

DATA ANALYSIS FOR MOLECULAR BIOLOGY AND BIOCHEMISTRY



Introductory data analysis focusing on molecular biology data sets and examples and including basic programming skills using Python and basic statistics skills using R. Prerequisite: MATH 12 or equivalent is recommended. Students with credit for MBB 243 may not take this course for further credit. CMPT 120 will be accepted in lieu of MBB 110.

TOPICS

- Flavours of data in molecular biology and biochemistry
- Genomic data
- Fundamentals of R and Python
- Regular expressions and patterns
- Quantitative DNA/RNA sequence analysis
- Exploratory data analysis in the Tidyverse
- Generic visualization methods using ggplot2
- Advanced visualizations for molecular biology

INSTRUCTORS

Sophie Sneddon



Bioinformatician, video game enthusiast, and cat person

Ryan Morin



Photograph may not be up to date



JULIA EVANS @bork

sort & uniq

16

sort sorts
its input

\$ sort names.txt

the default sort is alphabetical.

sort -n	
numeric sort	
'sort'order	`sort -n' order
12	12
15000	48 🖔
,, 48	96 😃
6020	6020
96	15000

sort -h: human sort

'sort -n' order | 'sort -h' order

156 | 45 K

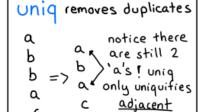
11 30 M | 30 M | 11

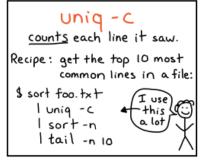
11 45 K | 156 | 11

2006 | 2006

Useful example:

du -sh * | sort -h





Group Data

matching lines

dplyr::group_by(iris, Species)

Group data into rows with the same value of Species.

dplyr::ungroup(iris)

Remove grouping information from data frame.

iris %>% group_by(Species) %>% summarise(...)
Compute separate summary row for each group.

