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Introduction to R



What is R?

As stated in the [manual for beginners](#) found at the **C**omprehensive **R**Archive **N**etwork, **R** is

- An effective data handling and storage facility,
- a suite of operators for calculations on arrays, in particular matrices
- a large, coherent, integrated collection of intermediate tools for data analysis,
- graphical facilities for data analysis and display either directly at the computer or on hard-copy,
- a well developed simple and effective programming language (called **S**) which includes conditionals, loops, user defined recursive functions and input and output facilities



What is R?

- It is **free** and available for all major OS.
- **R** is an **interpreted** language.
- **R** itself is a rather 'modern' **GNU** implementation of **S** language, which in turn was developed by *Bell Laboratories*. The first stable beta version was released around 2000.
- Part of **R** functions are written in **R** itself and make it easy to understand, part are written in **C**, **FORTRAN** and other languages for the sake of the performance (mostly computational parts). There is a toolchain for building **R** \leftrightarrow **C** interoperable libraries.
- **R** can be extended through *packages* that can be installed from different centralized repositories (**CRAN**, **MRAN** and mirrors), directly from directories on local machine, from ^{*}**git** repos and from many other sources.



Basic concepts

- There are several base types in R, **integer**, **double**, **character**, **complex**, **factor** and **logical**. It is safe to say that everything in R is a first-class object
- The simplest collection is a **vector** - a group of objects of the same type. Any standalone primitive object (like number 2.0) is itself a **vector** of size **1**.
- Matrices are arrays of higher order (rectangular), lists are general collections of any objects of any types, data frames are data-base like tables with columns of equal length but different type. Another type, part of the **tidyverse** package, **tibble** extends functions of data frames.
- Higher order types can be defined using class systems (I am aware of several, **S3**, **S4**, **Reference classes** and **S6**)



Examples: Run interactively

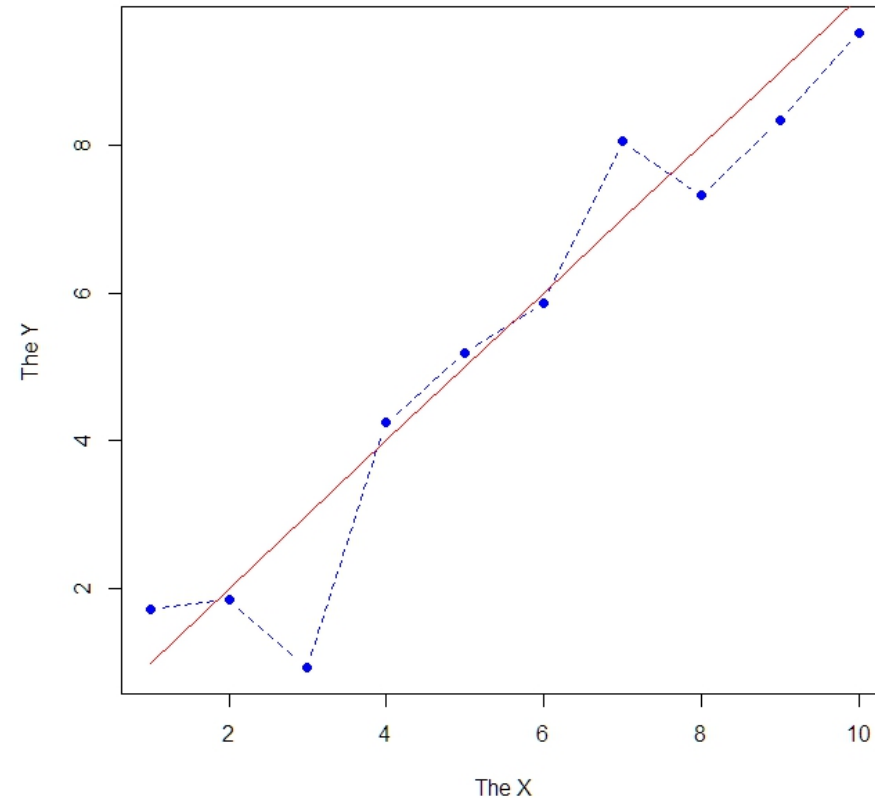
```
> x <- 5 # Assigns 5 (double) to x
> x      # Prints x, equivalent to print(x)
# [1] 5
> x[1]   # Accesses 1st element of a vector
# [1] 5
> y <- list(x, "Hello world") # A list
> y
# [[1]]
# [1] 5
#
# [[2]]
# [1] "Hello world"
> a <- 1:7 # Integer range from 1 to 7
> a ^ 2    # Vectorized power operation
# [1] 1 4 9 16 25 36 49
> mean(a ^ 2) # Calls built-in stat. mean
# [1] 38.5
> sd(a ^ 2) # Standard deviation
# [1] 34.17358
```

```
> sprintf("Compare %3.2f to %3.2e", 0.0123, 0.0123)
# [1] "Compare 0.01 to 1.23e-02"
> quantile(rnorm(n = 100, mean = 0, sd = 3),
+ probs = c(0.16, 0.84))
#          16%          84%
# -3.191771  2.949907
> f <- function(x) {
+ mn <- mean(x)
+ s <- sd(x)
+ rng <- range(x)
+ return(c(mn, s, rng))
+}
> f(c(10, 20, 30, 123, 0.06, -100))
# [1] 13.84333 71.24490 -100.00000 123.00000
> z <- 0
> for (i in 1:100) z <- z + i; print(z)
# [1] 5050
> sum(1:100)
# [1] 5050
```



Examples: Get help & make a plot

```
> f # Type the name of previously defined func
# function(x) {
# mn <- mean(x)
# s <- sd(x)
# rng <- range(x)
# return(c(mn, s, rng))
# }
> ?mean # Opens local web page with manual
> data <- data.frame(x = 1:10,
+ y = 1:10 + rnorm(10))
> plot(data$x, data$y, type = "b",
+ pch = 19, lty = 2, col = "blue",
+ xlab = "The X", ylab = "The Y")
> lines(1:10, 1:10, lty = 1, col = "#FF0000")
> data[1:2, ]
#   x     y
# 1 1 1.721106
# 2 2 1.848209
> quit("no")
```





Examples: Install **JAGS** and get some packages

- **JAGS** can be found [here](#)
- We need **rjags** to run **JAGS** from **R**, **tidyverse** (optional) to use fancy data manipulation facilities.

```
> install.packages(c("rjags", "tidyverse")) # Installs two packages  
> library(rjags)      # Loads rjags, can be used without quotes  
> library(tidyverse) # All useful data processing tools
```

- Now check out code examples (demo)



More useful examples

Assignment operators:

```
<- and -> are default operators  
= is used with named func. arguments  
<<- and ->> are global assignment ops.  
> getwd() # Returns current working directory  
# [1] "\path\to\current\dir"  
> setwd("\path\to\new\dir") # Sets work dir
```

Loop alternatives

```
apply, lapply, sapply, vapply  
> sapply(1:5, function(x) (x + 1) ^ 2)  
# [1] 4 9 16 25 36
```

```
> library(foreach)  
> foreach(i = 1:2) %do% { i ^ 2}  
# [[1]]  
# [1] 1  
#  
# [[2]]  
# [1] 4  
> library(parallel)  
> library(doSNOW)  
> cl <- makeCluster(2, "SOCK")  
> registerDoSNOW(cl)  
> foreach(...) %dopar% {...}  
> stopCluster(cl)
```

```
> file.path("dir1", "dir2")  
# "dir1/dir2"
```




More useful examples

```
> paste("d", c("x", "y", "z"), sep = "_")
# [1] "d_x" "d_y" "d_z"
> substring("Hello World!", 7, 11)
# [1] "World"
> grepl("[W|w]o.*!$",
+ c("World!", "world!", "world"))
# [1] TRUE TRUE FALSE
> c(5 / 2, 5 %% 2, 5 %/% 2)
# [1] 2.5 1.0 2.0
> c(1, 5, 11) %in% (1:10)
# [1] TRUE TRUE FALSE
> setNames(c(1, 2, 3), c("a", "b", "c"))
# a b c
# 1 2 3
> order(c(20, 50, 10, 90, 30))
# [1] 3 1 5 2 4
```

```
> sink("filename.dat") # Output goes to file
> print("Hello World!")
# Nothing in terminal, line appears in file
> sink() # Closes last sink
> pdf("file.pdf", width = 7, height = 5)
> plot(1:5, 5:1) # Plot goes into pdf file
> dev.off() # Closes current device !important!
> tryCatch(stop("Sample error"),
+ error = function(e) print("Err handled"),
+ finally = print("Cleanup"))
# [1] "Cleanup"
# [1] "Err handled"
```



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Thank you!