

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

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“Front matter”: Meta data
about the document goes
here

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