

Name: Sara St. Clair

Score = 22.5 /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	3	
Effective use of comments within code chunks	2	2	
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: Are weed lifeform and biovigalence status independent from each other?

Workflow checklist

☒ 1. Plot data ☐ 2. Guess relationships

☒ 3. Create model: Chi-square test of independence

☒ 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: Are weeds in different habitats occurring independent of biogeographic realm?

Workflow checklist

- |   |  |
|---|--|
| <input checked="" type="checkbox"/> 1. Plot data                          | <input type="checkbox"/> 2. Guess relationships                |
| <input checked="" type="checkbox"/> 3. Create model: <u>chi-square</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 between lifeform and conservation status (Red\_UK)?

Workflow checklist

- |   |  |
|---|--|
| <input checked="" type="checkbox"/> 1. Plot data                          | <input type="checkbox"/> 2. Guess relationships                |
| <input checked="" type="checkbox"/> 3. Create model: <u>chi square</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

Overall, good, though as you know, rather repetitive. To extend your learning, I'd like you to learn how to conduct post-hoc tests with this sort of analysis and I've provided a link with information.

Please carefully read and apply the feedback in the prelim-analysis-feedback file.

You'll note that I didn't check any of the boxes for "guess relationships" - I'd like to see more biological thinking a priori about which groups you might expect to see have certain relationships as part of your alternative hypothesis statements.