VERSION 2

JAN 08, 2019

OPEN ACCESS

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External link:

<http://hurwitzlab.org>

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Protocol status: Working
We use this protocol and it's working

Created: Jan 08, 2019

Last Modified: Jan 08, 2019

PROTOCOL integer ID:
19098

Create a class Github repository V.2

Version 1 is forked from [Create "abe487" Github repo](#)

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Biosys-analytics

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ABSTRACT

The github repository will serve as a location for you to turn-in your homework, and allow the instructors to push back code modifications and suggestions.

BEFORE START INSTRUCTIONS

Sign up for a free Github account: <http://github.com>

- 1 Login to <http://github.com>
- 2 Go to the class Github repository: <https://github.com/hurwitzlab/biosys-analytics>
- 3 Fork (or "copy") the class repository into your own Github account. This creates a copy of the repository in your own account that you can edit and update. Select your Github account as the place to "fork" the repository.

Forking (or copying the biosys-analytics repository). All homework assignments will be in this repository.

- 4 Go to your Github account. You will see a directory called biosys-analytics. This is your copy.
<http://github.com/my-username> . (replace my-username with yours!)

Search or jump to... / Pull requests Issues Marketplace Explore

Your pinned repositories have been updated.

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Overview Repositories **4** Settings

Pinned repositories

- biosys-analytics**
Forked from hurwitzlab/biosys-analytics
BE434/535 Biosystems Analytics
Haskell

- 5 Click on your "biosys-analytics" repository. Click on the "Settings" tab on the top bar, click on the "Collaborators" tab on the side bar, and search for & add the instructors as collaborators. The instructors Github usernames are: bhurwitz33 and kycklar.

Adding collaborators (the instructors) to YOUR Github repository called biosys-analytics