



# BIMM 143

## Course Summary & GitHub Portfolio

### Class 20

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UC San Diego

[https://marcos-diazg.github.io/BIMM143\\_SP23/](https://marcos-diazg.github.io/BIMM143_SP23/)

# Today's Menu

- Summary of major learning goals
- Course discussion and feedback ( [https://board.net/p/BIMM143\\_SP23\\_DiazGay](https://board.net/p/BIMM143_SP23_DiazGay) )
- Final exam
  - ➔ Test structure, guidelines and rules
  - ➔ Topics and example questions
  - ➔ Exam preparation, discussion and open study
- Polish our GitHub content and publish your own website portfolios
- SET evaluation ( [Link](#) )
- **Bioinformatics in industry session! (1:30pm)**

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- **Bioinformatics in industry session! (1:30pm)**



[https://bioboot.github.io/bimm143\\_F22/class-material/BIMM143\\_exam\\_guidlines.pdf](https://bioboot.github.io/bimm143_F22/class-material/BIMM143_exam_guidlines.pdf)

The screenshot shows a web browser window with the URL [https://bioboot.github.io/bimm143\\_S18/class-material/BIMM143\\_exam\\_guidlines.pdf](https://bioboot.github.io/bimm143_S18/class-material/BIMM143_exam_guidlines.pdf) in the address bar. The page content is a PDF document titled "BIMM-143: INTRODUCTION TO BIOINFORMATICS" with the URL <http://thegrantlab.org/bimm143>. Below the title, there is a section titled "Preparing for the Final Exam". The text discusses the final exam format, which is open-book and open-notes, lasting 150 minutes and consisting of 35 questions. It specifies that most questions are short answer (worth 2 points) and includes some longer answer questions (worth 5 points). The number of points for each question is indicated at the beginning of each question. A total of 80 points are available. The document notes that no questions from lecture 10 (git version control system) will be asked, but major points from all other lecture material are examinable.

**BIMM-143: INTRODUCTION TO BIOINFORMATICS**  
<http://thegrantlab.org/bimm143>

**Preparing for the Final Exam**

**Overview:** The final exam for BIMM-143 will be an open-book, open-notes 150-minute test consisting of 35 questions.

Questions will be predominantly short answer (typically worth 2 points) with a number of more involved longer answer questions (typically worth 5 points).

The number of points for each question is indicated at the beginning of each question. There are 80 total points on offer.

There will be no questions covering the material from lecture 10 (the git version control system). However, major points from all other lecture material are examinable

**Q1.** Did you enjoy this course in relation to others you have experienced at UCSD?

**Q2.** Should this course be offered again? Still in-person or fully remotely?

**Q3.** If so what changes would you recommend for this course? (e.g., more/less DataCamp & Projects)

**Q4.** Was the course effectively organized (lecture and lab material online vs handout or Canvas site)?

**Q5.** What advice would you give to another student who is considering taking this course?

**Q6.** Considering both the limitations and possibilities of the subject matter and the course, how would you rate the overall effectiveness of this course and the instructional team?

**Q7.** Do you agree or disagree - The course developed my abilities and skills for the subject?

**Q8.** On average, how many hours per week have you spent on this course? (including attending classes, doing homeworks, and assignments)

**Q9.** Any other comments you would like to share?

EtherPad Version: [https://board.net/p/BIMM143\\_SP23\\_DiazGay](https://board.net/p/BIMM143_SP23_DiazGay)

Form Version: <https://forms.gle/phWPQtPpknDgoCMa6>

Thank you very much!

# GitHub Spit & Polish

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bioboot/bimm143\_serina\_f18: Serina's Fall 2018 class repository fork Serina's BIMM 143 Class Repository | Serina's Bioinformatics Class (BIMM143,...) Bioinformatics Class BIMM-143 | Introduction to Bioinformatics (BIMM143) +

# Introduction to Bioinformatics



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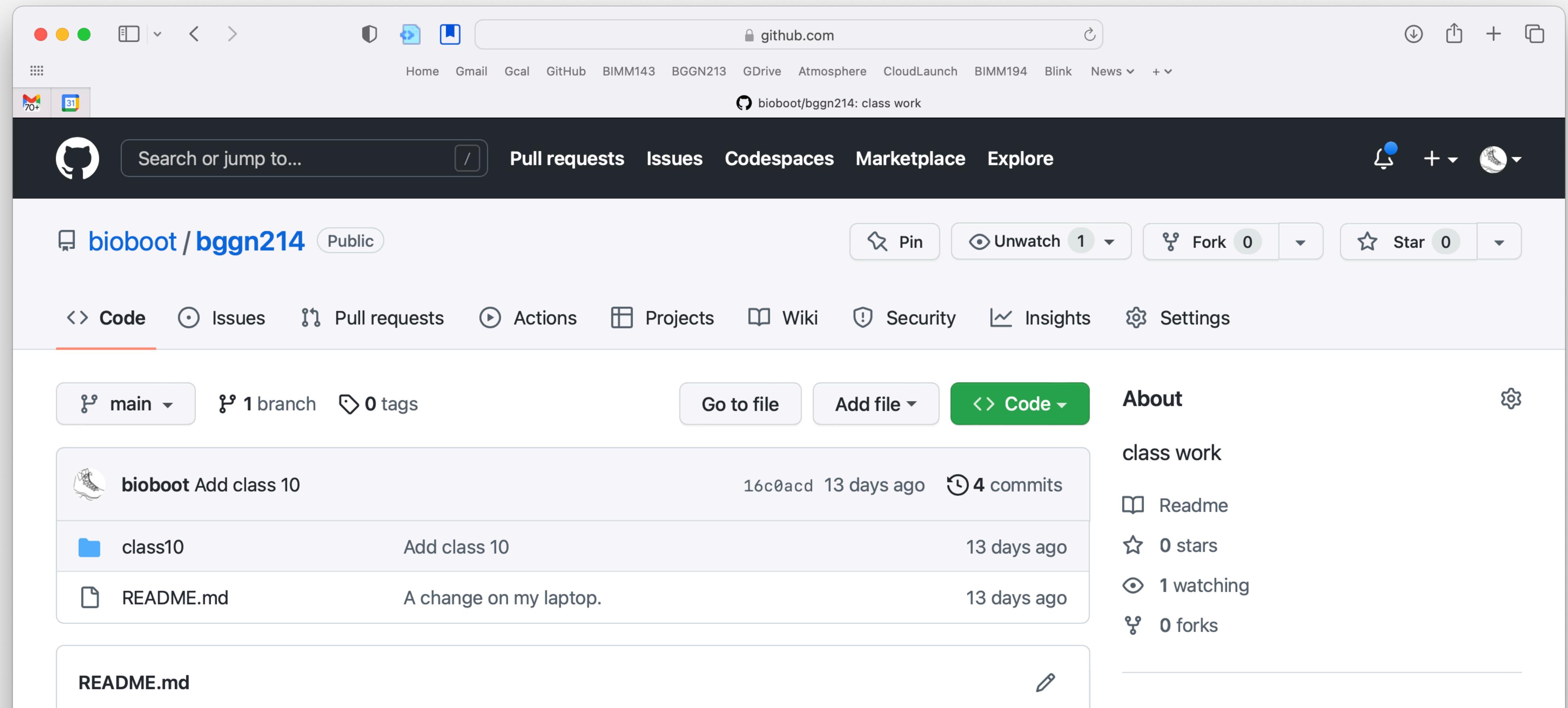
Class 14 - [Transcriptomics and RNA-Seq Analysis](#)

**In your web browser navigate to your GitHub class repository <<https://github.com/>>**

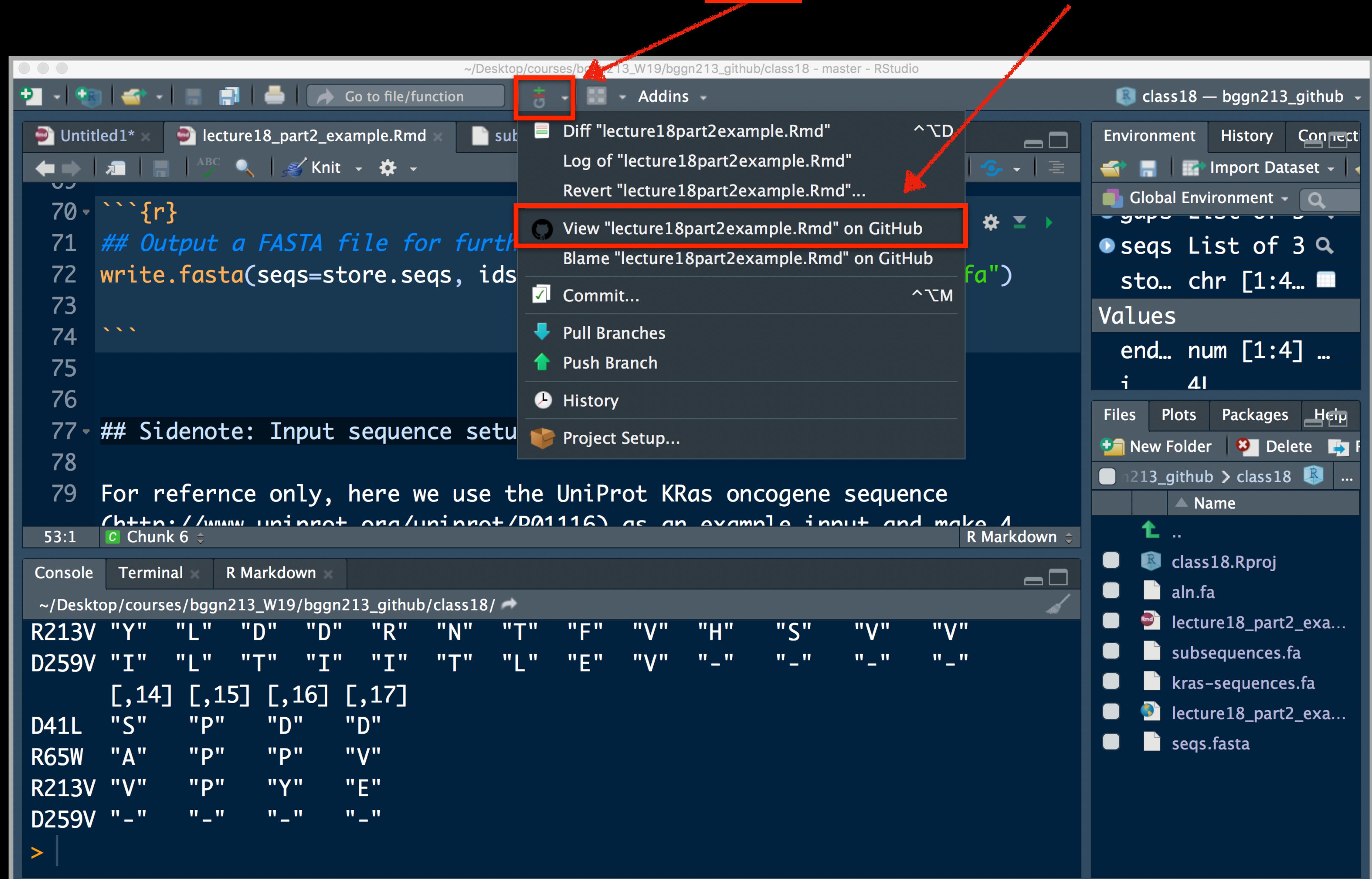
**Side-note:**

**To find the link to your GitHub repository from RStudio, open one of your past class projects and in the terminal type:**

**git remote -v**



Or for a given GitHub tracked file click **GIT** icon and "VIEW on GITHUB"

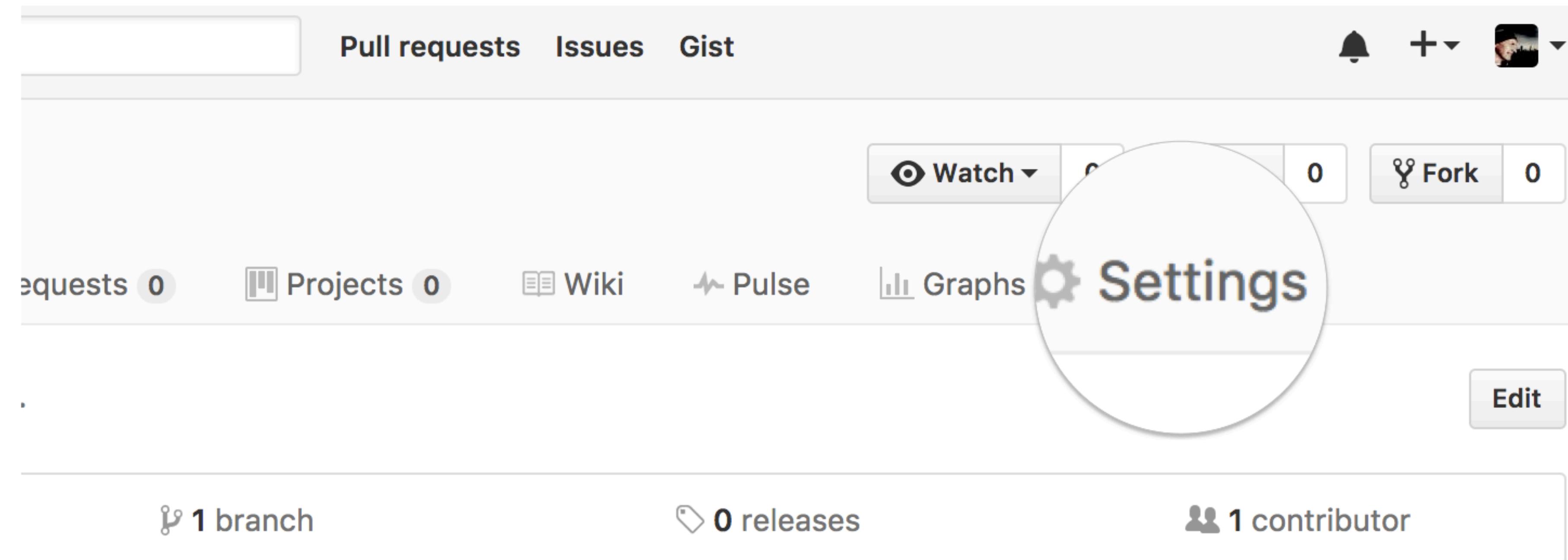


1

# Repository Settings

Head over to your **GitHub** class repository (where you are “pushing” all your class work)

Click on the **Settings** tab.



## Theme chooser

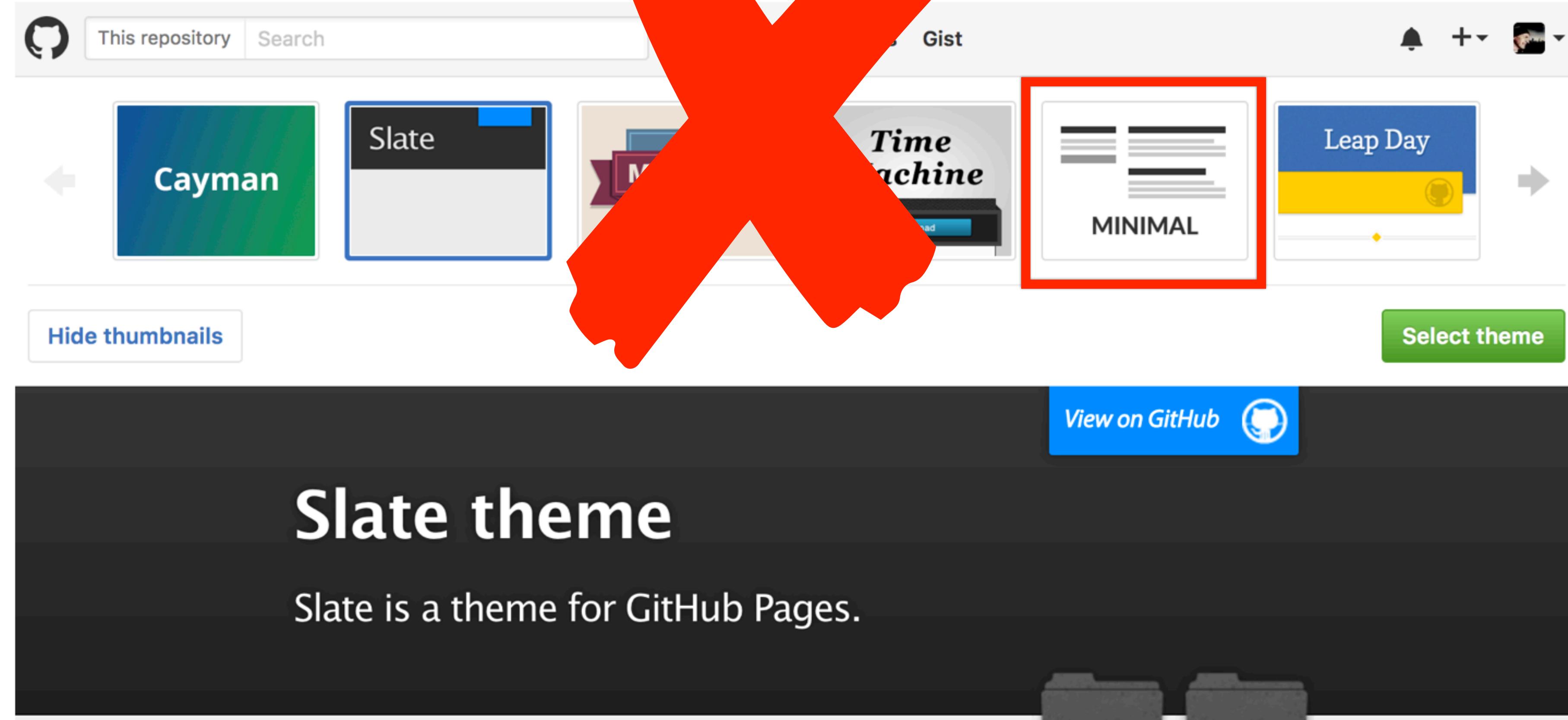
Go to the **Pages** section in the sidebar. Then change the **Branch** from “None” to “**main**”.

The screenshot shows the GitHub Pages settings page. On the left, there's a sidebar with various repository management options: General, Access, Collaborators, Moderation options, Code and automation (with sub-options like Branches, Tags, Actions, Webhooks, Environments, and Codespaces), and Pages. The Pages option is highlighted with a red box. The main content area is titled "GitHub Pages" and contains sections for "Access", "Collaborators", "Moderation options", "Build and deployment", "Source", and "Branch". The "Source" section has a dropdown menu set to "Deploy from a branch". Below it, the "Branch" section says "GitHub Pages is currently disabled. Select a source below to enable GitHub Pages for this repository." It features a dropdown menu where "main" is selected and highlighted with a red box. There are also buttons for "Save" and another for "Publish privately to people with read access to this repository".

## Pick a theme

Choose one of the themes from the carousel at the top.

When you're done, click **Select theme** on the right.



## Side-note:

**Scroll down again to the [GitHub Pages](#) section to find the link to your new website.  
Open this link in a [New Tab](#) of your browser:**

### GitHub Pages

GitHub Pages is designed to host your personal, organization, or project pages from a GitHub repository.

Your site is ready to be published at [https://bioboot.github.io/bimm143\\_serina\\_f18/](https://bioboot.github.io/bimm143_serina_f18/).

#### Source

Your GitHub Pages site is currently being built from the `master` branch. [Learn more.](#)

[master branch ▾](#)

[Save](#)

#### Theme Chooser

Select a theme to publish your site with a Jekyll theme. [Learn more.](#)

Your site is currently using the Minimal theme.

[Change theme](#)

## Edit content

**Back on the repository main page use the [GitHub](#) online editor to add content.  
In particular, add links to each classes .MD file**

The screenshot shows the GitHub repository interface for 'jldec / new-pages-site'. The top navigation bar includes the repository name, a 'Watch' button (0), a 'Star' button (0), and a 'Fork' button (0). Below the navigation bar are links for 'Issues 0', 'Pull requests 0', 'Projects 0', 'Wiki', 'Pulse', 'Graphs', and 'Settings'. The main content area shows the 'new-pages-site / README.md' file being edited. The file contains the following Markdown content:

```
1 ## Welcome to GitHub Pages
2
3 You can use the [editor on GitHub](https://github.com/jldec/new-pages-site/edit/master/README.md) to maintain and preview the content
for your website in Markdown files.
4
5 Whenever you commit to this repository, GitHub Pages will run [Jekyll](https://jekyllrb.com/) to rebuild the pages in your site, from
the content in your Markdown files.
6
7 ### Markdown
8
9 Markdown is a lightweight and easy-to-use syntax for styling your writing. It includes conventions for
10
11 ````markdown
12 Syntax highlighted code block
13
14 # Header 1
```

 [bioboot / bimm143\\_serina\\_f18](#)  
forked from [serinahuang/bimm143](#)

[Unwatch](#) 1 [Star](#) 0 [Fork](#) 1

[Code](#) [Pull requests 0](#) [Projects 0](#) [Wiki](#) [Insights](#) [Settings](#)

Branch: [master](#) [bimm143\\_serina\\_f18 / \\_config.yml](#) [Find file](#) [Copy path](#)

 **bioboot** Update \_config.yml 3b72493 just now

1 contributor

4 lines (3 sloc) | 151 Bytes [Raw](#) [Blame](#) [History](#) [!\[\]\(694dc821ccdd019462bbe776fc2013d6\_img.jpg\)](#) [!\[\]\(fb072c6d3f639f477372c36606365f1a\_img.jpg\)](#) [!\[\]\(81aee2fece3f9c930db8cc14ef658a9a\_img.jpg\)](#)

```
1 theme: jekyll-theme-minimal
2 logo: https://bioboot.github.io/bimm143_F18/assets/img/logo.png
3 title: Serina's Bioinformatics Class (BIMM143, Fall 2018)
```

(4.1) Edited config.yml (adding logo and title), (4.2) Edited README.md

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Class 14 - [Transcriptomics and RNA-Seq Analysis](#)

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The screenshot shows a GitHub repository page for 'bioboot / tmp\_test'. The repository has 1 unwatched star and 0 forks. The 'Code' tab is selected, showing the contents of the 'tmp\_test' branch's README.md file. The file content is as follows:

```
1 # BIMM143 Classwork
2
3 This is a store of my class-work for [BIMM143 Winter 2019](https://bioboot.github.io/bimm143\_W19/) at UC San Diego.
4
5 ## Content
6 - Class05: [R fundamentals](https://github.com/bioboot/tmp\_test/blob/master/class05/class05.md)
7 - Class06: [R graphics]()
8 - Class07: R Functions
9 - Class08: R packages from CRAN, Bioconductor and GitHub
10 - Class09: Introduction to machine learning
11 - Class10: Some thing else
12 - Class11: (Structural Bioinformatics)(https://github.com/bioboot/tmp\_test/blob/master/class11/class11.md)
13 - Class12: etc. etc.
14
```

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# Commit

Enter a commit comment and click on **Commit changes** below the editor.

```
35  ### Support or Contact  
36  
37  Having trouble with Pages? Check out our \[documentation\]\(https://help.github.com/categories/gi\)  
\(\) and we'll help you sort it out.  
38
```



## Commit changes

Add content to new pages site

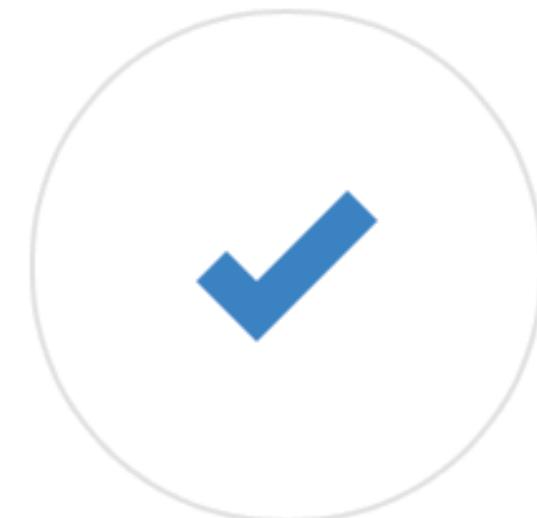
Add an optional extended description...

Commit directly to the `master` branch.

6

...and you're done!

Fire up a browser and go to **<http://username.github.io/repository>**.



Thank you very much!

# Thank you very much!

Post your GitHub Pages portfolio URL to GradeScope!

Remember: Late Due Date for all past assignments -> Friday, June 16 at 12pm

# Bonus:

# Going Further With DataCamp

[https://marcos-diazg.github.io/BIMM143\\_SP23/class-material/class20/datacamp\\_extras.pdf](https://marcos-diazg.github.io/BIMM143_SP23/class-material/class20/datacamp_extras.pdf)

The screenshot shows a web browser displaying the DataCamp website at [www.datacamp.com/tracks/custom-bioinformatics-extension](https://www.datacamp.com/tracks/custom-bioinformatics-extension). The page title is "CUSTOM TRACK Bioinformatics Extension". The main content area describes Barry's suggested DataCamp courses for delving deeper into R, Git, the Shell, data visualization, and Bioinformatics data analysis generally. It includes a yellow "Enroll" button and navigation links for R Language, Shell, Git, Spreadsheets, 111 hours, and 26 Courses. The top navigation bar includes links for Home, Gmail, Gcal, GitHub, BIMM143, BGNN213, Atmosphere, BIMM194, Blink, News, and a user profile icon.

# SET evaluations

Please fill out the new Student Evaluations of Teaching (SET) if you have a chance.  
It is important to the courses we offer in the future and how we teach them!

**Bonus:**  
**Bioinformatics & Genomics**  
**in Industry (2020 edition)**  
Live Stream Video

Enjoy a set of short interviews from leading genomic scientists at **Illumina Inc.**, **Synthetic Genomics Inc.**, and the **La Jolla Institute for Immunology**



# The End!

[https://marcos-diazg.github.io/BIMM143\\_SP23/](https://marcos-diazg.github.io/BIMM143_SP23/)



# Thank You!

[https://marcos-diazg.github.io/BIMM143\\_SP23/](https://marcos-diazg.github.io/BIMM143_SP23/)