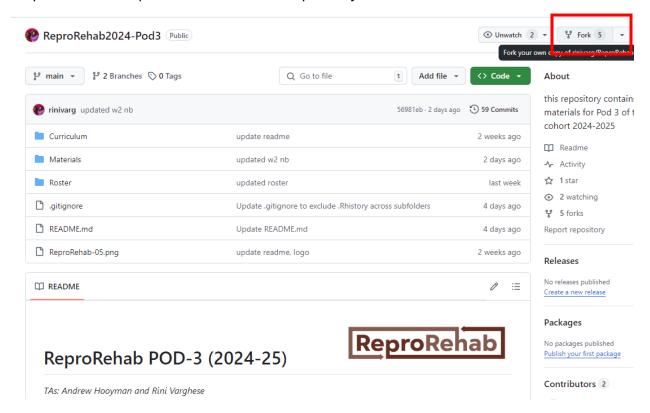
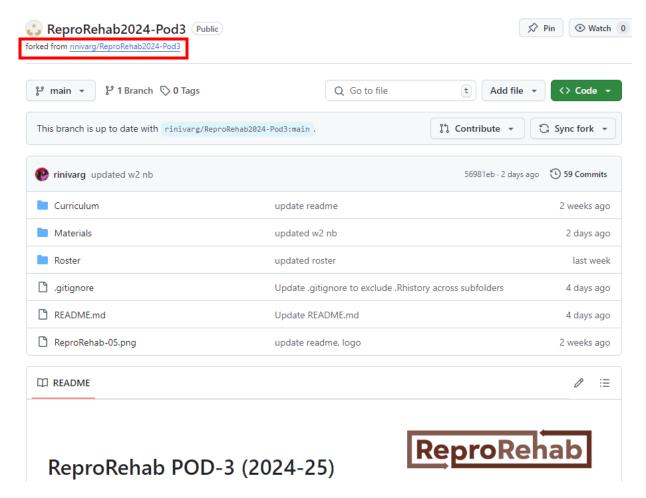
## Step 1: Create a github account

Step 2: Fork the ReproRehab2024-Pod3 Repository in Github

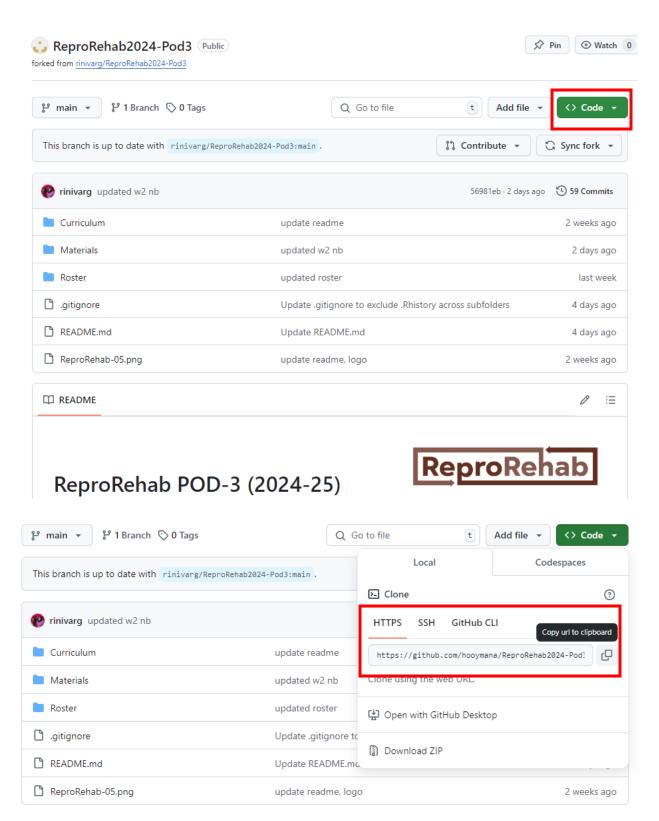


This will create a "forker" (copied) repository in your github account. See mine below



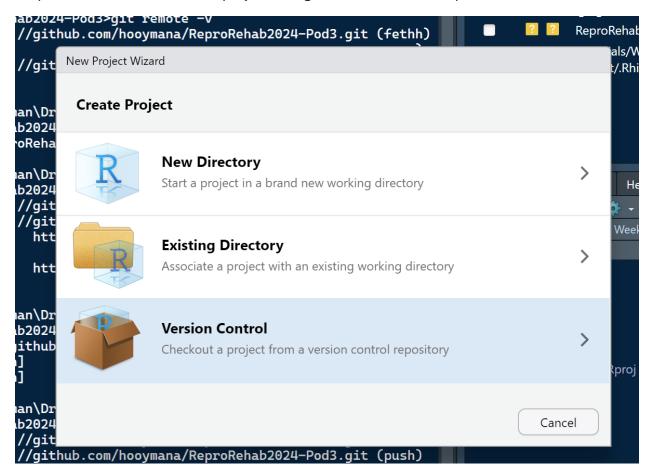
Now we want to "clone" (download) this repository onto our machine.

Step 3: Clone the repository, click on the green code button and copy the repository link

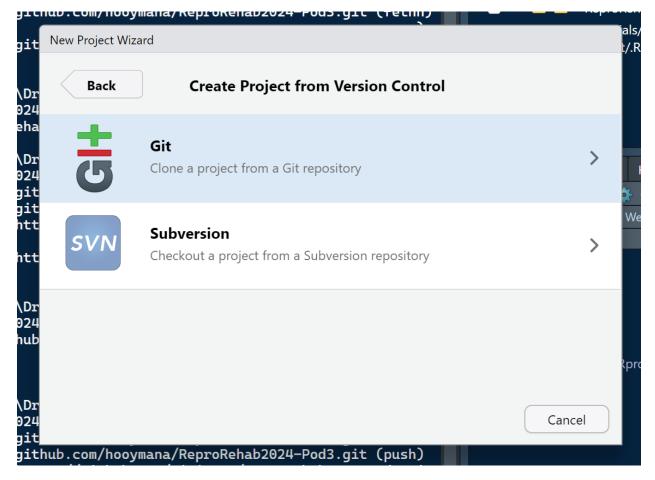


With this copied url, now go to R studio.

Step 4: We will create a new R project using the version control option



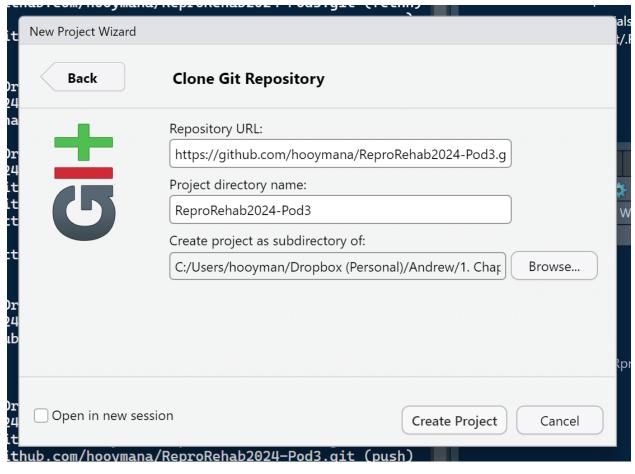
Select Git (if you don't see this option you may need to install git onto your computer).



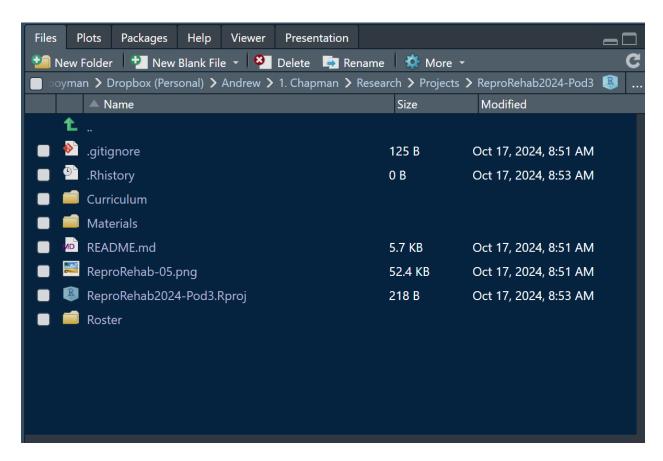
Paste the copied forked repository URL from your account into the Repository URL field. This will also create the directory name, and make sure the subdirectory is in the right place.

Click create project.

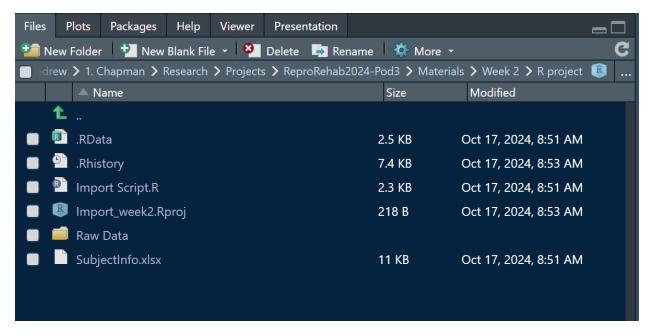
This will then "clone" (download) the entire repro onto your local machine and store it as an R project in that subdirectory.



The cloned repo now looks like any other type of folder and file system in your computer.



You can now navigate into materials and use the code and data in the repo for each week.



IMPORTANT NOTE: THIS ESTABLISHES A REMOTE CONNECTION TO YOUR FORKED REPO. YOU CAN NOW CHANGE THE CODE AND MAKE COMMITS/PUSHES TO YOUR

FORKED REPO AND IT WILL NOT MODIFY THE MAIN REPO THAT YOU ORIGINALLY FORKED FROM.

**FAQ** 

Q. Do I now have to continue to fork and clone the main repo everytime there are changes?

A. No. You can sync any new commits to the main branch into your fork which should then update the project on your local machine.

Check out the below link for a great tutorial on how to handle the majority of use cases when using git in Rstudio and Github.

https://docs.github.com/en/pull-requests/collaborating-with-pull-requests/working-with-forks/syncing-a-fork