

# One Page R Reference Card

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c(a, b): concatenate, vectorize # comment char  
install.packages("package1"): install package package1  
is.na(x1): TRUE if x1 == NA  
library(package1): load package package1  
NA: missing value 3e3:  $3 \times 10^3 = 3000$  (e-notation)  
options(scipen=100): get rid of e-notation  
<- or assign(): assign  
q("no"): quit R, do not save environment  
history(Inf); savehistory(file1): look on command history;  
    save history (not on macOS GUI)  
; used to separate commands :: used as package::command()  
Tab; Up; Ctrl+U: complete; repeat command; delete command  
    (not on macOS GUI)

## Help

example(com1): run examples for command com1  
help(com1) or ?com1: help about command com1  
help(package=rpart): help for the package, e.g. rpart  
function1; methods(function2); getAnywhere(method2): look  
    on the function1 and 2 codes  
??topic1": finds topic1 in all help files (slow!)

## Entering and saving data

dir(...): list files in directory, go to another  
read.table("file1", h=T, sep=";", as.is=T): read data into  
    data frame from file1 which has header and semicolon as  
    separator; do not convert variables into factors  
scan("file1", what="char"): read one series of character codes  
    from disk into variable  
sink("file1", split=TRUE): output to file1 and to the terminal  
    until sink()  
source("file1.r"): run commands from file file1.r  
write.table(x1, "file1"): write object x1 to the file file1

## Manage variables and objects

1:3 or c(1, 2, 3): concatenate 1, 2, 3 into vector  
as.data.frame(x1), as.matrix(x1): conversion  
cbind(a1, b1, c1) or rbind(a1, b1, c1): join columns or  
    rows into matrix  
cut(v1, 2, labels=c("small", "big")): split vector v1 in  
    two intervals  
data.frame(v1, v2): list from same-length vectors v1 and v2  
df1\$a1: variable (column) named a1 from data frame df1  
dimnames(mat1), or names(df1) and row.names(df1): names of  
    rows and columns of mat1 or df1

droplevels(factor1): drop unused factor levels  
grep("str1", x1): search str1 in x1  
gsub("str1", "str2", x1): replace str1 to the str2 in x1  
head(df1): first rows of data frame  
length(v1), nrow(mat1), ncol(df1): sizes  
list1[[-5]]: all list elements except 5th  
ls(): list all active objects

mat1[, 2:5] or mat1[, c(2, 3, 4, 5)]: columns from 2nd to  
    5th

matrix(vector1, r1, c1): transform vector1 into matrix with  
    r1 rows and c1 columns, columnwise

merge(df1, df2): merge two data frames  
paste("cow", "boy", sep=""): outputs "cowboy"

rep(x1, n1): repeat vector x1 n1 times  
sample(x1, n1): sample n1 elements from x1 without replacement

seq(n1, n2, n3): sequence from n1 to n2 by n3 steps  
stack() and unstack(): convert from short to long form and  
    back again

str(obj1): structure of object obj1  
t(mat1): rotate 90° matrix or data frame

with(x1, ...): do something within x1

## Cycles, conditions and functions

plot(.., pch=..): 1 ○ 2 △ 3 + 4 × 5 ◇ 6 ▽ 7 ☒ 8 \* 9 ◆ 10 ⊕ 11 ☀ 12田 13⊗ 14▣ 15■

16● 17▲ 18◆ 19● 20● 21○ 22□ 23◇ 24△ 25▽ \* . a a ? ? 0 □

for (i1 in sequence1) dosomething : cycle  
fun1 <- function(args1) dosomething : define function  
if (condition1) ...else ...: single condition  
ifelse(condition1, yes, no): vectorized condition

## Logic and math

is.factor(obj1), is.atomic(obj1), is.data.frame(obj1):  
    check the type of object obj1  
mat1[mat1 > 0]: elements of mat1 which are positive  
!<, &, |, ==: "not less", "and", "or", "equal"  
cumsum(x1); diff(x1); prod(x1); sum(x1): vector math  
round(x1): round  
unique(x1): list unique elements of x1 (could be sparse)  
\*, ^, sqrt(pi), abs(-3), log(1): multiplication, degree,  $\sqrt{\pi}$ ,  
    3, natural logarithm  
x1 %in% x2: which elements of x1 are in x2  
which(logic1): indexes of all TRUE's

## Descriptive statistics

aggregate(...): pivot table  
apply(x1, n1, function): apply function to all rows (if n1  
    = 1) or columns (n1 = 2)  
colSums(mat1): calculate sums of every column  
rev(x1), order(x1), scale(x1), sort(x1): reverse, sorting  
    indexes, scale and center, (ascending) sort  
sapply(); lapply(); do.call(); replicate(): vectorize  
summary(x1); IQR(x1); fivenum(x1); mad(x1); max(x1);  
    mean(x1); median(x1); min(x1); sd(x1); var(x1): de-  
    scriptive statistics  
table(x1, x2): cross-tabulation  
tapply(x1, list1, f1): apply function f1 to x1 grouping by  
    list1

## Inferential statistics

chisq.test(tab1):  $\chi^2$ -test for table tab1  
cor(df1): (Pearson) correlations between all columns of the data  
    frame  
cor.test(x1, x2): (Pearson) correlation test  
ks.test(...); t.test(...), wilcox.test(...): other tests  
lm(...); glm(...); aov(...); anova(...): linear and non-  
    linear models, analyses of variation (ANOVA)  
predict(model1): predict from model  
lm(y ~ x + z, data=...): formula interface to the additive lin-  
    ear model, y responses on two variables, x and z

## Multivariate statistics

dist(...): distance calculation  
cmdscale(...): metric multidimensional scaling (MDA)  
hclust(...): hierarchical cluster analysis  
princomp(...); prcomp(...): principal component analyses  
    (PCA)

## Plots

boxplot(...), dotchart(...), hist(...): useful plots  
identify(...): reveal information from points using mouse  
legend("topleft", legend="..."): add legend to the top left  
    corner  
lines(...); points(...); text(...): add lines, then points,  
    then text  
pdf("file1.pdf"): draw into file1.pdf until dev.off()  
oldpar <- par(mfrow=c(2,1)): plots will be stacked until  
    par(oldpar)  
oldpar <- par(mar=c(0,0,0,0)): all plot margins set to zero  
    until par(oldpar)  
plot(.., cex=1|2): normal dot size, double dot size  
plot(.., col=0|1|2|3): white, black, red, green color  
plot(.., lty=0|1|2): no lines, straight line, dashed line  
plot(.., type="p|l|s|n"): points, lines, stairs and no plot  
qqnorm(vec1); qqline(vec1): check normality