

Markdown file structure

```
---  
title: "genome569"  
output: html_document  
---
```

Course description ← Section header

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++.

← Paragraph

Instructional staff

* [Cole Trapnell](<http://cole-trapnell-lab.github.io/>)

* Teaching assistant: April Lo

← Hyperlink

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Bulleted list. Enumerated lists also easy.