the distribution function  $F$  of a noncentral  $t$  random variable with `IDF` degrees of freedom and noncentrality parameter `DELTA`; that is, with  $v = \text{IDF}$ ,  $\delta = \text{DELTA}$ , and  $t_0 = T$ ,

$$F(t_0) = \int_{-\infty}^{t_0} \frac{v^{v/2} e^{-\delta^2/2}}{\sqrt{\pi} \Gamma(v/2) (v+x^2)^{(v+1)/2}} \sum_{i=0}^{\infty} \Gamma((v+i+1)/2) \left(\frac{\delta^i}{i!}\right) \left(\frac{2x^2}{v+x^2}\right)^{i/2} dx$$

where  $\Gamma(\cdot)$  is the gamma function. The value of the distribution function at the point  $t_0$  is the probability that the random variable takes a value less than or equal to  $t_0$ .

The noncentral  $t$  random variable can be defined by the distribution function above, or alternatively and equivalently, as the ratio of a normal random variable and an independent chi-squared random variable. If  $w$  has a normal distribution with mean  $\delta$  and variance equal to one,  $u$  has an independent chi-squared distribution with  $v$  degrees of freedom, and

$$x = w / \sqrt{u/v}$$

then  $x$  has a noncentral  $t$  distribution with degrees of freedom and noncentrality parameter  $\delta$ .

The distribution function of the noncentral  $t$  can also be expressed as a double integral involving a normal density function (see, for example, Owen 1962, page 108). The function `TNDF` uses the method of Owen (1962, 1965), which uses repeated integration by parts on that alternate expression for the distribution function.



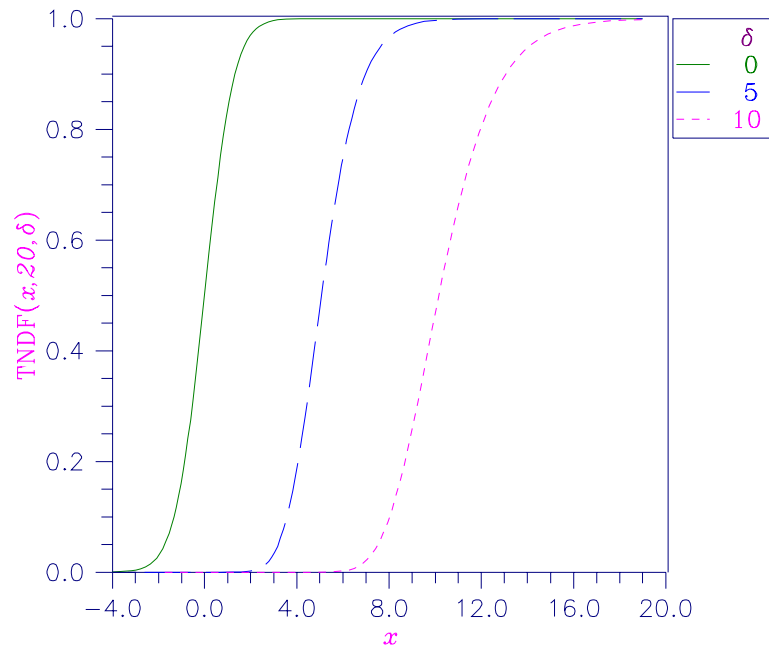


Figure 17-13 Noncentral Student's  $t$  Distribution Function

### Example

Suppose  $T$  is a noncentral  $t$  random variable with 6 degrees of freedom and noncentrality parameter 6. In this example, we find the probability that  $T$  is less than 12.0. (This can be checked using the table on page 111 of Owen 1962, with  $\eta = 0.866$ , which yields  $\lambda = 1.664$ .)

```

INTEGER      IDF, NOUT
REAL         DELTA, P, T, TNDF
EXTERNAL     TNDF, UMACH
C
CALL UMACH (2, NOUT)
IDF = 6
DELTA = 6.0
T = 12.0
P = TNDF(T, IDF, DELTA)
WRITE (NOUT, 99999) P
99999 FORMAT (' The probability that T is less than 12.0 is ', F6.4)
END

```

### Output

The probability that T is less than 12.0 is 0.9501

---

## TNIN/DTNIN (Single/Double precision)

Evaluate the inverse of the noncentral Student's  $t$  distribution function.

### Usage

TNIN( $P$ ,  $IDF$ ,  $DELTA$ )

### Arguments

$P$  — Probability for which the inverse of the noncentral Student's  $t$  distribution function is to be evaluated. (Input)

$P$  must be in the open interval (0.0, 1.0).

$IDF$  — Number of degrees of freedom of the noncentral Student's  $t$  distribution. (Input)  $IDF$  must be positive.

$DELTA$  — The noncentrality parameter. (Input)

$TNIN$  — Function value. (Output)

The probability that a noncentral Student's  $t$  random variable takes a value less than or equal to  $TNIN$  is  $P$ .

### Comments

Informational error

Type	Code	
4	1	Over 100 iterations have occurred without convergence. Convergence is assumed.

### Algorithm

Function  $TNIN$  evaluates the inverse distribution function of a noncentral  $t$  random variable with  $IDF$  degrees of freedom and noncentrality parameter  $DELTA$ ; that is, with  $P = P$ ,  $\nu = IDF$ , and  $\delta = DELTA$ , it determines  $t_0 (= TNIN(P, IDF, DELTA))$ , such that

$$P = \int_{-\infty}^{t_0} \frac{\nu^{v/2} e^{-\delta^2/2}}{\sqrt{\pi} \Gamma(\nu/2) (\nu + x^2)^{(\nu+1)/2}} \sum_{i=0}^{\infty} \Gamma((\nu + i + 1) / 2) \left(\frac{\delta^i}{i!}\right) \left(\frac{2x^2}{\nu + x^2}\right)^{i/2} dx$$

where  $\Gamma(\cdot)$  is the gamma function. The probability that the random variable takes a value less than or equal to  $t_0$  is  $P$ . See  $TNDF$  (page 1146) for an alternative definition in terms of normal and chi-squared random variables. The function  $TNIN$  uses bisection and modified regula falsi to invert the distribution function, which is evaluated using routine  $TNDF$ .

### Example

In this example, we find the 95-th percentage point for a noncentral  $t$  random variable with 6 degrees of freedom and noncentrality parameter 6.

```

      INTEGER      IDF, NOUT
      REAL        DELTA, P, T, TNIN
      EXTERNAL    TNIN, UMACH
C
      CALL UMACH (2, NOUT)
      IDF      = 6
      DELTA    = 6.0
      P        = 0.95
      T        = TNIN(P, IDF, DELTA)
      WRITE (NOUT, 99999) T
C
99999 FORMAT (' The 0.05 noncentral t critical value is ', F6.3,
&           ' .')
C
      END

```

### Output

The 0.05 noncentral t critical value is 11.995.

---

## GCDF/DGCDF (Single/Double precision)

Evaluate a general continuous cumulative distribution function given ordinates of the density.

### Usage

GCDF(*X0*, *IOPT*, *M*, *X*, *F*)

### Arguments

*X0* — Point at which the distribution function is to be evaluated. (Input)

*IOPT* — Indicator of the method of interpolation. (Input)

#### *IOPT* Interpolation Method

- 1 Linear interpolation with equally spaced abscissas.
- 2 Linear interpolation with possibly unequally spaced abscissas.
- 3 A cubic spline is fitted to equally spaced abscissas.
- 4 A cubic spline is fitted to possibly unequally spaced abscissas.

*M* — Number of ordinates of the density supplied. (Input)

*M* must be greater than 1 for linear interpolation (*IOPT* = 1 or 2) and greater than 3 if a curve is fitted through the ordinates (*IOPT* = 3 or 4).

*X* — Array containing the abscissas or the endpoints. (Input)

If *IOPT* = 1 or 3, *X* is of length 2. If *IOPT* = 2 or 4, *X* is of length *M*. For *IOPT* = 1 or 3, *X*(1) contains the lower endpoint of the support of the distribution and *X*(2) is the upper endpoint. For *IOPT* = 2 or 4, *X* contains, in strictly increasing order, the abscissas such that *X*(*I*) corresponds to *F*(*I*).

*F* — Vector of length *M* containing the probability density ordinates corresponding to increasing abscissas. (Input)

If  $IOPT = 1$  or  $3$ , for  $I = 1, 2, \dots, M$ ,  $F(I)$  corresponds to  $x(1) + (I - 1) * (x(2) - x(1))/(M - 1)$ ; otherwise,  $F$  and  $x$  correspond one for one.

**GCDF** — Function value, the probability that a random variable whose density is given in  $F$  takes a value less than or equal to  $x_0$ . (Output)

### Comments

If  $IOPT = 3$ , automatic workspace usage is

GCDF	6 * M units, or
DGCDF	11 * M units.

If  $IOPT = 4$ , automatic workspace usage is

GCDF	5 * M units, or
DGCDF	9 * M units.

Workspace may be explicitly provided, if desired, by the use of  $G4DF/DG4DF$ . The reference is

```
G4DF(P, IOPT, M, X, F, WK, IWK)
```

The arguments in addition to those of  $GCDF$  are

**WK** — Work vector of length  $5 * M$  if  $IOPT = 3$ , and of length  $4 * M$  if  $IOPT = 4$ .

**IWK** — Work vector of length  $M$ .

### Algorithm

Function  $GCDF$  evaluates a continuous distribution function, given ordinates of the probability density function. It requires that the range of the distribution be specified in  $x$ . For distributions with infinite ranges, endpoints must be chosen so that most of the probability content is included. The function  $GCDF$  first fits a curve to the points given in  $x$  and  $F$  with either a piecewise linear interpolant or a  $C^1$  cubic spline interpolant based on a method by Akima (1970). Function  $GCDF$  then determines the area,  $A$ , under the curve. (If the distribution were of finite range and if the fit were exact, this area would be 1.0.) Using the same fitted curve,  $GCDF$  next determines the area up to the point  $x_0 (= x_0)$ . The value returned is the area up to  $x_0$  divided by  $A$ . Because of the scaling by  $A$ , it is not assumed that the integral of the density defined by  $x$  and  $F$  is 1.0. For most distributions, it is likely that better approximations to the distribution function are obtained when  $IOPT$  equals 3 or 4, that is, when a cubic spline is used to approximate the function. It is also likely that better approximations can be obtained when the abscissas are chosen more densely over regions where the density and its derivatives (when they exist) are varying greatly.

### Example

In this example, we evaluate the beta distribution function at the point 0.6. The probability density function of a beta random variable with parameters  $p$  and  $q$  is

$$f(x) = \frac{\Gamma(p+q)}{\Gamma(p)\Gamma(q)} x^{p-1}(1-x)^{q-1} \quad \text{for } 0 \leq x \leq 1$$

where  $\Gamma(\cdot)$  is the gamma function. The density is equal to 0 outside the interval  $[0, 1]$ . We compute a constant multiple (we can ignore the constant gamma functions) of the density at 300 equally spaced points and input this information in  $X$  and  $F$ . Knowing that the probability density of this distribution is very peaked in the vicinity of 0.5, we could perhaps get a better fit by using unequally spaced abscissas, but we will keep it simple. Note that this is the same example as one used in the description of routine `BETDF` (page 1125). The result from `BETDF` would be expected to be more accurate than that from `GCDF` since `BETDF` is designed specifically for this distribution.

```

INTEGER      M
PARAMETER   (M=300)
C
INTEGER      I, IOPT, NOUT
REAL         F(M), GCDF, H, P, PIN1, QIN1, X(2), X0, XI
EXTERNAL     GCDF, UMACH
C
CALL UMACH (2, NOUT)
X0 = 0.6
IOPT = 3
C
C                                     Initializations for a beta(12,12)
C                                     distribution.
PIN1 = 11.0
QIN1 = 11.0
XI = 0.0
H = 1.0/(M-1.0)
X(1) = XI
F(1) = 0.0
XI = XI + H
C
C                                     Compute ordinates of the probability
C                                     density function.
DO 10 I=2, M - 1
    F(I) = XI**PIN1*(1.0-XI)**QIN1
    XI = XI + H
10 CONTINUE
X(2) = 1.0
F(M) = 0.0
P = GCDF(X0, IOPT, M, X, F)
WRITE (NOUT, 99999) P
99999 FORMAT (' The probability that X is less than 0.6 is ', F6.4)
END

```

### Output

The probability that X is less than 0.6 is 0.8364

---

## GCIN/DGCIN (Single/Double precision)

Evaluate the inverse of a general continuous cumulative distribution function given ordinates of the density.

## Usage

GCIN(P, IOPT, M, X, F)

## Arguments

**P** —Probability for which the inverse of the distribution function is to be evaluated. (Input)

**P** must be in the open interval (0.0, 1.0).

**IOPT** — Indicator of the method of interpolation. (Input)

### IOPT Interpolation Method

- 1 Linear interpolation with equally spaced abscissas.
- 2 Linear interpolation with possibly unequally spaced abscissas.
- 3 A cubic spline is fitted to equally spaced abscissas.
- 4 A cubic spline is fitted to possibly unequally spaced abscissas.

**M** —Number of ordinates of the density supplied. (Input)

**M** must be greater than 1 for linear interpolation (**IOPT** = 1 or 2) and greater than 3 if a curve is fitted through the ordinates (**IOPT** = 3 or 4).

**X** —Array containing the abscissas or the endpoints. (Input)

If **IOPT** = 1 or 3, **X** is of length 2. If **IOPT** = 2 or 4, **X** is of length **M**. For **IOPT** = 1 or 3, **X**(1) contains the lower endpoint of the support of the distribution and **X**(2) is the upper endpoint. For **IOPT** = 2 or 4, **X** contains, in strictly increasing order, the abscissas such that **X**(**I**) corresponds to **F**(**I**).

**F** —Vector of length **M** containing the probability density ordinates corresponding to increasing abscissas. (Input)

If **IOPT** = 1 or 3, for **I** = 1, 2, ..., **M**, **F**(**I**) corresponds to  $X(1) + (I - 1) * (X(2) - X(1)) / (M - 1)$ ; otherwise, **F** and **X** correspond one for one.

**GCIN** — Function value. (Output)

The probability that a random variable whose density is given in **F** takes a value less than or equal to **GCIN** is **P**.

## Comments

If **IOPT** = 3, automatic workspace usage is

GCIN 6 \* **M** units, or  
DGCIN 11 \* **M** units.

If **IOPT** = 4, automatic workspace usage is

GCIN 5 \* **M** units, or  
DGCIN 9 \* **M** units.

Workspace may be explicitly provided, if desired, by the use of **G3IN/DG3IN**. The reference is

G3IN(P, IOPT, M, X, F, WK, IWK)

The arguments in addition to those of **GCIN** are

**WK** — Work vector of length  $5 * M$  if  $IOPT = 3$ , and of length  $4 * M$  if  $IOPT = 4$ .

**IWK** — Work vector of length  $M$ .

### Algorithm

Function **GCIN** evaluates the inverse of a continuous distribution function, given ordinates of the probability density function. The range of the distribution must be specified in **x**. For distributions with infinite ranges, endpoints must be chosen so that most of the probability content is included.

The function **GCIN** first fits a curve to the points given in **x** and **F** with either a piecewise linear interpolant or a  $C^1$  cubic spline interpolant based on a method by Akima (1970). Function **GCIN** then determines the area,  $A$ , under the curve. (If the distribution were of finite range and if the fit were exact, this area would be 1.0.) It next finds the maximum abscissa up to which the area is less than  $AP$  and the minimum abscissa up to which the area is greater than  $AP$ . The routine then interpolates for the point corresponding to  $AP$ . Because of the scaling by  $A$ , it is not assumed that the integral of the density defined by **x** and **F** is 1.0.

For most distributions, it is likely that better approximations to the distribution function are obtained when  $IOPT$  equals 3 or 4, that is, when a cubic spline is used to approximate the function. It is also likely that better approximations can be obtained when the abscissas are chosen more densely over regions where the density and its derivatives (when they exist) are varying greatly.

### Example

In this example, we find the 90-th percentage point for a beta random variable with parameters 12 and 12. The probability density function of a beta random variable with parameters  $p$  and  $q$  is

$$f(x) = \frac{\Gamma(p+q)}{\Gamma(p)\Gamma(q)} x^{p-1} (1-x)^{q-1} \quad \text{for } 0 \leq x \leq 1$$

where  $\Gamma(\cdot)$  is the gamma function. The density is equal to 0 outside the interval  $[0, 1]$ . With  $p = q$ , this is a symmetric distribution. Knowing that the probability density of this distribution is very peaked in the vicinity of 0.5, we could perhaps get a better fit by using unequally spaced abscissas, but we will keep it simple and use 300 equally spaced points. Note that this is the same example that is used in the description of routine **BETIN** (page 1127). The result from **BETIN** would be expected to be more accurate than that from **GCIN** since **BETIN** is designed specifically for this distribution.

```
C
INTEGER      M
PARAMETER   (M=300)

INTEGER      I, IOPT, NOUT
REAL         BETA, C, F(M), GCIN, H, P, PIN, PIN1, QIN, QIN1,
&            X(2), X0, XI
EXTERNAL    BETA, GCIN, UMACH
```

```

C
CALL UMACH (2, NOUT)
P = 0.9
IOPT = 3
C
C
C Initializations for a beta(12,12)
C distribution.
PIN = 12.0
QIN = 12.0
PIN1 = PIN - 1.0
QIN1 = QIN - 1.0
C = 1.0/BETA(PIN,QIN)
XI = 0.0
H = 1.0/(M-1.0)
X(1) = XI
F(1) = 0.0
XI = XI + H
C
C Compute ordinates of the probability
C density function.
DO 10 I=2, M - 1
    F(I) = C*XI**PIN1*(1.0-XI)**QIN1
    XI = XI + H
10 CONTINUE
X(2) = 1.0
F(M) = 0.0
X0 = GCIN(P, IOPT, M, X, F)
WRITE (NOUT, 99999) X0
99999 FORMAT (' X is less than ', F6.4, ' with probability 0.9.')
END

```

### Output

X is less than 0.6304 with probability 0.9.

---

## GFNIN/DGFNIN (Single/Double precision)

Evaluate the inverse of a general continuous cumulative distribution function given in a subprogram.

### Usage

GFNIN(F, P, EPS, GUESS)

### Arguments

**F** — User-supplied FUNCTION to be inverted. F must be continuous and strictly monotone. The form is F(X), where

X — The argument to the function. (Input)

F — The value of the function at X. (Output)

F must be declared EXTERNAL in the calling program.

**P** — The point at which the inverse of F is desired. (Input)

**EPS** — Convergence criterion. (Input)

When the relative change in GFNIN from one iteration to the next is less than



EPS, convergence is assumed. A common value for EPS is 0.0001. Another common value is 100 times the machine epsilon.

**GUESS** — An initial estimate of the inverse of  $F$  at  $P$ . (Input)

**GFNIN** — The inverse of the function  $F$  at the point  $P$ . (Output)  
 $F(\text{GFNIN})$  is “close” to  $P$ .

### Comments

1. Informational errors

Type	Code	
4	1	After 100 attempts, a bound for the inverse cannot be determined. Try again with a different initial estimate.
4	2	No unique inverse exists.
4	3	Over 100 iterations have occurred without convergence. Convergence is assumed.
2. The function to be inverted need not be a distribution function, it can be any continuous, monotonic function.

### Algorithm

Function **GFNIN** evaluates the inverse of a continuous, strictly monotone function. Its most obvious use is in evaluating inverses of continuous distribution functions that can be defined by a FORTRAN function. If the distribution function cannot be specified in a FORTRAN function, but the density function can be evaluated at a number of points, then routine **GCIN** (page 1152) can be used.

Function **GFNIN** uses regula falsi and/or bisection, possibly with the Illinois modification (see Dahlquist and Bjorck 1974). A maximum of 100 iterations are performed.

### Example

In this example, we find the 99–th percentage point for an  $F$  random variable with 1 and 7 degrees of freedom. (This problem could be solved easily using routine **FIN** (page 1139). Compare the example for **FIN**). The function to be inverted is the  $F$  distribution function, for which we use routine **FDF** (page 1137). Since **FDF** requires the degrees of freedom in addition to the point at which the function is evaluated, we write another function **F** that receives the degrees of freedom via a common block and then calls **FDF**. The starting point (initial guess) is taken as two standard deviations above the mean (since this would be a good guess for a normal distribution). It is not necessary to supply such a good guess. In this particular case, an initial estimate of 1.0, for example, yields the same answer in essentially the same number of iterations. (In fact, since the  $F$  distribution is skewed, the initial guess, 7.0, is really not that close to the final answer.)

```

      INTEGER      NOUT
      REAL         DFD, DFN, EPS, F, F0, GFNIN, GUESS, P, SQRT
      COMMON      /FCOM/ DFN, DFD
      INTRINSIC   SQRT
      EXTERNAL    F, GFNIN, UMACH
C
      CALL UMACH (2, NOUT)
      P = 0.99
      DFN = 1.0
      DFD = 7.0
C
      Compute GUESS as two standard
      deviations above the mean.
C
      GUESS = DFD/(DFD-2.0) + 2.0*SQRT(2.0*DFD*DFD*(DFN+DFD-2.0)/(DFN*
&      (DFD-2.0)**2*(DFD-4.0)))
      EPS = 0.00001
      F0 = GFNIN(F,P,EPS,GUESS)
      WRITE (NOUT,99999) F0
99999 FORMAT (' The F(1,7) 0.01 critical value is ', F6.3)
      END
C
      REAL FUNCTION F (X)
      REAL      X
C
      REAL      DFD, DFN, FDF
      COMMON    /FCOM/ DFN, DFD
      EXTERNAL  FDF
C
      F = FDF(X,DFN,DFD)
      RETURN
      END

```

### Output

The F(1,7) 0.01 critical value is 12.246

# Chapter 18: Random Number Generation

---

## Routines

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---

## Usage Notes

In the following discussions, the phrases “random numbers,” “random deviates,” “deviates,” and “variates” are used interchangeably. The phrase “pseudorandom” is sometimes used to emphasize that the numbers generated are not really “random” since they result from a deterministic process. The usefulness of pseudorandom numbers derives from the similarity, in a statistical sense, of samples of the pseudorandom numbers to samples of observations from the specified distributions. In short, while the pseudorandom numbers are completely deterministic and repeatable, they *simulate* the realizations of independent and identically distributed random variables.

## The Basic Uniform Generators

The random number generators in this chapter use either a multiplicative congruential method or a generalized feedback shift register (GFSR) method. The selection of the type of generator is made by calling the routine `RNOPT` (page 1165). If no selection is made explicitly, a multiplicative generator (with multiplier 16807) is used. Whatever distribution is being simulated, uniform (0, 1) numbers are first generated and then transformed if necessary. The generation of the uniform (0, 1) numbers is done by the routine `RNUN` (page 1171), or by its function analog `RNUNF`. These routines are *portable* in the sense that, given the same seed and for a given type of generator, they produce the same sequence in all computer/compiler environments. There are many other issues that must be considered in developing programs for the methods described below (see Gentle 1981 and 1990).

## The Multiplicative Congruential Generators

The form of the multiplicative congruential generators is

$$x_i \equiv cx_{i-1} \pmod{2^{31} - 1}$$

Each  $x_i$  is then scaled into the unit interval (0, 1). If the multiplier,  $c$ , is a primitive root modulo  $2^{31} - 1$  (which is a prime), then the generator will have maximal period of  $2^{31} - 2$ . There are several other considerations, however. The lattice structure induced by congruential generators (see Marsaglia 1968) can be assessed by the lattice test of Marsaglia (1972) or the spectral test of Coveyou and MacPherson (1967) (see also Knuth 1981, pages 89–113). Also, empirical studies, such as by Fishman and Moore (1982 and 1986), indicate that different values of multipliers, all of which perform well under the lattice test and the spectral test, may yield quite different performances where the criterion is similarity of samples generated to samples from a true uniform distribution.

There are three possible choices for  $c$  in the IMSL generators: 16807 (which is  $7^5$ ), 397204094 (which is  $2 \cdot 7^2 \cdot 4053103$ ), and 950706376 (which is  $2^3 \cdot 118838297$ ). The selection is made by the routine `RNOPT` (page 1165). The choice of 16807 will result in the fastest execution time (see Gentle 1981), but Fishman and Moore's studies would seem to indicate that the performance of 950706376 is best among these three choices. If no selection is made explicitly, the routines use the multiplier 16807, which has been in use for some time (Lewis, Goodman, and Miller 1969). It is the "minimal standard generator" discussed by Park and Miller (1988).

The user can also select a shuffled version of the multiplicative congruential generators using `RNOPT` (page 1165). The shuffled generators use a scheme due to Learmonth and Lewis (1973a). In this scheme, a table is filled with the first 128 uniform (0, 1) numbers resulting from the simple multiplicative congruential generator. Then, for each  $x_i$  from the simple generator, the low-

order bits of  $x_j$  are used to select a random integer,  $j$ , from 1 to 128. The  $j$ -th entry in the table is then delivered as the random number; and  $x_j$ , after being scaled into the unit interval, is inserted into the  $j$ -th position in the table.

## The Generalized Feedback Shift Register Generator

The GFSR generator uses the recursion  $X_t = X_{t-1563} \oplus X_{t-96}$ . This generator, which is different from earlier GFSR generators, was proposed by Fushimi (1990), who discusses the theory behind the generator and reports on several empirical tests of it. Background discussions on this type of generator can be found in Kennedy and Gentle (1980), pages 150–162.

## Setting the Seed

The seed of the generator can be set in `RNSET` (page 1167) and can be retrieved by `RNGET` (page 1167). Prior to invoking any generator in this chapter, the user can call `RNSET` to initialize the seed, which is an integer variable taking a value between 1 and 2147483646. If it is not initialized by `RNSET`, a random seed is obtained from the system clock. Once it is initialized, the seed need not be set again. The seed is updated and passed from one routine to another by means of a named `COMMON` block, `R2NCOM`.

If the user wishes to restart a simulation, `RNGET` can be used to obtain the final seed value of one run to be used as the starting value in a subsequent run. Also, if two random number streams are desired in one run, `RNSET` and `RNGET` can be used before and after the invocations of the generators in each stream. If a shuffled generator or the GFSR generator is used, in addition to resetting the seed, the user must also reset some values in a table. For the shuffled generators, this is done using the routines `RNGES` (page 1167) and `RNSES` (page 1167); and for the GFSR generator, the table is retrieved and set by the routines `RNGEF` (page 1167) and `RNSEF` (page 1167). The tables for the shuffled generators are separate for single and double precision; so, if precisions are mixed in a program, it is necessary to manage each precision separately for the shuffled generators.

## Timing Considerations

The generation of the uniform (0,1) numbers is done by the routine `RNUN` (page 1171) or by its function analog `RNUNF` (page 1172). The particular generator selected in `RNOPT` (page 1165), that is, the value of the multiplier and whether shuffling is done or whether the GFSR generator is used, affects the speed of `RNUN` and `RNUNF`. The smaller multiplier (16807, selected by `IOPT = 1`) is faster than the other multipliers. The multiplicative congruential generators that do not shuffle are faster than the ones that do. The GFSR generator is roughly as fast as the fastest multiplicative congruential generator, but the initialization for it (required only on the first invocation) takes longer than the

generation of thousands of uniform random numbers. Precise statements of relative speeds depend on the computing system.

Whether `RNUN` or `RNUNF` is used also has an effect on the speed due to the overhead in invoking an external routine, or due to the program's inability to optimize computations by holding some operands in registers. This effect, of course, may be different in different environments. On an array processor or other computers with pipelined instructions, `RNUN` is likely to be considerably faster than `RNUNF` when several random numbers are to be generated at one time. In the case of array processors, the multiplicative congruential generators in `RNUN` are coded to generate subsequences in larger blocks (see Gentle 1990).

## Use of Customized Uniform Generators

The basic uniform (0, 1) generators `RNUN` or `RNUNF` are used by all other routines in this chapter. If, for some reason, the user would prefer a different basic uniform generator, routines named “`RNUN`” and “`RNUNF`” can be written so that they include the named `COMMON`, through which the seed is passed, and that calls the user's custom generator. The named `COMMON` is

```
COMMON /R2NCOM/ D2P31A, DSEED, D2P31R, DWK, DINTTB, INDCTR,  
&      INTTB, WK, ICEED, IDSTFS, INTFS, ISRCFS, S2P31R, IWFS  
DOUBLE PRECISION D2P31A, D2P31R, DSEED, DWK(128)  
REAL      S2P31R, WK(128)  
INTEGER  ICEED, IDSTFS, INDCTR, ISRCFS, IWFS(1563)  
LOGICAL  DINTTB, INTTB, INTFS  
SAVE    /R2NCOM/
```

The user's “`RNUN`” and “`RNUNF`” can pass the seed through any of the variables, but the routines `RNSET` (page 1167) and `RNGET` (page 1167) expect the seed to be in `ICEED`. (The user should not expect to use any utility routines other than `RNSET` and `RNGET` if customized versions of `RNUN` or `RNUNF` are used.) The double precision versions of the nonuniform generators, such as `DRNBET` (page 1191) and `DRNGAM` (page 1198), use the double precision versions of the uniform generators, `DRNUN` (page 1171) and `DRNUNF` (page 1172), so to use the double precision nonuniform generators with customized uniform generators, the user would supply routines to replace `DRNUN` and `DRNUNF`.

## Distributions Other than the Uniform

The nonuniform generators use a variety of transformation procedures. All of the transformations used are exact (mathematically). The most straightforward transformation is the *inverse CDF technique*, but it is often less efficient than others involving *acceptance/rejection* and *mixtures*. See Kennedy and Gentle (1980) for discussion of these and other techniques.

Many of the nonuniform generators in this chapter use different algorithms depending on the values of the parameters of the distributions. This is particularly true of the generators for discrete distributions. Schmeiser (1983) gives an overview of techniques for generating deviates from discrete distributions.

Although, as noted above, the uniform generators yield the same sequences on different computers, because of rounding, the nonuniform generators that use acceptance/rejection may occasionally produce different sequences on different computer/compiler environments.

Although the generators for nonuniform distributions use fast algorithms, if a very large number of deviates from a fixed distribution are to be generated, it might be worthwhile to consider a table sampling method, as implemented in the routines `RNGDA` (page 1174), `RNGDS` (page 1177), `RNGDT` (page 1181), `RNGCS` (page 1200), and `RNGCT` (page 1202). After an initialization stage, which may take some time, the actual generation may proceed very fast.

## Order Statistics and Antithetic Variates

For those generators, such as `RNCHY` (page 1194) and `RNNOR` (page 1208), that use the inverse CDF technique, it is possible to generate any set of order statistics directly by use of a customized uniform generator, as discussed above, by generating order statistics in a custom “`RNUN`” or “`RNUNF`”. In some routines that employ an inverse CDF technique, such as `RNEXP` (page 1196) and `RNWIB` (page 1214), instead of directly using the uniform (0, 1) deviate  $u$  from `RNUN` (page 1171), the uniform (0, 1) deviate  $1 - u$  is used. In such routines the  $i$ -th order statistic from the uniform will yield the  $(n + 1 - i)$ -th order statistic from the nonuniform distribution.

A similar technique can be used to get antithetic variates. For each uniform deviate  $u$ , a second deviate  $1 - u$  could be produced by a custom “`RNUN`” or “`RNUNF`”. As with order statistics, this technique would only be reasonable for routines that use the inverse CDF technique.

## Tests

Extensive empirical tests of some of the uniform random number generators available in `RNUN` (page 1171) and `RNUNF` (page 1172) are reported by Fishman and Moore (1982 and 1986). Results of tests on the generator using the multiplier 16807 with and without shuffling are reported by Learmonth and Lewis (1973b). If the user wishes to perform additional tests, the routines in Chapter 7, “Tests of Goodness of Fit and Randomness,” may be of use. The user may also wish to compute some basic statistics or to make some plots of the output of the random number generator being used. The routines in Chapter 1, “Basic Statistics,” and Chapter 16, “Line Printer Graphics,” may be used for this purpose. Often in Monte Carlo applications, it is appropriate to construct an ad hoc test that is sensitive to departures that are important in the given application. For example, in using Monte Carlo methods to evaluate a one-dimensional integral, autocorrelations of order one may not be harmful, but they may be disastrous in evaluating a two-dimensional integral. Although generally the routines in this chapter for generating random deviates from nonuniform distributions use exact methods, and, hence, their quality depends almost solely on the quality of the underlying uniform generator, it is often advisable to



employ an ad hoc test of goodness of fit for the transformations that are to be applied to the deviates from the nonuniform generator.

### Other Notes on Usage

The generators for continuous distributions are available in both single and double precision versions. This is merely for the convenience of the user; the double precision versions should not be considered more “accurate,” except possibly for the multivariate distributions.

The names of all of the routines for random number generation begin with “RN” for single precision and “DRN” for double precision. In most routines, the first argument, *NR*, is the number of variates to generate; and the last variable, either *R* or *IR*, is the vector of random variates.

Error handling and workspace allocation in the routines for random number generation are done somewhat differently than in most other IMSL routines. In general, there is less error checking than in other routines since there is more emphasis on speed in the random number generation routines. Simple checks for gross errors are made in all routines; and the routines for setup do complete checking since it is assumed that they would not be called frequently. Some routines, such as those that construct tables or interpolate from tables, require that the user explicitly provide some work arrays.

---

## Random Number Generation Utility Routines

All of the random number generators in this chapter depend on the generation of uniform (0, 1) numbers, which is done by the routine *RNUN* (page 1171), or by its function analog *RNUNF* (page 1172). These basic generators use either a multiplicative congruential method or a generalized feedback shift register (GFSR) method to yield a subsequence of a fixed cyclic sequence. The beginning of the subsequence is determined by the seed.

The utility routines for the random number generators allow the user to select the type of the generator (or to determine the type of the generator being used) and to set or retrieve the seed.

### Selection of the Type of the Generator

For generating uniform (0, 1) random numbers either a multiplicative congruential or a GFSR method is used. In the multiplicative congruential method one of three different multipliers, with or without shuffling, can be chosen. The selection of the type of generator is made by calling the routine *RNOPT*, choosing one of seven different options. The usage is

```
CALL RNOPT ( IOPT )
```

The argument is

*IOPT* — The indicator of the generator. (Input)

### **IOPT Generator**

- 1 Congruential, with multiplier 16807 is used.
- 2 Congruential, with multiplier 16807 is used with shuffling.
- 3 Congruential, with multiplier 397204094 is used.
- 4 Congruential, with multiplier 397204094 is used with shuffling.
- 5 Congruential, with multiplier 950706376 is used.
- 6 Congruential, with multiplier 950706376 is used with shuffling.
- 7 GFSR, with the recursion  $X_t = X_{t-1563} \oplus X_{t-96}$  is used.

If no selection is made explicitly, a multiplicative generator (with multiplier 16807) is used (equivalent to `IOPT = 1`).

The type of generator being used can be determined by calling the routine `RNOPG`. The usage is

```
CALL RNOPG ( IOPT )
```

`IOPT` is an output variable in `RNOPG`.

### **Setting the Seed**

Before using any of the random number generators, the generator must be initialized by selecting a *seed*, or starting value. The user does not have to do this, but it can be done by calling the routine `RNSET`. If the user does not select a seed, one is generated using the system clock. A seed needs to be selected only once in a program unless there is some desire to maintain two separate streams of random numbers. The usage is

```
CALL RNSET ( ISEED )
```

The argument is

***ISEED*** — The seed of the random number generator. (Input)

`ISEED` must be in the range (0, 2147483646). If `ISEED` is zero (or if `RNSET` is not called before the generation of random numbers begins), a value is computed using the system clock; and, hence, the results of programs using the IMSL random number generators will be different at different times.

### **Stopping and Restarting Simulations and Controlling More Than One Stream of Random Numbers**

For most purposes, even if several simulations are being run in the same program or if the simulation is being conducted in blocks, it is best to use the sequence of uniform random deviates just as produced by `RNUN` (page 1171) or `RNUNF` (page 1172) without concern for from where in the underlying cyclic sequence the numbers are coming.

If, however, the simulations are being conducted incrementally or if simulations are being run in parallel, it may be necessary to exercise more control over the sequence. The routines that are used in stopping and restarting simulations come in pairs, one to get the current value and one to set the value. The

argument for each pair is the same within the pair; it is output in one case and input in the other. (*RNSET* is an exception since it is often used at the beginning of a simulation before any seed is ever set.) If a nonshuffled form of the multiplicative congruential generators is used (that is *IOPT* in *RNOPT*, page 1165, is 1, 3, or 5), the only thing that must be controlled is the seed, so in this case the only routines needed are

***RNGET*** — Retrieve the current value of the seed

***RNSET*** — Initialize the seed used in the generators

The usages are

```
CALL RNGET ( ISEED ) ( ISEED is output.)
```

```
CALL RNSET ( ISEED ) ( ISEED is input.)
```

*ISEED* is an integer in the range 1 to 2147483646 (except, as noted above, it can be input to *RNSET* as 0 to indicate that the system clock is to be used to generate a seed).

If a shuffled generator or the GFSR generator is used, in addition to controlling the seed as described above, another array must be maintained if the user wishes to stop and restart the simulation. It is a floating-point array for the shuffled generators and an integer array for the GFSR generator. The routines are

***RNGES*** — Retrieve the current table used in the shuffled generators

***RNSES*** — Initialize the table used in the shuffled generators

***RNGEF*** — Retrieve the current table used in the GFSR generator

***RNSEF*** — Initialize the table used in the GFSR generator

There are different tables used in the single and double precision versions of the shuffled generators, so *RNGES* and *RNSES* have double precision counterparts, *DRNGES* and *DRNSES*, respectively.

The usages are

```
CALL RNGES ( TABLE ) ( TABLE is output.)
```

```
CALL RNSES ( TABLE ) ( TABLE is input.)
```

```
CALL RNGEF ( IARRAY ) ( IARRAY is output.)
```

```
CALL RNSEF ( IARRAY ) ( IARRAY is input.)
```

The arguments are

***TABLE*** — Array of length 128 used in the shuffled generators.

***IARRAY*** — Array of length 1565 used in the GFSR generators.

The values in both *TABLE* and *IARRAY* are initialized by the IMSL random number generators. The values are all positive in both arrays except if the user wishes to reinitialize the array, in which case the first element of the array is input as a nonpositive value. (Usually, one should avoid reinitializing these arrays, but it might be necessary sometimes in restarting a simulation.) If the first element of *TABLE* or *IARRAY* is set to a nonpositive value on the call to *RNSES* or *RNSEF*, on the next invocation of a routine to generate random

numbers using shuffling (if *RNSES*) or a GFSR method (if *RNSEF*), the appropriate array will be reinitialized.

In addition to controlling separate streams of random numbers, sometimes it is desirable to insure from the beginning that two streams do not overlap. This can be done with the congruential generators that do not do shuffling by using *RNISR* to get a seed that will generate random numbers beginning 100,000 numbers farther along.

The usage is

```
CALL RNISR ( ISEED1, ISEED2 )
```

The arguments are

***ISEED1*** — The seed that yields the first stream. (Input)

***ISEED2*** — The seed that yields a stream beginning 100,000 numbers beyond the stream that begins with *ISEED1*. (Output)

Given a seed, *ISEED1*, *RNISR* determines another seed, *ISEED2*, such that if one of the IMSL multiplicative congruential generators, using no shuffling, went through 100,000 generations starting with *ISEED1*, the next number in that sequence would be the first number in the sequence that begins with the seed *ISEED2*. This can be described more simply by stating that *RN1* and *RN2* in the following sequence of FORTRAN are assigned the same values.

```
CALL RNISR( ISEED1, ISEED2 )
CALL RNSET( ISEED1 )
DO 10 I = 1, 100000
    RN1 = RNUNF( )
10 CONTINUE
RN1 = RNUNF( )
CALL RNSET( ISEED2 )
RN2 = RNUNF( )
```

To obtain seeds that generate sequences with beginning values separated by 200,000 numbers, call *RNISR* twice:

```
CALL RNISR( ISEED1, ISEED2 )
CALL RNISR( ISEED2, ISEED2 )
```

Note that *RNISR* works only when a multiplicative congruential generator without shuffling is used. This means that either the routine *RNOPT* (page 1165) has not been called at all or that it has been last called with *IOPT* taking a value of 1, 3, or 5.

For many of the IMSL generators for nonuniform distributions that do not use the inverse CDF method, the distance between the sequences generated starting with *ISEED1* and starting with *ISEED2* may be less than 100,000. This is because the nonuniform generators that use other techniques may require more than one uniform deviate for each output deviate.

The reason that one may want two seeds that generate sequences a known distance apart is for blocking Monte Carlo experiments or for running parallel streams.

## Example 1

### Selecting the Type of Generator and Stopping and Restarting the Simulations

In this example, three separate simulation streams are used, each with a different form of the generator. Each stream is stopped and restarted. (Although this example is obviously an artificial one, there may be reasons for maintaining separate streams and stopping and restarting them because of the nature of the usage of the random numbers coming from the separate streams.)

```
INTEGER      I, IARRAY(1565), ISEED1, ISEED2, ISEED7, NOUT, NR
REAL         R(5), TABLE(128)
EXTERNAL     RNGEF, RNGES, RNGET, RNOPT, RNSEF, RNSES, RNSET,
&           RNUN, UMACH
C
CALL UMACH (2, NOUT)
NR      = 5
ISEED1 = 123457
ISEED2 = 123457
ISEED7 = 123457
C
C                               Begin first stream, IOPT = 1 (by
C                               default)
CALL RNSET (ISEED1)
CALL RNUN (NR, R)
CALL RNGET (ISEED1)
WRITE (NOUT,99997) (R(I),I=1,NR), ISEED1
C                               Begin second stream, IOPT = 2
CALL RNOPT (2)
CALL RNSET (ISEED2)
CALL RNUN (NR, R)
CALL RNGET (ISEED2)
CALL RNGES (TABLE)
WRITE (NOUT,99998) (R(I),I=1,NR), ISEED2
C                               Begin third stream, IOPT = 7
CALL RNOPT (7)
CALL RNSET (ISEED7)
CALL RNUN (NR, R)
CALL RNGET (ISEED7)
CALL RNGEF (IARRAY)
WRITE (NOUT,99999) (R(I),I=1,NR), ISEED7
C                               Reinitialize seed
C                               Resume first stream
CALL RNOPT (1)
CALL RNSET (ISEED1)
CALL RNUN (NR, R)
CALL RNGET (ISEED1)
WRITE (NOUT,99997) (R(I),I=1,NR), ISEED1
C                               Reinitialize seed and table for
C                               shuffling
C                               Resume second stream
CALL RNOPT (2)
CALL RNSET (ISEED2)
CALL RNSES (TABLE)
CALL RNUN (NR, R)
CALL RNGET (ISEED2)
WRITE (NOUT,99998) (R(I),I=1,NR), ISEED2
```

```

C                               Reinitialize seed and table for GFSR
C                               Resume third stream
      CALL RNOPT (7)
      CALL RNSET (ISEED7)
      CALL RNSEF (IARRAY)
      CALL RNUN (NR, R)
      CALL RNGET (ISEED7)
      WRITE (NOUT,99999) (R(I),I=1,NR), ISEED7
C
99997 FORMAT (/, ' First stream ', 5F8.4, /, ' Output seed = ',
&           I11)
99998 FORMAT (/, ' Second stream ', 5F8.4, /, ' Output seed = ',
&           I11)
99999 FORMAT (/, ' Third stream ', 5F8.4, /, ' Output seed = ',
&           I11)
C
      END

```

### Output

```

First stream  0.9662  0.2607  0.7663  0.5693  0.8448
Output seed =    123457
Second stream 0.7095  0.1861  0.4794  0.6038  0.3790
Output seed = 1435003364
Third stream  0.7095  0.1861  0.4794  0.6038  0.3790
Output seed =    123457
First stream  0.9662  0.2607  0.7663  0.5693  0.8448
Output seed =    123457
Second stream 0.8662  0.4786  0.2062  0.2092  0.9154
Output seed = 1435003364
Third stream  0.8662  0.4786  0.2062  0.2092  0.9154
Output seed =    123457

```

### Example 2

#### Determining Seeds for Separate Streams

In this example, RNISD (page 1168) is used to determine seeds for 4 separate streams, each 200,000 numbers apart, for a multiplicative congruential generator without shuffling. (Since RNOPT is not invoked to select a generator, the multiplier is 16807.) To get each seed requires two invocations of RNISD. All of the streams are non-overlapping, since the period of the underlying generator is 2,147,483,646.

```

      INTEGER    ISEED1, ISEED2, ISEED3, ISEED4, NOUT
      EXTERNAL  RNISD, UMACH
C
      CALL UMACH (2, NOUT)
      ISEED1 = 123457
      CALL RNISD (ISEED1, ISEED2)
      CALL RNISD (ISEED2, ISEED2)
      CALL RNISD (ISEED2, ISEED3)
      CALL RNISD (ISEED3, ISEED3)
      CALL RNISD (ISEED3, ISEED4)
      CALL RNISD (ISEED4, ISEED4)
      WRITE (NOUT,99999) ISEED1, ISEED2, ISEED3, ISEED4
C

```

```
99999 FORMAT ( ' Seeds for four separate streams: ', /, ' ', 4I11)
C
      END
```

### Output

```
Seeds for four separate streams:
123457 2016130173 85016329 979156171
```

---

## RNUN/DRNUN (Single/Double precision)

Generate pseudorandom numbers from a uniform (0, 1) distribution.

### Usage

```
CALL RNUN (NR, R)
```

### Arguments

*NR* — Number of random numbers to generate. (Input)

*R* — Vector of length *NR* containing the random uniform (0, 1) deviates.  
(Output)

### Comments

The routine *RNSET* (page 1167) can be used to initialize the seed of the random number generator. The routine *RNOPT* (page 1165) can be used to select the form of the generator.

### Algorithm

Routine *RNUN* generates pseudorandom numbers from a uniform (0, 1) distribution using one of the algorithms described in the introduction to the chapter on random number generation. The algorithm used is determined by *RNOPT* (page 1165). The values returned in *R* by *RNUN* are positive and less than 1.0. Values in *R* may be smaller than the smallest relative spacing, however. Hence, it may be the case that some value  $R(i)$  is such that  $1.0 - R(i) = 1.0$ .

Deviates from the distribution with uniform density over the interval (A, B) can be obtained by scaling the output from *RNUN*. The following statements (in single precision) would yield random deviates from a uniform (A, B) distribution:

```
CALL RNUN (NR, R)
CALL SSCAL (NR, B-A, R, 1)
CALL SADD (NR, A, R, 1)
```

### Example

In this example, *RNUN* is used to generate five pseudorandom uniform numbers. Since *RNOPT* (page 1165) is not called, the generator used is a simple multiplicative congruential one with a multiplier of 16807.

```

      INTEGER      ISEED, NOUT, NR
      REAL         R(5)
      EXTERNAL    RNSET, RNUN, UMACH
C
      CALL UMACH (2, NOUT)
      NR      = 5
      ISEED = 123457
      CALL RNSET (ISEED)
      CALL RNUN (NR, R)
      WRITE (NOUT,99999) R
99999 FORMAT ('          Uniform random deviates: ', 5F8.4)
      END

```

### Output

```
Uniform random deviates:   .9662   .2607   .7663   .5693   .8448
```

---

## RNUNF/DRNUNF (Single/Double precision)

Generate a pseudorandom number from a uniform (0, 1) distribution.

### Usage

RNUNF( )

### Argument

**RNUNF** — Function value, a random uniform (0, 1) deviate. (Output)

### Comments

1. Routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.
2. This function has a side effect: it changes the value of the seed, which is passed through a common block.

### Algorithm

Routine RNUNF is the function form of RNUN (page 1171). The routine RNUNF generates pseudorandom numbers from a uniform (0, 1) distribution. The algorithm used is determined by RNOPT (page 1165). The values returned by RNUNF are positive and less than 1.0.

If several uniform deviates are needed, it may be more efficient to obtain them all at once by a call to RNUN rather than by several references to RNUNF.

### Example

In this example, RNUNF is used to generate five pseudorandom uniform numbers. Since RNOPT (page 1165) is not called, the generator used is a simple multiplicative congruential one with a multiplier of 16807.



```

      INTEGER    I, ISEED, NOUT
      REAL       R(5), RNUNF
      EXTERNAL   RNSET, RNUNF, UMACH
C
      CALL UMACH (2, NOUT)
      ISEED = 123457
      CALL RNSET (ISEED)
      DO 10 I=1, 5
         R(I) = RNUNF()
10 CONTINUE
      WRITE (NOUT,99999) R
99999 FORMAT ('          Uniform random deviates: ', 5F8.4)
      END

```

### Output

```
Uniform random deviates:   0.9662  0.2607  0.7663  0.5693  0.8448
```

## RNBIN

Generate pseudorandom numbers from a binomial distribution.

### Usage

```
CALL RNBIN (NR, N, P, IR)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**N** — Number of Bernoulli trials. (Input)

**P** — Probability of success on each trial. (Input)

**P** must be greater than 0.0 and less than 1.0.

**IR** — Vector of length NR containing the random binomial deviates. (Output)

### Comments

The IMSL routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNBIN generates pseudorandom numbers from a binomial distribution with parameters N and P. N and P must be positive, and P must be less than 1. The probability function (with  $n = N$  and  $p = P$ ) is

$$f(x) = \binom{n}{x} p^x (1-p)^{n-x}$$

for  $x = 0, 1, 2, \dots, n$ .

The algorithm used depends on the values of  $n$  and  $p$ . If  $np < 10$  or if  $p$  is less than a machine epsilon (AMACH(4) (Reference Material)), the inverse CDF technique is used; otherwise, the BTPE algorithm of Kachitvichyanukul and Schmeiser (see Kachitvichyanukul 1982) is used. This is an acceptance/rejection method using a composition of four regions. (TPE = Triangle, Parallelogram, Exponential, left and right.)

### Example

In this example, RNBIN is used to generate five pseudorandom binomial variates with parameters 20 and 0.5.

```

C      INTEGER      NR
      PARAMETER    (NR=5)

C      INTEGER      IR(NR), ISEED, N, NOUT
      REAL          P
      EXTERNAL      RNBIN, RNSET, UMACH

C      CALL UMACH (2, NOUT)
      N          = 20
      P          = 0.5
      ISEED = 123457
      CALL RNSET (ISEED)
      CALL RNBIN (NR, N, P, IR)
      WRITE (NOUT,99999) IR
99999 FORMAT (' Binomial (20, 0.5) random deviates: ', 5I4)
      END

```

### Output

```
Binomial (20, 0.5) random deviates:   14   9  12  10  12
```

---

## RNGDA/DRNGDA (Single/Double precision)

Generate pseudorandom numbers from a general discrete distribution using an alias method.

### Usage

```
CALL RNGDA (NR, IOPT, IMIN, NMASS, PROBS, IWK, WK, IR)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**IOPT** — Indicator of whether the alias vectors are to be initialized. (Input)

#### IOPT Action

0 The alias vectors are to be initialized using the probabilities in PROBS. IOPT is set to 0 on the first call to RNGDA.

1 The alias vectors IWK and WK are used but PROBS is not used.

**IMIN** — Smallest value the random deviate can assume. (Input)  
This is the value corresponding to the probability in PROBS(1).

**NMASS** — Number of mass points in the discrete distribution. (Input)

**PROBS** — Vector of length NMASS containing probabilities associated with the individual mass points. (Input)  
The elements of PROBS must be nonnegative and must sum to 1.0.

**IWK** — Index vector of length NMASS. (Input, if IOPT = 1; output, if IOPT = 0)  
IWK is a work vector.

**WK** — Index vector of length NMASS. (Input, if IOPT = 1; output, if IOPT = 0)  
WK is a work vector.

**IR** — Vector of length NR containing the random discrete deviates. (Output)

### Comments

1. In the interest of efficiency, this routine does only limited error checking when IOPT = 1.
2. The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNGDA generates pseudorandom numbers from a discrete distribution with probability function given in the vector PROBS; that is

$$\Pr(X = i) = p_j$$

for  $i = i_{\min}, i_{\min} + 1, \dots, i_{\min} + n_m - 1$  where  $j = i - i_{\min} + 1$ ,  $p_j = \text{PROBS}(j)$ ,  $i_{\min} = \text{IMIN}$ , and  $n_m = \text{NMASS}$ .

The algorithm is the *alias* method, due to Walker (1974), with modifications suggested by Kronmal and Peterson (1979). The method involves a setup phase, in which the vectors IWK and WK are filled. After the vectors are filled, the generation phase is very fast.

### Example 1

In this example, RNGDA is used to generate five pseudorandom variates from the discrete distribution:

$$\Pr(X = 1) = .05$$

$$\Pr(X = 2) = .45$$

$$\Pr(X = 3) = .31$$

$$\Pr(X = 4) = .04$$

$$\Pr(X = 5) = .15$$

When RNGDA is called the first time, IOPT is input as 0. This causes the work arrays to be initialized. In the next call, IOPT is 1, so the setup phase is bypassed.

```

INTEGER      NMASS, NR
PARAMETER   (NMASS=5, NR=5)
C
INTEGER      IMIN, IOPT, IR(NR), ISEED, IWK(NMASS), NOUT
REAL         PROBS(NMASS), WK(NMASS)
EXTERNAL     RNGDA, RNSET, UMACH
C
CALL UMACH (2, NOUT)
IMIN        = 1
PROBS(1)    = 0.05
PROBS(2)    = 0.45
PROBS(3)    = 0.31
PROBS(4)    = 0.04
PROBS(5)    = 0.15
IOPT        = 0
ISEED       = 123457
CALL RNSET (ISEED)
CALL RNGDA (NR, IOPT, IMIN, NMASS, PROBS, IWK, WK, IR)
WRITE (NOUT,99998) IR
99998 FORMAT ('          Random deviates: ', 5I4)
IOPT = 1
CALL RNGDA (NR, IOPT, IMIN, NMASS, PROBS, IWK, WK, IR)
WRITE (NOUT,99999) IR
99999 FORMAT ('          ', 5I4)
END

```

### Output

```

Random deviates:   3   2   2   3   5
                   1   3   4   5   3

```

### Example 2

In this example, RNGDA is used to generate five pseudorandom binomial variates with parameters 20 and 0.5.

```

INTEGER      NMASS, NR
PARAMETER   (NMASS=21, NR=5)
C
INTEGER      IMIN, IOPT, IR(NR), ISEED, IWK(NMASS), K, N, NOUT
REAL         BINPR, P, PROBS(NMASS), WK(NMASS)
EXTERNAL     BINPR, RNGDA, RNSET, UMACH
C
CALL UMACH (2, NOUT)
N           = 20
P           = 0.5
IMIN        = 0
DO 10 K=1, NMASS
    PROBS(K) = BINPR(K-1,N,P)
10 CONTINUE
IOPT        = 0
ISEED       = 123457
CALL RNSET (ISEED)
CALL RNGDA (NR, IOPT, IMIN, NMASS, PROBS, IWK, WK, IR)

```

```

WRITE (NOUT,99999) IR
99999 FORMAT (' Binomial (20, .5) deviates: ', 5I4)
END

```

### Output

```
Binomial (20, .5) deviates: 12 10 16 12 11
```

## RNGDS/DRNGDS (Single/Double precision)

Set up table to generate pseudorandom numbers from a general discrete distribution.

### Usage

```
CALL RNGDS (PRF, IOPT, DEL, NNDX, IMIN, NMASS, CUMPR,
           LCUMPR)
```

### Arguments

**PRF** — User-supplied FUNCTION to compute the probability associated with each mass point of the distribution. The form is PRF(IX), where  
 IX — Point at which the probability function is to be evaluated. (Input)  
 IX can range from IMIN to the value at which the cumulative probability is greater than or equal to 1.0 – DEL.  
 PRF — Value of the probability function at IX. (Output)  
 PRF must be declared EXTERNAL in the calling program.

**IOPT** — Indicator of the extent to which CUMPR is initialized prior to calling RNGDS. (Input)

#### IOPT Action

0 RNGDS fills all of CUMPR, using PRF.  
 1 RNGDS fills only the index portion of CUMPR, using the values in the first NMASS positions. PRF is not used and may be a dummy function; also, IMIN and DEL are not used.

**DEL** — Maximum absolute error allowed in computing the cumulative probability. (Input)

Probabilities smaller than DEL are ignored; hence, DEL should be a small positive number. If DEL is too small, however, CUMPR(NMASS) must be exactly 1.0 since that value is compared to 1.0 – DEL.

**NNDX** — The number of elements of CUMPR available to be used as indexes. (Input)

NNDX must be greater than or equal to 1. In general, the larger NNDX is, to within sixty or seventy percent of NMASS, the more efficient the generation of random numbers using RNGDS will be.

**IMIN** — Smallest value the random deviate can assume. (Input/Output)  
 IMIN is not used if IOPT = 1. If IOPT = 0, PRF is evaluated at IMIN. If this

value is less than DEL, IMIN is incremented by 1 and again PRF is evaluated at IMIN. This process is continued until  $\text{PRF}(\text{IMIN}) \geq \text{DEL}$ . IMIN is output as this value and CUMPR(1) is output as PRF(IMIN).

**NMASS** — The number of mass points in the distribution. (Input, if IOPT = 1; output, if IOPT = 0)

If IOPT = 0, NMASS is the smallest integer such that  $\text{PRF}(\text{IMIN} + \text{NMASS} - 1) > 1.0 - \text{DEL}$ . NMASS does include the points  $\text{IMIN}(\text{in}) + j$  for which  $\text{PRF}(\text{IMIN}(\text{in}) + j) < \text{DEL}$ , for  $j = 0, 1, \dots, \text{IMIN}(\text{out}) - \text{IMIN}(\text{in})$ , where IMIN(in) denotes the input value of IMIN and IMIN(out) denotes its output value.

**CUMPR** — Vector of length NMASS + NNDX containing in the first NMASS positions, the cumulative probabilities and in some of the remaining positions, indexes to speed access to the probabilities. (Output, if IOPT = 0; input/output, otherwise)

CUMPR(NMASS + 1) + 1 is the actual number of index positions used.

**LCUMPR** — Dimension of CUMPR exactly as specified in the dimension statement in the calling program. (Input)

Since the logical length of CUMPR is determined in RNGDS, LCUMPR is used for error checking.

### Comments

1. Informational error
 

Type	Code	
3	1	For some I, CUMPR(I) is computed to be less than $1.0 - \text{DEL}$ , and yet $\text{CUMPR}(I + 1) - 1.0$ is greater than $1.0 - \text{CUMPR}(I + 1)$ . In this case, the maximum value that the random variable is allowed to take on is I; that is, CUMPR(I) is set to 1.0.
2. The routine RNGDT (page 1181) uses the table set up by RNGDS to generate random numbers from the distribution with CDF represented in CUMPR.

### Algorithm

Routine RNGDS sets up a table that routine RNGDT (page 1181) uses to generate pseudorandom deviates from a discrete distribution. The distribution can be specified either by its probability function PRF or by a vector of values of the cumulative probability function. Note that PRF is *not* the cumulative probability distribution function. If the cumulative probabilities are already available in CUMPR, the only reason to call RNGDS is to form an index vector in the upper portion of CUMPR so as to speed up the generation of random deviates by the routine RNGDT.

### Example 1

In this example, RNGDS is used to set up a table to generate pseudorandom variates from the discrete distribution:

$$\Pr(X = 1) = .05$$

$$\Pr(X = 2) = .45$$

$$\Pr(X = 3) = .31$$

$$\Pr(X = 4) = .04$$

$$\Pr(X = 5) = .15$$

In this simple example, we input the cumulative probabilities directly in CUMPR and request 3 indexes to be computed (NNDX = 4). Since the number of mass points is so small, the indexes would not have much effect on the speed of the generation of the random variates.

```
INTEGER      LCUMPR
PARAMETER    (LCUMPR=9)
C
INTEGER      IMIN, IOPT, NMASS, NNDX, NOUT
REAL         CUMPR(LCUMPR), DEL, PRF
EXTERNAL     PRF, RNGDS, UMACH
C
CALL UMACH (2, NOUT)
NMASS       = 5
CUMPR(1)    = 0.05
CUMPR(2)    = 0.50
CUMPR(3)    = 0.81
CUMPR(4)    = 0.85
CUMPR(5)    = 1.00
IOPT        = 1
NNDX        = 4
DEL         = 0.00001
CALL RNGDS (PRF, IOPT, DEL, NNDX, IMIN, NMASS, CUMPR, LCUMPR)
WRITE (NOUT,99999) CUMPR
99999 FORMAT (' Cumulative probabilities and indexes: ', /, 9F6.2)
END
C
C                                     Dummy function
REAL FUNCTION PRF (IX)
INTEGER      IX
C
RETURN
END
```

### Output

```
Cumulative probabilities and indexes:
0.05  0.50  0.81  0.85  1.00  3.00  1.00  2.00  5.00
```

## Example 2

This example, RNGDS is used to set up a table to generate binomial variates with parameters 20 and 0.5. The routine BINPR (page 1110) is used to compute the probabilities.

```
INTEGER      LCUMPR
PARAMETER   (LCUMPR=33)
C
INTEGER      I, IMIN, IOPT, N, NMASS, NNDX, NOUT
REAL        CUMPR(LCUMPR), DEL, P, PRF
COMMON      /BINCOM/ N, P
EXTERNAL    PRF, RNGDS, UMACH
C
CALL UMACH (2, NOUT)
N          = 20
P          = 0.5
IMIN       = 0
IOPT       = 0
NNDX       = 12
DEL        = 0.00001
CALL RNGDS (PRF, IOPT, DEL, NNDX, IMIN, NMASS, CUMPR, LCUMPR)
WRITE (NOUT,99998) IMIN, NMASS
99998 FORMAT (' The smallest point with positive probability using ',
&           /, ' the given DEL is ', I1, ' and all points after ', /,
&           ' point number ', I2, ' (counting from the input value ',
&           /, ' of IMIN) have zero probability.')
WRITE (NOUT,99999) (CUMPR(I),I=1,NMASS+NNDX)
99999 FORMAT (' Cumulative probabilities and indexes: ', /, (5X,8F8.4))
END
C
C                                     Compute binomial probabilities
REAL FUNCTION PRF (IX)
INTEGER      IX
C
INTEGER      N
REAL        BINPR, P
COMMON      /BINCOM/ N, P
EXTERNAL    BINPR
C
PRF = BINPR(IX,N,P)
RETURN
END
```

## Output

The smallest point with positive probability using  
the given DEL is 1 and all points after  
point number 19 (counting from the input value  
of IMIN) have zero probability.

Cumulative probabilities and indexes:

0.0000	0.0002	0.0013	0.0059	0.0207	0.0577	0.1316	0.2517
0.4119	0.5881	0.7483	0.8684	0.9423	0.9793	0.9941	0.9987
0.9998	1.0000	1.0000	11.0000	1.0000	7.0000	8.0000	9.0000
9.0000	10.0000	11.0000	11.0000	12.0000	13.0000	19.0000	



---

## RNGDT/DRNGDT (Single/Double precision)

Generate pseudorandom numbers from a general discrete distribution using a table lookup method.

### Usage

```
CALL RNGDT (NR, IMIN, NMASS, CUMPR, IR)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**IMIN** — Smallest value the random deviate can assume. (Input)  
This is the value corresponding to the probability in **CUMPR(1)**.

**NMASS** — Number of mass points in the discrete distribution. (Input)

**CUMPR** — Vector of length at least **NMASS + 1** containing in the first **NMASS** positions the cumulative probabilities and, possibly, indexes to speed access to the probabilities. (Input)

IMSL routine **RNGDS** (page 1177) can be used to initialize **CUMPR** properly. If no elements of **CUMPR** are used as indexes, **CUMPR(NMASS + 1)** is 0.0 on input. The value in **CUMPR(1)** is the probability of **IMIN**. The value in **CUMPR(NMASS)** must be exactly 1.0 (since this is the CDF at the upper range of the distribution).

**IR** — Vector of length **NR** containing the random discrete deviates. (Output)

### Comments

1. Informational error  

Type	Code	
3	1	The value in <b>CUMPR(NMASS)</b> is not exactly 1.0, but it was considered close enough to 1.0 that it was set to that value.
2. In the interest of efficiency, this routine does only limited error checking. If **CUMPR** is generated by the routine **RNGDS** (page 1177), the error checking is sufficient.
3. The routine **RNSET** (page 1167) can be used to initialize the seed of the random number generator. The routine **RNOPT** (page 1165) can be used to select the form of the generator.

### Algorithm

Routine **RNGDT** generates pseudorandom deviates from a discrete distribution, using the table **CUMPR**, which contains the cumulative probabilities of the distribution and, possibly, indexes to speed the search of the table. The routine **RNGDS** (page 1177) can be used to set up the table **CUMPR**. **RNGDT** uses the inverse CDF method to generate the variates.

### Example 1

These examples are the same ones used for the routine RNGDS (page 1177). In this first example, RNGDS is used to set up a table and then RNGDT is used to generate five pseudorandom variates from the discrete distribution:

$$\Pr(X = 1) = .05$$

$$\Pr(X = 2) = .45$$

$$\Pr(X = 3) = .31$$

$$\Pr(X = 4) = .04$$

$$\Pr(X = 5) = .15$$

The cumulative probabilities are input directly in CUMPR, and three indexes are computed by RNGDS (NNDX = 4). Since the number of mass points is so small, the indexes would not have much effect on the speed of the generation of the random variates.

```
INTEGER      LCUMPR, NR
PARAMETER   (LCUMPR=9, NR=5)
C
INTEGER      IMIN, IOPT, IR(NR), ISEED, NMASS, NNDX, NOUT
REAL        CUMPR(LCUMPR), DEL, PRF
EXTERNAL    PRF, RNGDS, RNGDT, RNSET, UMACH
C
CALL UMACH (2, NOUT)
IMIN       = 1
NMASS     = 5
CUMPR(1)  = 0.05
CUMPR(2)  = 0.50
CUMPR(3)  = 0.81
CUMPR(4)  = 0.85
CUMPR(5)  = 1.00
IOPT      = 1
NNDX     = 4
DEL       = 0.00001
C
                                Set up table
CALL RNGDS (PRF, IOPT, DEL, NNDX, IMIN, NMASS, CUMPR, LCUMPR)
ISEED = 123457
CALL RNSET (ISEED)
C
                                Generate variates
CALL RNGDT (NR, IMIN, NMASS, CUMPR, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Discrete random deviates: ', 5I4)
END
C
C
                                Dummy function
REAL FUNCTION PRF (IX)
INTEGER      IX
C
RETURN
END
```

### Output

Discrete random deviates: 5 2 3 3 4

## Example 2

In this example, RNGDS (page 1177) is used to set up a table and then RNGDT is used to generate five pseudorandom variates from the binomial distribution with parameters 20 and 0.5. The routine BINPR (page 1110) is used to compute the probabilities.

```
INTEGER      LCUMPR, NR
PARAMETER   (LCUMPR=33, NR=5)
C
INTEGER      IMIN, IOPT, IR(NR), ISEED, NMASS, NNDX, NOUT
REAL        CUMPR(LCUMPR), DEL, PRF
EXTERNAL    PRF, RNGDS, RNGDT, RNSET, UMACH
C
CALL UMACH (2, NOUT)
IMIN = 0
NMASS = 21
IOPT = 0
NNDX = 12
DEL = 0.00001
C
                                Set up table
CALL RNGDS (PRF, IOPT, DEL, NNDX, IMIN, NMASS, CUMPR, LCUMPR)
ISEED = 123457
CALL RNSET (ISEED)
C
                                Generate variates
CALL RNGDT (NR, IMIN, NMASS, CUMPR, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Binomial (20, 0.5) random deviates: ', 5I4)
END
C
                                Compute binomial probabilities
REAL FUNCTION PRF (IX)
INTEGER      IX
C
REAL        BINPR
EXTERNAL    BINPR
C
PRF = BINPR (IX, 20, 0.5)
RETURN
END
```

## Output

```
Binomial (20, 0.5) random deviates:   14   9  12  10  12
```

---

# RNGEO

Generate pseudorandom numbers from a geometric distribution.

## Usage

```
CALL RNGEO (NR, P, IR)
```

## Arguments

*NR* — Number of random numbers to generate. (Input)

$P$  — Probability of success on each trial. (Input)  
 $P$  must be positive and less than 1.0.

$IR$  — Vector of length  $NR$  containing the random geometric deviates. (Output)

### Comments

The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

### Algorithm

Routine `RNGEO` generates pseudorandom numbers from a geometric distribution with parameter  $P$ , where  $P$  is the probability of getting a success on any trial. A geometric deviate can be interpreted as the number of trials until the first success (including the trial in which the first success is obtained). The probability function is

$$f(x) = P(1 - P)^{x-1}$$

for  $x = 1, 2, \dots$  and  $0 < P < 1$

The geometric distribution as defined above has mean  $1/P$ .

The  $i$ -th geometric deviate is generated as the smallest integer not less than  $\log(U_i)/\log(1 - P)$ , where the  $U_i$  are independent uniform (0, 1) random numbers (see Knuth, 1981).

The geometric distribution is often defined on 0, 1, 2, ..., with mean  $(1 - P)/P$ . Such deviates can be obtained by subtracting 1 from each element of  $IR$ .

### Example

In this example, `RNGEO` is used to generate five pseudorandom deviates from a geometric distribution with parameter  $P$  equal to 0.3.

```
INTEGER      NR
PARAMETER   (NR=5)
C
INTEGER      IR(NR), ISEED, NOUT
REAL         P
EXTERNAL     RNGEO, RNSET, UMACH
C
CALL UMACH (2, NOUT)
P          = 0.3
ISEED     = 123457
CALL RNSET (ISEED)
CALL RNGEO (NR, P, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Geometric(0.3) random deviates: ', 5I8)
END
```

## Output

Geometric(0.3) random deviates:            1            4            1            2            1

---

# RNHYP

Generate pseudorandom numbers from a hypergeometric distribution.

## Usage

CALL RNHYP (NR, N, M, L, IR)

## Arguments

**NR** — Number of random numbers to generate. (Input)

**N** — Number of items in the sample. (Input)  
N must be positive.

**M** — Number of special items in the population, or lot. (Input)  
M must be positive.

**L** — Number of items in the lot. (Input)  
L must be greater than both N and M.

**IR** — Vector of length NR containing the random hypergeometric deviates. (Output)

Each element of IR can be considered to be the number of special items in a sample of size N drawn without replacement from a population of size L that contains M such special items.

## Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNHYP generates pseudorandom numbers from a hypergeometric distribution with parameters  $N$ ,  $M$ , and  $L$ . The hypergeometric random variable  $X$  can be thought of as the number of items of a given type in a random sample of size  $N$  that is drawn without replacement from a population of size  $L$  containing  $M$  items of this type. The probability function is

$$f(x) = \frac{\binom{M}{x} \binom{L-M}{N-x}}{\binom{L}{N}}$$

for  $x = \max(0, N - L + M), 1, 2, \dots, \min(N, M)$

If the hypergeometric probability function with parameters  $N$ ,  $M$ , and  $L$  evaluated at  $N - L + M$  (or at 0 if this is negative) is greater than the machine epsilon (`AMACH(4)` (Reference Material)), and less than 1.0 minus the machine epsilon, then `RNHYP` uses the inverse CDF technique. The routine recursively computes the hypergeometric probabilities, starting at  $x = \max(0, N - L + M)$  and using the ratio  $f(X = x + 1)/f(X = x)$  (see Fishman 1978, page 457).

If the hypergeometric probability function is too small or too close to 1.0, then `RNHYP` generates integer deviates uniformly in the interval  $[1, L - i]$ , for  $i = 0, 1, \dots$ ; and at the  $i$ -th step, if the generated deviate is less than or equal to the number of special items remaining in the lot, the occurrence of one special item is tallied and the number of remaining special items is decreased by one. This process continues until the sample size or the number of special items in the lot is reached, whichever comes first. This method can be much slower than the inverse CDF technique. The timing depends on  $N$ . If  $N$  is more than half of  $L$  (which in practical examples is rarely the case), the user may wish to modify the problem, replacing  $N$  by  $L - N$ , and to consider the deviates in `IR` to be the number of special items *not* included in the sample.

### Example

In this example, `RNHYP` is used to generate five pseudorandom deviates from a hypergeometric distribution to simulate taking random samples of size 4 from a lot containing 20 items of which 12 are defective. The resulting hypergeometric deviates represent the numbers of defectives in each of the five samples of size 4.

```

INTEGER      NR
PARAMETER   (NR=5)
C
INTEGER      IR(NR), ISEED, L, M, N, NOUT
EXTERNAL     RNHYP, RNSET, UMACH
C
CALL UMACH (2, NOUT)
N           = 4
M           = 12
L           = 20
ISEED      = 123457
CALL RNSET (ISEED)
CALL RNHYP (NR, N, M, L, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Hypergeometric random deviates: ', 5I8)
END

```

### Output

```
Hypergeometric random deviates:      4      2      3      3      3
```

---

## RNLGR

Generate pseudorandom numbers from a logarithmic distribution.

## Usage

CALL RNLGR (NR, A, IR)

## Arguments

**NR** — Number of random numbers to generate. (Input)

**A** — Parameter of the logarithmic distribution. (Input)  
A must be positive and less than 1.0.

**IR** — Vector of length NR containing the random logarithmic deviates. (Output)

## Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNLGR generates pseudorandom numbers from a logarithmic distribution with parameter A. The probability function is

$$f(x) = -\frac{a^x}{x \ln(1-a)}$$

for  $x = 1, 2, 3, \dots$ , and  $0 < a < 1$ .

The methods used are described by Kemp (1981) and depend on the value of A. If A is less than 0.95, Kemp's algorithm LS, which is a "chop-down" variant of an inverse CDF technique, is used. Otherwise, Kemp's algorithm LK, which gives special treatment to the highly probable values of 1 and 2, is used.

## Example

In this example, RNLGR is used to generate 5 pseudo-random deviates from a logarithmic distribution with parameter A equal to 0.3.

```
C      INTEGER      NR
      PARAMETER    (NR=5)

C      INTEGER      IR(NR), ISEED, NOUT
      REAL          A
      EXTERNAL      RNLGR, RNSET, UMACH

C      CALL UMACH (2, NOUT)
      A      = 0.3
      ISEED = 123457
      CALL RNSET (ISEED)
      CALL RNLGR (NR, A, IR)
      WRITE (NOUT,99999) IR
99999 FORMAT (' Logarithmic (0.3) random deviates: ', 5I8)
      END
```

## Output

Logarithmic (0.3) random deviates:            2            1            1            1            2

---

# RNNBN

Generate pseudorandom numbers from a negative binomial distribution.

## Usage

CALL RNNBN (NR, RK, P, IR)

## Arguments

**NR** — Number of random numbers to generate. (Input)

**RK** — Negative binomial parameter. (Input)  
RK must be positive.

**P** — Probability of success on each trial. (Input)  
P must be greater than the machine epsilon, `AMACH(4)` (Reference Material) and less than 1.0.

**IR** — Vector of length NR containing the random negative binomial deviates. (Output)

## Comments

1. The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.
2. If RK is an integer, the deviates in IR can be thought of as the number of failures in a sequence of Bernoulli trials before RK successes occur.

## Algorithm

Routine RNNBN generates pseudorandom numbers from a negative binomial distribution with parameters RK and P. RK and P must be positive and P must be less than 1. The probability function (with  $r = RK$  and  $p = P$ ) is

$$f(x) = \binom{r+x-1}{x} (1-p)^r p^x$$

for  $x = 0, 1, 2, \dots$

If  $r$  is an integer, the distribution is often called the Pascal distribution and can be thought of as modeling the length of a sequence of Bernoulli trials until  $r$  successes are obtained, where  $p$  is the probability of getting a success on any trial. In this form, the random variable takes values  $r, r+1, r+2, \dots$  and can be obtained from the negative binomial random variable defined above by adding  $r$



to the negative binomial variable. This latter form is also equivalent to the sum of  $r$  geometric random variables defined as taking values 1, 2, 3, ... .

If  $rp/(1-p)$  is less than 100 and  $(1-p)^r$  is greater than the machine epsilon, RNNBN uses the inverse CDF technique; otherwise, for each negative binomial deviate, RNNBN generates a gamma  $(r, p/(1-p))$  deviate  $Y$  and then generates a Poisson deviate with parameter  $Y$ .

### Example

In this example, RNNBN is used to generate five pseudorandom deviates from a negative binomial (Pascal) distribution with parameter  $r$  equal to 4 and  $p$  equal to 0.3.

```
INTEGER      NR
PARAMETER   (NR=5)
C
INTEGER      IR(NR), ISEED, NOUT
REAL        P, RK
EXTERNAL    RNNBN, RNSET, UMACH
C
CALL UMACH (2, NOUT)
P          = 0.3
RK         = 4.0
ISEED     = 123457
CALL RNSET (ISEED)
CALL RNNBN (NR, RK, P, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Negative binomial (4.0, 0.3) random deviates: ', 5I4)
END
```

### Output

```
Negative binomial (4.0, 0.3) random deviates:    5    1    3    2    3
```

---

## RNPOI

Generate pseudorandom numbers from a Poisson distribution.

### Usage

```
CALL RNPOI (NR, THETA, IR)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**THETA** — Mean of the Poisson distribution. (Input)

THETA must be positive.

**IR** — Vector of length NR containing the random Poisson deviates. (Output)

## Comments

The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

## Algorithm

Routine `RNPOI` generates pseudorandom numbers from a Poisson distribution with parameter `THETA`. `THETA`, which is the mean of the Poisson random variable, must be positive. The probability function (with  $\theta = \text{THETA}$ ) is

$$f(x) = e^{-\theta} \theta^x / x!$$

for  $x = 0, 1, 2, \dots$

If `THETA` is less than 15, `RNPOI` uses an inverse CDF method; otherwise the `PTPE` method of Schmeiser and Kachitvichyanukul (1981) (see also Schmeiser 1983) is used.

The `PTPE` method uses a composition of four regions, a triangle, a parallelogram, and two negative exponentials. In each region except the triangle, acceptance/rejection is used. The execution time of the method is essentially insensitive to the mean of the Poisson.

## Example

In this example, `RNPOI` is used to generate five pseudorandom deviates from a Poisson distribution with mean equal to 0.5.

```
INTEGER      NR
PARAMETER   (NR=5)
C
INTEGER      IR(NR), ISEED, NOUT
REAL        THETA
EXTERNAL    RNPOI, RNSET, UMACH
C
CALL UMACH (2, NOUT)
THETA = 0.5
ISEED = 123457
CALL RNSET (ISEED)
CALL RNPOI (NR, THETA, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Poisson(0.5) random deviates: ', 5I8)
END
```

## Output

```
Poisson(0.5) random deviates:      2      0      1      0      1
```

---

# RNUND

Generate pseudorandom numbers from a discrete uniform distribution.

## Usage

CALL RNUND (NR, K, IR)

## Arguments

**NR** — Number of random numbers to generate. (Input)

**K** — Parameter of the discrete uniform distribution. (Input)

The integers 1, 2, ...,  $\kappa$  occur with equal probability.  $\kappa$  must be positive.

**IR** — Vector of length NR containing the random discrete uniform deviates. (Output)

## Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNUND generates pseudorandom numbers from a discrete uniform distribution over the integers 1, 2, ...,  $\kappa$ . A random integer is generated by multiplying  $\kappa$  by a uniform (0, 1) random number, adding 1.0, and truncating the result to an integer. This, of course, is equivalent to sampling with replacement from a finite population of size  $\kappa$ . To do the equivalent of sampling without replacement, the routine RNSRI (page 1241) can be used.

## Example

In this example, RNUND is used to generate five pseudorandom deviates from a discrete uniform distribution over the integers from 1 to 6.

```
INTEGER      IR(5), ISEED, K, NOUT, NR
EXTERNAL    RNSET, RNUND, UMACH
C
CALL UMACH (2, NOUT)
K          = 6
NR         = 5
ISEED     = 123457
CALL RNSET (ISEED)
CALL RNUND (NR, K, IR)
WRITE (NOUT,99999) IR
99999 FORMAT (' Discrete uniform (1,6) random deviates: ', 5I7)
END
```

## Output

```
Discrete uniform (1,6) random deviates:      6      2      5      4      6
```

---

# RNBET/DRNBET (Single/Double precision)

Generate pseudorandom numbers from a beta distribution.

## Usage

CALL RNBET (NR, PIN, QIN, R)

## Arguments

**NR** — Number of random numbers to generate. (Input)

**PIN** — First beta distribution parameter. (Input)  
PIN must be positive.

**QIN** — Second beta distribution parameter. (Input)  
QIN must be positive.

**R** — Vector of length NR containing the random standard beta deviates. (Output)

## Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNBET generates pseudorandom numbers from a beta distribution with parameters PIN and QIN, both of which must be positive. With  $p = \text{PIN}$  and  $q = \text{QIN}$ , the probability density function is

$$f(x) = \frac{\Gamma(p+q)}{\Gamma(p)\Gamma(q)} x^{p-1}(1-x)^{q-1} \quad \text{for } 0 \leq x \leq 1$$

where  $\Gamma(\cdot)$  is the gamma function.

The algorithm used depends on the values of  $p$  and  $q$ . Except for the trivial cases of  $p = 1$  or  $q = 1$ , in which the inverse CDF method is used, all of the methods use acceptance/rejection. If  $p$  and  $q$  are both less than 1, the method of Johnk (1964) is used; if either  $p$  or  $q$  is less than 1 and the other is greater than 1, the method of Atkinson (1979) is used; if both  $p$  and  $q$  are greater than 1, algorithm BB of Cheng (1978), which requires very little setup time, is used if NR is less than 4; and algorithm B4PE of Schmeiser and Babu (1980) is used if NR is greater than or equal to 4. Note that for  $p$  and  $q$  both greater than 1, calling RNBET in a loop getting less than 4 variates on each call will not yield the same set of deviates as calling RNBET once and getting all the deviates at once.

The values returned in R are less than 1.0 and greater than  $\epsilon$ , where  $\epsilon$  is the smallest positive number such that  $1.0 - \epsilon$  is less than 1.0.

## Example

In this example, RNBET is used to generate five pseudorandom beta (3, 2) variates.

```

      INTEGER      NR
      PARAMETER   (NR=5)
C
      INTEGER      ISEED, NOUT
      REAL         PIN, QIN, R(NR)
      EXTERNAL     RNBET, RNSET, UMACH
C
      CALL UMACH (2, NOUT)
      PIN  = 3.0
      QIN  = 2.0
      ISEED = 123457
      CALL RNSET (ISEED)
      CALL RNBET (NR, PIN, QIN, R)
      WRITE (NOUT,99999) R
99999 FORMAT ('      Beta (3,2) random deviates: ', 5F7.4)
      END

```

### Output

```
Beta (3,2) random deviates:  0.2814 0.9483 0.3984 0.3103 0.8296
```

---

## RNCHI/DRNCHI (Single/Double precision)

Generate pseudorandom numbers from a chi-squared distribution.

### Usage

```
CALL RNCHI (NR, DF, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**DF** — Degrees of freedom. (Input)  
DF must be positive.

**R** — Vector of length NR containing the random chi-squared deviates. (Output)

### Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNCHI generates pseudorandom numbers from a chi-squared distribution with DF degrees of freedom. If DF is an even integer less than 17, the chi-squared deviate  $r$  is generated as

$$r = -2 \ln\left(\prod_{i=1}^n u_i\right)$$

where  $n = \text{DF}/2$  and the  $u_i$  are independent random deviates from a uniform (0, 1) distribution. If  $\text{DF}$  is an odd integer less than 17, the chi-squared deviate is generated in the same way, except the square of a normal deviate is added to the expression above. If  $\text{DF}$  is greater than 16 or is not an integer, and if it is not too large to cause overflow in the gamma random number generator, the chi-squared deviate is generated as a special case of a gamma deviate, using routine `RNGAM` (page 1198). If overflow would occur in `RNGAM`, the chi-squared deviate is generated in the manner described above, using the logarithm of the product of uniforms, but scaling the quantities to prevent underflow and overflow.

### Example

In this example, `RNCHI` is used to generate five pseudorandom chi-squared deviates with 5 degrees of freedom.

```

INTEGER      ISEED, NOUT, NR
REAL         DF, R(5)
EXTERNAL     RNCHI, RNSET, UMACH
C
CALL UMACH (2, NOUT)
DF      = 5.0
NR      = 5
ISEED = 123457
CALL RNSET (ISEED)
CALL RNCHI (NR, DF, R)
WRITE (NOUT,99999) R
99999 FORMAT (' Chi-squared random deviates with 5 df: ', 5F7.3)
END

```

### Output

```
Chi-squared random deviates with 5 df:  12.090  0.481  1.798 14.871  1.748
```

---

## RNCHY/DRNCHY (Single/Double precision)

Generate pseudorandom numbers from a Cauchy distribution.

### Usage

```
CALL RNCHY (NR, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**R** — Vector of length `NR` containing the random Cauchy deviates. (Output)

### Comments

The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNCHY generates pseudorandom numbers from a standard Cauchy distribution. The probability density function is

$$f(x) = \frac{1}{\pi(1+x^2)}$$

Use of the inverse CDF technique would yield a Cauchy deviate from a uniform (0, 1) deviate,  $u$ , as  $\tan[\pi(u - .5)]$ . Rather than evaluating a tangent directly, however, RNCHY generates two uniform (-1, 1) deviates,  $x_1$  and  $x_2$ . These values can be thought of as sine and cosine values. If

$$x_1^2 + x_2^2$$

is less than or equal to 1, then  $x_1/x_2$  is delivered as the Cauchy deviate; otherwise,  $x_1$  and  $x_2$  are rejected and two new uniform (-1, 1) deviates are generated. This method is also equivalent to taking the ratio of two independent normal deviates.

Deviates from the Cauchy distribution with median  $T$  and first quartile  $T - S$ , that is, with density

$$f(x) = \frac{S}{\pi[S^2 + (x - T)^2]}$$

can be obtained by scaling the output from RNCHY. The following statements (in single precision) would yield random deviates from this Cauchy distribution.

```
CALL RNCHY (NR, R)
CALL SSCAL (NR, S, R, 1)
CALL SADD (NR, T, R, 1)
```

The Cauchy distribution is a member of the symmetric stable family of distributions. The routine RNSTA (page 1209) can be used to generate deviates from this more general family of distributions or even from the stable family not requiring symmetry.

## Example

In this example, RNCHY is used to generate five pseudorandom deviates from a Cauchy distribution.

```
C
INTEGER      ISEED, NOUT, NR
REAL         R(5)
EXTERNAL     RNCHY, RNSET, UMACH

CALL UMACH (2, NOUT)
NR         = 5
ISEED     = 123457
CALL RNSET (ISEED)
CALL RNCHY (NR, R)
WRITE (NOUT, 99999) R
```

```
99999 FORMAT ( '          Cauchy random deviates: ', 5F8.4)
      END
```

### Output

```
Cauchy random deviates:   3.5765  0.9353 15.5797  2.0815 -0.1333
```

---

## RNEXP/DRNEXP (Single/Double precision)

Generate pseudorandom numbers from a standard exponential distribution.

### Usage

```
CALL RNEXP (NR, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**R** — Vector of length NR containing the random standard exponential deviates. (Output)

### Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNEXP generates pseudorandom numbers from a standard exponential distribution. The probability density function is  $f(x) = e^{-x}$ ; for  $x > 0$ . RNEXP uses an antithetic inverse CDF technique; that is, a uniform random deviate  $U$  is generated and the inverse of the exponential cumulative distribution function is evaluated at  $1.0 - U$  to yield the exponential deviate.

Deviates from the exponential distribution with mean THETA can be generated by using RNEXP and then multiplying each entry in R by THETA. The following statements (in single precision using the routine SSCAL (Reference Material)) would yield random deviates from such a distribution:

```
CALL RNEXP (NR, R)
CALL SSCAL (NR, THETA, R, 1)
```

### Example

In this example, RNEXP is used to generate five pseudorandom deviates from a standard exponential distribution.

```
INTEGER   ISEED, NOUT, NR
REAL      R(5)
EXTERNAL  RNEXP, RNSET, UMACH
```

C



```

CALL UMACH (2, NOUT)
NR      = 5
ISEED = 123457
CALL RNSET (ISEED)
CALL RNEXP (NR, R)
WRITE (NOUT,99999) R
99999 FORMAT ('      Exponential random deviates: ', 5F8.4)
END

```

### Output

Exponential random deviates: 0.0344 1.3443 0.2662 0.5633 0.1686

---

## RNEXT/DRNEXT (Single/Double precision)

Generate pseudorandom numbers from a mixture of two exponential distributions.

### Usage

```
CALL RNEXT (NR, THETA1, THETA2, P, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**THETA1** — Mean of the exponential distribution that has the larger mean. (Input)

**THETA2** — Mean of the exponential distribution that has the smaller mean. (Input)

THETA2 must be positive and less than or equal to THETA1.

**P** — Mixing parameter. (Input)

P must be nonnegative and less than or equal to THETA1/(THETA1 - THETA2).

**R** — Vector of length NR containing the random deviates from a mixture of exponentials. (Output)

### Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNEXT generates pseudorandom numbers from a mixture of two exponential distributions. The probability density function is

$$f(x) = \frac{p}{\theta_1} e^{-x/\theta_1} + \frac{1-p}{\theta_2} e^{-x/\theta_2} \quad \text{for } x > 0$$

where  $p = P$ ,  $\theta_1 = \text{THETA1}$ , and  $\theta_2 = \text{THETA2}$ .

In the case of a convex mixture, that is, the case  $0 < p < 1$ , the mixing parameter  $p$  is interpretable as a probability; and `RNEXT` with probability  $p$  generates an exponential deviate with mean  $\theta_1$ , and with probability  $1 - p$  generates an exponential with mean  $\theta_2$ . When  $p$  is greater than 1, but less than  $\theta_1/(\theta_1 - \theta_2)$ , then either an exponential deviate with mean  $\theta_1$  or the sum of two exponentials with means  $\theta_1$  and  $\theta_2$  is generated. The probabilities are  $q = p - (p - 1)\theta_1/\theta_2$  and  $1 - q$ , respectively, for the single exponential and the sum of the two exponentials.

### Example

In this example, `RNEXT` is used to generate five pseudorandom deviates from a mixture of exponentials with means 2 and 1, respectively, and with mixing parameter 0.5.

```
INTEGER      ISEED, NOUT, NR
REAL         P, R(5), THETA1, THETA2
EXTERNAL    RNEXT, RNSET, UMACH
C
CALL UMACH (2, NOUT)
THETA1 = 2.0
THETA2 = 1.0
P      = 0.5
NR     = 5
ISEED  = 123457
CALL RNSET (ISEED)
CALL RNEXT (NR, THETA1, THETA2, P, R)
WRITE (NOUT,99999) R
99999 FORMAT (' Random deviates from a mixture of exponentials: ', /,
&           5X, 5F8.4)
END
```

### Output

```
Random deviates from a mixture of exponentials:
0.0700 1.3024 0.6301 1.9756 0.3716
```

---

## RNGAM/DRNGAM (Single/Double precision)

Generate pseudorandom numbers from a standard gamma distribution.

### Usage

```
CALL RNGAM (NR, A, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**A** — The shape parameter of the gamma distribution. (Input)  
This parameter must be positive.

**R** — Vector of length NR containing the random standard gamma deviates.  
(Output)

### Comments

The routine RNSSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNGAM generates pseudorandom numbers from a gamma distribution with shape parameter  $a$  and unit scale parameter. The probability density function is

$$f(x) = \frac{1}{\Gamma(a)} x^{a-1} e^{-x} \quad \text{for } x \geq 0$$

Various computational algorithms are used depending on the value of the shape parameter  $a$ . For the special case of  $a = 0.5$ , squared and halved normal deviates are used; and for the special case of  $a = 1.0$ , exponential deviates (from IMSL routine RNEXP, page 1196) are used. Otherwise, if  $a$  is less than 1.0, an acceptance-rejection method due to Ahrens, described in Ahrens and Dieter (1974), is used; if  $a$  is greater than 1.0, a ten-region rejection procedure developed by Schmeiser and Lal (1980) is used.

Deviates from the two-parameter gamma distribution with shape parameter  $a$  and scale parameter  $b$  can be generated by using RNGAM and then multiplying each entry in R by  $b$ . The following statements (in single precision) would yield random deviates from a gamma ( $a, b$ ) distribution.

```
CALL RNGAM (NR, A, R)
CALL SSCAL (NR, B, R, 1)
```

The Erlang distribution is a standard gamma distribution with the shape parameter having a value equal to a positive integer; hence, RNGAM generates pseudorandom deviates from an Erlang distribution with no modifications required.

### Example

In this example, RNGAM is used to generate five pseudorandom deviates from a gamma (Erlang) distribution with shape parameter equal to 3.0.

```
C
INTEGER      NR
PARAMETER   (NR=5)

C
INTEGER      ISEED, NOUT
REAL         A, R(NR)
EXTERNAL    RNGAM, RNSSET, UMACH

C
CALL UMACH (2, NOUT)
A          = 3.0
```

```

ISEED = 123457
CALL RNSET (ISEED)
CALL RNGAM (NR, A, R)
WRITE (NOUT,99999) R
99999 FORMAT (' Gamma(3) random deviates: ', 5F8.4)
END

```

### Output

```
Gamma(3) random deviates: 6.8428 3.4452 1.8535 3.9992 0.7794
```

## RNGCS/DRNGCS (Single/Double precision)

Set up table to generate pseudorandom numbers from a general continuous distribution.

### Usage

```
CALL RNGCS (CDF, IOPT, NDATA, TABLE, LD_TBL)
```

### Arguments

**CDF** — User-supplied FUNCTION to compute the cumulative distribution function. The form is CDF(X), where

X — Point at which the distribution function is to be evaluated. (Input)

CDF — Value of the distribution function at X. (Output)

CDF must be declared EXTERNAL in the calling program.

**IOPT** — Indicator of the extent to which TABLE is initialized prior to calling RNGCS. (Input)

#### IOPT Action

0 RNGCS fills the last four columns of TABLE. The user inputs the points at which the CDF is to be evaluated in the first column of TABLE. These must be in ascending order.

1 RNGCS fills the last three columns of TABLE. CDF is not used and may be a dummy function; instead, the cumulative distribution function is specified in the first two columns of TABLE. The abscissas (in the first column) must be in ascending order and the function must be strictly monotonically increasing.

**NDATA** — Number of points at which the CDF is evaluated for interpolation. (Input)

NDATA must be greater than or equal to 4.

**TABLE** — NDATA by 5 table to be used for interpolation of the cumulative distribution function. (Input and output)

The first column of TABLE contains abscissas of the cumulative distribution function in ascending order, the second column contains the values of the CDF (which must be strictly increasing), and the remaining columns contain values

used in interpolation. The first row of `TABLE` corresponds to the left limit of the support of the distribution and the last row corresponds to the right limit of the support; that is, `TABLE(1, 2) = 0.0` and `TABLE(NDATA, 2) = 1.0`.

***LDTABL*** — Leading dimension of `TABLE` exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Informational error
 

Type	Code	
3	1	The values in <code>TABLE(1, 2)</code> and/or <code>TABLE(NDATA, 2)</code> are not exactly 0.0 and 1.0, respectively, but they are considered close enough to these values that they are set to these values.
  
2. The routine `RNGCT` (page 1202) uses the table set up by `RNGCS` to generate random numbers from the distribution with CDF represented in `TABLE`.

### Algorithm

Routine `RNGCS` sets up a table that routine `RNGCT` (page 1202) can use to generate pseudorandom deviates from a continuous distribution. The distribution is specified by its cumulative distribution function, which can be supplied either in tabular form in `TABLE` or by a FORTRAN function `CDF`. See the documentation for the routine `RNGCT` for a description of the method.

### Example

In this example, `RNGCS` is used to set up a table to generate pseudorandom variates from a beta distribution. This example is continued in the documentation for routine `RNGCT` (page 1202) to generate the random variates.

```

INTEGER      LDATABL
PARAMETER   (LDATABL=100)
C
INTEGER      I, IOPT, NINT, NOUT
REAL         CDF, PIN, QIN, TABLE(LDATABL,5), X
COMMON      /BCOM/ PIN, QIN
EXTERNAL    CDF, RNGCS, UMACH
C
CALL UMACH (2, NOUT)
PIN  = 3.0
QIN  = 2.0
IOPT = 0
NINT = 100
X    = 0.0
C
C                                     Fill the first column of the table
C                                     with abscissas for interpolation.
DO 10 I=1, NINT
   TABLE(I,1) = X
   X           = X + 0.01
10 CONTINUE
CALL RNGCS (CDF, IOPT, NINT, TABLE, LDATABL)

```

```

        WRITE (NOUT,99999) (TABLE(I,1),TABLE(I,2),I=1,10)
99999 FORMAT ('  First few elements of the table: ', F4.2, F8.4, /,
&          (36X,F4.2,F8.4))
        END
C
C          Beta distribution function
C
      REAL FUNCTION CDF (X)
      REAL      X
C
      REAL      BETDF, PIN, QIN
      COMMON    /BCOM/ PIN, QIN
      EXTERNAL  BETDF
C
      CDF = BETDF(X,PIN,QIN)
      RETURN
      END

```

### Output

```

*** WARNING  ERROR 1 from RNGCS.  The values of the CDF in the second
***          column of TABLE did not begin at 0.0 and end at 1.0, but they
***          have been adjusted. Prior to adjustment,
***          TABLE(1,2) = 0.000000E+00 and TABLE(NDATA,2) = 9.994079E-01.
First few elements of the table:  0.00  0.0000
                                0.01  0.0000
                                0.02  0.0000
                                0.03  0.0001
                                0.04  0.0002
                                0.05  0.0005
                                0.06  0.0008
                                0.07  0.0013
                                0.08  0.0019
                                0.09  0.0027

```

---

## RNGCT/DRNGCT (Single/Double precision)

Generate pseudorandom numbers from a general continuous distribution.

### Usage

```
CALL RNGCT (NR, NDATA, TABLE, LDTABL, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**NDATA** — Number of points at which the cumulative distribution function is evaluated for interpolation. (Input)  
 NDATA must be greater than or equal to 4.

**TABLE** — NDATA by 5 table to be used for interpolation of the cumulative distribution function. (Input)

The first column of **TABLE** contains abscissas of the cumulative distribution function in ascending order, the second column contains the values of the CDF (which must be strictly increasing beginning with 0.0 and ending at 1.0) and the

remaining columns contain values used in interpolation. This table is set up using routine RNGCS (page 1200).

**LDTABL** — Leading dimension of TABLE exactly as specified in the dimension statement in the calling program. (Input)

**R** — Vector of length NR containing the random deviates. (Output)

### Comments

1. The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.
2. In the interest of efficiency, this routine does only limited error checking. If TABLE is generated by the routine RNGCS (page 1200), the error checking is sufficient.

### Algorithm

Routine RNGCT generates pseudorandom numbers from a continuous distribution using the inverse CDF technique, by interpolation of points of the distribution function given in TABLE, which is set up by routine RNGCS (page 1200). A strictly monotone increasing distribution function is assumed. The interpolation is by an algorithm attributable to Akima (1970), using piecewise cubics. The use of this technique for generation of random numbers is due to Guerra, Tapia, and Thompson (1976), who give a description of the algorithm and accuracy comparisons between this method and linear interpolation. The relative errors using the Akima interpolation are generally considered very good.

### Example

In this example, RNGCS (page 1200) is used to set up a table for generation of beta pseudorandom deviates. The CDF for this distribution is computed by the routine BETDF (page 1125). The table contains 100 points at which the CDF is evaluated and that are used for interpolation.

```
C      INTEGER      LDTABL, NR
      PARAMETER    (LDTABL=100, NR=5)

C      INTEGER      I, IOPT, ISEED, NINT, NOUT
      REAL          CDF, PIN, QIN, R(NR), TABLE(LDTABL,5), X
      COMMON       /BCOM/ PIN, QIN
C      EXTERNAL    CDF, RNGCS, RNGCT, RNSET, UMACH

C      CALL UMACH (2, NOUT)
      PIN = 3.0
      QIN = 2.0
      IOPT = 0
      NINT = 100
      X = 0.0

C                                     Fill the first column of the table
C                                     with abscissas for interpolation.
      DO 10 I=1, NINT
```

```

        TABLE(I,1) = X
        X           = X + 0.01
10 CONTINUE
    CALL RNGCS (CDF, IOPT, NINT, TABLE, LDTABL)
C                                     Initialize seed of random number
C                                     generator.
    ISEED = 123457
    CALL RNSET (ISEED)
C                                     Now generate the random deviates.
    CALL RNGCT (NR, NINT, TABLE, LDTABL, R)
    WRITE (NOUT,99999) R
99999 FORMAT (' Beta (3,2) random deviates: ', 5F7.4)
    END
C
C                                     Beta distribution function
    REAL FUNCTION CDF (X)
    REAL          X
C
    REAL          BETDF, PIN, QIN
    COMMON        /BCOM/ PIN, QIN
    EXTERNAL      BETDF
C
    CDF = BETDF(X,PIN,QIN)
    RETURN
    END

```

### Output

```

*** WARNING  ERROR 1 from RNGCS.  The values of the CDF in the second
***          column of TABLE did not begin at 0.0 and end at 1.0, but they
***          have been adjusted. Prior to adjustment,
***          TABLE(1,2) = 0.000000E+00 and TABLE(NDATA,2) = 9.994079E-01.
Beta (3,2) random deviates:  0.9208 0.4641 0.7668 0.6536 0.8171

```

---

## RNLNL/DRNLNL (Single/Double precision)

Generate pseudorandom numbers from a lognormal distribution.

### Usage

```
CALL RNLNL (NR, XM, S, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**XM** — Mean of the underlying normal distribution. (Input)

**S** — Standard deviation of the underlying normal distribution. (Input)  
S must be positive.

**R** — Vector of length NR containing the random lognormal deviates. (Output)  
The log of each element of R has a normal distribution with mean XM and standard deviation S.



## Comments

The routine RNSSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNLNL generates pseudorandom numbers from a lognormal distribution with parameters  $\mu$  and  $\sigma$ . The scale parameter in the underlying normal distribution,  $\sigma$ , must be positive. The method is to generate normal deviates with mean  $\mu$  and standard deviation  $\sigma$  and then to exponentiate the normal deviates.

With  $\mu = \mu$  and  $\sigma = \sigma$ , the probability density function for the lognormal distribution is

$$f(x) = \frac{1}{\sigma x \sqrt{2\pi}} \exp\left[-\frac{1}{2\sigma^2} (\ln x - \mu)^2\right] \quad \text{for } x > 0$$

The mean and variance of the lognormal distribution are  $\exp(\mu + \sigma^2/2)$  and  $\exp(2\mu + 2\sigma^2) - \exp(2\mu + \sigma^2)$ , respectively.

## Example

In this example, RNLNL is used to generate five pseudorandom lognormal deviates with  $\mu = 0$  and  $\sigma = 1$ .

```
C      INTEGER      NR
      PARAMETER    (NR=5)

C      INTEGER      ISEED, NOUT
      REAL          R(NR), S, XM
      EXTERNAL      RNLNL, RNSET, UMACH

C      CALL UMACH (2, NOUT)
      XM      = 0.0
      S      = 1.0
      ISEED = 123457
      CALL RNSET (ISEED)
      CALL RNLNL (NR, XM, S, R)
      WRITE (NOUT,99999) R
99999 FORMAT (' Lognormal random deviates: ', 5F8.4)
      END
```

## Output

```
Lognormal random deviates:  7.7801  2.9543  1.0861  3.5885  0.2935
```

---

## RNNOA/DRNNOA (Single/Double precision)

Generate pseudorandom numbers from a standard normal distribution using an acceptance/rejection method.

## Usage

CALL RNNOA (NR, R)

## Arguments

*NR* — Number of random numbers to generate. (Input)

*R* — Vector of length *NR* containing the random standard normal deviates.  
(Output)

## Comments

The routine *RNSET* (page 1167) can be used to initialize the seed of the random number generator. The routine *RNOPT* (page 1165) can be used to select the form of the generator.

## Algorithm

Routine *RNNOA* generates pseudorandom numbers from a standard normal (Gaussian) distribution using an acceptance/rejection technique due to Kinderman and Ramage (1976). In this method, the normal density is represented as a mixture of densities over which a variety of acceptance/rejection methods due to Marsaglia (1964), Marsaglia and Bray (1964), and Marsaglia, MacLaren, and Bray (1964) are applied. This method is faster than the inverse CDF technique used in *RNNOR* (page 1208) to generate standard normal deviates.

Deviates from the normal distribution with mean *XM* and standard deviation *XSTD* can be obtained by scaling the output from *RNNOA*. The following statements (in single precision) would yield random deviates from a normal (*XM*, *XSTD\*\*2*) distribution.

```
CALL RNNOA (NR, R)
CALL SSCAL (NR, XSTD, R, 1)
CALL SADD (NR, XM, R, 1)
```

## Example

In this example, *RNNOA* is used to generate five pseudorandom deviates from a standard normal distribution.

```
C
INTEGER      ISEED, NOUT, NR
REAL         R(5)
EXTERNAL     RNNOA, RNSET, UMACH

CALL UMACH (2, NOUT)
NR      = 5
ISEED = 123457
CALL RNSET (ISEED)
CALL RNNOA (NR, R)
WRITE (NOUT,99999) R
99999 FORMAT (' Standard normal random deviates: ', 5F8.4)
END
```

## Output

Standard normal random deviates: 2.0516 1.0833 0.0826 1.2777 -1.2260

---

# RNNOF/DRNNOF (Single/Double precision)

Generate a pseudorandom number from a standard normal distribution.

## Usage

RNNOF ( )

## Argument

*RNNOF* — Function value, a random standard normal deviate. (Output)

## Comments

1. The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.
2. This function has a side effect: it changes the value of the seed, which is passed through a common block.

## Algorithm

Routine RNNOF is the function form of RNNOR. If several standard normal deviates are needed, it may be more efficient to obtain them all at once by a call to RNNOR, rather than by several references to RNNOF.

## Example

In this example, RNNOF is used to generate five pseudorandom standard normal numbers.

```
INTEGER    I, ISEED, NOUT, NR
REAL      R(5), RNNOF
EXTERNAL  RNNOF, RNSET, UMACH
C
CALL UMACH (2, NOUT)
ISEED = 123457
CALL RNSET (ISEED)
DO 10 I=1, NR
    R(I) = RNNOF()
10 CONTINUE
WRITE (NOUT,99999) R
99999 FORMAT (' Standard normal random deviates: ', 5F8.4)
END
```

## Output

Standard normal random deviates: 0.0000 0.0000 0.0000 0.0000 0.0000

---

## RNNOR/DRNNOR (Single/Double precision)

Generate pseudorandom numbers from a standard normal distribution using an inverse CDF method.

### Usage

```
CALL RNNOR (NR, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**R** — Vector of length NR containing the random standard normal deviates. (Output)

### Comments

The routine RNSSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNNOR generates pseudorandom numbers from a standard normal (Gaussian) distribution using an inverse CDF technique. In this method, a uniform (0,1) random deviate is generated and then the inverse of the normal distribution function is evaluated at that point, using the routine ANORIN (page 1124). This method is slower than the acceptance/rejection technique used in the routine RNNOA (page 1205) to generate standard normal deviates. Deviates from the normal distribution with mean XM and standard deviation XSTD can be obtained by scaling the output from RNNOR. The following statements (in single precision, using the routines SSCAL (IMSL MATH/LIBRARY) and SADD (IMSL MATH/LIBRARY).) would yield random deviates from a normal (XM, XSTD\*\*2) distribution.

```
CALL RNNOR (NR, R)
CALL SSCAL (NR, XSTD, R, 1)
CALL SADD (NR, XM, R, 1)
```

### Example

In this example, RNNOR is used to generate five pseudorandom deviates from a standard normal distribution.

```
C
INTEGER    ISEED, NOUT, NR
REAL       R(5)
EXTERNAL   RNNOR, RNSSET, UMACH

CALL UMACH (2, NOUT)
NR       = 5
ISEED   = 123457
CALL RNSSET (ISEED)
```

```

CALL RNNOR (NR, R)
WRITE (NOUT,99999) R
99999 FORMAT (' Standard normal random deviates: ', 5F8.4)
END

```

### Output

Standard normal random deviates: 1.8279 -0.6412 0.7266 0.1747 1.0145

## RNSTA/DRNSTA (Single/Double precision)

Generate pseudorandom numbers from a stable distribution.

### Usage

```
CALL RNSTA (NR, ALPHA, BPRIME, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**ALPHA** — Characteristic exponent of the stable distribution. (Input)  
This parameter must be positive and less than or equal to 2.

**BPRIME** — Skewness parameter of the stable distribution. (Input)  
When **BPRIME** = 0, the distribution is symmetric. Unless **ALPHA** = 1, **BPRIME** is not the usual skewness parameter of the stable distribution. **BPRIME** must be greater than or equal to -1 and less than or equal to 1.

**R** — Vector of length **NR** containing the random stable deviates. (Output)

### Comments

The routine **RNSET** (page 1167) can be used to initialize the seed of the random number generator. The routine **RNOPT** (page 1165) can be used to select the form of the generator.

### Algorithm

Routine **RNSTA** generates pseudorandom numbers from a stable distribution with parameters **ALPHA** and **BPRIME**. **ALPHA** is the usual characteristic exponent parameter  $\alpha$  and **BPRIME** is related to the usual skewness parameter  $\beta$  of the stable distribution. With the restrictions  $0 < \alpha \leq 2$  and  $-1 \leq \beta \leq 1$ , the characteristic function of the distribution is

$$\varphi(t) = \exp[-|t|^\alpha \exp(-\pi i \beta (1 - |1 - \alpha| \text{sign}(t)/2))] \quad \text{for } \alpha \neq 1$$

and

$$\varphi(t) = \exp[-|t| (1 + 2i\beta \ln|t|) \text{sign}(t)/\pi] \quad \text{for } \alpha = 1$$

When  $\beta = 0$ , the distribution is symmetric. In this case, if  $\alpha = 2$ , the distribution is normal with mean 0 and variance 2; and if  $\alpha = 1$ , the distribution is Cauchy.

The parameterization using `BPRIME` and the algorithm used here are due to Chambers, Mallows, and Stuck (1976). The relationship between  $BPRIME = \beta'$  and the standard  $\beta$  is

$$\beta' = -\tan(\pi(1 - \alpha)/2) \tan(-\pi\beta(1 - |1 - \alpha|)/2) \quad \text{for } \alpha \neq 1$$

and

$$\beta' = \beta \quad \text{for } \alpha = 1$$

The algorithm involves formation of the ratio of a uniform and an exponential random variate.

### Example

In this example, `RNSTA` is used to generate five pseudorandom symmetric stable variates with characteristic exponent 1.5. The tails of this distribution are heavier than those of a normal distribution, but not so heavy as those of a Cauchy distribution. The variance of this distribution does not exist, however. (This is the case for any stable distribution with characteristic exponent less than 2.)

```

C      INTEGER      NR
      PARAMETER    (NR=5)

C      INTEGER      ISEED, NOUT
      REAL          ALPHA, BPRIM, R(NR)
      EXTERNAL      RNSET, RNSTA, UMACH

C      CALL UMACH (2, NOUT)
      ALPHA = 1.5
      BPRIM = 0.0
      ISEED = 123457
      CALL RNSET (ISEED)
      CALL RNSTA (NR, ALPHA, BPRIM, R)
      WRITE (NOUT,99999) R
99999 FORMAT (' Stable random deviates: ', 5F9.4)
      END

```

### Output

```
Stable random deviates:    4.4091    1.0564    2.5463    5.6724    2.1656
```

---

## RNSTT/DRNSTT (Single/Double precision)

Generate pseudorandom numbers from a Student's  $t$  distribution.

### Usage

```
CALL RNSTT (NR, DF, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**DF** — Degrees of freedom. (Input)  
DF must be positive.

**R** — Vector of length NR containing the random Student's *t* deviates. (Output)

### Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNSTT generates pseudo-random numbers from a Student's *t* distribution with DF degrees of freedom, using a method suggested by Kinderman, Monahan, and Ramage (1977). The method ("TMX" in the reference) involves a representation of the *t* density as the sum of a triangular density over  $(-2, 2)$  and the difference of this and the *t* density. The mixing probabilities depend on the degrees of freedom of the *t* distribution. If the triangular density is chosen, the variate is generated as the sum of two uniforms; otherwise, an acceptance/rejection method is used to generate a variate from the difference density.

For degrees of freedom less than 100, RNSTT requires approximately twice the execution time as routine RNNOA (page 1205) which generates pseudorandom normal deviates. The execution time of RNSTT increases very slowly as the degrees of freedom increase. Since for very large degrees of freedom the normal distribution and the *t* distribution are very similar, the user may find that the difference in the normal and the *t* does not warrant the additional generation time required to use RNSTT instead of RNNOA.

### Example

In this example, RNSTT is used to generate 5 pseudo-random *t* variates with 10 degrees of freedom.

```
C      INTEGER      NR
      PARAMETER   (NR=5)

C      INTEGER      ISEED, NOUT
      REAL         DF, R(NR)
      EXTERNAL     RNSET, RNSTT, UMACH

C      CALL UMACH(2, NOUT)
      DF = 10.0
      ISEED = 123457
      CALL RNSET(ISEED)
      CALL RNSTT(NR, DF, R)
      WRITE(NOUT, 99999) R
99999 FORMAT (' t (10) random deviates: ', 5F8.4)
      END
```

## Output

t (10) random deviates: 0.6152 1.1528 0.0881 1.3382 -0.9893

---

# RNTRI/DRNTRI (Single/Double precision)

Generate pseudorandom numbers from a triangular distribution on the interval (0, 1).

## Usage

CALL RNTRI (NR, R)

## Arguments

**NR** — Number of random numbers to generate. (Input)

**R** — Vector of length NR containing the random triangular deviates. (Output)

## Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNTRI generates pseudorandom numbers from a triangular distribution over the unit interval. The probability density function is  $f(x) = 4x$ , for  $0 \leq x \leq .5$ , and  $f(x) = 4(1 - x)$ , for  $.5 < x \leq 1$ . RNTRI uses an inverse CDF technique.

## Example

In this example, RNTRI is used to generate five pseudorandom deviates from a triangular distribution.

```
INTEGER      ISEED, NOUT, NR
REAL         R(5)
EXTERNAL     RNSET, RNTRI, UMACH
C
CALL UMACH (2, NOUT)
NR         = 5
ISEED     = 123457
CALL RNSET (ISEED)
CALL RNTRI (NR, R)
WRITE (NOUT,99999) R
99999 FORMAT ('      Triangular random deviates: ', 5F8.4)
END
```

## Output

Triangular random deviates: 0.8700 0.3610 0.6581 0.5360 0.7215



---

## RNVMS/DRNVMS (Single/Double precision)

Generate pseudorandom numbers from a von Mises distribution.

### Usage

```
CALL RNVMS (NR, C, R)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**C** — Parameter of the von Mises distribution. (Input)

This parameter must be greater than one half of machine epsilon. (On many machines, the lower bound for  $C$  is  $10^{-3}$ .)

**R** — Vector of length **NR** containing the random von Mises deviates. (Output)

### Comments

The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

### Algorithm

Routine `RNVMS` generates pseudorandom numbers from a von Mises distribution with parameter  $C$ , which must be positive. With  $c = C$ , the probability density function is

$$f(x) = \frac{1}{2\pi I_0(c)} \exp[c \cos(x)] \text{ for } -\pi < x < \pi$$

where  $I_0(c)$  is the modified Bessel function of the first kind of order 0. The probability density equals 0 outside the interval  $(-\pi, \pi)$ .

The algorithm is an acceptance/rejection method using a wrapped Cauchy distribution as the majorizing distribution. It is due to Best and Fisher (1979).

### Example

In this example, `RNVMS` is used to generate five pseudorandom von Mises variates with  $c = 1$ .

```
C
INTEGER      NR
PARAMETER   (NR=5)

C
INTEGER      ISEED, NOUT
REAL         C, R(NR)
EXTERNAL    RNSET, RNVMS, UMACH

C
CALL UMACH (2, NOUT)
C          = 1.0
```

```

ISEED = 123457
CALL RNSET (ISEED)
CALL RNVMS (NR, C, R)
WRITE (NOUT,99999) R
99999 FORMAT (' Von Mises random deviates: ', 5F8.4)
END

```

### Output

Von Mises random deviates: 0.2472 -2.4326 -1.0216 -2.1722 -0.5029

## RNWIB/DRNWIB (Single/Double precision)

Generate pseudorandom numbers from a Weibull distribution.

### Usage

CALL RNWIB (NR, A, R)

### Arguments

**NR** — Number of random numbers to generate. (Input)

**A** — The shape parameter of the Weibull distribution. (Input)  
This parameter must be positive.

**R** — Vector of length NR containing the random Weibull deviates. (Output)

### Comments

1. Informational error  

Type	Code	
3	1	The value of A is so small that the proportion of values from the Weibull that are too large to represent is greater than machine epsilon.
2. The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNWIB generates pseudorandom numbers from a Weibull distribution with shape parameter A and unit scale parameter. The probability density function is

$$f(x) = Ax^{A-1}e^{-x^A} \quad \text{for } x \geq 0$$

Routine RNWIB uses an antithetic inverse CDF technique to generate a Weibull variate; that is, a uniform random deviate  $U$  is generated and the inverse of the Weibull cumulative distribution function is evaluated at  $1.0 - U$  to yield the Weibull deviate.

Deviates from the two-parameter Weibull distribution with shape parameter  $A$  and scale parameter  $B$  can be generated by using `RNWIB` and then multiplying each entry in `R` by  $B$ . The following statements (using routine `SSCAL` (IMSL MATH/LIBRARY) in single precision) would yield random deviates from a two-parameter Weibull distribution.

```
CALL RNWIB (NR, A, R)
CALL SSCAL (NR, B, R, 1)
```

The Rayleigh distribution with probability density function,

$$r(x) = \frac{1}{\alpha^2} x e^{-(x^2/2\alpha^2)} \quad \text{for } x \geq 0$$

is the same as a Weibull distribution with shape parameter  $A$  equal to 2 and scale parameter  $B$  equal to

$$\sqrt{2}\alpha$$

hence, `RNWIB` and `SSCAL` (or simple multiplication) can be used to generate Rayleigh deviates.

### Example

In this example, `RNWIB` is used to generate five pseudorandom deviates from a two-parameter Weibull distribution with shape parameter equal to 2.0 and scale parameter equal to 6.0, a Rayleigh distribution with parameter

$$\alpha = 3\sqrt{2}$$

```

C
INTEGER      ISEED, NOUT, NR
REAL         A, B, R(5)
EXTERNAL     RNSET, RNWIB, SSCAL, UMACH

CALL UMACH (2, NOUT)
A          = 2.0
B          = 6.0
NR         = 5
ISEED     = 123457
CALL RNSET (ISEED)
CALL RNWIB (NR, A, R)
CALL SSCAL (NR, B, R, 1)
WRITE (NOUT,99999) R
99999 FORMAT ('          Weibull(2,6) random deviates: ', 5F8.4)
END
```

### Output

```
Weibull(2,6) random deviates:  1.1122  6.9568  3.0959  4.5031  2.4638
```

---

## RNCOR/DRNCOR (Single/Double precision)

Generate a pseudorandom orthogonal matrix or a correlation matrix.

## Usage

CALL RNCOR (N, IOPT, EV, A, LDA, COR, LDCOR)

## Arguments

*N* — The order of the matrices to be generated. (Input)

*N* must be at least two.

*IOPT* — Option indicator. (Input)

### **IOPT**    **Action**

0        A random orthogonal matrix is generated in *A*.

1        A random correlation matrix is generated in *COR*. (*A* is used as workspace.)

2        A random correlation matrix is generated in *COR* using the orthogonal matrix input in *A*.

*EV* — If *IOPT* = 1 or 2, a vector of length *N* containing the eigenvalues of the correlation matrix to be generated. (Input, if *IOPT* = 1 or 2; not used otherwise.) The elements of *EV* must be positive, they must sum to *N*, and they cannot all be equal.

*A* — *N* by *N* random orthogonal matrix. (Output, if *IOPT* = 0; workspace if *IOPT* = 1; input/output, if *IOPT* = 2. If *IOPT* = 2, *A* is destroyed.)

*LDA* — Leading dimension of *A* exactly as specified in the dimension statement in the calling program. (Input)

*COR* — *N* by *N* random correlation matrix. (Output, if *IOPT* = 1 or 2; not used otherwise.)

*LDCOR* — Leading dimension of *COR* exactly as specified in the dimension statement in the calling program. (Input)

## Comments

1.        Automatic workspace usage is

RNCOR    4 \* *N* units, or

DRNCOR  5 \* *N* units.

Workspace may be explicitly provided, if desired, by use of

R2COR/DR2COR. The reference is

```
CALL R2COR (N, IOPT, EV, A, LDA, COR, LDCOR, IWK,  
            WK)
```

The additional arguments are as follows:

*IWK* — Work vector of length 3 \* *N*.

*WK* — Work vector of length *N*.

2.        Informational error

Type    Code

- 3            1            Considerable loss of precision occurred in the rotations used to form the correlation matrix. Some of the diagonals of COR differ from 1.0 by more than the machine epsilon.
3.            The routine RNSSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNCOR generates a pseudorandom orthogonal matrix A from the invariant Haar measure. For each column of A, a random vector from a uniform distribution on a hypersphere is selected and then is projected onto the orthogonal complement of the columns of A already formed. The method is described by Heiberger (1978). (See also Tanner and Thisted 1982.)

A correlation matrix is formed by applying a sequence of planar rotations to the matrix  $A^T D A$ , where  $D = \text{diag}(\text{EV}(1), \dots, \text{EV}(N))$ , so as to yield ones along the diagonal. The planar rotations are applied in such an order that in the two by two matrix that determines the rotation, one diagonal element is less than 1.0 and one is greater than 1.0. This method is discussed by Bendel and Mickey (1978) and by Lin and Bendel (1985).

The distribution of the correlation matrices produced by this method is not known. Bendel and Mickey (1978) and Johnson and Welch (1980) discuss the distribution.

For larger matrices, rounding can become severe; and the double precision results may differ significantly from single precision results.

### Example

In this example, RNCOR is used to generate a 4 by 4 pseudorandom correlation matrix with eigenvalues in the ratio 1:2:3:4. (Note that the eigenvalues must sum to 4.) Routines MXTXF (IMSL MATH/LIBRARY) and EVCSF (IMSL MATH/LIBRARY) are used to check the output.

```

INTEGER      I, IOPT, ISEED, J, LDA, LDCOR, N, NOUT
REAL         A(4,4), COR(4,4), EV(4), EVAL(4), EVEC(4,4), FLOAT,
&            SSUM, SUM, XID(4,4)
INTRINSIC    FLOAT
EXTERNAL     EVCSF, MXTXF, RNCOR, RNSSET, SSCAL, SSUM, UMACH

C
CALL UMACH (2, NOUT)
N          = 4
LDA        = 4
LDCOR      = 4
EV(1)      = 1.0
EV(2)      = 2.0
EV(3)      = 3.0
EV(4)      = 4.0

C                               Scale the eigenvalues to sum to N.
```

```

SUM = SSUM(N,EV,1)
CALL SSCAL (N, FLOAT(N)/SUM, EV, 1)
ISEED = 123457
CALL RNSET (ISEED)
C
C          Generate an orthogonal matrix.
IOPT = 0
CALL RNCOR (N, IOPT, EV, A, LDA, COR, LDCOR)
WRITE (NOUT,99996) ((A(I,J),J=1,N),I=1,N)
99996 FORMAT (' A random orthogonal matrix: ', /, (5X,4F8.4))
C          Check it for orthogonality.
CALL MXTXF (N, N, A, LDA, N, XID, LDA)
WRITE (NOUT,99997) ((XID(I,J),J=1,N),I=1,N)
99997 FORMAT (' The identity matrix?: ', /, (5X,4F8.4))
C
C          Now get a correlation matrix using
C          the orthogonal matrix in A, which
C          will be destroyed.
IOPT = 2
CALL RNCOR (N, IOPT, EV, A, LDA, COR, LDCOR)
WRITE (NOUT,99998) ((COR(I,J),J=1,N),I=1,N)
99998 FORMAT (' A random correlation matrix: ', /, (5X,4F8.4))
C          Check the eigenvalues.
CALL EVCSF (N, COR, LDCOR, EVAL, EVEC, LDCOR)
WRITE (NOUT,99999) (EVAL(I),I=1,N)
99999 FORMAT (' The computed eigenvalues:', 4F8.4)
END

```

### Output

```

A random orthogonal matrix:
-0.8804 -0.2417  0.4065 -0.0351
 0.3088 -0.3002  0.5520  0.7141
-0.3500  0.5256 -0.3874  0.6717
-0.0841 -0.7584 -0.6165  0.1941
The identity matrix?:
 1.0000  0.0000  0.0000  0.0000
 0.0000  1.0000  0.0000  0.0000
 0.0000  0.0000  1.0000  0.0000
 0.0000  0.0000  0.0000  1.0000
A random correlation matrix:
 1.0000 -0.2358 -0.3258 -0.1101
-0.2358  1.0000  0.1906 -0.0172
-0.3258  0.1906  1.0000 -0.4353
-0.1101 -0.0172 -0.4353  1.0000
The computed eigenvalues:  1.6000  1.2000  0.8000  0.4000

```

---

## RNDAT/DRNDAT (Single/Double precision)

Generate pseudorandom numbers from a multivariate distribution determined from a given sample.

### Usage

```
CALL RNDAT (IDO, NR, K, NSAMP, X, LDX, NN, R, LDR)
```

## Arguments

**IDO** — Generator option. (Input)

<b>IDO</b>	<b>Action</b>
0	This is the only invocation of <code>RNDAT</code> with the sample in <code>X</code> and all desired pseudorandom numbers are to be generated in this call.
1	This is the first invocation, and additional calls to <code>RNDAT</code> will be made to generate additional random numbers using the same given sample.
2	This is an intermediate invocation of <code>RNDAT</code> . The work vectors have been set up in a previous call, but they are not to be released because additional calls will be made.
3	This is the final invocation of <code>RNDAT</code> . The work vectors have been set up in a previous call and they are to be released.

**NR** — Number of random multivariate vectors to generate. (Input)

If `NR = 0`, only initialization or wrap up operations are performed. (This would make sense only if `IDO = 1` or `3`.)

**K** — The length of the multivariate vectors, that is, the number of dimensions. (Input)

**NSAMP** — Number of given data points from the distribution to be simulated. (Input)

**X** — `NSAMP` by `K` matrix containing the given sample. (Input/Output)

If `IDO = 0` or `1`, on output the rows of `X` are rearranged by routine `QUADT` to form a  $k$ - $d$  tree.

**LDX** — Leading dimension of `X` exactly as specified in the dimension statement in the calling program. (Input)

**NN** — Number of nearest neighbors of the randomly selected point in `X` that are used to form the output point in `R`. (Input)

**R** — `NR` by `K` matrix containing the random multivariate vectors in its rows. (Output)

**LDR** — Leading dimension of `R` exactly as specified in the dimension statement of the calling program. (Input)

## Comments

1. Automatic workspace usage is

`RNDAT`  $4 * NSAMP + 3 * LEN + 2 * K * LEN + 2 * K + 3 * NN$  units, or  
`DRNDAT`  $6 * NSAMP + 3 * LEN + 4 * K * LEN + 3 * K + 5 * NN$  units.

Here, `LEN` is the greatest integer in  $\ln(NSAMP)/\ln(2) + 4$ . Workspace may be explicitly provided, if desired, by use of `R2DAT/DR2DAT`. The reference is

```
CALL R2DAT (IDO, NR, K, NSAMP, X, LDX, NN, R, LDR,  
           IWK, WK)
```

The additional arguments are as follows:

**IWK** — Work vector of length equal to  $2 * \text{NSAMP} + 3 * \text{LEN} + \text{K} + \text{NN}$ .

**WK** — Work vector of length equal to  $2 * \text{NSAMP} + 2 * \text{K} * \text{LEN} + \text{K} + 2 * \text{NN}$ .

R2DAT allows alternating calls for two different populations (see Comment 3). **Warning:** R2DAT does no error checking.

2. The rows of X are rearranged on output from either RNDAT or R2DAT.
3. When more than one call is to be made to RNDAT to generate more than one R matrix using the same sample in X, IDO should be set to 1 for the first call, to 2 for all subsequent calls except the last one, and to 3 for the last call. If more than one population is to be simulated (that is, there is more than one sample, X), it is necessary to generate all of the observations from each population at one time because data is stored in the work vectors. If the user provides work vectors for each population to be simulated, R2DAT can be used to simulate different population alternatively.
4. The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Given a sample of size  $n$  ( $= \text{NSAMP}$ ) of observations of a  $k$ -variate random variable, RNDAT generates a pseudorandom sample with approximately the same moments as the given sample. The sample obtained is essentially the same as if sampling from a Gaussian kernel estimate of the sample density. (See Thompson 1989.) Routine RNDAT uses methods described by Taylor and Thompson (1986).

Assume that the (vector-valued) observations  $x_i$  are in the rows of X. An observation,  $x_j$ , is chosen randomly; its nearest  $m$  ( $= \text{NN}$ ) neighbors,

$$x_{j_1}, x_{j_2}, \dots, x_{j_m}$$

are determined; and the mean

$$\bar{x}_j$$

of those nearest neighbors is calculated. Next, a random sample

$u_1, u_2, \dots, u_m$  is generated from a uniform distribution with lower bound

$$\frac{1}{m} - \sqrt{\frac{3(m-1)}{m^2}}$$

and upper bound



$$\frac{1}{m} + \sqrt{\frac{3(m-1)}{m^2}}$$

The random variate delivered is

$$\sum_{l=1}^m u_l (x_{jl} - \bar{x}_j) + \bar{x}_j$$

The process is then repeated until NR such simulated variates are generated and stored in the rows of R.

When RNDAT is invoked for the first time for a given sample, a search tree is computed for the rows of X. During the generation process, this tree is used to find the nearest neighbors of the randomly selected row. The argument IDO is used to determine whether or not the tree must be computed and whether workspace has to be allocated to store the tree.

### Example

In this example, RNDAT is used to generate 5 pseudorandom vectors of length 4 using the initial and final systolic pressure and the initial and final diastolic pressure from Data Set A in Afifi and Azen (1979) as the fixed sample from the population to be modeled. (Values of these four variables are in the seventh, tenth, twenty-first, and twenty-fourth columns of data set number nine in routine GDATA, page 1302)

```

INTEGER      LDR, LDRDAT, LDX, NDR, NDRDAT, NDX
PARAMETER   (LDR=5, LDRDAT=113, LDX=113, NDR=4, NDRDAT=34, NDX=4)
C
INTEGER      IDO, ISEED, K, NN, NR, NRCOL, NRROW, NSAMP
REAL         R(LDR,NDR), RDATA(LDRDAT,NDRDAT), X(LDX,NDX)
EXTERNAL     GDATA, RNDAT, RNSET, SCOPY, WRRRL
C
CALL GDATA (9, 0, NRROW, NRCOL, RDATA, LDRDAT, NDRDAT)
CALL SCOPY (NRROW, RDATA(1,7), 1, X(1,1), 1)
CALL SCOPY (NRROW, RDATA(1,10), 1, X(1,2), 1)
CALL SCOPY (NRROW, RDATA(1,21), 1, X(1,3), 1)
CALL SCOPY (NRROW, RDATA(1,24), 1, X(1,4), 1)
C
ISEED = 123457
CALL RNSET (ISEED)
C
                                Set input values
IDO   = 0
NR    = 5
K     = 4
NSAMP = 113
NN    = 5
C
                                Generate random variates
CALL RNDAT (IDO, NR, K, NSAMP, X, LDX, NN, R, LDR)
C
                                Print results
CALL WRRRL ('Random variates', NR, K, R, LDR, 0, '(F15.4)',
&          'NUMBER', 'NUMBER')
C
END

```

## Output

	Random variates			
	1	2	3	4
1	162.7668	90.5057	153.7173	104.8768
2	153.3533	78.3180	176.6643	85.2155
3	93.6958	48.1675	153.5495	71.3688
4	101.7508	54.1855	113.1215	56.2916
5	91.7403	58.7684	48.4368	28.0994

---

## RNMTN

Generate pseudorandom numbers from a multinomial distribution.

### Usage

CALL RNMTN (NR, N, K, P, IR, LDIR)

### Arguments

**NR** — Number of random multinomial vectors to generate. (Input)

**N** — Multinomial parameter indicating the number of independent trials. (Input)

**K** — The number of mutually exclusive outcomes on any trial. (Input)

**K** is the length of the multinomial vectors. **K** must be greater than or equal to 2.

**P** — Vector of length **K** containing the probabilities of the possible outcomes.

(Input)

The elements of **P** must be positive and must sum to 1.0.

**I** — **NR** by **K** matrix containing the random multinomial vectors in its rows.

(Output)

**LDIR** — Leading dimension of **IR** exactly as specified in the dimension statement of the calling program. (Input)

### Comments

The routine **RNSET** (page 1167) can be used to initialize the seed of the random number generator. The routine **RNOPT** (page 1165) can be used to select the form of the generator.

### Algorithm

Routine **RNMTN** generates pseudorandom numbers from a **K**-variate multinomial distribution with parameters **N** and **P**. **K** and **N** must be positive. Each element of **P** must be positive and the elements must sum to 1. The probability function (with  $n = N$ ,  $k = K$ , and  $p_i = P(I)$ ) is

$$f(x_1, x_2, \dots, x_k) = \frac{n!}{x_1! x_2! \dots x_k!} p_1^{x_1} p_2^{x_2} \dots p_k^{x_k}$$

for  $x_i \geq 0$  and

$$\sum_{i=1}^k x_i = n$$

The deviate in each row of IR is produced by generation of the binomial deviate  $x_1$  with parameters  $n$  and  $p_1$  and then by successive generations of the conditional binomial deviates  $x_j$  given  $x_1, x_2, \dots, x_{j-1}$  with parameters  $n - x_1 - x_2 - \dots - x_{j-1}$  and  $p_j / (1 - p_1 - p_2 - \dots - p_{j-1})$ .

### Example

In this example, RNMTN is used to generate five pseudorandom 3-dimensional multinomial variates with parameters  $N = 20$  and  $P = (0.1, 0.3, 0.6)$ .

```
INTEGER      K, LDIR
PARAMETER   (K=3, LDIR=5)
C
INTEGER      I, IR(LDIR,K), ISEED, J, N, NOUT, NR
REAL         P(K)
EXTERNAL     RNMTN, RNSET, UMACH
C
CALL UMACH (2, NOUT)
N           = 20
P(1)       = 0.1
P(2)       = 0.3
P(3)       = 0.6
NR         = 5
ISEED      = 123457
CALL RNSET (ISEED)
CALL RNMTN (NR, N, K, P, IR, LDIR)
WRITE (NOUT,99999) ((IR(I,J),J=1,K),I=1,NR)
99999 FORMAT (' Multinomial random deviates: ', 3I4, /, (30X,3I4))
END
```

### Output

```
Multinomial random deviates:   5   4  11
                                3   6  11
                                3   3  14
                                5   5  10
                                4   5  11
```

---

## RNMVN/DRNMVN (Single/Double precision)

Generate pseudorandom numbers from a multivariate normal distribution.

### Usage

```
CALL RNMVN (NR, K, RSIG, LDRSIG, R, LDR)
```

## Arguments

**NR** — Number of random multivariate normal vectors to generate. (Input)

**K** — Length of the multivariate normal vectors. (Input)

**RSIG** — Upper triangular matrix,  $K$  by  $K$ , containing the Cholesky factor of the variance-covariance matrix. (Input)

The variance-covariance matrix is equal to the product of the transpose of **RSIG** and **RSIG**. **RSIG** can be obtained from the variance-covariance matrix using routine **CHFAC**.

**LDRSIG** — Leading dimension of **RSIG** exactly as specified in the dimension statement in the calling program. (Input)

**R** —  $NR$  by  $K$  matrix containing the random multivariate normal vectors in its rows. (Output)

**LDR** — Leading dimension of **R** exactly as specified in the dimension statement of the calling program. (Input)

## Comments

The routine **RNSET** (page 1167) can be used to initialize the seed of the random number generator. The routine **RNOPT** (page 1165) can be used to select the form of the generator.

## Algorithm

Routine **RNMVN** generates pseudorandom numbers from a multivariate normal distribution with mean vector consisting of all zeroes and variance-covariance matrix whose Cholesky factor (or “square root”) is **RSIG**; that is, **RSIG** is an upper triangular matrix such that the transpose of **RSIG** times **RSIG** is the variance-covariance matrix. First, independent random normal deviates with mean 0 and variance 1 are generated, and then the matrix containing these deviates is postmultiplied by **RSIG**. The independent normals are generated into the columns of a matrix, which has  $NR$  rows; hence, if **RNSET** is called with different values of  $NR$ , the output is different even if the seed is the same in the calls.

Deviates from a multivariate normal distribution with means other than zero can be generated by using **RNMVN** and then by adding the vector of means to each row of **R**.

## Example

In this example, **RNMVN** is used to generate five pseudorandom multivariate normal vectors of length 2 with variance-covariance matrix equal to

$$\begin{array}{cc} 0.500 & 0.375 \\ 0.375 & 0.500 \end{array}$$

The routine CHFAC is first called to compute the Cholesky factorization of the variance-covariance matrix.

```

INTEGER      I, IRANK, ISEED, J, K, LDR, LDRSIG, NOUT, NR
REAL         COV(2,2), R(5,2), RSIG(2,2)
EXTERNAL     CHFAC, RNMVN, RNSET, UMACH
C
CALL UMACH (2, NOUT)
NR          = 5
K           = 2
LDRSIG      = 2
LDR         = 5
COV(1,1)    = 0.5
COV(1,2)    = 0.375
COV(2,1)    = 0.375
COV(2,2)    = 0.5
C
CALL CHFAC (K, COV, 2, 0.00001, IRANK, RSIG, LDRSIG)
C
C           Obtain the Cholesky factorization.
C           Initialize seed of random number
C           generator.
ISEED = 123457
CALL RNSET (ISEED)
CALL RNMVN (NR, K, RSIG, LDRSIG, R, LDR)
WRITE (NOUT,99999) ((R(I,J),J=1,K),I=1,NR)
99999 FORMAT ('   Multivariate normal random deviates: ', /,
&            (1X,2F8.4))
END

```

### Output

```

Multivariate normal random deviates:
1.4507  1.2463
0.7660 -0.0429
0.0584 -0.6692
0.9035  0.4628
-0.8669 -0.9334

```

---

## RNSPH/DRNSPH (Single/Double precision)

Generate pseudorandom points on a unit circle or  $K$ -dimensional sphere.

### Usage

```
CALL RNSPH (NR, K, Z, LDZ)
```

### Arguments

**NR** — Number of random numbers to generate. (Input)

**K** — Dimension of the circle ( $K = 2$ ) or of the sphere. (Input)

**Z** —  $NR$  by  $K$  matrix containing the random Cartesian coordinates on the unit circle or sphere. (Output)

**LDZ** — Leading dimension of  $Z$  exactly as specified in the dimension statement of the calling program. (Input)

## Comments

The routine RNSSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNSPH generates pseudorandom coordinates of points that lie on a unit circle or a unit sphere in  $\kappa$ -dimensional space. For points on a circle ( $\kappa = 2$ ), pairs of uniform  $(-1, 1)$  points are generated and accepted only if they fall within the unit circle (the sum of their squares is less than 1), in which case they are scaled so as to lie on the circle.

For spheres in three or four dimensions, the algorithms of Marsaglia (1972) are used. For three dimensions, two independent uniform  $(-1, 1)$  deviates  $U_1$  and  $U_2$  are generated and accepted only if the sum of their squares  $S_1$  is less than 1. Then, the coordinates

$$Z_1 = 2U_1\sqrt{1-S_1}, Z_2 = 2U_2\sqrt{1-S_1}, \text{ and } Z_3 = 1-2S_1$$

are formed. For four dimensions,  $U_1, U_2$ , and  $S_1$  are produced as described above. Similarly,  $U_3, U_4$ , and  $S_2$  are formed. The coordinates are then

$$Z_1 = U_1, Z_2 = U_2, Z_3 = U_3\sqrt{(1-S_1)/S_2}$$

and

$$Z_4 = U_4\sqrt{(1-S_1)/S_2}$$

For spheres in higher dimensions,  $\kappa$  independent normal deviates are generated and scaled so as to lie on the unit sphere in the manner suggested by Muller (1959).

## Example

In this example, RNSPH is used to generate two uniform random deviates from the surface of the unit sphere in three space.

```
INTEGER      K, LDZ
PARAMETER   (K=3, LDZ=2)
C
INTEGER      I, ISEED, J, NOUT, NR
REAL         Z(LDZ,K)
EXTERNAL    RNSSET, RNSPH, UMACH
C
CALL UMACH (2, NOUT)
NR          = 2
ISEED      = 123457
CALL RNSSET (ISEED)
CALL RNSPH (NR, K, Z, LDZ)
WRITE (NOUT,99999) ((Z(I,J),J=1,K),I=1,NR)
99999 FORMAT ('      Coordinates of first point: ', 3F8.4, /,
```

```
&          '      Coordinates of second point:', 3F8.4)
END
```

### Output

```
Coordinates of first point:   0.8893  0.2316  0.3944
Coordinates of second point:  0.1901  0.0396 -0.9810
```

---

## RNTAB

Generate a pseudorandom two-way table.

### Usage

```
CALL RNTAB (IDO, NROW, NCOL, NRTOT, NCTOT, ITAB, LDITAB)
```

### Arguments

**IDO** — Generator option. (Input)

<b>IDO</b>	<b>Action</b>
------------	---------------

- |   |   |
|---|---|
| 0 | This is the only invocation of RNTAB with these input specifications of the two-way table.  |
| 1 | This is the first invocation, and additional calls to RNTAB will be made to generate random tables with the same specifications.  |
| 2 | This is an intermediate invocation of RNTAB. The work vectors have been set up in a previous call, but they are not to be released because additional calls will be made. |
| 3 | This is the final invocation of RNTAB. The work vectors have been set up in a previous call and they are to be released.  |

**NROW** — Number of rows in the table. (Input)

**NCOL** — Number of columns in the table. (Input)

**NRTOT** — Vector of length NROW containing the row totals. (Input)

**NCTOT** — Vector of length NCOL containing the column totals. (Input)  
The elements of NRTOT and NCTOT must be nonnegative and must sum to the same quantity.

**ITAB** — NROW by NCOL random matrix with the given row and column totals. (Output)

**LDITAB** — Leading dimension of ITAB exactly as specified in the dimension statement in the calling program. (Input)

### Comments

- Let  $IRSUM =$  the sum of the elements in NRTOT. If  $IRSUM + 1$  is less than  $2 * NROW * NCOL$ , automatic workspace usage is  $IRSUM$ ; otherwise, automatic workspace usage is  $2 * IRSUM + 1$  because a different algorithm is used. Workspace may be explicitly provided, if

desired, by use of `R2TAB`. `R2TAB` allows selection of the algorithm to be used and it allows alternating calls for two different problems (see Comment 3). The reference is

```
CALL R2TAB (IDO, NROW, NCOL, NRTOT, NCTOT, ITAB,  
           LDITAB, IOPT, IRSUM, IWK, WK).
```

The additional arguments are as follows:

***IOPT*** — Option indicator. (Input)

If `IOPT = 1`, Boyette's method is used.

If `IOPT = 2`, Patefield's method is used.

***IRSUM*** — Sum of the elements in `NRTOT`. (Output)

***IWK*** — Work vector of length equal to the sum of the elements in `NRTOT`.

***WK*** — Work vector of length equal to the sum of the elements in `NRTOT` plus one, used only if `IOPT = 2`.

WARNING: `R2TAB` does no error checking.

2. Informational error  
Type Code  
3 1 The values of `NRTOT` and/or of `NCTOT` are such that the probability distribution of tables is degenerate, that is, only one such table is possible.
3. When more than one table with the same marginal totals is to be generated, `IDO` should be set to 1 for the first call, to 2 for all subsequent calls except the last one, and to 3 for the last call. If several tables of different sizes or with different marginal totals are to be generated, it is necessary to generate all of each type together because of the data stored in the work vectors. If the user provides work vectors for each type of table to be generated, `R2TAB` can be used to generate different types of tables alternatively.
4. The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

### Algorithm

Routine `RNTAB` generates pseudorandom entries for a two-way contingency table with fixed row and column totals. The method depends on the size of the table and the total number of entries in the table. If the total number of entries is less than twice the product of the number of rows and columns, the method described by Boyette (1979) and by Agresti, Wackerly, and Boyette (1979) is used. In this method, a work vector is filled with row indices so that the number of times each index appears equals the given row total. This vector is then randomly permuted and used to increment the entries in each row so that the given row total is attained.



For tables with larger numbers of entries, the method of Patefield (1981) is used. This method can be considerably faster in these cases. The method depends on the conditional probability distribution of individual elements, given the entries in the previous rows. The probabilities for the individual elements are computed starting from their conditional means.

On the first call to RNTAB with a given set of row and column totals, certain checking is done, and the work vector is allocated and initialized. On the final call, the work vector is released. The argument IDO indicates the nature of the call. In a simulation study, RNTAB would typically be called first with IDO = 1, then would be called several times with IDO = 2, and then finally would be called with IDO = 3. If only one table is needed, IDO should be set to 0.

### Example

In this example, RNTAB is used to generate a two by three table with row totals 3 and 5, and column totals 2, 4, and 2.

```

INTEGER      I, IDO, ISEED, ITAB(2,3), IWK, J, LDITAB, NCOL,
&            NCTOT(3), NOUT, NROW, NRTOT(2)
EXTERNAL     RNSET, RNTAB, UMACH
C
CALL UMACH (2, NOUT)
IDO          = 0
NROW        = 2
NCOL        = 3
NRTOT(1)    = 3
NRTOT(2)    = 5
NCTOT(1)    = 2
NCTOT(2)    = 4
NCTOT(3)    = 2
LDITAB      = 2
ISEED       = 123457
CALL RNSET (ISEED)
CALL RNTAB (IDO, NROW, NCOL, NRTOT, NCTOT, ITAB, LDITAB)
WRITE (NOUT,99999) ((ITAB(I,J),J=1,NCOL),I=1,NROW)
99999 FORMAT (' A random contingency table with fixed marginal totals:'
&            , /, (5X,3I5))
END

```

### Output

A random contingency table with fixed marginal totals:

```

  0   2   1
  2   2   1

```

---

## RNNOS/DRNNOS (Single/Double precision)

Generate pseudorandom order statistics from a standard normal distribution.

### Usage

```
CALL RNNOS (IFIRST, ILAST, N, R)
```

## Arguments

**IFIRST** — First order statistic to generate. (Input)

**ILAST** — Last order statistic to generate. (Input)

ILAST must be greater than or equal to IFIRST. The full set of order statistics from IFIRST to ILAST is generated. If only one order statistic is desired, set ILAST = IFIRST.

**N** — Size of the sample from which the order statistics arise. (Input)

**R** — Vector of length ILAST + 1 - IFIRST containing the random order statistics in ascending order. (Output)

The first element of R is the IFIRST-th order statistic in a random sample of size N from the standard normal distribution.

## Comments

The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

## Algorithm

Routine RNNOS generates the IFIRST through the ILAST order statistics from a pseudorandom sample of size N from a normal (0, 1) distribution. Routine RNNOS uses the routine RNUNO (page 1231) to generate order statistics from the uniform (0, 1) distribution and then obtains the normal order statistics using the inverse CDF transformation.

Each call to RNNOS yields an independent event so order statistics from different calls may not have the same order relations with each other.

## Example

In this example, RNNOS is used to generate the fifteenth through the nineteenth order statistics from a sample of size twenty.

```
INTEGER    IFIRST, ILAST, ISEED, N, NOUT
REAL      R(5)
EXTERNAL  RNNOS, RNSET, UMACH
C
CALL UMACH (2, NOUT)
IFIRST = 15
ILAST  = 19
N      = 20
C                                     Initialize seed of random number
C                                     generator.
ISEED = 123457
CALL RNSET (ISEED)
CALL RNNOS (IFIRST, ILAST, N, R)
WRITE (NOUT,99999) R
99999 FORMAT (' The 15th through the 19th order statistics from a',
&           /, ' random sample of size 20 from a normal distribution'
&           , /, 5F8.4)
```

END

### Output

The 15th through the 19th order statistics from a random sample of size 20 from a normal distribution  
0.4056 0.4681 0.4697 0.9067 0.9362

---

## RNUNO/DRNUNO (Single/Double precision)

Generate pseudorandom order statistics from a uniform (0, 1) distribution.

### Usage

```
CALL RNUNO (IFIRST, ILAST, N, R)
```

### Arguments

**IFIRST** — First order statistic to generate. (Input)

**ILAST** — Last order statistic to generate. (Input)

**ILAST** must be greater than or equal to **IFIRST**. The full set of order statistics from **IFIRST** to **ILAST** is generated. If only one order statistic is desired, set **ILAST** = **IFIRST**.

**N** — Size of the sample from which the order statistics arise. (Input)

**R** — Vector of length **ILAST** + 1 - **IFIRST** containing the random order statistics in ascending order. (Output)

The first element of **R** is the **IFIRST**-th order statistic in a random sample of size **N** from the uniform (0, 1) distribution.

### Comments

The routine **RNSET** (page 1167) can be used to initialize the seed of the random number generator. The routine **RNOPT** (page 1165) can be used to select the form of the generator.

### Algorithm

Routine **RNUNO** generates the **IFIRST** through the **ILAST** order statistics from a pseudorandom sample of size **N** from a uniform (0, 1) distribution. Depending on the values of **IFIRST** and **ILAST**, different methods of generation are used to achieve greater efficiency. If **IFIRST** = 1 and **ILAST** = **N**, that is, if the full set of order statistics are desired, the spacings between successive order statistics are generated as ratios of exponential variates. If the full set is not desired, a beta variate is generated for one of the order statistics, and the others are generated as extreme order statistics from conditional uniform distributions. Extreme order statistics from a uniform distribution can be obtained by raising a uniform deviate to an appropriate power.

Each call to `RNUNO` yields an independent event. This means, for example, that if on one call the fourth order statistic is requested and on a second call the third order statistic is requested, the “fourth” may be smaller than the “third”. If both the third and fourth order statistics from a given sample are desired, they should be obtained from a single call to `RNUNO` (by specifying `IFIRST` less than or equal to 3 and `ILAST` greater than or equal to 4).

### Example

In this example, `RNUNO` is used to generate the fifteenth through the nineteenth order statistics from a sample of size twenty.

```

INTEGER      IFIRST, ILAST, ISEED, N, NOUT
REAL         R(5)
EXTERNAL     RNSET, RNUNO, UMACH
C
CALL UMACH (2, NOUT)
IFIRST = 15
ILAST  = 19
N      = 20
C
C                                     Initialize seed of random number
C                                     generator.
ISEED = 123457
CALL RNSET (ISEED)
CALL RNUNO (IFIRST, ILAST, N, R)
WRITE (NOUT,99999) R
99999 FORMAT (' The 15th through the 19th order statistics from a ',
&           /, ' random sample of size 20 from a uniform ',
&           'distribution', /, 5F8.4)
END

```

### Output

```

The 15th through the 19th order statistics from a
random sample of size 20 from a uniform distribution
0.6575  0.6802  0.6807  0.8177  0.8254

```

---

## RNARM/DRNARM (Single/Double precision)

Generate a time series from a specified ARMA model.

### Usage

```

CALL RNARM (NW, CONST, NPAR, PAR, LAGAR, NPMA, PMA, LAGMA,
            IADIST, AVAR, A, WI, W)

```

### Arguments

**NW** — Number of observations of the time series to generate. (Input)  
**NW** must be greater than or equal to one.

**CONST** — Overall constant. (Input)  
See Comments.

**NPAR** — Number of autoregressive parameters. (Input)  
NPAR must be greater than or equal to zero.

**PAR** — Vector of length NPAR containing the autoregressive parameters. (Input)

**LAGAR** — Vector of length NPAR containing the order of the autoregressive parameters. (Input)  
The elements of LAGAR must be greater than or equal to one.

**NPMA** — Number of moving average parameters. (Input)  
NPMA must be greater than or equal to zero.

**PMA** — Vector of length NPMA containing the moving average parameters. (Input)

**LAGMA** — Vector of length NPMA containing the order of the moving average parameters. (Input)  
The elements of LAGMA must be greater than or equal to one.

**IADIST** — Option for normally distributed innovations. (Input)

**IADIST Action**

- 0 Innovations are generated from a normal distribution (white noise) with mean 0 and variance AVAR.
- 1 Innovations are specified by the user.

**AVAR** — Variance of the normal distribution, if used. (Input)  
For IADIST = 0, AVAR is input; and for IADIST = 1, AVAR is unused.

**A** — Vector of length NW + max(LAGMA(j)) containing the innovations. (Input or output)  
For IADIST = 1, A is input; and for IADIST = 0, A is output.

**WI** — Vector of length max(LAGAR(i)) containing the initial values of the time series. (Input)

**W** — Vector of length NW containing the generated time series. (Output)

**Comments**

1. The time series is generated according to the following model:

$$X(i) = \text{CONST} + \text{PAR}(1) * X(i - \text{LAGAR}(1)) + \dots + \text{PAR}(\text{NPAR}) * X(i - \text{LAGAR}(\text{NPAR})) + A(i) - \text{PMA}(1) * A(i - \text{LAGMA}(1)) - \dots - \text{PMA}(\text{NPMA}) * A(i - \text{LAGAR}(\text{NPMA}))$$

where

$$X(t) = W(t), \quad t = 1, 2, \dots, \text{NW}$$

and

$$W(t) = \text{WI}(t + p), \quad t = 1 - p, 2 - p, \dots, -1, 0$$

with  $p = \max(\text{LAGAR}(k))$ .

The constant is related to the mean of the series, WMEAN, as follows:

$$\text{CONST} = \text{WMEAN} * (1 - \text{PAR}(1) - \dots - \text{PAR}(\text{NPAR}))$$

2. Time series whose innovations have a nonnormal distribution may be simulated by setting IADIST = 1 and by providing the appropriate innovations in A and start values in WI.
3. The routine RNSET (page 1167) can be used to initialize the seed of the random number generator. The routine RNOPT (page 1165) can be used to select the form of the generator.

### Algorithm

Routine RNARM simulates an ARMA( $p, q$ ) process,  $\{W_t\}$  for  $t = 1, 2, \dots, n$  (with  $n = \text{NW}$ ,  $p = \text{NPAR}$ , and  $q = \text{NPMA}$ ). The model is

$$\phi(B)W_t = \theta_0 + \theta(B)A_t \quad t \in \mathbb{Z}$$

where  $B$  is the backward shift operator,

$$\phi(B) = 1 - \phi_1 B - \phi_2 B^2 - \dots - \phi_p B^p$$

$$\theta(B) = 1 - \theta_1 B - \theta_2 B^2 - \dots - \theta_q B^q$$

Let  $\mu$  be the mean of the time series  $\{W_t\}$ . The overall constant  $\theta_0$  (CONST) is

$$\theta_0 = \begin{cases} \mu & p = 0 \\ \mu(1 - \sum_{i=1}^p \phi_i) & p > 0 \end{cases}$$

### Example 1

In this example, RNARM is used to generate a time series of length five, using an ARMA model with three autoregressive parameters and two moving average parameters. The start values are 0.1000, 0.0500, and 0.0375.

```

C      INTEGER      NPAR, NPMA, NW
      PARAMETER    (NPAR=3, NPMA=2, NW=5)

C      INTEGER      I, IADIST, ISEED, LAGAR(NPAR), LAGMA(NPMA), NOUT
      REAL          A(NW+2), AVAR, CONST, PAR(NPAR), PMA(NPMA), W(NW),
&                WI(3)
C      EXTERNAL    RNARM, RNSET, UMACH

      CALL UMACH (2, NOUT)
      LAGAR(1) = 1
      LAGAR(2) = 2
      LAGAR(3) = 3
      PAR(1)   = 0.500
      PAR(2)   = 0.250
      PAR(3)   = 0.125
      LAGMA(1) = 1
      LAGMA(2) = 2
      PMA(1)   = -0.500

```

```

PMA(2) = -0.250
IADIST = 0
CONST = 1.0
AVAR = 0.1
WI(1) = 0.1
WI(2) = 0.05
WI(3) = 0.0375
ISEED = 123457
CALL RNSET (ISEED)
CALL RNARM (NW, CONST, NPAR, PAR, LAGAR, NPMA, PMA, LAGMA,
&          IADIST, AVAR, A, WI, W)
WRITE (NOUT,99999) (W(I),I=1,NW)
99999 FORMAT (' Simulated ARMA(3,2) series ', 5F7.4)
END

```

### Output

Simulated ARMA(3,2) series 1.4033 2.2200 2.2864 2.8878 2.8322

### Example 2

In this example, 500 observations from an ARMA(2, 2) process are simulated using RNARM; and then routine NSPE (page 664) is used to estimate the parameters of the model. The model is used as an example by Priestley (1981), page 139.

```

C
INTEGER      NPAR, NPMA, NW
PARAMETER    (NPAR=2, NPMA=2, NW=500)

C
INTEGER      IADIST, ISEED, LAGAR(NPAR), LAGMA(NPMA)
REAL         A(NW+2), AVAR, AVAR1, CONST, CONST1, PAR(NPAR),
&           PAR1(NPAR), PMA(NPMA), PMA1(NPMA), W(NW), WI(2), WMEAN
C
EXTERNAL     NSPE, RNARM, RNSET

LAGAR(1) = 1
LAGAR(2) = 2
PAR(1) = -1.4
PAR(2) = -0.5
LAGMA(1) = 1
LAGMA(2) = 2
PMA(1) = 0.2
PMA(2) = 0.1
IADIST = 0
CONST = 0.0
AVAR = 1.0
WI(1) = 0.0
WI(2) = 0.0
ISEED = 123457
CALL RNSET (ISEED)
CALL RNARM (NW, CONST, NPAR, PAR, LAGAR, NPMA, PMA, LAGMA,
&          IADIST, AVAR, A, WI, W)
CALL NSPE (NW, W, 1, 1, WMEAN, NPAR, NPMA, 0.0, 0, CONST1, PAR1,
&          PMA1, AVAR1)
END

```

### Output

Results from NSPE/N2PE  
WMEAN = 2.26275E-02

```

CONST =      7.19606E-02
AVAR  =      1.12579
      PAR
      1      2
-1.536 -0.645
      PMA
      1      2
0.0631 0.1405

```

---

## RNNPP/DRNNPP (Single/Double precision)

Generate pseudorandom numbers from a nonhomogeneous Poisson process.

### Usage

```
CALL RNNPP (TIMBEG, TIMEND, FTHETA, THEMIN, THEMAX, NEUB,
           NE, R)
```

### Arguments

**TIMBEG** — Lower endpoint of the time interval of the process. (Input)  
TIMBEG must be nonnegative. Usually, TIMBEG = 0.

**TIMEND** — Upper endpoint of the time interval of the process. (Input)  
TIMEND must be greater than TIMBEG.

**FTHETA** — User-supplied FUNCTION to provide the value of the rate of the process as a function of time. This function must be defined over the interval from TIMBEG to TIMEND and must be nonnegative in that interval. The form is FTHETA(TIME), where

TIME — Time at which the rate function is evaluated. (Input)

FTHETA — Value of the rate function. (Output)

FTHETA must be declared EXTERNAL in the calling program.

**THEMIN** — Minimum value of the rate function FTHETA in the interval (TIMBEG, TIMEND). (Input)

If the actual minimum is unknown, set THEMIN = 0.0.

**THEMAX** — Maximum value of the rate function FTHETA in the interval (TIMBEG, TIMEND). (Input)

If the actual maximum is unknown, set THEMAX to a known upper bound of the maximum. The efficiency of RNNPP is less the greater THEMAX exceeds the true maximum.

**NEUB** — Upper bound on the number of events to be generated. (Input)

In order to be reasonably sure that the full process through time TIMEND is generated, calculate NEUB as  $NEUB = X + 10.0 * \text{SQRT}(X)$ , where  $X = \text{THEMAX} * (\text{TIMEND} - \text{TIMBEG})$ . The only penalty in setting NEUB too large is that the output vector must be dimensioned of length NEUB.



$NE$  — Number of events actually generated. (Output)

If  $NE$  is less than  $NEUB$ , the time  $TIMEND$  is reached before  $NEUB$  events are realized.

$R$  — Vector of length  $NE$  containing the times to events. (Output)

$R$  must be dimensioned to be of length  $NEUB$ .

### Comments

The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

### Algorithm

Routine `RNNPP` simulates a one-dimensional nonhomogeneous Poisson process with rate function  $\text{THETA}$  in a fixed interval  $(\text{TIMBEG}, \text{TIMEND}]$ .

Let  $\lambda(t)$  be the rate function and  $t_0 = \text{TIMBEG}$  and  $t_1 = \text{TIMEND}$ . Routine `RNNPP` uses a method of thinning a nonhomogeneous Poisson process  $\{N^*(t), t \geq t_0\}$  with rate function  $\lambda^*(t) \geq \lambda(t)$  in  $(t_0, t_1]$ , where the number of events,  $N^*$ , in the interval  $(t_0, t_1]$  has a Poisson distribution with parameter

$$\mu_0 = \int_{t_0}^{t_1} \lambda(t) dt$$

The function

$$\Lambda(t) = \int_0^t \lambda(t) dt$$

is called the *integrated rate function*.) In `RNNPP`,  $\lambda^*(t)$  is taken to be a constant  $\lambda^*$  ( $= \text{THEMAX}$ ) so that at time  $t_i$ , the time of the next event  $t_{i+1}$  is obtained by generating and cumulating exponential random numbers

$$E_{1,i}^*, E_{2,i}^*, \dots,$$

with parameter  $\lambda^*$ , until for the first time

$$u_{j,i} \leq (t_i + E_{1,i}^* + \dots + E_{j,i}^*) / \lambda^*$$

where the  $u_{j,i}$  are independent uniform random numbers between 0 and 1. This process is continued until the specified number of events,  $NEUB$ , is realized or until the time,  $TIMEND$ , is exceeded. This method is due to Lewis and Shedler (1979), who also review other methods. The most straightforward (and most efficient) method is by inverting the integrated rate function, but often this is not possible.

If  $\text{THEMAX}$  is actually greater than the maximum of  $\lambda(t)$  in  $(t_0, t_1]$ , the routine will work, but less efficiently. Also, if  $\lambda(t)$  varies greatly within the interval, the

efficiency is reduced. In that case, it may be desirable to divide the time interval into subintervals within which the rate function is less variable. This is possible because the process is without memory.

If no time horizon arises naturally, TIMEND must be set large enough to allow for the required number of events to be realized. Care must be taken, however, that FTHETA is defined over the entire interval.

After simulating a given number of events, the next event can be generated by setting TIMBEG to the time of the last event (the sum of the elements in R) and calling RNNPP again. Cox and Lewis (1966) discuss modeling applications of nonhomogeneous Poisson processes.

### Example 1

In this example, RNNPP is used to generate the first five events in the time 0 to 20 (if that many events are realized) in a nonhomogeneous process with rate function

$$\lambda(t) = 0.6342 e^{0.001427 t}$$

for  $0 < t \leq 20$ .

Since this is a monotonically increasing function of  $t$ , the minimum is at  $t = 0$  and is 0.6342, and the maximum is at  $t = 20$  and is  $0.6342 e^{0.02854} = 0.652561$ .

```

INTEGER      NEUB
PARAMETER   (NEUB=5)
C
INTEGER      I, ISEED, NE, NOUT
REAL         FTHETA, R(NEUB), THEMAX, THEMIN, TIMBEG, TIMEND
EXTERNAL     FTHETA, RNNPP, RNSET, UMACH
C
CALL UMACH (2, NOUT)
TIMBEG = 0.0
TIMEND = 20.0
THEMIN = 0.6342
THEMAX = 0.652561
ISEED = 123457
CALL RNSET (ISEED)
CALL RNNPP (TIMBEG, TIMEND, FTHETA, THEMIN, THEMAX, NEUB, NE, R)
WRITE (NOUT,99999) NE, (R(I),I=1,NE)
99999 FORMAT (' Inter-event times for the first ', I1, ' events', /,
&           ' in the process: ', 5F7.4)
END
C
REAL FUNCTION FTHETA (T)
REAL      T
C
REAL      EXP
INTRINSIC EXP
C
FTHETA = 0.6342*EXP(0.001427*T)
RETURN
END

```

## Output

Inter-event times for the first 5 events  
in the process: 0.0527 0.4080 0.2584 0.0198 0.1676

## Example 2

As it turns out in the simulation above, the first five events are realized before time equals 20. If it is desired to continue the simulation to time equals 20, setting NEUB to 49 (that is,

$$\lambda^*(t_1 - t_0) + 10\sqrt{\lambda^*(t_1 - t_0)}$$

would likely ensure that the time is reached. In the following example, we see that there are twelve events realized by time equals 20.

```
INTEGER      NEUB
PARAMETER    (NEUB=49)

C
INTEGER      ISEED, NE, NOUT
REAL         FTHETA, R(NEUB), SSUM, T, THEMAX, THEMIN, TIMBEG,
&           TIMEND
EXTERNAL     FTHETA, RNNPP, RNSET, SSUM, UMACH

C
CALL UMACH (2, NOUT)
TIMBEG = 0.0
TIMEND = 20.0
THEMIN = 0.6342
THEMAX = 0.652561
ISEED = 123457
CALL RNSET (ISEED)
CALL RNNPP (TIMBEG, TIMEND, FTHETA, THEMIN, THEMAX, NEUB, NE, R)
T = TIMBEG + SSUM(NE,R,1)
IF (NE .LT. NEUB) THEN
99998  WRITE (NOUT,99998) NE, T
&     FORMAT (' Only ', I2, ' events occurred before the time',
&           /, ' limit expired. The last event occurred at', /,
&           ' time = ', F6.3)
ELSE
99999  WRITE (NOUT,99999) NE, T
&     FORMAT (' Possibly more than ', I2, ' events would have',
&           /, ' occurred before the time limit expired.', /,
&           ' The last event occurred at time = ', F6.3)
END IF
END

C
REAL FUNCTION FTHETA (T)
REAL      T

C
REAL      EXP
INTRINSIC EXP

C
FTHETA = 0.6342*EXP(0.001427*T)
RETURN
END
```

## Output

Only 12 events occurred before the time limit expired. The last event occurred at time = 18.809

---

# RNPER

Generate a pseudorandom permutation.

## Usage

```
CALL RNPER (K, IPER)
```

## Arguments

**K** — Number of integers to be permuted. (Input)

**IPER** — Vector of length  $K$  containing the random permutation of the integers from 1 to  $K$ . (Output)

## Comments

The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.

## Algorithm

Routine `RNPER` generates a pseudorandom permutation of the integers from 1 to  $K$ . It begins by filling a vector of length  $K$  with the consecutive integers 1 to  $K$ . Then, with  $M$  initially equal to  $K$ , a random index  $J$  between 1 and  $M$  (inclusive) is generated. The element of the vector with the index  $M$  and the element with index  $J$  swap places in the vector.  $M$  is then decremented by 1 and the process repeated until  $M = 1$ .

## Example

In this example, `RNPER` is called to produce a pseudorandom permutation of the integers from 1 to 10.

```
INTEGER      IPER(10), ISEED, K, NOUT
EXTERNAL     RNPER, RNSET, UMACH
C
CALL UMACH (2, NOUT)
K = 10
C                                     Initialize seed of random number
C                                     generator.
ISEED = 123457
CALL RNSET (ISEED)
CALL RNPER (K, IPER)
WRITE (NOUT,99999) IPER
99999 FORMAT (' Random permutation of the integers from 1 to 10', /,
&           10I5)
```

END

### Output

Random permutation of the integers from 1 to 10  
5 9 2 8 1 6 4 7 3 10

---

## RNSRI

Generate a simple pseudorandom sample of indices.

### Usage

```
CALL RNSRI (NSAMP, NPOP, INDEX)
```

### Arguments

*NSAMP* — Sample size desired. (Input)

*NPOP* — Number of items in the population. (Input)

*INDEX* — Vector of length *NSAMP* containing the indices of the sample.  
(Output)

*INDEX* is a random sample (without replacement) of the integers from 1 to *NPOP*, in increasing order.

### Comments

1. The routine *RNSET* (page 1167) can be used to initialize the seed of the random number generator. If *NSAMP* is greater than *NPOP*/2, *RNSRI* uses two different generators in an algorithm due to Ahrens and Dieter (1985). The routine *RNOPT* (page 1165) can be used to select the form of the generator used for uniform deviates in the algorithm. The generator used for exponential deviates in the algorithm is a nonshuffled generator that is different from the one for the uniform. If *IOPTU* is the option indicator for the uniform generator (see documentation for *RNOPT*, page 1165), then the option indicator for the exponential generator is  $\text{MOD}((2 * \text{INT}((\text{IOPTU} + 1)/2) + 1), 6)$ .
2. The routine *RNSRS* (page 1242) can be used to select a sample from a population of unknown size.

### Algorithm

Routine *RNSRI* generates the indices of a pseudorandom sample, without replacement, of size *NSAMP* numbers from a population of size *NPOP*. If *NSAMP* is greater than *NPOP*/2, the integers from 1 to *NPOP* are selected sequentially with a probability conditional on the number selected and the number remaining to be considered. If, when the *i*-th population index is considered, *j* items have been included in the sample, then the index *i* is included with probability  $(\text{NSAMP} - j) / (\text{NPOP} + 1 - i)$ .

If `NSAMP` is not greater than `NPOP/2`, a  $O(NSAMP)$  algorithm due to Ahrens and Dieter (1985) is used. Of the methods discussed by Ahrens and Dieter, the one called `SG*` is used in `RNSRI`. It involves a preliminary selection of  $q$  indices using a geometric distribution for the distances between each index and the next one. If the preliminary sample size  $q$  is less than `NSAMP`, a new preliminary sample is chosen, and this is continued until a preliminary sample greater in size than `NSAMP` is chosen. This preliminary sample is then thinned using the same kind of sampling as described above for the case in which the sample size is greater than half of the population size. Routine `RNSRI` does not store the preliminary sample indices, but rather restores the state of the generator used in selecting the sample initially, and then passes through once again, making the final selection as the preliminary sample indices are being generated.

### Example

In this example, `RNSRI` is used to generate the indices of a pseudorandom sample of size 5 from a population of size 100.

```

C
INTEGER      INDEX(5), ISEED, NOUT, NPOP, NSAMP
EXTERNAL     RNSRI, UMACH

CALL UMACH (2, NOUT)
NSAMP = 5
NPOP = 100
ISEED = 123457
CALL RNSRI (ISEED)
CALL RNSRI (NSAMP, NPOP, INDEX)
WRITE (NOUT,99999) INDEX
99999 FORMAT ('      Random sample: ', 5I4)
END

```

### Output

```
Random sample:      2      4      6      8     10
```

---

## RNSRS/DRNSRS (Single/Double precision)

Generate a simple pseudorandom sample from a finite population.

### Usage

```
CALL RNSRS (IDO, NROW, NVAR, POP, LDPOP, NSAMP, NPOP, SAMP,
           LDSAMP, INDEX)
```

### Arguments

**IDO** — Processing option. (Input)

#### IDO Action

- 0 This is the only invocation of `RNSRS` for this data set, and the entire population is input at once.
- 1 This is the first invocation, and additional calls to `RNSRS` will be made. Initialization and updating for the subpopulation in `POP` are performed.

2 This is an additional invocation of `RNSRS`, and updating for the subpopulation in `POP` is performed.

***NROW*** — Number of rows of data currently input in `POP`. (Input)  
`NROW` must be nonnegative.

***NVAR*** — Number of variables in the population and in the sample. (Input)

***POP*** — `NROW` by `NVAR` matrix containing the population to be sampled. (Input)  
If `IDO = 0`, `POP` contains the entire population; otherwise, `POP` contains a different part of the population on each invocation of `RNSRS`.

***LDPOP*** — Leading dimension of `POP` exactly as specified in the dimension statement in the calling program. (Input)

***NSAMP*** — The sample size desired. (Input)

***NPOP*** — The number of items in the population. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2`.)

If `IDO = 0`, `NPOP = NROW` on output. If the population is input a few items at a time, it is not necessary to know the number of items in the population in advance. `NPOP` is used to cumulate the population size and should not be changed between calls to `RNSRS`. If, on output, `NPOP` is greater than or equal to `NSAMP`, the sampling can be considered complete for a population of size `NPOP`.

***SAMP*** — `NSAMP` by `NVAR` matrix containing the sample. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2`.)

***LDSAMP*** — Leading dimension of `SAMP` exactly as specified in the dimension statement in the calling program. (Input)

***INDEX*** — Vector of length `NSAMP` containing the indices of the sample in the population. (Output, if `IDO = 0` or `1`; input/output, if `IDO = 2`.) The `INDEX(I)`-th item in the population is the `I`-th item in the sample. `INDEX` is not necessarily in increasing order.

### Comments

1. The routine `RNSET` (page 1167) can be used to initialize the seed of the random number generator. The routine `RNOPT` (page 1165) can be used to select the form of the generator.
2. The routine `RNSRI` (page 1241) can be used to select a sample of indices in increasing order.

### Algorithm

Routine `RNSRS` generates a pseudorandom sample from a given population, without replacement, using an algorithm due to McLeod and Bellhouse (1983).

The first `NSAMP` items in the population are included in the sample. Then, for each successive item from the population, a random item in the sample is replaced by that item from the population with probability equal to the sample

size divided by the number of population items that have been encountered at that time.

### Example 1

In this example, RNSRS is used to generate a sample of size 5 from a population stored in the matrix POP. All of the data are available at once, so IDO is set to 0.

```

INTEGER      I, IDO, INDEX(5), ISEED, J, LDPOP, LDSAMP, NOUT,
&            NPOP, NROW, NSAMP, NVAR
REAL         POP(176,2), SAMP(5,2)
EXTERNAL     GDATA, RNSET, RNSRS, UMACH

C
CALL UMACH (2, NOUT)
LDPOP = 176
NSAMP = 5
LDSAMP = 5

C                               Get Wolfer sunspot data to use
C                               as "population".
CALL GDATA (2, 0, NROW, NVAR, POP, 176, 2)
IDO = 0

C                               Initialize seed of random number
C                               generator.
ISEED = 123457
CALL RNSET (ISEED)
CALL RNSRS (IDO, NROW, NVAR, POP, LDPOP, NSAMP, NPOP, SAMP,
&          LDSAMP, INDEX)
WRITE (NOUT,99999) NPOP, INDEX, ((SAMP(I,J),I=1,5),J=1,2)
99999 FORMAT ('      The population size is ', I5, /, '      Indices of ',
&            'random sample: ', 5I8, /, '                        The sample: '
&            , 5F8.0, /, '                        ', 5F8.0)
END

```

### Output

```

The population size is 176
Indices of random sample:      16      80      175      25      21
The sample:      1764.    1828.    1923.    1773.    1769.
                   36.     62.     6.     35.     106.

```

### Example 2

Routine RNSRS is now used to generate a sample of size 5 from the same population as in the example above except the data are input to RNSRS one observation at a time. This is the way RNSRS may be used to sample from a file on disk or tape. Notice that the number of records need not be known in advance.

```

INTEGER      ISEED, NOUT, IDO, NROW, NVAR, LDPOP, NSAMP, NPOP,
&            LDSAMP, INDEX(5)
REAL         POP(176,2), SAMP(5,2), X(2)
CALL UMACH(2, NOUT)
LDPOP = 176
NSAMP = 5
LDSAMP = 5

C                               Get Wolfer sunspot data to use
C                               as "population".

```



```

CALL GDATA (2, 0, NROW, NVAR, POP, 176, 2)
C                                     Initialize seed of random number
C                                     generator.
ISEED = 123457
CALL RNSET(ISEED)
IDO = 1
DO 10 I=1,176
C                                     In this DO-loop, the data would
C                                     generally be read from a file,
C                                     one observation at a time. This
C                                     program simulates this by copying
C                                     the observations one at a time into
C                                     X from POP.
X(1) = POP(I,1)
X(2) = POP(I,2)
CALL RNSRS (IDO, 1, NVAR, X, 1, NSAMP, NPOP, SAMP, LDSAMP,
&          INDEX)
&          IDO = 2
10 CONTINUE
WRITE(NOUT, 20) NPOP, INDEX, ((SAMP(I,J),I=1,5),J=1,2)
20 FORMAT ('      The population size is ', I5,/,
&          '      Indices of random sample: ', 5I8,/,
&          '      The sample: ', 5F8.0,/,
&          '      ', 5F8.0)
END

```

### Output

```

The population size is 176
Indices of random sample: 16      80      175      25      21
The sample: 1764.  1828.  1923.  1773.  1769.
                36.    62.    6.    35.   106.

```

# Chapter 19: Utilities

---

## Routines

<b>19.1. Print</b>		
Real rectangular matrix with integer row and column labels .....	WRRRN	1248
Real rectangular matrix with given format and labels .....	WRRRL	1250
Integer rectangular matrix with integer row and column labels .....	WRIRN	1253
Integer rectangular matrix with given format and labels .....	WRIRL	1254
Set or retrieve options for printing a matrix .....	WROPT	1257
Set or retrieve page width and length .....	PGOPT	1263
<b>19.2. Permute</b>		
Elements of a vector .....	PERMU	1265
Rows/Columns of a matrix .....	PERMA	1266
Rows/Columns of a symmetric matrix .....	RORDM	1268
Move any rows with NaN to the last rows of the matrix .....	MVNAN	1269
<b>19.3. Sort</b>		
Real vector by algebraic value .....	SVRGN	1273
Real vector by algebraic value 0 and permutations returned .....	SVRGP	1274
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---

## WRRRN/DWRRRN (Single/Double precision)

Print a real rectangular matrix with integer row and column labels.

### Usage

CALL WRRRN (TITLE, NRA, NCA, A, LDA, ITRING)

### Arguments

**TITLE** — Character string specifying the title. (Input)

TITLE set equal to a blank character(s) suppresses printing of the title. Use “%*n*” within the title to create a new line. Long titles are automatically wrapped.

**NRA** — Number of rows. (Input)

**NCA** — Number of columns. (Input)

**A** — NRA by NCA matrix to be printed. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**ITRING** — Triangle option. (Input)

### ITRING Action

0 Full matrix is printed.

1 Upper triangle of A is printed, including the diagonal.

2 Upper triangle of A excluding the diagonal of A is printed.

-1 Lower triangle of A is printed, including the diagonal.

-2 Lower triangle of A excluding the diagonal of A is printed.

## Comments

1. A single D, E, or F format is chosen automatically in order to print 4 significant digits for the largest element of A in absolute value. Routine WROPT (page 1257) can be used to change the default format.
2. Horizontal centering, a method for printing large matrices, paging, printing a title on each page, and many other options can be selected by invoking WROPT.
3. A page width of 78 characters is used. Page width and page length can be reset by invoking PGOPT (page 1263).
4. Output is written to the unit specified by UMACH (Reference Material).

## Algorithm

Routine WRRRN prints a real rectangular matrix with the rows and columns labeled 1, 2, 3, and so on. WRRRN can restrict printing to the elements of the upper or lower triangles of matrices via the ITRING option. Generally, ITRING  $\neq$  0 is used with symmetric matrices.

In addition, one-dimensional arrays can be printed as column or row vectors. For a column vector, set NRA to the length of the array and set NCA = 1. For a row vector, set NRA = 1 and set NCA to the length of the array. In both cases, set LDA = NRA and set ITRING = 0.

## Example

The following example prints all of a  $3 \times 4$  matrix A where  $a_{ij} = i + j/10$ .

```
C      INTEGER      ITRING, LDA, NCA, NRA
      PARAMETER    ( ITRING=0, LDA=10, NCA=4, NRA=3 )
C
      INTEGER      I, J
      REAL         A(LDA,NCA)
      EXTERNAL     WRRRN
C
      DO 20 I=1, NRA
        DO 10 J=1, NCA
          A(I,J) = I + J*0.1
10     CONTINUE
20    CONTINUE
C
                                     Write A matrix.
      CALL WRRRN ('A', NRA, NCA, A, LDA, ITRING)
      END
```

## Output

```

          A
         1  2  3  4
1  1.100  1.200  1.300  1.400
2  2.100  2.200  2.300  2.400
3  3.100  3.200  3.300  3.400
```

---

## WRRRL/DWRRRL (Single/Double precision)

Print a real rectangular matrix with a given format and labels.

### Usage

```
CALL WRRRL (TITLE, NRA, NCA, A, LDA, ITRING, FMT, RLABEL,  
           CLABEL)
```

### Arguments

**TITLE** — Character string specifying the title. (Input)

TITLE set equal to a blank character(s) suppresses printing of the title.

**NRA** — Number of rows. (Input)

**NCA** — Number of columns. (Input)

**A** — NRA by NCA matrix to be printed. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

**ITRING** — Triangle option. (Input)

#### ITRING Action

0 Full matrix is printed.

1 Upper triangle of A is printed, including the diagonal.

2 Upper triangle of A excluding the diagonal of A is printed.

-1 Lower triangle of A is printed, including the diagonal.

-2 Lower triangle of A excluding the diagonal of A is printed.

**FMT** — Character string containing formats. (Input)

If FMT is set to a blank character(s), the format used is specified by WROPT (page 1257). Otherwise, FMT must contain exactly one set of parentheses and one or more edit descriptors. For example, FMT = '(F10.3)' specifies this F format for the entire matrix. FMT = '(2E10.3, 3F10.3)' specifies an E format for columns 1 and 2 and an F format for columns 3, 4 and 5. If the end of FMT is encountered and if some columns of the matrix remain, format control continues with the first format in FMT. Even though the matrix A is real, an I format can be used to print the integer part of matrix elements of A. The most useful formats are special formats, called the "v and w formats," that can be used to specify pretty formats automatically. Set FMT = '(V10.4)' if you want a single D, E, or F format selected automatically with field width 10 and with 4 significant digits. Set FMT = '(W10.4)' if you want a single D, E, F, or I format selected automatically with field width 10 and with 4 significant digits. While the v format prints trailing zeroes and a trailing decimal point, the w format does not. See Comment 4 for general descriptions of the v and w formats. FMT may contain only D, E, F, G, I, V, or W edit descriptors, e.g., the X descriptor is not allowed.

**RLABEL** — CHARACTER\*(\*) vector of labels for rows of A. (Input)

If rows are to be numbered consecutively 1, 2, ..., NRA, use

RLABEL(1) = 'NUMBER'. If no row labels are desired, use RLABEL(1) = 'NONE'.

Otherwise, RLABEL is a vector of length NRA containing the labels.

**CLABEL** — CHARACTER\*(\*) vector of labels for columns of A. (Input)

If columns are to be numbered consecutively 1, 2, ..., NCA, use

CLABEL(1) = 'NUMBER'. If no column labels are desired, use

CLABEL(1) = 'NONE'. Otherwise, CLABEL(1) is the heading for the row labels,

and either CLABEL(2) must be 'NUMBER' or 'NONE', or CLABEL must be a vector of length NCA + 1 with CLABEL(1 + j) containing the column heading for the j-th column.

### Comments

1. Automatic workspace is used only if all of the following three conditions are met: (1) FMT contains v or w edit descriptors. (2) FMT is not a single v or w format with no repetition factor. (3) WROPT has previously been invoked with IOPT = -2 and ISET = 0. In this case, workspace usage is

WRRRL 10 \* NCA character units, or

DWRRRL 10 \* NCA character units.

Workspace may be explicitly provided, if desired, by use of

W2RRL/DW2RRL. The reference is

```
CALL W2RRL (TITLE, NRA, NCA, A, LDA, ITRING, FMT,  
           RLABEL, CLABEL, CHWK)
```

The additional argument is

**CHWK** — CHARACTER \* 10 work vector of length NCA. This workspace is referenced only if all three conditions indicated at the beginning of this remark are met. Otherwise, CHWK is not referenced and can be a CHARACTER \* 10 vector of length one.

2. The output appears in the following form:

```
                TITLE  
CLABEL(1)  CLABEL(2)  CLABEL(3)  CLABEL(4)  
RLABEL(1)      XXXXX      XXXXX      XXXXX  
RLABEL(2)      XXXXX      XXXXX      XXXXX
```

3. Use “%/” within titles or labels to create a new line. Long titles or labels are automatically wrapped.
4. For printing numbers whose magnitudes are unknown, the G format in FORTRAN is useful; however, the decimal points will generally not be aligned when printing a column of numbers. The v and w formats are special formats used by this routine to select a D, E, F, or I format so

that the decimal points will be aligned. The *v* and *w* formats are specified as *Vn.d* and *Wn.d*. Here, *n* is the field width and *d* is the number of significant digits generally printed. Valid values for *n* are 3, 4, ..., 40. Valid values for *d* are 1, 2, ..., *n* - 2. If *FMT* specifies one format and that format is a *v* or *w* format, all elements of the matrix *A* are examined to determine one FORTRAN format for printing. If *FMT* specifies more than one format, FORTRAN formats are generated separately from each *v* or *w* format.

5. A page width of 78 characters is used. Page width and page length can be reset by invoking *PGOPT* (page 1263).
6. Horizontal centering, method for printing large matrices, paging, method for printing NaN (not a number), printing a title on each page, and many other options can be selected by invoking *WROPT* (page 1257).
7. Output is written to the unit specified by *UMACH* (Reference Material).

### Algorithm

Routine *WRRRL* prints a real rectangular matrix (stored in *A*) with row and column labels (specified by *RLABEL* and *CLABEL*, respectively) according to a given format (stored in *FMT*). *WRRRL* can restrict printing to the elements of upper or lower triangles of matrices via the *ITRING* option. Generally, *ITRING* ≠ 0 is used with symmetric matrices.

In addition, one-dimensional arrays can be printed as column or row vectors. For a column vector, set *NRA* to the length of the array and set *NCA* = 1. For a row vector, set *NRA* = 1 and set *NCA* to the length of the array. In both cases, set *LDA* = *NRA*, and set *ITRING* = 0

### Example

The following example prints all of a 3 × 4 matrix *A* where  $a_{ij} = (i + j/10)10^{j-3}$ .

```

C      INTEGER      ITRING, LDA, NCA, NRA
      PARAMETER    (ITRING=0, LDA=10, NCA=4, NRA=3)

C      INTEGER      I, J
      REAL          A(LDA,NCA)
      CHARACTER     CLABEL(5)*5, FMT*8, RLABEL(3)*5
      EXTERNAL      WRRRL

C      DATA FMT/'(W10.6)'/
      DATA CLABEL/' ', 'Col 1', 'Col 2', 'Col 3', 'Col 4'/
      DATA RLABEL/'Row 1', 'Row 2', 'Row 3'/

C      DO 20 I=1, NRA
          DO 10 J=1, NCA
              A(I,J) = (I+J*0.1)*10.0**(J-3)
10      CONTINUE
20 CONTINUE

```

```

C                                     Write A matrix.
CALL WRRRL ('A', NRA, NCA, A, LDA, ITRING, FMT, RLABEL, CLABEL)
END

```

### Output

	A			
	Col 1	Col 2	Col 3	Col 4
Row 1	0.011	0.120	1.300	14.000
Row 2	0.021	0.220	2.300	24.000
Row 3	0.031	0.320	3.300	34.000

---

## WRIRN

Print an integer rectangular matrix with integer row and column labels.

### Usage

```
CALL WRIRN (TITLE, NRMAT, NCMAT, MAT, LDMAT, ITRING)
```

### Arguments

**TITLE** — Character string specifying the title. (Input)

TITLE set equal to a blank character(s) suppresses printing of the title. Use “%l” within the title to create a new line. Long titles are automatically wrapped.

**NRMAT** — Number of rows. (Input)

**NCMAT** — Number of columns. (Input)

**MAT** — NRMAT by NCMAT matrix to be printed. (Input)

**LDMAT** — Leading dimension of MAT exactly as specified in the dimension statement in the calling program. (Input)

**ITRING** — Triangle option. (Input)

### ITRING Action

- 0 Full matrix is printed.
- 1 Upper triangle of MAT is printed, including the diagonal.
- 2 Upper triangle of MAT excluding the diagonal of MAT is printed.
- 1 Lower triangle of MAT is printed, including the diagonal.
- 2 Lower triangle of MAT excluding the diagonal of MAT is printed.

### Comments

1. All the entries in MAT are printed using a single I format. The field width is determined by the largest absolute entry.
2. Horizontal centering, a method for printing large matrices, paging, printing a title on each page, and many other options can be selected by invoking WROPT (page 1257).



3. A page width of 78 characters is used. Page width and page length can be reset by invoking PGOPT (page 1263).
4. Output is written to the unit specified by UMACH (Reference Material).

### Algorithm

Routine WRIRN prints an integer rectangular matrix with the rows and columns labeled 1, 2, 3, and so on. WRIRN can restrict printing to elements of the upper and lower triangles of matrices via the ITRING option. Generally, ITRING  $\neq$  0 is used with symmetric matrices. In addition, one-dimensional arrays can be printed as column or row vectors. For a column vector, set NRMAT to the length of the array and set NCMAT = 1. For a row vector, set NRMAT = 1 and set NCMAT to the length of the array. In both cases, set LDMAT = NRMAT and set ITRING = 0.

### Example

The following example prints all of a  $3 \times 4$  matrix  $A = MAT$  where  $a_{ij} = 10i + j$ .

```

INTEGER      ITRING, LDMAT, NCMAT, NRMAT
PARAMETER   (ITRING=0, LDMAT=10, NCMAT=4, NRMAT=3)
C
INTEGER      I, J, MAT(LDMAT,NCMAT)
EXTERNAL     WRIRN
C
DO 20 I=1, NRMAT
  DO 10 J=1, NCMAT
    MAT(I,J) = I*10 + J
10  CONTINUE
20 CONTINUE
C
                                Write MAT matrix.
CALL WRIRN ('MAT', NRMAT, NCMAT, MAT, LDMAT, ITRING)
END

```

### Output

```

          MAT
         1  2  3  4
1  11  12  13  14
2  21  22  23  24
3  31  32  33  34

```

---

## WRIRL

Print an integer rectangular matrix with a given format and labels.

### Usage

```

CALL WRIRL (TITLE, NRMAT, NCMAT, MAT, LDMAT, ITRING, FMT,
            RLABEL, CLABEL)

```

## Arguments

**TITLE** — Character string specifying the title. (Input)

TITLE set equal to a blank character(s) suppresses printing of the title.

**NRMAT** — Number of rows. (Input)

**NCMAT** — Number of columns. (Input)

**MAT** — NRMAT by NCMAT matrix to be printed. (Input)

**LDMAT** — Leading dimension of MAT exactly as specified in the dimension statement in the calling program. (Input)

**ITRING** — Triangle option. (Input)

### ITRING Action

0 Full matrix is printed.

1 Upper triangle of MAT is printed, including the diagonal.

2 Upper triangle of MAT excluding the diagonal of MAT is printed.

-1 Lower triangle of MAT is printed, including the diagonal.

-2 Lower triangle of MAT excluding the diagonal of MAT is printed.

**FMT** — Character string containing formats. (Input)

If FMT is set to a blank character(s), the format used is a single I format with field width determined by the largest absolute entry. Otherwise, FMT must contain exactly one set of parentheses and one or more I edit descriptors. For example, FMT = '(I10)' specifies this I format for the entire matrix. FMT = '(2I10, 3I5)' specifies an I10 format for columns 1 and 2 and an I5 format for columns 3, 4 and 5. If the end of FMT is encountered and if some columns of the matrix remain, format control continues with the first format in FMT. FMT may only contain the I edit descriptor, e.g., the X edit descriptor is not allowed.

**RLABEL** — CHARACTER \* (\*) vector of labels for rows of MAT. (Input)

If rows are to be numbered consecutively 1, 2, ..., NRMAT, use

RLABEL(1) = 'NUMBER'. If no row labels are desired, use

RLABEL(1) = 'NONE'. Otherwise, RLABEL is a vector of length NRMAT containing the labels.

**CLABEL** — CHARACTER \* (\*) vector of labels for columns of MAT. (Input)

If columns are to be numbered consecutively 1, 2, ..., NCMAT, use

CLABEL(1) = 'NUMBER'. If no column labels are desired, use

CLABEL(1) = 'NONE'. Otherwise, CLABEL(1) is the heading for the row labels,

and either CLABEL(2) must be 'NUMBER' or 'NONE', or CLABEL must be a vector of length NCMAT + 1 with CLABEL(1 + j) containing the column heading for the j-th column.

## Comments

1. The output appears in the following form:

TITLE			
CLABEL(1)	CLABEL(2)	CLABEL(3)	CLABEL(4)
RLABEL(1)	XXXXX	XXXXX	XXXXX
RLABEL(2)	XXXXX	XXXXX	XXXXX

2. Use “%/” within titles or labels to create a new line. Long titles or labels are automatically wrapped.
3. A page width of 78 characters is used. Page width and page length can be reset by invoking PGOPT (page 1263).
4. Horizontal centering, a method for printing large matrices, paging, printing a title on each page, and many other options can be selected by invoking WROPT (page 1257).
5. Output is written to the unit specified by UMACH (Reference Material).

### Algorithm

Routine WRIRL prints an integer rectangular matrix (stored in MAT) with row and column labels (specified by RLABEL and CLABEL, respectively), according to a given format (stored in FMT). WRIRL can restrict printing to the elements of upper or lower triangles of matrices via the ITRING option. Generally, ITRING ≠ 0 is used with symmetric matrices.

In addition, one-dimensional arrays can be printed as column or row vectors. For a column vector, set NRMAT to the length of the array and set NCMAT = 1. For a row vector, set NRMAT = 1 and set NCMAT to the length of the array. In both cases, set LDMAT = NRMAT, and set ITRING = 0.

### Example

The following example prints all of a  $3 \times 4$  matrix  $A = MAT$  where  $a_{ij} = 10i + j$ .

```

INTEGER      ITRING, LDMAT, NCMAT, NRMAT
PARAMETER   (ITRING=0, LDMAT=10, NCMAT=4, NRMAT=3)
C
INTEGER      I, J, MAT(LDMAT,NCMAT)
CHARACTER    CLABEL(5)*5, FMT*8, RLABEL(3)*5
EXTERNAL     WRIRL
C
DATA FMT/'(I2)'/
DATA CLABEL/'      ', 'Col 1', 'Col 2', 'Col 3', 'Col 4'/
DATA RLABEL/'Row 1', 'Row 2', 'Row 3'/
C
DO 20 I=1, NRMAT
  DO 10 J=1, NCMAT
    MAT(I,J) = I*10 + J
10  CONTINUE
20  CONTINUE
C
                                Write MAT matrix.
CALL WRIRL ('MAT', NRMAT, NCMAT, MAT, LDMAT, ITRING, FMT,
&          RLABEL, CLABEL)
END

```

## Output

	MAT			
	Col 1	Col 2	Col 3	Col 4
Row 1	11	12	13	14
Row 2	21	22	23	24
Row 3	31	32	33	34

---

# WROPT

Set or retrieve an option for printing a matrix.

## Usage

CALL WROPT ( IOPT, ISET, ISCOPE )

## Arguments

**IOPT** — Indicator of option type. (Input)

<b>IOPT</b>	<b>Description of Option Type</b>
-1, 1	Horizontal centering or left justification of matrix to be printed
-2, 2	Method for printing large matrices
-3, 3	Paging
-4, 4	Method for printing NaN (not a number), and negative and positive machine infinity.
-5, 5	Title option
-6, 6	Default format for real and complex numbers
-7, 7	Spacing between columns
-8, 8	Maximum horizontal space reserved for row labels
-9, 9	Indentation of continuation lines for row labels
-10, 10	Hot zone option for determining line breaks for row labels
-11, 11	Maximum horizontal space reserved for column labels
-12, 12	Hot zone option for determining line breaks for column labels
-13, 13	Hot zone option for determining line breaks for titles
-14, 14	Option for the label that appears in the upper left hand corner that can be used as a heading for the row numbers or a label for the column headings for WR**N routines
-15, 15	Option for skipping a line between invocations of WR**N routines, provided a new page is not to be issued
-16, 16	Option for vertical alignment of the matrix values relative to the associated row labels that occupy more than one line
0	Reset all the current settings saved in internal variables back to their last setting made with an invocation of WROPT with ISCOPE = 1. (This option is used internally by routines printing a matrix and is not useful otherwise.)

If IOPT is negative, ISET and ISCOPE are input and are saved in internal variables. If IOPT is positive, ISET is output and receives the currently active

setting for the option (if `ISCOPE = 0`) or the last global setting for the option (if `ISCOPE = 1`). If `IOPT = 0`, `ISET` and `ISCOPE` are not referenced.

**ISET** — Setting for option selected by `IOPT`. (Input, if `IOPT` is negative; output, if `IOPT` is positive; not referenced if `IOPT = 0`)

<b>IOPT</b>	<b>ISET</b>	<b>Meaning</b>
-1, 1	0	Matrix is left justified.
-1		Matrix is centered horizontally on page.
-2, 2	0	A complete row is printed before the next row is printed. Wrapping is used if necessary.
	<i>m</i>	Here, <i>m</i> is a positive integer. Let $n_1$ be the maximum number of columns beginning with column 1 that fit across the page (as determined by the widths of the printing formats). First, columns 1 through $n_1$ are printed for rows 1 through <i>m</i> . Let $n_2$ be the maximum number of columns beginning with column $n_1 + 1$ that fit across the page. Second, columns $n_1 + 1$ through $n_1 + n_2$ are printed for rows 1 through <i>m</i> . This continues until the last columns are printed for rows 1 through <i>m</i> . Printing continues in this fashion for the next <i>m</i> rows, etc.
-3, 3	-2	Printing begins on the next line, and no paging occurs.
	-1	Paging is on. Every invocation of a <code>WR***</code> routine begins on a new page, and paging occurs within each invocation as is needed.
	0	Paging is on. The first invocation of a <code>WR***</code> routine begins on a new page, and subsequent paging occurs as is needed. With this option, every invocation of a <code>WR***</code> routine ends with a call to <code>WROPT</code> to reset this option to <i>k</i> , a positive integer giving the number of lines printed on the current page.

IOPT	ISET	Meaning
	$k$	Here, $k$ is a positive integer. Paging is on, and $k$ lines have been printed on the current page. If $k$ is less than the page length $IPAGE$ (see $PGOPT$ , page 1263), then $IPAGE - k$ lines are printed before a new page instruction is issued. If $k$ is greater than or equal to $IPAGE$ , then the first invocation of a $WR^{***}$ routine begins on a new page. In any case, subsequent paging occurs as is needed. With this option, every invocation of a $WR^{***}$ routine ends with a call to $WROPT$ to reset the value of $k$ .
-4, 4	0	NaN is printed as a series of decimal points, negative machine infinity is printed as a series of minus signs, and positive machine infinity is printed as a series of plus signs.
	1	NaN is printed as a series of blank characters, negative machine infinity is printed as a series of minus signs, and positive machine infinity is printed as a series of plus signs.
	2	NaN is printed as "NaN," negative machine infinity is printed as "-Inf" and positive machine infinity is printed as "Inf."
	3	NaN is printed as a series of blank characters, negative machine infinity is printed as "-Inf," and positive machine infinity is printed as "Inf."
-5, 5	0	Title appears only on first page.
	1	Title appears on the first page and all continuation pages.
-6, 6	0	Format is (w10.4). See Comment 2.
	1	Format is (w12.6). See Comment 2.
	2	Format is (1PE12.5).
	3	Format is $Vn.4$ where the field width $n$ is determined. See Comment 2.
	4	Format is $Vn.6$ where the field width $n$ is determined. See Comment 2.
	5	Format is $1PEN.d$ where $n = d + 7$ , and $d + 1$ is the maximum number of significant digits.
-7, 7	$k_1$	Number of characters left blank between columns. $k_1$ must be between 0 and 5, inclusively.
-8, 8	$k_2$	Maximum width (in characters) reserved for row labels. $k_2 = 0$ means use the default.

<b>IOPT</b>	<b>ISET</b>	<b>Meaning</b>
-9, 9	$k_3$	Number of characters used to indent continuation lines for row labels. $k_3$ must be between 0 and 10, inclusively.
-10, 10	$k_4$	Width (in characters) of the hot zone where line breaks in row labels can occur. $k_4 = 0$ means use the default. $k_4$ must not exceed 50.
-11, 11	$k_5$	Maximum width (in characters) reserved for column labels. $k_5 = 0$ means use the default.
-12, 12	$k_6$	Width (in characters) of the hot zone where line breaks in column labels can occur. $k_6 = 0$ means use the default. $k_6$ must not exceed 50.
-13, 13	$k_7$	Width (in characters) of the hot zone where line breaks in titles can occur. $k_7$ must be between 1 and 50, inclusively.
-14	0	There is no label in the upper left hand corner.
	1	The label in the upper left hand corner is "Component" if a row vector or column vector is printed; the label is "Row/Column" if both the number of rows and columns are greater than one; otherwise, there is no label.
-15	0	A blank line is printed on each invocation of a WR**N routine before the matrix title provided a new page is not to be issued.
	1	A blank line is not printed on each invocation of a WR**N routine before the matrix title.
-16, 16	0	The matrix values are aligned vertically with the last line of the associated row label for the case IOPT = 2 and ISET is positive.
	1	The matrix values are aligned vertically with the first line of the associated row label.

**ISCOPE** — Indicator of the scope of the option. (Input if IOPT is nonzero; not referenced if IOPT = 0)

**ISCOPE Action**

- 0 Setting is temporarily active for the next invocation of a WR\*\*\* matrix printing routine.
- 1 Setting is active until it is changed by another invocation of WROPT.

**Comments**

1. This program can be invoked repeatedly before using a WR\*\*\* routine to print a matrix. The matrix printing routines retrieve these settings to

determine the printing options. It is not necessary to call `WROPT` if a default value of a printing option is desired. The defaults are as follows.

<b>IOPT</b>	<b>Default Value for ISET</b>	<b>Meaning</b>
1	0	Left justified.
2	1000000	Number lines before wrapping.
3	-2	No paging.
4	2	NaN is printed as "NaN," negative machine infinity is printed as "-Inf" and positive machine infinity is printed as "Inf."
5	0	Title only on first page.
6	3	Default format is <i>Vn.4</i> .
7	2	2 spaces between columns.
8	0	Maximum row label width $MAXRLW = 2 * IPAGEW/3$ if matrix has one column; $MAXRLW = IPAGEW/4$ otherwise.
9	3	3 character indentation of row labels continued beyond one line.
10	0	Width of row label hot zone is $MAXRLW/3$ characters.
11	0	Maximum column label width $MAXCLW = \min\{\max(NW + NW/2, 15), 40\}$ for integer and real matrices, where $NW$ is the field width for the format corresponding to the particular column. $MAXCLW = \min\{\max(NW + NW/2, 15), 83\}$ for complex matrices, where $NW$ is the sum of the two field widths for the formats corresponding to the particular column plus 3.
12	0	Width of column label hot zone is $MAXCLW/3$ characters.
13	10	Width of hot zone for titles is 10 characters.
14	0	There is no label in the upper left hand corner.
15	0	Blank line is printed.



<b>IOPT</b>	<b>Default Value for ISET</b>	<b>Meaning</b>
16	0	The matrix values are aligned vertically with the last line of the associated row label. For IOPT = 8, the default depends on the current value for the page width, IPAGEW (see PGOPT, page 1263).

- The *v* and *w* formats are special formats that can be used to select a *D*, *E*, *F*, or *I* format so that the decimal points will be aligned. The *v* and *w* formats are specified as *Vn.d* and *Wn.d*. Here, *n* is the field width and *d* is the number of significant digits generally printed. Valid values for *n* are 3, 4, ..., 40. Valid values for *d* are 1, 2, ..., *n* - 2. While the *v* format prints trailing zeroes and a trailing decimal point, the *w* format does not.

### Algorithm

Routine `WROPT` allows the user to set or retrieve an option for printing a matrix. The options controlled by `WROPT` include the following: horizontal centering, a method for printing large matrices, paging, method for printing NaN (not a number) and positive and negative machine infinities, printing titles, default formats for numbers, spacing between columns, maximum widths reserved for row and column labels, indentation of row labels that continue beyond one line, widths of hot zones for breaking of labels and titles, the default heading for row labels, whether to print a blank line between invocations of routines, and vertical alignment of matrix entries with respect to row labels continued beyond one line. (NaN and positive and negative machine infinities can be retrieved by `AMACH` and `DMACH` that are documented in the section “Machine-Dependent Constants” in the Reference Material.) Options can be set globally (`ISCOPE = 1`) or temporarily for the next call to a printing routine (`ISCOPE = 0`).

### Example

The following example illustrates the effect of `WROPT` when printing a  $3 \times 4$  real matrix *A* with `WRRRN` (page 1248) where  $a_{ij} = i + j/10$ . The first call to `WROPT` sets horizontal printing so that the matrix is first printed horizontally centered on the page. In the next invocation of `WRRRN`, the left-justification option has been set via routine `WROPT` so the matrix is left justified when printed. Finally, because the scope of left justification was only for the next call to a printing routine, the last call to `WRRRN` results in horizontally centered printing.

```

C
INTEGER      ITRING, LDA, NCA, NRA
PARAMETER   (ITRING=0, LDA=10, NCA=4, NRA=3)

C
INTEGER      I, IOPT, ISCOPE, ISET, J
REAL        A(LDA,NCA)
EXTERNAL    WROPT, WRRRN

C
DO 20 I=1, NRA
  DO 10 J=1, NCA
```

```

          A(I,J) = I + J*0.1
10      CONTINUE
20      CONTINUE
C
C          Activate centering option.
C          Scope is global.
      IOPT = -1
      ISET = 1
      ISCOPE = 1
C
      CALL WROPT (IOPT, ISET, ISCOPE)
C          Write A matrix.
      CALL WRRRN ('A', NRA, NCA, A, LDA, ITRING)
C          Activate left justification.
C          Scope is local.
      IOPT = -1
      ISET = 0
      ISCOPE = 0
      CALL WROPT (IOPT, ISET, ISCOPE)
      CALL WRRRN ('A', NRA, NCA, A, LDA, ITRING)
      CALL WRRRN ('A', NRA, NCA, A, LDA, ITRING)
END

```

### Output

```

          A
          1      2      3      4
      1  1.100  1.200  1.300  1.400
      2  2.100  2.200  2.300  2.400
      3  3.100  3.200  3.300  3.400

      A
      1      2      3      4
1  1.100  1.200  1.300  1.400
2  2.100  2.200  2.300  2.400
3  3.100  3.200  3.300  3.400

          A
          1      2      3      4
      1  1.100  1.200  1.300  1.400
      2  2.100  2.200  2.300  2.400
      3  3.100  3.200  3.300  3.400

```

---

## PGOPT

Set or retrieve page width and length for printing.

### Usage

```
CALL PGOPT (IOPT, IPAGE)
```

### Arguments

*IOPT* — Page attribute option. (Input)

<i>IOPT</i>	Description of Attribute
-1, 1	Page width.

-2, 2 Page length.

Negative values of IOPT indicate the setting IPAGE is input. Positive values of IOPT indicate the setting IPAGE is output.

**IPAGE** — Value of page attribute. (Input, if IOPT is negative; output, if IOPT is positive.)

**IOPT Description of Attribute Settings for IPAGE**

-1, 1 Page width (in characters) 10, 11, ...

-2, 2 Page length (in lines) 10, 11, ...

**Algorithm**

Routine PGOPT is used to set or retrieve the page width or the page length for routines that perform printing.

**Example**

The following example illustrates the use of PGOPT to set the page width at 20 characters. Routine WRRRN (page 1248) is then used to print a  $3 \times 4$  matrix  $A$  where  $a_{ij} = i + j/10$ .

```
INTEGER      ITRING, LDA, NCA, NRA
PARAMETER    (ITRING=0, LDA=10, NCA=4, NRA=3)
C
INTEGER      I, IOPT, IPAGE, J
REAL         A(LDA,NCA)
EXTERNAL     PGOPT, WRRRN
C
DO 20 I=1, NRA
  DO 10 J=1, NCA
    A(I,J) = I + J*0.1
10  CONTINUE
20  CONTINUE
C
                                Set page width.
IOPT = -1
IPAGE = 20
CALL PGOPT (IOPT, IPAGE)
C
                                Print the matrix A.
CALL WRRRN ('A', NRA, NCA, A, LDA, ITRING)
END
```

**Output**

```
      A
      1      2
1  1.100  1.200
2  2.100  2.200
3  3.100  3.200

      3      4
1  1.300  1.400
2  2.300  2.400
3  3.300  3.400
```

---

# PERMU/DPERMU (Single/Double precision)

Rearrange the elements of an array as specified by a permutation.

## Usage

```
CALL PERMU (N, X, IPERMU, IPATH, XPERMU)
```

## Arguments

*N* — Length of the arrays *X* and *XPERMU*. (Input)

*X* — Real vector of length *N* containing the array to be permuted. (Input)

*IPERMU* — Integer vector of length *N* containing a permutation *IPERMU*(1), ..., *IPERMU*(*N*) of the integers 1, ..., *N*. (Input)

*IPATH* — Integer flag. (Input)

*IPATH* = 1 means *IPERMU* represents a forward permutation, i.e., *X*(*IPERMU*(*I*)) is moved to *XPERMU*(*I*). *IPATH* = 2 means *IPERMU* represents a backward permutation, i.e., *X*(*I*) is moved to *XPERMU*(*IPERMU*(*I*)).

*XPERMU* — Real vector of length *N* containing the array *X* permuted. (Output)  
If *X* is not needed, *X* and *XPERMU* can share the same storage locations.

## Algorithm

Routine *PERMU* rearranges the elements of an array according to a permutation vector. It has the option to do both forward and backward permutations.

## Example

This example rearranges the array *X* using *IPERMU*; forward permutation is performed.

```
C                                     Declare variables
C   INTEGER      IPATH, N
C   PARAMETER    ( IPATH=1, N=4)
C
C   INTEGER      IPERMU(N), J, NOUT
C   REAL         X(N), XPERMU(N)
C   EXTERNAL     PERMU, UMACH
C                                     Set values for X, IPERMU
C
C                                     X = ( 5.0  6.0  1.0  4.0 )
C                                     IPERMU = ( 3 1 4 2 )
C
C   DATA X/5.0, 6.0, 1.0, 4.0/, IPERMU/3, 1, 4, 2/
C                                     Permute X into XPERMU
C   CALL PERMU (N, X, IPERMU, IPATH, XPERMU)
C                                     Get output unit number
C   CALL UMACH (2, NOUT)
C
C                                     Print results
C   WRITE (NOUT,99999) (XPERMU(J),J=1,N)
```

```
C
99999 FORMAT (' The output vector is:', /, 10(1X,F10.2))
      END
```

### Output

```
The output vector is:
      1.00      5.00      4.00      6.00
```

---

## PERMA/DPERMA (Single/Double precision)

Permute the rows or columns of a matrix.

### Usage

```
CALL PERMA (NRA, NCA, A, LDA, IPERMU, IPATH, APER, LDAPER)
```

### Arguments

**NRA** — Number of rows. (Input)

**NCA** — Number of columns. (Input)

**A** — NRA by NCA matrix to be permuted. (Input)

**LDA** — Leading dimension of A exactly as specified in the dimension statement of the calling program. (Input)

**IPERMU** — Vector of length  $\kappa$  containing a permutation  $IPERMU(1), \dots, IPERMU(\kappa)$  of the integers  $1, \dots, \kappa$  where  $\kappa = NRA$  if the rows of A are to be permuted and  $\kappa = NCA$  if the columns of A are to be permuted. (Input)

**IPATH** — Option parameter. (Input)

$IPATH = 1$  means the rows of A will be permuted.  $IPATH = 2$  means the columns of A will be permuted.

**APER** — NRA by NCA matrix containing the permuted matrix. (Output)

If A is not needed, A and APER can share the same storage locations.

**LDAPER** — Leading dimension of APER exactly as specified in the dimension statement of the calling program. (Input)

### Comments

Automatic workspace usage is

```
PERMA  NCA units, or
DPERMA 2 * NCA units.
```

Workspace may be explicitly provided, if desired, by use of P2RMA/DP2RMA. The reference is

```
CALL P2RMA (NRA, NCA, A, LDA, IPERMU, IPATH, APER, LDAPER,
           WORK)
```

The additional argument is

**WORK** — Real work vector of length NCA.

### Algorithm

Routine PERMA interchanges the rows or columns of a matrix using a permutation vector.

The routine PERMA permutes a column (row) at a time by calling PERMU (page 1265). This process is continued until all the columns (rows) are permuted. On completion, let  $B = APER$  and  $p_i = IPERMU(I)$ , then

$$B_{ij} = A_{p_i j}$$

for all  $i, j$ .

### Example

This example permutes the columns of a matrix A.

```
C                               Declare variables
C   INTEGER   IPATH, LDA, LDAPER, NCA, NRA
C   PARAMETER (IPATH=2, LDA=3, LDAPER=3, NCA=5, NRA=3)
C
C   INTEGER   I, IPERMU(5), J, NOUT
C   REAL      A(LDA,NCA), APER(LDAPER,NCA)
C   EXTERNAL  PERMA, UMACH
C                               Set values for A, IPERMU
C   A = ( 3.0  5.0  1.0  2.0  4.0 )
C         ( 3.0  5.0  1.0  2.0  4.0 )
C         ( 3.0  5.0  1.0  2.0  4.0 )
C
C                               IPERMU = ( 3 4 1 5 2 )
C
C   DATA A/3*3.0, 3*5.0, 3*1.0, 3*2.0, 3*4.0/, IPERMU/3, 4, 1, 5, 2/
C                               Perform column permutation on A,
C                               giving APER
C   CALL PERMA (NRA, NCA, A, LDA, IPERMU, IPATH, APER, LDAPER)
C                               Get output unit number
C   CALL UMACH (2, NOUT)
C                               Print results
C   WRITE (NOUT,99999) ((APER(I,J),J=1,NCA),I=1,NRA)
C
C   99999 FORMAT (' The output matrix is:', /, 3(5F8.1,/))
C   END
```

### Output

```
The output matrix is:
1.0    2.0    3.0    4.0    5.0
1.0    2.0    3.0    4.0    5.0
1.0    2.0    3.0    4.0    5.0
```

---

## RORDM/DRORDM (Single/Double precision)

Reorder rows and columns of a symmetric matrix.

### Usage

```
CALL RORDM (NAA, AA, LDAA, NA, INDAA, A, LDA)
```

### Arguments

**NAA** — Order of the matrix AA. (Input)

**AA** — NAA by NAA symmetric matrix to be reordered. (Input)  
Only elements in the upper triangle of AA are referenced.

**LDAA** — Leading dimension of AA exactly as specified in the dimension statement in the calling program. (Input)

**NA** — Order of the reordered matrix A. (Input)  
NA must be less than or equal to NAA.

**INDAA** — Index vector of length NA containing the indices of the rows/columns of AA that are being selected for inclusion into A. (Input)  
 $INDAA(I) = J$  means the J-th row and column of AA will be the I-th row and column of A.

**A** — NAA by NAA matrix containing the reordered AA. (Output)  
The first NA rows and columns of A are those specified by INDAA. The remaining elements of A contain the rows and columns not specified in INDAA

**LDA** — Leading dimension of A exactly as specified in the dimension statement in the calling program. (Input)

### Comments

Automatic workspace usage is

```
RORDM NAA units, or  
DRORDM NAA units.
```

Workspace may be explicitly provided, if desired, by use of R2RDM/DR2RDM. The reference is

```
CALL R2RDM (NAA, AA, LDAA, NA, INDAA, A, LDA, IWK)
```

The additional argument is

**IWK** — Work vector of length NAA indicating how the entire AA matrix has been reordered and returned in A.  $IWK(I) = J$  means the J-th row and column of AA are returned as the I-th row and column of A.

## Algorithm

Routine RORDM reorders the rows and columns of a symmetric matrix. Frequently in practice a sum of squares and crossproducts matrix is first computed for all variables in a data set. Then, a sum of squares and crossproducts matrix is needed for some subset of the data set variables. Alternatively, a specific order for the selected variables may be required for input into an analysis routine. For example, in regression, IMSL routine RCOV requires the sum of squares and crossproducts matrix for the independent variables and the dependent variables. Sums of squares and crossproducts for the independent variables must appear first, followed by entries for the dependent variables. Variables not in the regression analysis, but in the data set, can appear last. RORDM can be used to reorder the sum of squares and crossproducts matrix for input to RCOV.

## Example

A  $4 \times 4$  symmetric matrix AA is reordered so that row/column 4, 3, and 1 of AA correspond to row/ column 1, 2, and 3 of A, respectively.

```
C
INTEGER   LDA, LDAA, NA, NAA
PARAMETER (NA=3, NAA=4, LDA=NAA, LDAA=NAA)

C
INTEGER   INDAA(NA)
REAL      A(LDA,NAA), AA(LDAA,NAA)
EXTERNAL  RORDM, WRRRN

C
DATA (AA(1,J),J=1,NAA)/10., 20., 40., 70./
DATA (AA(2,J),J=1,NAA)/20., 30., 50., 80./
DATA (AA(3,J),J=1,NAA)/40., 50., 60., 90./
DATA (AA(4,J),J=1,NAA)/70., 80., 90., 100./
DATA INDAA/4, 3, 1/

C
CALL RORDM (NAA, AA, LDAA, NA, INDAA, A, LDA)
CALL WRRRN ('A', NAA, NAA, A, LDA, 0)
END
```

## Output

		A			
		1	2	3	4
1	100.0	90.0	70.0	80.0	
2	90.0	60.0	40.0	50.0	
3	70.0	40.0	10.0	20.0	
4	80.0	50.0	20.0	30.0	

---

## MVNAN/DMVNAN (Single/Double precision)

Move any rows of a matrix with the IMSL missing value code NaN (not a number) in the specified columns to the last rows of the matrix.



## Usage

CALL MVNAN (NROW, NCOL, IIND, IND, X, LDX, ISWAP, NRMISS)

## Arguments

**NROW** — Number of rows. (Input)

**NCOL** — Number of columns. (Input)

**IIND** — Index vector option. (Input)

### **IIND** Meaning

< 0 The first  $-IIND$  columns of  $X$  are checked for NaN.

> 0 The  $IIND$  columns of  $X$  given by  $IND$  are checked for NaN.

**IND** — Index vector of length  $IIND$  containing the column numbers of  $X$  that are to be checked for NaN. (Input if  $IIND$  is positive)

If  $IIND$  is negative,  $IND$  is not referenced and can be a vector of length one.

**X** —  $NROW$  by  $NCOL$  matrix whose rows are checked for NaN (not a number). (Input/Output)

On output, the rows of  $X$  containing NaN are the last  $NRMISS$  rows of  $X$ .

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement of the calling program. (Input)

**ISWAP** — Vector of length  $NROW$  specifying the rows that were exchanged (swapped). (Output)

The number of nonzero elements in  $ISWAP$  is the number of swaps that took place.  $ISWAP(I) = J$  ( $J$  greater than zero) means that rows  $I$  and  $J$  of  $X$  were swapped, i.e., row  $I$  of the input  $X$  is row  $J$  of the output  $X$  and row  $J$  of the input  $X$  is row  $I$  of the output  $X$ .

**NRMISS** — Number of rows that contained NaN in the specified columns of  $X$ . (Output)

## Algorithm

Routine `MVNAN` moves any rows containing the IMSL missing value code NaN (not a number) in the specified columns to the last rows of the matrix. In single precision, NaN can be assigned to the element of a matrix by using routine `AMACH`. In double precision, routine `DMACH` should be used instead. (See the following examples and the section “Machine-Dependent Constants” in the Reference Material.)

The columns that are checked for NaN are specified through arguments  $IIND$  and  $IND$ . Let  $c_1, c_2, \dots, c_k$  be the column numbers specified. The main steps performed by `MVNAN` are as follows:

1. Initialize the  $NROW$  elements of  $ISWAP$  to zero.
2. Exchange (swap) the first row of  $X$  containing NaN in at least one of the specified columns  $c_1, c_2, \dots, c_k$  with the last row of  $X$  that does not

- contain NaN in any of the specified columns  $c_1, c_2, \dots, c_k$ . Let  $i$  and  $j$  ( $i < j$ ) correspond to the rows that are exchanged. Set the  $i$ -th element of ISWAP to  $j$ .
3. Starting with the  $(i + 1)$ -st row of  $x$ , exchange the first row containing NaN with the last row not containing NaN, beginning from row  $j - 1$  and continuing backward. Reset  $i$  and  $j$  ( $i < j$ ) to correspond to the rows exchanged.
  4. Continue in this manner until all rows have been examined for NaN.

### Example 1

In this example, MVNAN is used to move rows containing NaN in columns 1 and 2 of a 5 by 3 matrix  $x$  to the last rows.

```

INTEGER      LDX, NCOL, NROW
PARAMETER   (NCOL=3, NROW=5, LDX=NROW)
C
INTEGER      IIND, IND(1), ISWAP(NROW), NOUT, NRMISS
REAL         AMACH, X(LDX,NCOL)
EXTERNAL     AMACH, MVNAN, UMACH, WRIRN, WRRRN
C
DATA (X(1,J),J=1,NCOL)/1.0, 10.0, 100.0/
DATA (X(2,J),J=1,NCOL)/2.0, 20.0, 200.0/
DATA (X(3,J),J=1,NCOL)/3.0, 30.0, 300.0/
DATA (X(4,J),J=1,NCOL)/4.0, 40.0, 400.0/
DATA (X(5,J),J=1,NCOL)/5.0, 50.0, 500.0/
C
X(2,2) = AMACH(6)
X(4,1) = AMACH(6)
IIND   = -2
CALL WRRRN ('Input X', NROW, NCOL, X, LDX, 0)
CALL MVNAN (NROW, NCOL, IIND, IND, X, LDX, ISWAP, NRMISS)
CALL WRRRN ('Output X', NROW, NCOL, X, LDX, 0)
CALL WRIRN ('ISWAP', NROW, 1, ISWAP, NROW, 0)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) ' '
WRITE (NOUT,*) 'NRMISS = ', NRMISS
END

```

### Output

```

      Input X
      1      2      3
1     1.0    10.0   100.0
2     2.0     NaN   200.0
3     3.0    30.0   300.0
4     NaN    40.0   400.0
5     5.0    50.0   500.0

      Output X
      1      2      3
1     1.0    10.0   100.0
2     5.0    50.0   500.0
3     3.0    30.0   300.0
4     NaN    40.0   400.0

```

```
5      2.0      NaN      200.0
```

```
ISWAP
```

```
1      0
```

```
2      5
```

```
3      0
```

```
4      0
```

```
5      0
```

```
NRMISS =      2
```

## Example 2

In this example, MVNAN is used to move rows containing NaN in column 1 and 3 of a 5 by 3 matrix X to the last rows.

```
INTEGER      LDX, NCOL, NROW
PARAMETER    (NCOL=3, NROW=5, LDX=NROW)
C
INTEGER      IIND, IND(2), ISWAP(NROW), NOUT, NRMISS
REAL         AMACH, X(LDX,NCOL)
EXTERNAL     AMACH, MVNAN, UMACH, WRIRN, WRRRN
C
DATA (X(1,J),J=1,NCOL)/1.0, 10.0, 100.0/
DATA (X(2,J),J=1,NCOL)/2.0, 20.0, 200.0/
DATA (X(3,J),J=1,NCOL)/3.0, 30.0, 300.0/
DATA (X(4,J),J=1,NCOL)/4.0, 40.0, 400.0/
DATA (X(5,J),J=1,NCOL)/5.0, 50.0, 500.0/
DATA IND/1, 3/
C
X(2,2) = AMACH(6)
X(4,1) = AMACH(6)
IIND   = 2
CALL WRRRN ('Input X', NROW, NCOL, X, LDX, 0)
CALL MVNAN (NROW, NCOL, IIND, IND, X, LDX, ISWAP, NRMISS)
CALL WRRRN ('Output X', NROW, NCOL, X, LDX, 0)
CALL WRIRN ('ISWAP', NROW, 1, ISWAP, NROW, 0)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) ' '
WRITE (NOUT,*) 'NRMISS = ', NRMISS
END
```

## Output

```
      Input X
      1      2      3
1      1.0     10.0    100.0
2      2.0      NaN    200.0
3      3.0     30.0    300.0
4      NaN     40.0    400.0
5      5.0     50.0    500.0
```

```
      Output X
      1      2      3
1      1.0     10.0    100.0
2      2.0      NaN    200.0
3      3.0     30.0    300.0
4      5.0     50.0    500.0
```

```
5      NaN      40.0      400.0
```

```
ISWAP
1      0
2      0
3      0
4      5
5      0
NRMISS =      1
```

---

## SVRGN/DSVRGN (Single/Double precision)

Sort a real array by algebraically increasing value.

### Usage

```
CALL SVRGN (N, RA, RB)
```

### Arguments

*N* — Number of elements in the array to be sorted. (Input)

*RA* — Vector of length *N* containing the array to be sorted. (Input)

*RB* — Vector of length *N* containing the sorted array. (Output)

If *RA* is not needed, *RA* and *RB* can share the same storage locations.

### Algorithm

Routine *SVRGN* sorts the elements of an array, *A*, into ascending order by algebraic value. The array *A* is divided into two parts by picking a central element *T* of the array. The first and last elements of *A* are compared with *T* and exchanged until the three values appear in the array in ascending order. The elements of the array are rearranged until all elements greater than or equal to the central element appear in the second part of the array and all those less than or equal to the central element appear in the first part. The upper and lower subscripts of one of the segments are saved, and the process continues iteratively on the other segment. When one segment is finally sorted, the process begins again by retrieving the subscripts of another unsorted portion of the array. On completion,  $A_j \leq A_i$  for  $j < i$ . For more details, see Singleton (1969), Griffin and Redish (1970), and Petro (1970).

### Example

This example sorts the 10-element array *RA* algebraically.

```
C                                     Declare variables
PARAMETER (N=10)
REAL      RA(N), RB(N)

C                                     Set values for RA
C      RA = ( -1.0  2.0  -3.0  4.0  -5.0  6.0  -7.0  8.0  -9.0  10.0 )
C
DATA RA/-1.0, 2.0, -3.0, 4.0, -5.0, 6.0, -7.0, 8.0, -9.0, 10.0/
```

```

C          Sort RA by algebraic value into RB
C          CALL SVRGN (N, RA, RB)
C          Print results
C          CALL UMACH (2,NOUT)
C          WRITE (NOUT, 99999) (RB(J),J=1,N)
C
99999 FORMAT (' The output vector is:', /, 10(1X,F5.1))
END

```

### Output

```

The output vector is:
-9.0 -7.0 -5.0 -3.0 -1.0  2.0  4.0  6.0  8.0 10.0

```

---

## SVRGP/DSVRGP (Single/Double precision)

Sort a real array by algebraically increasing value and return the permutation that rearranges the array.

### Usage

```
CALL SVRGP (N, RA, RB, IPERM)
```

### Arguments

*N* — Number of elements in the array to be sorted. (Input)

*RA* — Vector of length *N* containing the array to be sorted. (Input)

*RB* — Vector of length *N* containing the sorted array. (Output)

If *RA* is not needed, *RA* and *RB* can share the same storage locations.

*IPERM* — Vector of length *N*. (Input/Output)

On input, *IPERM* should be initialized to the values 1, 2, ..., *N*. On output, *IPERM* contains a record of permutations made on the vector *RA*.

### Comments

For wider applicability, integers (1, 2, ..., *N*) that are to be associated with *RA(I)* for *I* = 1, 2, ..., *N* may be entered into *IPERM(I)* in any order. Note that these integers must be unique.

### Algorithm

Routine *SVRGP* sorts the elements of an array, *A*, into ascending order by algebraic value, keeping a record in *P* of the permutations to the array *A*. That is, the elements of *P* are moved in the same manner as are the elements in *A* as *A* is being sorted. The routine *SVRGP* uses the algorithm discussed in *SVRGN* (page 1273). On completion,  $A_j \leq A_i$  for  $j < i$ .

### Example

This example sorts the 10-element array *RA* algebraically.

```

C                                     Declare variables
PARAMETER (N=10)
REAL      RA(N), RB(N)
INTEGER   IPERM(N)

C                                     Set values for RA and IPERM
C      RA      = ( 10.0  -9.0  8.0  -7.0  6.0  5.0  4.0  -3.0  -2.0  -1.0 )
C
C      IPERM = ( 1  2  3  4  5  6  7  8  9  10)
C
C      DATA RA/10.0, -9.0, 8.0, -7.0, 6.0, 5.0, 4.0, -3.0, -2.0, -1.0/
C      DATA IPERM/1, 2, 3, 4, 5, 6, 7, 8, 9, 10/
C                                     Sort RA by algebraic value into RB
CALL SVRGP (N, RA, RB, IPERM)

C                                     Print results
CALL UMACH (2,NOUT)
WRITE (NOUT, 99998) (RB(J),J=1,N)
WRITE (NOUT, 99999) (IPERM(J),J=1,N)

C
99998 FORMAT (' The output vector is:', /, 10(1X,F5.1))
99999 FORMAT (' The permutation vector is:', /, 10(1X,I5))
END

```

### Output

```

The output vector is:
-9.0  -7.0  -3.0  -2.0  -1.0  4.0  5.0  6.0  8.0  10.0
The permutation vector is:
  2    4    8    9    10    7    6    5    3    1

```

---

## SVIGN

Sort an integer array by algebraically increasing value.

### Usage

```
CALL SVIGN (N, IA, IB)
```

### Arguments

*N* — Number of elements in the array to be sorted. (Input)

*IA* — Integer vector of length *N* containing the array to be sorted. (Input)

*IB* — Integer vector of length *N* containing the sorted array. (Output)

If *IA* is not needed, *IA* and *IB* can share the same storage locations.

### Algorithm

Routine *SVIGN* sorts the elements of an integer array, *A*, into ascending order by algebraic value. The routine *SVIGN* uses the algorithm discussed in *SVRGN* (page 1273). On completion,  $A_j \leq A_i$  for  $j < i$ .

### Example

This example sorts the 10-element array *IA* algebraically.

```

C                                     Declare variables
C   PARAMETER (N=10)
C   INTEGER   IA(N), IB(N)
C                                     Set values for IA
C   IA = ( -1  2  -3  4  -5  6  -7  8  -9  10 )
C
C   DATA IA/-1, 2, -3, 4, -5, 6, -7, 8, -9, 10/
C                                     Sort IA by algebraic value into IB
C   CALL SVIGN (N, IA, IB)
C
C                                     Print results
C   CALL UMACH (2,NOUT)
C   WRITE (NOUT, 99999) (IB(J),J=1,N)
C
C 99999 FORMAT (' The output vector is:', /, 10(1X,I5))
C END

```

### Output

```

The output vector is:
-9  -7  -5  -3  -1  2  4  6  8  10

```

---

## SVIGP

Sort an integer array by algebraically increasing value and return the permutation that rearranges the array.

### Usage

```
CALL SVIGP (N, IA, IB, IPERM)
```

### Arguments

*N* — Number of elements in the array to be sorted. (Input)

*IA* — Integer vector of length *N* containing the array to be sorted. (Input)

*IB* — Integer vector of length *N* containing the sorted array. (Output)

If *IA* is not needed, *IA* and *IB* can share the same storage locations.

*IPERM* — Vector of length *N*. (Input/Output)

On input, *IPERM* should be initialized to the values 1, 2, ..., *N*. On output, *IPERM* contains a record of permutations made on the vector *IA*.

### Comments

For wider applicability, integers (1, 2, ..., *N*) that are to be associated with *IA(I)* for *I* = 1, 2, ..., *N* may be entered into *IPERM(I)* in any order. Note that these integers must be unique.

### Algorithm

Routine *SVIGP* sorts the elements of an integer array, *A*, into ascending order by algebraic value, keeping a record in *P* of the permutations to the array *A*. That is, the elements of *P* are moved in the same manner as are the elements in *A* as

$A$  is being sorted. The routine `SVIGP` uses the algorithm discussed in `SVRGN` (page 1273). On completion,  $A_j \leq A_i$  for  $j < i$ .

### Example

This example sorts the 10-element array `IA` algebraically.

```
C                               Declare variables
PARAMETER (N=10)
INTEGER   IA(N), IB(N), IPERM(N)
C                               Set values for IA and IPERM
C   IA   = ( 10  -9  8  -7  6  5  4  -3  -2  -1 )
C
C   IPERM = ( 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 )
C
DATA IA/10, -9, 8, -7, 6, 5, 4, -3, -2, -1/
DATA IPERM/1, 2, 3, 4, 5, 6, 7, 8, 9, 10/
C                               Sort IA by algebraic value into IB
CALL SVIGP (N, IA, IB, IPERM)
C                               Print results
CALL UMACH (2,NOUT)
WRITE (NOUT, 99998) (IB(J),J=1,N)
WRITE (NOUT, 99999) (IPERM(J),J=1,N)
C
99998 FORMAT (' The output vector is:', /, 10(1X,I5))
99999 FORMAT (' The permutation vector is:', /, 10(1X,I5))
END
```

### Output

```
The output vector is:
-9  -7  -3  -2  -1  4  5  6  8  10
The permutation vector is:
2  4  8  9  10  7  6  5  3  1
```

---

## SCOLR/DSCOLR (Single/Double precision)

Sort columns of a real rectangular matrix using keys in rows.

### Usage

```
CALL SCOLR (NRX, NCX, X, LDX, ICOMP, IORDR, IRET, NKEY,
            INDKEY, IPERM, NGROUP, NI)
```

### Arguments

**NRX** — Number of rows of  $X$ . (Input)

**NCX** — Number of columns of  $X$ . (Input)

**X** —  $NRX$  by  $NCX$  matrix. (Input, if `IRET = 1`; input/output if `IRET = 0`)

On input,  $X$  contains the matrix to be sorted. If `IRET = 0`, the output  $X$  contains the sorted matrix.



**LDX** — Leading dimension of  $x$  exactly as specified in the dimension statement in the calling program. (Input)

**ICOMP** — Option giving the method of comparison of the column vectors. (Input)

**ICOMP Action**

0 Elementwise, by algebraic values  
1 Elementwise, by absolute values

**IORDR** — Option giving the sorting order. (Input)

**IORDR Action**

0 Ascending  
1 Descending

**IRET** — Option for determining whether the columns of  $x$  are to be permuted. (Input)

**IRET Action**

0 The columns of  $x$  are sorted.  
1  $x$  is unchanged (detached key sort).

**NKEY** — Number of rows of  $x$  on which to sort. (Input)

**INDKEY** — Vector of length  $NKEY$  giving the row numbers of  $x$  which are to be used in the sort. (Input)

**IPERM** — Permutation vector of length  $NCX$  specifying the rearrangement of the columns. (Output)

$IPERM(I) = J$  means column  $I$  of the sorted  $x$  is column  $J$  of the unsorted  $x$ .

**NGROUP** — Number of groups. (Output)

The columns of the sorted  $x$  are partitioned into groups. A group contains columns that are equal with respect to the method of comparison.  $NGROUP$  is the number of groups of different columns.

**NI** — Vector of length  $NGROUP$  containing the number of columns in each group. (Output)

The first  $NI(1)$  columns of the sorted  $x$  are group number 1; the next  $NI(2)$  columns of the sorted  $x$  are group number 2;... the last  $NI(NGROUP)$  columns of the sorted  $x$  are group number  $NGROUP$ . If  $NGROUP$  is not known prior to the invocation of this routine,  $NCX$  (an upper bound for  $NGROUP$ ) can be used as the dimension of  $NI$ .

### Comments

1. Automatic workspace usage is

SCOLR  $3 * m + INT(2.8854 * \ln(m)) + 2$  units, or

DSCOLR  $5 * m + INT(2.8854 * \ln(m)) + 2$  units,

where  $m = \max(NRX, NCX)$ . Workspace may be explicitly provided, if desired, by use of  $S2OLR/DS2OLR$ . The reference is

```
CALL S2OLR (NRX, NCX, X, LDX, ICOMP, IORDR, IRET,
           NKEY, INDKEY, IPERM, NGROUP, NI, WK,
           IWK)
```

The additional arguments are as follows:

**WK** — Work vector of length  $2 * m$ .

**IWK** — Work vector of length  $m + \text{INT}(2.8854 \ln(m)) + 2$ .

- When  $X$  is sorted by algebraic value ( $\text{ICOMP} = 0$ ) in ascending order, the resulting array  $X$  is such that:

For  $i = 1, 2, \dots, \text{NCX} - 1$ ,  $X(\text{INDKEY}(1), i) \leq X(\text{INDKEY}(1), i + 1)$

For  $k = 2, \dots, \text{NKEY}$ , if  $X(\text{INDKEY}(j), i) = X(\text{INDKEY}(j), i + 1)$  for  $j = 1, 2, \dots, k - 1$ , then  $X(\text{INDKEY}(k), i) \leq X(\text{INDKEY}(k), i + 1)$ .

When  $\text{ICOMP} = 1$ , the absolute values are compared instead.

### Algorithm

Routine `SCOLR` sorts the columns of a real matrix  $X$  using particular rows in  $X$  as the keys. One of two methods for comparing the columns can be used for sorting.

- Algebraic with the first key as the most significant, the second key next most significant and so forth.
- Absolute values with the first key as the most significant, the second key next most significant and so forth.

The columns of  $X$  can be put in ascending or descending order.

The routine is useful for data containing classification variables. Routine `CSTAT` (page 54) can be used to form the cells and frequency counts for a multi-way table from data. The columns of the output matrix contain the values of each combination of values of the classification variables along with the tallies. `SCOLR` can then be used to sort the columns of this output matrix using the classification variables as keys.

`SCOLR` is based on a quicksort method given by Singleton (1969). Modifications by Griffin and Redish (1970) and Petro (1970) are incorporated.

### Example

The columns of a  $5 \times 10$  matrix  $X$  are sorted in descending order by absolute value using rows 1, 2, 3, and 5 as the keys. The permutations to put the columns of  $X$  in order are returned. The input matrix  $X$  is not changed.

```
C
INTEGER      LDX, NCX, NKEY, NRX
PARAMETER    (NCX=10, NKEY=4, NRX=5, LDX=NRX)

INTEGER      ICOMP, INDKEY(NKEY), IORDR, IPERM(NCX), IRET, NI(NCX),
&            NGROUP, NOUT
REAL         X(LDX,NCX)
CHARACTER    CLABEL(1)*10, FMT*10, RLABEL(1)*23
EXTERNAL     SCOLR, UMACH, WRIRL, WRRRL
```

```

C
  DATA CLABEL(1)/'NONE'/, RLABEL(1)/'NONE'/
  DATA X/-1.0, -10.0, -11.0, 10.0, -1.0, 2.0, 20.0, 22.0, -20.0,
&      -2.0, -3.0, -30.0, 33.0, 30.0, -3.0, 4.0, 40.0, 44.0,
&      -40.0, -4.0, -5.0, -50.0, 55.0, 50.0, -5.0, -1.0, 60.0,
&      -66.0, -60.0, 6.0, 2.0, -70.0, -77.0, 70.0, 7.0, -3.0,
&      -30.0, -88.0, 80.0, 8.0, 4.0, 40.0, -99.0, -90.0, 9.0,
&      -5.0, -50.0, -100.0, 100.0, 10.0/
  DATA INDKEY/1, 2, 3, 5/

C
  ICOMP = 1
  IORDR = 1
  IRET = 1
  CALL SCOLR (NRX, NCX, X, LDX, ICOMP, IORDR, IRET, NKEY, INDKEY,
&            IPERM, NGROUP, NI)

C
  FMT      = '(F6.1)'
  RLABEL(1) = 'NONE'
  CALL WRRRL ('X', NRX, NCX, X, LDX, 0, FMT, RLABEL, CLABEL)

C
  FMT      = '(I4)'
  RLABEL(1) = 'IPERM = '
  CALL WRIRL ('%/', 1, NCX, IPERM, 1, 0, FMT, RLABEL, CLABEL)

C
  CALL UMACH (2, NOUT)
  WRITE (NOUT,*)
  WRITE (NOUT,*) 'NGROUP = ', NGROUP

C
  RLABEL(1) = 'NI = '
  CALL WRIRL ('%/', 1, NGROUP, NI, 1, 0, FMT, RLABEL, CLABEL)

C
  END

```

### Output

	X									
-1.0	2.0	-3.0	4.0	-5.0	-1.0	2.0	-3.0	4.0	-5.0	
-10.0	20.0	-30.0	40.0	-50.0	60.0	-70.0	-30.0	40.0	-50.0	
-11.0	22.0	33.0	44.0	55.0	-66.0	-77.0	-88.0	-99.0	-100.0	
10.0	-20.0	30.0	-40.0	50.0	-60.0	70.0	80.0	-90.0	100.0	
-1.0	-2.0	-3.0	-4.0	-5.0	6.0	7.0	8.0	9.0	10.0	
IPERM =	10	5	9	4	8	3	7	2	6	1
NGROUP =	10									
NI =	1	1	1	1	1	1	1	1	1	1

---

## SROWR/DSROWR (Single/Double precision)

Sort rows of a real rectangular matrix using keys in columns.

### Usage

```

CALL SROWR (NROW, NCOL, X, LDX, ICOMP, IORDR, IRET, NKEY,
            INDKEY, IPERM, NGROUP, NI, NRMIS)

```

## Arguments

**NROW** — Number of rows of  $x$ . (Input)

**NCOL** — Number of columns of  $x$ . (Input)

**X** —  $NROW$  by  $NCOL$  matrix. (Input, if  $IRET = 1$ ; input/output if  $IRET = 0$ )

On input,  $x$  contains the matrix to be sorted. If  $IRET = 0$ , the output  $x$  contains the sorted matrix.

**LDX** — Leading dimension of  $x$  exactly as specified in the dimension statement of the calling program. (Input)

**ICOMP** — Option giving the method of comparison of the row vectors. (Input)

### **ICOMP Action**

0 Elementwise, by algebraic values

1 Elementwise, by absolute values

**IORDR** — Option giving the sorting order. (Input)

### **IORDR Action**

0 Ascending

1 Descending

**IRET** — Option to indicate information returned. (Input)

### **IRET Action**

0 The sorted  $x$  is returned along with  $NGROUP$  and  $NI$ .

1  $x$  is unchanged (detached key sort) and  $NGROUP$  and  $NI$  are returned.

2 The sorted  $x$  is returned, but  $NGROUP$  and  $NI$  are not returned.

3  $x$  is unchanged (detached key sort) and  $NGROUP$  and  $NI$  are not returned.

**NKEY** — Number of columns of  $x$  on which to sort. (Input)

**INDKEY** — Vector of length  $NKEY$  giving the column numbers of  $x$  which are to be used in the sort. (Input)

**IPERM** — Permutation vector of length  $NROW$  specifying the rearrangement of the rows. (Output)

$IPERM(I) = J$  means row  $I$  of the sorted  $x$  is row  $J$  of the unsorted  $x$ .

**NGROUP** — Number of groups. (Output, if  $IRET \leq 1$ )

The rows of the sorted  $x$  are partitioned into groups. A group contains rows that are equal with respect to the method of comparison.  $NGROUP$  is the number of groups of different rows.

**NI** — Vector of length  $NGROUP$  containing the number of rows in each group. (Output, if  $IRET \leq 1$ )

The first  $NI(1)$  rows of the sorted  $x$  are group number 1. The next  $NI(2)$  rows of the sorted  $x$  are group number 2. ... The last  $NI(NGROUP)$  rows of the sorted  $x$  are group number  $NGROUP$ . If  $NGROUP$  is not known prior to the invocation of

this routine, *NROW* (an upper bound for *NGROUP*) can be used as the dimension of *NI*. If *IRET*  $\geq 2$ , *NI* is not referenced and can be a vector of length one.

***NRMISS*** — Number of rows that contained NaN in the columns of *x* used in the sort. (Output)

These rows are considered as a separate group from the other *NGROUP* groups and are put as the last *NRMISS* rows of the sorted *x*.

### Comments

1. Automatic workspace usage is

*SROWR*  $3 * m + \text{INT}(2.8854 * \ln(m)) + 2$  units, or

*DSROWR*  $5 * m + \text{INT}(2.8854 * \ln(m)) + 2$  units,

where  $m = \max(\text{NROW}, \text{NCOL})$ . Workspace may be explicitly provided, if desired, by use of *S2OWR/DS2OWR*. The reference is

```
CALL S2OWR (NROW, NCOL, X, LDX, ICOMP, IORDER, IRET,
           NKEY, INDKEY, IPERM, NGROUP, NI, NRMISS,
           WK, IWK)
```

The additional arguments are as follows:

***WK*** — Work vector of length  $2 * m$ .

***IWK*** — Work vector of length  $m + \text{INT}(2.8854 * \ln(m)) + 2$ .

2. When *x* is sorted by algebraic values (*ICOMP* = 0), in ascending order, the resulting array *x* is such that:

For  $i = 1, 2, \dots, \text{NROW} - 1$ ,  $x(i, \text{INDKEY}(1)) \leq x(i + 1, \text{INDKEY}(1))$ .

For  $k = 2, \dots, \text{NKEY}$ , if  $x(i, \text{INDKEY}(j)) = x(i + 1, \text{INDKEY}(j))$  for  $j = 1, 2, \dots, k - 1$ ; then  $x(i, \text{INDKEY}(k)) \leq x(i + 1, \text{INDKEY}(k))$ .

When *ICOMP* = 1, the absolute values are compared instead.

### Algorithm

Routine *SROWR* sorts the rows of a real matrix *x* using particular rows in *x* as the keys. One of two methods for comparing the rows can be used for sorting.

1. Algebraic with the first key as the most significant, the second key next most significant and so forth.
2. Absolute values with the first key as the most significant, the second key next most significant and so forth.

The rows of *x* can be put in ascending or descending order.

The routine is useful for grouping data based on values of specified variables. The rows of *x* containing the IMSL missing value code NaN (not a number) in at least one of the specified columns are considered as an additional group of *NRMISS* rows. These rows are moved to the end of the sorted *x*. *SROWR* is based on a quicksort method given by Singleton (1969). Modifications by Griffin and Redish (1970) and Petro (1970) are incorporated.

### Example

The rows of a  $10 \times 3$  matrix  $x$  are sorted in ascending order by algebraic value using columns 2 and 3 as the keys. The permutations to put the rows of the input  $x$  into sorted order are returned along with the sorted  $x$ .

```
INTEGER      LDX, NCOL, NKEY, NROW
PARAMETER    (NCOL=3, NKEY=2, NROW=10, LDX=NROW)

C
INTEGER      ICOMP, INDKEY(NKEY), IORDR, IPERM(NROW), IRET,
&           NGROUP, NI(NROW), NOUT, NRMISS
REAL         AMACH, X(LDX,NCOL)
EXTERNAL     AMACH, SROWR, UMACH, WRIRN, WRRRN

C
DATA (X(1,J),J=1,3)/1.0, 1., 1./
DATA (X(2,J),J=1,3)/2.0, 2., 1./
DATA (X(3,J),J=1,3)/3.0, 1., 1./
DATA (X(4,J),J=1,3)/4.0, 1., 1./
DATA (X(5,J),J=1,3)/5.0, 2., 2./
DATA (X(6,J),J=1,3)/6.0, 1., 2./
DATA (X(7,J),J=1,3)/7.0, 1., 2./
DATA (X(8,J),J=1,3)/8.0, 1., 1./
DATA (X(9,J),J=1,3)/9.0, 2., 2./
DATA (X(10,J),J=1,3)/9.0, 1., 1./
DATA INDKEY/2, 3/

C
X(5,3) = AMACH(6)
X(7,2) = AMACH(6)
IRET   = 0
ICOMP  = 0
IORDR  = 0
CALL SROWR (NROW, NCOL, X, LDX, ICOMP, IORDR, IRET, NKEY,
&          INDKEY, IPERM, NGROUP, NI, NRMISS)
CALL WRRRN ('X', NROW, NCOL, X, LDX, 0)
CALL WRIRN ('IPERM', NROW, 1, IPERM, NROW, 0)
CALL WRIRN ('NI', NGROUP, 1, NI, NGROUP, 0)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) ' '
WRITE (NOUT,*) 'NRMISS = ', NRMISS
END
```

### Output

```
      X
      1      2      3
1  1.000  1.000  1.000
2  9.000  1.000  1.000
3  3.000  1.000  1.000
4  4.000  1.000  1.000
5  8.000  1.000  1.000
6  6.000  1.000  2.000
7  2.000  2.000  1.000
8  9.000  2.000  2.000
9  7.000   NaN  2.000
10 5.000  2.000   NaN
```

```
IPERM
1    1
```

```

2 10
3 3
4 4
5 8
6 6
7 2
8 9
9 7
10 5

```

```

      NI
1     5
2     1
3     1
4     1
NRMISS = 2

```

---

## SRCH/DSRCH (Single/Double precision)

Search a sorted vector for a given scalar and return its index.

### Usage

```
CALL SRCH (N, VALUE, X, INCX, INDEX)
```

### Arguments

*N* — Length of vector *Y*. (Input)

*VALUE* — Scalar to be searched for in *Y*. (Input)

*X* — Vector of length  $N * INCX$ . (Input)

*Y* is obtained from *x* for  $I = 1, 2, \dots, N$  by  $Y(I) = X(1 + (I - 1) * INCX)$ .  $Y(1), Y(2), \dots, Y(N)$  must be in ascending order.

*INCX* — Displacement between elements of *x*. (Input)

*INCX* must be greater than zero.

*INDEX* — Index of *Y* pointing to *VALUE*. (Output)

If *INDEX* is positive, *VALUE* is found in *Y*. If *INDEX* is negative, *VALUE* is not found in *Y*.

<b>INDEX</b>	<b>Location of VALUE</b>
1 thru N	VALUE = Y(INDEX)
-1	VALUE < Y(1) or N = 0
-N thru -2	$Y(-INDEX - 1) < VALUE < Y(-INDEX)$
-(N + 1)	VALUE > Y(N)

### Algorithm

Routine SRCH searches a real vector *x* (stored in *X*), whose *n* elements are sorted in ascending order for a real number *c* (stored in *VALUE*). If *c* is found in *x*, its

index  $i$  (stored in INDEX) is returned so that  $x_i = c$ . Otherwise, a negative number  $i$  is returned for the index. Specifically,

```

if  $1 \leq i \leq n$  then  $x_i = c$ 
if  $i = -1$  then  $c < x_1$  or  $n = 0$ 
if  $-n \leq i \leq -2$  then  $x_{-i-1} < c < x_{-i}$ 
if  $i = -(n+1)$  then  $c > x_n$ 

```

The argument INCX is useful if a row of a matrix, for example, row number I of a matrix X, must be searched. The elements of row I are assumed to be in ascending order. In this case, set INCX equal to the leading dimension of X exactly as specified in the dimension statement in the calling program. With X declared

```
REAL X(LDX,N)
```

the invocation

```
CALL SRCH (N, VALUE, X(I,1), LDX, INDEX)
```

returns an index that will reference a column number of X.

Routine SRCH performs a binary search. The routine is an implementation of Algorithm B discussed by Knuth (1973, pages 407–411).

### Example

This example searches a real vector sorted in ascending order for the value 653.0. The problem is discussed by Knuth (1973, pages 407–409).

```

C      INTEGER      N
C      PARAMETER   (N=16)

C      INTEGER      INCX, INDEX, NOUT
C      REAL         VALUE, X(N)
C      EXTERNAL     SRCH, UMACH

C      DATA X/61.0, 87.0, 154.0, 170.0, 275.0, 426.0, 503.0, 509.0,
C      &          512.0, 612.0, 653.0, 677.0, 703.0, 765.0, 897.0, 908.0/

C      INCX = 1
C      VALUE = 653.0
C      CALL SRCH (N, VALUE, X, INCX, INDEX)

C      CALL UMACH (2, NOUT)
C      WRITE (NOUT,*) 'INDEX = ', INDEX
C      END

```

### Output

```
INDEX = 11
```



---

# ISRCH

Search a sorted integer vector for a given integer and return its index.

## Usage

```
CALL ISRCH (N, IVALUE, IX, INCX, INDEX)
```

## Arguments

*N* — Length of vector *IY*. (Input)

*IVALUE* — Scalar to be searched for in *IY*. (Input)

*IX* — Vector of length  $N * INCX$ . (Input)

*IY* is obtained from *IX* for  $I = 1, 2, \dots, N$  by  $IY(I) = IX(1 + (I - 1) * INCX)$ .  
*IY*(1), *IY*(2), ..., *IY*(*N*) must be in ascending order.

*INCX* — Displacement between elements of *IX*. (Input)  
*INCX* must be greater than zero.

*INDEX* — Index of *IY* pointing to *IVALUE*. (Output)

If *INDEX* is positive, *IVALUE* is found in *IY*. If *INDEX* is negative, *IVALUE* is not found in *IY*.

<i>INDEX</i>	Location of <i>IVALUE</i>
1 thru <i>N</i>	$IVALUE = IY(INDEX)$
-1	$IVALUE < IY(1)$ or $N = 0$
- <i>N</i> thru -2	$IY(-INDEX - 1) < IVALUE < IY(-INDEX)$
-( <i>N</i> + 1)	$IVALUE > IY(N)$

## Algorithm

Routine *ISRCH* searches an integer vector  $x$  (stored in *IX*), whose  $n$  elements are sorted in ascending order for an integer  $c$  (stored in *IVALUE*). If  $c$  is found in  $x$ , its index  $i$  (stored in *INDEX*) is returned so that  $x_i = c$ . Otherwise, a negative number  $i$  is returned for the index.

Specifically,

if $1 \leq i \leq n$	then	$x_i = c$
if $i = -1$	then	$c < x_1$ or $n = 0$
if $-n \leq i \leq -2$	then	$x_{-i-1} < c < x_{-i}$
if $i = -(n + 1)$	then	$c > x_n$

The argument *INCX* is useful if a row of a matrix, for example, row number *I* of a matrix *IX*, must be searched. The elements of row *I* are assumed to be in ascending order. Here, set *INCX* equal to the leading dimension of *IX* exactly as specified in the dimension statement in the calling program. With *IX* declared

```
INTEGER IX(LDIX,N)
```

the invocation

```
CALL ISRCH (N, IVALUE, IX(I,1), LDIX, INDEX)
```

returns an index that will reference a column number of *IX*. The routine *ISRCH* performs a binary search. The routine is an implementation of Algorithm B discussed by Knuth (1973, pages 407–411).

### Example

This example searches an integer vector sorted in ascending order for the value 653. The problem is discussed by Knuth (1973, pages 407–409).

```
C
INTEGER      N
PARAMETER   (N=16)

C
INTEGER      INCX, INDEX, NOUT
INTEGER      IVALUE, IX(N)
EXTERNAL     ISRCH, UMACH

C
DATA IX/61, 87, 154, 170, 275, 426, 503, 509, 512, 612, 653, 677,
&          703, 765, 897, 908/

C
INCX = 1
IVALUE = 653
CALL ISRCH (N, IVALUE, IX, INCX, INDEX)

C
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'INDEX = ', INDEX
END
```

### Output

```
INDEX =    11
```

---

## SSRCH

Search a character vector, sorted in ascending ASCII order, for a given string and return its index.

### Usage

```
CALL SSRCH (N, STRING, CHX, INCX, INDEX)
```

### Arguments

*N* — Length of vector *CHX*. (Input)

*STRING* — Character string to be searched for in *CHX*. (Input)

*CHX* — Vector of length  $N * INCX$  containing character strings. (Input)  
*CHY* is obtained from *CHX* for  $I = 1, 2, \dots, N$  by  $CHY(I) = CHX(1 + (I - 1) * INCX)$ . *CHY*(1), *CHY*(2), ..., *CHY*(*N*) must be in ascending ASCII order.

**INCX** — Displacement between elements of CHX. (Input)  
INCX must be greater than zero.

**INDEX** — Index of CHY pointing to STRING. (Output)  
If INDEX is positive, STRING is found in CHY. If INDEX is negative, STRING is not found in CHY.

<b>INDEX</b>	<b>Location of STRING</b>
1 thru N	STRING = CHY(INDEX)
-1	STRING < CHY(1) or N = 0
-N thru -2	CHY(-INDEX - 1) < STRING < CHY(-INDEX)
-(N + 1)	STRING > CHY(N)

### Algorithm

Routine SSRCH searches a vector of character strings  $x$  (stored in CHX), whose  $n$  elements are sorted in ascending ASCII order, for a character string  $c$  (stored in STRING). If  $c$  is found in  $x$ , its index  $i$  (stored in INDEX) is returned so that  $x_i = c$ . Otherwise, a negative number  $i$  is returned for the index. Specifically,

if  $1 \leq i \leq n$  then  $x_i = c$   
if  $i = -1$  then  $c < x_1$  or  $n = 0$   
if  $-n \leq i \leq -2$  then  $x_{-i-1} < c < x_{-i}$   
if  $i = -(n + 1)$  then  $c > x_n$

Here, “<” and “>” are in reference to the ASCII collating sequence. For comparisons made between character strings  $c$  and  $x_i$  with different lengths, the shorter string is considered as if it were extended on the right with blanks to the length of the longer string. (SSRCH uses FORTRAN intrinsic functions LLT and LGT.)

The argument INCX is useful if a row of a matrix, for example, row number I of a matrix CHX, must be searched. The elements of row I are assumed to be in ascending ASCII order. In this case, set INCX equal to the leading dimension of CHX exactly as specified in the dimension statement in the calling program. With CHX declared

```
CHARACTER*7 CHX(LDCHX,N)
```

the invocation

```
CALL SSRCH (N, STRING, CHX(I,1), LDCHX, INDEX)
```

returns an index that will reference a column number of CHX.

Routine SSRCH performs a binary search. The routine is an implementation of Algorithm B discussed by Knuth (1973, pages 407–411).

### Example

This example searches a CHARACTER\*2 vector containing 9 character strings, sorted in ascending ASCII order, for the value 'CC'.

```
INTEGER      N
PARAMETER   (N=9)

C
INTEGER      INCX, INDEX, NOUT
CHARACTER    CHX(N)*2, STRING*2
EXTERNAL     SSRCH, UMACH

C
DATA CHX/'AA', 'BB', 'CC', 'DD', 'EE', 'FF', 'GG', 'HH',
&      'II'/

C
INCX      = 1
STRING    = 'CC'
CALL SSRCH (N, STRING, CHX, INCX, INDEX)

C
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'INDEX = ', INDEX
END
```

### Output

```
INDEX =      3
```

---

## ACHAR

Return a character given its ASCII value.

### Usage

```
ACHAR(I)
```

### Arguments

*I* — Integer ASCII value of the character desired. (Input)  
*I* must be greater than or equal to zero and less than or equal to 127.

*ACHAR* — CHARACTER \* 1 string containing the character in the *I*-th position of the ASCII collating sequence. (Output)

### Algorithm

Routine ACHAR returns the character of the input ASCII value. The input value should be between 0 and 127. If the input value is out of range, the value returned in ACHAR is machine dependent.

### Example

This example returns the character of the ASCII value 65.

```
INTEGER      I, NOUT
CHARACTER    ACHAR
```

```

EXTERNAL  ACHAR, UMACH
C
CALL UMACH (2, NOUT)
C                                     Get character for ASCII value
C                                     of 65 ('A')
      I = 65
      WRITE (NOUT,99999) I, ACHAR(I)
C
99999 FORMAT (' For the ASCII value of ', I2, ', the character is : ',
&           A1)
      END

```

### Output

For the ASCII value of 65, the character is : A

## IACHAR

Return the integer ASCII value of a character argument.

### Usage

IACHAR(CH)

### Arguments

*CH* — Character argument for which the integer ASCII value is desired. (Input)

*IACHAR* — Integer ASCII value for CH. (Output)

The character CH is in the *IACHAR*-th position of the ASCII collating sequence.

### Algorithm

Routine IACHAR returns the ASCII value of the input character.

### Example

This example gives the ASCII value of character A.

```

INTEGER  IACHAR, NOUT
CHARACTER CH
EXTERNAL IACHAR, UMACH
C
CALL UMACH (2, NOUT)
C                                     Get ASCII value for the character
C                                     'A'.
      CH = 'A'
      WRITE (NOUT,99999) CH, IACHAR(CH)
C
99999 FORMAT (' For the character ', A1, ' the ASCII value is : ',
&           I3)
      END

```

## Output

For the character A the ASCII value is : 65

---

# ICASE

Return the ASCII value of a character converted to uppercase.

## Usage

ICASE(CH)

## Arguments

*CH* — Character to be converted. (Input)

*ICASE* — Integer ASCII value for CH without regard to the case of CH. (Output)

Routine ICASE returns the same value as IACHAR (page 1290) for all but lowercase letters. For these, it returns the IACHAR value for the corresponding uppercase letter.

## Algorithm

Routine ICASE converts a character to its integer ASCII value. The conversion is case insensitive; that is, it returns the ASCII value of the corresponding uppercase letter for a lowercase letter.

## Example

This example shows the case insensitive conversion.

```
INTEGER      ICASE, NOUT
CHARACTER    CH
EXTERNAL     ICASE, UMACH
C
CALL UMACH (2, NOUT)           Get output unit number
C
C                               Get ASCII value for the character
C                               'a'.
CH = 'a'
WRITE (NOUT,99999) CH, ICASE(CH)
C
99999 FORMAT (' For the character ', A1, ' the ICASE value is : ',
&           I3)
END
```

## Output

For the character a the ICASE value is : 65

---

# IICSR

Compare two character strings using the ASCII collating sequence but without regard to case.

## Usage

IICSR(STR1, STR2)

## Arguments

**STR1** — First character string. (Input)

**STR2** — Second character string. (Input)

**IICSR** — Comparison indicator. (Output)

Let USTR1 and USTR2 be the uppercase versions of STR1 and STR2, respectively. The following table indicates the relationship between USTR1 and USTR2 as determined by the ASCII collating sequence.

### IICSR Meaning

-1	USTR1 precedes USTR2
0	USTR1 equals USTR2
1	USTR1 follows USTR2

## Comments

If the two strings, STR1 and STR2, are of unequal length, the shorter string is considered as if it were extended with blanks to the length of the longer string.

## Algorithm

Routine IICSR compares two character strings. It returns -1 if the first string is less than the second string, 0 if they are equal, and 1 if the first string is greater than the second string. The comparison is case insensitive.

## Example

This example shows different cases on comparing two strings.

```
INTEGER      IICSR, NOUT
CHARACTER    STR1*6, STR2*6
EXTERNAL     IICSR, UMACH
C
CALL UMACH (2, NOUT)
C
C
C      STR1 = 'ABc 1'
C      STR2 = ' '
C      WRITE (NOUT,99999) STR1, STR2, IICSR(STR1,STR2)
C
C
C      STR1 = 'AbC'
C      STR2 = 'ABc'
```

```

WRITE (NOUT,99999) STR1, STR2, IICSR(STR1,STR2)
C
C                               String1 is 'smaller' than String2
      STR1 = 'ABC'
      STR2 = 'aBC 1'
      WRITE (NOUT,99999) STR1, STR2, IICSR(STR1,STR2)
C
99999 FORMAT (' For String1 = ', A6, 'and String2 = ', A6,
&           ' IICSR = ', I2, '/')
      END

```

### Output

```

For String1 = ABC 1 and String2 =           IICSR = 1
For String1 = AbC   and String2 = ABC      IICSR = 0
For String1 = ABC   and String2 = aBC 1    IICSR = -1

```

---

## IIDEX

Determine the position in a string at which a given character sequence begins without regard to case.

### Usage

IIDEX(CHRSTR, KEY)

### Arguments

**CHRSTR** — Character string to be searched. (Input)

**KEY** — Character string that contains the key sequence. (Input)

**IIDEX** — Position in CHRSTR where KEY begins. (Output)

If KEY occurs more than once in CHRSTR, the starting position of the first occurrence is returned. If KEY does not occur in CHRSTR, then IIDEX returns a zero.

### Comments

If the length of KEY is greater than the length CHRSTR, IIDEX returns a zero.

### Algorithm

Routine IIDEX searches for a key string in a given string and returns the index of the starting element at which the key character string begins. It returns 0 if there is no match. The comparison is case insensitive. For a case-sensitive version, use the FORTRAN 77 intrinsic function INDEX.

### Example

This example locates a key string.

```

INTEGER      IIDEX, NOUT
CHARACTER    KEY*5, STRING*10

```



```

EXTERNAL  IINDEX, UMACH
C          Get output unit number
CALL UMACH (2, NOUT)
C          Locate KEY in STRING
STRING = 'alb2c3d4e5'
KEY     = 'C3d4E'
WRITE (NOUT,99999) STRING, KEY, IINDEX(STRING,KEY)
C
KEY = 'F'
WRITE (NOUT,99999) STRING, KEY, IINDEX(STRING,KEY)
C
99999 FORMAT (' For STRING = ', A10, ' and KEY = ', A5, ' IINDEX = ', I2,
&           /)
END

```

### Output

```

For STRING = alb2c3d4e5 and KEY = C3d4E IINDEX = 5
For STRING = alb2c3d4e5 and KEY = F      IINDEX = 0

```

## CVTSI

Convert a character string containing an integer number into the corresponding integer form.

### Usage

```
CALL CVTSI (STRING, NUMBER)
```

### Arguments

**STRING** — Character string containing an integer number. (Input)

**NUMBER** — The integer equivalent of **STRING**. (Output)

### Algorithm

Routine CVTSI converts a character string containing an integer to an INTEGER variable. Leading and trailing blanks in the string are ignored. If the string contains something other than an integer, a terminal error is issued. If the string contains an integer larger than can be represented by an INTEGER variable as determined from routine IMACH (Reference Material), a terminal error is issued.

### Example

The string “12345” is converted to an INTEGER variable.

```

INTEGER  NOUT, NUMBER
CHARACTER STRING*10
EXTERNAL CVTSI, UMACH
C
DATA STRING/'12345'/
C
CALL CVTSI (STRING, NUMBER)
C

```

```

CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'NUMBER = ', NUMBER
END

```

### Output

```
NUMBER = 12345
```

## CPSEC

Return CPU time used in seconds.

### Usage

```
CPSEC ( )
```

### Argument

*CPSEC* — CPU time used (in seconds) since first call to *CPSEC*. (Output)

### Comments

1. The first call to *CPSEC* returns 0.0.
2. The accuracy of this routine depends on the hardware and the operating system. On some systems, identical runs can produce timings differing by more than 10 percent.

### Algorithm

Function *CPSEC* is used to find the CPU time used in a particular section of FORTRAN code. The first call to *CPSEC* in a program returns *CPSEC*( ) = 0.0. Subsequent calls to *CPSEC* return the CPU time (in seconds) since the first call.

### Example

The following example uses *CPSEC* to return the CPU time to compute 1 million additions in a DO loop on a particular computer. The program was compiled without optimization.

```

C
INTEGER      I, NOUT
REAL         A, CPSEC, TIME0, TIME1, X, Y
EXTERNAL     CPSEC, UMACH

X           = 1.414
Y           = 3.142
TIME0 = CPSEC()
DO 10 I=1, 1000000
  A = X + Y
10 CONTINUE
TIME1 = CPSEC()
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'CPU time (seconds) = ', TIME1 - TIME0
END

```

## Output

CPU time (seconds) = 1.09000

---

# TIMDY

Get time of day.

## Usage

```
CALL TIMDY (Ihour, MINUTE, ISEC)
```

## Arguments

**Ihour** — Hour of the day. (Output)

Ihour is between 0 and 23 inclusive.

**MINUTE** — Minute within the hour. (Output)

MINUTE is between 0 and 59 inclusive.

**ISEC** — Second within the minute. (Output)

ISEC is between 0 and 59 inclusive.

## Algorithm

Routine TIMDY is used to retrieve the time of day.

## Example

The following example uses TIMDY to return the current time. Obviously, the output is dependent upon the time at which the program is run.

```
C
INTEGER      Ihour, IMIN, ISEC, NOUT
EXTERNAL     TIMDY, UMACH

CALL TIMDY (Ihour, IMIN, ISEC)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'Hour:Minute:Second = ', Ihour, ':', IMIN,
&              ':', ISEC
IF (Ihour .EQ. 0) THEN
  WRITE (NOUT,*) 'The time is ', IMIN, ' minute(s), ', ISEC,
&              ' second(s) past midnight.'
ELSE IF (Ihour .LT. 12) THEN
  WRITE (NOUT,*) 'The time is ', IMIN, ' minute(s), ', ISEC,
&              ' second(s) past ', Ihour, ' am.'
ELSE IF (Ihour .EQ. 12) THEN
  WRITE (NOUT,*) 'The time is ', IMIN, ' minute(s), ', ISEC,
&              ' second(s) past noon.'
ELSE
  WRITE (NOUT,*) 'The time is ', IMIN, ' minute(s), ', ISEC,
&              ' second(s) past ', Ihour-12, ' pm.'
END IF
END
```

### Output

```
Hour:Minute:Second = 16: 52: 29  
The time is 52 minute(s), 29 second(s) past 4 pm.
```

---

## TDATE

Get today's date.

### Usage

```
CALL TDATE (IDAY, MONTH, IYEAR)
```

### Arguments

**IDAY** — Day of the month. (Output)  
IDAY is between 1 and 31 inclusive.

**MONTH** — Month of the year. (Output)  
MONTH is between 1 and 12 inclusive.

**IYEAR** — Year. (Output)  
For example, IYEAR = 1985.

### Algorithm

Routine TDATE is used to retrieve today's date. Obviously, the output is dependent upon the date the program is run.

### Example

The following example uses TDATE to return today's date.

```
INTEGER      IDAY, IYEAR, MONTH, NOUT  
EXTERNAL    TDATE, UMACH  
C  
CALL TDATE (IDAY, MONTH, IYEAR)  
CALL UMACH (2, NOUT)  
WRITE (NOUT,*) 'Day-Month-Year = ', IDAY, '-', MONTH,  
&              '- ', IYEAR  
END
```

### Output

```
Day-Month-Year = 2- 4- 1991
```

---

## NDAYS

Compute the number of days from January 1, 1900, to the given date.

### Usage

```
NDAYS(IDAY, MONTH, IYEAR)
```

## Arguments

**IDAY** — Day of the input date. (Input)

**MONTH** — Month of the input date. (Input)

**IYEAR** — Year of the input date. (Input)

1950 would correspond to the year 1950 A.D. and 50 would correspond to year 50 A.D.

**NDAYS** — Function value. (Output)

If **NDAYS** is negative, it indicates the number of days prior to January 1, 1900.

## Comments

1. Informational error  
Type Code  
1 1 The Julian calendar, the first modern calendar, went into use in 45 B.C. No calendar prior to 45 B.C. was as universally used nor as accurate as the Julian. Therefore, it is assumed that the Julian calendar was in use prior to 45 B.C.
2. The number of days from one date to a second date can be computed by two references to **NDAYS** and then calculating the difference.
3. The beginning of the Gregorian calendar was the first day after October 4, 1582, which became October 15, 1582. Prior to that, the Julian calendar was in use. **NDAYS** makes the proper adjustment for the change in calendars.

## Algorithm

Function **NDAYS** returns the number of days from January 1, 1900, to the given date. The function **NDAYS** returns negative values for days prior to January 1, 1900. A negative **IYEAR** can be used to specify B.C. Input dates in year 0 and for October 5, 1582, through October 14, 1582, inclusive, do not exist; consequently, in these cases, **NDAYS** issues a terminal error.

## Example

The following example uses **NDAYS** to compute the number of days from January 15, 1986, to February 28, 1986:

```
C
INTEGER      IDAY, IYEAR, MONTH, NDAY0, NDAY1, NDAYS, NOUT
EXTERNAL    NDAYS, UMACH

IDAY  = 15
MONTH = 1
IYEAR = 1986
NDAY0 = NDAYS( IDAY, MONTH, IYEAR )
IDAY  = 28
MONTH = 2
IYEAR = 1986
```

```

NDAY1 = NDAYS(IDAY,MONTH,IYEAR)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'Number of days = ', NDAY1 - NDAY0
END

```

### Output

Number of days = 44

## NDYIN

Give the date corresponding to the number of days since January 1, 1900.

### Usage

```
CALL NDYIN (NDAYS, IDAY, MONTH, IYEAR)
```

### Arguments

*NDAYS* — Number of days since January 1, 1900. (Input)

*IDAY* — Day of the input date. (Output)

*MONTH* — Month of the input date. (Output)

*IYEAR* — Year of the input date. (Output)

1950 would correspond to the year 1950 A.D. and -50 would correspond to year 50 B.C.

### Comments

The beginning of the Gregorian calendar was the first day after October 4, 1582, which became October 15, 1582. Prior to that, the Julian calendar was in use. Routine NDYIN makes the proper adjustment for the change in calendars.

### Algorithm

Routine NDYIN computes the date corresponding to the number of days since January 1, 1900. For an input value of *NDAYS* that is negative, the date computed is prior to January 1, 1900. The routine NDYIN is the inverse of *NDAYS* (page 1297).

### Example

The following example uses NDYIN to compute the date for the 100th day of 1986. This is accomplished by first using *NDAYS* (see page 1297) to get the “day number” for December 31, 1985.

```

NDAY0 = NDAYS(31,12,1985)
CALL NDYIN (NDAY0+100, IDAY, MONTH, IYEAR)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'Day 100 of 1986 is (day-month-year) ', IDAY,
&          '- ', MONTH, '- ', IYEAR
END

```

## Output

Day 100 of 1986 is (day-month-year) 10- 4- 1986

---

# IDYWK

Compute the day of the week for a given date.

## Usage

IDYWK( IDAY, MONTH, IYEAR )

## Arguments

*IDAY* — Day of the input date. (Input)

*MONTH* — Month of the input date. (Input)

*IYEAR* — Year of the input date. (Input)

1950 would correspond to the year 1950 A.D. and 50 would correspond to year 50 A.D.

*IDYWK* — Function value. (Output)

The value of *IDYWK* ranges from 1 to 7, where 1 corresponds to Sunday and 7 corresponds to Saturday.

## Comments

1. Informational error  

Type	Code	
1	1	The Julian calendar, the first modern calendar, went into use in 45 B.C. No calendar prior to 45 B.C. was as universally used nor as accurate as the Julian. Therefore, it is assumed that the Julian calendar was in use prior to 45 B.C.
2. The beginning of the Gregorian calendar was the first day after October 4, 1582, which became October 15, 1582. Prior to that, the Julian calendar was in use. Function *IDYWK* makes the proper adjustment for the change in calendars.

## Algorithm

Function *IDYWK* returns an integer code that specifies the day of week for a given date. Sunday corresponds to 1, Monday corresponds to 2, and so forth. A negative *IYEAR* can be used to specify B.C. Input dates in year 0 and for October 5, 1582, through October 14, 1582, inclusive, do not exist; consequently, in these cases, *IDYWK* issues a terminal error.

### Example

The following example uses `IDYWK` to return the day of the week for February 24, 1963.

```
C
INTEGER    IDAY, IDYWK, IYEAR, MONTH, NOUT
EXTERNAL   IDYWK, UMACH

IDAY = 24
MONTH = 2
IYEAR = 1963
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'IDYWK (index for day of week) = ',
&              IDYWK(IDAY,MONTH,IYEAR)
END
```

### Output

```
IDYWK (index for day of week) = 1
```

---

## VERSL

Obtain STAT/LIBRARY-related version, system and serial numbers.

### Usage

```
VERSL( ISELECT)
```

### Arguments

*ISELECT* — Option for the information to retrieve. (Input)

**ISELECT** **VERSL**

- 1 IMSL STAT/LIBRARY version number
- 2 Operating system (and version number) for which the library was produced.
- 3 Fortran compiler (and version number) for which the library was produced.
- 4 IMSL STAT/LIBRARY serial number

*VERSL* — CHARACTER string containing information. (Output)

### Example

In this example, we print all of the information returned by `VERSL` on a particular machine. The output is omitted because the results are system dependent.

```
C
INTEGER    ISELECT, NOUT
CHARACTER  STRING(4)*50, TEMP*32, VERSL*32
EXTERNAL   UMACH, VERSL

STRING(1) = ((' IMSL STAT/LIBRARY Version Number: ', A)
STRING(2) = ((' Operating System ID Number: ', A)
STRING(3) = ((' Fortran Compiler Version Number: ', A)
```



```

STRING(4) = ((' IMSL STAT/LIBRARY Serial Number: ', A)
C
CALL UMACH (2, NOUT)          Print the versions and numbers.
DO 10 ISELCT=1, 4
    TEMP = VERSL(ISELCT)
    WRITE (NOUT,STRING(ISELCT)) TEMP
10 CONTINUE
END

```

---

## GDATA/DGDATA (Single/Double precision)

Retrieve a commonly analyzed data set.

### Usage

```
CALL GDATA (IDATA, IPRINT, NOBS, NVAR, X, LDX, NDX)
```

### Arguments

**IDATA** — Data set indicator. (Input)

<b>IDATA</b>	<b>NOBS</b>	<b>NVAR</b>	<b>Description of Data Set</b>
1	16	7	Longley
2	176	2	Wolfer sunspot
3	150	5	Fisher iris
4	144	1	Box and Jenkins Series G
5	13	5	Draper and Smith Appendix B
6	197	1	Box and Jenkins Series A
7	296	2	Box and Jenkins Series J
8	100	4	Robinson Multichannel Time Series
9	113	34	Afifi and Azen Data Set A

Set **IDATA** = 0 to print a description of all the data sets above. In this case, the remaining arguments are not referenced.

**IPRINT** — Printing option. (Input)

#### **IPRINT Action**

- 0 No printing is performed.
- 1 Rows 1 through 10 of **X** are printed.
- 2 All rows of **X** are printed.

When printing is performed, a header listing the data set name and a reference is printed.

**NOBS** — Number of observations or rows in the output matrix. (Output)

**NVAR** — Number of variables or columns in the output matrix. (Output)

**X** — **NOBS** by **NVAR** matrix containing the data set. (Output)

**LDX** — Leading dimension of  $x$  exactly as specified in the dimension statement in the calling program. (Input)

**NDX** — Second dimension of the matrix  $x$  exactly as specified in the dimension statement of the calling program. (Input)

### Algorithm

Routine `GDATA` retrieves a standard data set frequently cited in statistics textbooks or in this manual. The following table gives the references for each data set:

#### IDATA Reference

- 1 Longley (1967)
- 2 Anderson (1971, page 660)
- 3 Fisher (1936); Mardia, Kent, and Bibby (1979, Table 1.2.2)
- 4 Box and Jenkins (1976, page 531)
- 5 Draper and Smith (1981, pages 629–630)
- 6 Box and Jenkins (1976, page 525)
- 7 Box and Jenkins (1976, page 532–533)
- 8 Robinson (1967, page 204)
- 9 Afifi and Azen (1979, pages 16–22)

### Example

`GDATA` is used to copy the Longley data set into the matrix  $x$ .

```
C      INTEGER      LDX, NDX
      PARAMETER    (LDX=200, NDX=10)

C      INTEGER      IDATA, IPRINT, NOBS, NVAR
      REAL          X(LDX,NDX)
      EXTERNAL      GDATA

C      IDATA = 1
      IPRINT = 2
      CALL GDATA (IDATA, IPRINT, NOBS, NVAR, X, LDX, NDX)

C      END
```

### Output

The Longley data.

Longley, James W. (1967), An appraisal of least squares programs for the electronic computer from the point of view of the user, *Journal of the American Statistical Association*, 62, 819-841.

This data set consists of 16 observations on 7 variables.

X

	1	2	3	4	5	6
1	83.0	234289.0	2356.0	1590.0	107608.0	1947.0
2	88.5	259426.0	2325.0	1456.0	108632.0	1948.0
3	88.2	258054.0	3682.0	1616.0	109773.0	1949.0
4	89.5	284599.0	3351.0	1650.0	110929.0	1950.0
5	96.2	328975.0	2099.0	3099.0	112075.0	1951.0
6	98.1	346999.0	1932.0	3594.0	113270.0	1952.0
7	99.0	365385.0	1870.0	3547.0	115094.0	1953.0
8	100.0	363112.0	3578.0	3350.0	116219.0	1954.0
9	101.2	397469.0	2904.0	3048.0	117388.0	1955.0
10	104.6	419180.0	2822.0	2857.0	118734.0	1956.0
11	108.4	442769.0	2936.0	2798.0	120445.0	1957.0
12	110.8	444546.0	4681.0	2637.0	121950.0	1958.0
13	112.6	482704.0	3813.0	2552.0	123366.0	1959.0
14	114.2	502601.0	3931.0	2514.0	125368.0	1960.0
15	115.7	518173.0	4806.0	2572.0	127852.0	1961.0
16	116.9	554894.0	4007.0	2827.0	130081.0	1962.0

	7
1	60323.0
2	61122.0
3	60171.0
4	61187.0
5	63221.0
6	63639.0
7	64989.0
8	63761.0
9	66019.0
10	67857.0
11	68169.0
12	66513.0
13	68655.0
14	69564.0
15	69331.0
16	70551.0

# Chapter 20: Mathematical Support

---

## Routines

<b>20.1</b>	<b>Linear Systems</b>		
	Solve a triangular linear system given $R$ .....	GIRTS	1305
	Cholesky factorization $R^T R$ of a nonnegative definite matrix .....	CHFAC	1308
	Modified Cholesky factorization .....	MCHOL	1311
<b>20.2</b>	<b>Special Functions</b>		
	Expected value of a normal order statistic .....	ENOS	1314
	Mill's ratio .....	AMILLR	1315
<b>20.3</b>	<b>Nearest Neighbors</b>		
	Form a $k$ - $d$ tree .....	QUADT	1317
	Search a $k$ - $d$ tree for the $m$ nearest neighbors .....	NGHBR	1320

---

## GIRTS/DGIRTS (Single/Double precision)

Solve a triangular (possibly singular) set of linear systems and/or compute a generalized inverse of an upper triangular matrix.

### Usage

```
CALL GIRTS (N, R, LDR, NB, B, LDB, IPATH, IRANK, X, LDX,  
           RINV, LDRINV)
```

### Arguments

$N$  — Order of the upper triangular matrix  $R$ . (Input)

$R$  —  $N$  by  $N$  upper triangular matrix. (Input)

If  $R$  contains a zero along the diagonal, the remaining elements of the row must also be zero. Only the upper triangle of  $R$  is referenced.

**LDR** — Leading dimension of  $R$  exactly as specified in the dimension statement of the calling program. (Input)

**NB** — Number of columns in  $B$ . (Input)

$NB$  must be nonnegative. If  $NB$  is zero, no linear systems are solved.

**B** —  $N$  by  $NB$  matrix containing the right hand sides of the linear system. (Input, if  $NB > 0$ )

If  $NB = 0$ ,  $B$  is not referenced and can be a vector length one.

**LDB** — Leading dimension of  $B$  exactly as specified in the dimension statement of the calling program. (Input)

**IPATH** — Path option. (Input)

**IPATH Action**

- 1 Solve  $R * X = B$ .
- 2 Solve  $R^T * X = B$ .
- 3 Solve  $R * X = B$  and compute  $RINV$ .
- 4 Solve  $R^T * X = B$  and compute  $RINV$ .

**IRANK** — Rank of  $R$ . (Output)

**X** —  $N$  by  $NB$  matrix containing the solution matrix corresponding to the right hand side  $B$ . (Output, if  $NB > 0$ )

If  $B$  is not needed, then  $X$  and  $B$  can share the same storage locations. If  $NB = 0$ ,  $X$  is not referenced and can be a vector of length one.

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement of the calling program. (Input)

**RINV** —  $N$  by  $N$  upper triangular matrix that is the inverse of  $R$  when  $R$  is nonsingular. (Output, if  $IPATH$  equals 3 or 4)

(When  $R$  is singular,  $RINV$  is  $g_3$  inverse. See the Algorithm section for an explanation of  $g_3$  inverses.) If  $IPATH = 1$  or 2,  $RINV$  is not referenced and can be a vector of length one. If  $IPATH = 3$  or 4 and  $R$  is not needed, then  $R$  and  $RINV$  can share the same storage locations.

**LDRINV** — Leading dimension of  $RINV$  exactly as specified in the dimension statement of the calling program. (Input)

**Comments**

- 1. Informational error  
Type Code  
3 1 The linear system of equations is inconsistent.
- 2. Routine **GIRTS** assumes that a singular  $R$  is represented by zero rows in  $R$ . No other forms of singularity in  $R$  are allowed.

## Algorithm

Routine GIRTS solves systems of linear algebraic equations with a triangular coefficient matrix. Inversion of the coefficient matrix is an option. The coefficient matrix can contain a zero diagonal element, but if so, the remaining elements in the row must be zero also. (A terminal error message is issued if a nonzero element appears in the row of the coefficient matrix where a zero diagonal element appears.)

If solution of a linear system is requested (i.e.,  $NB > 0$ ) and row  $i$  of the coefficient matrix contains elements all equal to zero, the following action is taken:

- The  $i$ -th row of the solution  $X$  is set to zero.
- If  $IPATH$  is 1 or 3, a warning error is issued when the  $i$ -th row of the right-hand side  $B$  is not zero.
- If  $IPATH$  is 2 or 4, a warning error is issued when the  $i$ -th row of the reduced right-hand side (obtained after the first  $i - 1$  variables are eliminated from row  $i$ ) is not zero within a computed tolerance.

If an inverse of the coefficient matrix is requested and row  $i$  contains elements all equal to zero, row  $i$  and column  $i$  elements of  $RINV$  are set to zero. The resulting inverse is a  $g_3$  inverse of  $R$ . For a matrix  $G$  to be  $g_3$  inverse of a matrix  $A$ ,  $G$  must satisfy Conditions 1, 2, and 3 for the Moore-Penrose inverse but generally fail Condition 4. The four conditions for  $G$  to be a Moore-Penrose inverse of  $A$  are as follows:

1.  $AGA = A$
2.  $GAG = G$
3.  $AG$  is symmetric
4.  $GA$  is symmetric

For a detailed description of the algorithm, see Section 2 in Sallas and Lioni (1988).

## Example

The following example is taken from Maindonald (1984, pp. 102-105). A linear system  $Rx = B$  is solved, and a  $g_3$  inverse of  $R$  is computed.

```
C      INTEGER      LDB, LDR, LDRINV, LDX, N, NB
PARAMETER      (N=4, NB=1, LDB=N, LDR=N, LDRINV=N, LDX=N)
C
      INTEGER      IPATH, IRANK
      REAL         B(LDB,NB), R(LDR,N), RINV(LDRINV,N), X(LDX,NB)
      EXTERNAL    GIRTS, WRRRN
C
      DATA (R(1,J),J=1,N)/6.0, 2.0, 5.0, 1.0/, B(1,1)/3.0/
```

```

DATA (R(2,J),J=1,N)/0.0, 4.0,-2.0, 2.0/, B(2,1)/4.0/
DATA (R(3,J),J=1,N)/0.0, 0.0, 0.0, 0.0/, B(3,1)/0.0/
DATA (R(4,J),J=1,N)/0.0, 0.0, 0.0, 3.0/, B(4,1)/3.0/
C
  IPATH = 3
  CALL GIRTS (N, R, LDR, NB, B, LDB, IPATH, IRANK, X, LDX, RINV,
&            LDRINV)
C
  CALL WRRRN ('RINV', N, N, RINV, LDRINV, 0)
  CALL WRRRN ('X', N, NB, X, LDX, 0)

```

### Output

	RINV			
	1	2	3	4
1	0.1667	-0.0833	0.0000	0.0000
2	0.0000	0.2500	0.0000	-0.1667
3	0.0000	0.0000	0.0000	0.0000
4	0.0000	0.0000	0.0000	0.3333

	X
1	0.167
2	0.500
3	0.000
4	1.000

---

## CHFAC/DCHFAC (Single/Double precision)

Compute an upper triangular factorization of a real symmetric nonnegative definite matrix.

### Usage

```
CALL CHFAC (N, A, LDA, TOL, IRANK, R, LDR)
```

### Arguments

*N* — Order of the matrix. (Input)

*A* — *N* by *N* symmetric nonnegative definite matrix for which an upper triangular factorization is desired. (Input)

Only elements in the upper triangle of *A* are referenced.

*LDA* — Leading dimension of *A* exactly as specified in the dimension statement in the calling program. (Input)

*TOL* — Tolerance used in determining linear dependence. (Input)

For CHFAC,  $TOL = 100 * AMACH(4)$  is a common choice. For DCHFAC,  $TOL = 100 * DMACH(4)$  is a common choice. See documentation for routine AMACH.

*IRANK* — Rank of *A*. (Output)

$N - IRANK$  is the number of effective zero pivots.

**R** — N by N upper triangular matrix containing the *R* matrix from a Cholesky decomposition  $R^T R$  of *A*. (Output)

The elements of the appropriate rows of *R* are set to 0.0 if linear dependence of the columns of *A* is declared. (There are  $N - \text{IRANK}$  rows of *R* whose elements are set to 0.0.) If *A* is not needed, then *R* and *A* can share the same storage locations.

**LDR** — Leading dimension of *R* exactly as specified in the dimension statement in the calling program. (Input)

### Comments

1. Informational error  

Type	Code	
3	1	The input matrix is not nonnegative definite within the tolerance defined by TOL.
2. Elements of row *i* of *R* are set to 0.0 if a linear dependence is declared. Linear dependence is declared if

$$\left| a_{ii} - \sum_{j=1}^{i-1} r_{ji}^2 \right| \leq \text{TOL} * |a_{ii}|$$

### Algorithm

Routine CHFAC computes a Cholesky factorization  $R^T R = A$  of an  $n \times n$  symmetric nonnegative definite matrix *A*. The matrix *R* is taken to be an upper triangular matrix. The diagonal elements of *R* are taken to be nonnegative. If *A* is singular and has rank *r*,  $n - r$  rows of *R* have all their elements taken to be zero.

The algorithm is based on the work of Healy (1968). The algorithm proceeds sequentially by columns. The *i*-th column is declared to be linearly dependent on the first  $i - 1$  columns if

$$\left| a_{ii} - \sum_{j=1}^{i-1} r_{ji}^2 \right| \leq \epsilon |a_{ii}|$$

where  $\epsilon$  (stored in TOL) is the input tolerance. When a linear dependence is declared, all elements in the *i*-th row of *R* are set to zero.

Modifications due to Farebrother and Berry (1974) and Barrett and Healy (1978) for checking for matrices that are not nonnegative definite are also incorporated. Routine CHFAC declares *A* to not be nonnegative definite and issues an error message with an error code of 1 if either of the following conditions is satisfied:



1.  $a_{ii} - \sum_{j=1}^{i-1} r_{ji}^2 < -\epsilon|a_{ii}|$
2.  $r_{ii} = 0$  and  $|a_{ik} - \sum_{j=1}^{i-1} r_{ji}r_{jk}| > \epsilon\sqrt{a_{ii}a_{kk}}, k > i$

Healy's (1968) algorithm and CHFAC permit the matrices  $A$  and  $R$  to occupy the same storage locations. Barrett and Healy (1978) in their remark neglect this fact. Routine CHFAC uses

$$\sum_{j=1}^{i-1} r_{ji}^2$$

for  $a_{ii}$  in the Condition 2 above to remedy this problem.

### Example

A Cholesky factorization of a  $5 \times 5$  symmetric nonnegative definite matrix is computed. Maindonald (1984, pages 85–86) discusses in detail the computations for this problem.

```

C                               SPECIFICATIONS FOR PARAMETERS
INTEGER      LDA, LDR, N
PARAMETER    (N=5, LDA=N, LDR=N)

C
INTEGER      IRANK, NOUT
REAL         A(LDA,N), AMACH, R(LDR,N), TOL
EXTERNAL     AMACH, CHFAC, UMACH, WRRRN

C
DATA (A(1,J),J=1,N)/36.0, 12.0, 30.0, 6.0, 18.0/
DATA (A(2,J),J=1,N)/12.0, 20.0, 2.0, 10.0, 22.0/
DATA (A(3,J),J=1,N)/30.0, 2.0, 29.0, 1.0, 7.0/
DATA (A(4,J),J=1,N)/ 6.0, 10.0, 1.0, 14.0, 20.0/
DATA (A(5,J),J=1,N)/ 8.0, 22.0, 7.0, 20.0, 40.0/

C
TOL = 100.0*AMACH(4)

C
CALL CHFAC (N, A, LDA, TOL, IRANK, R, LDR)
CALL UMACH (2, NOUT)
WRITE (NOUT,*) 'IRANK = ', IRANK
CALL WRRRN ('R', N, N, R, LDR, 0)
END

```

### Output

```

IRANK = 4

      R
      1      2      3      4      5
1  6.000  2.000  5.000  1.000  3.000
2  0.000  4.000 -2.000  2.000  4.000
3  0.000  0.000  0.000  0.000  0.000
4  0.000  0.000  0.000  3.000  3.000
5  0.000  0.000  0.000  0.000  2.449

```

---

## MCHOL/DMCHOL (Single/Double precision)

Compute an upper triangular factorization of a real symmetric matrix  $A$  plus a diagonal matrix  $D$ , where  $D$  is determined sequentially during the Cholesky factorization in order to make  $A + D$  nonnegative definite.

### Usage

CALL MCHOL (N, A, LDA, TOL, IRANK, R, LDR, DMAX, IND)

### Arguments

**N** — Order of the matrix. (Input)

**A** —  $N$  by  $N$  symmetric matrix for which a Cholesky factorization is attempted. (Input)

Only elements in the upper triangle and diagonal of  $A$  are referenced.

**LDA** — Leading dimension of  $A$  exactly as specified in the dimension statement in the calling program. (Input)

**TOL** — Tolerance used in determining linear dependence. (Input)

For MCHOL,  $TOL = 100 * AMACH(4)$  is a common choice. For DMCHOL,  $TOL = 100 * DMACH(4)$  is a common choice. See documentation for routine AMACH.

**IRANK** — Rank of  $A + D$ . (Output)

**R** —  $N$  by  $N$  upper triangular matrix containing the  $R$  matrix from a Cholesky decomposition  $R^T R$  of  $A + D$ . (Output)

The lower triangle of  $R$  is not referenced. If  $A$  is not needed, then  $R$  and  $A$  can share the same storage locations.

**LDR** — Leading dimension of  $R$  exactly as specified in the dimension statement in the calling program. (Input)

**DMAX** — Largest diagonal element of  $D$ . (Output)

If  $DMAX$  equals 0.0, then  $A$  is nonnegative definite, and  $R$  is a Cholesky factorization of  $A$ . If  $DMAX$  is positive, then  $A$  is indefinite, i.e.,  $A$  has at least one negative eigenvalue. In this case,  $DMAX$  is an upper bound on the absolute value of the minimum eigenvalue.

**IND** — Index for subsequent computation of a descent direction in the case of a saddle point. (Output)

If  $IND = 0$ , then  $A$  is nonnegative definite. For positive  $IND$ , let  $e$  be a unit vector with  $IND$ -th element 1 and remaining elements 0. The solution  $s$  of  $Rs = e$  is a direction of negative curvature, i.e.,  $s^T A s$  is negative.

## Algorithm

Routine MCHOL computes a Cholesky factorization,  $R^T R$ , of  $A + D$  where  $A$  is symmetric and  $D$  is a diagonal matrix with sufficiently large diagonal elements such that  $A + D$  is nonnegative definite. The routine is similar to one described by Gill, Murray, and Wright (1981, pages 108–111). Here, though, we allow  $A + D$  to be singular.

The algorithm proceeds sequentially by rows. If  $A + D$  is singular, the Cholesky factor  $R$  is taken to have some rows that are entirely zero. The  $i$ -th row of  $A + D$  is declared to be linearly dependent on the first  $i - 1$  rows if the following two conditions are satisfied:

1.  $\left| a_{ii} - \sum_{j=1}^{i-1} r_{ji}^2 \right| \leq \epsilon |a_{ii}|$
2.  $\left| a_{ik} - \sum_{j=1}^{i-1} r_{ji} r_{jk} \right| \leq \epsilon \sqrt{a_{ii} a_{kk}}, k > i$

where  $\epsilon$  is the input argument TOL.

The routine MCHOL is often used to find a descent direction in a minimization problem. Let  $A$  and  $g$  be the current Hessian and gradient, respectively, associated with the minimization problem. The solution  $s$  of  $As = -g$  may not give a descent direction if  $A$  is not nonnegative definite. Instead, in order to guarantee a descent direction, a solution  $s$  of  $(A + D)s = -g$  can be found where  $A + D$  is nonnegative definite. Routine MCHOL is useful for computing the upper triangular Cholesky factor  $R$  of  $A + D$  so that routine GIRTS can be invoked to compute the descent direction  $s$  by solving successively the two triangular linear systems  $R^T x = -g$  and  $Rs = x$  for  $x$  and then  $s$ . Also if  $g = 0$  and  $A$  is not nonnegative definite, i.e., the current solution is a saddle point, GIRTS can be used to compute a descent direction  $s$  from the linear system  $Rs = e$  where  $e$  is a unit vector with

$$\epsilon_i = \begin{cases} 1 & \text{if } i = \text{IND} \\ 0 & \text{otherwise} \end{cases}$$

## Example 1

A Cholesky factorization of a  $5 \times 5$  symmetric nonnegative definite matrix is computed. Maindonald (1984, pages 85–86) discusses the example.

```
C                                     SPECIFICATIONS FOR PARAMETERS
      INTEGER      LDA, LDR, N
      PARAMETER    (N=5, LDA=N, LDR=N)

C
      INTEGER      IND, IRANK, NOUT
      REAL         A(LDA,N), DMAX, R(LDR,N), TOL
      EXTERNAL     MCHOL, UMACH, WRRRN
```

```

C
DATA (A(1,J),J=1,N)/36.0, 12.0, 30.0, 6.0, 18.0/
DATA (A(2,J),J=1,N)/12.0, 20.0, 2.0, 10.0, 22.0/
DATA (A(3,J),J=1,N)/30.0, 2.0, 29.0, 1.0, 7.0/
DATA (A(4,J),J=1,N)/6.0, 10.0, 1.0, 14.0, 20.0/
DATA (A(5,J),J=1,N)/8.0, 22.0, 7.0, 20.0, 40.0/

C
TOL = 0.00001
CALL MCHOL (N, A, LDA, TOL, IRANK, R, LDR, DMAX, IND)
CALL UMACH (2, NOUT)
WRITE (NOUT,99998) ' IRANK = ', IRANK
WRITE (NOUT,99999) ' DMAX = ', DMAX
WRITE (NOUT,99998) ' IND = ', IND
99998 FORMAT (A, I3)
99999 FORMAT (A, 1PE10.3)
CALL WRRRN ('R', N, N, R, LDR, 0)
END

```

### Output

```

IRANK = 4
DMAX = 0.000E+00
IND = 0

```

```

          R
         1      2      3      4      5
1  6.000  2.000  5.000  1.000  3.000
2  0.000  4.000 -2.000  2.000  4.000
3  0.000  0.000  0.000  0.000  0.000
4  0.000  0.000  0.000  3.000  3.000
5  0.000  0.000  0.000  0.000  2.449

```

### Example 2

A modified Cholesky factorization of a  $3 \times 3$  symmetric indefinite matrix  $A$  is computed. A solution of  $Rs = e_3$  is also obtained using routine GIRTS. Note that  $s^T A s$  is negative as verified by using routine BLINF (IMSL MATH/LIBRARY). Gill, Murray, and Wright (1981, page 111) discuss the example.

```

C
SPECIFICATIONS FOR PARAMETERS
INTEGER LDA, LDR, N
PARAMETER (N=3, LDA=N, LDR=N)

C
INTEGER IND, IRANK, NOUT
REAL A(LDA,N), BLINF, DMAX, E(N), R(LDR,N), S(N), SPAS, TOL
EXTERNAL BLINF, GIRTS, MCHOL, SSET, UMACH, WRRRN

C
DATA (A(1,J),J=1,N)/1, 1, 2/
DATA (A(2,J),J=1,N)/1, 1, 3/
DATA (A(3,J),J=1,N)/2, 3, 1/

C
TOL = 0.00001
CALL MCHOL (N, A, LDA, TOL, IRANK, R, LDR, DMAX, IND)
CALL UMACH (2, NOUT)

```

```

WRITE (NOUT,99998) ' IRANK = ', IRANK
WRITE (NOUT,99999) ' DMAX = ', DMAX
WRITE (NOUT,99998) ' IND = ', IND
CALL WRRRN ('R', N, N, R, LDR, 0)
IF (IND .GT. 0) THEN
  CALL SSET (N, 0.0, E, 1)
  E(IND) = 1.0
  CALL GIRTS (N, R, LDR, 1, E, N, 1, IRANK, S, N, R, LDR)
  SPAS = BLINF(N,N,A,LDA,S,S)
  WRITE (NOUT,*) ' '
  WRITE (NOUT,99999) ' trans(s)*A*s = ', SPAS
END IF
99998 FORMAT (A, I3)
99999 FORMAT (A, F10.3)
END

```

### Output

```

IRANK =    3
DMAX   =    5.016
IND    =    3

      R
      1      2      3
1  1.942  0.515  1.030
2  0.000  2.398  1.030
3  0.000  0.000  1.059

trans(s)*A*s =    -2.254

```

---

## ENOS/DENOS (Single/Double precision)

Evaluate the expected value of a normal order statistic.

### Usage

Usage ENOS(I, N)

### Arguments

*I*— Rank of the order statistic. (Input)

*N*— Sample size. (Input)

*ENOS*— Function value, the expected value of the *I*-th order statistic in a sample of size *N* from the standard normal distribution. (Output)

### Comments

Informational errors

Type	Code	
3	1	The rank of the order statistic is less than 1. A rank of 1 is assumed.

3      2      The rank of the order statistic is greater than sample size ( $n$ ). A rank of  $N$  is assumed.

### Algorithm

Let  $X_1 \leq X_2 \leq \dots \leq X_n$  be the order statistics of a random sample of size  $n$  from a standard normal distribution. The expected value of  $X_i$  is given by

$$\frac{n!}{(n-i)!(i-1)!} \int_{-\infty}^{\infty} x [\Phi(x)]^{i-1} [1 - \Phi(x)]^{n-i} \phi(x) dx$$

where  $\phi(x)$  and  $\Phi(x)$  are the standard normal density and cumulative distribution functions respectively (David 1981).

Function `ENOS` evaluates the integral using a trapezoidal rule after first making a logarithmic transformation. This is the method used by Harter (1961). Although the method permits computations for any value of  $n$ , extremely large values of  $n$  cannot be guaranteed to be as accurate as smaller values of  $n$ . For  $n > 2500$ , the method is inappropriate.

### Example

In this example, we compute the expected value of the first order statistic in a sample of size 5 from a standard normal distribution.

```

INTEGER      I, N, NOUT
REAL         ENOS, EX
EXTERNAL     ENOS, UMACH
C
CALL UMACH (2, NOUT)
I = 1
N = 5
EX = ENOS(I,N)
WRITE (NOUT,99999) EX
99999 FORMAT (' The expected value of the smallest order statistic',
&           /, ' in a normal sample of size 5 is ', F9.5)
END

```

### Output

```

The expected value of the smallest order statistic
in a normal sample of size 5 is  -1.16296

```

---

## AMILLR/DMILLR (Single/Double precision)

Evaluate Mill's ratio (the ratio of the ordinate to the upper tail area of the standardized normal distribution).

### Usage

`AMILLR(X)`

## Arguments

$X$  — Value at which Mill's ratio is evaluated. (Input)

In order to avoid overflow,  $x$  must be less than a bound that is machine dependent. On most machines, the bound is greater than  $-13$ . The function underflows (and is set to 0.0) for small values of  $x$ . On most machines, the underflow does not occur unless  $x$  is less than  $-13$ .

**AMILLR** — Function value, Mill's ratio. (Output)

## Comments

Informational error

Type Code

2 1 The function underflows because  $x$  is too small.

## Algorithm

Function **AMILLR** evaluates Mill's ratio, the *hazard rate* for the standard normal distribution. It is computed as the ratio of the ordinate to the upper tail area of the standard normal distribution, that is,  $\phi(x)/(1 - \Phi(x))$ , where  $\phi(x)$  and  $\Phi(x)$  are the standard normal density and cumulative distribution functions, respectively. The reciprocal of Mill's ratio is called the *failure rate* in reliability and life testing applications. As  $x$  becomes small, the ratio goes to zero. For large  $x$  (how large is machine dependent), the ratio cannot be computed. Function **AMILLR** computes  $1 - \Phi(x)$  using the complementary error function (IMSL 1991) rather than as one minus the normal distribution function, which would underflow sooner as  $x$  gets small.

## Example

In this example, we compute Mill's ratio at  $x = -1.0$ .

```
INTEGER      NOUT
REAL         AMILLR, R, X
EXTERNAL     AMILLR, UMACH
C
CALL UMACH (2, NOUT)
X = -1.0
R = AMILLR(X)
WRITE (NOUT,99999) R
99999 FORMAT (' Mill's ratio at -1.0 is ', F8.5)
END
```

## Output

```
Mill's ratio at -1.0 is  0.28760
```

---

## QUADT/DQUADT (Single/Double precision)

Form a  $k$ - $d$  tree.

### Usage

CALL QUADT (NROW, NVAR, NCOL, X, LDX, IND, NBUCK, IDISCR,  
PART)

### Arguments

**NROW** — Number of rows of  $X$  to be used in forming the  $k$ - $d$  tree. (Input)

**NVAR** — Number of variables to be used in forming the tree. (Input)

**NCOL** — Number of columns in  $X$ . (Input)

**X** — NROW by NCOL matrix containing the data to be used on this call.  
(Input/Output)

On output the rows of  $X$  have been rearranged to form the  $k$ - $d$  tree.  $X$  must not contain missing values (NaN).

**LDX** — Leading dimension of  $X$  exactly as specified in the dimension statement in the calling program. (Input)

**IND** — Vector of length NVAR containing the column numbers in  $X$  to be used in the forming the  $k$ - $d$  tree. (Input)

**NBUCK** — Bucket size. (Input)

NBUCK gives the maximum number of observations in a leaf of the  $k$ - $d$  tree. NBUCK = 3 is a common choice. NBUCK should be small when compared to NROW.

**IDISCR** — Vector of length NROW containing the element number in IND that points to the column of  $X$  to be used as the discriminator in the  $k$ - $d$  tree. (Output)  
IDISCR(I) = 0 if the observation is a terminal node. IND(IDISCR(I)) is the column number in  $X$  to be used as the discriminator.

**PART** — Vector of length NROW containing the value to be used in the partition for this observation. (Output)

### Comments

Automatic workspace usage is

QUADT  $2 * NROW + 2 * (\log_2(NROW) + 3)$  units, or

DQUADT  $3 * NROW + 2 * (\log_2(NROW) + 3)$  units.

Workspace may be explicitly provided, if desired, by use of Q2ADT/DQ2ADT. The reference is



```
CALL Q2ADT (NROW, NVAR, NCOL, X, LDX, IND, NBUCK, IDISCR,
           PART, ILOW, IHIGH, WK, IWK)
```

The additional arguments are as follows:

**ILOW** — Work vector of length  $\log_2(\text{NROW}) + 3$ .

**IHIGH** — Work vector of length  $\log_2(\text{NROW}) + 3$ .

**WK** — Work vector of length **NROW**.

**IWK** — Work vector of length **NROW**.

### Algorithm

Routine **QUADT** creates the data structure required for a *k-d* tree. A *k-d* tree is a multivariate form of B-tree that is especially useful for finding nearest neighbors but may be of use in other situations. Once the *k-d* tree has been formed, routine **NGHBR** (page 1320) may be used to find the nearest neighbors for any point in logarithmic time.

The basic algorithm is given by Friedman, Bentley, and Finkel (1977) and can be summarized as follows:

1. Let  $l = 1$  and  $h = \text{NROW}$ .
2. Let  $k = (l + h)/2$ .
3. Each column in **X** to be used in forming the *k-d* tree is examined over the range  $[l, h]$  in order to find the column with the maximum spread. Let  $j$  equal this column number.
4. The  $k$ -th element of **PART** is set to the median value in the range  $[l, h]$  of the  $j$ -th column in **X** while **IDISCR**( $k$ ) is set to the element in **IND** that points to this column.
5. The rows of **X** are interchanged so that all rows of **X** with values in column  $j$  less than or equal to the median value computed in Step 4 occur before (or at) the  $k$ -th element.
6. Go to Step 2 repeatedly with zero, one, or two submatrices in **X**. Go to Step 2 with the submatrix formed from rows  $l$  to  $k$  of **X** if  $k - l$  is greater than **NBUCK**. Go to Step 2 with the submatrix formed from rows  $k + 1$  to  $h$  of **X** if  $h - k - 1$  is greater than **NBUCK**.

The bucket size, **NBUCK**, is the maximum number of observations allowed in the lowest level of the *k-d* tree, i.e., in the leaves of the tree. The choice of **NBUCK** can affect the speed with which nearest neighbors are found. A value of 3 or 5 is a common choice, but if the number of nearest neighbors to be obtained is large, a larger value for **NBUCK** should probably be used.

## Example

The following example creates a  $k$ - $d$  tree from financial data collected for firms approximately 2 years prior to bankruptcy and for financially sound firms at about the same point in time. The data on five variables,  $X_1$  = (population),  $X_2$  = (cash flow)/(total debt),  $X_3$  = (net income)/(total assets),  $X_4$  = (current assets)/(current liabilities), and  $X_5$  = (current assets)/(net sales) are taken from Johnson and Wichern (1988, page 536).

```
C      INTEGER      LDX, NBUCK, NCOL, NROW, NVAR
      PARAMETER    (LDX=47, NBUCK=3, NCOL=5, NROW=47, NVAR=4)

C      INTEGER      IDISCR(NROW), IND(NVAR)
      REAL          PART(NROW), X(LDX,NCOL)
      EXTERNAL      QUADT, WRIRN, WRRRN

C      DATA IND/2, 3, 4, 5/
      DATA (X(I,1),I=1,47)/1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 1.,
& 1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 2., 2., 2., 2., 2.,
& 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2.,
& 2., 2., 2., 2., 2., 2./
      DATA (X(I,2),I=1,47)/-0.4485, -0.5633, 0.0643, -0.0721, -0.1002,
& -0.1421, 0.0351, -0.0653, 0.0724, -0.1353, -0.2298, 0.0713,
& 0.0109, -0.2777, 0.1454, 0.3703, -0.0757, 0.0451, 0.0115,
& 0.1227, -0.2843, 0.5135, 0.0769, 0.3776, 0.1933, 0.3248,
& 0.3132, 0.1184, -0.0173, 0.2169, 0.1703, 0.1460, -0.0985,
& 0.1398, 0.1379, 0.1486, 0.1633, 0.2907, 0.5383, -0.3330,
& 0.4785, 0.5603, 0.2029, 0.2029, 0.4746, 0.1661, 0.5808/
      DATA (X(I,3),I=1,47)/-0.4106, -0.3114, -0.3114, -0.0930,
& -0.0917, -0.0651, 0.0147, -0.0566, -0.0076, -0.1433,
& -0.2961, 0.0205, 0.0011, -0.2316, 0.0500, 0.1098, -0.0821,
& 0.0263, -0.0032, 0.1055, -0.2703, 0.1001, 0.0195, 0.1075,
& 0.0473, 0.0718, 0.0511, 0.0499, 0.0233, 0.0779, 0.0695,
& 0.0518, -0.0123, -0.0312, 0.0728, 0.0564, 0.0486, 0.0597,
& 0.1064, -0.0854, 0.0910, 0.1112, 0.0792, 0.0792, 0.1380,
& 0.0351, 0.0371/
      DATA (X(I,4),I=1,47)/1.0865, 1.5134, 1.0077, 1.4544, 1.5644,
& 0.7066, 1.5046, 1.3737, 1.3723, 1.4196, 0.3310, 1.3124,
& 2.1495, 1.1918, 1.8762, 1.9941, 1.5077, 1.6756, 1.2602,
& 1.1434, 1.2722, 2.4871, 2.0069, 3.2651, 2.2506, 4.2401,
& 4.4500, 2.5210, 2.0538, 2.3489, 1.7973, 2.1692, 2.5029,
& 0.4611, 2.6123, 2.2347, 2.3080, 1.8381, 2.3293, 3.0124,
& 1.2444, 4.2918, 1.9936, 1.9936, 2.9166, 2.4527, 5.0594/
      DATA (X(I,5),I=1,47)/0.4526, 0.1642, 0.3978, 0.2589, 0.6683,
& 0.2794, 0.7080, 0.4032, 0.3361, 0.4347, 0.1824, 0.2497,
& 0.6969, 0.6601, 0.2723, 0.3828, 0.4215, 0.9494, 0.6038,
& 0.1655, 0.5128, 0.5368, 0.5304, 0.3548, 0.3309, 0.6279,
& 0.6852, 0.6925, 0.3483, 0.3970, 0.5174, 0.5500, 0.5778,
& 0.2643, 0.5151, 0.5563, 0.1978, 0.3786, 0.4835, 0.4730,
& 0.1847, 0.4443, 0.3018, 0.3018, 0.4487, 0.1370, 0.1268/

C      CALL QUADT (NROW, NVAR, NCOL, X, LDX, IND, NBUCK, IDISCR, PART)
      CALL WRRRN ('first 10 rows of X after QUADT', 10, NCOL, X, LDX,
& 0)
      CALL WRRRN ('PART', 1, NROW, PART, 1, 0)
```

```

C      CALL WRIRN ('IDISCR', 1, NROW, IDISCR, 1, 0)
      END

```

### Output

```

first 10 rows of X after QUADT
      1      2      3      4      5
1  1.000 -0.230 -0.296  0.331  0.182
2  2.000  0.140 -0.031  0.461  0.264
3  1.000 -0.142 -0.065  0.707  0.279
4  1.000 -0.449 -0.411  1.087  0.453
5  1.000  0.064 -0.311  1.008  0.398
6  1.000  0.123  0.105  1.143  0.166
7  1.000 -0.284 -0.270  1.272  0.513
8  1.000 -0.278 -0.232  1.192  0.660
9  1.000  0.012 -0.003  1.260  0.604
10 1.000  0.071  0.021  1.312  0.250

```

```

PART
      1      2      3      4      5      6      7      8      9     10
0.000  0.461  0.857  0.000  0.064  1.168  0.000 -0.278  0.041  0.000

     11     12     13     14     15     16     17     18     19     20
0.072  1.373  0.000 -0.072  0.412  0.000  0.435 -0.015  0.000  1.876

     21     22     23     24     25     26     27     28     29     30
0.448  0.000  .708  1.994  0.000  0.203  2.152  0.000  2.308  0.390

     31     32     33     34     35     36     37     38     39     40
0.000  .550  0.147  0.000  0.217  2.453  0.000  2.521  0.128  0.000

     41     42     43     44     45     46     47
2.612  3.012  0.000  4.240  4.292  4.755  0.000

```

```

IDISCR
  1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20
  0  3  3  0  1  3  0  1  1  0  1  3  0  1  4  0  4  1  0  3

 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40
  4  0  4  3  0  1  3  0  3  4  0  4  1  0  1  3  0  3  1  0

 41 42 43 44 45 46 47
  3  3  0  3  3  3  0

```

---

## NGHBR/DNGHBR (Single/Double precision)

Search a  $k$ - $d$  tree for the  $k$  nearest neighbors of a key.

### Usage

```

CALL NGHBR (NVAR, XKEY, K, NROW, NCOL, X, LDX, IND,
            NBUCK, IDISCR, PART, METRIC, IPQR, PQD)

```

## Arguments

**NVAR** — Number of variables used to form the  $k$ - $d$  tree. (Input)

**XKEY** — Vector of length NVAR containing the key for which nearest neighbors are desired. (Input)

Note that the elements in XKEY are not arranged in the same manner as the columns in X.

**K** — Number of nearest neighbors to find. (Input)

**NROW** — Number of rows of X used to form the  $k$ - $d$  tree. (Input)

**NCOL** — Number of columns in X. (Input)

**X** — NROW by NCOL matrix containing the data used to form the  $k$ - $d$  tree as output from routine QUADT (page 1317). (Input)  
X must not contain missing values (NaN).

**LDX** — Leading dimension of X exactly as specified in the dimension statement in the calling program. (Input)

**IND** — Vector of length NVAR containing the column numbers in X used in forming the  $k$ - $d$  tree. (Input)

**NBUCK** — Bucket size. (Input)

NBUCK is the maximum number of observations in a leaf of the  $k$ - $d$  tree. The value of NBUCK should be the same as the value used in forming the  $k$ - $d$  tree (i.e. as input to the routine QUADT).

**IDISCR** — Vector of length NROW containing the element number in IND that points to the column of X to be used as the discriminator in the  $k$ - $d$  tree, as output from routine QUADT. (Input)

IDISCR(I) = 0 if the observation is a terminal node. IND(IDISCR(I)) is the column number in X to be used as the discriminator.

**PART** — Vector of length NROW containing the median value to be used for the partition, as output from routine QUADT. (Input)

**METRIC** — Metric to use in computing the  $k$  nearest neighbors. (Input)

### **METRIC** Metric used

0	Euclidean distance
1	$L_1$ norm
2	$L_\infty$ norm

**IPQR** — Vector of length K containing the indices of the nearest neighbors. (Output)

**PQD** — Vector of length K containing the nearest neighbor distances. (Output)

## Comments

1. Automatic workspace usage is

NGHBR  $(2 * NVAR + 3) * (\log_2(NROW) + 3)$  units, or

DNGHBR  $(4 * NVAR + 3) * (\log_2(NROW) + 3)$  units.

Workspace may be explicitly provided, if desired, by use of N2HBR/DN2HBR. The reference is

```
CALL N2HBR (NVAR, XKEY, K, NROW, NCOL, X, LDX, IND,  
           IDISCR, PART, METRIC, IPQR, PQD, ILOW,  
           IHIGH, ISIDE, BNDL, BNDH)
```

The additional arguments are as follows:

**ILOW** — Work vector of length  $\log_2(NROW) + 3$ .

**IHIGH** — Work vector of length  $\log_2(NROW) + 3$ .

**ISIDE** — Work vector of length  $\log_2(NROW) + 3$ .

**BNDL** — Work vector of length  $NVAR * (\log_2(NROW) + 3)$ .

**BNDH** — Work vector of length  $NVAR * (\log_2(NROW) + 3)$ .

2. Informational error

Type	Code	
4	1	The data structure input is not a $k$ - $d$ tree. Use routine QUADT to create the $k$ - $d$ tree.

## Algorithm

Routine NGHBR finds the  $k$  nearest neighbors in an input  $k$ - $d$  tree for an arbitrary key, XKEY in logarithmic time. A  $k$ - $d$  tree is a form of B-tree that is especially useful for finding nearest neighbors. The  $k$ - $d$  tree input into routine NGHBR should be produced by routine QUADT (page 1317). Three metrics, Euclidean,  $L_1$ , and  $L_\infty$ , are available for defining the nearest neighbors. The user should note that if the input key is a row of the  $k$ - $d$  tree, then the row will be returned as one of the nearest neighbors. In this case, only  $k - 1$  nearest neighbors will be found.

The algorithm is given by Friedman, Bentley, and Finkel (1977) and is summarized in the following. The basic idea is to traverse the  $k$ - $d$  tree in order to determine which leaves of the tree need to be examined for the nearest neighbor. The algorithm is efficient because most leaves are not examined.

1. Let  $l = 1$  and  $h = NROW$ .
2. Let  $k = (l + h)/2$ , and  $j$  and  $p$  be the  $k$ -th elements of IDISCR and PART, respectively.

3. If  $(h - l)$  is less than NBUCK, then go to Step 4. Otherwise, let  $m$  be the  $j$ -th element of IND. If the  $(k, m)$ -th element of  $x$  is greater than  $p$ , then let  $l = k + 1$  and go to Step 2. Otherwise, set  $h = k$  and go to Step 2.
4. Examine each row in  $x$  from row  $l$  to row  $h$  to determine if it is a nearest neighbor. Check to see if rows in  $x$  (leaves of the tree) adjacent to these rows need to be examined (see Friedman, Bentley, and Finkel (1977)). If necessary, examine the adjacent rows for nearest neighbors.

The value used for the bucket size, NBUCK, must be the same value as was used in routine QUADT when the  $k$ - $d$  tree was created. A common choice for NBUCK is three.

### Example

The following example creates a  $k$ - $d$  tree from financial data collected for firms approximately 2 years prior to bankruptcy and for financially sound firms at about the same point in time. The data on five variables,  $X_1$  = (population),  $X_2$  = (cash flow)/(total dept),  $X_3$  = (net income)/(total assets),  $X_4$  = (current assets)/(current liabilities), and  $X_5$  = (current assets)/(net sales) are taken from Johnson and Wichern, page 536. Routine NGHBR is then used to determine the 5 nearest neighbors of the first row in  $x$ . As expected, one of the nearest neighbors found is the key (the first row in  $x$ ).

```

INTEGER      K, LDX, METRIC, NBUCK, NCOL, NROW, NVAR
PARAMETER    (K=5, LDX=47, METRIC=1, NBUCK=3, NCOL=5, NROW=47,
&            NVAR=4)
C
INTEGER      I, IDISCR(NROW), IND(NVAR), IPQR(K)
REAL         PART(NROW), PQD(K), X(LDX,NCOL), XKEY(NVAR)
EXTERNAL     NGHBR, QUADT, WRIRN, WRRRN
C
DATA IND/2, 3, 4, 5/
DATA (X(I,1),I=1,47)/1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 1.,
& 1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 1., 2., 2., 2., 2., 2.,
& 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2., 2.,
& 2., 2., 2., 2., 2., 2./
DATA (X(I,2),I=1,47)/-0.4485, -0.5633, 0.0643, -0.0721, -0.1002,
& -0.1421, 0.0351, -0.0653, 0.0724, -0.1353, -0.2298, 0.0713,
& 0.0109, -0.2777, 0.1454, 0.3703, -0.0757, 0.0451, 0.0115,
& 0.1227, -0.2843, 0.5135, 0.0769, 0.3776, 0.1933, 0.3248,
& 0.3132, 0.1184, -0.0173, 0.2169, 0.1703, 0.1460, -0.0985,
& 0.1398, 0.1379, 0.1486, 0.1633, 0.2907, 0.5383, -0.3330,
& 0.4785, 0.5603, 0.2029, 0.2029, 0.4746, 0.1661, 0.5808/
DATA (X(I,3),I=1,47)/-0.4106, -0.3114, -0.3114, -0.0930,
& -0.0917, -0.0651, 0.0147, -0.0566, -0.0076, -0.1433,
& -0.2961, 0.0205, 0.0011, -0.2316, 0.0500, 0.1098, -0.0821,
& 0.0263, -0.0032, 0.1055, -0.2703, 0.1001, 0.0195, 0.1075,
& 0.0473, 0.0718, 0.0511, 0.0499, 0.0233, 0.0779, 0.0695,
& 0.0518, -0.0123, -0.0312, 0.0728, 0.0564, 0.0486, 0.0597,
& 0.1064, -0.0854, 0.0910, 0.1112, 0.0792, 0.0792, 0.1380,
& 0.0351, 0.0371/

```

```

DATA (X(I,4),I=1,47)/1.0865, 1.5134, 1.0077, 1.4544, 1.5644,
& 0.7066, 1.5046, 1.3737, 1.3723, 1.4196, 0.3310, 1.3124,
& 2.1495, 1.1918, 1.8762, 1.9941, 1.5077, 1.6756, 1.2602,
& 1.1434, 1.2722, 2.4871, 2.0069, 3.2651, 2.2506, 4.2401,
& 4.4500, 2.5210, 2.0538, 2.3489, 1.7973, 2.1692, 2.5029,
& 0.4611, 2.6123, 2.2347, 2.3080, 1.8381, 2.3293, 3.0124,
& 1.2444, 4.2918, 1.9936, 1.9936, 2.9166, 2.4527, 5.0594/
DATA (X(I,5),I=1,47)/0.4526, 0.1642, 0.3978, 0.2589, 0.6683,
& 0.2794, 0.7080, 0.4032, 0.3361, 0.4347, 0.1824, 0.2497,
& 0.6969, 0.6601, 0.2723, 0.3828, 0.4215, 0.9494, 0.6038,
& 0.1655, 0.5128, 0.5368, 0.5304, 0.3548, 0.3309, 0.6279,
& 0.6852, 0.6925, 0.3483, 0.3970, 0.5174, 0.5500, 0.5778,
& 0.2643, 0.5151, 0.5563, 0.1978, 0.3786, 0.4835, 0.4730,
& 0.1847, 0.4443, 0.3018, 0.3018, 0.4487, 0.1370, 0.1268/
C
C
C          Create the k-d tree
C
C          CALL QUADT (NROW, NVAR, NCOL, X, LDX, IND, NBUCK, IDISCR, PART)
C
C          DO 10 I=1, NVAR
C             XKEY(I) = X(1,IND(I))
10 CONTINUE
C
C          CALL NGHBR (NVAR, XKEY, K, NROW, NCOL, X, LDX, IND, NBUCK,
& IDISCR, PART, METRIC, IPQR, PQD)
C
C          CALL WRIRN ('Indices of the nearest neighbors, IPQR.', 1, K,
& IPQR, 1, 0)
C          CALL WRRRN ('Nearest neighbor distances, PQD.', 1, K, PQD, 1, 0)
C
C          END

```

### Output

Indices of the nearest neighbors, IPQR.

1	2	3	4	5
1	3	2	5	7

Nearest neighbor distances, PQD.

1	2	3	4	5
0.000	0.791	0.847	1.201	1.352

# Reference Material

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## User Errors

IMSL routines attempt to detect user errors and handle them in a way that provides as much information to the user as possible. To do this, we recognize various levels of severity of errors, and we also consider the extent of the error in the context of the purpose of the routine; a trivial error in one situation may be serious in another. IMSL routines attempt to report as many errors as they can reasonably detect. Multiple errors present a difficult problem in error detection because input is interpreted in an uncertain context after the first error is detected.

### What Determines Error Severity

In some cases, the user's input may be mathematically correct, but because of limitations of the computer arithmetic and of the algorithm used, it is not possible to compute an answer accurately. In this case, the assessed degree of accuracy determines the severity of the error. In cases where the routine computes several output quantities, if some are not computable but most are, an error condition exists. The severity depends on an assessment of the overall impact of the error.

### Terminal errors

If the user's input is regarded as meaningless, such as  $N = -1$  when "N" is the number of equations, the routine prints a message giving the value of the erroneous input argument(s) and the reason for the erroneous input. The routine will then cause the user's program to stop. An error in which the user's input is



meaningless is the most severe error and is called a *terminal error*. Multiple terminal error messages may be printed from a single routine.

### Informational errors

In many cases, the best way to respond to an error condition is simply to correct the input and rerun the program. In other cases, the user may want to take actions in the program itself based on errors that occur. An error that may be used as the basis for corrective action within the program is called an *informational error*. If an informational error occurs, a user-retrievable code is set. A routine can return at most one informational error for a single reference to the routine. The codes for the informational error codes are printed in the error messages.

### Other errors

In addition to informational errors, IMSL routines issue error messages for which no user-retrievable code is set. Multiple error messages for this kind of error may be printed. These errors, which generally are not described in the documentation, include terminal errors as well as less serious errors. Corrective action within the calling program is not possible for these errors.

## Kinds of Errors and Default Actions

Five levels of severity of errors are defined in the STAT/LIBRARY. Each level has an associated PRINT attribute and a STOP attribute. These attributes have default settings (YES or NO), but they may also be set by the user. The purpose of having multiple error severity levels is to provide independent control of actions to be taken for errors of different severity. Upon return from an IMSL routine, exactly one error state exists. (A code 0 “error” is no informational error.) Even if more than one informational error occurs, only one message is printed (if the PRINT attribute is YES). Multiple errors for which no corrective action within the calling program is reasonable or necessary result in the printing of multiple messages (if the PRINT attribute for their severity level is YES). Errors of any of the severity levels except level 5 may be informational errors.

**Level 1: Note.** A *note* is issued to indicate the possibility of a trivial error or simply to provide information about the computations. Default attributes: PRINT = NO, STOP = NO

**Level 2: Alert.** An *alert* indicates that the user should be advised about events occurring in the software. Default attributes: PRINT = NO, STOP = NO

**Level 3: Warning.** A *warning* indicates the existence of a condition that may require corrective action by the user or calling routine. A warning error may be issued because the results are accurate to only a few decimal places, because some of the output may be erroneous but most of the output is correct, or because some assumptions underlying the analysis

technique are violated. Often no corrective action is necessary and the condition can be ignored. Default attributes: PRINT = YES, STOP = NO

**Level 4: Fatal.** A *fatal* error indicates the existence of a condition that may be serious. In most cases, the user or calling routine must take corrective action to recover. Default attributes: PRINT = YES, STOP = YES

**Level 5: Terminal.** A *terminal* error is serious. It usually is the result of an incorrect specification, such as specifying a negative number as the number of equations. These errors may also be caused by various programming errors impossible to diagnose correctly in FORTRAN. The resulting error message may be perplexing to the user. In such cases, the user is advised to compare carefully the actual arguments passed to the routine with the dummy argument descriptions given in the documentation. Special attention should be given to checking argument order and data types.

A terminal error is not an informational error because corrective action within the program is generally not reasonable. In normal usage, execution is terminated immediately when a terminal error occurs. Messages relating to more than one terminal error are printed if they occur. Default attributes: PRINT = YES, STOP = YES

The user can set PRINT and STOP attributes by calling ERSET as described in "Routines for Error Handling."

## Errors in Lower-Level Routines

It is possible that a user's program may call an IMSL routine that in turn calls a nested sequence of lower-level IMSL routines. If an error occurs at a lower level in such a nest of routines and if the lower-level routine cannot pass the information up to the original user-called routine, then a traceback of the routines is produced. The only common situation in which this can occur is when an IMSL routine calls a user-supplied routine that in turn calls another IMSL routine.

## Routines for Error Handling

There are three ways in which the user may interact with the IMSL error handling system: (1) to change the default actions, (2) to retrieve the integer code of an informational error so as to take corrective action, and (3) to determine the severity level of an error. The routines to use are ERSET, IERCD, and N1RTY, respectively.

---

# ERSET

Change the default printing or stopping actions when errors of a particular error severity level occur.

## Usage

CALL ERSET ( IERSVR, IPACT, ISACT )

## Arguments

**IERSVR** — Error severity level indicator. (Input)

If IERSVR = 0, actions are set for levels 1 to 5. If IERSVR is 1 to 5, actions are set for errors of the specified severity level.

**IPACT** — Printing action. (Input)

### IPACT Action

- 1 Do not change current setting(s).
- 0 Do not print.
- 1 Print.
- 2 Restore the default setting(s).

**ISACT** — Stopping action. (Input)

### ISACT Action

- 1 Do not change current setting(s).
- 0 Do not stop.
- 1 Stop.
- 2 Restore the default setting(s).

---

## IERCD and N1RTY

The last two routines for interacting with the error handling system, IERCD and N1RTY, are INTEGER functions and are described in the following material.

IERCD retrieves the integer code for an informational error. Since it has no arguments, it may be used in the following way:

ICODE = IERCD( )

The function retrieves the code set by the most recently called IMSL routine.

N1RTY retrieves the error type set by the most recently called IMSL routine. It is used in the following way:

ITYPE = N1RTY(1)

ITYPE = 1, 2, 4, and 5 correspond to error severity levels 1, 2, 4, and 5, respectively. ITYPE = 3 and ITYPE = 6 are both warning errors, error severity level 3. While ITYPE = 3 errors are informational errors (IERCD( ) ≠ 0), ITYPE = 6 errors are not informational errors (IERCD( ) = 0).

For software developers requiring additional interaction with the IMSL error handling system, see Aird and Howell (1991).

## Examples

### Changes to default actions

Some possible changes to the default actions are illustrated below. The default actions remain in effect for the kinds of errors not included in the call to ERSET.

To turn off printing of warning error messages:

```
CALL ERSET (3, 0, -1)
```

To stop if warning errors occur:

```
CALL ERSET (3, -1, 1)
```

To print all error messages:

```
CALL ERSET (0, 1, -1)
```

To restore all default settings:

```
CALL ERSET (0, 2, 2)
```

### Use of informational error to determine program action

In the program segment below, the Cholesky factorization of a matrix is to be performed. If it is determined that the matrix is not nonnegative definite (and often this is not immediately obvious), the program is to take a different branch.

```

      .
      .
      .
      CALL CHFAC (N, A, TOL, IRANK, R, CDR)
      IF (IERCD() .EQ. 1) THEN
C      Handle matrix that is not nonnegative definite
      .
      .
      .
      END IF
```

### Examples of All Types of Errors

The program below illustrates each of the different types of errors detected by the STAT/LIBRARY routines. If the call to ERSET was not made, messages for errors of levels 1 and 2 would not be printed.

The error messages refer to the argument names that are used in the documentation for the routine, rather than the user's name of the variable used for the argument. In the messages generated by IMSL routine CHFAC in this example, references are made to LDA and LDR, whereas in the program literals were used for these arguments. Note that error codes are printed as part of the messages for informational errors.

```
C      Specifications for local variables
      INTEGER IDO, IOPT, IRANK, N, NMISS, NOBS, NPOP, NROW, NUM
      REAL A(2,2), CHSQ, CONPER, DF, PR, R(2,2), RCOEF, STAT(20),
      & SUMRY(11), TOL, X(10), XMEAN, Y(10)
C      Specifications for subroutines
```

```

EXTERNAL CHFAC, ERSET, LETTR, SMPRR
C                                     Specifications for functions
EXTERNAL CHIDF
REAL      CHIDF
C
DATA X/-5.0, -4.0, -3.0, -2.0, -1.0, 1.0, 2.0, 3.0, 4.0, 5.0/
DATA Y/3.0, 5.0, 4.0, 5.0, 6.0, 7.0, 6.0, 8.0, 7.0, 9.0/
DATA A/2.0, 0.0, 0.0, -3.0/
C                                     Turn on printing and turn off
C                                     stopping for all error types.
CALL ERSET (0, 1, 0)
C                                     Generate level 1 informational error.
DF = 1000.0
CHSQ = -1.0
PR = CHIDF(CHSQ,DF)
C                                     Generate level 2 informational error.
DF = 1000.0
CHSQ = 10.0
PR = CHIDF(CHSQ,DF)
C                                     Generate level 3 informational error.
NOBS = 10
NUM = 11
CALL LETTR (NOBS, X, NUM, SUMRY, NMISS)
C                                     Generate level 4 informational error.
N = 2
TOL = 0.0001
CALL CHFAC (N, A, 2, TOL, IRANK, R, 2)
C                                     Generate several level 5 errors.
CALL CHFAC (N, A, 1, TOL, IRANK, R, -2)
C                                     Generate several warning errors that
C                                     do not allow corrective action
C                                     (because no codes are listed for
C                                     these errors in the document for the
C                                     routine).
IDO = 0
NROW = 10
NPOP = 100
IOPT = 1
CONPER = 0.95
CALL SMPRR (IDO, NROW, X, Y, NPOP, IOPT, XMEAN, RCOEF, CONPER,
&          STAT)
END

```

### Output

```

*** NOTE      ERROR 1 from CHIDF. Since CHSQ = -1.000000E+00 is less than
***          zero, the distribution function is zero at CHSQ.
*** ALERT    ERROR 3 from CHIDF. The normal distribution is used for large
***          degrees of freedom. However, it has produced underflow.
***          Therefore, the probability is set to 0.
*** WARNING  ERROR 3 from LETTR. NUM = 11 and the number of observations =
***          10. Since NUM is greater than the number of observations, it
***          is likely that the results are not useful.
*** WARNING  ERROR 1 from CHFAC. The leading 2 by 2 submatrix of the input
***          matrix is not nonnegative definite within the tolerance
***          defined by TOL = 1.000000E-04.
*** TERMINAL ERROR 3 from CHFAC. N = 2 and LDA = 1. N must be less than or
***          equal to LDA.
*** TERMINAL ERROR 5 from CHFAC. LDR = -2. LDR must be greater than or

```

```

***          equal to 1.
*** WARNING  ERROR 1 from SMPRR. CONPER = 9.500000E-01. The confidence
***          percentage is less than 50.0. Commonly used confidence
***          percentages are: 90.0, 95.0 or 99.0.
*** WARNING  ERROR 3 from SMPRR. The sample size, STAT(19) = 10. This is
***          less than 30. The confidence limits, which are computed using
***          a normal approximation, may not be very accurate.
*** WARNING  ERROR 7 from SMPRR. The coefficient of variation of one or
***          both of the variables exceeds 10%. The confidence limits,
***          which are computed using a normal approximation, may not be
***          very accurate.

```

### Example of Traceback

The next program illustrates a situation in which a traceback is produced. Although the traceback shows an error code associated with a terminal error, this code has no meaning to the user; the printed message contains all relevant information and it is not assumed that the user would take corrective action based on knowledge of the code.

```

C          Specifications for local variables
C          REAL      A, B, ERRABS, ERRREL, RESULT, ERREST
C          Specifications for common variables
C          REAL      PIN, QIN, SAMP
C          COMMON    PIN, QIN, SAMP
C          Specifications for subroutines
C          EXTERNAL  QDAGS
C          Specifications for functions
C          EXTERNAL F
C          REAL F
C
C          Compute the expected value of the
C          maximum order statistic in a sample
C          of size SAMP from a beta distribution.
C          A = 0.0
C          B = 1.0
C          ERRABS = 0.0
C          ERRREL = 0.001
C
C          Initialize parameters for the beta
C          order statistic of interest.
C          SAMP = 10.0
C          PIN = 2.0
C          QIN = -3.0
C
C          The parameters for the beta must be
C          nonnegative -- hence, the preceding
C          assignment causes an error.
C          CALL QDAGS (F, A, B, ERRABS, ERRREL, RESULT, ERREST)
C
C          WRITE (*, *) RESULT, ERREST
C          END
C
C          REAL FUNCTION F (X)
C          REAL      X, PIN, QIN, SAMP
C          COMMON    PIN, QIN, SAMP
C
C          F = X*BETDF(X,PIN,QIN)**(SAMP-1.0)
C          RETURN
C          END

```

### Output

```
*** TERMINAL ERROR 4 from BETDF.   QIN = -3.000000E+00 must be greater than
***                               0.0.
```

Here is a traceback of subprogram calls in reverse order:

Routine name	Error type	Error code
BETDF	5	4
Q2AGS	0	0 (Called internally)
QDAGS	0	0
USER	0	0

---

## Automatic Workspace Allocation

FORTRAN subroutines that work with arrays as input and output often require extra arrays for use as workspace while doing computations or moving around data. IMSL routines generally do not require the user explicitly to allocate such arrays for use as workspace. On most systems the workspace allocation is handled transparently. The only limitation is the actual amount of memory available on the system.

On some systems the workspace is allocated out of a stack that is passed as a FORTRAN array in a named common block `WORKSP`. A very similar use of a workspace stack is described by Fox et al. (1978, pages 116–121). (For compatibility with older versions of the IMSL Libraries, space is allocated from the `COMMON` block, if possible.)

The arrays for workspace appear as arguments in lower-level routines. For example, the IMSL routine `FREQ` (page 13), which computes frequency tabulations, needs arrays for workspace. `FREQ` allocates arrays from the common area and passes them to the lower-level routine `F2EQ`, which does the computations. In the “Comments” section of the documentation for `FREQ`, the amount of workspace is noted, and the call to `F2EQ` is described. This scheme for using lower-level routines is followed throughout the IMSL Libraries. The names of these routines have a “2” in the second position (or in the third position in double precision routines having a “D” prefix). The user can provide workspace explicitly and call directly the “2-level” routine, which is documented along with the main routine. In a very few cases, the 2-level routine allows additional options that the main routine does not allow.

Prior to returning to the calling program, a routine that allocates workspace generally deallocates that space, so that it becomes available for use in other routines. There are some exceptions to this, as noted in the section “IDO Routines” which follows later in this chapter.

### Changing the Amount of Space Allocated

*This section is relevant only to those systems on which the transparent workspace allocator is not available.*

By default, the total amount of space allocated in the common area for storage of numeric data is 5000 numeric storage units. (A numeric storage unit is the

amount of space required to store an integer or a real number. By comparison, a double precision unit is twice this amount. Therefore the total amount of space allocated in the common area for storage of numeric data is 2500 double precision units.) This space is allocated as needed for INTEGER, REAL, or other numeric data. For larger problems in which the default amount of workspace is insufficient, the user can change the allocation by supplying the FORTRAN statements to define the array in the named common block and by informing the IMSL workspace allocation system of the new size of the common array. To request 7000 units, the statements are

```
COMMON /WORKSP/ RWKSP
REAL RWKSP(7000)
CALL IWKIN(7000)
```

If an IMSL routine attempts to allocate workspace in excess of the amount available in the common stack, the routine issues a fatal error message that indicates how much space is needed and prints statements like those above to guide the user in allocating the necessary amount. The program below uses IMSL routine PERMA (page 1266) to permute rows or columns of a matrix. This routine requires workspace equal to the number of columns, which in this example is too large. (Note that the work vector RWKSP must also provide extra space for bookkeeping.)

```
C                               Specifications for local variables
INTEGER   NRA, NCA, LDA, IPERMU(6000), IPATH
REAL A(2,6000)

C                               Specifications for subroutines
EXTERNAL PERMA

C
NRA = 2
NCA = 6000
LDA = 2

C                               Initialize permutation index
DO 10 I = 1, NCA
    IPERMU(I) = NCA + 1 - I
10 CONTINUE
IPATH = 2
CALL PERMA (NRA, NCA, A, LDA, IPERMU, IPATH, A, LDA)
END
```

### Output

```
*** TERMINAL ERROR 10 from PERMA.  Insufficient workspace for current
*** allocation(s).  Correct by calling IWKIN from main program with
*** the three following statements: (REGARDLESS OF PRECISION)
***     COMMON /WORKSP/ RWKSP
***     REAL RWKSP(6018)
***     CALL IWKIN(6018)
*** TERMINAL ERROR 10 from PERMA.  Workspace allocation was based on NCA =
*** 6000.
```

In most cases, the amount of workspace is dependent on the parameters of the problem so the amount needed is known exactly. In a few cases, however, the amount of workspace is dependent on the data (for example, if it is necessary to count all of the unique values in a vector), so the IMSL routine cannot tell in



advance exactly how much workspace is needed. In such cases the error message printed is an estimate of the amount of space required.

### IDO Routines

Some routines with an argument named "IDO" allocate workspace automatically and store intermediate results in elements of workspace that are referenced in subsequent calls. Typically, these routines are called in a loop. With each call, some rows of the data set are input to the routine and statistics stored in workspace are updated. In this case, the workspace must be preserved between calls.

For these routines, when IDO indicates this is the first call, the routine allocates workspace; when IDO indicates this is the last call, the routine deallocates the workspace. Because of the way this workspace is allocated and deallocated, no IMSL routine requiring additional automatic workspace can be used between these two calls. If it is necessary to call additional routines requiring workspace, use the 2-level routines and explicitly allocate the work arrays.

Not all IDO routines require workspace to be preserved between their first and last call. Some may not even use workspace. Others may allocate and deallocate workspace with each call. The statement "workspace should not be changed between calls" will be in the description of the "IDO" routine that requires that workspace be preserved. (This statement will occur in the description of one or more of the workspace arguments for the 2-level routine.)

### Character Workspace

Since character arrays cannot be equivalenced with numeric arrays, a separate named common block WKSPCH is provided for character workspace. In most respects this stack is managed in the same way as the numeric stack. The default size of the character workspace is 2000 character units. (A character unit is the amount of space required to store one character.) The routine analogous to IWKIN used to change the default allocation is IWKCIN.

---

## Machine-Dependent Constants

The function subprograms in this section return machine-dependent information and can be used to enhance portability of programs between different computers. The routines IMACH, AMACH and DMACH describe the computer's arithmetic. The routine UMACH describes the input, output, and error output unit numbers.

```
INTEGER FUNCTION IMACH(I)
```

IMACH retrieves machine integer constants that define the arithmetic used by the computer.

IMACH(1) = Number of bits per integer storage unit.

IMACH(2) = Number of characters per integer storage unit.

Integers are represented in *M*-digit, base *A* form as

$$\sigma \sum_{k=0}^M x_k A^k$$

where  $\sigma$  is the sign and  $0 \leq x_k < A$ ,  $k = 0, \dots, M$ . Then,

IMACH(3) =  $A$ , the base.

IMACH(4) =  $M$ , the number of base- $A$  digits.

IMACH(5) =  $AM - 1$ , the largest integer.

The machine model assumes that floating-point numbers are represented in normalized  $N$ -digit, base  $B$  form as

$$\sigma B^E \sum_{k=1}^N x_k B^{-k}$$

where  $\sigma$  is the sign,  $0 < x_1 < B$ ,  $0 \leq x_k < B$ ,  $k = 2, \dots, N$  and

$$E_{\min} \leq E \leq E_{\max}$$

Then,

IMACH(6) =  $B$ , the base.

IMACH(7) =  $N_s$ , the number of base- $B$  digits in single precision

IMACH(8) =  $E_{\min_s}$ , the smallest single precision exponent

IMACH(9) =  $E_{\max_s}$ , the largest single precision exponent

IMACH(10) =  $N_d$ , the number of base- $B$  digits in double precision

IMACH(11) =  $E_{\min_d}$ , the smallest double precision exponent

IMACH(12) =  $E_{\max_d}$ , the number of base- $B$  digits in double precision

REAL FUNCTION AMACH(I)

The function subprogram AMACH retrieves real machine constants that define the computer's real or single-precision arithmetic. Such floating-point numbers are represented in normalized  $N_s$ -digit, base  $B$  form as

$$\sigma B^E \sum_{k=1}^{N_s} x_k B^{-k}$$

where  $\sigma$  is the sign,  $0 < x_1 < B$ ,  $0 \leq x_k < B$ ,  $k = 2, \dots, N_s$  and

$$E_{\min_s} \leq E \leq E_{\max_s}$$

Note that  $B = \text{IMACH}(6)$ ,  $N_s = \text{IMACH}(7)$ ,

$$E_{\min_s} = \text{IMACH}(8), \text{ and } E_{\max_s} = \text{IMACH}(9)$$

The IEEE standard for binary arithmetic (see IEEE 1985) specifies *quiet* NaN (not a number) as the result of various invalid or ambiguous operations, such as 0/0. The intent is that AMACH(6) return a *signaling* NaN. On IEEE format computers that do not support signaling NaN, a quiet NaN is returned. If the

machine does not support a NaN, a special value near AMACH(2) is returned for AMACH(6). On computers that do not have a special representation for infinity, AMACH(7) returns the same value as AMACH(2).

AMACH is defined by the following table:

AMACH(1) =  $B^{E_{\min_s} - 1}$ , the smallest normalized positive number.

AMACH(2) =  $B^{E_{\max_s}} (1 - B^{-N_s})$ , the largest number.

AMACH(3) =  $B^{-N_s}$ , the smallest relative spacing.

AMACH(4) =  $B^{1 - N_s}$ , the largest relative spacing.

AMACH(5) =  $\log_{10} B$

AMACH(6) = NaN (signaling not a number).

AMACH(7) = positive machine infinity.

AMACH(8) = negative machine infinity.

DOUBLE PRECISION FUNCTION DMACH(I)

The function subprogram DMACH retrieves real machine constants that define the computer's double precision arithmetic. Such double-precision floating-point numbers are represented in normalized  $Nd$ -digit, base  $B$  form as

$$\sigma B^E \sum_{k=1}^{N_d} x_k B^{-k}$$

where  $\sigma$  is the sign,  $0 < x_1 < B$ ,  $0 \leq x_k < B$ ,  $k = 2, \dots, Nd$  and

$$E_{\min_d} \leq E \leq E_{\max_d}$$

Note that  $B = \text{IMACH}(6)$ ,  $Nd = \text{IMACH}(10)$ ,

$E_{\min_d} = \text{IMACH}(11)$ , and  $E_{\max_d} = \text{IMACH}(12)$

The IEEE standard for binary arithmetic (see IEEE 1985) specifies quiet NaN (not a number) as the result of various invalid or ambiguous operations, such as 0/0. The intent is that DMACH(6) return a *signaling* NaN. On IEEE format computers that do not support signaling NaN, a quiet NaN is returned. If the machine does not support a NaN, a special value near DMACH(2) is returned for DMACH(6). On computers that do not have a special representation for infinity, DMACH(7) returns the same value as DMACH(2).

DMACH is defined by the following table:

DMACH(1) =  $B^{E_{\min_d} - 1}$  the smallest normalized positive number.

DMACH(2) =  $B^{E_{\max_d}} (1 - B^{-N_d})$ , the largest number.

DMACH(3) =  $B^{-N_d}$ , the smallest relative spacing.

DMACH(4) =  $B^{1-N_d}$ , the largest relative spacing.

DMACH(5) =  $\log_{10} B$

DMACH(6) = NaN (signaling not a number).

DMACH(7) = positive machine infinity.

DMACH(8) = negative machine infinity.

LOGICAL FUNCTION IFNAN(X), DIFNAN(DX)

The logical function IFNAN checks if the REAL argument X is NaN (not a number). Similarly, DIFNAN checks if the DOUBLE PRECISION argument DX is NaN.

The functions IFNAN and DIFNAN are provided to facilitate the transfer of programs across computer systems. This is because the check for NaN can be tricky and not portable across computer systems that do not adhere to the IEEE standard. For example, on computers that support the IEEE standard for binary arithmetic (see IEEE 1985), NaN is specified as a bit format not equal to itself. Thus, the check is performed as

```
IFNAN = X .NE. X
```

On other computers that do not use IEEE floating-point format, the check can be performed in single precision as

```
IFNAN = X .EQ. AMACH(6)
```

The function IFNAN or DIFNAN is equivalent to the specification of the function Isnan listed in the Appendix, (IEEE 1985). The following example illustrates the use of IFNAN. If X is NaN, a message is printed instead of X. (Routine UMACH, which is described in the following section, is used to retrieve the output unit number for printing the message.)

```
INTEGER      NOUT
REAL         AMACH, X
LOGICAL      IFNAN
EXTERNAL     AMACH, IFNAN, UMACH

C
CALL UMACH (2, NOUT)

C
X = AMACH(6)
IF (IFNAN(X)) THEN
    WRITE (NOUT,*) ' X is NaN (not a number).'
ELSE
    WRITE (NOUT,*) ' X = ', X
END IF

C
```

END

### Output

X is NaN (not a number).

```
SUBROUTINE UMACH(N, NUNIT)
```

Routine UMACH sets or retrieves the input, output, or error output device unit numbers. UMACH is set automatically so that the default FORTRAN unit numbers for standard input, standard output, and standard error are used. These unit numbers can be changed by inserting a call to UMACH at the beginning of the main program that calls STAT/LIBRARY routines. If these unit numbers are changed from the standard values, the user should insert an appropriate OPEN statement in the calling program. The calling sequence for UMACH is

```
CALL UMACH (N, NUNIT)
```

where NUNIT is the input, output, or error output unit number that is either retrieved or set, depending on which value of N is selected.

The arguments are summarized by the following table:

N	Effect
1	Retrieves input unit number in NUNIT.
2	Retrieves output unit number in NUNIT.
3	Retrieves error output unit number in NUNIT.
-1	Sets the input unit number to NUNIT.
-2	Sets the output unit number to NUNIT.
-3	Sets the error output unit number to NUNIT.

If the value of N is negative, the input, output, or error output unit number is reset to NUNIT. If the value of N is positive, the input, output, or error output unit number is returned in NUNIT.

In the following example, a terminal error is issued from the STAT/LIBRARY AMACH function since the argument is invalid. With a call to UMACH, the error message will be written to a local file named "CHECKERR".

```
INTEGER      N, AMACH
REAL         X
EXTERNAL     AMACH, UMACH
C
C                                     Set Parameter
N = 0
C
CALL UMACH (-3, 9)
OPEN (UNIT=9,FILE='CHECKERR')
X = AMACH(N)
END
```

The output from this example, written to 'CHECKERR' is:

```
*** TERMINAL ERROR 5 from AMACH. The argument must be between 1 and 8
*** inclusive. N = 0
```

## Missing Values

Missing values in IMSL routines are always indicated by NaN (not a number). This is AMACH(6) in single precision and DMACH(6) in double precision. There is no missing-value indicator for FORTRAN integers. Users will almost always have to convert from their missing value indicators to NaN. This is easily accomplished, as the following example illustrates. In this example, “-9.9” in the input represents a missing value, which is converted to NaN (not a number). After the input matrix is printed with missing values as “NaN”, IMSL routine UVSTA is called to obtain univariate summary statistics. When a frequency is missing, the entire row is omitted from the analysis, whereas only the data element in question is omitted otherwise.

```
INTEGER IDO, IFRQ, IPRINT, IWT, LDSTAT, LDX, MOPT, NROW, NVAR
REAL CONPRM, CONPRV
PARAMETER (CONPRM=95.0, CONPRV=90.0, IDO=0, IFRQ=3, IPRINT=1,
&          IWT=0, LDSTAT=15, LDX=10, MOPT=1, NROW=10, NVAR=2)
C
  INTEGER I, J, NRMIS
  REAL AMACH, STAT(LDSTAT,NVAR), X(LDX,3)
  EXTERNAL AMACH, UVSTA, WRRRN
C
  DATA X/1.2, 1.6, -9.9, 1.6, 1.3, -9.9, 1.1, 1.3, 1.5, 1.2, 8.2,
& 7.6, 10.3, -9.9, 8.2, 11.3, -9.9, 7.4, -9.9, -9.9, 10, 7,
& 4, -9.9, 10, 11, 4, 5, -9.9, 10/
C
  DO 20 I=1, NROW
    DO 10 J=1, 3
      IF (X(I,J) .EQ. -9.9) X(I,J) = AMACH(6)
10 CONTINUE
20 CONTINUE
C
  CALL WRRRN ('X', NROW, 3, X, LDX, 0)
C
  CALL UVSTA (IDO, NROW, NVAR, X, LDX, IFRQ, IWT, MOPT, CONPRM,
&            CONPRV, IPRINT, STAT, LDSTAT, NRMIS)
C
  END
```

## Output

	X		
	1	2	3
1	1.20	8.20	10.00
2	1.60	7.60	7.00
3	NaN	10.30	4.00
4	1.60	NaN	NaN
5	1.30	8.20	10.00
6	NaN	11.30	11.00
7	1.10	NaN	4.00
8	1.30	7.40	5.00
9	1.50	NaN	NaN
10	1.20	NaN	10.00

Univariate Statistics from UVSTA					
Variable	Mean	Variance	Std. Dev.	Skewness	Kurtosis
1	1.2848	0.02176	0.1475	1.2727	0.5878

2	8.9298	2.26170	1.5039	0.7405	-1.1927
Variable	Minimum	Maximum	Range	Coef. Var.	Count
1	1.1000	1.6000	0.5000	0.1148	46.0000
2	7.4000	11.3000	3.9000	0.1684	47.0000
Variable	Lower CLM	Upper CLM	Lower CLV	Upper CLV	
1	1.2410	1.3286	0.01588	0.03199	
2	8.4882	9.3713	1.65588	3.30921	

---

## Matrix Storage Modes

In this section, the word *matrix* will be used to refer to a mathematical object, and the word *array* will be used to refer to its representation as a FORTRAN data structure.

### General Mode

A *general* matrix is an  $N \times N$  matrix  $A$ . It is stored in a FORTRAN array that is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter `LDA` is called the *leading dimension* of  $A$ . It must be at least as large as  $N$ . IMSL general matrix subprograms only refer to values  $A_{ij}$  for  $i = 1, \dots, N$  and  $j = 1, \dots, N$ . The data type of a general array can be one of `REAL`, `DOUBLE PRECISION`, or `COMPLEX`. If your FORTRAN compiler allows, the nonstandard data type `DOUBLE COMPLEX` can also be declared.

### Rectangular Mode

A *rectangular* matrix is an  $M \times N$  matrix  $A$ . It is stored in a FORTRAN array that is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter `LDA` is called the *leading dimension* of  $A$ . It must be at least as large as  $M$ . IMSL rectangular matrix subprograms only refer to values  $A_{ij}$  for  $i = 1, \dots, M$  and  $j = 1, \dots, N$ . The data type of a rectangular array can be `REAL`, `DOUBLE PRECISION`, or `COMPLEX`. If your FORTRAN compiler allows, you can declare the nonstandard data type `DOUBLE COMPLEX`.

### Symmetric Mode

A symmetric matrix is a square  $N \times N$  matrix  $A$ , such that  $AT = A$ . ( $AT$  is the transpose of  $A$ .) It is stored in a FORTRAN array that is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter `LDA` is called the *leading dimension* of  $A$ . It must be at least as large as  $N$ . IMSL symmetric matrix subprograms only refer to the upper or to the lower half of  $A$  (i.e., to values  $A_{ij}$  for  $i = 1, \dots, N$  and  $j = 1, \dots, N$ , or  $A_{ij}$  for

$j = 1, \dots, N$  and  $i = j, \dots, N$ . The data type of a symmetric array can be one of REAL or DOUBLE PRECISION. Use of the upper half of the array is denoted in the BLAS that compute with symmetric matrices using the CHARACTER\*1 flag UPLO = 'U'. Otherwise, UPLO = 'L' denotes that the lower half of the array is used.

## Hermitian Mode

A *Hermitian* matrix is a square  $N \times N$  matrix  $A$ , such that

$$\overline{A}^T = A$$

The matrix

$$\overline{A}$$

is the complex conjugate of  $A$  and

$$A^H \equiv \overline{A}^T$$

is the conjugate transpose of  $A$ . For Hermitian matrices,  $A^H = A$ . The matrix is stored in a FORTRAN array that is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter LDA is called the *leading dimension* of  $A$ . It must be at least as large as  $N$ . IMSL Hermitian matrix subprograms only refer to the upper or to the lower half of  $A$  (i.e., to values  $A_{ij}$  for  $i = 1, \dots, N$  and  $j = i, \dots, N$ , or  $A_{ij}$  for  $j = 1, \dots, N$  and  $i = j, \dots, N$ ). Use of the upper half of the array is denoted in the BLAS that compute with Hermitian matrices using the CHARACTER\*1 flag UPLO = 'U'. Otherwise, UPLO = 'L' denotes that the lower half of the array is used. The data type of a Hermitian array can be COMPLEX or, if your FORTRAN compiler allows, the nonstandard data type DOUBLE COMPLEX.

## Triangular Mode

A *triangular* matrix is a square  $N \times N$  matrix  $A$  such that values  $A_{ij} = 0$  for  $i < j$  or  $A_{ij} = 0$  for  $i > j$ . The first condition defines a *lower* triangular matrix while the second condition defines an *upper* triangular matrix. A lower triangular matrix  $A$  is stored in the lower triangular part of a FORTRAN array  $A$ . An upper triangular matrix is stored in the upper triangular part of a FORTRAN array. Triangular matrices are called *unit* triangular whenever  $A_{jj} = 1, j = 1, \dots, N$ . For unit triangular matrices, only the strictly lower or upper parts of the array are referenced. This is denoted in the BLAS that compute with triangular matrices using the CHARACTER\*1 flag DIAG = 'U'. Otherwise, DIAG = 'N' denotes that the diagonal array terms should be used. For unit triangular matrices, the diagonal terms are each used with the mathematical value 1. The array diagonal term does not need to be 1.0 in this usage. Use of the upper half of the array is denoted in the BLAS that compute with triangular matrices, IMSL MATH/LIBRARY, using the CHARACTER\*1 flag UPLO = 'U'. Otherwise,



UPLO = 'L' denotes that the lower half of the array is used. The data type of an array that contains a triangular matrix can be one of REAL, DOUBLE PRECISION, or COMPLEX. If your FORTRAN compiler allows, the nonstandard data type DOUBLE COMPLEX can also be declared.

## Band Storage Mode

A *band matrix* is an  $M \times N$  matrix  $A$  with all of its nonzero elements “close” to the main diagonal. Specifically, values  $A_{ij} = 0$  if  $i - j > NLCA$  or  $j - i > NUCA$ . The integers NLCA and NUCA are the *lower* and *upper* band widths. The integer  $m = NLCA + NUCA + 1$  is the total band width. The diagonals, other than the main diagonal, are called *codiagonals*. While any  $M \times N$  matrix is a band matrix, the band matrix mode is most useful only when the number of nonzero codiagonals is much less than  $m$ .

In the band storage mode, the NLCA lower codiagonals and NUCA upper codiagonals are stored in the rows of a FORTRAN array of dimension  $m \times N$ . The elements are stored in the same column of the array as they are in the matrix. The values  $A_{ij}$  inside the band width are stored in array positions  $(i - j + NUCA + 1, j)$ . This array is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter LDA is called the *leading dimension* of  $A$ . It must be at least as large as  $m$ . The data type of a band matrix array can be one of REAL, DOUBLE PRECISION, COMPLEX or, if your FORTRAN compiler allows, the nonstandard data type DOUBLE COMPLEX. Use of the CHARACTER\*1 flag TRANS = 'N' in the BLAS specifies that the matrix  $A$  is used. The flag value TRANS = 'T' uses  $A^T$  while TRANS = 'C' uses

$$\overline{A}^T$$

For example, consider a real  $5 \times 5$  band matrix with 1 lower and 2 upper codiagonals, stored in the FORTRAN array declared by the following statements:

```
PARAMETER (N=5, NLCA=1, NUCA=2)
REAL A(NLCA+NUCA+1, N)
```

The matrix  $A$  has the form

$$A = \begin{pmatrix} A_{11} & A_{12} & A_{13} & 0 & 0 \\ A_{21} & A_{22} & A_{23} & A_{24} & 0 \\ 0 & A_{32} & A_{33} & A_{34} & A_{35} \\ 0 & 0 & A_{43} & A_{44} & A_{45} \\ 0 & 0 & 0 & A_{54} & A_{55} \end{pmatrix}$$

As a FORTRAN array, it is

$$A = \begin{array}{ccccc} \times & \times & A_{13} & A_{24} & A_{35} \\ \times & A_{12} & A_{23} & A_{34} & A_{45} \\ A_{11} & A_{22} & A_{33} & A_{44} & A_{45} \\ A_{21} & A_{32} & A_{43} & A_{54} & \times \end{array}$$

The entries marked with an  $\times$  in the above array are not referenced by the IMSL band symmetric subprograms.

## Band Symmetric Storage Mode

A *band symmetric* matrix is a band matrix that is also symmetric. The band symmetric storage mode is similar to the band mode except only the lower or upper codiagonals are stored.

In the band symmetric storage mode, the `NCODA` upper codiagonals are stored in the rows of a FORTRAN array of dimension  $(\text{NCODA} + 1) \times N$ . The elements are stored in the same column of the array as they are in the matrix. Specifically, values  $A_{ij}$ ,  $j \geq i$  inside the band are stored in array positions  $(i - j + \text{NCODA} + 1, j)$ . This is the storage mode designated by using the CHARACTER\*1 flag `UPLO = 'U'` in Level 2 BLAS that compute with band symmetric matrices. Alternatively,  $A_{ij}$ ,  $j \leq i$ , inside the band, are stored in array positions  $(i - j + 1, j)$ . This is the storage mode designated by using the CHARACTER\*1 flag `UPLO = 'L'` in these Level 2 BLAS. The array is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter `LDA` is called the *leading dimension* of  $A$ . It must be at least as large as `NCODA + 1`. The data type of a band symmetric array can be `REAL` or `DOUBLE PRECISION`. For example, consider a real  $5 \times 5$  band matrix with 2 codiagonals. Its FORTRAN declaration is

```
PARAMETER (N=5, NCODA=2)
REAL A(NCODA+1, N)
```

The matrix  $A$  has the form

$$A = \begin{array}{ccccc} A_{11} & A_{12} & A_{13} & 0 & 0 \\ A_{12} & A_{22} & A_{23} & A_{24} & 0 \\ A_{13} & A_{23} & A_{33} & A_{34} & A_{35} \\ 0 & A_{24} & A_{34} & A_{44} & A_{45} \\ 0 & 0 & A_{35} & A_{45} & A_{55} \end{array}$$

Since  $A$  is symmetric, the values  $A_{ij} = A_{ji}$ . In the FORTRAN array, it is

$$A = \begin{pmatrix} \times & \times & A_{13} & A_{24} & A_{35} \\ \times & A_{12} & A_{23} & A_{34} & A_{45} \\ A_{11} & A_{22} & A_{33} & A_{44} & A_{45} \end{pmatrix}$$

The entries marked with an  $\times$  in the above array are not referenced by the IMSL band symmetric subprograms.

An alternate storage mode for band symmetric matrices is designated using the CHARACTER\*1 flag UPLO = 'L' in Level 2 BLAS that compute with band symmetric matrices. In that case, the example matrix is represented as

$$A = \begin{pmatrix} A_{11} & A_{22} & A_{33} & A_{44} & A_{55} \\ A_{12} & A_{23} & A_{34} & A_{45} & \times \\ A_{13} & A_{24} & A_{35} & \times & \times \end{pmatrix}$$

## Band Hermitian Storage Mode

A *band Hermitian* matrix is a band matrix that is also Hermitian. The band Hermitian mode is a complex analogue of the band symmetric mode.

In the band Hermitian storage mode, the NCODA upper codiagonals are stored in the rows of a FORTRAN array of dimension  $(\text{NCODA} + 1) \times N$ . The elements are stored in the same column of the array as they are in the matrix. In the Level 2 BLAS, IMSL MATH/LIBRARY, this is denoted by using the CHARACTER\*1 flag UPLO = 'U'. The array is declared by the following statement:

```
DIMENSION A(LDA,N)
```

The parameter LDA is called the *leading dimension* of A. It must be at least as large as  $(\text{NCODA} + 1)$ . The data type of a band Hermitian array can be COMPLEX or, if your FORTRAN compiler allows, the nonstandard data type DOUBLE COMPLEX.

For example, consider a complex  $5 \times 5$  band matrix with 2 codiagonals. Its FORTRAN declaration is

```
PARAMETER (N=5, NCODA = 2)
COMPLEX A(NCODA + 1, N)
```

The matrix A has the form

$$A = \begin{pmatrix} A_{11} & A_{12} & A_{13} & 0 & 0 \\ \bar{A}_{12} & A_{22} & A_{23} & A_{24} & 0 \\ \bar{A}_{13} & \bar{A}_{23} & A_{33} & A_{34} & A_{35} \\ 0 & \bar{A}_{24} & \bar{A}_{34} & A_{44} & A_{45} \\ 0 & 0 & \bar{A}_{35} & \bar{A}_{45} & A_{55} \end{pmatrix}$$

where the value

$$\bar{A}_{ij}$$

is the complex conjugate of  $A_{ij}$ . This matrix represented as a FORTRAN array is

$$A = \begin{pmatrix} \times & \times & A_{13} & A_{24} & A_{35} \\ \times & A_{12} & A_{23} & A_{34} & A_{45} \\ A_{11} & A_{22} & A_{33} & A_{44} & A_{55} \end{pmatrix}$$

The entries marked with an  $\times$  in the above array are not referenced by the IMSL band Hermitian subprograms.

An alternate storage mode for band Hermitian matrices is designated using the CHARACTER\*1 flag UPLO = 'L' in Level 2 BLAS that compute with band Hermitian matrices. In that case, the example matrix is represented as

$$A = \begin{pmatrix} A_{11} & A_{22} & A_{33} & A_{44} & A_{55} \\ \bar{A}_{12} & \bar{A}_{23} & \bar{A}_{34} & \bar{A}_{45} & \times \\ \bar{A}_{13} & \bar{A}_{24} & \bar{A}_{35} & \times & \times \end{pmatrix}$$

## Band Triangular Storage Mode

A *band triangular* matrix is a band matrix that is also triangular. In the band triangular storage mode, the NCODA codiagonals are stored in the rows of a FORTRAN array of dimension  $(\text{NCODA} + 1) \times N$ . The elements are stored in the same column of the array as they are in the matrix. For usage in the Level 2 BLAS, the CHARACTER\*1 flag DIAG has the same meaning as used in section "Triangular Storage Mode". The flag UPLO has the meaning analogous with its usage in the section "Band Symmetric Storage Mode". This array is declared by the following statement:

```
DIMENSION A(LDA, N)
```

The parameter LDA is called the *leading dimension* of A. It must be at least as large as  $(\text{NCODA} + 1)$ .

For example, consider a  $5 \times 5$  band upper triangular matrix with 2 codiagonals. Its FORTRAN declaration is

```
PARAMETER (N = 5, NCODA = 2)
COMPLEX A(NCODA + 1, N)
```

The matrix A has the form

$$A = \begin{pmatrix} A_{11} & A_{12} & A_{13} & 0 & 0 \\ 0 & A_{22} & A_{23} & A_{24} & 0 \\ 0 & 0 & A_{33} & A_{34} & A_{35} \\ 0 & 0 & 0 & A_{44} & A_{45} \\ 0 & 0 & 0 & 0 & A_{55} \end{pmatrix}$$

This matrix represented as a FORTRAN array is

$$A = \begin{pmatrix} \times & \times & A_{13} & A_{24} & A_{35} \\ \times & A_{12} & A_{23} & A_{34} & A_{45} \\ A_{11} & A_{22} & A_{33} & A_{44} & A_{55} \end{pmatrix}$$

This corresponds to the CHARACTER\*1 flags DIAG = 'N' and UPLO = 'U'.

The matrix  $AT$  is represented as the FORTRAN array

$$A = \begin{pmatrix} A_{11} & A_{22} & A_{33} & A_{44} & A_{55} \\ A_{12} & A_{23} & A_{34} & A_{45} & \times \\ A_{13} & A_{24} & A_{35} & \times & \times \end{pmatrix}$$

This corresponds to the CHARACTER\*1 flags DIAG = 'N' and UPLO = 'L'. In both examples, the entries indicated with an  $\times$  are not referenced by IMSL subprograms.

## Codiagonal Band Symmetric Storage Mode

This is an alternate storage mode for band symmetric matrices. It is not used by any of the BLAS. Storing data in a form transposed from the **Band Symmetric Storage Mode** maintains unit spacing between consecutive referenced array elements. This data structure is used to get good performance in the Cholesky decomposition algorithm that solves positive definite symmetric systems of linear equations  $Ax = b$ . The data type can be REAL or DOUBLE PRECISION. In the codiagonal band symmetric storage mode, the NCODA upper codiagonals and right-hand-side are stored in columns of this FORTRAN array. This array is declared by the following statement:

```
DIMENSION A(LDA, NCODA + 2)
```

The parameter LDA is the *leading positive dimension* of A. It must be at least as large as  $N + NCODA$ .

Consider a real symmetric  $5 \times 5$  matrix with 2 codiagonals

$$A = \begin{pmatrix} A_{11} & A_{12} & A_{13} & 0 & 0 \\ A_{12} & A_{22} & A_{23} & A_{24} & 0 \\ A_{13} & A_{23} & A_{33} & A_{34} & A_{35} \\ 0 & A_{24} & A_{34} & A_{44} & A_{45} \\ 0 & 0 & A_{35} & A_{45} & A_{55} \end{pmatrix}$$

and a right-hand-side vector

$$b = \begin{pmatrix} b_1 \\ b_2 \\ b_3 \\ b_4 \\ b_5 \end{pmatrix}$$

A FORTRAN declaration for the array to hold this matrix and right-hand-side vector is

```
PARAMETER (N = 5, NCODA = 2, LDA = N + NCODA)
REAL A(LDA, NCODA + 2)
```

The matrix and right-hand-side entries are placed in the FORTRAN array *A* as follows:

$$A = \begin{pmatrix} \times & \times & \times & \times \\ \times & \times & \times & \times \\ A_{11} & \times & \times & b_1 \\ A_{22} & A_{12} & \times & b_2 \\ A_{33} & A_{23} & A_{13} & b_3 \\ A_{44} & A_{34} & A_{24} & b_4 \\ A_{55} & A_{45} & A_{35} & b_5 \end{pmatrix}$$

Entries marked with a  $\times$  do not need to be defined. Certain of the IMSL band symmetric subprograms will initialize and use these values during the solution process. When a solution is computed, the  $b_i$ ,  $i = 1, \dots, 5$ , are replaced by  $x_i$ ,  $i = 1, \dots, 5$ .

The nonzero  $A_{ij}$ ,  $j \geq i$ , are stored in array locations  $A(j + NCODA, (j - i) + 1)$ . The right-hand-side entries  $b_j$  are stored in locations  $A(j + NCODA, NCODA + 2)$ . The solution entries  $x_j$  are returned in  $A(j + NCODA, NCODA + 2)$ .

### Codiagonal Band Hermitian Storage Mode

This is an alternate storage mode for band Hermitian matrices. It is not used by any of the BLAS, IMSL MATH/LIBRARY. In the codiagonal band Hermitian storage mode, the real and imaginary parts of the  $2 * NCODA + 1$  upper codiagonals and right-hand-side are stored in columns of a FORTRAN array. Note that there is no explicit use of the COMPLEX or the nonstandard data type DOUBLE COMPLEX data type in this storage mode.

For *Hermitian* complex matrices,

$$A = U + \sqrt{-1}V$$

where  $U$  and  $V$  are real matrices. They satisfy the conditions  $U = UT$  and  $V = -VT$ . The right-hand-side

$$b = c + \sqrt{-1}d$$

where  $c$  and  $d$  are real vectors. The solution vector is denoted as

$$x = u + \sqrt{-1}v$$

where  $u$  and  $v$  are real. The storage is declared with the following statement

```
DIMENSION A(LDA, 2*NCODA + 3)
```

The parameter LDA is the *leading positive dimension* of A. It must be at least as large as  $N + NCODA$ .

The diagonal terms  $U_{jj}$  are stored in array locations  $A(j + NCODA, 1)$ . The diagonal  $V_{jj}$  are zero and are not stored. The nonzero  $U_{ij}, j > i$ , are stored in locations  $A(j + NCODA, 2 * (j - i))$ .

The nonzero  $V_{ij}$  are stored in locations  $A(j + NCODA, 2 * (j - i) + 1)$ . The right side vector  $b$  is stored with  $c_j$  and  $d_j$  in locations  $A(j + NCODA, 2 * NCODA + 2)$  and  $A(j + NCODA, 2 * NCODA + 3)$  respectively. The real and imaginary parts of the solution,  $u_j$  and  $v_j$ , respectively overwrite  $c_j$  and  $d_j$ .

Consider a complex hermitian  $5 \times 5$  matrix with 2 codiagonals

$$A = \begin{pmatrix} U_{11} & U_{12} & U_{13} & 0 & 0 \\ U_{12} & U_{22} & U_{23} & U_{24} & 0 \\ U_{13} & U_{23} & U_{33} & U_{34} & U_{35} \\ 0 & U_{24} & U_{34} & U_{44} & U_{45} \\ 0 & 0 & U_{35} & U_{45} & U_{55} \end{pmatrix} + \sqrt{-1} \begin{pmatrix} 0 & V_{12} & V_{13} & 0 & 0 \\ -V_{12} & 0 & V_{23} & V_{24} & 0 \\ -V_{13} & -V_{23} & 0 & V_{34} & V_{35} \\ 0 & -V_{24} & -V_{34} & 0 & V_{45} \\ 0 & 0 & -V_{35} & -V_{45} & 0 \end{pmatrix}$$

and a right-hand-side vector

$$b = \begin{pmatrix} c_1 \\ c_2 \\ c_3 \\ c_4 \\ c_5 \end{pmatrix} + \sqrt{-1} \begin{pmatrix} d_1 \\ d_2 \\ d_3 \\ d_4 \\ d_5 \end{pmatrix}$$

A FORTRAN declaration for the array to hold this matrix and right-hand-side vector is

```
PARAMETER (N = 5, NCODA = 2, LDA = N + NCODA)
REAL A(LDA, 2*NCODA + 3)
```

The matrix and right-hand-side entries are placed in the FORTRAN array A as follows:

$$A = \begin{array}{cccccccc}
\times & \times & \times & \times & \times & \times & \times & \times \\
\times & \times & \times & \times & \times & \times & \times & \times \\
U_{11} & \times & \times & \times & \times & c_1 & d_1 \\
U_{22} & U_{12} & V_{12} & \times & \times & c_2 & d_2 \\
U_{33} & U_{23} & V_{23} & U_{13} & V_{13} & c_3 & d_3 \\
U_{44} & U_{34} & V_{34} & U_{24} & V_{24} & c_4 & d_4 \\
U_{55} & U_{45} & V_{45} & U_{35} & V_{35} & c_5 & d_5
\end{array}$$

Entries marked with an  $\times$  do not need to be defined.

---

## Reserved Names

When writing programs accessing the STAT/LIBRARY, the user should choose FORTRAN names that do not conflict with names of IMSL subroutines, functions, or named common blocks, such as the workspace common block WORKSP (page 1332). The user needs to be aware of two types of name conflicts that can arise. The first type of name conflict occurs when a name (technically a *symbolic name*) is not uniquely defined within a program unit (either a main program or a subprogram). For example, such a name conflict exists when the name RCURV is used to refer both to a type REAL variable and to the IMSL subroutine RCURV in a single program unit. Such errors are detected during compilation and are easy to correct. The second type of name conflict, which can be more serious, occurs when names of program units and named common blocks are not unique. For example, such a name conflict would be caused by the user defining a subroutine named WORKSP and also referencing an STAT/LIBRARY subroutine that uses the named common block WORKSP. Likewise, the user must not define a subprogram with the same name as a subprogram in the STAT/LIBRARY, that is referenced directly by the user's program or is referenced indirectly by other STAT/LIBRARY subprograms.

The STAT/LIBRARY consists of many routines, some that are described in the *User's Manual* and others that are not intended to be called by the user and, hence, that are not documented. If the choice of names were completely random over the set of valid FORTRAN names, and if a program uses only a small subset of the STAT/LIBRARY, the probability of name conflicts is very small. Since names are usually chosen to be mnemonic, however, the user may wish to take some precautions in choosing FORTRAN names.

Many IMSL names consist of a root name that may have a prefix to indicate the type of the routine. For example, the IMSL single precision subroutine for fitting a polynomial by least squares has the name RCURV, which is the root name, and the corresponding IMSL double precision routine has the name DRCURV. Associated with these two routines are R2URV and DR2URV. RCURV and DRCURV are listed in the Alphabetical Index of Routines, but R2URV and DR2URV are not.



The user of `RCURV` must consider both names `RCURV` and `R2URV` to be reserved; likewise, the user of `DRCURV` must consider both names `DRCURV` and `DR2URV` to be reserved. The names of *all* routines and named common blocks that are used by the `STAT/LIBRARY` and that do not have a numeral in the second position of the root name are listed in the Alphabetical Summary of Routines.

The careful user can avoid any conflicts with `IMSL` names if the following rules are observed:

- Do not choose a name that appears in the Alphabetical Summary of Routines in the *User's Manual*.
- Do not choose a name of three or more characters with a numeral in the second or third position.

These simplified rules include many combinations that are, in fact, allowable. However, if the user selects names that conform to these rules, no conflict will be encountered.

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## Deprecated and Renamed Routines

The routines in the following list are being deprecated in Version 2.0 of `STAT/LIBRARY`. A deprecated routine is one that is no longer used by anything in the library but is being included in the product for those users who may be currently referencing it in their application. However, any future versions of `STAT/LIBRARY` will not include these routines. If any of these routines are being called within an application, it is recommended that you change your code or retain the deprecated routine before replacing this library with the next version. Most of these routines were called by users only when they needed to set up their own workspace. Thus, the impact of these changes should be limited.

DHOUAP  
DHOUTR  
DG2DF  
DG2IN  
DG3DF  
G2DF  
G2IN  
G3DF  
SHOUAP  
SHOUTR

The following routines have been renamed due to naming conflicts with other software manufacturers.

`CTIME` — replaced with `CPSEC`

`DTIME` — replaced with `TIMDY`

`PAGE` — replaced with `PGOPT`

# Appendix A: GAMS Index

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## Description

This index lists routines in STAT/LIBRARY by a tree-structured classification scheme known as GAMS. Boisvert, Howe, Kahaner, and Springmann (1990) give the GAMS classification scheme. The classification scheme given here is Version 2.0.

The first level of the full classification scheme is denoted by a letter A thru Z as follows:

- A. Arithmetic, Error Analysis
- B. Number Theory
- C. Elementary and Special Functions
- D. Linear Algebra
- E. Interpolation
- F. Solution of Nonlinear Equations
- G. Optimization
- H. Differentiation and Integration
- I. Differential and Integral Equations
- J. Integral Transforms
- K. Approximation
- L. Statistics, Probability
- M. Simulation, Stochastic Modeling
- N. Data Handling
- O. Symbolic Computation
- P. Computational Geometry
- Q. Graphics
- R. Service Routines
- S. Software Development Tools
- Z. Other

There are seven levels in the classification scheme. Classes in the first level are identified by a capital letter as is given above. Classes in the remaining levels are identified by alternating letter-and-number combinations. A single letter (a–z) is used with the odd-numbered levels. A number (1–26) is used within the even-numbered levels.

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# IMSL STAT/LIBRARY

C ..... ELEMENTARY AND SPECIAL FUNCTIONS (*search also class L5*)

C3 ..... Polynomials

C3a..... Orthogonal

OPOLY Generate orthogonal polynomials with respect to  $x$  values and specified weights.

C7 ..... Gamma

C7e..... Incomplete gamma

CHIDF Evaluate the chi-squared distribution function.

CHIIN Evaluate the inverse of the chi-squared distribution function.

GAMDF Evaluate the gamma distribution function.

GAMIN Evaluate the inverse of the gamma distribution function.

C7f..... Incomplete gamma

BETDF Evaluate the beta probability distribution function.

BETIN Evaluate the inverse of the beta distribution function.

C8 ..... Error functions

C8a..... Error functions, their inverses, integrals, including the normal distribution function

ANORDF Evaluate the standard normal (Gaussian) distribution function.

ANORIN Evaluate the inverse of the standard normal (Gaussian) distribution function.

K ..... APPROXIMATION (*search also class L8*)

K1 ..... Least squares ( $L_2$ ) approximation

K1a ..... Linear least squares (*search also classes D5, D6, D9*)

K1a1 .... Unconstrained

RCOV Fit a multiple linear regression model given the variance-covariance matrix.

RGIVN Fit a multivariate linear regression model via fast Givens transformations.

RGLM Fit a multivariate general linear model.

RLSE Fit a multiple linear regression model using least squares.

K1a1a... Univariate data (curve fitting)

K1a1a2.. Polynomials

RCURV Fit a polynomial curve using least squares.

RFORP Fit an orthogonal polynomial regression model.

RPOLY Analyze a polynomial regression model.

K1a2 .... Constrained

K1a2a .. Linear constraints

RLEQU Fit a multivariate linear regression model with linear equality restrictions  $HB = G$  imposed on the regression parameters given results from IMSL routine RGIVN after  $IDO = 1$  and  $IDO = 2$  and prior to  $IDO = 3$ .

K1b ..... Nonlinear least squares

K1b1 .... Unconstrained

K1b1a .. Smooth functions

K1b1a1 User provides no derivatives

RNLIN Fit a nonlinear regression model.

K1b1a2 User provides first derivatives

RNLIN Fit a nonlinear regression model.

K2 ..... Minimax ( $L_\infty$ ) approximation

RLMV Fit a multiple linear regression model using the minimax criterion.

K3 ..... Least absolute value ( $L_1$ ) approximation

RLLP Fit a multiple linear regression model using the  $L_p$  norm criterion.

K4 ..... Other analytic approximations (e.g., Taylor polynomial, Pade)

RLLP Fit a multiple linear regression model using the  $L_p$  norm criterion.

L ..... STATISTICS, PROBABILITY

L1 ..... Data summarization

L1a ..... One-dimensional data

L1a1 ..... Raw data

EQTIL Compute empirical quantiles.

LETRR Produce a letter value summary.

ORDST Determine order statistics.

L1a1a... Location

UVSTA Compute basic univariate statistics.

L1a1b... Dispersion

UVSTA Compute basic univariate statistics.

L1a1c... Shape

UVSTA Compute basic univariate statistics.

L1a1e... Ties

NTIES Compute tie statistics for a sample of observations.

- L1a3..... Grouped data
  - GRPES Compute basic statistics from grouped data.
- L1c..... Multi-dimensional data
  - L1c1..... Raw data
    - CSTAT Compute cell frequencies, cell means, and cell sums of squares for multivariate data.
  - L1c1b... Covariance, correlation
    - CORVC Compute the variance-covariance or correlation matrix.
    - PCORR Compute partial correlations or covariances from the covariance or correlation matrix.
    - RBCOV Compute a robust estimate of a covariance matrix and mean vector.
- L2..... Data manipulation
  - L2a..... Transform (*search also classes L10a, N6, and N8*)
    - BCTR Perform a forward or an inverse Box-Cox (power) transformation.
    - GCSCP Generate centered variables, squares, and crossproducts.
    - OPOLY Generate orthogonal polynomials with respect to  $x$  values and specified weights.
    - RANKS Compute the ranks, normal scores, or exponential scores for a vector of observations.
  - L2b..... Tally
    - CSTAT Compute cell frequencies, cell means, and cell sums of squares for multivariate data.
    - FREQ Tally multivariate observations into a multi-way frequency table.
    - OWFRQ Tally observations into a one-way frequency table.
    - TWFRQ Tally observations into a two-way frequency table.
  - L2e..... Construct new variables (e.g., indicator variables)
    - GRGLM Generate regressors for a general linear model.
- L3..... Elementary statistical graphics (*search also class Q*)
  - L3a..... One-dimensional data
    - L3a1..... Histograms
      - HHSTP Print a horizontal histogram.
      - VHSTP Print a vertical histogram.
    - L3a2..... Frequency, cumulative frequency, percentile plots
      - CDFP Print a sample cumulative distribution function (CDF), a theoretical CDF, and confidence band information.
    - L3a3..... EDA graphics (e.g., box plots)
      - BOXP Print boxplots for one or more samples.
      - STMLP Print a stem-and-leaf plot.

L3a4..... Bar charts  
 HHSTP Print a horizontal histogram.  
 VHSTP Print a vertical histogram.

L3b ..... Two-dimensional data (*search also class L3e*)

L3b1 .... Histograms (superimposed and bivariate)  
 VHS2P Print a vertical histogram with every bar subdivided into two parts.

L3b2 .... Frequency, cumulative frequency  
 CDF2P Print a plot of two sample cumulative distribution functions.

L3e..... Multi-dimensional data

L3e3..... Scatter diagrams

L3e3a... Superimposed  $Y$  vs.  $X$   
 PLOTP Print a plot of up to ten sets of points.  
 SCTP Print a scatterplot of several groups of data.

L3e4..... EDA  
 BOXP Print boxplots for one or more samples.

L4 ..... Elementary data analysis

L4a..... One-dimensional data

L4a1 .... Raw data

L4a1a... Parametric analysis  
 CDFP Print a sample cumulative distribution function (CDF), a theoretical CDF, and confidence band information.

L4a1a2. Probability plots

L4a1a2e Exponential, extreme value  
 PROBP Print a probability plot.

L4a1a2h Halfnormal  
 PROBP Print a probability plot.

L4a1a2l Lambfa, logistic, lognormal  
 PROBP Print a probability plot.

L4a1a2n Negative binomial, normal  
 PROBP Print a probability plot.

L4a1a2w Weibull  
 PROBP Print a probability plot.

L4a1a4. Parameter estimates and tests

L4a1a4b Binomial  
 BINES Estimate the parameter  $p$  of the binomial distribution.

#### L4a1a4p Poisson

POIES Estimate the parameter of the Poisson distribution.

#### L4a1b... Nonparametric analysis

##### L4a1b1. Estimates and test regarding location (e.g., median), dispersion and shape

SIGNT Perform a sign test of the hypothesis that a given value is a specified quantile of a distribution.

SNRNK Perform a Wilcoxon signed rank test.

##### L4a1b2. Density function estimation

DESKN Perform nonparametric probability density function estimation by the kernel method.

DESPL Perform nonparametric probability density function estimation by the penalized likelihood method.

DESPT Estimate a probability density function at specified points using linear or cubic interpolation.

DNFFT Compute Gaussian kernel estimates of a univariate density via the fast Fourier transform over a fixed interval.

##### L4a1c ... Goodness-of-fit tests

CHIGF Perform a chi-squared goodness-of-fit test.

KSONE Perform a Kolmogorov-Smirnov one-sample test for continuous distributions.

LILLF Perform Lilliefors test for an exponential or normal distribution.

SPWLK Perform a Shapiro-Wilk  $W$ -test for normality.

##### L4ald.... Analysis of a sequence of numbers (*search also class L10a*)

DCUBE Perform a triplets test.

DSQAR Perform a D-square test.

NCTRD Perform the Noether test for cyclical trend.

PAIRS Perform a pairs test.

RUNS Perform a runs up test.

SDPLC Perform the Cox and Stuart sign test for trends in dispersion and location.

##### L4a3..... Grouped (and/or censored) data

GRPES Compute basic statistics from grouped data.

NRCES Compute maximum likelihood estimates of the mean and variance from grouped and/or censored normal data.

##### L4a4..... Data sampled from a finite population

SMPPR Compute statistics for inferences regarding the population proportion and total, given proportion data from a simple random sample.

SMPPS Compute statistics for inferences regarding the population proportion and total, given proportion data from a stratified random sample.

- SMPSC Compute statistics for inferences regarding the population mean and total using single-stage cluster sampling with continuous data.
  - SMPSR Compute statistics for inferences regarding the population mean and total, given data from a simple random sample.
  - SMPSS Compute statistics for inferences regarding the population mean and total, given data from a stratified random sample.
  - SMPST Compute statistics for inferences regarding the population mean and total, given continuous data from a two-stage sample with equisized primary units.
- L4b ..... Two dimensional data (*search also class L4c*)
- L4b1 .... Pairwise independent data
- L4b1a... Parametric analysis
- L4b1a4. Parameter estimates and hypothesis tests
- TWOMV Compute statistics for mean and variance inferences using samples from two normal populations.
- L4b1b .. Nonparametric analysis (e.g., tests based on ranks)
- CNCRD Calculate and test the significance of the Kendall coefficient of concordance.
  - INCLD Perform an inclusion test.
  - KENDL Compute and test Kendall's rank correlation coefficient.
  - RNKSM Perform the Wilcoxon rank sum test.
- L4b1c... Goodness-of-fit tests
- KSTWO Perform a Kolmogorov-Smirnov two-sample test.
- L4b4 .... Pairwise dependent grouped data
- CTRHO Estimate the bivariate normal correlation coefficient using a contingency table.
  - TETCC Categorize bivariate data and compute the tetrachoric correlation coefficient.
- L4b5 .... Data sampled from a finite population
- SMPRR Compute statistics for inferences regarding the population mean and total using ratio or regression estimation, or inferences regarding the population ratio, given a simple random sample.
  - SMPRS Compute statistics for inferences regarding the population mean and total using ratio or regression estimation, given continuous data from a stratified random sample.
- L4c..... Multi-dimensional data (*search also classes L4b and L7a1*)
- L4c1 ..... Independent data
- L4c1b... Nonparametric analysis
- BHAKV Perform a Bhapkar V test.



- KRSKL Perform a Kruskal-Wallis test for identical population medians.
  - KTRND Perform a  $k$ -sample trends test against ordered alternatives.
  - MVMMT Compute Mardia's multivariate measures of skewness and kurtosis and test for multivariate normality.
  - QTEST Perform a Cochran  $Q$  test for related observations.
- L4e..... Multiple multi-dimensional data sets
- MVIND Compute a test for the independence of  $k$  sets of multivariate normal variables.
- L5..... Function evaluation (*search also class C*)
- L5a..... Univariate
- L5a1 ..... Cumulative distribution functions, probability density functions
- L5a1b... Beta, binomial
- BETDF Evaluate the beta probability distribution function.
  - BINDF Evaluate the binomial distribution function.
  - BINPR Evaluate the binomial probability function.
- L5a1c ... Cauchy, chi-squared
- CHIDF Evaluate the chi-squared distribution function.
  - CSNDF Evaluate the noncentral chi-squared distribution function.
- L5a1f....  $F$  distribution
- FDF Evaluate the  $F$  distribution function.
- L5a1g... Gamma, general, geometric
- GAMDF Evaluate the gamma distribution function.
  - GCDF Evaluate a general continuous cumulative distribution function given ordinates of the density.
- L5a1h... Halfnormal, hypergeometric
- HYPDF Evaluate the hypergeometric distribution function.
  - HYPFR Evaluate the hypergeometric probability function.
- L5a1k... Kendall  $F$  statistic, Kolmogorov-Smirnov
- AKS1DF Evaluate the distribution function of the one-sided Kolmogorov-Smirnov goodness-of-fit  $D^+$  or  $D^-$  test statistic based on continuous data for one sample.
  - AKS2DF Evaluate the distribution function of the Kolmogorov-Smirnov goodness-of-fit  $D$  test statistic based on continuous data for two samples.
  - KENDP Compute the frequency distribution of the total score in Kendall's rank correlation coefficient.
- L5a1n... Negative binomial, normal
- ANORDF Evaluate the standard normal (Gaussian) distribution function.

L5a1p... Pareto, Poisson  
 POIDF Evaluate the Poisson distribution function.  
 POIPR Evaluate the Poisson probability function.

L5a1t.... *t* distribution  
 TDF Evaluate the Student's *t* distribution function.  
 TNDF Evaluate the noncentral Student's *t* distribution function.

L5a2..... Inverse cumulative distribution functions, sparsity functions

L5a2b... Beta, binomial  
 BETIN Evaluate the inverse of the beta distribution function.

L5a2c ... Cauchy, chi-squared  
 CHIIN Evaluate the inverse of the chi-squared distribution function.  
 CSNIN Evaluate the inverse of the noncentral chi-squared function.

L5a2f ... *F* distribution  
 FIN Evaluate the inverse of the *F* distribution function.

L5a2g... Gamma, general, geometric  
 GAMIN Evaluate the inverse of the gamma distribution function.  
 GCIN Evaluate the inverse of a general continuous cumulative distribution function given ordinates of the density.  
 GFNIN Evaluate the inverse of a general continuous cumulative distribution function given in a subprogram.

L5a2t.... *t* distribution  
 TIN Evaluate the inverse of the Student's *t* distribution function.  
 TNIN Evaluate the inverse of the noncentral Student's *t* distribution function.

L5b ..... Multivariate

L5b1 .... Cumulative distribution functions, probability density functions

L5b1n... Normal  
 BNRDF Evaluate the bivariate normal distribution function.

L6 ..... Random number generation

L6a..... Univariate

L6a2..... Beta, binomial, Boolean  
 RNBET Generate pseudorandom numbers from a beta distribution.  
 RNBIN Generate pseudorandom numbers from a binomial distribution.

L6a3..... Cauchy, chi-squared  
 RNCHI Generate pseudorandom numbers from a chi-squared distribution.

- RNCHY Generate pseudorandom numbers from a Cauchy distribution.
- L6a5..... Exponential, extreme value
- RNEXP Generate pseudorandom numbers from a standard exponential distribution.
- RNEXT Generate pseudorandom numbers from a mixture of two exponential distributions.
- L6a7..... Gamma, general (continuous, discrete), geometric
- RNGAM Generate pseudorandom numbers from a standard gamma distribution.
- RNGCS Set up table to generate pseudorandom numbers from a general continuous distribution.
- RNGCT Generate pseudorandom numbers from a general continuous distribution.
- RNGDA Generate pseudorandom numbers from a general discrete distribution using an alias method.
- RNGDS Set up table to generate pseudorandom numbers from a general discrete distribution.
- RNGDT Generate pseudorandom numbers from a general discrete distribution using a table lookup method.
- RNGEO Generate pseudorandom numbers from a geometric distribution.
- L6a8..... Halfnormal, hypergeometric
- RNHYP Generate pseudorandom numbers from a hypergeometric distribution.
- L6a12... Lambda, logistic, lognormal
- RNLGR Generate pseudorandom numbers from a logarithmic distribution.
- RNLNL Generate pseudorandom numbers from a lognormal distribution.
- L6a14... Negative binomial, normal, normal order statistics
- RNNBN Generate pseudorandom numbers from a negative binomial distribution.
- RNNOA Generate pseudorandom numbers from a standard normal distribution using an acceptance/rejection method.
- RNNOF Generate a pseudorandom number from a standard normal distribution.
- RNNOR Generate pseudorandom numbers from a standard normal distribution using an inverse CDF method.
- RNNOS Generate pseudorandom order statistics from a standard normal distribution.
- L6a16... Pareto, Pascal, permutations, Poisson
- RNNPP Generate pseudorandom numbers from a nonhomogeneous Poisson process.

- RNPER Generate a pseudorandom permutation.
- RNPOI Generate pseudorandom numbers from a Poisson distribution.
- L6a19... Samples, stable distribution
- RNSRI Generate a simple pseudorandom sample of indices.
- RNSRS Generate a simple pseudorandom sample from a finite population.
- RNSTA Generate pseudorandom numbers from a stable distribution.
- L6a20...  $t$  distribution, time series, triangular
- RNARM Generate a time series from a specified ARMA model.
- RNNPP Generate pseudorandom numbers from a nonhomogeneous Poisson process.
- RNSTT Generate pseudorandom numbers from a Student's  $t$  distribution.
- RNTRI Generate pseudorandom numbers from a triangular distribution on the interval (0,1).
- L6a21... Uniform (continuous, discrete), uniform order statistics
- RNUN Generate pseudorandom numbers from a uniform (0,1) distribution.
- RNUND Generate pseudorandom numbers from a discrete uniform distribution.
- RNUNF Generate a pseudorandom number from a uniform (0, 1) distribution.
- RNUNO Generate pseudorandom order statistics from a uniform (0, 1) distribution.
- L6a22... Von Mises
- RNVMS Generate pseudorandom numbers from a von Mises distribution.
- L6a23... Weibull
- RNWIB Generate pseudorandom numbers from a Weibull distribution.
- L6b ..... Multivariate
- RNDAT Generate pseudorandom numbers from a multivariate distribution determined from a given sample.
- L6b3 .... Contingency table, correlation matrix
- RNCOR Generate a pseudorandom orthogonal matrix or a correlation matrix.
- RNTAB Generate a pseudorandom two-way table.
- L6b13 .. Multinomial
- RNMTN Generate pseudorandom numbers from a multinomial distribution.

- L6b14... Normal
  - RNMVN Generate pseudorandom numbers from a multivariate normal distribution.
- L6b15... Orthogonal matrix
  - RNCOR Generate a pseudorandom orthogonal matrix or a correlation matrix.
- L6b21... Uniform
  - RNSPH Generate pseudorandom points on a unit circle or  $\kappa$ -dimensional sphere.
- L6c..... Service routines (e.g., seed)
  - RNGEF Retrieve the current value of the array used in the IMSL GFSR random number generator.
  - RNGES Retrieve the current value of the table in the IMSL random number generators that use shuffling.
  - RNGET Retrieve the current value of the seed used in the IMSL random number generators.
  - RNISD Determine a seed that yields a stream beginning 100,000 numbers beyond the beginning of the stream yielded by a given seed used in IMSL multiplicative congruential generators (with no shufflings).
  - RNOPG Retrieve the indicator of the type of uniform random number generator.
  - RNOPT Select the uniform (0, 1) multiplicative congruential pseudorandom number generator.
  - RNSEF Initialize the array used in the IMSL GFSR random number generator.
  - RNSES Initialize the table in the IMSL random number generators that use shuffling.
  - RNSET Initialize a random seed for use in the IMSL randomnumber generators.
- L7..... Analysis of variance (including analysis of covariance)
  - L7a..... One-way
    - L7a1 ..... Parametric
      - AONEC Analyze a one-way classification model with covariates.
      - AONEW Analyze a one-way classification model.
      - CTRST Compute contrast estimates and sums of squares.
      - SCIPM Compute simultaneous confidence intervals on all pairwise differences of means.
      - SNKMC Perform Student-Newman-Keuls multiple comparison test.
  - L7b..... Two-way (*search also class L7d*)
    - ATWOB Analyze a randomized block design or a two-way balanced design.

- FRDMN Perform Friedman's test for a randomized complete block design.
- MEDPL Compute a median polish of a two-way table.
- L7c..... Three-way (e.g., Latin squares) (*search also class L7d*)
  - ALATN Analyze a Latin square design.
- L7d ..... Multi-way
  - L7d1 .... Balanced complete data (e.g., factorial designs)
    - ABALD Analyze a balanced complete experimental design for a fixed, random, or mixed model.
    - ANEST Analyze a completely nested random model with possibly unequal numbers in the subgroups.
    - ANWAY Analyze a balanced  $n$ -way classification model with fixed effects.
    - CIDMS Compute a confidence interval on a variance component estimated as proportional to the difference in two mean squares in a balanced complete experimental design.
    - ROREX Reorder the responses from a balanced complete experimental design.
  - L7d2 .... Balanced incomplete data (e.g., fractional factorial designs)
    - ABIBD Analyze a balanced incomplete block design or a balanced lattice design.
  - L7d3 .... General linear models (unbalanced data)
    - ANEST Analyze a completely nested random model with possibly unequal numbers in the subgroups.
    - RGLM Fit a multivariate general linear model.
- L7e..... Multivariate
  - RGLM Fit a multivariate general linear model.
- L7f..... Generate experimental designs
  - RCOMP Generate an orthogonal central composite design.
- L8 ..... Regression (*search also classes D5, D6, D9, G, K*)
  - L8a..... Simple linear (e.g.,  $y = \beta_0 + \beta_1 x + \epsilon$ )
    - L8a1 .... Ordinary least squares
      - RONE Analyze a simple linear regression model.
    - L8a1a ... Parameter estimation
      - L8a1a1 . Unweighted data
        - RLINE Fit a line to a set of data points using least squares.
      - L8a1d... Inference (e.g., calibration) (*search also class L8a1a*)
        - RINCF Perform response control given a fitted simple linear regression model.
        - RINPF Perform inverse prediction given a fitted simple linear regression model.

- L8a2.....  $L_p$  for  $p$  different from 2 (e.g., least absolute value, minimax)
- RLAV Fit a multiple linear regression model using the least absolute values criterion.
  - RLLP Fit a multiple linear regression model using the  $L_p$  norm criterion.
  - RLMV Fit a multiple linear regression model using the minimax criterion.
- L8b..... Polynomial (e.g.,  $y = \beta_0 + \beta_1 x + \beta_2 x^2 + \epsilon$ ) (*search also class L8c*)
- L8b1..... Ordinary least squares
- L8b1a... Degree determination
- RFORP Fit an orthogonal polynomial regression model.
  - RPOLY Analyze a polynomial regression model.
- L8b1b... Parameter estimation
- L8b1b2. Using orthogonal polynomials
- RCURV Fit a polynomial curve using least squares.
  - RFORP Fit an orthogonal polynomial regression model.
  - RPOLY Analyze a polynomial regression model.
- L8b1c... Analysis (*search also class L8b1b*)
- RCASP Compute case statistics for a polynomial regression model given the fit based on orthogonal polynomials.
  - RPOLY Analyze a polynomial regression model.
  - RSTAP Compute summary statistics for a polynomial regression model given the fit based on orthogonal polynomials.
- L8b1d... Inference (*search also class L8b1b*)
- RCASP Compute case statistics for a polynomial regression model given the fit based on orthogonal polynomials.
  - RPOLY Analyze a polynomial regression model.
  - RSTAP Compute summary statistics for a polynomial regression model given the fit based on orthogonal polynomials.
- L8c..... Multiple linear (e.g.,  $y = \beta_0 + \beta_1 x_1 + \dots + \beta_k x_k + \epsilon$ )
- L8c1..... Ordinary least squares
- L8c1a... Variable selection
- L8c1a2. Using correlation or covariance data
- GSWEP Perform a generalized sweep of a row of a nonnegative definite matrix.
  - RBEST Select the best multiple linear regression models.
  - RSTEP Build multiple linear regression models using forward selection, backward selection, or stepwise selection.

L8c1b... Parameter estimation (*search also class L8c1a*)

L8c1b1. Using raw data

- RGIVN Fit a multivariate linear regression model via fast Givens transformations.
- RGLM Fit a multivariate general linear model.
- RLSE Fit a multiple linear regression model using least squares.

L8c1b2. Using correlation data

- RCOV Fit a multiple linear regression model given the variance-covariance matrix.

L8c1c... Analysis (*search also classes L8c1a and L8c1b*)

- RCASE Compute case statistics and diagnostics given data points, coefficient estimates  $\hat{\beta}$ , and the  $R$  matrix for a fitted general linear model.
- RCOVB Compute the estimated variance-covariance matrix of the estimated regression coefficients given the  $R$  matrix.
- RLOFE Compute a lack-of-fit test based on exact replicates for a fitted regression model.
- RLOFN Compute a lack-of-fit test based on near replicates for a fitted regression model.
- ROTIN Compute diagnostics for detection of outliers and influential data points given residuals and the  $R$  matrix for a fitted general linear model.
- RSTAT Compute statistics related to a regression fit given the coefficient estimates  $\hat{\beta}$  and the  $R$  matrix.

L8c1d... Inference (*search also classes L8c1a and L8c1b*)

- CESTI Construct an equivalent completely testable multivariate general linear hypothesis  $HBV = G$  from a partially testable hypothesis  $H_pBV = G_p$ .
- RCASE Compute case statistics and diagnostics given data points, coefficient estimates  $\hat{\beta}$ , and the  $R$  matrix for a fitted general linear model.
- RHPSS Compute the matrix of sums of squares and crossproducts for the multivariate general linear hypothesis  $HBV = G$  given the coefficient estimates  $\hat{B}$  and the  $R$  matrix.
- RHPTE Perform tests for a multivariate general linear hypothesis  $HBV = G$  given the hypothesis sums of squares and crossproducts matrix  $S_H$  and the error sums of squares and crossproducts matrix  $S_E$ .
- RSTAT Compute statistics related to a regression fit given the coefficient estimates  $\hat{\beta}$  and the  $R$  matrix.

L8c3.....  $L_p$  for  $p$  different from 2

- RLAV Fit a multiple linear regression model using the least absolute values criterion.



- RLLP Fit a multiple linear regression model using the  $L_p$  norm criterion.
- RLMV Fit a multiple linear regression model using the minimax criterion.
- L8d..... Polynomial in several variables
  - RCOMP Generate an orthogonal central composite design.
  - TCSCP Transform coefficients from a quadratic regression model generated from squares and crossproducts of centered variables to a model using uncentered variables.
- L8e..... Nonlinear (i.e.,  $y = f(X; \theta) + \epsilon$ )
- L8e1..... Ordinary least squares
- L8e1b... Parameter estimation
  - RNLIN Fit a nonlinear regression model.
- L8f..... Simultaneous (i.e.,  $Y = XB + \epsilon$ )
  - RCOV Fit a multiple linear regression model given the variance-covariance matrix.
  - RGIVN Fit a multivariate linear regression model via fast Givens transformations.
  - RGLM Fit a multivariate general linear model.
  - RHPSS Compute the matrix of sums of squares and crossproducts for the multivariate general linear hypothesis  $HBU = G$  given the coefficient estimates  $\hat{B}$  and the  $R$  matrix.
  - RHPTF Perform tests for a multivariate general linear hypothesis  $HBU = G$  given the hypothesis sums of squares and crossproducts matrix  $S_H$  and the error sums of squares and crossproducts matrix  $S_E$ .
  - RLEQU Fit a multivariate linear regression model with linear equality restrictions  $HB = G$  imposed on the regression parameters given results from IMSL routine RGIVN after  $IDO = 1$  and  $IDO = 2$  and prior to  $IDO = 3$ .
- L8i..... Service routines (e.g., matrix manipulation for variable selection)
  - GCLAS Get the unique values of each classification variable.
  - GCSCP Generate centered variables, squares, and crossproducts.
  - GRGLM Generate regressors for a general linear model.
  - RORDM Reorder rows and columns of a symmetric matrix.
  - RSUBM Retrieve a symmetric submatrix from a symmetric matrix.
- L9..... Categorical data analysis
  - CTGLM Analyze categorical data using logistic, Probit, Poisson, and other generalized linear models.
  - CTRAN Perform generalized Mantel-Haenszel tests in a stratified contingency table.
- L9a..... 2-by-2 tables

- CTTWO Perform a chi-squared analysis of a 2 by 2 contingency table.
- L9b ..... Two-way tables (*search also class L9d*)
- CTCHI Perform a chi-squared analysis of a two-way contingency table.
- CTEPR Compute Fisher's exact test probability and a hybrid approximation to the Fisher exact test probability for a contingency table using the network algorithm.
- CTPRB Compute exact probabilities in a two-way contingency table.
- CTRHO Estimate the bivariate normal correlation coefficient using a contingency table.
- CTWLS Perform a generalized linear least squares analysis of transformed probabilities in a two-dimensional contingency table.
- MEDPL Compute a median polish of a two-way table.
- TWFRQ Tally observations into a two-way frequency table.
- L9c..... Log-linear model
- CTASC Compute partial association statistics for log-linear models in a multidimensional contingency table.
- CTLLN Compute model estimates and associated statistics for a hierarchical log-linear model.
- CTPAR Compute model estimates and covariances in a fitted log-linear model.
- CTSTP Build hierarchical log-linear models using forward selection, backward selection, or stepwise selection.
- PRPFT Perform iterative proportional fitting of a contingency table using a loglinear model.
- L9d ..... EDA (e.g., median polish)
- MEDPL Compute a median polish of a two-way table.
- L10 ..... Time series analysis (*search also class J*)
- L10a..... Univariate
- L10a1... Transformations
- L10a1b. Stationarity (*search also class L8a1*)
- BCTR Perform a forward or an inverse Box-Cox (power) transformation.
- L10a1c. Filters
- L10a1c1 Difference (nonseasonal and seasonal)
- DIFF Difference a time series.
- L10a2... Time domain analysis
- L10a2a . Summary statistics

- L10a2a1 Autocovariances and autocorrelations
  - ACF Compute the sample autocorrelation function of a stationary time series.
  - LOFCF Perform lack-of-fit test for a univariate time series or transfer function given the appropriate correlation function.
- L10a2a2 Partial autocorrelations
  - PACF Compute the sample partial autocorrelation function of a stationary time series.
- L10a2c . Autoregressive models
  - SPWF Compute the Wiener forecast operator for a stationary stochastic process.
- L10a2d . ARMA and ARIMA models (including Box-Jenkins methods)
  - L10a2d2 Parameter estimation
    - ARMME Compute method of moments estimates of the autoregressive parameters of an ARMA model.
    - MAMME Compute method of moments estimates of the moving average parameters of an ARMA model.
    - NSLSE Compute least squares estimates of parameters for a nonseasonal ARMA model.
    - NSPE Compute preliminary estimates of the autoregressive and moving average parameters of an ARMA model.
  - L10a2d3 Forecasting
    - NSBJF Compute Box-Jenkins forecasts and their associated probability limits for a nonseasonal ARMA model.
- L10a2e . State-space analysis (e.g., Kalman filtering)
  - KALMN Perform Kalman filtering and evaluate the likelihood function for the state-space model.
- L10a3 ... Frequency domain analysis (*search also class J1*)
  - L10a3a . Spectral Analysis
    - L10a3a2 Periodogram analysis
      - PFFT Compute the periodogram of a stationary time series using a fast Fourier transform.
    - L10a3a3 Spectrum estimation using the periodogram
      - SSWD Estimate the nonnormalized spectral density of a stationary time series using a spectral window given the time series data.
      - SSWP Estimate the nonnormalized spectral density of a stationary time series using a spectral window given the periodogram.

- SWED Estimation of the nonnormalized spectral density of a stationary time series based on specified periodogram weights given the time series data.
  - SWEP Estimation of the nonnormalized spectral density of a stationary time series based on specified periodogram weights given the periodogram.
- L10a3a6 Spectral windows
- DIRIC Compute the Dirichlet kernel.
  - FEJER Compute the Fejér kernel.
- L10b .... Two time series (*search also classes L10c and L10d*)
- L10b2 .. Time domain analysis
- L10b2a. Summary statistics (e.g., cross-correlations)
- CCF Compute the sample cross-correlation function of two stationary time series.
- L10b2b Transfer function models
- IRNSE Compute estimates of the impulse response weights and noise series of a univariate transfer function model.
  - TFPE Compute preliminary estimates of parameters for a univariate transfer function model.
- L10b3 .. Frequency domain analysis (*search also class J1*)
- L10b3a. Cross-spectral analysis
- L10b3a3 Cross-spectrum estimation using the cross-periodogram
- CSSWD Estimate the nonnormalized cross-spectral density of two stationary time series using a spectral window given the time series data.
  - CSSWP Estimate the nonnormalized cross-spectral density of two stationary time series using a spectral window given the spectral densities and cross periodogram.
  - CSWED Estimate the nonnormalized cross-spectral density of two stationary time series using a weighted cross periodogram given the time series data.
  - CSWEP Estimate the nonnormalized cross-spectral density of two stationary time series using a weighted cross periodogram given the spectral densities and cross periodogram.
- L10c..... Multivariate time series (*search also classes J1, L3e3 and L10b*)
- KALMN Perform Kalman filtering and evaluate the likelihood function for the state-space model.
- L10d .... Two multi-channel time series
- MCCF Compute the multichannel cross-correlation function of two mutually stationary multichannel time series.

- MLSE Compute least squares estimates of a linear regression model for a multichannel time series with a specified base channel.
- MWFE Compute least squares estimates of the multichannel Wiener filter coefficients for two mutually stationary multichannel time series.
- L11..... Correlation analysis (*search also classes L4 and L13c*)
  - BSCAT Compute the biserial correlation coefficient for a dichotomous variable and a classification variable.
  - BSPBS Compute the biserial and point-biserial correlation coefficients for a dichotomous variable and a numerically measurable classification variable.
  - CORVC Compute the variance-covariance or correlation matrix.
  - COVPL Compute a pooled variance-covariance matrix from the observations.
  - CTRHO Estimate the bivariate normal correlation coefficient using a contingency table.
  - KENDP Compute the frequency distribution of the total score in Kendall's rank correlation coefficient.
  - PCORR Compute partial correlations or covariances from the covariance or correlation matrix.
  - RBCOV Compute a robust estimate of a covariance matrix and mean vector.
  - TETCC Categorize bivariate data and compute the tetrachoric correlation coefficient.
- L12..... Discriminant analysis
  - DMSCR Use Fisher's linear discriminant analysis method to reduce the number of variables.
  - DSCRM Perform a linear or a quadratic discriminant function analysis among several known groups.
  - NNBRD Perform a  $k$  nearest neighbor discrimination.
- L13..... Covariance structures models
  - L13a..... Factor analysis
    - FACTR Extract initial factor-loading estimates in factor analysis.
    - FCOEF Compute a matrix of factor score coefficients for input to the following IMSL routine (FSCOR).
    - FDOBL Compute a direct oblimin rotation of a factor-loading matrix.
    - FGCRF Compute direct oblique rotation according to a generalized fourth-degree polynomial criterion.
    - FHARR Compute an oblique rotation of an unrotated factor-loading matrix using the Harris-Kaiser method.
    - FIMAG Compute the image transformation matrix.
    - FOPCS Compute an orthogonal Procrustes rotation of a factor-loading matrix using a target matrix.

- FPRMX Compute an oblique Promax or Procrustes rotation of a factor-loading matrix using a target matrix, including pivot and power vector options.
  - FRESI Compute commonalities and the standardized factor residual correlation matrix.
  - FROTA Compute an orthogonal rotation of a factor-loading matrix using a generalized orthomax criterion, including quartimax, varimax, and equamax rotations.
  - FRVAR Compute the factor structures and the variance explained by each factor.
  - FSCOR Compute a set of factor scores given the factor score coefficient matrix.
- L13b .... Principal components analysis
- KPRIN Maximum likelihood or least-squares estimates for principle components from one or more matrices.
  - PRINC Compute principal components from a variance-covariance matrix or a correlation matrix.
- L13c..... Canonical correlation
- CANCR Perform canonical correlation analysis from a data matrix.
  - CANVC Perform canonical correlation analysis from a variance-covariance matrix or a correlation matrix.
- L14 ..... Cluster analysis
- L14a..... One-way
- L14a1... Unconstrained
- L14a1a. Nested
- L14a1a1 Joining (e.g., single link)
- CLINK Perform a hierarchical cluster analysis given a distance matrix.
- L14a1b. Non-nested (e.g.,  $K$  means)
- KMEAN Perform a  $K$ -means (centroid) cluster analysis.
- L14c..... Display
- TREEP Print a binary tree.
- L14d .... Service routines (e.g., compute distance matrix)
- CDIST Compute a matrix of dissimilarities (or similarities) between the columns (or rows) of a matrix.
  - CNUMB Compute cluster membership for a hierarchical cluster tree.
- L15 ..... Life testing, survival analysis
- ACTBL Produce population and cohort life tables.
  - HAZEZ Perform nonparametric hazard rate estimation using kernel functions. Easy-to-use version of the previous IMSL subroutine (HAZRD).

HAZRD Perform nonparametric hazard rate estimation using kernel functions and quasi-likelihoods.

HAZST Perform hazard rate estimation over a grid of points using a kernel function.

KAPMR Compute Kaplan-Meier estimates of survival probabilities in stratified samples.

KTBLE Print Kaplan-Meier estimates of survival probabilities in stratified samples.

NRCES Compute maximum likelihood estimates of the mean and variance from grouped and/or censored normal data.

PHGLM Analyze time event data via the proportional hazards model.

STBLE Estimate survival probabilities and hazard rates for various parametric models.

SVGLM Analyze censored survival data using a generalized linear model.

TRNBL Compute Turnbull's generalized Kaplan-Meier estimates of survival probabilities in samples with interval censoring.

L16..... Multidimensional scaling

MSDBL Obtain normalized product-moment (double centered) matrices from dissimilarity matrices.

MSDST Compute distances in a multidimensional scaling model.

MSIDV Perform individual-differences multidimensional scaling for metric data using alternating least squares.

MSINI Compute initial estimates in multidimensional scaling models.

MSSTN Transform dissimilarity/similarity matrices and replace missing values by estimates to obtain standardized dissimilarity matrices.

MSTRS Compute various stress criteria in multidimensional scaling.

L17..... Statistical data sets

GDATA Retrieve a commonly analyzed data set.

N ..... DATA HANDLING (*search also class L2*)

N1 ..... Input, output

PGOPT Set or retrieve page width and length for printing.

WRIRL Print an integer rectangular matrix with a given format and labels.

WRIRN Print an integer rectangular matrix with integer row and column labels.

WROPT Set or retrieve an option for printing a matrix.

WRRRL Print a real rectangular matrix with a given format and labels.

- WRRRN Print a real rectangular matrix with integer row and column labels.
- N3..... Character manipulation
- ACHAR Return a character given its ASCII value.
  - CVTSI Convert a character string containing an integer number into the corresponding integer form.
  - IACHAR Return the integer ASCII value of a character argument.
  - ICASE Return the ASCII value of a character converted to uppercase.
  - IICSR Compare two character strings using the ASCII collating sequence without regard to case.
  - IIDEX Determine the position in a string at which a given character sequence begins without regard to case.
- N5..... Searching
- N5a ..... Extreme value
- EQTIL Compute empirical quantiles.
  - ORDST Determine order statistics.
- N5b ..... Insertion position
- ISRCH Search a sorted integer vector for a given integer and return its index.
  - SRCH Search a sorted vector for a given scalar and return its index.
  - SSRCH Search a character vector, sorted in ascending ASCII order, for a given string and return its index.
- N5c ..... On a key
- IIDEX Determine the position in a string at which a given character sequence begins without regard to case.
  - ISRCH Search a sorted integer vector for a given integer and return its index.
  - SRCH Search a sorted vector for a given scalar and return its index.
  - SSRCH Search a character vector, sorted in ascending ASCII order, for a given string and return its index.
- N6..... Sorting
- N6a ..... Internal
- N6a1 .... Passive (i.e., construct pointer array, rank)
- N6a1a .. Integer
- SVIGP Sort an integer array by algebraic value and return the permutations.
- N6a1b .. Real
- RANKS Compute the ranks, normal scores, or exponential scores for a vector of observations.



- SCOLR Sort columns of a real rectangular matrix using keys in rows.
- SROWR Sort rows of a real rectangular matrix using keys in columns.
- SVRGP Sort a real array by algebraic value and return the permutations.
- N6a2 .... Active
- N6a2a... Integer
- SVIGN Sort an integer array by algebraic value.
- SVIGP Sort an integer array by algebraically increasing value and return the permutation that rearranges the array.
- N6a2b .. Real
- SCOLR Sort columns of a real rectangular matrix using keys in rows.
- SROWR Sort rows of a real rectangular matrix using keys in columns.
- SVRGN Sort a real array by algebraic value.
- SVRGP Sort a real array by algebraic value and return the permutations.
- N8 ..... Permuting
- MVNAN Move any rows of a matrix with the IMSL missing value code NaN (not a number) in the specified columns to the last rows of the matrix.
- PERMA Permute the rows or columns of a matrix.
- PERMU Rearrange the elements of an array as specified by a permutation.
- RORDM Reorder rows and columns of a symmetric matrix.
- Q ..... GRAPHICS *(search also classes L3)*
- BOXP Print boxplots for one or more samples.
- CDF2P Print a plot of two sample cumulative distribution functions.
- CDFP Print a sample cumulative distribution function (CDF), a theoretical CDF, and confidence band information.
- HHSTP Print a horizontal histogram.
- PLOTP Print a plot of up to ten sets of points.
- PROBP Print a probability plot.
- SCTP Print a scatterplot of several groups of data.
- STMLP Print a stem-and-leaf plot.
- TREEP Print a binary tree.
- VHS2P Print a vertical histogram with every bar subdivided into two parts.
- VHSTP Print a vertical histogram.
- R ..... SERVICE ROUTINES
- IDYWK Compute the day of the week for a given date.

NDAYS Compute the number of days from January 1, 1900, to the given date.  
 NDYIN Give the date corresponding to the number of days since January 1, 1900.  
 TDATE Get today's date.  
 TIMDY Get time of day.  
 VERSL Obtain STAT/LIBRARY-related version, system and license numbers.

R1 ..... Machine-dependent constants

AMACH Retrieve machine constants.  
 IFNAN Check if a floating-point number is NaN (not a number).  
 IMACH Retrieve integer machine constants.  
 UMACH Set or retrieve input or output device unit numbers.

R3 ..... Error handling

R3b ..... Set unit number for error messages

UMACH Set or retrieve input or output device unit numbers.

R3c ..... Other utilities

ERSET Set error handler default print and stop actions.  
 IERCD Retrieve the code for an informational error.  
 N1RTY Retrieve an error type for the most recently called IMSL routine.

S..... SOFTWARE DEVELOPMENT TOOLS

CPSEC Return CPU time used in seconds.

# Appendix B: Alphabetical Summary of Routines

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## IMSL STAT/LIBRARY

ABALD	396	Analyze a balanced complete experimental design for a fixed, random, or mixed model.
ABIBD	380	Analyze a balanced incomplete block design or a balanced lattice design.
ACF	637	Compute the sample autocorrelation function of a stationary time series.
ACHAR	1289	Return a character given its ASCII value.
ACTBL	992	Produce population and cohort life tables.
AKS1DF	1117	Evaluate the distribution function of the one-sided Kolmogorov-Smirnov goodness-of-fit $D^+$ or $D^-$ test statistic based on continuous data for one sample.
AKS2DF	1120	Evaluate the distribution function of the Kolmogorov-Smirnov goodness-of-fit $D$ test statistic based on continuous data for two samples.
ALATN	386	Analyze a Latin square design.
AMACH	1336	Retrieve machine constants.
AMILLR	1315	Evaluate Mill's ratio (the ratio of the ordinate to the upper tail area of the standardized normal distribution).
ANEST	409	Analyze a completely nested random model with possibly unequal numbers in the subgroups.
ANORDF	1122	Evaluate the standard normal (Gaussian) distribution function.
ANORIN	1124	Evaluate the inverse of the standard normal (Gaussian) distribution function.

ANWAY	390	Analyze a balanced $n$ -way classification model with fixed effects.
AONEC	364	Analyze a one-way classification model with covariates.
AONEW	362	Analyze a one-way classification model.
ARMME	657	Compute method of moments estimates of the autoregressive parameters of an ARMA model.
ATWOB	375	Analyze a randomized block design or a two-way balanced design.
BCTR	629	Perform a forward or an inverse Box-Cox (power) transformation.
BETDF	1125	Evaluate the beta probability distribution function.
BETIN	1127	Evaluate the inverse of the beta distribution function.
BHAKV	566	Perform a Bhapkar $V$ test.
BINDF	1108	Evaluate the binomial distribution function.
BINES	44	Estimate the parameter $p$ of the binomial distribution.
BINPR	1110	Evaluate the binomial probability function.
BNRDF	1128	Evaluate the bivariate normal distribution function.
BOXP	1083	Print boxplots for one or more samples.
BSCAT	348	Compute the biserial correlation coefficient for a dichotomous variable and a classification variable.
BSPBS	346	Compute the biserial and point-biserial correlation coefficients for a dichotomous variable and a numerically measurable classification variable.
CANCR	844	Perform canonical correlation analysis from a data matrix.
CANVC	857	Perform canonical correlation analysis from a variance-covariance matrix or a correlation matrix.
CCF	644	Compute the sample cross-correlation function of two stationary time series.
CDF2P	1090	Print a plot of two sample cumulative distribution functions.
CDFP	1087	Print a sample cumulative distribution function (CDF), a theoretical CDF, and confidence band information.
CDIST	889	Compute a matrix of dissimilarities (or similarities) between the columns (or rows) of a matrix.
CESTI	157	Construct an equivalent completely testable multivariate general linear hypothesis $HBU = G$ from a partially testable hypothesis $H_pBU = G_p$ .

CHFAC	1308	Compute an upper triangular factorization of a real symmetric nonnegative definite matrix.
CHIDF	1129	Evaluate the chi-squared distribution function.
CHIGF	584	Perform a chi-squared goodness-of-fit test.
CHIIIN	1132	Evaluate the inverse of the chi-squared distribution function.
CIDMS	426	Compute a confidence interval on a variance component estimated as proportional to the difference in two mean squares in a balanced complete experimental design.
CLINK	892	Perform a hierarchical cluster analysis given a distance matrix.
CNCRD	350	Calculate and test the significance of the Kendall coefficient of concordance.
CNUMB	897	Compute cluster membership for a hierarchical cluster tree.
CORVC	314	Compute the variance-covariance or correlation matrix.
COVPL	322	Compute a pooled variance-covariance matrix from the observations.
CPFFT	750	Compute the cross periodogram of two stationary time series using a fast Fourier transform.
CPSEC	1295	Return CPU time used in seconds.
CSNDF	1133	Evaluate the noncentral chi-squared distribution function.
CSNIN	1136	Evaluate the inverse of the noncentral chi-squared function.
CSSWD	757	Estimate the nonnormalized cross-spectral density of two stationary time series using a spectral window given the time series data.
CSSWP	767	Estimate the nonnormalized cross-spectral density of two stationary time series using a spectral window given the spectral densities and cross periodogram.
CSTAT	54	Compute cell frequencies, cell means, and cell sums of squares for multivariate data.
CSWED	773	Estimate the nonnormalized cross-spectral density of two stationary time series using a weighted cross periodogram given the time series data.
CSWEP	782	Estimate the nonnormalized cross-spectral density of two stationary time series using a weighted cross periodogram given the spectral densities and cross periodogram.
CTASC	482	Compute partial association statistics for log-linear models in a multidimensional contingency table.

CTCHI	446	Perform a chi-squared analysis of a two-way contingency table.
CTEPR	459	Compute Fisher's exact test probability and a hybrid approximation to the Fisher exact test probability for a contingency table using the network algorithm.
CTGLM	510	Analyze categorical data using logistic, Probit, Poisson, and other generalized linear models.
CTLLN	467	Compute model estimates and associated statistics for a hierarchical log-linear model.
CTPAR	476	Compute model estimates and covariances in a fitted log-linear model.
CTPRB	456	Compute exact probabilities in a two-way contingency table.
CTRAN	502	Perform generalized Mantel-Haenszel tests in a stratified contingency table.
CTRHO	339	Estimate the bivariate normal correlation coefficient using a contingency table.
CTRST	417	Compute contrast estimates and sums of squares.
CTSTP	489	Build hierarchical log-linear models using forward selection, backward selection, or stepwise selection.
CTTWO	436	Perform a chi-squared analysis of a 2 by 2 contingency table.
CTWLS	526	Perform a generalized linear least squares analysis of transformed probabilities in a two-dimensional contingency table.
CVTSI	1294	Convert a character string containing an integer number into the corresponding integer form.
DCUBE	609	Perform a triplets test.
DESKN	1044	Perform nonparametric probability density function estimation by the kernel method.
DESPL	1040	Perform nonparametric probability density function estimation by the penalized likelihood method.
DESPT	1052	Estimate a probability density function at specified points using linear or cubic interpolation.
DIFF	633	Difference a time series.
DIRIC	719	Compute the Dirichlet kernel.
DMSCR	876	Use Fisher's linear discriminant analysis method to reduce the number of variables.

DNFFT	1047	Compute Gaussian kernel estimates of a univariate density via the fast Fourier transform over a fixed interval.
DSCRM	863	Perform a linear or a quadratic discriminant function analysis among several known groups.
DSQAR	607	Perform a D-square test.
ENOS	1314	Evaluate the expected value of a normal order statistic.
EQTIL	35	Compute empirical quantiles.
ERSET	1327	Set error handler default print and stop actions.
FACTR	801	Extract initial factor-loading estimates in factor analysis.
FCOEF	833	Compute a matrix of factor score coefficients for input to the following IMSL routine (FSCOR).
FDF	1137	Evaluate the $F$ distribution function.
FDOBL	815	Compute a direct oblimin rotation of a factor-loading matrix.
FEJER	721	Compute the Fejér kernel.
FGCRF	825	Compute direct oblique rotation according to a generalized fourth-degree polynomial criterion.
FHARR	822	Compute an oblique rotation of an unrotated factor-loading matrix using the Harris-Kaiser method.
FIMAG	829	Compute the image transformation matrix.
FIN	1139	Evaluate the inverse of the $F$ distribution function.
FOPCS	812	Compute an orthogonal Procrustes rotation of a factor-loading matrix using a target matrix.
FPRMX	818	Compute an oblique Promax or Procrustes rotation of a factor-loading matrix using a target matrix, including pivot and power vector options.
FRDMN	568	Perform Friedman's test for a randomized complete block design.
FREQ	13	Tally multivariate observations into a multi-way frequency table.
FRESI	840	Compute commonalities and the standardized factor residual correlation matrix.
FROTA	809	Compute an orthogonal rotation of a factor-loading matrix using a generalized orthomax criterion, including quartimax, varimax, and equamax rotations.
FRVAR	831	Compute the factor structures and the variance explained by each factor.

FSCOR	838	Compute a set of factor scores given the factor score coefficient matrix.
GAMDF	1140	Evaluate the gamma distribution function.
GAMIN	1142	Evaluate the inverse of the gamma distribution function.
GCDF	1150	Evaluate a general continuous cumulative distribution function given ordinates of the density.
GCIN	1152	Evaluate the inverse of a general continuous cumulative distribution function given ordinates of the density.
GCLAS	207	Get the unique values of each classification variable.
GCSCP	272	Generate centered variables, squares, and crossproducts.
GDATA	1302	Retrieve a commonly analyzed data set.
GFNIN	1155	Evaluate the inverse of a general continuous cumulative distribution function given in a subprogram.
GIRTS	1305	Solve a triangular (possibly singular) set of linear systems and/or compute a generalized inverse of an upper triangular matrix.
GRGLM	210	Generate regressors for a general linear model.
GRPES	51	Compute basic statistics from grouped data.
GSWEP	230	Perform a generalized sweep of a row of a nonnegative definite matrix.
HAZEZ	1061	Perform nonparametric hazard rate estimation using kernel functions. Easy-to-use version of the previous IMSL subroutine (HAZRD).
HAZRD	1054	Perform nonparametric hazard rate estimation using kernel functions and quasi-likelihoods.
HAZST	1069	Perform hazard rate estimation over a grid of points using a kernel function.
HHSTP	1078	Print a horizontal histogram.
HYPDF	1111	Evaluate the hypergeometric distribution function.
HYPPR	1113	Evaluate the hypergeometric probability function.
IACHAR	1290	Return the integer ASCII value of a character argument.
ICASE	1291	Return the ASCII value of a character converted to uppercase.
IDYWK	1300	Compute the day of the week for a given date.
IERCD	1328	Retrieve the code for an informational error.
IFNAN	1337	Check if a floating-point number is NaN (not a number).



IICSR	1292	Compare two character strings using the ASCII collating sequence without regard to case.
IIDEX	1293	Determine the position in a string at which a given character sequence begins without regard to case.
IMACH	1335	Retrieve integer machine constants.
INCLD	561	Perform an inclusion test.
IRNSE	685	Compute estimates of the impulse response weights and noise series of a univariate transfer function model.
ISRCH	1286	Search a sorted integer vector for a given integer and return its index.
KALMN	705	Perform Kalman filtering and evaluate the likelihood function for the state-space model.
KAPMR	938	Compute Kaplan-Meier estimates of survival probabilities in stratified samples.
KENDL	353	Compute and test Kendall's rank correlation coefficient.
KENDP	357	Compute the frequency distribution of the total score in Kendall's rank correlation coefficient.
KMEAN	900	Perform a <i>K</i> -means (centroid) cluster analysis.
KPRIN	797	Maximum likelihood or least-squares estimates for principle components from one or more matrices.
KRSKL	564	Perform a Kruskal-Wallis test for identical population medians.
KSONE	580	Perform a Kolmogorov-Smirnov one-sample test for continuous distributions.
KSTWO	598	Perform a Kolmogorov-Smirnov two-sample test.
KTBLE	942	Print Kaplan-Meier estimates of survival probabilities in stratified samples.
KTRND	574	Perform a <i>k</i> -sample trends test against ordered alternatives.
LETTR	29	Produce a letter value summary.
LILLF	591	Perform Lilliefors test for an exponential or normal distribution.
LOFCF	716	Perform lack-of-fit test for a univariate time series or transfer function given the appropriate correlation function.
MAMME	660	Compute method of moments estimates of the moving average parameters of an ARMA model.
MCCF	649	Compute the multichannel cross-correlation function of two mutually stationary multichannel time series.

MCHOL	1311	Compute an upper triangular factorization of a real symmetric matrix $A$ plus a diagonal matrix $D$ , where $D$ is determined sequentially during the Cholesky factorization in order to make $A + D$ nonnegative definite.
MEDPL	59	Compute a median polish of a two-way table.
MLSE	694	Compute least squares estimates of a linear regression model for a multichannel time series with a specified base channel.
MSDBL	1024	Obtain normalized product-moment (double centered) matrices from dissimilarity matrices.
MSDST	1017	Compute distances in a multidimensional scaling model.
MSIDV	1003	Perform individual-differences multidimensional scaling for metric data using alternating least squares.
MSINI	1028	Compute initial estimates in multidimensional scaling models.
MSSTN	1020	Transform dissimilarity/similarity matrices and replace missing values by estimates to obtain standardized dissimilarity matrices.
MSTRS	1035	Compute various stress criteria in multidimensional scaling.
MVIND	842	Compute a test for the independence of $k$ sets of multivariate normal variables.
MVMMT	594	Compute Mardia's multivariate measures of skewness and kurtosis and test for multivariate normality.
MVNAN	1269	Move any rows of a matrix with the IMSL missing value code NaN (not a number) in the specified columns to the last rows of the matrix.
MWFE	700	Compute least squares estimates of the multichannel Wiener filter coefficients for two mutually stationary multichannel time series.
N1RTY	1328	Retrieve an error type for the most recently called IMSL routine.
NCTRD	548	Perform the Noether test for cyclical trend.
NDAYS	1297	Compute the number of days from January 1, 1900, to the given date.
NDYIN	1299	Give the date corresponding to the number of days since January 1, 1900.
NGHBR	1320	Search a $k$ - $d$ tree for the $k$ nearest neighbors of a key.
NNBRD	880	Perform a $k$ nearest neighbor discrimination.

NRCES	48	Compute maximum likelihood estimates of the mean and variance from grouped and/or censored normal data.
NSBJF	680	Compute Box-Jenkins forecasts and their associated probability limits for a nonseasonal ARMA model.
NLSLE	669	Compute least squares estimates of parameters for a nonseasonal ARMA model.
NSPE	664	Compute preliminary estimates of the autoregressive and moving average parameters of an ARMA model.
NTIES	555	Compute tie statistics for a sample of observations.
OPOLY	269	Generate orthogonal polynomials with respect to $x$ values and specified weights.
ORDST	31	Determine order statistics.
OWFRQ	3	Tally observations into a one-way frequency table.
PACF	641	Compute the sample partial autocorrelation function of a stationary time series.
PAIRS	604	Perform a pairs test.
PCORR	327	Compute partial correlations or covariances from the covariance or correlation matrix.
PERMA	1266	Permute the rows or columns of a matrix.
PERMU	1265	Rearrange the elements of an array as specified by a permutation.
PFFT	723	Compute the periodogram of a stationary time series using a fast Fourier transform.
PGOPT	1263	Set or retrieve page width and length for printing.
PHGLM	951	Analyze time event data via the proportional hazards model.
PLOTP	1096	Print a plot of up to ten sets of points.
POIDF	1114	Evaluate the Poisson distribution function.
POIES	46	Estimate the parameter of the Poisson distribution.
POIPR	1115	Evaluate the Poisson probability function.
PRINC	793	Compute principal components from a variance-covariance matrix or a correlation matrix.
PROBP	1092	Print a probability plot.
PRPFT	463	Perform iterative proportional fitting of a contingency table using a loglinear model.
QTEST	572	Perform a Cochran $Q$ test for related observations.

QUADT	1317	Form a $k$ - $d$ tree.
RANKS	24	Compute the ranks, normal scores, or exponential scores for a vector of observations.
RBCOV	331	Compute a robust estimate of a covariance matrix and mean vector.
RBEST	214	Select the best multiple linear regression models.
RCASE	191	Compute case statistics and diagnostics given data points, coefficient estimates $\hat{\beta}$ , and the $R$ matrix for a fitted general linear model.
RCASP	263	Compute case statistics for a polynomial regression model given the fit based on orthogonal polynomials.
RCOMP	248	Generate an orthogonal central composite design.
RCOV	104	Fit a multiple linear regression model given the variance-covariance matrix.
RCOVB	152	Compute the estimated variance-covariance matrix of the estimated regression coefficients given the $R$ matrix.
RCURV	237	Fit a polynomial curve using least squares.
RFORP	252	Fit an orthogonal polynomial regression model.
RGIVN	107	Fit a multivariate linear regression model via fast Givens transformations.
RGLM	117	Fit a multivariate general linear model.
RHPSS	163	Compute the matrix of sums of squares and crossproducts for the multivariate general linear hypothesis $HBU = G$ given the coefficient estimates $\hat{B}$ and the $R$ matrix.
RHPTE	170	Perform tests for a multivariate general linear hypothesis $HBU = G$ given the hypothesis sums of squares and crossproducts matrix $S_H$ and the error sums of squares and crossproducts matrix $S_E$ .
RINCF	90	Perform response control given a fitted simple linear regression model.
RINPF	94	Perform inverse prediction given a fitted simple linear regression model.
RLAV	293	Fit a multiple linear regression model using the least absolute values criterion.
RLEQU	131	Fit a multivariate linear regression model with linear equality restrictions $HB = G$ imposed on the regression parameters given results from IMSL routine RGIVN after $IDO = 1$ and $IDO = 2$ and prior to $IDO = 3$ .

RLEQU	131	Fit a multivariate linear regression model with linear equality restrictions $HB = G$ imposed on the regression parameters given results from IMSL routine RGIWN after IDO = 1 and IDO = 2 and prior to IDO = 3.
RLINE	79	Fit a line to a set of data points using least squares.
RLLP	297	Fit a multiple linear regression model using the $L_p$ norm criterion.
RLMV	308	Fit a multiple linear regression model using the minimax criterion.
RLOFE	176	Compute a lack-of-fit test based on exact replicates for a fitted regression model.
RLOFN	182	Compute a lack-of-fit test based on near replicates for a fitted regression model.
RLSE	98	Fit a multiple linear regression model using least squares.
RNARM	1232	Generate a time series from a specified ARMA model.
RNBET	1191	Generate pseudorandom numbers from a beta distribution.
RNBIN	1173	Generate pseudorandom numbers from a binomial distribution.
RNCHI	1193	Generate pseudorandom numbers from a chi-squared distribution.
RNCHY	1194	Generate pseudorandom numbers from a Cauchy distribution.
RNCOR	1215	Generate a pseudorandom orthogonal matrix or a correlation matrix.
RNDAT	1218	Generate pseudorandom numbers from a multivariate distribution determined from a given sample.
RNEXP	1196	Generate pseudorandom numbers from a standard exponential distribution.
RNEXT	1197	Generate pseudorandom numbers from a mixture of two exponential distributions.
RNGAM	1198	Generate pseudorandom numbers from a standard gamma distribution.
RNGCS	1200	Set up table to generate pseudorandom numbers from a general continuous distribution.
RNGCT	1202	Generate pseudorandom numbers from a general continuous distribution.
RNGDA	1174	Generate pseudorandom numbers from a general discrete distribution using an alias method.

RNGDS	1177	Set up table to generate pseudorandom numbers from a general discrete distribution.
RNGDT	1181	Generate pseudorandom numbers from a general discrete distribution using a table lookup method.
RNGEF	1167	Retrieve the current value of the array used in the IMSL GFSR random number generator.
RNGEO	1183	Generate pseudorandom numbers from a geometric distribution.
RNGES	1167	Retrieve the current value of the table in the IMSL random number generators that use shuffling.
RNGET	1167	Retrieve the current value of the seed used in the IMSL random number generators.
RNHYP	1185	Generate pseudorandom numbers from a hypergeometric distribution.
RNISD	1168	Determine a seed that yields a stream beginning 100,000 numbers beyond the beginning of the stream yielded by a given seed used in IMSL multiplicative congruential generators (with no shufflings).
RNKSM	557	Perform the Wilcoxon rank sum test.
RNLGR	1186	Generate pseudorandom numbers from a logarithmic distribution.
RNLIN	280	Fit a nonlinear regression model.
RNLNL	1204	Generate pseudorandom numbers from a lognormal distribution.
RNMTN	1222	Generate pseudorandom numbers from a multinomial distribution.
RNMVN	1223	Generate pseudorandom numbers from a multivariate normal distribution.
RNNBN	1188	Generate pseudorandom numbers from a negative binomial distribution.
RNNOA	1205	Generate pseudorandom numbers from a standard normal distribution using an acceptance/rejection method.
RNNOF	1207	Generate a pseudorandom number from a standard normal distribution.
RNNOR	1208	Generate pseudorandom numbers from a standard normal distribution using an inverse CDF method.
RNNOS	1229	Generate pseudorandom order statistics from a standard normal distribution.

RNNPP	1236	Generate pseudorandom numbers from a nonhomogeneous Poisson process.
RNOPG	1166	Retrieve the indicator of the type of uniform random number generator.
RNOPT	1165	Select the uniform (0, 1) multiplicative congruential pseudorandom number generator.
RNPER	1240	Generate a pseudorandom permutation.
RNPOI	1189	Generate pseudorandom numbers from a Poisson distribution.
RNSEF	1167	Initialize the array used in the IMSL GFSR random number generator.
RNSES	1167	Initialize the table in the IMSL random number generators that use shuffling.
RNSET	1167	Initialize a random seed for use in the IMSL randomnumber generators.
RNSPH	1225	Generate pseudorandom points on a unit circle or $\kappa$ -dimensional sphere.
RNSRI	1241	Generate a simple pseudorandom sample of indices.
RNSRS	1242	Generate a simple pseudorandom sample from a finite population.
RNSTA	1209	Generate pseudorandom numbers from a stable distribution.
RNSTT	1210	Generate pseudorandom numbers from a Student's $t$ distribution.
RNTAB	1227	Generate a pseudorandom two-way table.
RNTRI	1212	Generate pseudorandom numbers from a triangular distribution on the interval (0,1).
RNUN	1171	Generate pseudorandom numbers from a uniform (0,1) distribution.
RNUND	1190	Generate pseudorandom numbers from a discrete uniform distribution.
RNUNF	1172	Generate a pseudorandom number from a uniform (0, 1) distribution.
RNUNO	1231	Generate pseudorandom order statistics from a uniform (0, 1) distribution.
RNVMS	1213	Generate pseudorandom numbers from a von Mises distribution.
RNWIB	1214	Generate pseudorandom numbers from a Weibull distribution.

RONE	82	Analyze a simple linear regression model.
RORDM	1268	Reorder rows and columns of a symmetric matrix.
ROREX	429	Reorder the responses from a balanced complete experimental design.
ROTIN	201	Compute diagnostics for detection of outliers and influential data points given residuals and the $R$ matrix for a fitted general linear model.
RPOLY	241	Analyze a polynomial regression model.
RSTAP	258	Compute summary statistics for a polynomial regression model given the fit based on orthogonal polynomials.
RSTAT	141	Compute statistics related to a regression fit given the coefficient estimates $\hat{\beta}$ and the $R$ matrix.
RSTEP	221	Build multiple linear regression models using forward selection, backward selection, or stepwise selection.
RSUBM	233	Retrieve a symmetric submatrix from a symmetric matrix.
RUNS	600	Perform a runs up test.
SCIPM	419	Compute simultaneous confidence intervals on all pairwise differences of means.
SCOLR	1277	Sort columns of a real rectangular matrix using keys in rows.
SCTP	1081	Print a scatterplot of several groups of data.
SDPLC	551	Perform the Cox and Stuart sign test for trends in dispersion and location.
SIGNT	542	Perform a sign test of the hypothesis that a given value is a specified quantile of a distribution.
SMPPR	906	Compute statistics for inferences regarding the population proportion and total, given proportion data from a simple random sample.
SMPPS	909	Compute statistics for inferences regarding the population proportion and total, given proportion data from a stratified random sample.
SMPRR	911	Compute statistics for inferences regarding the population mean and total using ratio or regression estimation, or inferences regarding the population ratio, given a simple random sample.
SMPRS	918	Compute statistics for inferences regarding the population mean and total using ratio or regression estimation, given continuous data from a stratified random sample.



SMFSC	923	Compute statistics for inferences regarding the population mean and total using single-stage cluster sampling with continuous data.
SMPSR	927	Compute statistics for inferences regarding the population mean and total, given data from a simple random sample.
SMPSS	930	Compute statistics for inferences regarding the population mean and total, given data from a stratified random sample.
SMPST	933	Compute statistics for inferences regarding the population mean and total, given continuous data from a two-stage sample with equisized primary units.
SNKMC	424	Perform Student-Newman-Keuls multiple comparison test.
SNRNK	544	Perform a Wilcoxon signed rank test.
SPWF	677	Compute the Wiener forecast operator for a stationary stochastic process.
SPWLK	589	Perform a Shapiro-Wilk $W$ -test for normality.
SRCH	1284	Search a sorted vector for a given scalar and return its index.
SROWR	1280	Sort rows of a real rectangular matrix using keys in columns.
SSRCH	1287	Search a character vector, sorted in ascending ASCII order, for a given string and return its index.
SSWD	729	Estimate the nonnormalized spectral density of a stationary time series using a spectral window given the time series data.
SSWP	736	Estimate the nonnormalized spectral density of a stationary time series using a spectral window given the periodogram.
STBLE	985	Estimate survival probabilities and hazard rates for various parametric models.
STMLP	1085	Print a stem-and-leaf plot.
SVGLM	967	Analyze censored survival data using a generalized linear model.
SVIGN	1275	Sort an integer array by algebraic value.
SVIGP	1276	Sort an integer array by algebraic value and return the permutations.
SVRGN	1273	Sort a real array by algebraic value.
SVRGP	1274	Sort a real array by algebraic value and return the permutations.

SWED	741	Estimation of the nonnormalized spectral density of a stationary time series based on specified periodogram weights given the time series data.
SWEP	747	Estimation of the nonnormalized spectral density of a stationary time series based on specified periodogram weights given the periodogram.
TCSCP	277	Transform coefficients from a quadratic regression model generated from squares and crossproducts of centered variables to a model using uncentered variables.
TDATE	1297	Get today's date.
TDF	1143	Evaluate the Student's $t$ distribution function.
TETCC	342	Categorize bivariate data and compute the tetrachoric correlation coefficient.
TFPE	689	Compute preliminary estimates of parameters for a univariate transfer function model.
TIMDY	1296	Get time of day.
TIN	1145	Evaluate the inverse of the Student's $t$ distribution function.
TNDF	1146	Evaluate the noncentral Student's $t$ distribution function.
TNIN	1149	Evaluate the inverse of the noncentral Student's $t$ distribution function.
TREEP	1098	Print a binary tree.
TRNBL	946	Compute Turnbull's generalized Kaplan-Meier estimates of survival probabilities in samples with interval censoring.
TWFRQ	7	Tally observations into a two-way frequency table.
TWOMV	37	Compute statistics for mean and variance inferences using samples from two normal populations.
UMACH	1338	Set or retrieve input or output device unit numbers.
UVSTA	16	Compute basic univariate statistics.
VERSL	1301	Obtain STAT/LIBRARY-related version, system and license numbers.
VHS2P	1076	Print a vertical histogram with every bar subdivided into two parts.
VHSTP	1074	Print a vertical histogram.
WRIRL	1254	Print an integer rectangular matrix with a given format and labels.
WRIRN	1253	Print an integer rectangular matrix with integer row and column labels.

WROPT	1257	Set or retrieve an option for printing a matrix.
WRRRL	1250	Print a real rectangular matrix with a given format and labels.
WRRRN	1248	Print a real rectangular matrix with integer row and column labels.

# Appendix C: References

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