Biostatistics:	Problem	Set 3 –	Introductory	Statistics

Name: Matt Serrano				Score = 21	/25
GitHub repo:	https:/	/github.cor	m/mattserrai	no1/Serrano-PS	S3.git
Submitted on			\bigcap N		

		Pts	
Project element	Value	earned	Comments
Successfully fork a GitHub repository and create a new RStudio project from fork • Project called "Lastname-PS3"	1	1	thanks for naming project correctly
Set up project and workspace, pull in and examine data, fix mistakes Lastname-PS3.qmd Use at least 2 functions Assign data types Error checking	2	2	Good job checking both factor and numeric variables
Analyze Q1: Does body mass differ b/w these 5 species of bats, and if so, how does body mass differ b/w species? Nature of P and R vars Analysis method explained More polished figure Clear, written interpretation	4	3.25	Why did you choose 1-way ANOVA? Should write that in narrative. Should address response and pred var types. Line 114 - what does QQ plot help you understand? Line 149 - looking for a nice final plot, not a nice model. Plot should show means, seee fig 5.11.
Analyze Q2: Does body length differ b/w species and, if so, how? Nature of P and R vars Analysis method explained More polished figure Clear, written interpretation	4	3.25	Same feedback as for Q1 - need a final plot and more on why you chose ANOVA. Also null hypotheses would be good for both.
Analyze Q3: Is the number of ticks found on the bats associated with their sex or age? Nature of P and R vars Analysis method explained More polished figure Clear, written interpretation	4	2.5	Selected incorrect statistical test - should be chi-square test of association (= test of indpendence). discrete var against 2 categorcial vars.

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	1	1	
Analyze Q4: Disregarding species, is there a relationship in bats b/w tail length and body length? • Nature of P and R vars • Analysis method explained • More polished figure • Clear, written interpretation	4	3.5	What is is about your two variables that makes theses data a good fit for linear regression? geom_smooth(method = lm) not model How do you interpret adjusted R2 value?
Thought processes are well documented outside of code blocks, code is well commented, all steps prior to data analysis	4	4	good
Successfully open a pull request to add your changes to the forked repository Commit changes Open PR Link pasted in Canvas	1	0.5	You did not make a pull request
Code represents material we have covered in GSWR Chs 3-5 and not elsewhere	1	1	
Additional feedback			

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