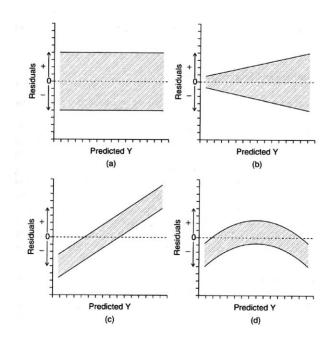
(370	pts	total):
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Section 1 – Short answer

1. (15 pts) Let's say an ecologist wants to model the presence or absence (binary 0/1) of a species as a function of Elevation and Rainfall. Write the appropriate equation(s) to completely describe this model.

2. (15 pts) In Week #8, we discussed several different coding schemes for linear models (dummy coding, effect coding, Helmert contrast coding, etc.). In 1-2 sentences, explain why a modeler may choose to select a particular coding scheme for an analysis. (In other words, why have so many different ways of writing down the same model?)

3. (16 pts) Below are four plots depicting the residuals of a linear model plotted as a function of \hat{Y} .



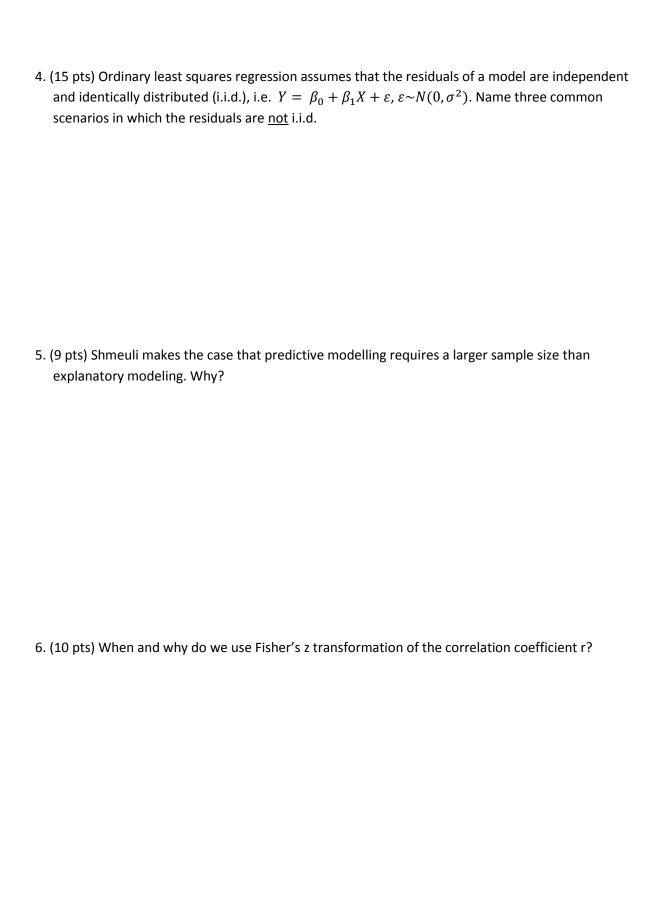
For each panel, state whether the model violates any of the assumptions of linear regression and, if yes, which assumption(s) of linear regression are violated.

Panel a)

Panel b)

Panel c)

Panel d)



7. (15 pts) Fill in the three empty boxes.

Test	Hypothesis (assuming two- tailed tests)	Test statistic T	$f(T H_0)$ (Distribution of T under H_0)	Assumptions
Two sample paired t-test	H_0 : $\mu_A = \mu_B$ H_A : $\mu_A \neq \mu_B$			

8. (15 pts)

a. What is the difference between AIC and likelihood (both mathematically and in terms of how they assess model fit)?

b. How do you use AIC to calculate model weights (include the appropriate equation)?

9. (10 pts) What is the difference between a data point with high leverage a influence?	nd a data point that has high
10. (10 pts) Let's say you have a two-factor crossed ANOVA	
$Y_{ijk} \sim \mu + A_i + B_j + AB_{ij} + \varepsilon_{ijk}$	
and the ANOVA finds that the interaction term is statistically significant. In with interpret a statistically significant interaction between the two factors \boldsymbol{A} and	

Section 2 - Long answer

11. (85 pts)

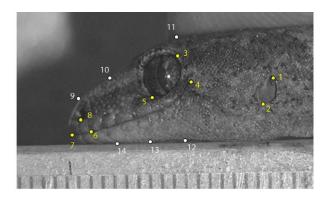
Sexual size and shape dimorphism and allometric scaling patterns in head traits in the New Zealand common gecko Woodworthia maculatus



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Sexual dimorphism (the existence of different sizes or shapes for male vs. female members of the same species) is studied in the New Zealand common gecko by measuring particular points along the head (see Figure).



Let's say the authors were interested in studying head width (HW) as a function of body size (BS). One model that they might use would be the following equation:

$$HW = \alpha \times BS^{\beta}$$

We will call this the "Allometric model".

Another approach would be to take the log of both sides of this equation and fit this equation:

$$\log(HW) = \log(\alpha) + \beta\log(BS)$$

We will call this the "Log-linear model".



c)	Would 'Individual be considered a fixed or a random effect and why?
d)	State the null hypothesis being tested with regards to the effect of 'Individual'.
e)	Why might the researchers want to replicate the measurements on each gecko?
f)	Why would it be wrong to analyze all 40 male and 40 female head width measurements together using a simple t-test?

g) Fill in the gray squares to complete the appropriate ANOVA table for the analysis of HW ~ Sex + Individual. (Since you don't have data, just fill in the appropriate equations in the ANOVA table.) For the p-value column, use the indicated space below the table to fill in the R code you would need to calculate the associated p-value.

Source	Degrees of freedom	SS	MS (leave as ratio)	F-ratio (leave as ratio)	p-value
					write answer below
					write answer below
			N/A	N/A	N/A
Total			N/A	N/A	N/A

p-value for the	first row of	the ANOVA	table =

p-value for the second row of the ANOVA table =

12. (75 pts)

Consider the following model for the relationship between the *Sex Ratio* ("SR"=response) of eggs in a clutch, *Temperature* and *Maternal Age* ("T" and "MA" = covariates). You examined 50 clutches of eggs in the field.

In all cases

$$SR_i \sim Binomial(n_i, p_i)$$

where i represents the ith clutch and n_i is assumed to be known (fixed value, counted in the field).

Model 1: $logit(p) \sim \beta_0 + \beta_1 * T + \beta_2 * MA$; Log-likelihood (Model 1) = -92

Model 2: $logit(p) \sim \beta_0 + \beta_1 * T$; Log-likelihood (Model 2) = -95

Model 3: $ogit(p) \sim \beta_0 + \beta_2 * MA$; Log-likelihood (Model 3) = -99

Model 4: $ogit(p) \sim \beta_0$; Log-likelihood (Model 4) = -100

A) (15 pts) Which combinations of models can be compared by a likelihood ratio test?

B) (15 pts) What is the test statistic and distribution under the null hypothesis for the likelihood ratio test?

C) (24 pts) Fill in the AIC model selection table below (for the last column, feel free to leave the answer as a mathematical expression).

Model	# parameters	AIC	AICc
1			
2			
3			
4			

D) (21 pts) Discuss briefly the pros and cons of using the likelihood ratio test vs. an Information Theoretic approach such as AIC.

13. (40 points) In a linear regression, we model

$$Y_i = \beta_0 + \beta_1 * X_i + \varepsilon_i$$
$$\varepsilon_i \sim N(0, \sigma^2)$$

a. Derive the maximum likelihood estimate of $\widehat{\beta_1}$. (Note: Substantial credit will be awarded for correctly setting up the solution.)

(TOP I	PORTION OF PAGE LE	FT BLANK AS EXTRA SPAC	CE FOR SHOWING YOU!	R WORK.)
b.		s that could be used to e		ure to include the entire R its to the R function.
b.				

Section 3 – Essay (40 pts) Each question can and should be answered succinctly in 4-5 sentences max.)
14a. In Hurlbert (1984), Cox suggests that one possible method of overcoming segregated designs after randomization is to "rerandomize" until some pre-specified amount of interspersion has been achieved. Why would this suggestion be so problematic for Fisher?

14b. What is the difference between the *pre-layout* and *layout specific* alpha? Which is of greater interest to experimenters and why?