

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

Bulleted list. Enumerated lists also easy.

Markdown key features