

Biostatistics: Problem Set 2 – Git and Beginning R

Name: **Matt Serrano** Score = **21.5** /25

GitHub repo: <https://github.com/mattserrano1/PS2-trick-or-treat-personal.git>

Submitted on time? ☒ Y ☐ N

Project element	Value	Pts earned	Comments
Successfully fork a GitHub repository and create a new RStudio project from fork <ul style="list-style-type: none"> Project called “Lastname-Trick-or-Treat” 	3	2.5	You did not name your github repo or project according to the instructions
Set up project and workspace, pull in and examine data, fix mistakes Lastname-PS2.qmd <ul style="list-style-type: none"> Use at least 2 functions Assign data types Fix errors (factors, plots) 	3	2.5	You never did any error checking for numeric variables.
Manipulate data and use it to correctly create Figure 1	3	3	good but notice you have different levels of the factor than I did.
Manipulate data and use it to correctly create figure 2	3	2.5	You did not adjust axis labels
Create a correct summary data table that opens in a new view <ul style="list-style-type: none"> Sugar high, mean age, sd age, mean g candy, sd g candy 	3	2	You did not get mean g candy or sd g candy in your code.
Manipulate data and use it to correctly create Figure 3	3	3	good but again notice that you don't have same factor levels that I do
Thought processes are well documented outside of code blocks	3	2	I'd like you to work more on this - esp. w/ regard to remarking on the outcome of running code chunks.
Successfully open a pull request to add your changes to the forked repository <ul style="list-style-type: none"> Commit changes Open PR Link pasted in Canvas 	2	2	good
Code represents material we have covered in GSWR Chs 2 - 4 and not elsewhere	2	2	good