

Quick Introduction to R

All variables, data, functions, results etc. are stored as so-called *objects* which have their own *names* and other *attributes* such as length, dimensions (e.g. in the case of matrices), names of columns (e.g. in the case of data frame) etc. ... In R we do not declare objects - they are created at the moment of their first appearance and they have a type accordingly to the type of the assigned value.

- **Types of data (modes):**

`numeric`
`character` - values: strings in single or double quotes
`complex`
`logical` - values: `FALSE`, `TRUE`

- **Some data structures (objects):**

- **vector**

Creating a vector:

`c(...)` - combines its arguments to form a vector,
`:` - colon operator, creates a regular sequence with an increment equal to 1, e.g. `1:4` is equal to `c(1,2,3,4)`,
`seq()` - generates a regular sequence,
`rep(x,n)` - creates a vector of n elements equal to x ,
`vector()` - creates a vector.

Subsetting a vector:

`a[1]` - first element of vector `a`,
`a[c(1,2,5)]` - first, second and fifth elements of vector `a`,
`a[a>10]` - the elements of vector `a` that are greater than 10.

- **factor** - categorical variable.

Creating a factor: `factor()`.

Function `levels()` provides information about all possible values (`levels`) of the factor.

- **array** - table. Special case of table is matrix (`matrix`) - two-dimensional table.

All elements of vectors, factors and arrays are of the same type.

- **data frame** - table composed with one or several vectors and/or factors all of the same length but possibly of different modes.

Useful functions:

`data=data.frame(parameters)` - creates data frame named `data`,
`data(data_name, package="package_name")` - loads data frame from R library,
`data=read.table("file", header=FALSE, col.names=c(V1,V2,...), ...)` - reads a data frame from a file outside R and names it `data`. Here default values of some parameters are given.
`dim(data_frame)` - dimensions of a data frame,

Subsetting data frame:

`a[,1]` - vector made of elements contained in the first column of data frame `a`,
`a$V1` - column (vector) named `V1` of a data frame `a`,
`a[, 'V1']` - data frame containing column `V1` of data frame `a`,
`attach(data_frame)` - gives a direct access to names of variables in a data frame.
`detach(data_frame)` - detaches data.

- **list** - contains elements/objects of any type. Elements of a list may be for example: statistical models, functions, expressions...

Creating a list: `list()`,

Subsetting a list: `a[[2]]` - second element of a list `a`,

`a$b` - element named `b` of a list `a`.

Function `names()` gives names of the elements of a list.

Example: function `lm()` builds a linear model and returns an object which is a list containing elements such as: `coefficients`, `residuals`, `fitted.values`, `df.residual`, ...

- **Random numbers:**

Normal distribution:

`rnorm(n, mean, sd)` - random generation of n numbers

`dnorm(x, mean, sd)` - density function

`pnorm(p, mean, sd)` - distribution function,

`qnorm(q, mean, sd)` - quantile function,

Exponential distribution:

`rexp(n, rate)` - random generation of n numbers

`dexp(x, rate)` - density function

`pexp(p, rate)` - distribution function,

`qexp(q, rate)` - quantile function,

And so on...

- **Numerical summaries:**

`summary()` - provides basic numerical summaries of data such as minimal and maximal values, mean, median, quantiles, ...

`mean()` - mean value of vector,

`median()` - median,

`range()` - returns a vector containing the minimum and maximum of all the given arguments,

`quantile()` - sample quantiles corresponding to the given probabilities,

`min()`, `max()` - minimal and maximal value of all the given arguments,

`var(x)`, `sd(x)`, `cov(x,y)`, `cor(x,y)` - computes the variance of x , standard deviation of x and the covariance or correlation of x and y if these are vectors.

- **Graphics:**

`hist()` - draws a histogram of the given data,

`boxplot()` - produce box-and-whisker plot(s) of the given (grouped) values,

`plot(x,y,type="p",...)` - scatterplot,

`curve(function_name,xlim=c(a,b))` - draws a curve defined as a function in the range of argument from a to b ,

`points(x,...)` - draws a sequence of points at the specified coordinates,

`lines(x,...)` - joins the corresponding points with line segments,

`abline(a,b)` - adds a line with intercept a and slope b to the current plot,

`qqnorm(x)` - produces a normal QQ plot of the values x ,

`par(mfrow=c(2,2))` - divides a graphics window into two rows and two columns (enables to produce several plots in one display),

`get("windows")()` - opens a new graphics window (without closing the current one).

- **Useful commands:**

`source("file_name")` - loads a file containing source code,

`getwd()`, `setwd()` - gives the name of working directory, changes working directory,

`library(package_name)` - loads a package,

`?name`, `help(name)`, `help("name")` - help about object named $name$

`as.numeric(object_name)`, `as.logical(object_name)`, `as.character(object_name)`, ... - conversion functions

`table(V1,V2)` - creates contingency table for variables `V1` and `V2`.

`sort()` - orders data,

`NA` - Not Available (missing data), `Inf` - Infinity, `NaN` - Not a Number