GEOG 400/500 Project 2

Name:

First step: Simple linear regressions. Again.

Put an "X" next to your hypothesized direction of association between each Xi and Y (direct or inverse):

Y and X1	Direct	]	Inverse	%cen cty 99 $\rightarrow$ Gunmurd97%				
Y and X2	Direct		Inverse	%gbn Bush→ Gunmurd97%				
Y and X3	Direct	]	Inverse	2K per cap $\rightarrow$ Gunmurd97%				
Y and X4	Direct	]	Inverse	%>24:BA $\rightarrow$ Gunmurd97%				
Y and X5	Direct	]	Inverse	NRA carry99 → Gunmurd97%				
Y and X6	Direct	]	Inverse	% yngmen $\rightarrow$ Gunmurd97%				
Alpha you plan to use to test the null hypotheses, "there is no association between Xi & Y"								
Alpha =								
Justification for the <i>alpha</i> you picked:								
Which associations turned out significant at your chosen alpha level? (mark with an "X")								
And which of the <b>significant</b> associations had the same direction you predicted (direct or inverse)? Mark with an "X." Your								
	Calculated P-Value		Is P-Value < alpha?	Do significant associations	predic- tion			
	4 decimal places		Y or N	match your prediction? Y or N?	I or D?			
Y and X1	Prob-value		Significant?	Prediction correct?				

Y and X2	Prob-value		Significant?		P	rediction correct?	
Y and X3	Prob-value		Significant?		F	rediction correct?	
Y and X4	Prob-value		Significant?		F	rediction correct?	
Y and X5	Prob-value		Significant?		P	rediction correct?	
Y and X6	Prob-value		Significant?		P	rediction correct?	
Interpretation	n/speculations:						
Kitchen sink multiple regression model $\rightarrow$ Y variable is							
				Best X			Ξ
Name the two variables in your <b>best</b> simple linear regression model above: $\rightarrow$ Best X <sub>i</sub> Below, compare the multiple regression model with the best simple linear model (3 decimal places works here):							
Below, comp					ks nere).		
Multiple regression R			Best bivariate	model's R			
Multiple regression R <sup>2</sup> <sub>adj</sub> vs.			Best bivariate model's R <sup>2</sup>				
By comparing movement in R <sup>2</sup> /R <sup>2</sup> <sub>adj</sub> , did you produce a <b>much</b> better explanation of Y with 6 X <sub>i</sub> s? Y or N							]
At 4 decimal places of accuracy, do you see a noticeable improvement in the significance of the model?							]
Whether it was worth your bother or not, please write down your model, showing a and b coëfficients at <b>4</b> decimal places: Don't forget the signs of the b coëfficients and placing the X <sub>1</sub> , X <sub>2</sub> ,X <sub>6</sub> after the coëfficients and before the sign of the next term.							
Y =							
							_

Looking at the t-scores and prob-values for each variable in the kitchen sink model and comparing them with their corresponding values in the simple linear regressions you did earlier, which (if any) have significant p-values once they're allowed to interact?

X <sub>1</sub>	X <sub>2</sub>	X <sub>3</sub>	X <sub>4</sub>	X <sub>5</sub>	X <sub>6</sub>

Put an "X" below the variable(s).

Why is it that several of the variables are significant considered alone (bivariate simple linear regressions) but drop out of significance when they are all put together in a common multiple regression model?

Prune the model of all X variables that have p-values larger than your pre-selected alpha standard. Names of rejected variables:

## Refining the multiple regression model through backwards elimination:

Rerun the multiple regression, but ONLY with the X, variables that still have p-values smaller than alpha in the multiple regression

While R<sup>2</sup> can be expected to decline, the key diagnostics are the changes in significance and in the F statistic that defines it.

Did F:

 Increase
 Decrease
 Stay roughly the same-ish

 Write down your new model (4 decimal places, with proper signs and the original X<sub>i</sub> numbers: e.g., Y= 0.0571 + 0.3452X<sub>2</sub> - 0.3671X<sub>6</sub>

 Y =

 Now, re-examine the new t-scores and p-values. The new regression will have altered them from either the simple linear

regression or the kitchen sink everything-in-it regression. Did the interactions among variables in the new model cause any of your X variables to generate new p-values **higher** than your *alpha* standard (meaning you throw them out in any further round)?

Yes		No		Which?
-	-		-	

Dump any X<sub>i</sub> variable with effects that are no longer significant in comparison with your *alpha* standard. Now, rerun the regression.

Write down your newest model:						
Y =						
What happened to the new(est) R <sup>2</sup> <sub>adj</sub> ?						
Increased	Decreased	Stayed roughly the same				
Dealing with an outlier						
After doing your six scatterplots of each $X_{i}$	on Y, identify the outlier record:					
Redo the kitchen-sink multiple regression (	all X <sub>i</sub> variables at once) but with the outlier	removed.				
Write down this newest kitchen-sink maste	rpiece, leaving X <sub>i</sub> in their original order, co	efficients at 4 decimal places, proper signs:				
Y =						
As before, refine through backwards elimination. Throw out all X <sub>i</sub> variables with p-values above your <i>alpha</i> standard.						
Write down the newest model.						
Y =						
Which of the two outlier-free models (kitchen-sink or the backwards pruned one) has the best F score/significance value?						
Interpretation of the performance of your models, both with and without the outlier. Use sheet below, if necessary.						

Overflow if you need more space: