

## R Quick Start Guide

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### R Language Tips:

- Expressions and commands are CaSe-SenSiTiVe
- Anything following the pound character (#) R ignores as a comment.
- Assignments to variables are done with either an '=' sign or the '<-' character sequence.
- Assignment to parameters for function calls are done using the equal (=) sign.
- An object name must start with an alphabetical character, but may contain numeric characters thereafter.
- A period may also form part of the name of an object. For example, x.1 is a valid name for an object in R.  
Spaces, 'x 1' for example, are not valid in an object name.
- **List:** To list the objects you have created in a session enter the commands `objects()` or `ls()` (both do the same thing).
- **Delete:** To remove all the objects you have created, enter the command `rm(list = ls(all = TRUE))`.
- **Save:** To save all the objects you have created in your current session, use the "File" menu (Save As()) or use the `save.image()` command.

### R Basic Commands

- **Help**
  - `Help.search('search term')`: get help when you are not sure of the name of the command or package. (e.g. `help.search('graphics')`)
  - `??'search term'`: get help when you are not sure what the name of the command or package. R will return results related to your search term.
  - `help(command)`: get help on using a certain command when you know the name of the command. (e.g. `help(print)`)
  - `? 'command'`: get the help content on a command (e.g. `?print`);
  - `args(command)`: gets help with the command arguments when you know the name of the command.
- **Demo**
  - `demo('graphics')`: demo the graphics capabilities of R
  - `demo(lm.glm)`: demo the linear modeling capability of R.
  - `example(command)`: Will execute all the example code from the 'command'
- **Data**
  - `X<-5` or `X=5`: Assign a single value to an object.

- `c()` : To create a vector of numbers (e.g. `my.vector <- c(1, 2, 3, 4, 5)`) Can also be used to create a vector of character strings (e.g. `c("cat", "dog", "fish")`) or True/False values (e.g. `c(T, F, T, T)`)
- `:` or `seq()` : To create a sequence of number use the colon `:` or the `seq()` command (e.g. `my.sequence<-0:10` or `my.sequence2<-seq(0,10, by = 2)`)
- **Named Vector:** Create a named vector (e.g. `aNamedVector <- c(type =1, count =7 ,max =10)` or `bNamed Vector <- c("a type" = 1, "b%type" = 7)`
  - The values can be integer, numeric, character, logical (T/F) or complex values
- `[]` : To reference specific elements use square brackets (`blob<=foo[4]`)
  - Element numbering starts with 1 not 0. (`my.vector[3]=3`)
  - Use the name of the element (e.g. `aNamedVector[["type"]]` or `aNamedVector[c("type", "max")]`)
- `indicator`: create a vector of TRUE, FALSE values based upon a conditional (e.g. `indicator <- y <4`; Which values in the vector `y` are less than 4?)
- `rep(value, number of repeats)` : repeat function (e.g. `rep(1:5, 2)` repeat the vector 1 to 5 twice)
- `%in%` : Find a specific value (substring) in another string (e.g. `y %in% c("cat", "dog")`)
- `!` (Negation) : Negate a T/F vector (e.g. `!z`)
- `&` (AND) : 'And' the values of at T/F vector (e.g. `x & y`)
- `|` (OR) : 'OR' the values of a T/F vector (e.g. `x | y`)
- **Quotes:** R will accept both double and single quotes for all character strings. It is a way of embedding quoted values within another string. (e.g. `x<- "A String with 'inner quotes'"`)
- `print(y)` : print the contents of the variable `y` to the screen
- `cat()` : prints text to the screen (e.g. `cat("Here is the value of ", x, "\n")`)
- `load('filename')` : Load a data file from local directory.
- `attach(dataset)` : load named data set (`attach(mtcars)`)
- `str(y)` : show the structure of a variable `y`
- `length(y)` : show the number of elements in `y`
- `head(y)` : show the first few members of the variable `y`
- `tail(y)` : show the last few members of the variable `y`
- `which(y)` : returns the indices of a T/F vector which are TRUE
- `any(y)` : returns TRUE if any of the values of the T/F vector are TRUE
- `all(y)` : returns TRUE if all of the values of a T/F vector are TRUE
- **Dataframe:** A basic unit of data in R is the data frame. It is typically a matrix of data with named columns much like an Excel spreadsheet
  - Example: The `mtCars` data set. It has the following columns:
    - `mpg, cyl, disp: num, hp, drat, wt, qsec, vs.am, gear, carb`
- **Column Reference:** To reference a named member of a data frame use the dollar sign (`$`)

- e.g. `mtCars$mpg`, `mtCar$qsec`.
- `nrow(dataset)` : Display the number of rows in a named data frame (e.g. `nrow(mtcars)`)
- `ncol(dataset)` : Display the number of rows in a named data frame (e.g. `ncol(mtcars)`)
- `nchar(x)` : Count the number of characters in the character vector `x`
- **Matrix:** A matrix is similar to a dataframe except that all the elements in a matrix must be the *same* type but the type does not matter (integer, numeric, character, logical, or complex)
  - Matrices are created using the `matrix()` command. (e.g. `m1<-matrix(1:15, nrow = 5, ncol =3)`). They are loaded column-wise by default
  - All the standard matrix operations are available in R including transpose (`t(m1)`), Inner product, outer product and multiplication (all use `'%*%'`; R returns the correct value depending upon the data type.), inversion (`solve(m1)`), eigenvectors (`eigen(m1)`) and determinant (`deter(m1)`)
- It is possible to convert from a formal matrix to a dataframe (and back) as long as there is only one datatype.
- `rownames(m1)` : Extract the row names from a matrix
- `colnames(m1)` : Extract the column names from a matrix
- **Lists:** A list is a collection of R objects which do not have to be of any particular type or size and can even be other lists.
  - Lists are created using the `list()` command. (e.g. `aList <- list(a = 1:5, b = rep(TRUE, 2), c = letters[1:3])`)
- `names(l1)` : Extract the names from a list
- **Basic Calculator Commands**
  - Addition: `'+'` sign (e.g. `7+3`)
  - Subtraction: `'-'` sign (e.g. `7-3`)
  - Multiplication: `'*'` sign (e.g. `7*3`)
  - Division: `'/'` sign (e.g. `7/3`)
  - Integer Division: `'%/%'` (e.g. `7%/% 3`)
  - Division Remainder: `'%%'` (e.g. `7%%3`)
  - Exponentiation: `exp(#)` (e.g. `exp(7)` = the constant `e` = 2.718282.. raised to the power of 7 )
  - Natural Logarithm: `log(#)` (e.g. `log(7)`)
  - Base 10 Log: `log10(#)` (e.g. `log10(1000)`)
  - Square Root: `sqrt(#)` (e.g. `sqrt(16)`)
  - Cosine: `cos(radian)` (e.g. `cos(pi)` where `pi` is a defined constant = 3.141519...radians)
  - Large Numbers: `#e#` (e.g. `1.7e+05`)
  - Small Numbers: `#e#` (e.g. `1.7e-03`)
  - Division by 0: `#/0` (e.g. `1/0 = NaN`)

- **Basic Statistical Commands**
  - `^`: raise to the power (e.g. `y<-x^2` ( $y = x^2$ ))
  - `mean(y)`: calculate and show the mean of the vector `y`
  - `var(y)`: calculate and show the variance of the vector `y`
  - `sd(y)`: calculate and show the standard deviation for a vector `y`
  - `sum(y)`: calculate and show the sum of the vector(`y`) for a numerical vector or the number of true values in a T/F vector `y`
  - `min(y)`: calculate and show the minimum of a vector `y`
  - `max(y)`: calculate and show the maximum of vector `y`
  - `quantile(y, .25)`: calculate the 25<sup>th</sup> percentile of a vector `y`
  - `quantile(y, 0.75)`: calculate and show the 75<sup>th</sup> percentile of the vector `y`
  - `summary(y)`: calculate and show basic statistical measures of the variable `y`
  - `lm()`: fit a linear regression model to the data (e.g. `lm_1<- lm(y ~ x)`  $y = f(x)$ )
- **Input/Output**
  - `read.csv()`: read a comma-separated-value (csv) file (e.g. `read.csv("nf-week2-sample.csv")`);
  - `save()`: Save an R binary object to disk for use later in the session or a new session. Used primarily for objects which take some time to compute. (e.g. `x <- runif(20); y <- list(a = 1, b = TRUE, c = "oops"); save(x, y, file = "xy.Rdata")`)
  - `unlink()`: Remove a previously saved R binary data object from disk (e.g. `unlink("xy.Rdata")`)
- **Library Commands**
  - `install.packages("package")`: install for use in the working directory a new R package (e.g. `install.packages("ggplot2")`);). To install a new R package for use in specified location use the argument *lib* to specify such as `install.packages("package", lib="location")` (e.g. `install.packages("ggplot2", lib="/data/Rpackages/")`);
  - `library(library)`: loads library for use (e.g. `library(lattice)`);
  - `view(package)`: Show contents of package (e.g. `view(mtcars)`)
  - `attach(package)`: Load package (e.g. `load(mtcars)`);
- **Directory Commands**
  - `ls()`: Show all variables in the current environment
  - `dir.create('location')`: Create a local directory (e.g. `dir.create("data/artifacts")`)
  - `dir('directory')`: look at the contents of a directory
  - `dir()`: look at the contents of the current working directory
  - `setwd('directory')`: set working directory
  - `getwd()`: get working directory
  - `source('script_file.R', echo = TRUE)`: run an entire R script file and show the commands and the results in the console window

- **Plotting**
  - `hist()`: make a histogram plot (e.g. `hist(mtCars$mpg)`)
  - `plot()`: plot an x vs y plot (e.g. `plot(mtCars$mpg, mtCars$cyl)`)
  - `ggplot2()`: plotting command from the `ggplot` package (see `??ggplot` for details).
- **Functions**
  - Functions in R have inputs (arguments) and outputs(values)
  - Arguments can be named or un-named
  - Examples:
    - Simple: `simple<- function(x,y) { return x+y-7) }`
    - Named argument: `simpl <- function(x, method = "sum") {if (method == "sum") {out <- sum(x)}else if (method == "prod") {out <- prod(x)}else {stop("'method' must be 'sum' or 'prod'") }return(out)}`

## Web Sites:

- Introduction to R, CRAN: <http://cran.r-project.org/doc/manuals/R-intro.html>
- R for Beginners: [http://cran.r-project.org/doc/contrib/Paradis-rdebuts\\_en.pdf](http://cran.r-project.org/doc/contrib/Paradis-rdebuts_en.pdf)
- R Tutorial: <http://www.r-tutor.com/>
- RStudio: <http://www.rstudio.com>
- CRAN: <http://cran.us.r-project.org>
- DataDR: <http://tesseractdata.org/datadr/>
- Trelliscope: <http://tesseractdata.org/trelliscope/>
- Bootcamp code and description: <http://tesseractdata.org/docs-r-intro-bootcamp/>
- VAST Challenge Data exploration <http://tesseractdata.org/example-vast-challenge/>