

# Rpad and R Reference Card

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## Getting help

Most R functions have online documentation.

**help(topic)** documentation on *topic*

**?topic** id.

**help.search("topic")** search the help system

**apropos("topic")** the names of all objects in the search list matching the regular expression "topic"

**help.start()** start the HTML version of help

**str(a)** display the internal \*str\*ucture of an R object

**summary(a)** gives a "summary" of *a*, usually a statistical summary but it is *generic* meaning it has different operations for different classes of *a*

**ls()** show objects in the search path; specify *pat*="pat" to search on a pattern

**ls.str()** str() for each variable in the search path

**dir()** show files in the current directory

**methods(a)** shows S3 methods of *a*

**methods(class=class(a))** lists all the methods to handle objects of class *a*

## Input and output

**load()** load the datasets written with *save*

**data(x)** loads specified data sets

**library(x)** load add-on packages

**read.table(file)** reads a file in table format and creates a data frame from it; the default separator *sep*=" " is any whitespace; use *header*=TRUE to read the first line as a header of column names; use *as.is*=TRUE to prevent character vectors from being converted to factors; use *comment.char*=" " to prevent "#" from being interpreted as a comment; use *skip*=*n* to skip *n* lines before reading data; see the help for options on row naming, NA treatment, and others

**read.csv("filename",header=TRUE)** id. but with defaults set for reading comma-delimited files

**read.delim("filename",header=TRUE)** id. but with defaults set for reading tab-delimited files

**read.fwf(file,widths,header=FALSE,sep="^\t",as.is=FALSE)** read a table of *fixed width formatted data* into a 'data.frame'; *widths* is an integer vector, giving the widths of the fixed-width fields

**save(file,...)** saves the specified objects (...) in the XDR platform-independent binary format

**save.image(file)** saves all objects

**cat(...,file="",sep=" ")** prints the arguments after coercing to character; *sep* is the character separator between arguments

**print(a,...)** prints its arguments; *generic*, meaning it can have different methods for different objects

**format(x,...)** format an R object for pretty printing

**write.table(x,file="",row.names=TRUE,col.names=TRUE,sep=" ")** prints *x* after converting to a data frame; if *quote* is TRUE, character or factor columns are surrounded by quotes (""); *sep* is the

field separator; *eol* is the end-of-line separator; *na* is the string for missing values; use *col.names*=NA to add a blank column header to get the column headers aligned correctly for spreadsheet input

**sink(file)** output to *file*, until *sink()*

Most of the I/O functions have a *file* argument. This can often be a character string naming a file or a connection. *file*="" means the standard input or output. Connections can include files, pipes, zipped files, and R variables.

On windows, the file connection can also be used with *description* = "clipboard". To read a table copied from Excel, use

```
x <- read.delim("clipboard")
```

To write a table to the clipboard for Excel, use

```
write.table(x,"clipboard",sep="\t",col.names=NA)
```

For database interaction, see packages RODBC, DBI, RMySQL, RPgSQL, and ROracle. See packages XML, hdf5, netCDF for reading other file formats.

## Data creation

**c(...)** generic function to combine arguments with the default forming a vector; with *recursive*=TRUE descends through lists combining all elements into one vector

**from:to** generates a sequence; ":" has operator priority; 1:4 + 1 is "2,3,4,5"

**seq(from,to)** generates a sequence *by*= specifies increment; *length*= specifies desired length

**seq(along=x)** generates 1, 2, ..., length(*along*); useful for loops

**rep(x,times)** replicate *x* *times*; use *each*= to repeat "each" element of *x* each *times*; rep(c(1,2,3),2) is 1 2 3 1 2 3; rep(c(1,2,3),each=2) is 1 1 2 2 3 3

**data.frame(...)** create a data frame of the named or unnamed arguments; data.frame(v=1:4,ch=c("a","B","c","d"),n=10); shorter vectors are recycled to the length of the longest

**list(...)** create a list of the named or unnamed arguments; list(a=c(1,2),b="hi",c=3i);

**array(x,dim=)** array with data *x*; specify dimensions like dim=c(3,4,2); elements of *x* recycle if *x* is not long enough

**matrix(x,nrow=,ncol=)** matrix; elements of *x* recycle

**factor(x,levels=)** encodes a vector *x* as a factor

**gl(n,k,length=n\*k,labels=1:n)** generate levels (factors) by specifying the pattern of their levels; *k* is the number of levels, and *n* is the number of replications

**expand.grid()** a data frame from all combinations of the supplied vectors or factors

**rbind(...)** combine arguments by rows for matrices, data frames, and others

**cbind(...)** id. by columns

## Slicing and extracting data

Indexing vectors

```
x[n]          nth element
x[-n]         all but the nth element
x[1:n]        first n elements
x[-(1:n)]     elements from n+1 to the end
x[c(1,4,2)]   specific elements
x["name"]     element named "name"
x[x > 3]      all elements greater than 3
x[x > 3 & x < 5] all elements between 3 and 5
x[x %in% c("a","and","the")] elements in the given set
```

Indexing lists

```
x[n]          list with elements n
x[[n]]        nth element of the list
x[["name"]]   element of the list named "name"
x$name        id.
```

Indexing matrices

```
x[i,j]        element at row i, column j
x[i,]         row i
x[,j]         column j
x[,c(1,3)]    columns 1 and 3
x["name",]    row named "name"
```

Indexing data frames (matrix indexing plus the following)

```
x[["name"]]   column named "name"
x$name        id.
```

## Variable conversion

**as.array(x)**, **as.data.frame(x)**, **as.numeric(x)**,  
**as.logical(x)**, **as.complex(x)**, **as.character(x)**,  
... convert type; for a complete list, use methods(*as*)

## Variable information

**is.na(x)**, **is.null(x)**, **is.array(x)**, **is.data.frame(x)**,  
**is.numeric(x)**, **is.complex(x)**, **is.character(x)**,  
... test for type; for a complete list, use methods(*is*)

**length(x)** number of elements in *x*

**dim(x)** Retrieve or set the dimension of an object; dim(x) <- c(3,2)

**dimnames(x)** Retrieve or set the dimension names of an object

**nrow(x)** number of rows; NROW(x) is the same but treats a vector as a one-row matrix

**ncol(x)** and NCOL(x) id. for columns

**class(x)** get or set the class of *x*; class(x) <- "myclass"

**unclass(x)** remove the class attribute of *x*

**attr(x,which)** get or set the attribute *which* of *x*

**attributes(obj)** get or set the list of attributes of *obj*

## Data selection and manipulation

**which.max(x)** returns the index of the greatest element of *x*

**which.min(x)** returns the index of the smallest element of *x*

**rev(x)** reverses the elements of *x*

**sort(x)** sorts the elements of *x* in increasing order; to sort in decreasing order: rev(sort(x))

**cut(x,breaks)** divides *x* into intervals (factors); *breaks* is the number of cut intervals or a vector of cut points

**match(x,y)** returns a vector of the same length than *x* with the elements of *x* which are in *y* (NA otherwise)

**which(x == a)** returns a vector of the indices of *x* if the comparison operation is true (TRUE), in this example the values of *i* for which *x*[*i*] == *a* (the argument of this function must be a variable of mode logical)

**choose(n,k)** computes the combinations of *k* events among *n* repetitions = *n!*/[(*n*-*k*)!*k*!]

**na.omit(x)** suppresses the observations with missing data (NA) (suppresses the corresponding line if *x* is a matrix or a data frame)

**na.fail(x)** returns an error message if *x* contains at least one NA

**unique(x)** if *x* is a vector or a data frame, returns a similar object but with the duplicate elements suppressed

**table(x)** returns a table with the numbers of the different values of *x* (typically for integers or factors)

**subset(x, ...)** returns a selection of *x* with respect to criteria (...), typically comparisons: `x$V1 < 10`; if *x* is a data frame, the option `select` gives the variables to be kept or dropped using a minus sign

**sample(x, size)** resample randomly and without replacement `size` elements in the vector *x*, the option `replace = TRUE` allows to resample with replacement

**prop.table(x, margin=)** table entries as fraction of marginal table

## Math

**sin, cos, tan, asin, acos, atan, atan2, log, log10, exp**

**max(x)** maximum of the elements of *x*

**min(x)** minimum of the elements of *x*

**range(x)** id. then `c(min(x), max(x))`

**sum(x)** sum of the elements of *x*

**diff(x)** lagged and iterated differences of vector *x*

**prod(x)** product of the elements of *x*

**mean(x)** mean of the elements of *x*

**median(x)** median of the elements of *x*

**quantile(x, probs=)** sample quantiles corresponding to the given probabilities (defaults to 0, .25, .5, .75, 1)

**weighted.mean(x, w)** mean of *x* with weights *w*

**rank(x)** ranks of the elements of *x*

**var(x)** or `cov(x)` variance of the elements of *x* (calculated on  $n - 1$ ); if *x* is a matrix or a data frame, the variance-covariance matrix is calculated

**sd(x)** standard deviation of *x*

**cor(x)** correlation matrix of *x* if it is a matrix or a data frame (1 if *x* is a vector)

**var(x, y)** or `cov(x, y)` covariance between *x* and *y*, or between the columns of *x* and those of *y* if they are matrices or data frames

**cor(x, y)** linear correlation between *x* and *y*, or correlation matrix if they are matrices or data frames

**round(x, n)** rounds the elements of *x* to *n* decimals

**log(x, base)** computes the logarithm of *x* with base *base*

**scale(x)** if *x* is a matrix, centers and reduces the data; to center only use the option `center=FALSE`, to reduce only `scale=FALSE` (by default `center=TRUE`, `scale=TRUE`)

**pmin(x, y, ...)** a vector which *i*th element is the minimum of `x[i]`, `y[i]`, ...

**pmax(x, y, ...)** id. for the maximum

**cumsum(x)** a vector which *i*th element is the sum from `x[1]` to `x[i]`

**cumprod(x)** id. for the product

**cummin(x)** id. for the minimum

**cummax(x)** id. for the maximum

**union(x, y)**, **intersect(x, y)**, **setdiff(x, y)**, **setequal(x, y)**, **is.element(e1, set)** “set” functions

**Re(x)** real part of a complex number

**Im(x)** imaginary part

**Mod(x)** modulus; `abs(x)` is the same

**Arg(x)** angle in radians of the complex number

**Conj(x)** complex conjugate

**convolve(x, y)** compute the several kinds of convolutions of two sequences

**fft(x)** Fast Fourier Transform of an array

**mvfft(x)** FFT of each column of a matrix

**filter(x, filter)** applies linear filtering to a univariate time series or to each series separately of a multivariate time series

Many math functions have a logical parameter `na.rm=FALSE` to specify missing data (NA) removal.

## Matrices

**t(x)** transpose

**diag(x)** diagonal

**%\*%** matrix multiplication

**solve(a, b)** solves  $a \%*\% x = b$  for *x*

**solve(a)** matrix inverse of *a*

**rowsum(x)** sum of rows for a matrix-like object; **rowSums(x)** is a faster version

**colsum(x)**, **colSums(x)** id. for columns

**rowMeans(x)** fast version of row means

**colMeans(x)** id. for columns

## Advanced data processing

**apply(X, INDEX, FUN=)** a vector or array or list of values obtained by applying a function *FUN* to margins (*INDEX*) of *X*

**lapply(X, FUN)** apply *FUN* to each element of the list *X*

**tapply(X, INDEX, FUN=)** apply *FUN* to each cell of a ragged array given by *X* with indexes *INDEX*

**by(data, INDEX, FUN)** apply *FUN* to data frame *data* subsetted by *INDEX*

**merge(a, b)** merge two data frames by common columns or row names

**xtabs(a, b, data=x)** a contingency table from cross-classifying factors

**aggregate(x, by, FUN)** splits the data frame *x* into subsets, computes summary statistics for each, and returns the result in a convenient form; *by* is a list of grouping elements, each as long as the variables in *x*

**stack(x, ...)** transform data available as separate columns in a data frame or list into a single column

**unstack(x, ...)** inverse of `stack()`

**reshape(x, ...)** reshapes a data frame between ‘wide’ format with repeated measurements in separate columns of the same record and ‘long’ format with the repeated measurements in separate records; use `(direction=“wide”)` or `(direction=“long”)`

## Strings

**paste(...)** concatenate vectors after converting to character; `sep=` is the string to separate terms (a single space is the default); `collapse=` is an optional string to separate “collapsed” results

**substr(x, start, stop)** substrings in a character vector; can also assign, as `substr(x, start, stop) <- value`

**strsplit(x, split)** split *x* according to the substring *split*

**grep(pattern, x)** searches for matches to *pattern* within *x*; see `?regex`

**gsub(pattern, replacement, x)** replacement of matches determined by regular expression matching `sub()` is the same but only replaces the first occurrence.

**tolower(x)** convert to lowercase

**toupper(x)** convert to uppercase

**match(x, table)** a vector of the positions of first matches for the elements of *x* among *table*

**x %in% table** id. but returns a logical vector

**pmatch(x, table)** partial matches for the elements of *x* among *table*

**nchar(x)** number of characters

## Dates and Times

The class `Date` has dates without times. `POSIXct` has dates and times, including time zones. Comparisons (e.g. `>`), `seq()`, and `difftime()` are useful. `Date` also allows `+` and `-`. `?DateTimeClasses` gives more information. See also package `chron`.

**as.Date(s)** and **as.POSIXct(s)** convert to the respective class; `format(dt)` converts to a string representation. The default string format is “2001-02-21”. These accept a second argument to specify a format for conversion. Some common formats are:

`%a`, `%A` Abbreviated and full weekday name.

`%b`, `%B` Abbreviated and full month name.

`%d` Day of the month (01–31).

`%H` Hours (00–23).

`%I` Hours (01–12).

`%j` Day of year (001–366).

`%m` Month (01–12).

`%M` Minute (00–59).

`%p` AM/PM indicator.

`%S` Second as decimal number (00–61).

`%U` Week (00–53); the first Sunday as day 1 of week 1.

`%w` Weekday (0–6, Sunday is 0).

`%W` Week (00–53); the first Monday as day 1 of week 1.

`%y` Year without century (00–99). Don’t use.

`%Y` Year with century.

`%z` (output only.) Offset from Greenwich; `-0800` is 8 hours west of.

`%Z` (output only.) Time zone as a character string (empty if not available).

Where leading zeros are shown they will be used on output but are optional on input. See `?strftime`.

## Plotting

**plot(x)** plot of the values of *x* (on the *y*-axis) ordered on the *x*-axis

**plot(x, y)** bivariate plot of *x* (on the *x*-axis) and *y* (on the *y*-axis)

**hist(x)** histogram of the frequencies of *x*

**barplot(x)** histogram of the values of *x*; use `horiz=FALSE` for horizontal bars

**dotplot(x)** if *x* is a data frame, plots a Cleveland dot plot (stacked plots line-by-line and column-by-column)

**piechart(x)** circular pie-chart

**boxplot(x)** “box-and-whiskers” plot

**sunflowerplot(x, y)** id. than `plot()` but the points with similar coordinates are drawn as flowers which petal number represents the number of points

**stripplot(x)** plot of the values of *x* on a line (an alternative to `boxplot()` for small sample sizes)

**coplot(x~y | z)** bivariate plot of *x* and *y* for each value or interval of values of *z*

**interaction.plot(f1, f2, y)** if *f1* and *f2* are factors, plots the means of *y* (on the *y*-axis) with respect to the values of *f1* (on the *x*-axis) and of *f2* (different curves); the option `fun` allows to choose the summary statistic of *y* (by default `fun=mean`)

**matplot(x, y)** bivariate plot of the first column of *x* vs. the first one of *y*, the second one of *x* vs. the second one of *y*, etc.

**fourfoldplot(x)** visualizes, with quarters of circles, the association between two dichotomous variables for different populations ( $x$  must be an array with  $\text{dim}=\text{c}(2, 2, k)$ , or a matrix with  $\text{dim}=\text{c}(2, 2)$  if  $k=1$ )

**assocplot(x)** Cohen–Friendly graph showing the deviations from independence of rows and columns in a two dimensional contingency table

**mosaicplot(x)** ‘mosaic’ graph of the residuals from a log-linear regression of a contingency table

**pairs(x)** if  $x$  is a matrix or a data frame, draws all possible bivariate plots between the columns of  $x$

**plot.ts(x)** if  $x$  is an object of class "ts", plot of  $x$  with respect to time,  $x$  may be multivariate but the series must have the same frequency and dates

**ts.plot(x)** id. but if  $x$  is multivariate the series may have different dates and must have the same frequency

**qqnorm(x)** quantiles of  $x$  with respect to the values expected under a normal law

**qqplot(x, y)** quantiles of  $y$  with respect to the quantiles of  $x$

**contour(x, y, z)** contour plot (data are interpolated to draw the curves),  $x$  and  $y$  must be vectors and  $z$  must be a matrix so that  $\text{dim}(z)=\text{c}(\text{length}(x), \text{length}(y))$  ( $x$  and  $y$  may be omitted)

**filled.contour(x, y, z)** id. but the areas between the contours are coloured, and a legend of the colours is drawn as well

**image(x, y, z)** id. but with colours (actual data are plotted)

**persp(x, y, z)** id. but in perspective (actual data are plotted)

**stars(x)** if  $x$  is a matrix or a data frame, draws a graph with segments or a star where each row of  $x$  is represented by a star and the columns are the lengths of the segments

**symbols(x, y, ...)** draws, at the coordinates given by  $x$  and  $y$ , symbols (circles, squares, rectangles, stars, thermometres or “boxplots”) which sizes, colours ... are specified by supplementary arguments

**termpart(mod.obj)** plot of the (partial) effects of a regression model (mod.obj)

The following parameters are common to many plotting functions:

**add=FALSE** if TRUE superposes the plot on the previous one (if it exists)

**axes=TRUE** if FALSE does not draw the axes and the box

**type="p"** specifies the type of plot, "p": points, "l": lines, "b": points connected by lines, "o": id. but the lines are over the points, "h": vertical lines, "s": steps, the data are represented by the top of the vertical lines, "S": id. but the data are represented by the bottom of the vertical lines

**xlim=, ylim=** specifies the lower and upper limits of the axes, for example with  $\text{xlim}=\text{c}(1, 10)$  or  $\text{xlim}=\text{range}(x)$

**xlab=, ylab=** annotates the axes, must be variables of mode character

**main=** main title, must be a variable of mode character

**sub=** sub-title (written in a smaller font)

## Low-level plotting commands

**points(x, y)** adds points (the option `type=` can be used)

**lines(x, y)** id. but with lines

**text(x, y, labels, ...)** adds text given by `labels` at coordinates ( $x,y$ ); a typical use is: `plot(x, y, type="n"); text(x, y, names)`

**mtext(text, side=3, line=0, ...)** adds text given by `text` in the margin specified by `side` (see `axis()` below); `line` specifies the line from the plotting area

**segments(x0, y0, x1, y1)** draws lines from points ( $x_0,y_0$ ) to points ( $x_1,y_1$ )

**arrows(x0, y0, x1, y1, angle= 30, code=2)** id. with arrows at points ( $x_0,y_0$ ) if `code=2`, at points ( $x_1,y_1$ ) if `code=1`, or both if `code=3`; `angle` controls the angle from the shaft of the arrow to the edge of the arrow head

**abline(a,b)** draws a line of slope  $b$  and intercept  $a$

**abline(h=y)** draws a horizontal line at ordinate  $y$

**abline(v=x)** draws a vertical line at abscissa  $x$

**abline(lm.obj)** draws the regression line given by `lm.obj`

**rect(x1, y1, x2, y2)** draws a rectangle which left, right, bottom, and top limits are  $x_1, x_2, y_1$ , and  $y_2$ , respectively

**polygon(x, y)** draws a polygon linking the points with coordinates given by  $x$  and  $y$

**legend(x, y, legend)** adds the legend at the point ( $x,y$ ) with the symbols given by `legend`

**title()** adds a title and optionally a sub-title

**axis(side, vect)** adds an axis at the bottom (`side=1`), on the left (`2`), at the top (`3`), or on the right (`4`); `vect` (optional) gives the abscissa (or ordinates) where tick-marks are drawn

**rug(x)** draws the data  $x$  on the  $x$ -axis as small vertical lines

**locator(n, type="n", ...)** returns the coordinates ( $x,y$ ) after the user has clicked  $n$  times on the plot with the mouse; also draws symbols (`type="p"`) or lines (`type="l"`) with respect to optional graphic parameters (...); by default nothing is drawn (`type="n"`)

## Graphical parameters

These can be set globally with `par(...)`; many can be passed as parameters to plotting commands.

**adj** controls text justification (0 left-justified, 0.5 centred, 1 right-justified)  
**bg** specifies the colour of the background (ex. : `bg="red"`, `bg="blue"`, ... the list of the 657 available colours is displayed with `colors()`)

**bty** controls the type of box drawn around the plot, allowed values are: "o", "l", "7", "c", "u" ou "j" (the box looks like the corresponding character); if `bty="n"` the box is not drawn

**cex** a value controlling the size of texts and symbols with respect to the default; the following parameters have the same control for numbers on the axes, `cex.axis`, the axis labels, `cex.lab`, the title, `cex.main`, and the sub-title, `cex.sub`

**col** controls the color of symbols and lines; use color names: "red", "blue" see `colors()` or as "#RRGGBB"; see `rgb()`, `hsv()`, `gray()`, and `rainbow()`; as for `cex` there are: `col.axis`, `col.lab`, `col.main`, `col.sub`

**font** an integer which controls the style of text (1: normal, 2: italics, 3: bold, 4: bold italics); as for `cex` there are: `font.axis`, `font.lab`, `font.main`, `font.sub`

**las** an integer which controls the orientation of the axis labels (0: parallel to the axes, 1: horizontal, 2: perpendicular to the axes, 3: vertical)

**lty** controls the type of lines, can be an integer or string (1: "solid", 2: "dashed", 3: "dotted", 4: "dotdash", 5: "longdash", 6: "twodash", or a string of up to eight characters (between "0" and "9") which specifies alternatively the length, in points or pixels, of the drawn elements and the blanks, for example `lty="44"` will have the same effect than `lty=2`

**lwd** a numeric which controls the width of lines, default 1

**mar** a vector of 4 numeric values which control the space between the axes and the border of the graph of the form `c(bottom, left, top, right)`, the default values are `c(5.1, 4.1, 4.1, 2.1)`

**mfcol** a vector of the form `c(nr,nc)` which partitions the graphic window as a matrix of `nr` lines and `nc` columns, the plots are then drawn in columns

**mfrow** id. but the plots are drawn by row

**pch** controls the type of symbol, either an integer between 1 and 25, or any single character within ""

1 ○ 2 △ 3 + 4 × 5 ◇ 6 ▽ 7 ☒ 8 \* 9 ⊕ 10 ⊕ 11 ☒ 12 ⊕ 13 ☒ 14 □ 15 ■  
16 ● 17 ▲ 18 ♦ 19 ● 20 ● 21 ○ 22 □ 23 ◇ 24 △ 25 ▽ \* . . . X X a a ? ?

**ps** an integer which controls the size in points of texts and symbols

**pty** a character which specifies the type of the plotting region, "s": square, "m": maximal

**tck** a value which specifies the length of tick-marks on the axes as a fraction of the smallest of the width or height of the plot; if `tck=1` a grid is drawn

**tcl** a value which specifies the length of tick-marks on the axes as a fraction of the height of a line of text (by default `tcl=-0.5`)

**xaxt** if `xaxt="n"` the  $x$ -axis is set but not drawn (useful in conjunction with `axis(side=1, ...)`)

**yaxt** if `yaxt="n"` the  $y$ -axis is set but not drawn (useful in conjunction with `axis(side=2, ...)`)

## Lattice (Trellis) graphics

**barchart(y~x)** histogram of the values of  $y$  with respect to those of  $x$

**bwplot(y~x)** “box-and-whiskers” plot

**densityplot(~x)** density functions plot

**dotplot(y~x)** Cleveland dot plot (stacked plots line-by-line and column-by-column)

**histogram(~x)** histogram of the frequencies of  $x$

**qqmath(~x)** quantiles of  $x$  with respect to the values expected under a theoretical distribution

**stripplot(y~x)** single dimension plot,  $x$  must be numeric,  $y$  may be a factor

**qq(y~x)** quantiles to compare two distributions,  $x$  must be numeric,  $y$  may be numeric, character, or factor but must have two ‘levels’

**xyplot(y~x)** bivariate plots (with many functionalities)

**levelplot(z~x\*y)** coloured plot of the values of  $z$  at the coordinates given by  $x$  and  $y$  ( $x, y$  and  $z$  are all of the same length)

**splo(m~x)** matrix of bivariate plots

**parallel(~x)** parallel coordinates plot

## Optimization and model fitting

**optim(par, fn, method = c("Nelder-Mead", "BFGS", "CG", "L-BFGS-B", "SANN"))** general-purpose optimization; `par` is initial values, `fn` is function to optimize (normally minimize)

**nlm(f,p)** minimize function  $f$  using a Newton-type algorithm with starting values  $p$

**lm(formula)** fit linear models; `formula` is typically of the form  $\text{response} = \text{termA} + \text{termB} + \dots$ ; use  $I(x*y)$  +  $I(x^2)$  for terms made of nonlinear components



**glm(formula, family=)** fit generalized linear models, specified by giving a symbolic description of the linear predictor and a description of the error distribution; *family* is a description of the error distribution and link function to be used in the model; see ?family

**nls(formula)** nonlinear least-squares estimates of the nonlinear model parameters

**approx(x,y=)** linearly interpolate given data points; *x* can be an xy plotting structure

**spline(x,y=)** cubic spline interpolation

**loess(formula)** fit a polynomial surface using local fitting

Many of the formula-based modeling functions have several common arguments: *data*= the data frame for the formula variables, *subset*= a subset of variables used in the fit, *na.action*= action for missing values: "na.fail", "na.omit", or a function. The following generics often apply to model fitting functions:

**predict(fit,...)** predictions from *fit* based on input data

**df.residual(fit)** returns the number of residual degrees of freedom

**coef(fit)** returns the estimated coefficients (sometimes with their standard-errors)

**residuals(fit)** returns the residuals

**deviance(fit)** returns the deviance

**fitted(fit)** returns the fitted values

**logLik(fit)** computes the logarithm of the likelihood and the number of parameters

**AIC(fit)** computes the Akaike information criterion or AIC

## Statistics

**aov(formula)** analysis of variance model

**anova(fit,...)** analysis of variance (or deviance) tables for one or more fitted model objects

**density(x)** kernel density estimates of *x*

**binom.test(), pairwise.t.test(), power.t.test(), prop.test(), t.test(), ...** use `help.search("test")`

## Distributions

**rnorm(n, mean=0, sd=1)** Gaussian (normal)

**rexp(n, rate=1)** exponential

**rgamma(n, shape, scale=1)** gamma

**rpois(n, lambda)** Poisson

**rweibull(n, shape, scale=1)** Weibull

**rcauchy(n, location=0, scale=1)** Cauchy

**rbeta(n, shape1, shape2)** beta

**rt(n, df)** 'Student' (*t*)

**rf(n, df1, df2)** Fisher–Snedecor (*F*) ( $\chi^2$ )

**rchisq(n, df)** Pearson

**rbinom(n, size, prob)** binomial

**rgeom(n, prob)** geometric

**rhyper(nn, m, n, k)** hypergeometric

**rlogis(n, location=0, scale=1)** logistic

**rlnorm(n, meanlog=0, sdlog=1)** lognormal

**rnbinom(n, size, prob)** negative binomial

**runif(n, min=0, max=1)** uniform

**rwilcox(nn, m, n), rsignrank(nn, n)** Wilcoxon's statistics

All these functions can be used by replacing the letter *r* with *d*, *p* or *q* to get, respectively, the probability density (*dfunc(x, ...)*), the cumulative probability density (*pfunc(x, ...)*), and the value of quantile (*qfunc(p, ...)*), with  $0 < p < 1$ .

## Programming

**function( arglist ) expr** function definition

**return(value)**

**if(cond) expr**

**if(cond) cons.expr else alt.expr**

**for(var in seq) expr**

**while(cond) expr**

**repeat expr**

**break**

**next**

Use braces `{}` around statements

**ifelse(test, yes, no)** a value with the same shape as *test* filled with elements from either *yes* or *no*

**do.call(funname, args)** executes a function call from the name of the function and a list of arguments to be passed to it

## Rpad HTML utilities

**HTML(x)** outputs an HTML representation of an object (uses package R2HTML)

**RpadDir** current Rpad directory

**HTMLon()** turn on HTML mode (the default is text)

**HTMLoff()** turn off HTML mode

**ROutputMode()** changes how R behaves with automatic printing; possible values are: "text" (the default), "html", or "none". "text" is like the command line: values returned in the script are automatically printed (without an explicit `print` statement) in standard text format. With "html", values returned are automatically printed, but HTML output is generated by using the HTML method instead of the `print` method. `ROutputMode` applies to all subsequent Rpad input section, including a rollover back to the beginning when a page is run several times.

**HTMLh1(text), HTMLh2(text), ...** output HTML headers H1, H2, ...

**HTMLargs(...)** returns a string with the arguments as "a='arg1' b='arg2'", and so on

**HTMLelement(tagName, ...)** output HTML tag "tagName" with arguments given by ...

**HTMLradio(variableName, commonName="radio", text="")** output HTML for a radio input with an R variable name *variableName*; *commonName* links it with other radio elements; *text* specifies an adjacent label

**HTMLcheckbox(name, text="", checked=FALSE)** output HTML for a checkbox input with an R variable name *name*

**HTMLselect(name, text, default=1, size=1)** output HTML for a select box with an R variable name *name* with the options *text*

**HTMLinput(name, value="", rpadtype="Rvariable")** output HTML for an input box with an R variable name *name* and default value; *rpadtype* can be "Rstring" or "Rvariable"

**HTMLlink(url, text)** output HTML for a link wrapped around *text*

**HTMLimg(filename)** output HTML for an `img`

**HTMLembed(filename,width=600,height=600)** output HTML for an `embed` (useful for pdf or svg)

## Rpad plotting utilities

**newgraph(name="",...)** sets up the graphics device (not needed unless you want to change parameters)

**showgraph()** generates the HTML to show the graph and runs `newgraph` to advance to the next graphics file; *link*=TRUE creates a link to the EPS file

**graphoptions(...)** changes the defaults for subsequent graphs `newgraph` and `graphoptions` have the following options with the defaults given: `type="pngalpha"`, `res=120`, `width=3.5`, `height=`, `pointsize=10`, `sublines=0`, `toplines=.6`, `ratio=4/3`, `leftlines=0`, `lwd=0.6`