

# R quick reference guide

## Creating a vector

```
x <- 10
```

|                        |             |
|------------------------|-------------|
| a <- c(2, 3, 4)        | 2 3 4       |
| a <- 2:6               | 2 3 4 5 6   |
| a <- seq(2, 3, by=0.5) | 2.0 2.5 3.0 |
| a <- rep(1:2, times=3) | 1 2 1 2 1 2 |
| a <- rep(1:2, each=3)  | 1 1 1 2 2 2 |

## Vector Functions

### sort(x)

Return x sorted.

### table(x)

See counts of values.

### rev(x)

Return x reversed.

### unique(x)

See unique values.

## Using Libraries

```
library(dplyr)
# including the package during the session

dplyr::select
# include any particular function from the package

data(iris)
# loading a built-in dataset into the environment
```

## Selecting Vector Elements

### By Position

**x[4]** The fourth element.

**x[-4]** All but the fourth.

**x[2:4]** Elements two to four.

**x[-(2:4)]** All elements except two to four.

**x[c(1, 5)]** Elements one and five.

### By Value

**x[x == 10]** Elements which are equal to 10.

**x[x < 0]** All elements less than zero.

**x[x %in% c(1, 2, 5)]** Elements in the set 1, 2, 5.

### Named Vectors

**x['apple']** Element with name 'apple'.

## For loop

```
a <- 0
for (i in 0:10){
  a <- a + 1
}
```

## If loop

```
if (i > 3){
  print('Yes')
} else {
  print('No')
}
```

## While loop

```
i <- 0
while (i < 5){
  i <- i + 1
}
```

## Function

```
square <- function(x){
  squared <- x*x
  return(squared)
}
```

## Read and write to file

|                              |                               |  |
|------------------------------|-------------------------------|--|
| df <- read.table('file.txt') | write.table(df, 'file.txt')   | Read and write a delimited text file.  |
| df <- read.csv('file.csv')   | write.csv(df, 'file.csv')     | Read and write a comma separated value file. This is a special case of read.table/write.table. |
| load('file.RData')           | save(df, file = 'file.Rdata') | Read and write an R data file, a file type special for R.                                      |

## Matrices

```
m <- matrix(x, nrow = 3, ncol = 3)
```

Create a matrix from x.



`m[2, ]` - Select a row



`m[, 1]` - Select a column



`m[2, 3]` - Select an element

`t(m)`

Transpose

`m %*% n`

Matrix Multiplication

`solve(m, n)`

Find x in:  $m * x = n$

## Mathematical Functions

|                           |                                 |                          |                         |
|---------------------------|---------------------------------|--------------------------|-------------------------|
| <code>log(x)</code>       | Natural log.                    | <code>sum(x)</code>      | Sum.                    |
| <code>exp(x)</code>       | Exponential.                    | <code>mean(x)</code>     | Mean.                   |
| <code>max(x)</code>       | Largest element.                | <code>median(x)</code>   | Median.                 |
| <code>min(x)</code>       | Smallest element.               | <code>quantile(x)</code> | Percentage quantiles.   |
| <code>round(x, n)</code>  | Round to n decimal places.      | <code>rank(x)</code>     | Rank of elements.       |
| <code>signif(x, n)</code> | Round to n significant figures. | <code>var(x)</code>      | The variance.           |
| <code>cor(x, y)</code>    | Correlation.                    | <code>sd(x)</code>       | The standard deviation. |

## Lists

```
l <- list(x = 1:5)
# create a list of vectors from 1 to 5
```

`l[[2]]`

Second element of l.

`l[1]`

New list with only the first element.

`l$x`

Element named x.

`l['y']`

New list with only element named y.

## Statistics

`lm(x ~ y, data=df)`  
Linear model.

`glm(x ~ y, data=df)`  
Generalised linear model.

`summary`

Get more detailed information out a model.

`t.test(x, y)`  
Preform a t-test for difference between means.

`pairwise.t.test`  
Preform a t-test for paired data.

`prop.test`  
Test for a difference between proportions.

`aov`  
Analysis of variance.

## Dataframe

```
df <- data.frame(x = 1:3, y = c('a', 'b', 'c'))
```

A special case of a list where all elements are the same length.

| x | y |
|---|---|
| 1 | a |
| 2 | b |
| 3 | c |

### List subsetting



`df$x`



`df[[2]]`

Understanding a data frame

`View(df)` See the full data frame.

`head(df)` See the first 6 rows.

### Matrix subsetting

`df[, 2]`



`df[2, ]`



`df[2, 2]`



`nrow(df)`  
Number of rows.

`ncol(df)`  
Number of columns.

`dim(df)`  
Number of columns and rows.

`cbind` - Bind columns.



`rbind` - Bind rows.



```
attach(df_name)
```

```
# this adds the dataframe or the list to R's search path so if you want to access a column
# of the dataframe you don't need to do mean(df$column_name) , you do mean(column_name)
```

```
names() # This method in R can be used in two ways one to set the names of the objects.
# second to get the name of the objects.
```

```
names(x) <- ("name1" , "name2")
```

```
names(df) # returns the column names of the dataframe.
```

## Conditions

|                     |           |                       |              |                        |                          |                         |            |
|---------------------|-----------|-----------------------|--------------|------------------------|--------------------------|-------------------------|------------|
| <code>a == b</code> | Are equal | <code>a &gt; b</code> | Greater than | <code>a &gt;= b</code> | Greater than or equal to | <code>is.na(a)</code>   | Is missing |
| <code>a != b</code> | Not equal | <code>a &lt; b</code> | Less than    | <code>a &lt;= b</code> | Less than or equal to    | <code>is.null(a)</code> | Is null    |