

Biostatistics: Problem Set 2 – Git and Beginning R

Name: Chris Augustine

Score = 22 /25

GitHub repo: <https://github.com/chrisaugustine123/PS2-trick-or-treat>

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 | Project/Repo not named "Lastname-Trick-or-Treat" |
| 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 | Why did you make costume a factor w/ 42 levels? Also, you didn't do any error checking on any of the numeric vars. |
| Manipulate data and use it to correctly create Figure 1 | 3 | 2.5 | I can see from your figure legend that your factor levels for costume type don't match mine. Subtle issue. |
| Manipulate data and use it to correctly create figure 2 | 3 | 3 | I'm not taking off any points this time, but your use of theme_minimal() causes your fig to look different from mine. |
| 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 | 3 | good |
| Manipulate data and use it to correctly create Figure 3 | 3 | 3 | Thanks for filling me in on why you adjusted the theme. |
| Thought processes are well documented outside of code blocks | 3 | 2 | I want to see more about what you learn from your code when you run it. For example, what did you learn about the data from running summary()? |
| 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. And I appreciate how much you teach yourself! Great. |