

Biostatistics I: Introduction to R

Introduction

Eleni-Rosalina Andrinopoulou

Department of Biostatistics, Erasmus Medical Center

✉ e.andrinopoulou@erasmusmc.nl

🐦 [@erandrinopoulou](https://twitter.com/erandrinopoulou)

Introduction to R

- ▶ **R** is a great tool to explore and investigate the data
- ▶ Several statistical methods can be performed with **R**
- ▶ It is important to understand the methods before applying them in **R**

How to use

R uses packages that perform specific tasks

- ▶ Install package only once
- ▶ Load package every time you open **R**

Introduction to R

- ▶ For this course: Rstudio (<http://www.rstudio.org/>)
 - ▶ free
 - ▶ works fine in Windows, MacOS and Linux
 - ▶ helpful with errors
 - ▶ alternative output options

Introduction to R

Basic functions

- ▶ `getwd()`, `setwd()`,
- ▶ `is.na()`,
`is.finite()`,
`is.null()`

Import/Export

- ▶ `read.csv()`, `write.csv()`
- ▶ `read.xlsx()`, `write.xlsx()`
- ▶ `read.table()`, `write.table()`

Save/Load

- ▶ `save()`, `saveRDS()`
- ▶ `load()`, `readRDS()`

Data Types/Structures

The simplest data types are:

- ▶ **numeric** : quantitative data
- ▶ **character** : qualitative data
- ▶ **integer** : whole numbers
- ▶ **logical** : TRUE or FALSE
- ▶ **factors** : qualitative data (levels)

Data Types/Structures

The most important data structures are:

- ▶ **Scalar** a single element
- ▶ **Vectors** have the same type of elements
- ▶ **Matrices** have the same type of elements with the same length
- ▶ **Arrays** have the same type of elements with the same length but can store the data in more than two dimensions
- ▶ **Data frames** have elements of different type with the same length
- ▶ **Lists** have elements of different type and length

Data Types/Structures

Data types

- ▶ `is.numeric() / as.numeric()`
- ▶ `is.character() / as.character()`
- ▶ `is.integer() / as.integer()`
- ▶ `is.logical / as.logical()`
- ▶ `is.factor() / as.factor()`
- ▶ `str(), mode()`

Data structures

- ▶ `c()`
- ▶ `matrix()`
- ▶ `array()`
- ▶ `data.frame()`
- ▶ `list()`

Other

- ▶ `ls(), objects()`

Indexing/Subsetting

- ▶ This can be done using square bracket (**[]**) notation and indices.
- ▶ Three basic types
 - ▶ position indexing
 - ▶ logical indexing
 - ▶ name indexing

Indexing/Subsetting

Vectors

- ▶ `[]`
- ▶ `[""]` - for categorical variables

Matrices

- ▶ `[,]`
- ▶ `[[]], []`

Arrays

- ▶ `[, ,]`

Data frames

- ▶ `[,]`
- ▶ `[[]], []`
- ▶ `$`

Lists

- ▶ `[]`
- ▶ `[[]]`
- ▶ `$`

Data Transformation/Exploration/Visualization

Transformation

- ▶ `round()`
- ▶ `factor()`
- ▶ `order()`
- ▶ `reshape()`

Exploration

- ▶ `mean(), sd()`
- ▶ `median(), IQR()`
- ▶ `table()`

Visualization

- ▶ `plot(), legend()`
- ▶ `hist()`
- ▶ `barchart()`
- ▶ `boxplot()`
- ▶ `xyplot(), ggplot()`
- ▶ `par()`

Correlation

Pearson correlation

- ▶ magnitude of association
- ▶ linear association
- ▶ direction of the relationship

A relationship is linear when a change in one variable is associated with a proportional change in the other variable

Correlation

Spearman correlation

- ▶ direction of the relationship
- ▶ monotonic relationship

In a monotonic relationship, the variables tend to change together, but not always at a constant rate (as in the linear case)

Test hypothesis

- ▶ **parametric** (assumptions about the distribution) / **non-parametric** (distribution-free)
- ▶ **one sample** / **two samples** / .. / **M samples**
 - ▶ compare one group with a value
 - ▶ compare two groups paired / unpaired
 - ▶ compare M groups
- ▶ **one-sided (one-tailed)** / **two-sided (two-tailed)**

$$H_0 : \theta = \theta_0$$

$$H_1 : \theta \neq \theta_0 \text{ (two-sided)}$$

$$H_1 : \theta > \theta_0 \text{ (one-sided)}$$

$$H_1 : \theta < \theta_0 \text{ (one-sided)}$$

Test hypothesis

- ▶ Choose a null hypothesis H_0 and an alternative hypothesis H_1
- ▶ Collect and visualize the data
- ▶ Choose and calculate the test statistic, which is a numerical summary of the data
- ▶ Determine the sampling distribution under the condition that the null-hypothesis holds
- ▶ Choose the type I error (significant level) α , usually $\alpha=0.05$
- ▶ Determine the corresponding critical value(s)
- ▶ Compare the test statistic with critical value(s) and reject or not