Markdown file structure

title: "genome569"

output: html_document

Course description

Programming skills and software tools for building automated bioinformatics pipelines and computational biology analyses. Emphasis on UNIX tools and R libraries for distilling raw sequencing data into interpretable results. This course is aimed at students familiar with UNIX and with some programming experience in python, R, or C/C++.

Instructional staff

- * [Cole Trapnell](http://cole-trapnell-lab.github.io/)
- * Teaching assistant: April Lo

"Front matter": Meta data about the document goes here

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