

# R cheat sheet

Copied in part from <https://cran.r-project.org/doc/contrib/Short-refcard.pdf>

## Getting help

?**topic** gives documentation on given topic.

## Input & output

**install.packages("name")** installs the given package to this computer

**library(name)** 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

**write.table(x, file="<file>")** saves data frame to file

## Data creation

**c(...)** generic function to combine arguments into a vector

**from:to** generates a sequence

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

**seq(along=x)** generates 1, 2, ..., length(along)

**rep(x,times)** replicate x times; use `each=` to repeat "each" element of x each times

**data.frame(...)** create a data frame of the named or unnamed arguments; shorter vectors are recycled to the length of the longest

**list(...)** create a list of the named or unnamed arguments

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

**gl(n,k, labels=<labels>)** generate levels (factors) by specifying the pattern of their levels; n is the number of levels, and k is the number of replications

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

**cbind(...)** idem by columns

## Slicing & extracting data

### Indexing vectors

**x[n]** n<sup>th</sup> element  
**x[-n]** all but the n<sup>th</sup> 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]]** n<sup>th</sup> element of the list  
**x[["name"]]** element of the list named "name"  
**x\$name** idem

### Indexing data frames

**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"  
**x[["name"]]** column named "name"  
**x\$name** idem

## 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

**nrow(x)** number of rows

**ncol(x)** number of columns

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

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

## Data selection and manipulation

**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)

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

**which(expr on x)** returns a vector of the indices of x if the comparison operation is true

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

**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 (...); if x is a data frame, the option `select` gives the variables to be kept (or dropped using a minus sign)

**sample(x, size)** sample randomly from x

## Math

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

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

**range(x)** gives `c(min(x), max(x))`

**sum(x)** sum 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)

**var(x)** or `cov(x)` variance of the elements of x (calculated on n-1);

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

**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

## Advanced data processing

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

**lapply(X, FUN)** apply FUN to each element of the list X; **sapply** is a wrapper for lapply that tries to simplify the result.

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

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

**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

## Strings

**paste(...)** concatenate vectors after converting to character  
**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  
**x %in% table** a logical vector of the matches for the elements of x among table  
**nchar(x)** number of characters

## Plotting

**plot(x)** plots 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  
**pie(x)** circular pie-chart  
**boxplot(x)** "box-and-whiskers" plot  
**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  
**pairs(x)** if x is a matrix or a data frame, draws all possible bivariate plots between the columns of x  
**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  
**matplot(x, y)** plot the columns of one matrix against the columns of another, or the columns of a matrix (or dataframe).

*The following parameters are common to many plotting functions:*

**add=FALSE** if TRUE superposes the plot on the previous one  
**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  
**xlab=, ylab=** annotates the axes with labels  
**main=** main title ; **sub=** sub-title

## 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)  
**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  
**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

## Graphical parameters

These can be set globally with `par(...)`; many can be passed as parameters to plotting commands. Type ?par for more info  
**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 "]" (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; as for cex there are: font.axis, font.lab, font.main, font.sub  
**las** an integer which controls the orientation of the axis labels  
**lty** controls the type of lines  
**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: c(bottom, left, top, right)  
**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 ""

## Optimization and model fitting

**lm(formula)** fit linear models  
**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

## 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)  
**rbinom(n, size, prob)** binomial  
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).  
**NB:** There are many more distributions!

## Programming

**function( arglist ) expr** function definition  
**return(value)**  
**if(cond) expr**  
**if(cond) cons.expr else alt.expr**  
**for(var in seq) expr** loop over seq and put the elements in var  
**while(cond) expr** loop while condition is true  
**break** breaks out of loop iteration  
**next** goes to next loop iteration  
Use braces {} around statements  
**ifelse(test, yes, no)** a value with the same shape as test filled with elements from either yes or no