

BMS 353

Bionformatics for Biomedical Sciences

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An introduction to the tools we will be using

R vs the old way

R is a free language and environment for statistical computing and graphics

Very expensive
Popular with industry



Not surprisingly R is far more popular now! http://r4stats.com/articles/popularity/

The number of jobs requiring R has increased too!



Packages

These save time by using other people's code

- Comprehensive R archive Network (CRAN)
 9662 packages (November 2016)
 https://cran.r-project.org
- Bioconductor 1211 packages (November 2016) http://bioconductor.org/

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Jupyter notebooks





Combine live computer code, data, tetx, and mathematics in an interactive document.



Academic papers only gve a description of your analyis.

Jupyter notebooks contain both the description and the analysis.



- Collaborative computational mathematics
- All open source environment for running Jupyter notebooks (and others) in the cloud
- Dedicated Virtual Machine for this course on the Google Cloud Platform
- As powerful as a node on the Sheffield University Supercomputer: Iceberg

Log into CoCalc https://cocalc.com



Handy tips for the notebooks

Ctrl/Cmd + y changes the cell to code mode

Ctrl/Cmd + m changes cell to markdown

Shift + Enter evaluates the cell

To add Latex in a markdown cell make sure to enclose in \$\$ or \$

Also, we put together a **mini** CoCal tutorial that you can check at https://tutorial.cocalc.com/students/