Final Project: Data Analysis Report

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#Q1 The relationship between body mass and length seems to be linear. Mass is positive correl ated with length.
#Q2 The data, especially body length, does not appear normally-distributed because the histogr ams are not symmetrical.
#Q3 I think the (unconditioned) body masses and body length are not normally-distributed (hist ograms are skewed, p-values < 0.05). My visual assessment of normality matched the results of the numerical normality tests.
#Q4 D. dorsalis seems to weigh more than D. sublineatus and male seems to weigh more than f emale, but not by much. There is no significant difference.

#Q5 Based on the numerical and graphical diagnostics, all models fail to fulfill the assumption of normality of the residuals.
#Q6 Violations of the normality assumption are not equally severe for all the models. The violati on is much more severe in model 1.

#Q7 0.8754988
#Q8 163.6745 mm
76.1246565+(0.8754988*100)

[1] 163.6745

#Q9 76.12466 mm
76.1246565+(0.8754988*0)

[1] 76.12466

#Q10 female
#Q11 Delomys dorsalis
#Q12 male
#Q13 Delomys dorsalis

#Q14 Sex and species are significant predictors for body mass.
#Q15 There is not a significant interaction.
#Q16 The significance level of sex and the significance level of species do not change much among the different models. All corresponding p-values are very small.

#Q17 model 4 and 5
#Q18 I would select model 4 because it is the best fit (lowest AIC), and an additive model is easi er to understand and explain than an interactive model.