

Godin_review

I was able to get your markdown file to run. My comments on the repository are as follows:

- The readme file and the repository itself are very well organized, everything was easy to find.
- The rmarkdown file was also well organized and documented.
- Your plots are also very well formatted. However, I think you are missing a visualization question. The assignment asks for a plot that analyzes SNP distribution across groups. You have the plot for samples with the homo, hetero, and missing data classes. However, I think you also need a plot where the x-axis is divided into groups like ZMLR, TRIPS, etc.
- Your decreasing position files appear to have replaced the SNPs with numbers. If you look at the difference in output between the files, it should be clear what I mean. Having played around with the code, the problem appears to stem from this line, `mutate(across(4:ncol(.), ~ ifelse(. == "?/?", "-/-", .))) %>%`. I would take a look at its output to see why it is giving this behavior. I assume it is do to some type of datatype handling error. You may want to use `gsub` to substitute the `?` for `-`.
- I'm not sure why you included the `testing.R` file in the repo. This looks like a file you could have included in the gitignore. It is not mentioned in the readme, so I think it is not needed.