R Quick Reference, by E. Slate and E. Hill, adapted from the "R Reference Card" by Jonathan Baron. Parentheses are for functions, brackets are for indicating the position of items in a vector or matrix.

Miscellaneous

q()	quit
options()	view/set global options, e.g. number of digits
history()	view past commands you've issued
<-	assignment
INSTALL package1	install package1
m1[,2]	column 2 of matrix m1
m1[,2:5] or	columns 2–5
m1[,c(2,3,4,5)]	
m1\$a1	variable a1 in data frame or list m1
NA	missing data
is.na()	true if data missing
library(mva)	load the package (e.g.) mva
require(mva)	load the package (e.g.) mva, if not already loaded
NaN	not a number
Inf	infinity
data()	available data sets
demo()	run demos

Help

help(command1)	get help with command1 (use this command for more detail than is provided here)
help.start()	start browser help
help(package=mva)	help with (e.g.) package mva
apropos("topic1")	commands relevant to topic1
example(command1)	examples of command1
args(fn1)	show arguments for function fn1

Input and output

source("file1")	run the commands in file1
read.table("file1"),	read in data from file1
read.csv("file1"), read.delim("f	ile1")
data.entry()	spreadsheet
scan("file1")	read from file1 (primitive)
download.file(url1)	from internet
url.show(url1),	remote input
read.table.url(url1)	
sink("file1")	output to file1, until sink()
write(object, "file1")	writes an object to file1
<pre>write.table(dataframe1, "file1")</pre>	writes a table to file1

Arithmetic

8*8	matrix multiplication
%/%, ^, %%, sqrt()	integer division, power, modulus, square root
outer	outer "product" function

Managing variables and objects

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attach(x1)	put variables in x1 in search path
detach(x1)	remove from search path
search()	view the search path
ls()	lists all the active objects
rm(object1)	removes object1
<pre>save(obj), save.image(fname)</pre>	save the workspace
load(fname)	load the workspace in fname
dim(matrix1)	dimensions of matrix1
dimnames(x1)	names of dimensions of x1
names(df1)	variable names in data frame df1
length(vector1)	length of vector1
1:3	the vector 1, 2, 3
c(1,2,3)	creates the same vector
seq(from, to, by),	create a sequence
seq(from, to, length)	
rep(x1,n1)	repeats the vector x1 n1 times
cbind(a1,b1,c1),	binds columns or rows into a matrix
rbind(a1,b1,c1)	
merge(df1,df2)	merge data frames
matrix(vector1,r1,c1)	make vector1 into a matrix with r1
	rows and c1 columns
data.frame(var1=v1,var2=v2)	make a data frame from vectors v1
	and v2
as.factor(), as.matrix(),	conversion
as.vector()	
<pre>is.factor(), is.matrix(),</pre>	what it is—returns TRUE/FALSE
is.vector()	
t()	transpose
which(x1==a1)	returns indices of $x1$ where $x1==a1$
unique(x1)	returns the unique elements in x1
is.element(e, x1),	set operations
union(x1,x2), setdiff(x1,x2),	
intersect(x1,x2)	

Control flow

for (i1 in vector1) {}	repeat length (vector1) times
<pre>if (condition1) { } else { }</pre>	conditional
while (condition) {}	do while condition is TRUE

Logic

! x	NOT x, elementwise
х & у	elementwise AND, all elements evaluated
х && У	sequential AND, only first elements of x, y used
x y, x y	elementwise and sequential OR
xor(x, y)	elementwise exclusive OR

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Numerical summaries

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<pre>max(), min(), mean(), quantile(), median(), sum(), var(), cor()</pre>	as named
summary(data.frame)	prints statistics
rank(), sort(), order()	ranking and sorting
ave(x1,y1)	averages of x1 grouped by factor y1
by()	apply function to data frame by factor
apply(x1,n1,function1)	apply function1 (e.g. mean) to x by
	rows (n1=1) or columns (n2=2)
tapply(x1,list1,function1)	apply function to x1 by list1
lapply(list1, function1)	apply function1 to the elements of
sapply(list1, function1)	list1; sapply simplifies
table()	make a table
tabulate()	tabulate a vector

Basic statistical analysis

Dasic statistical alialysis		
aov(), anova(), lm(), glm(),	linear and nonlinear models, anova	
nls(), nlm()		
<pre>more for regression: confint(), deviance(), df.residual(),</pre>		
rstandard(), rstudent(), dffits(), dfbetas(),		
<pre>cooks.distance(), hatvalues(), vcov(), predict()</pre>		
t.test()	t test	
<pre>prop.test(), binom.test()</pre>	proportions tests	
chisq.test(matrix1)	chi-square test on matrix1 columns	
fisher.test()	Fisher exact test	
cor(a)	show correlations	
cor.test(a,b)	test correlation	
friedman.test()	Friedman test	
<pre>runif(), rnorm(), rgamma(),</pre>	probability distributions; prefix is	
<pre>rbeta(), rchisq(), rbinom(),</pre>	r = random, p = prob, d = density,	
rt(), rpois(), etc.	q = quantile	
optim	general purpose optimization	
contrasts()	set dummy coding for factor variables	
library(help="stats")	additional standard stat methods	

Some statistics in mva package

	p
prcomp()	principal components
kmeans()	kmeans cluster analysis
factanal()	factor analysis
cancor()	canonical correlation

Graphics

<pre>windows(), postscript(), pdf()</pre>	new graphics device
<pre>plot(x,y), barplot(),</pre>	basic plots
<pre>boxplot(), stem(), hist()</pre>	
matplot(xmat, ymat)	<pre>matrix plot ymat[,i] on xmat[,i]</pre>
pairs(matrix)	scatterplots of all pairs of columns
scatter.smooth()	scatterplot with smooth trend

coplot()	conditional plot
stripplot()	strip plot (lattice)
qqplot()	quantile-quantile plot
qqnorm(), qqline()	fit normal distribution
<pre>contour(), persp()</pre>	plots for 3D data
heatmap()	
<pre>points(), lines(), segments(),</pre>	
<pre>arrows(), text(), polygon(),</pre>	add to current plot
<pre>symbols(), abline()</pre>	
mfrow(), layout()	multiple plotting regions per page
par()	View/set graphics parameters
<pre>lattice package: xyplot(),</pre>	
<pre>bwplot(), densityplot(),</pre>	trollis graphics
<pre>cloud(), wireframe(), splom(),</pre>	trellis graphics
parallel()	

Programmimg

<pre>print(), cat(),</pre>	debugging
traceback(),	
options(error =	
dump.frames),	
debugger()	
substitute()	read the help carefully!
missing()	check for missing function arguments
	additional name = value type arguments
match.call()	expand function arguments
do.call()	evaluate a constructed function call
R CMD	DOS/unix command line interface

Useful packages

Oscial packages	
lattice	trellis graphics
survival	survival analyses
mvtnorm	multivariate normal and t distributions
maps	displaying geographical data
boot	bootstrapping
nnet	neural nets
nlme, lme4	linear and nonlinear mixed models
hmisc, xtable	exporting tables to LateX, HTML, plus others
coda	handling output from BUGS (see also the BOA code)
ellipse	for elliptical confidence regions in plots
rggobi	graphical data exploration
gvlma	a new diagnostic for linear models
cluster	various clustering methods
mclust	model-based (i.e. Gaussian) clustering
gam	generalized additive models
BRugs, rbugs, R2WinBUGS	linking R and Win/OpenBUGS

See also <u>www.r-project.org</u>, <u>http://cran.us.r-project.org/</u>, <u>http://www.bioconductor.org/</u>