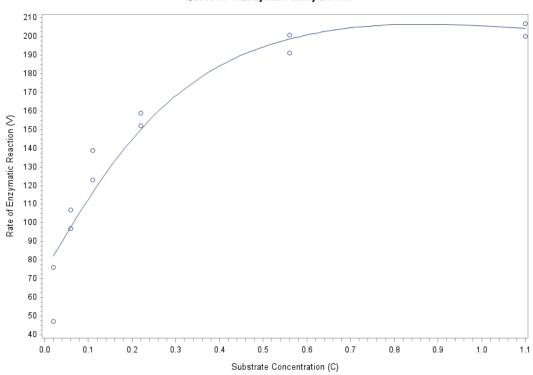
### STAT 512: Homework 3

## Name: Lei Nie

- 1. Consider the following data set that describes the relationship between the rate of an enzymatic reaction (V) and the substrate concentration (C). A common model used to describe the relationship between rate and concentration is the Michaelis-Menten model where,  $\theta_1$  is the maximum rate of the reaction and  $\theta_2$  describes how quickly the reaction will reach its maximum rate. With this mode,  $\frac{1}{V}$  can be written as a linear model with explanatory variable.
  - a. Generate a scatterplot of V vs C. Comment on the shape.

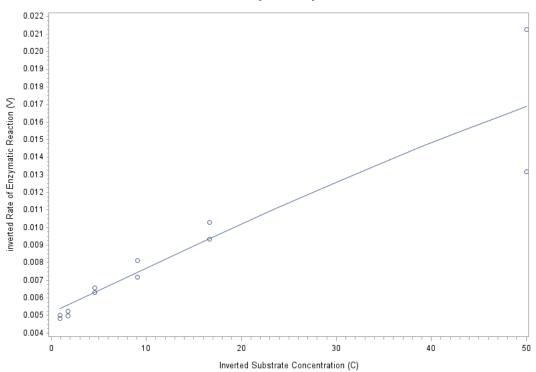
## Scatter plot of Rate(V) vs Concentration(C) (sm=70) Question 1: Enzymatic study-Lei Nie



The shape of the plot does not look linear at all. V increases slower when C is large.

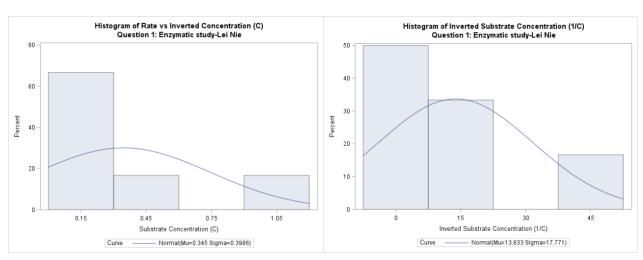
b. Define new variables for  $\frac{1}{V}$  and  $\frac{1}{C}$  in SAS and generate a scatterplot of the new variables. Does the fit appear linear? Do any assumptions appear to be violated?

# Scatter plot of Inverted Rate vs Inverted Concentration (sm=90) Question 1: Enzymatic study-Lei Nie



The line seems to be linear. However, the constant variance assumption is violated since the difference of residuals are wider when  $\frac{1}{c}$  is larger. When  $\frac{1}{c} = 1$ , the residuals are closer than those when  $\frac{1}{c} = 50$ .

c. How is the distribution of  $C^{-1}$  different from the distribution of C? Are there any points that may be more influential in determining the fit?

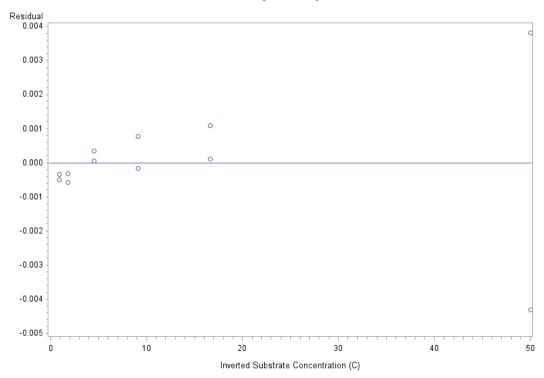


The distribution of C has a mean of 0.345, and standard deviation of 0.3986. The distribution of  $C^{-1}$  has a mean of 13.833, and standard deviation of 17.771. Both distributions are right-skewed, but the distribution of C is more skewed than that of  $C^{-1}$ . There are two influential observations for both histogram. They are corresponding to the two extreme values with C = 1.1 on the left plot, and  $C^{-1} = 50$  on the right plot. However, since  $C^{-1}$  has a smoother distribution than C, the influence of two extreme observations is smaller on the distribution of  $C^{-1}$  than on that of C.

d. Determine the least squares regression line for  $\frac{1}{V}vs\frac{1}{C}$ . Save the residuals and predicted values. Does the residual plot suggest any problems?

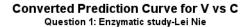
Parameter Estimates					
Variable	DF	Parameter Estimate	Standard Error	t Value	<b>Pr</b> >  t
Intercept	1	0.00511	0.00070400	7.25	<.0001
cinv	1	0.00024722	0.00003210	7.70	<.0001

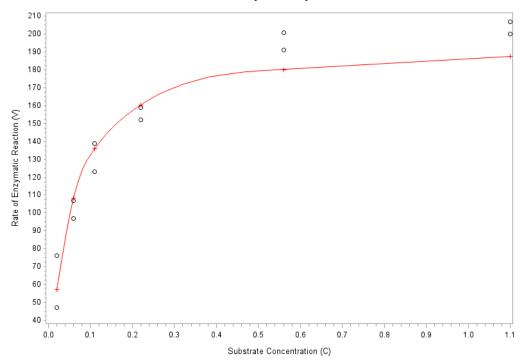
#### Residual Plot for 1/V vs 1/C Question 1: Enzymatic study-Lei Nie



The fitted model is  $\frac{\hat{1}}{v} = 0.00511 + 0.00024722 * \frac{1}{c}$ . The residual plot shows violation to constant variance assumption.

e. Convert this regression line back into the original nonlinear model and plot the predicted curve on a scatterplot of V vs C. Comment on the fit.





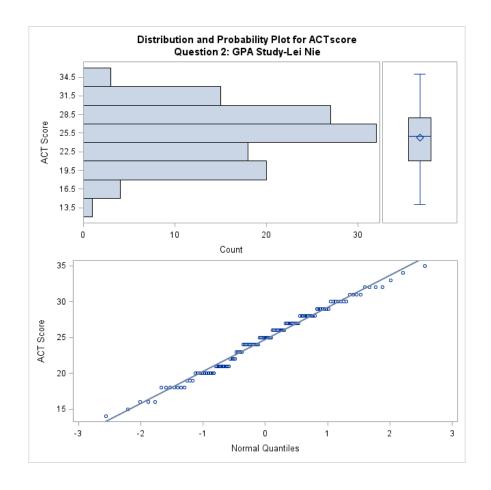
The prediction curve fits the original data pretty well when C is small. When C is large, the prediction value tends to smaller than the observation value.

2. Describe the distribution of the explanatory variable. Show the plots and output that were helpful in learning about this variable.

The UNIVARIATE Procedure Variable: ACTscore (ACT Score)

Basic Statistical Measures				
Location		Variability		
Mean	24.72500	Std Deviation	4.47207	
Median	25.00000	Variance	19.99937	
Mode	24.00000	Range	21.00000	
		Interquartile Range	7.00000	

Extreme Observations			
Lowest		Highest	
Value	Obs	Value	Obs
14	2	32	84
15	48	32	104
16	119	33	15
16	52	34	80
16	32	35	106



There are 120 observed ACT scores in total with range = 21, mean = 24.725, median = 25, standard deviation = 4.47207. There are no extreme influential observations based on the SAS output. The histogram and the box plot both show that the distribution of ACT scores is approximately symmetric and normal. The QQ plot shows the same trend.

- 3. Run the linear regression to predict GPA from the entrance test score and obtain the residuals (do not include a list of the residuals in your solution).
  - a. Verify that the sum of the residuals is zero by running proc univariate with the output from the regression.

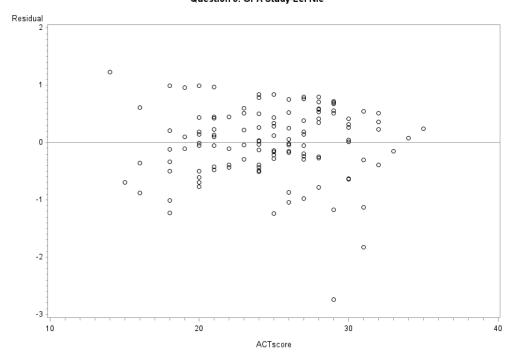
The UNIVARIATE Procedure Variable: resid (Residual)

	variable, resid (residual)			
Moments				
N	120	Sum Weights	120	
Mean	0	Sum Observations	0	
Std Deviation	0.62050134	Variance	0.38502191	
Skewness	-1.0067279	Kurtosis	2.50187662	
Uncorrected SS	45.8176078	Corrected SS	45.8176078	
Coeff Variation		Std Error Mean	0.05664376	

The residual sums to zero.

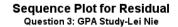
b. Plot the residuals versus the explanatory variable and briefly describe the plot noting any unusual patterns or points.

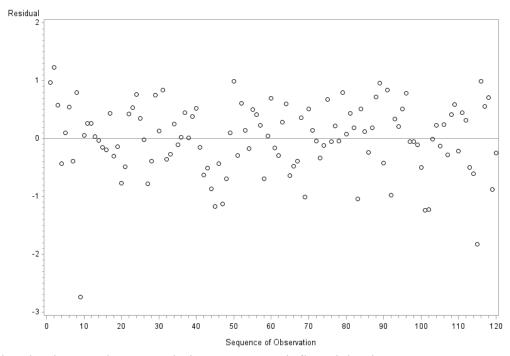
Residual Plot for GPA vs ACTscore Question 3: GPA Study-Lei Nie



The residual plot doesn't show any obvious pattern or influential points.

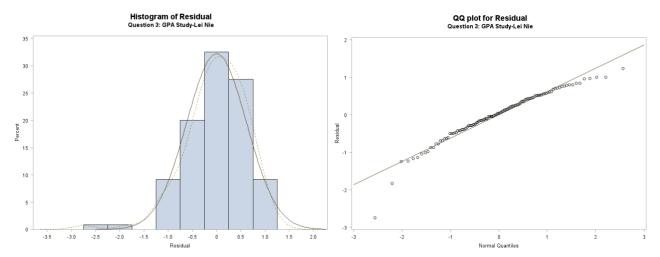
c. Plot the residuals versus the order in which the data appear in the data file and briefly describe the plot noting any unusual patterns or points.





The plot doesn't show any obvious pattern or influential points.

d. Examine the distribution of the residuals by getting a histogram and a normal probability plot of the residuals by using the histogram and qq-plot statements in proc univariate. What do you conclude?



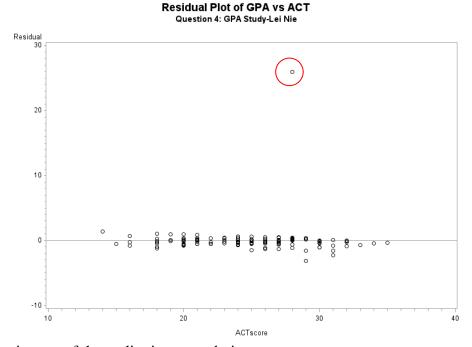
The residual plot seems approximately normal and reasonably symmetric. The QQ plot seems reasonably linear.

- 4. Change the data set by changing the value of the GPA for the last observation from 2.948 to 29.48 (e.g., a typo).
  - a. Make a table comparing the results of this analysis with the results of the analysis of the original data. Include in the table the following: fitted equation, t-test for the slope, with standard error and p-value,  $R^2$ , and the estimate of  $\sigma^2$ . Summarize the differences.

	Original	Туро
fitted equation	$\hat{Y} = 2.114 + 0.0388X$	$\hat{Y} = 1.432 + 0.0753X$
t-test for the slope	3.04	1.48
Standard Error for the slope	0.0128	0.0509
P-value for the slope	0.0029 (Reject)	0.1414 (Fail to reject)
$\mathbb{R}^2$	0.0726	0.0182
Estimate of $\sigma^2$	0.388	6.163

The outlier doubles the slope value and turns the slope from being significant to being insignificant. The  $R^2$  and  $\widehat{\sigma^2}$  change a lot after adding the outlier into the dataset. With the increase of the error variance,  $R^2$  becomes very small.

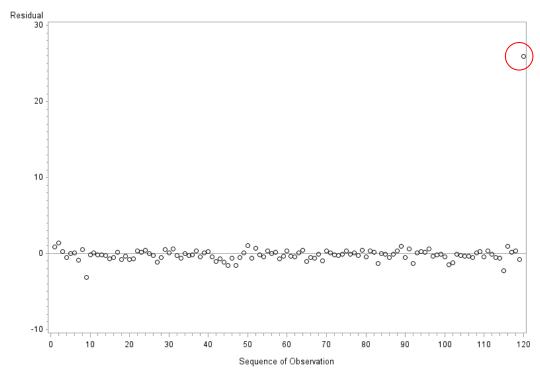
b. Plot the residuals versus the explanatory variable and briefly describe the plot noting any unusual patterns or points.



The existence of the outlier is pretty obvious.

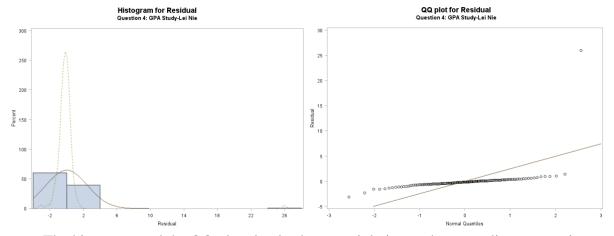
c. Plot the residuals versus the order in which the data appear in the data file and briefly describe the plot noting any unusual patterns or points.





The sequence plot shows that the last observation is the outlier identified in the residual plot above.

d. Examine the distribution of the residuals by getting a histogram and a normal probability plot of the residuals by using the histogram and qq-plot statements in proc univariate. What do you conclude?



The histogram and the QQ plot clearly shows a violation to the normality assumption.

#### Appendix: SAS code

```
*Question 1;
                                           title1 'Histogram of Inverted
data data1;
                                           Substrate Concentration (1/C)';
input C V;
                                           title2 'Question 1: Enzymatic
                                           study-Lei Nie';
cards;
0.02 76
                                           proc univariate data=data1b;
                                           var cinv;
                                         histogram cinv/normal
proc print data=data1;run;
                                          odstitle=title1 odstitle2=title2;
*Question 1a;
                                           label cinv='Inverted Substrate
                                        Concentration (1/C) ';run;
title1 'Scatter plot of Rate(V) vs
Concentration(C) (sm=70)';
                                         ods graphics off;
title2 'Question 1: Enzymatic
study-Lei Nie';
                                           *Ouestion 1d;
axis1 label=('Substrate
                                           proc reg data=data1b;
Concentration (C)');
                                          model vinv=cinv;
Concentration (C)'); model vinv=cinv; axis2 label=(angle=90 'Rate of output out=out1 r=resid p=pred;run; Enzymatic Reaction (V)'); symboll v=circle i=rl;
symbol1 v=circle i=sm70;
                                         title1 'Residual Plot for 1/V vs
proc gplot data=data1;
                                          1/C';
plot V*C/haxis=axis1
                                         title2 'Question 1: Enzymatic
vaxis=axis2;run;
                                         study-Lei Nie';
                                          axis1 label=('Inverted Substrate
*Question 1b;
                                          Concentration (C)');
data data1b;
                                         proc gplot data=out1;
set data1;
                                          plot resid*cinv/vref=0
vinv=1/V;
                                          haxis=axis1;run;
cinv=1/C; run;
title1 'Scatter plot of Inverted
                                         *Question 1e;
Rate vs Inverted Concentration
                                         data invert;
(sm=90)';
                                           set out1;
title2 'Question 1: Enzymatic
                                         predv = 1/pred;
                                           symbol1 v = circle i = none c =
study-Lei Nie';
axis1 label=('Inverted Substrate
                                         black;
Concentration (C)');
                                          symbol2 v = plus i = sm5 c = red;
axis2 label=(angle=90 'inverted
                                          title1 'Converted Prediction Curve
Rate of Enzymatic Reaction (V)');
                                          for V vs C';
                                           title2 'Question 1: Enzymatic
symbol1 v=circle i=sm90;
proc gplot data=data1b;
                                          study-Lei Nie';
plot vinv*cinv/haxis=axis1
                                          axis1 label=('Substrate
vaxis=axis2;run;
                                           Concentration (C)');
                                           axis2 label=(angle=90 'Rate of
*Ouestion 1c;
                                           Enzymatic Reaction (V)');
ods graphics on;
                                           proc gplot data = invert;
title1 'Histogram of Substrate
                                           plot V*C predv*C /haxis=axis1
Concentration (C)';
                                           vaxis=axis2 overlay;run;
title2 'Question 1: Enzymatic
study-Lei Nie';
proc univariate data=data1;
                                           *Ouestion 2/3/4;
var C;
                                           data data2;
histogram C/normal odstitle=title1
                                           input GPA ACTscore;
odstitle2=title2;
                                           seq=_n_;
label C='Substrate Concentration
                                           datalines;
(C) '; run;
                                           3.897 21
```

```
2.948 28
                                           *Question 4a;
                                          proc reg data=data4;
 proc print data=data2;run;
                                          model GPA=ACTscore;
                                          output out=out4 r=resid p=pred;run;
*Ouestion 2;
ods graphics on;
                                          *Question 4b;
proc univariate data=data2
                                          title1 'Residual Plot of GPA vs
plot(odstitle2='Question 2: GPA
                                          ACT':
                                          title2 'Question 4: GPA Study-Lei
Study-Lei Nie');
var ACTscore;
                                          Nie';
label ACTscore='ACT Score';run;
                                          proc gplot data=out4;
ods graphics off;
                                          plot resid*ACTscore / vref=0;run;
*Question 3a;
                                          *Ouestion 4c;
proc reg data=data2;
                                          title1 'Sequence Plot for Residual';
model GPA=ACTscore;
                                          title2 'Question 4: GPA Study-Lei
output out=out2 r=resid p=pred;run;
                                          Nie';
proc univariate data=out2;
                                          axis1 label=('Sequence of
var resid; run;
                                          Observation');
                                          proc gplot data=out4;
*Question 3b;
                                          plot resid*seq/ haxis=axis1
title1 'Residual Plot for GPA vs
                                          vref=0;run;
ACTscore';
                                          *Question 4d;
title2 'Question 3: GPA Study-Lei
Nie';
                                          title1 'QQ plot for Residual';
                                          title2 'Question 4: GPA Study-Lei
proc gplot data=out2;
plot resid*ACTscore / vref=0;run;
                                          proc univariate data=out4;
                                          var resid;
                                          histogram resid / normal
*Ouestion 3c;
                                          kernel(L=2);
title1 'Sequence Plot for Residual';
title2 'Question 3: GPA Study-Lei
                                          qqplot resid /normal (L=1 mu=est
Nie';
                                          sigma=est);run;
axis1 label=('Sequence of
Observation');
proc gplot data=out2;
plot resid*seq/ haxis=axis1
vref=0;run;
*Question 3d;
title1 'QQ plot for Residual';
title2 'Question 3: GPA Study-Lei
proc univariate data=out2;
var resid;
histogram resid / normal
kernel(L=2);
qqplot resid /normal (L=1 mu=est
sigma=est);run;
*Ouestion 4;
data data4;
set data2;
if seq eq 120 then GPA = 29.48;
proc print data=data4;run;
```