

Name: Alexis Parent

Score = 19 /25

Submitted on time? ☒ Y ☐ N

**GENERAL REQUIREMENTS (10 POINTS):**

Element	Points	Score	Feedback
Effective git/GitHub	1	1	
Well-organized	1	1	
Strong commentary outside of code chunks	3	2.5	
Effective use of comments within code chunks	2	1.5	
Code provides correct values and reduces “human intervention”	2	2	
Link on Canvas	1	1	

**STATISTICAL ANALYSES (15 POINTS):**

☐ Took initiative to learn new methods as appropriate

☒ Generally followed the our workflow:

Plot -> Guess -> Create model -> Check assumptions -> Interpret -> Final plot

**Statistical analysis 1:**

Question: Does MAOM change as a function of years between soil rotation?

Workflow checklist

☒ 1. Plot data ☐ 2. Guess relationships

☒ 3. Create model: 1-way anova (which is a form of lm) - but backwards

☐ Correct model?

☒ 4. Check model assumptions, if needed

☒ 6. Replot

☒ 5. Interpret model

☒ 7. Clear results statement

☒ Interpretation is correct

☒ In prose

☒ Outside of code chunk

**Statistical analysis 2:**

Question: Does POM change as a function of soil rotation?

Workflow checklist

- |   |  |
|---|--|
| <input checked="" type="checkbox"/> 1. Plot data                          | <input type="checkbox"/> 2. Guess relationships                |
| <input checked="" type="checkbox"/> 3. Create model: <u>1-way anova</u>   |  |
| <input checked="" type="checkbox"/> Correct model?                        |  |
| <input checked="" type="checkbox"/> 4. Check model assumptions, if needed | <input checked="" type="checkbox"/> 6. Replot                  |
| <input checked="" type="checkbox"/> 5. Interpret model                    | <input checked="" type="checkbox"/> 7. Clear results statement |
| <input checked="" type="checkbox"/> Interpretation is correct             | <input checked="" type="checkbox"/> In prose                   |
|   | <input checked="" type="checkbox"/> Outside of code chunk      |

**Statistical analysis 3:**

Question: Is there an association b/w SOM and soil Nitrogen?

Workflow checklist

- |   |  |
|---|--|
| <input checked="" type="checkbox"/> 1. Plot data                              | <input type="checkbox"/> 2. Guess relationships                |
| <input checked="" type="checkbox"/> 3. Create model: <u>linear regression</u> |  |
| <input checked="" type="checkbox"/> Correct model?                            |  |
| <input checked="" type="checkbox"/> 4. Check model assumptions, if needed     | <input checked="" type="checkbox"/> 6. Replot                  |
| <input checked="" type="checkbox"/> 5. Interpret model                        | <input checked="" type="checkbox"/> 7. Clear results statement |
| <input checked="" type="checkbox"/> Interpretation is correct                 | <input checked="" type="checkbox"/> In prose                   |
|   | <input checked="" type="checkbox"/> Outside of code chunk      |

Additional feedback

You are well on your way, but need to improve your technical writing about these analyses and include biological information in your justification for alternative hypotheses and in your interpretation of results.

See more detailed notes in [prelim-analysis-feedback.qmd](#)