

# BIMM 143

## Course Summary & GitHub Portfolio

### Class 20

Marcos Díaz-Gay  
UC San Diego

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



# Today's Menu

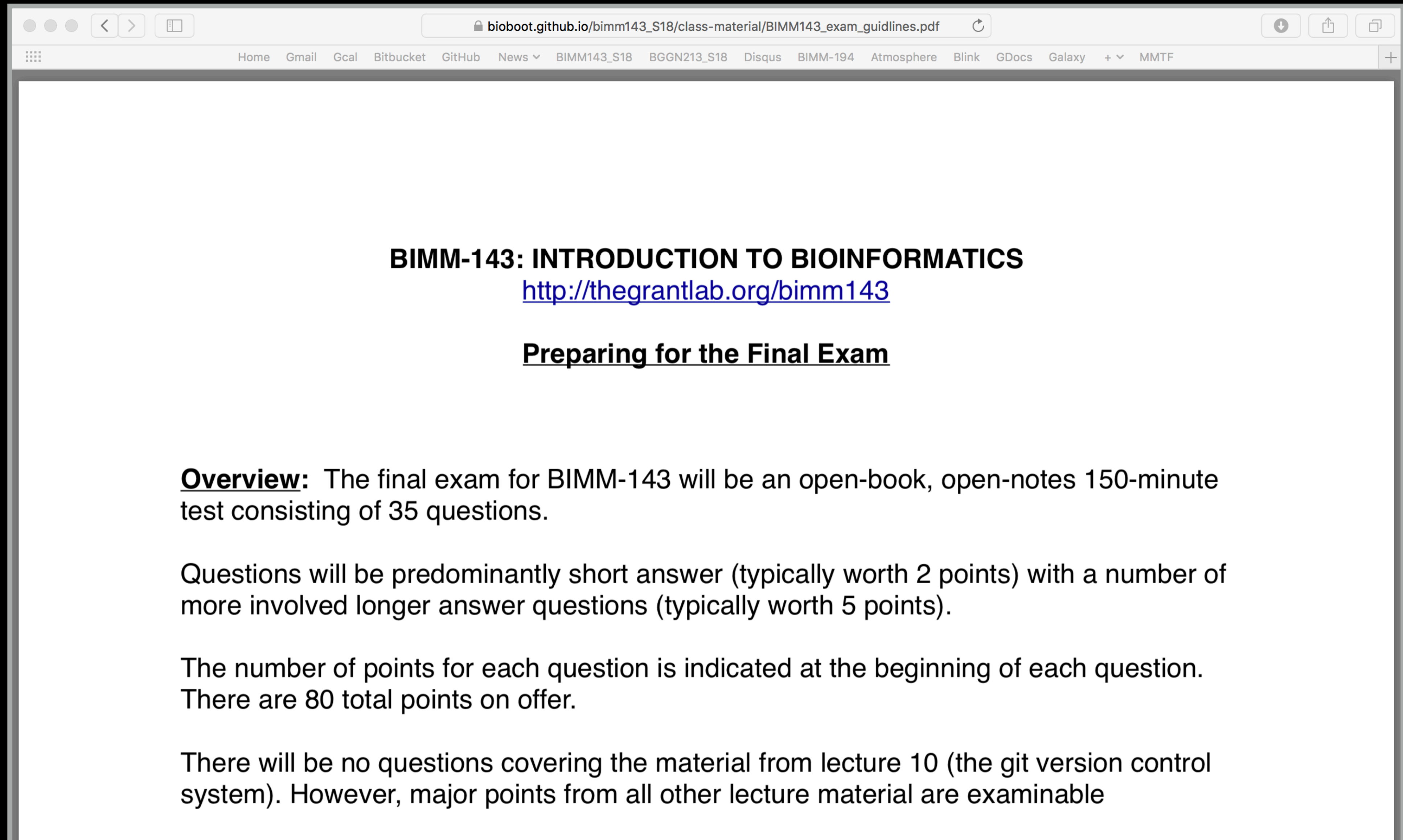
- Summary of major learning goals
- Course discussion and feedback ( [https://board.net/p/BIMM143\\_SP23\\_DiazGay](https://board.net/p/BIMM143_SP23_DiazGay) )
- Final exam
  - ➡ Test structure, guidelines and rules
  - ➡ Topics and example questions
  - ➡ Exam preparation, discussion and open study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

# Today's Menu

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[https://bioboot.github.io/bimm143\\_F22/class-material/BIMM143\\_exam\\_guidelines.pdf](https://bioboot.github.io/bimm143_F22/class-material/BIMM143_exam_guidelines.pdf)



The screenshot shows a web browser window with the address bar displaying the URL: [bioboot.github.io/bimm143\\_S18/class-material/BIMM143\\_exam\\_guidelines.pdf](https://bioboot.github.io/bimm143_S18/class-material/BIMM143_exam_guidelines.pdf). The browser's tab bar shows several open tabs: Home, Gmail, Gcal, Bitbucket, GitHub, News, BIMM143\_S18, BGGN213\_S18, Disqus, BIMM-194, Atmosphere, Blink, GDocs, Galaxy, and MMTF. The main content area of the browser displays the following text:

**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



# Introduction to Bioinformatics



A demo site of students cool class web site

[View the Project on GitHub](#)  
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This project is maintained by [bioboot](#)

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# Bioinformatics Class

This is my repository for my Bioinformatics class from UC San Diego in S18.

## Index of Material

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## Using R and Other Tools for Bioinformatics Analysis

## Class 8 - An Introduction to Machine Learning (Heirarchical Clustering)

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## Class 11 - Structural Bioinformatics: Analyzing Protein Structure and Function

## Class 12 - Drug Discovery: Techniques and Analysis

## Class 13 - Genome Informatics and High Throughput Sequencing (NGS, RNA-Seq, and FastQC)

## 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
```

The screenshot shows a web browser window displaying the GitHub repository page for `bioboot/bggn214`. The browser's address bar shows `github.com`. The repository is public and has 0 stars, 0 forks, and 1 watch. The repository contains a `main` branch with 1 branch and 0 tags. The commit history shows a commit titled "Add class 10" by `bioboot` 13 days ago, with 4 commits in total. The commit message is "A change on my laptop." The repository also has a `README.md` file. The right sidebar shows the repository name "class work" and a list of files: `Readme`, `0 stars`, `1 watching`, and `0 forks`.





View "lecture18part2example.Rmd" on GitHub

Blame "lecture18part2example.Rmd" on GitHub

 Commit...

## ↓ Pull Branches

## ↑ Push Branch

 History Project Setup...

```
70 ~~~{r}
71 ## Output a FASTA file for further analysis
72 write.fasta(seqs=store.seqs, ids=store.ids, file="sequences.fasta")
73 ~~~
74
75
76
77 ## Sidenote: Input sequence setup
78
79 For reference only, here we use the UniProt KRas oncogene sequence
80 (http://www.uniprot.org/uniprot/P01116) as an example input and m
```

```
~/Desktop/courses/bggg213_W19/bggg213_github/class18/ ➜  
R213V "Y" "L" "D" "D" "R" "N" "T" "F" "V" "H" "S" "V" "V"  
D259V "I" "L" "T" "I" "I" "T" "L" "E" "V" "_" "_" "_" "  
      [,14] [,15] [,16] [,17]  
D41L  "S"  "P"  "D"  "D"  
R65W  "A"  "P"  "P"  "V"  
R213V "V"  "P"  "Y"  "E"  
D259V "_"  "_"  "_"  "  
>
```

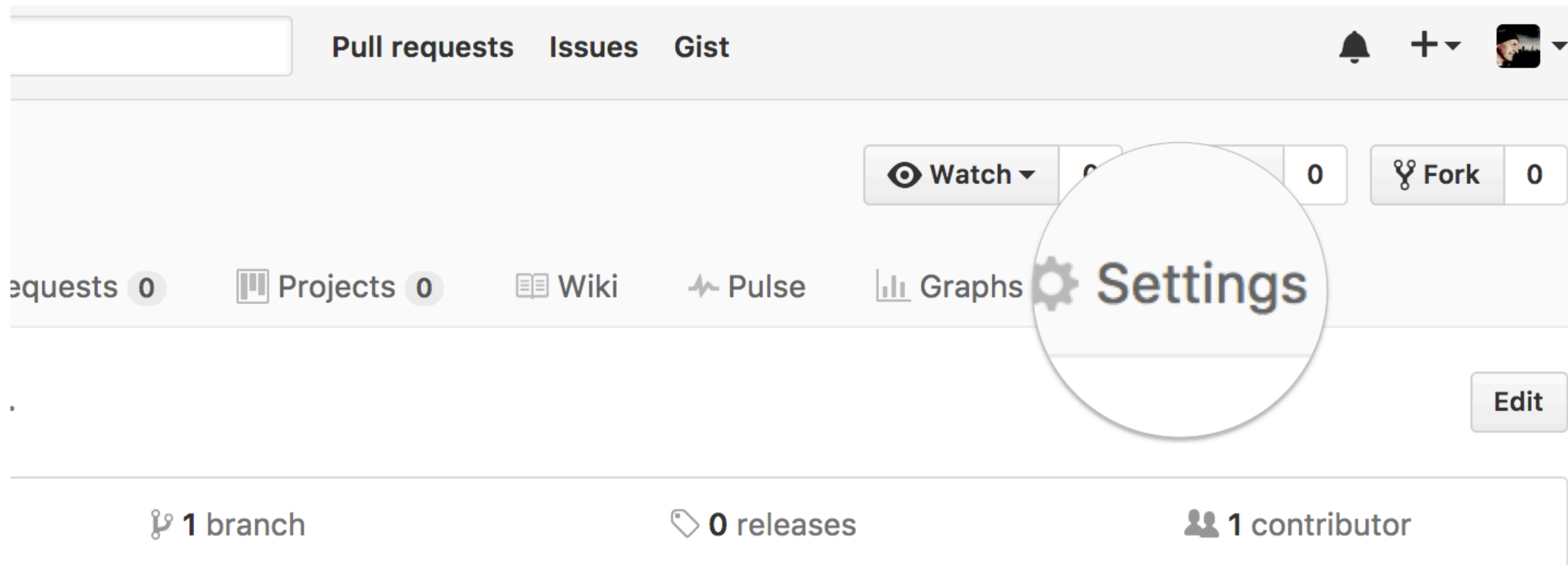


1

# 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”.



**General**

**Access**

- Collaborators
- Moderation options

**Code and automation**

- Branches
- Tags
- Actions
- Webhooks
- Environments
- Codespaces
- Pages**

## GitHub Pages

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

### Build and deployment

**Source**

Deploy from a branch

**Branch**

GitHub Pages is currently disabled. Select a source below to enable GitHub Pages for this repository. [Learn more.](#)

**main** / (root) Save

**Publish privately to people with read access to this repository**

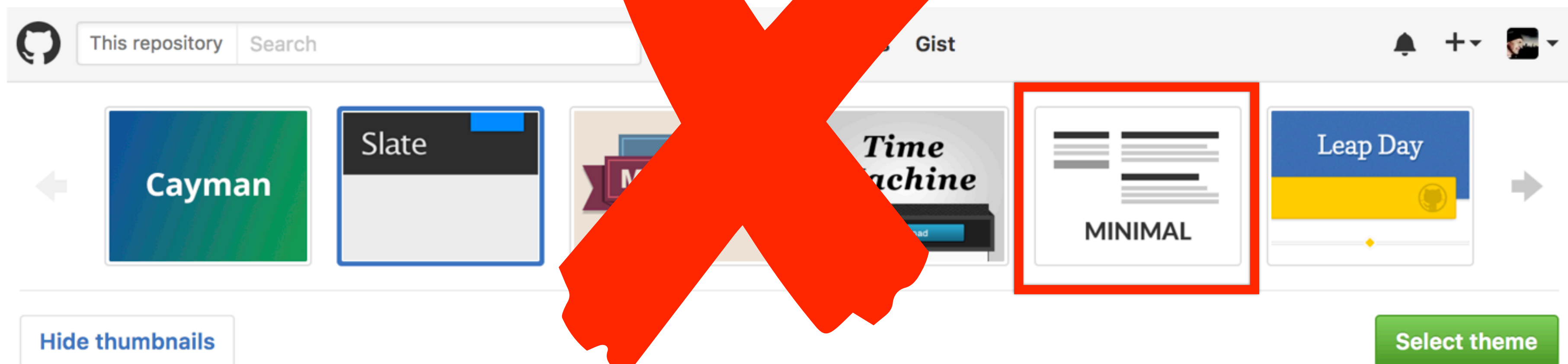
Try risk-free for 30 days using a GitHub Enterprise organization, or [learn more about changing the visibility of your GitHub Pages site.](#) ✕



## Pick a theme

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

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



## Slate theme

Slate is a theme for GitHub Pages.

[View on GitHub](#)



## 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 interface for the repository 'jldec / new-pages-site'. At the top, there are buttons for 'Watch' (0), 'Star' (0), and 'Fork' (0). Below these are tabs for 'Code', 'Issues' (0), 'Pull requests' (0), 'Projects' (0), 'Wiki', 'Pulse', 'Graphs', and 'Settings'. The 'Code' tab is selected, and the file 'README.md' is shown. The editor interface includes a toolbar with 'Edit file' and 'Preview changes' tabs, and settings for 'Spaces' (2) and 'Soft wrap'. The code content is as follows:

```
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
```



↔ 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



```
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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bioboot / tmp\_test

Unwatch 1 Star 0 Fork 0

<> Code Issues 0 Pull requests 0 Projects 0 Wiki Insights Settings

tmp\_test / README.md or cancel

<> Edit file Preview changes Spaces 2 Soft wrap


```
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/github-pages-faq)(<https://help.github.com/categories/github-pages-faq>)  
(<https://github.com/contact>) and we'll help you sort it out.

38



## Commit changes

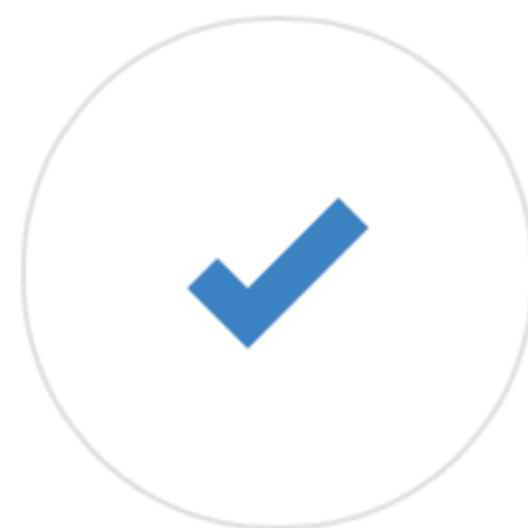
Add content to new pages site|

Add an optional extended description...

☒ Commit directly to the `master` branch.

...and you're done!

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





Thank you very much!

# Thank you very much!

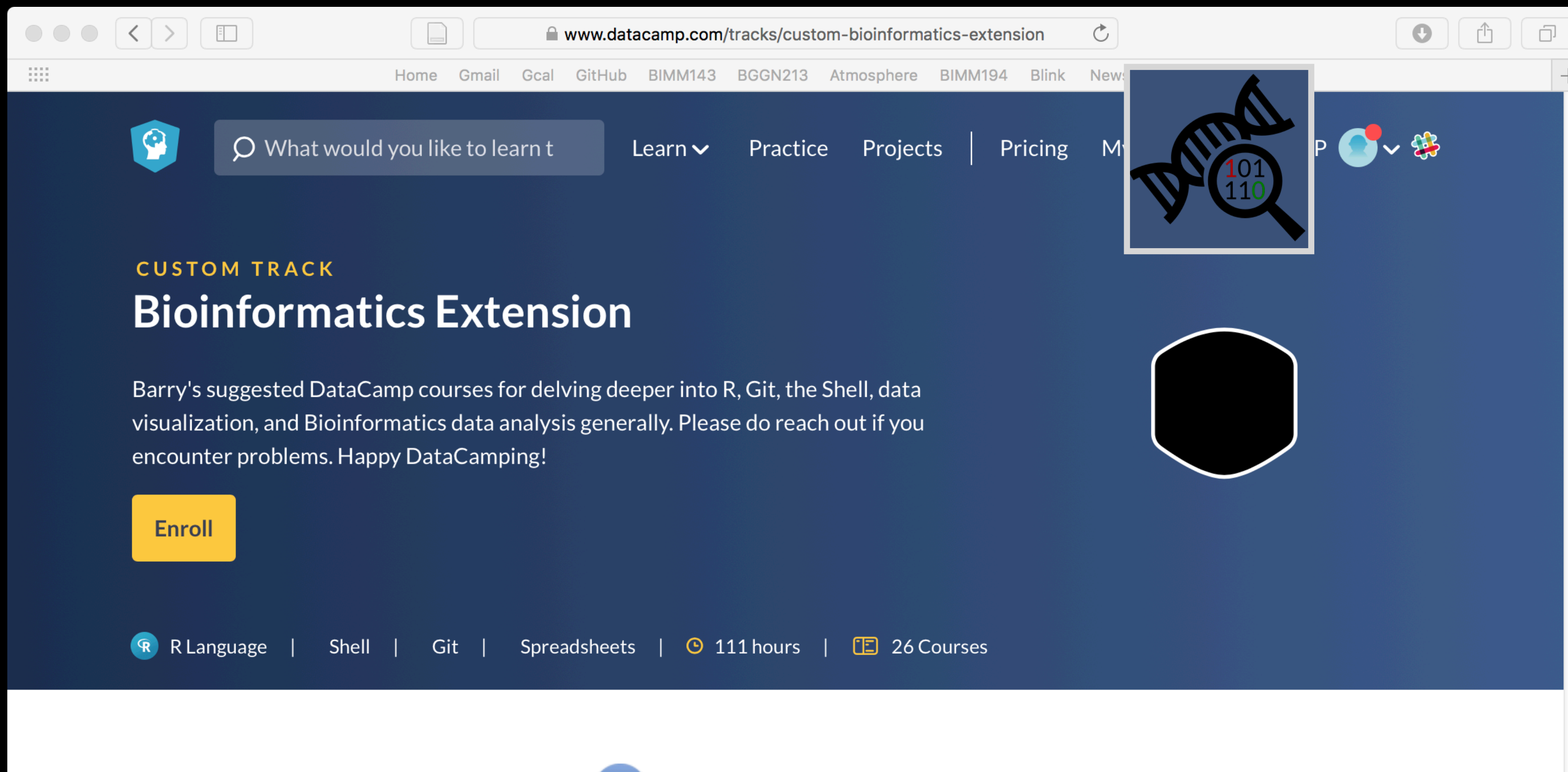
Post your GitHub Pages portfolio URL to GradeScope!

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)





# 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!

Announcement!

**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/)



The background of the slide is an abstract composition of numerous overlapping, semi-transparent spheres. These spheres are rendered in a variety of colors including shades of purple, magenta, blue, yellow, and orange. They are arranged in a way that creates a sense of depth and movement, with some spheres appearing closer and larger than others. The overall effect is a vibrant, textured field of color.

# Thank You!

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