# Scripting in a reproducible and collaborative framework using Github and RStudio

## 17/03/2022

## Group 2

Team: Adikrishna (s212940), Oriade (s172084), Prince (s220520), Razia (s151549), Yuan (s212420)

### Learning Objectives

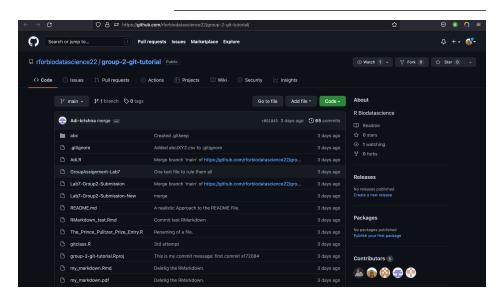
A student who has met the objectives of the session will be able to:

- Explain why reproducible data analysis is important
- Identify associated relevant challenges
- Explain the ability to replicate versus reproducibility
- Describe the components of a reproducible data analysis
- Use RStudio and GitHub (git) for collaborative bio data science projects

#### **Task 2.6:**

Your GROUP ASSIGNMENT is any written document (might be even a simple ".txt" file) containing the link to your repository and name of the file used in this exercise.

We can inspect it by using blame feature, which shows who committed and what was committed.



 $The \ link \ to \ the \ repository \ is: \ https://github.com/rforbiodatascience 22/group-2-git-tutorial/research and the repository is: \ https://github.com/rforbiodatascience 22/group-2-git-tutorial/research and the re$ 

The name of the files used in this exercise are: lab-7-group2-assignment and README.md