

# Introduction to R

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Biomedical Data Sciences

# Statistical programming language

- ▶ A language for data analysis and graphics
- ▶ Ross Ihaka and Robert Gentleman (1993), based on the statistical programming language S (Chambers, 1976)
- ▶ Currently maintained by a large groups of primarily statisticians all over the world

# Open source

- ▶ Supports various operating systems
- ▶ Command-line and many graphical user interface, i.e. RStudio
- ▶ Easily extendable using packages available at CRAN
- ▶ Computationally-intensive tasks can be written in C, C++ and Fortran code
- ▶ Large and active community; R journal, Annual conference, specialist mailing list, etc.

# Many application domains

- ▶ Econometrics, Genetics, Pharmacokinetics, Social Sciences, Bioinformatics, etc
- ▶ Can handle almost any data format: .xlsx, .sav, html, xml, images, binary data formats, connect to databases, etc.
- ▶ Can generate figures in any format: png, jpg, pdf, etc.
- ▶ Can generate reports in any format: pdf, markdown, html, docx, etc.
- ▶ Easy parallelization and efficient reading of large data

# Bioconductor

- ▶ Repository specific for bioinformatic analysis
- ▶ Expression, DNA methylation, copy number, proteomics, metabolomics, genetics, etc.
- ▶ Not only methods but annotation and data as well
- ▶ > 1000 interrelated packages
- ▶ High quality software (reviewed), daily build system (guarantees that software works) and obligatory documentation
- ▶ Mailing list, example workflows, course material, etc.

## Some disadvantages

- ▶ Syntax not always intuitive
- ▶ Dynamic language; continuously changing (often more efficient and additional features)
- ▶ Biannually new releases (incl. bug fixes) so keep uptodate
- ▶ Many solutions for the same problem often one robust and most efficient (requires experience)
- ▶ Many functions for problems (use Google to find them)

## Some nice examples

- ▶ R graph gallery: <http://www.r-graph-gallery.com/>
- ▶ R web application: <https://shiny.rstudio.com/gallery/>
- ▶ Bioconductor workflows:  
<https://www.bioconductor.org/help/workflows>

# Learning R

- ▶ Online course material, for example:  
<https://www.datacamp.com/>
- ▶ Books
- ▶ R course LUMC
- ▶ Practical (brief introduction)