INTRODUCTION TO R

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COURSE OVERVIEW

- 2-day Introduction to R aim to get everyone at the same level
- Two lectures today: Introduction to R and RStudio Cloud set up this and Discord
- Three practicals

1

- reproducible work
- importing data
- data manipulation

2

- data visualization
- analysis
- writing functions

3

- bioconductor
- genomic ranges
- basic analysis

WHAT IS R?

- Different from most statistical software you can't sidestep the blinking cursor
- Developed by Ross Ihaka and Robert Gentlemen
- Language for statistical analysis and graphics
- Inspired by the programming language, S
- Maintained by R Development Core Team
- Large group of primarily statisticians



Source - R. Ihaka

- Free especially attractive compared to other software
- No hidden costs

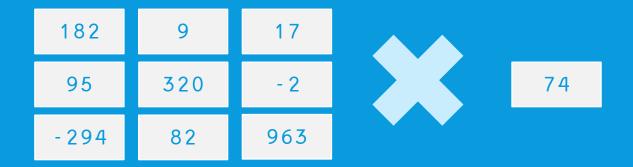
- Open source source code can be viewed and modified by anyone
- Actively maintained
- Stable and reliable
- Instead of relying on a small team of developers
- Bugs and issues can be detected and resolved far quicker

- Compiles and runs on multiple operating systems
- Can even port it to different hardware
- Large and active community
- Happy to provide support and information
- Scripting language write your own functions
- Novel methods quickly developed into packages by users
- Versatile and stays up-to-date other programs may never get niche methods

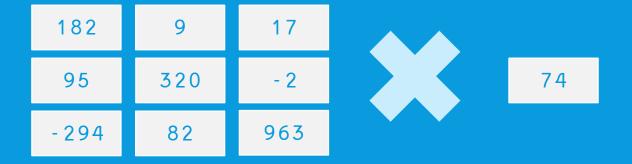


- Integrative plays well with others
- Individuals extended R to nest into their current workflows
- Can read in multiple data types html, sav, databases, IDAT
- Implement other languages C, Python, Java no need to import / export
- Output many file formats markdown, sav, pdfs

- Easy parallelization meaning faster calculations
- Optimized for **vector** operations better than scalar languages or loops

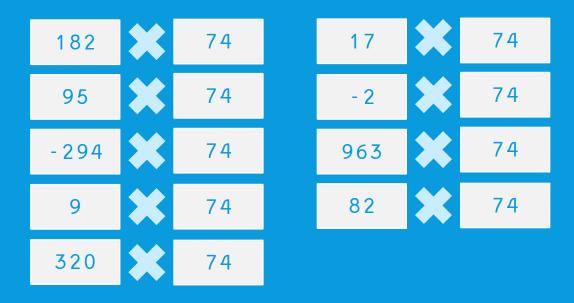


• CPU: 1 instruction per cycle, 3 operands per instruction



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- Scalar language or loop:

9 instructions = 9 cycles



- CPU: 1 instruction per cycle, 3 operands per instruction
- Vectorized language:

1 instruction

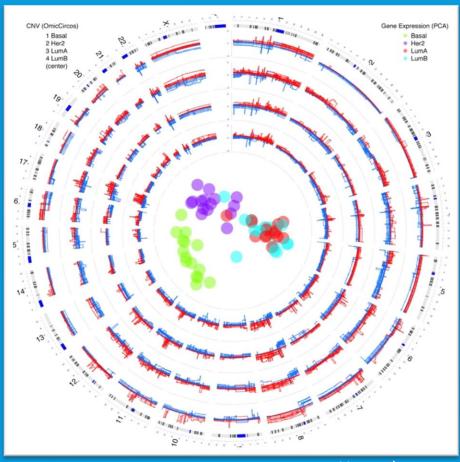
3 times as fast!

[182, 95, -294, 9, 320, 17, -2, 963, 82] 9 operands = 3 cycles [182, 95, -294] [9, 320, 17]

[182, 95, -294]

- Stunning graphics e.g. geo-spatial data
- Bioinformatics packages e.g. OmicCircos
- Used by Google, Twitter, and Facebook
- Build interactive web applications with Shiny





THE DISADVANTAGES OF R

- Working in R can be quite frustrating
- Syntax unintuitive and inconsistent
- Steep learning curve command-line type user interface
- Requires experience multiple options for any problem
- Continuously changing need to stay up-to-date
- Nothing to stop packages using the same named functions
- No official support sometimes poor documentation



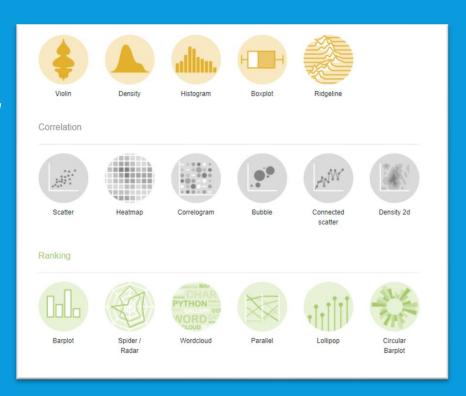
BIOCONDUCTOR



- Software repository over 1,500 bioinformatics packages
- · Analyse, annotate, and visualize high-throughput -omic data
- Have some quality standards
- Obligatory vignettes high quality documentation
- Daily build system guarantees software works
- Basics introduced in the 3rd practical

RESOURCES

- R graph gallery: http://www.r-graph-gallery.com/
- R web application: https://shiny.rstudio.com/gallery/
- Bioconductor: https://bioconductor.org/
- Online courses: https://www.datacamp.com/
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