Special Topics in Biostatistics and Bioinformatics Meet-and-greet

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3 March 2022



Ege Ülgen

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- Office: Block A, 6th floor (Dept. of Biostatistics and Medical Informatics)
- Office hours: by appointment

Lecture Notes and Scripts

Available on GitHub:

https://github.com/egeulgen/BB512 BB612 21 22

Syllabus

Week 1 – Introduction

Week 2 – Exploratory Data Analysis

Week 3 – Dimension Reduction/Preprocessing and Normalization

Week 4 – Review of HW1

Week 5 – Batch effects and Confounders

Week 6 – Inference

Week 7 – Review of HW2

Syllabus (cont.)

Week 8 – Gene Set Enrichment

Week 9 – Gene Expression Analysis

Week 10 – Review of HW3

Week 11 – Next Generation Genomics Sequencing Analysis

Week 14 – Review of Final Projects

Course Evaluation

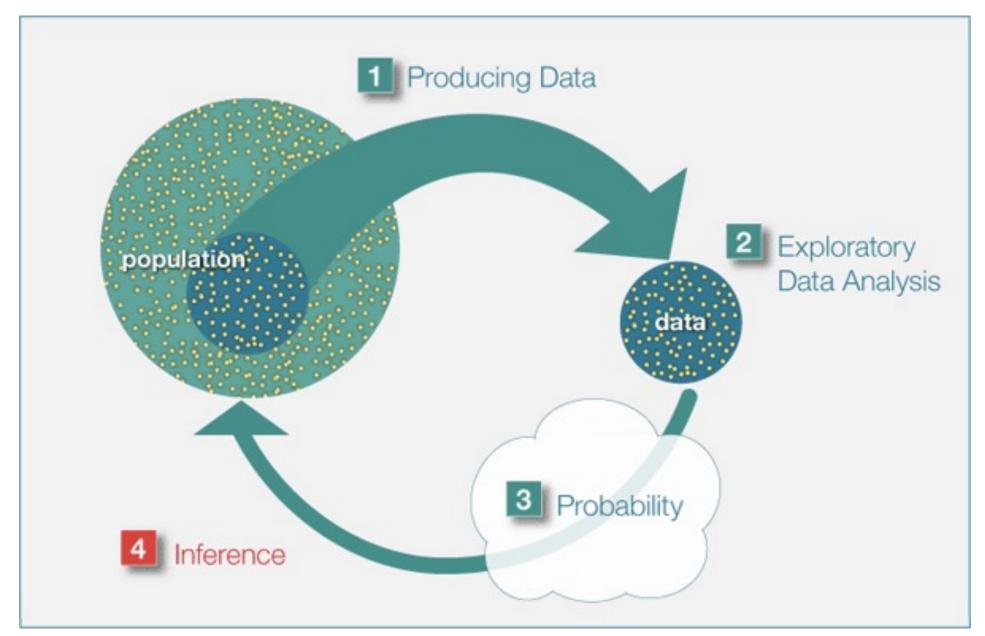
- Homeworks 30%
- Final Project 70%

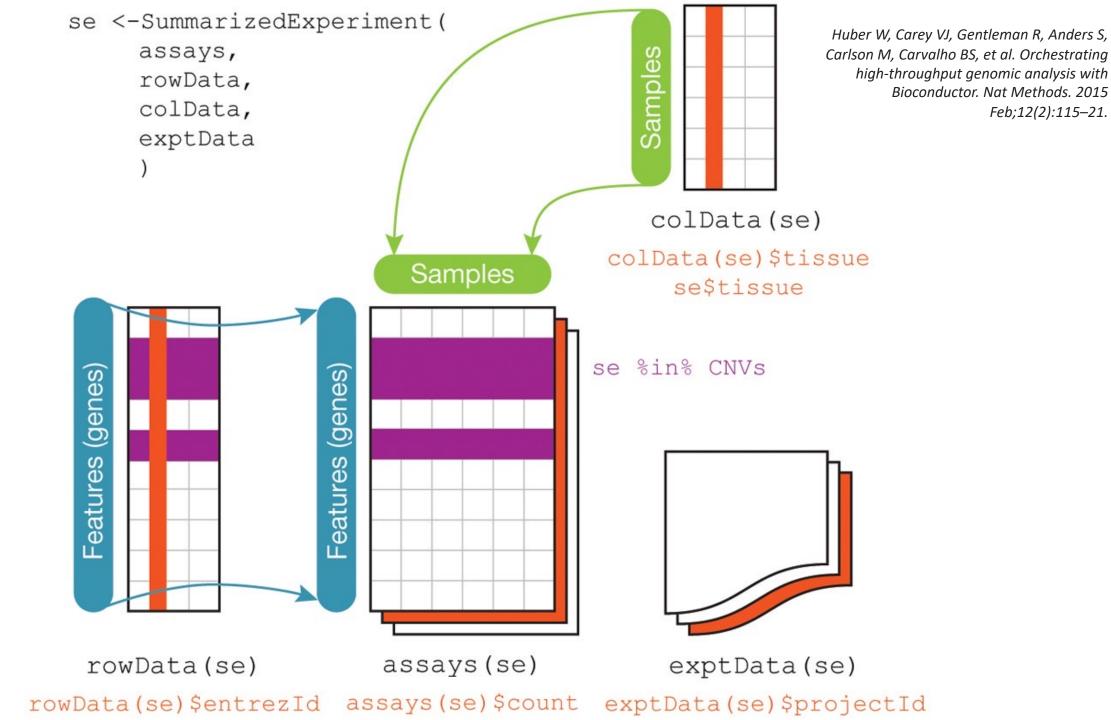
Software





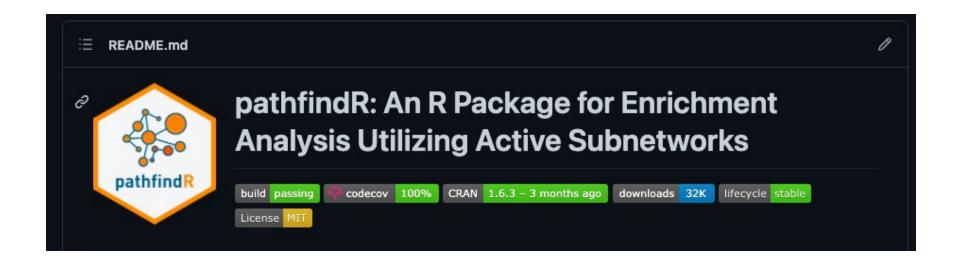
Some Introduction





Where to find trustworthy methods

- Articles
- GitHub
- Forums
- •



Where to get help

- If you can't figure out how to solve an issue, Google is your friend. e.g., "how to calculate empirical p values"
- If you get an error and cannot fix it. C/P the error into Google.
 Someone else most likely had a similar problem

- Some good sources for asking about bioinformatics issues:
 - Biostars https://www.biostars.org/
 - Bioconductor Support https://support.bioconductor.org/
 - Stack overflow https://stackoverflow.com/

Reproducibility

- Clean data
- A document with
 - variable names
 - descriptions
 - units ...
- (preferably Rmd or similar) Code to analyze raw input data and generate the output
 - With clear descriptions/comments

«Clean» Data

country	year	cases	population
Afghanstan	100	45	18.57071
Afghanistan	2000	2666	20!95360
Brazil	1999	37737	172006362
Brazil	2000	80488	174!04898
China	1999	212258	1272915272
Chin	200	21 66	1280 28583
variables			





values