

MA4710
FINAL PROJECT
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12/13/2022

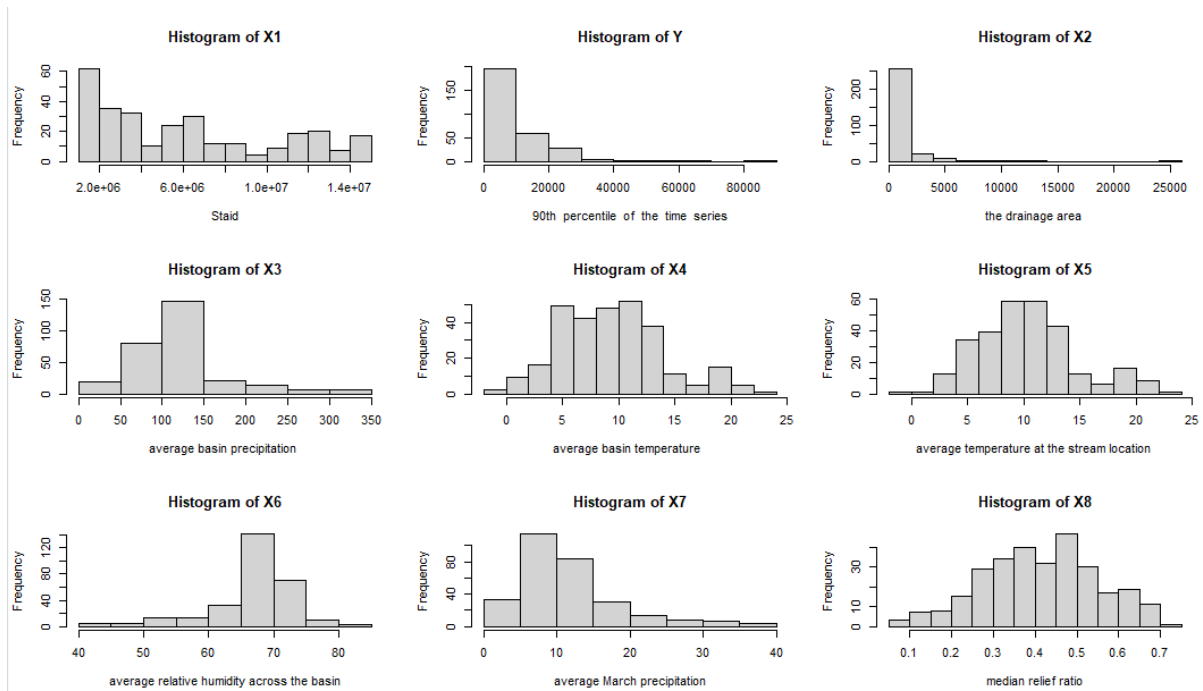
1. INTRODUCTION:

We were provided with a dataset called **streamflow.csv** which contains the 90th percentile maximum streamflow. It has 294 variables and 9 variables. The following is the description of the variables of the dataset:

- The response variable, Y (max90), is the 90th percentile of the time series of annual daily maxima.
- X1 (STAID) is, the stream identification number.
- X2 (DRAIN_SQKM) the drainage area.
- X3(PPTAVG_BASIN) the average basin precipitation.
- X4(T_AVG_BASIN) the average basin temperature.
- X5(T_AVG_SITE) the average temperature at the stream location.
- X6(RH_BASIN) the average relative humidity across the basin.
- X7(MAR_PPT7100_CM) the average March precipitation.
- X8(RRMEDIAN) the median relief ratio.

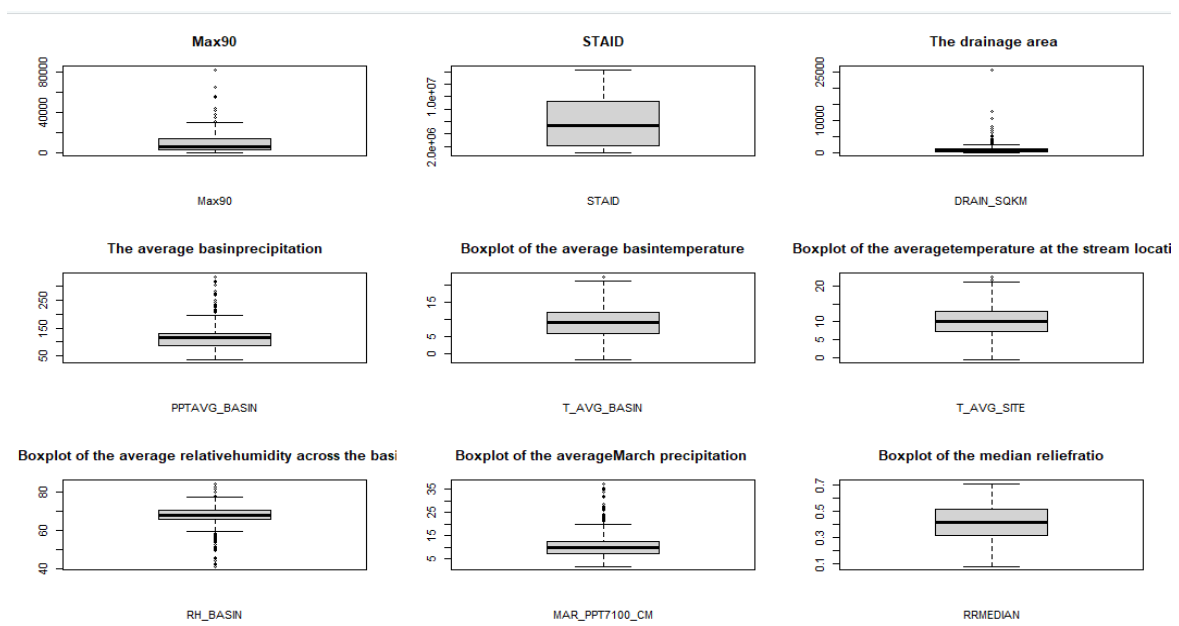
```
< summary(streamflow) ~ summary(Y, X1, X2, X3, X4, X5, X6, X7, X8)
> summary(streamflow)
      x1      Y      x2      x3
Min.   : 1013500 Min.   : 16.03 Min.   : 5.377 Min.   : 37.78
1st Qu.: 2065500 1st Qu.: 2231.00 1st Qu.: 208.686 1st Qu.: 88.46
Median : 5362000 Median : 5646.00 Median : 450.199 Median :114.68
Mean   : 5940630 Mean   : 9272.69 Mean   : 1102.691 Mean   :120.17
3rd Qu.: 9223000 3rd Qu.:13670.00 3rd Qu.: 1151.567 3rd Qu.:131.41
Max.   :14325000 Max.   :81900.00 Max.   :25791.040 Max.   :334.17
      x4      x5      x6      x7
Min.   : -1.580 Min.   : -0.40 Min.   :41.11 Min.   : 1.739
1st Qu.: 5.908 1st Qu.: 7.30 1st Qu.:65.74 1st Qu.: 7.304
Median : 9.044 Median :10.00 Median :67.79 Median : 9.876
Mean   : 9.415 Mean   :10.34 Mean   :66.69 Mean   :11.408
3rd Qu.:12.189 3rd Qu.:12.90 3rd Qu.:70.24 3rd Qu.:12.261
Max.   :22.500 Max.   :22.50 Max.   :84.20 Max.   :37.370
      x8
Min.   :0.08042
1st Qu.:0.31652
Median :0.41379
Mean   :0.41466
3rd Qu.:0.51370
Max.   :0.71084
> |
```

Above is the summary of all the numeric variables (Y and X1 through X8) provided in the data set streamflow.

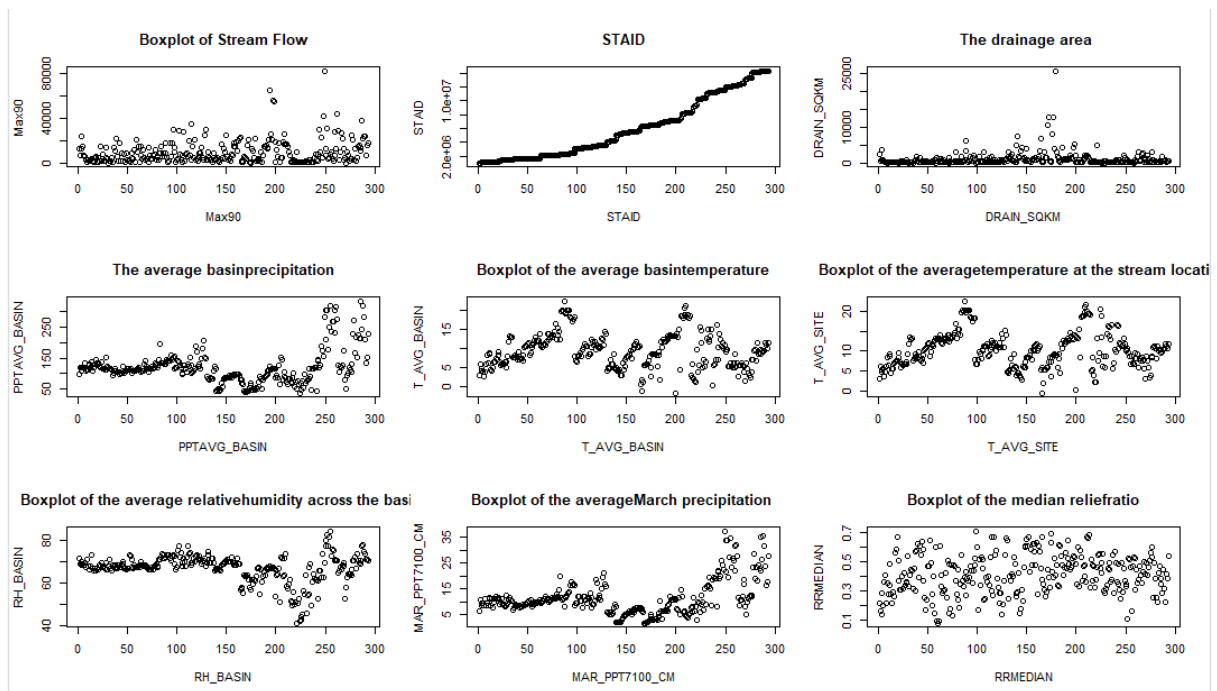


From the above figures we can interpret that,

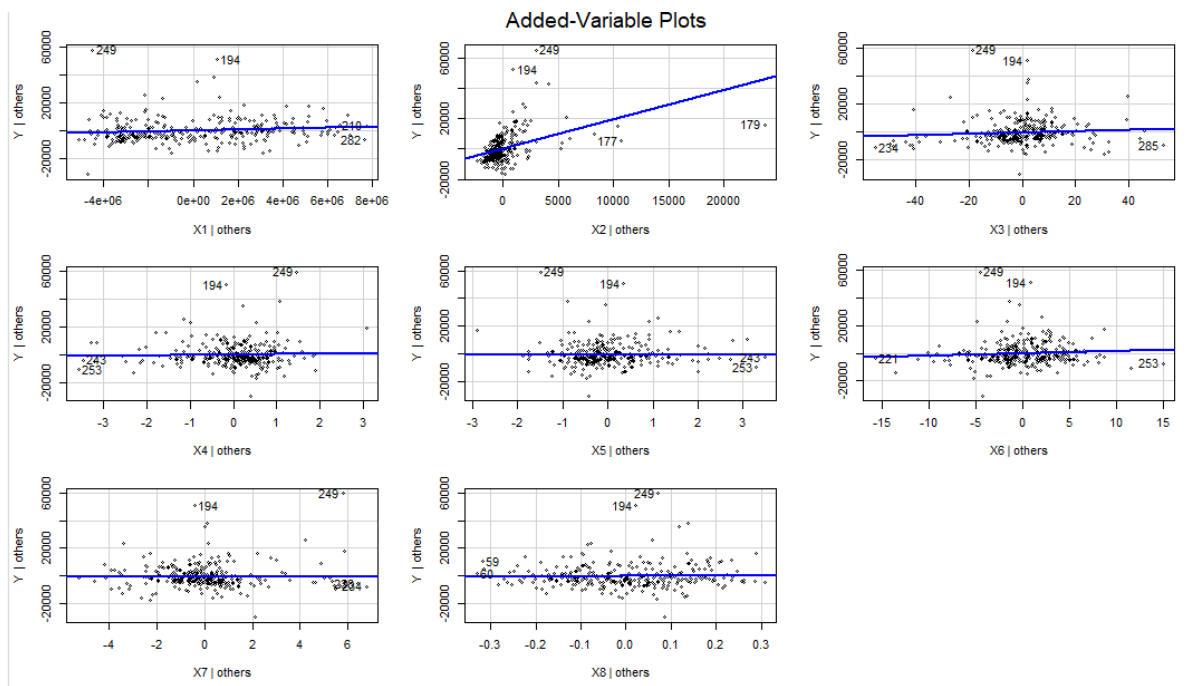
- The response variable Y has a distinct left skew.
- The predictor variable X1, has a uniform skew.
- The predictor variable X2, has a strong left skew.
- The predictor variable X3, has a left skew.
- The predictor variable X4, has a normal distribution.
- The predictor variable X5, has a symmetric distribution.
- The predictor variable X6, has a strong right skew.
- The predictor variable X7, has a left skew.
- The predictor variable X8 has a normal distribution.



Above are the box plots of all the numeric variables (Y and X1 through X8) provided in the data set streamflow.



Above are the scatter plots of all the numeric variables (Y and X1 through X8) provided in the data set streamflow.

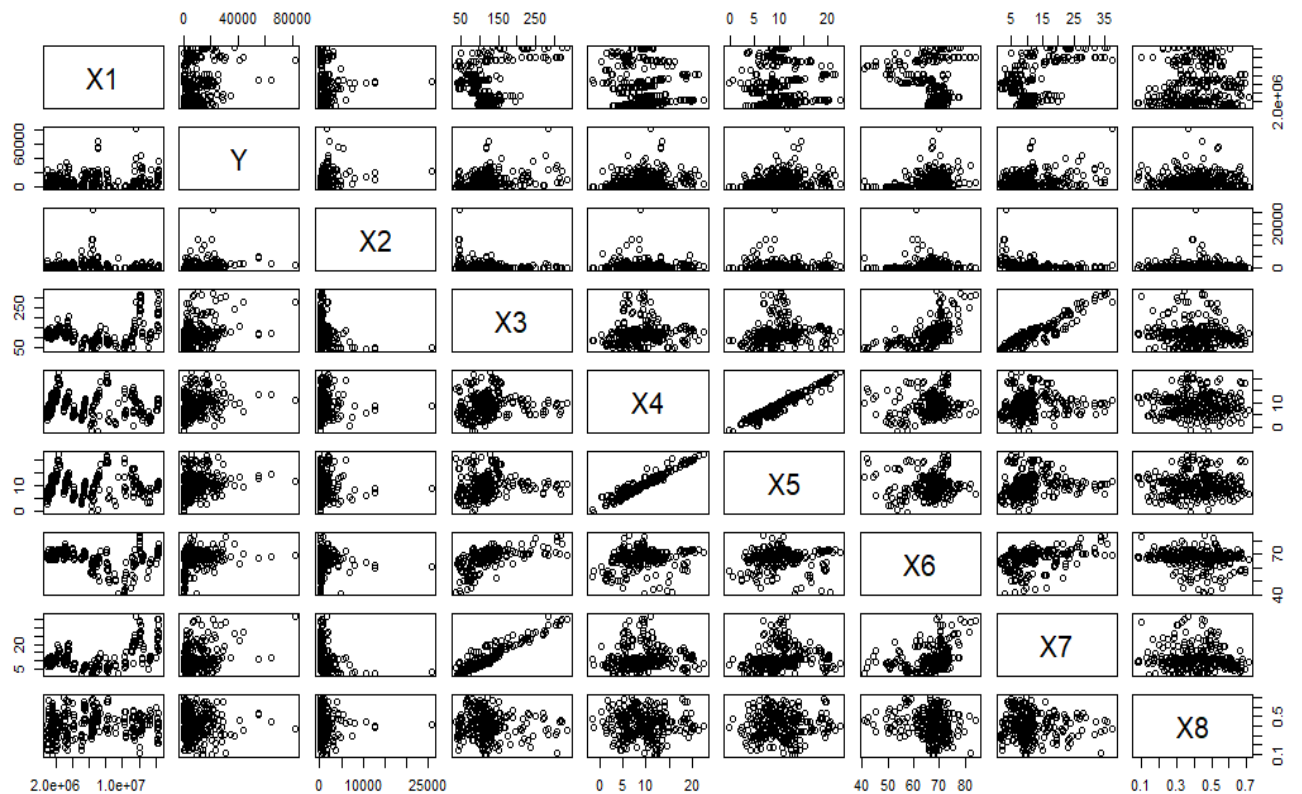


Above are the added-variable plots of all the numeric variables (Y and X1 through X8) provided in the data set streamflow.

```
> cor(streamflow)
```

	x1	Y	x2	x3	x4	x5	x6
x1	1.00000000	0.1926035	0.02097545	0.30978548	-0.15557066	-0.04759166	-0.21430059
Y	0.19260350	1.00000000	0.31807850	0.29602977	0.20550773	0.21113865	0.21802869
x2	0.02097545	0.3180785	1.00000000	-0.24704239	-0.03020605	-0.04769574	-0.08566400
x3	0.30978548	0.2960298	-0.24704239	1.00000000	0.07752031	0.10881444	0.55728901
x4	-0.15557066	0.2055077	-0.03020605	0.07752031	1.00000000	0.96818515	0.19074913
x5	-0.04759166	0.2111386	-0.04769574	0.10881444	0.96818515	1.00000000	0.09235669
x6	-0.21430059	0.2180287	-0.08566400	0.55728901	0.19074913	0.09235669	1.00000000
x7	0.48388068	0.2829461	-0.25355037	0.92688029	0.08247133	0.14754361	0.35554378
x8	0.11320656	-0.0116222	-0.02146808	-0.11035338	-0.01089296	0.02374341	-0.16804586

	x7	x8
x1	0.48388068	0.11320656
Y	0.28294614	-0.01162220
x2	-0.25355037	-0.02146808
x3	0.92688029	-0.11035338
x4	0.08247133	-0.01089296
x5	0.14754361	0.02374341
x6	0.35554378	-0.16804586
x7	1.00000000	-0.10848844
x8	-0.10848844	1.00000000



From the above correlation matrix and the plot, we can observe that there is not extreme multicollinearity problem between the variables.

2. MODELS AND METHODS:

Now we can fit our preliminary model. Our preliminary model will simply be our response variable Y (max90) regressed against all the predictor variables in our data set.

```
Call:
lm(formula = Y ~ X1 + X2 + X3 + X4 + X5 + X6 + X7 + X8, data = streamflow)

Residuals:
    Min       1Q   Median       3Q      Max
-29981  -4579  -1538    2868   59389

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -1.765e+04  8.178e+03  -2.159   0.0317 *
X1           3.323e-04  1.745e-04   1.904   0.0579 .
X2           1.942e+00  2.532e-01   7.671 2.75e-13 ***
X3           4.969e+01  3.527e+01   1.409   0.1600
X4           3.440e+02  5.819e+02   0.591   0.5549
X5           1.287e+02  6.068e+02   0.212   0.8322
X6           1.603e+02  1.305e+02   1.228   0.2203
X7           5.115e+01  2.751e+02   0.186   0.8526
X8           2.405e+03  4.039e+03   0.595   0.5521
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 8999 on 284 degrees of freedom
Multiple R-squared:  0.3009,    Adjusted R-squared:  0.2812
F-statistic: 15.28 on 8 and 284 DF,  p-value: < 2.2e-16
```

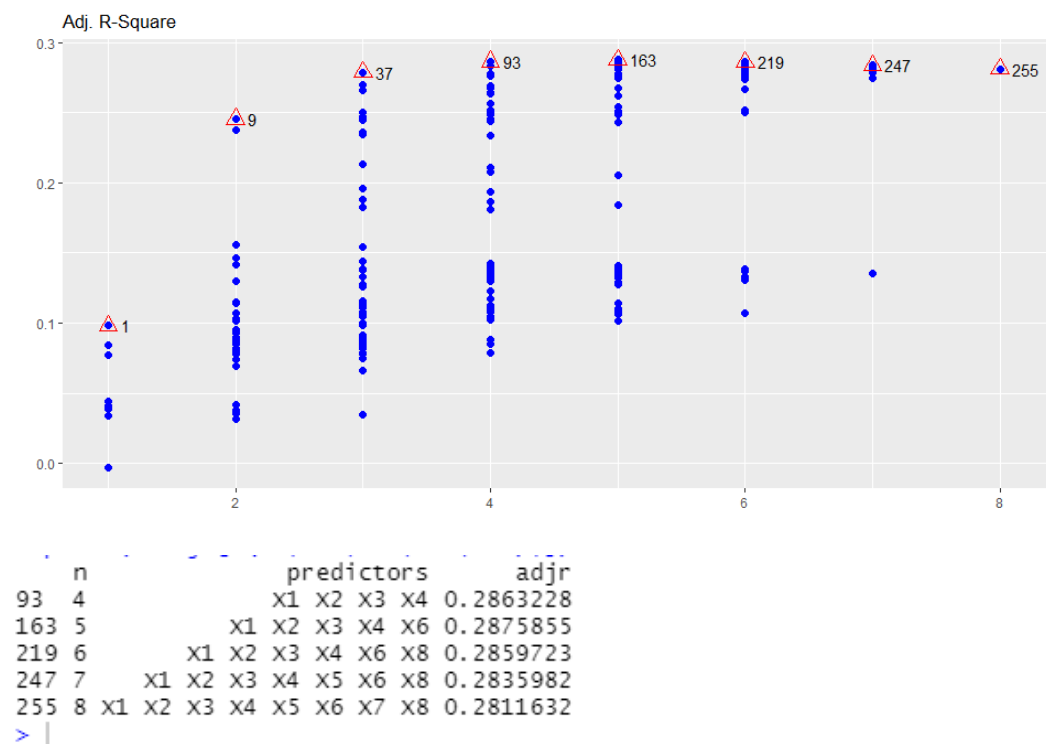
```
> anova(fitstream)
Analysis of Variance Table
```

```
Response: Y
      Df    Sum Sq   Mean Sq F value    Pr(>F)
X1      1 1.2203e+09 1220348895 15.0689 0.000129 ***
X2      1 3.2457e+09 3245732542 40.0783 9.495e-10 ***
X3      1 3.9125e+09 3912466049 48.3112 2.494e-11 ***
X4      1 1.3622e+09 1362215144 16.8206 5.370e-05 ***
X5      1 6.7370e+06   6736980   0.0832 0.773233
X6      1 1.2057e+08  120570374   1.4888 0.223414
X7      1 5.1717e+05    517174   0.0064 0.936363
X8      1 2.8703e+07   28703189   0.3544 0.552092
Residuals 284 2.3000e+10  80984724
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

We can see that the model is significant, but some of the individual predictors are not significant. Running a best subset and stepwise regression on this full model results in the following. First, we will look at the best subset for each number of predictor variables selected based on the highest R2 adj.

R2 Adjusted:

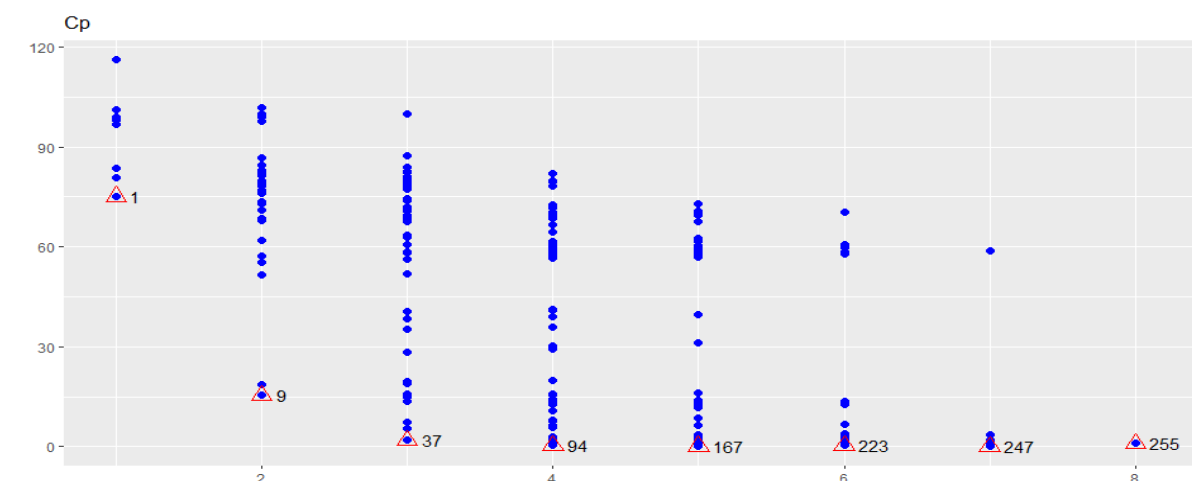
To get a better idea of where the R2 adj peaks we can look at a plot of the R2 adj against the number of predictors.



Now, let's perform the same by using CP, AIC, BIC.

CP:

Then plotting the CP against the number of predictors.



Then getting our best subset.

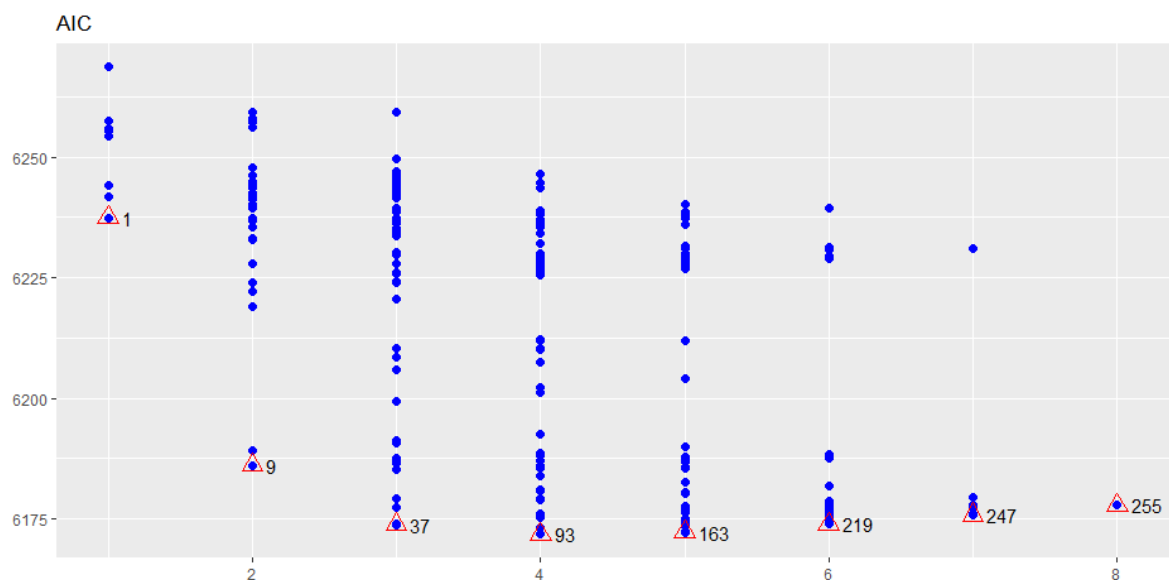
```

n      predictors      cp
93  4      x1 x2 x3 x4 2.932805
163 5      x1 x2 x3 x4 x6 3.435856
219 6      x1 x2 x3 x4 x6 x8 5.086642
247 7      x1 x2 x3 x4 x5 x6 x8 7.034584
255 8      x1 x2 x3 x4 x5 x6 x7 x8 9.000000

```

AIC:

Plotting AIC against number of predictors.



Then getting our best subset.

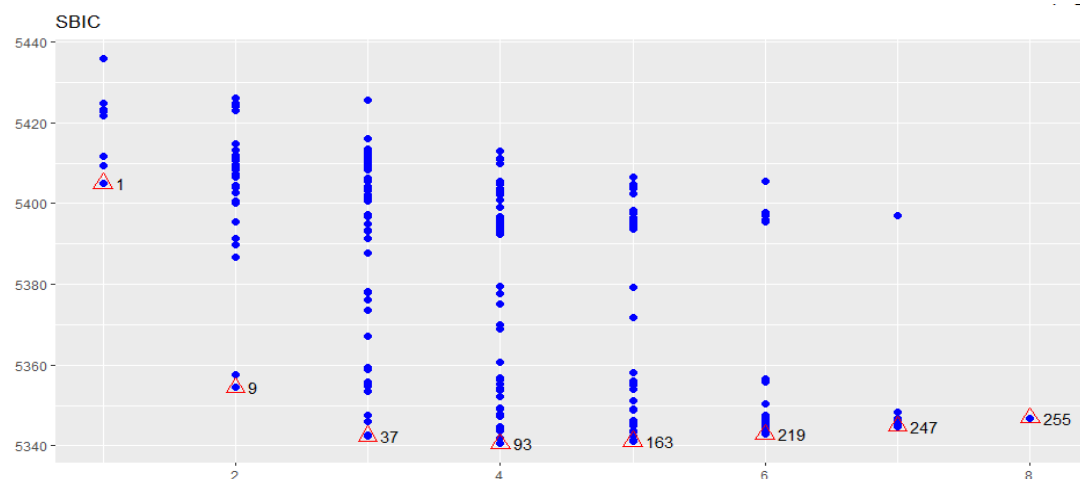
```

n      predictors      aic
93  4      x1 x2 x3 x4 6171.807
163 5      x1 x2 x3 x4 x6 6172.269
219 6      x1 x2 x3 x4 x6 x8 6173.909
247 7      x1 x2 x3 x4 x5 x6 x8 6175.855
255 8      x1 x2 x3 x4 x5 x6 x7 x8 6177.820
> |

```

BIC:

Plotting BIC against number of predictors.



Then getting our best subset.

```

      n      predictors      bic
93  4      x1 x2 x3 x4 5340.555
163 5      x1 x2 x3 x4 x6 5341.131
219 6      x1 x2 x3 x4 x6 x8 5342.848
247 7      x1 x2 x3 x4 x5 x6 x8 5344.861
255 8      x1 x2 x3 x4 x5 x6 x7 x8 5346.890
> |

```

And finally, a stepwise regression

We can see that all our procedures agree on a model. The best subset model based on R2 adj, CP, AIC, and BIC contain predictors X2, X3 and X5.

we are selecting variables based on p value...

Stepwise Selection: Step 1

+ x2

Model Summary							
R	0.318	RMSE				10080.200	
R-Squared	0.101	Coef. Var				108.708	
Adj. R-Squared	0.098	MSE				101610439.565	
Pred R-Squared	0.048	MAE				6897.294	
RMSE: Root Mean Square Error							
MSE: Mean Square Error							
MAE: Mean Absolute Error							
ANOVA							
	Sum of Squares	DF	Mean Square	F	Sig.		
Regression	3328314064.255	1	3328314064.255	32.756	0.0000		
Residual	29568637913.443	291	101610439.565				
Total	32896951977.699	292					
Parameter Estimates							
model	Beta	Std. Error	Std. Beta	t	sig	lower	upper
(Intercept)	7581.312	658.885		11.506	0.000	6284.527	8878.097
x2	1.534	0.268	0.318	5.723	0.000	1.006	2.061

Stepwise Selection: Step 2

+ x3

Model Summary							
R	0.501	RMSE				9219.937	
R-Squared	0.251	Coef. Var				99.431	
Adj. R-Squared	0.245	MSE				85007236.614	
Pred R-Squared	0.161	MAE				6158.189	
RMSE: Root Mean Square Error							
MSE: Mean Square Error							
MAE: Mean Absolute Error							
ANOVA							
	Sum of Squares	DF	Mean Square	F	Sig.		
Regression	8244853359.704	2	4122426679.852	48.495	0.0000		
Residual	24652098617.994	290	85007236.614				
Total	32896951977.699	292					
Parameter Estimates							
model	Beta	Std. Error	Std. Beta	t	sig	lower	upper
(Intercept)	-2286.243	1430.630		-1.598	0.111	-5101.977	529.492
x2	2.009	0.253	0.417	7.942	0.000	1.511	2.507
x3	77.753	10.224	0.399	7.605	0.000	57.631	97.876

Stepwise Selection: Step 3

+ X5

Model Summary							
R	0.535	RMSE		9013.625			
R-Squared	0.286	Coef. Var		97.206			
Adj. R-Squared	0.279	MSE		81245443.753			
Pred R-Squared	0.193	MAE		5926.228			
RMSE: Root Mean Square Error							
MSE: Mean Square Error							
MAE: Mean Absolute Error							
ANOVA							
	Sum of Squares	DF	Mean Square	F	Sig.		
Regression	9417018733.194	3	3139006244.398	38.636	0.0000		
Residual	23479933244.504	289	81245443.753				
Total	32896951977.699	292					
Parameter Estimates							
model	Beta	Std. Error	Std. Beta	t	sig	lower	upper
(Intercept)	-6708.252	1819.746		-3.686	0.000	-10289.887	-3126.617
x2	2.029	0.247	0.421	8.204	0.000	1.543	2.516
x3	73.928	10.046	0.379	7.359	0.000	54.156	93.700
x5	470.125	123.771	0.190	3.798	0.000	226.518	713.731

Since, we do not have any multicollinearity problem with the variables, we can proceed with the final model.

No more variables to be added/removed.

Final Model output

Model Summary							
R	0.535	RMSE		9013.625			
R-Squared	0.286	Coef. Var		97.206			
Adj. R-Squared	0.279	MSE		81245443.753			
Pred R-Squared	0.193	MAE		5926.228			
RMSE: Root Mean Square Error							
MSE: Mean Square Error							
MAE: Mean Absolute Error							
ANOVA							
	Sum of Squares	DF	Mean Square	F	Sig.		
Regression	9417018733.194	3	3139006244.398	38.636	0.0000		
Residual	23479933244.504	289	81245443.753				
Total	32896951977.699	292					
Parameter Estimates							
model	Beta	Std. Error	Std. Beta	t	sig	lower	upper
(Intercept)	-6708.252	1819.746		-3.686	0.000	-10289.887	-3126.617
x2	2.029	0.247	0.421	8.204	0.000	1.543	2.516
x3	73.928	10.046	0.379	7.359	0.000	54.156	93.700
x5	470.125	123.771	0.190	3.798	0.000	226.518	713.731

```

Call:
lm(formula = Y ~ X2 + X3 + X5, data = streamflow)

Residuals:
    Min       1Q   Median       3Q      Max
-31394  -4965  -1404    2619   59010

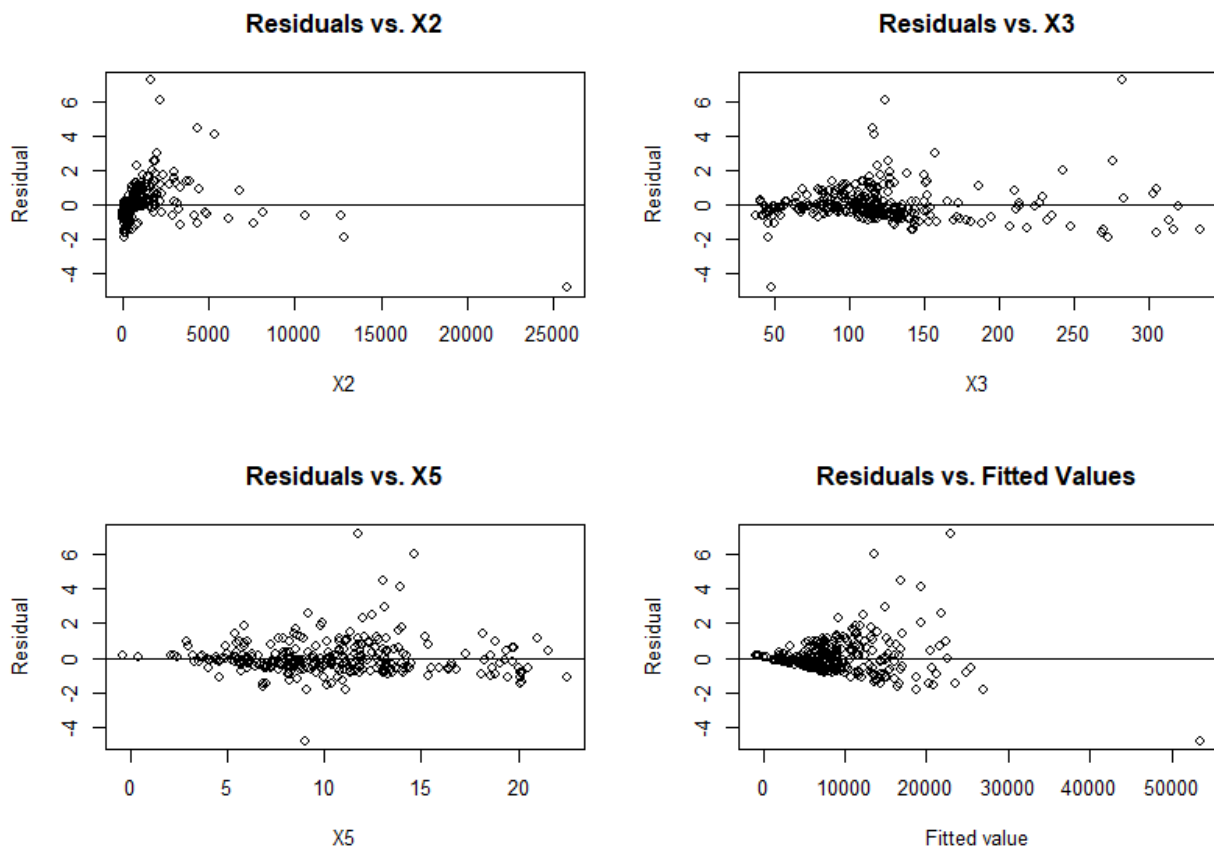
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -6708.2520  1819.7455  -3.686 0.000272 ***
X2           2.0295    0.2474   8.204 7.73e-15 ***
X3          73.9282    10.0458   7.359 1.93e-12 ***
X5         470.1248    123.7708   3.798 0.000178 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 9014 on 289 degrees of freedom
Multiple R-squared:  0.2863,    Adjusted R-squared:  0.2788
F-statistic: 38.64 on 3 and 289 DF,  p-value: < 2.2e-16

```

There is not much multicollinearity in the final model as we can observe.

Now, lets perform regression diagnostics and get the residual plots.



By using the vif built-in function we can confirm that there is not multicollinearity in the reduced model (X2+X3+X5) as well

```

      x2      x3      x5
1.065494 1.075808 1.012455

```

To confirm what we see in the above plots we can run a Breusch-Pagan test and Durbin-Watson test, where the p value in both cases is not greater than 0.05 so, we cannot retain our null assumption of independence.

```
> bptest(reduced.lmfit)
```

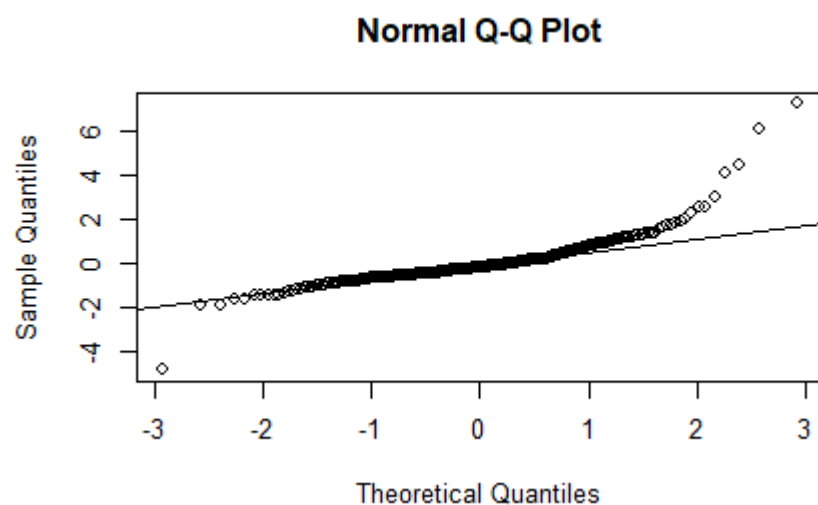
```
studentized Breusch-Pagan test
```

```
data: reduced.lmfit
BP = 39.762, df = 3, p-value = 1.197e-08
```

```
Durbin-watson test
```

```
data: fitstream
DW = 1.3968, p-value = 4.424e-08
alternative hypothesis: true autocorrelation is not 0
```

To assess normality we will start by looking at a Q-Q Plot of our residuals.



We can clearly see that there is a right skew primarily on the right side of the distribution. By using a Shapiro-Wilk test we can test for the likelihood of this distribution under the assumption of normality.

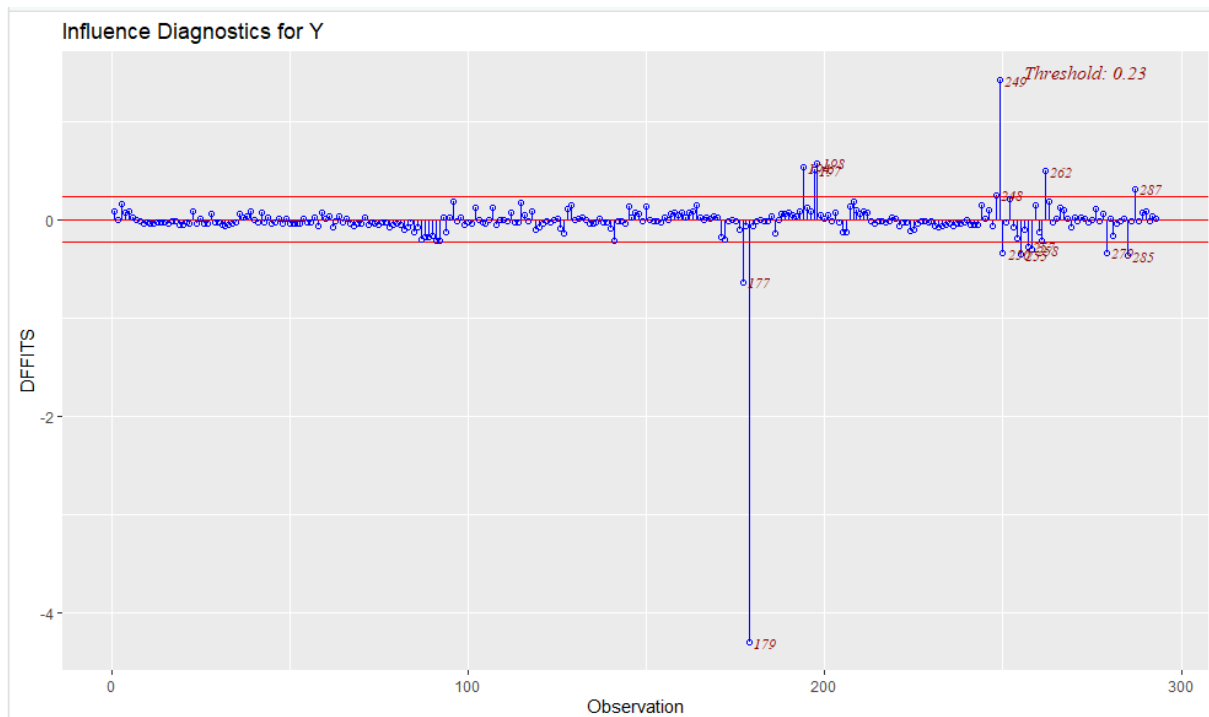
```
shapiro-wilk normality test
```

```
data: res
W = 0.80758, p-value < 2.2e-16
```

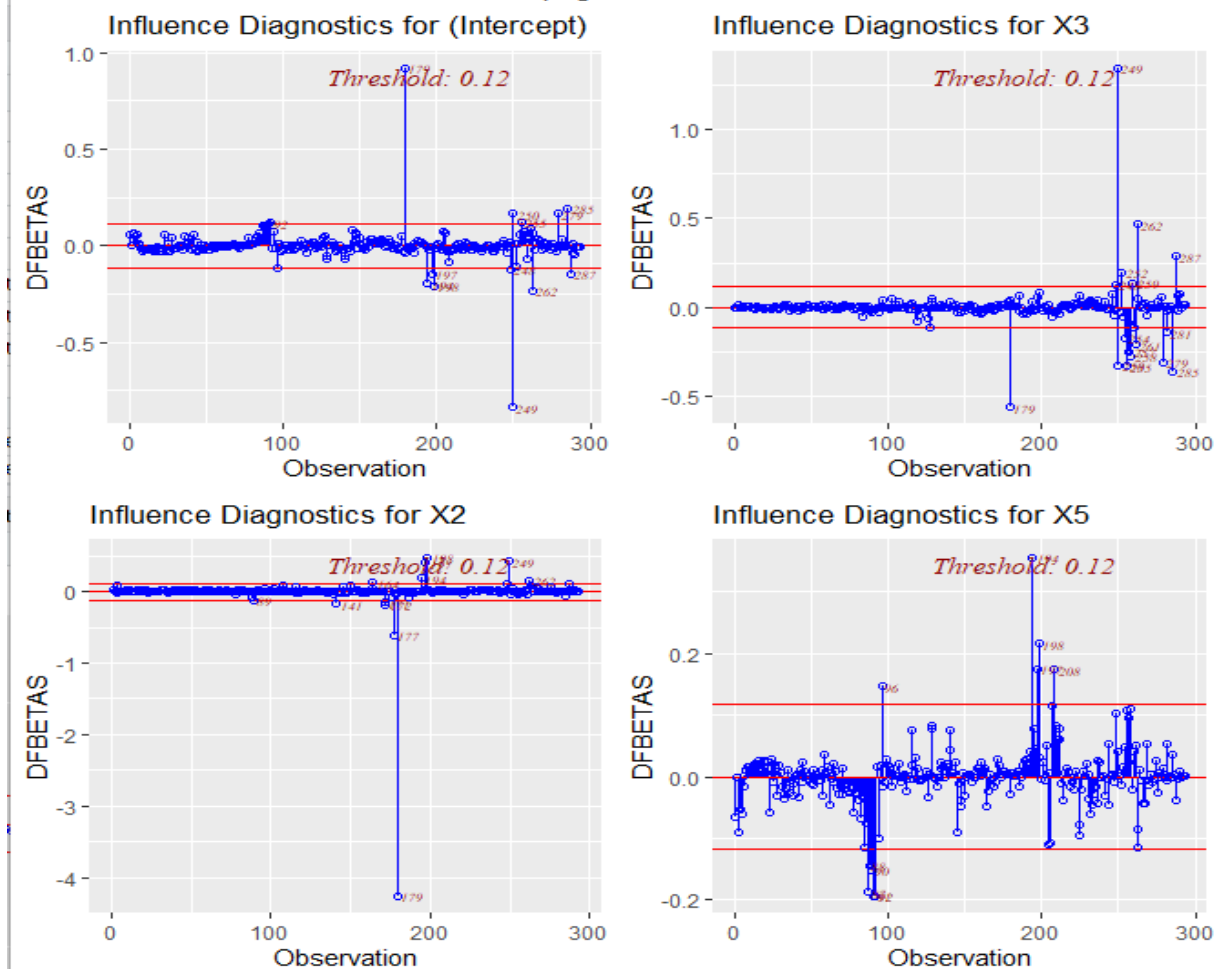
The p-value of 0.000000000000000022 means that this distribution has a 0.000000000000000022% chance of occurring if the population is normally distributed.

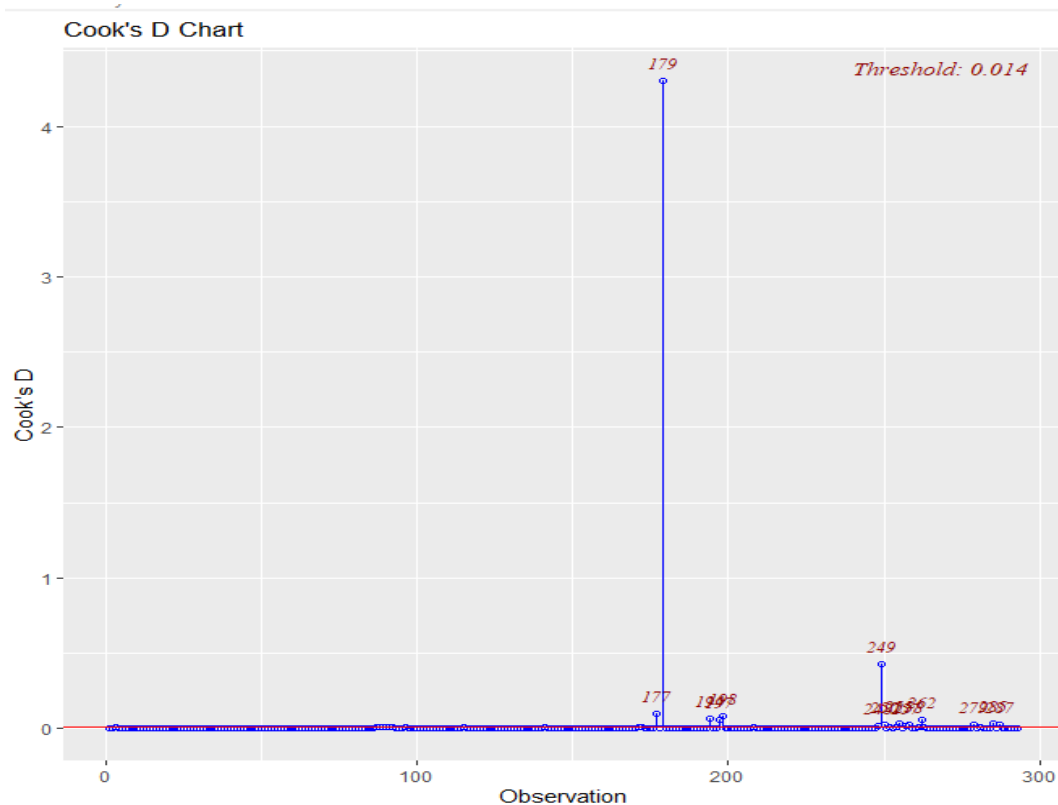
There is clearly a problem with our assumption of normality.

Finally, we will look at outliers using DFBETAS, DFFITS, and Cook's Distance plots.



page 1 of 1





Within our outliers we have observations 179 and 249 which are also two of the points that heavily influence the right skew previously seen in our Q-Q plot.

We have one assumption to remedy with our model, which is our assumption of normality. We will attempt to resolve this by transforming our model with a **Box-Cox Transformation**.

```
> lambda
[1] 0.2792849
```

We get that our optimized model has $\lambda = 0.2792849$. We then raise our response variable Y to λ and fit our model with our transformed Y.

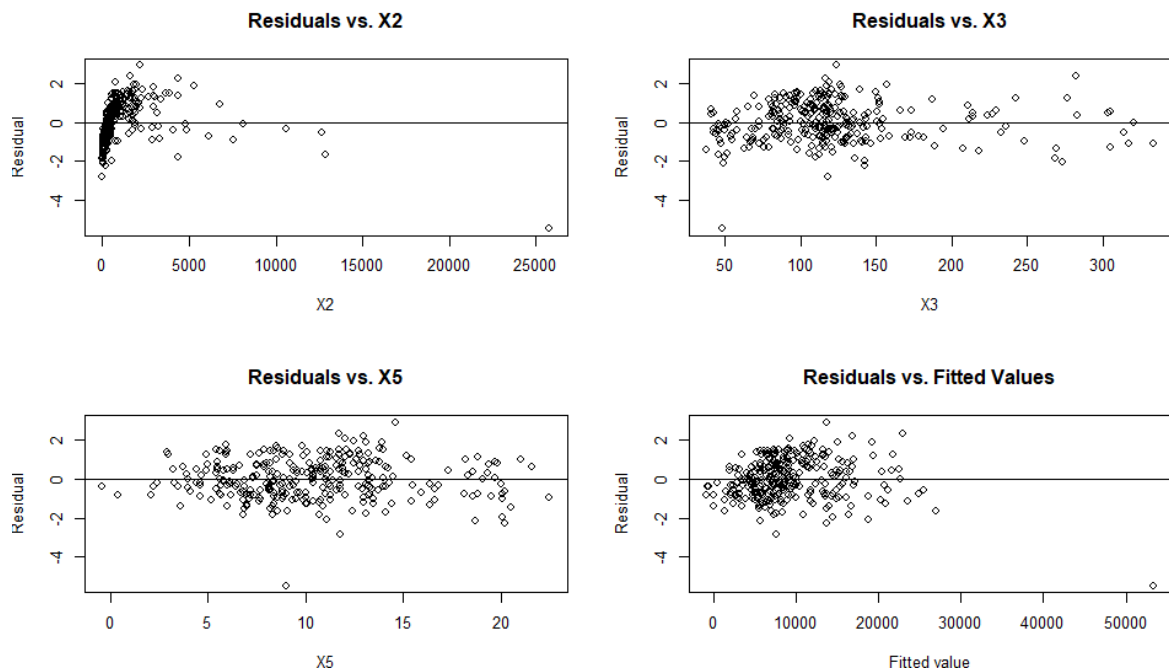
```
Call:
lm(formula = trans.Y ~ X2 + X3 + X5, data = streamflow)

Residuals:
    Min       1Q   Median       3Q      Max
-12.0998  -2.2829  -0.1921   2.1780   8.9177

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  5.043e+00  6.227e-01   8.098 1.58e-14 ***
X2           7.833e-04  8.466e-05   9.253 < 2e-16 ***
X3           2.766e-02  3.438e-03   8.046 2.24e-14 ***
X5           2.058e-01  4.236e-02   4.860 1.93e-06 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.085 on 289 degrees of freedom
Multiple R-squared:  0.3414,    Adjusted R-squared:  0.3346
F-statistic: 49.94 on 3 and 289 DF,  p-value: < 2.2e-16
```

The R^2 adj went up, but if our model satisfies all necessary assumptions it is a better model. Let's take a look at assumption diagnostics again, starting with constancy of variance.



studentized Breusch-Pagan test

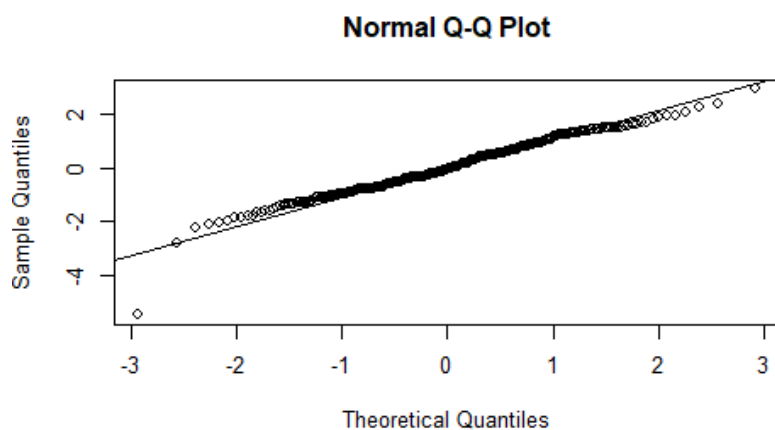
```
data: boxcox.lmfit
BP = 67.678, df = 3, p-value = 1.341e-14
```

Our graphs show fairly even spread. But a quick Breusch-Pagan test shows that our assumption of constancy of variance error is reasonable.

Durbin-Watson test

```
data: boxcox.lmfit
DW = 1.3658, p-value = 2.635e-08
alternative hypothesis: true autocorrelation is not 0
```

Even spread of residuals against an index and a high p-value in a Durbin-Watson test shows that our assumption of independence is reasonable.



```

      .
      .
      .
      shapiro-wilk normality test

data:  boxcox.res
W = 0.97508, p-value = 5.551e-05
      .

```

Our assumption of normality, which was violated in our previous model, is satisfied here. The p-value from the Shapiro-Wilk test is now much higher, and our Q-Q plot shows that the assumption of normality is reasonable.

3. RESULTS:

The summary of the final optimal model is:

$$Y = 5.043e+00 + 7.833e-04X_2 + 2.766e-02X_3 + 2.058e-01X_5$$

The predictor variables of the model are DRAIN_SQKM, PPTAVG_BASIN, T_AVG_SITE.

The value of the **F-statistic** is 49.94 on six predictor variables, and the **p-value** is 2.2e-16; Therefore, there is an overall significant relationship between the response variable and the predictor variables.

```

Call:
lm(formula = trans.Y ~ X2 + X3 + X5, data = streamflow)

Residuals:
    Min       1Q   Median       3Q      Max
-12.0998  -2.2829  -0.1921   2.1780   8.9177

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  5.043e+00  6.227e-01   8.098 1.58e-14 ***
X2           7.833e-04  8.466e-05   9.253 < 2e-16 ***
X3           2.766e-02  3.438e-03   8.046 2.24e-14 ***
X5           2.058e-01  4.236e-02   4.860 1.93e-06 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.085 on 289 degrees of freedom
Multiple R-squared:  0.3414,    Adjusted R-squared:  0.3346
F-statistic: 49.94 on 3 and 289 DF,  p-value: < 2.2e-16

> anova(boxcox.lmfit)
Analysis of Variance Table

Response: trans.Y
      Df Sum Sq Mean Sq F value    Pr(>F)
X2      1  501.01   501.01  52.655 3.678e-12 ***
X3      1  699.79   699.79  73.547 6.074e-16 ***
X5      1  224.71   224.71  23.616 1.932e-06 ***
Residuals 289 2749.81    9.51
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

From the summary statistics and ANOVA table, the results of each p-value are less than the significant level 0.05.

4. CONCLUSION:

To find the optimal model for the STREAMFLOW dataset, we performed exploratory data analysis and reduced the model based on the significant level of 0.05. To validate it, we chose the method for model selection and applied diagnostic measures to ensure the model is fitted perfectly. Later, to improve it further, we transformed the model, and there is a significant improvement in the Adjusted R-squared of the final model. We chose the transformed model, which has the better improvement.

5. APPENDIX:

#Data Summarization

```
library(readr)
```

```
streamflow <- read_csv("streamflow.csv")
```

```
View(streamflow)
```

```
colnames(streamflow) <- c("X1","Y","X2","X3","X4","X5","X6","X7","X8")
```

```
summary(streamflow)
```

HISTOGRAMS

```
par(mfrow=c(3,3))
```

```
hist(streamflow$X1,main="Histogram of X1",xlab="Staid")
```

```
hist(streamflow$Y,main="Histogram of Y",xlab="90th percentile of the time series")
```

```
hist(streamflow$X2,main="Histogram of X2",xlab="the drainage area ")
```

```
hist(streamflow$X3,main="Histogram of X3",xlab="average basin precipitation ")
```

```
hist(streamflow$X4,main="Histogram of X4",xlab="average basin temperature ")
```

```
hist(streamflow$X5,main="Histogram of X5",xlab="average temperature at the stream location ")
```

```
hist(streamflow$X6,main="Histogram of X6",xlab="average relative humidity across the basin")
```

```
hist(streamflow$X7,main="Histogram of X7",xlab="average March precipitation")
```

```
hist(streamflow$X8,main="Histogram of X8",xlab="median relief ratio")
```

BOX PLOTS

```
boxplot(streamflow$Y, xlab="Max90", main="Max90")
```

```
boxplot(streamflow$X1, xlab="STAID", main="STAID")
```

```
boxplot(streamflow$X2, xlab="DRAIN_SQKM", main=" The drainage area")
```



```
boxplot(streamflow$X3, xlab="PPTAVG_BASIN", main="The average
basinprecipitation")
```

```
boxplot(streamflow$X4, xlab="T_AVG_BASIN", main="Boxplot of the average
basintemperature")
```

```
boxplot(streamflow$X5, xlab="T_AVG_SITE", main="Boxplot of the
averagetemperature at the stream location")
```

```
boxplot(streamflow$X6, xlab="RH_BASIN", main="Boxplot of the average
relativehumidity across the basin ")
```

```
boxplot(streamflow$X7, xlab="MAR_PPT7100_CM", main="Boxplot of the
averageMarch precipitation ")
```

```
boxplot(streamflow$X8, xlab="RRMEDIAN", main="Boxplot of the median
reliefratio")
```

SCATTER PLOTS

```
plot(streamflow$Y, xlab="Max90",ylab="Max90", main="Boxplot of Stream Flow")
```

```
plot(streamflow$X1, xlab="STAID",ylab="STAID", main="STAID")
```

```
plot(streamflow$X2, xlab="DRAIN_SQKM", ylab="DRAIN_SQKM", main=" The
drainage area")
```

```
plot(streamflow$X3, xlab="PPTAVG_BASIN",ylab="PPTAVG_BASIN", main="The
average basinprecipitation")
```

```
plot(streamflow$X4, xlab="T_AVG_BASIN",ylab="T_AVG_BASIN", main="Boxplot
of the average basintemperature")
```

```
plot(streamflow$X5, xlab="T_AVG_SITE", ylab="T_AVG_SITE", main="Boxplot of
the averagetemperature at the stream location")
```

```
plot(streamflow$X6, xlab="RH_BASIN",ylab="RH_BASIN", main="Boxplot of the
average relativehumidity across the basin ")
```

```
plot(streamflow$X7, xlab="MAR_PPT7100_CM",ylab="MAR_PPT7100_CM",
main="Boxplot of the averageMarch precipitation ")
```

```
plot(streamflow$X8, xlab="RRMEDIAN",ylab="RRMEDIAN", main="Boxplot of the
median reliefratio")
```

ADDED VARIABLE PLOTS

```
library(car)
```

```
avPlots(fitstream)
```

CORRELATION MATRIX

```
cor(streamflow)
```

```

pairs(streamflow)
##checking for multicollinearity.
eigen(cor(streamflow))$values

## MODELS AND METHODS

fitstream<-lm(Y ~X1+X2+X3+X4+X5+X6+X7+X8,data=streamflow)
fitstream
summary(fitstream)

##ANOVA t-test
anova(fitstream)

library(MASS)
## Model Selection
library(olsrr)

##### Print all possible regression models in terms of adjr, Cp, AIC, and BIC.

par(mfrow=c(1,1))
b <- ols_step_all_possible(fitstream)
plot(b)
##### Adjusted R2 #####

b.adj_r = data.frame(n=b$n,predictors=b$predictors,adj_r=b$adj_r)
print(b.adj_r)
print(b.adj_r[c(93,163,219,247,255),])

##### Cp #####

b.cp = data.frame(n=b$n,predictors=b$predictors,cp=b$cp)
print(b.cp)

```

```
print(b.cp[c(93,163,219,247,255),])
```

```
#### AIC ####
```

```
b.aic = data.frame(n=b$n,predictors=b$predictors,aic=b$aic)
```

```
print(b.aic)
```

```
print(b.aic[c(93,163,219,247,255),])
```

```
#### BIC ####
```

```
b.bic = data.frame(n=b$n,predictors=b$predictors,bic=b$bic)
```

```
print(b.bic)
```

```
print(b.bic[c(93,163,219,247,255),])
```

```
#### PRESS ####
```

```
b.press = data.frame(n=b$n,predictors=b$predictors,press=b$msep)
```

```
print(b.press)
```

```
print(b.press[c(93,163,219,247,255),])
```

```
#### Stepwise Regression ####
```

```
k <- ols_step_both_p(fitstream,pent=0.10,prem=0.1,details=TRUE)
```

```
plot(k)
```

```
#### Final Model? ####
```

```
reduced.lmfit <- lm(Y ~ X2 + X3+X5, data=streamflow)
```

```
summary(reduced.lmfit)
```

```
##### Regression Diagnostics #####
```

```
res <- rstudent(reduced.lmfit)
fitted.y <- fitted(reduced.lmfit)
```

```
##### Residual Plots #####
```

```
par(mfrow=c(2,2))
```

```
plot(res ~ streamflow$X2, xlab="X2", ylab="Residual", main="Residuals vs. X2")
abline(h=0)
```

```
plot(res ~ streamflow$X3, xlab="X3", ylab="Residual", main="Residuals vs. X3")
abline(h=0)
```

```
plot(res ~ streamflow$X5, xlab="X5", ylab="Residual", main="Residuals vs. X5")
abline(h=0)
```

```
plot(res ~ fitted.y, xlab="Fitted value", ylab="Residual", main="Residuals vs. Fitted
Values")
abline(h=0)
```

```
##### Multicollinearity #####
```

```
vif(reduced.lmfit)
```

```
##### Constancy of Error Variances #####
```

```
library(lmtest)
bptest(reduced.lmfit)
```

```
#Durbin-Watson
```

```
#install lmtest
```

```
library(lmtest)
```

```
dwtest(fitstream, alternative="two.sided")
```

Normality

qqnorm(res);qqline(res)

#####Shapiro test#####

shapiro.test(res)

#DFFITS values

library(olsrr)

ols_plot_dffits(reduced.lmfit)

#DFBETAS values

ols_plot_dfbetas(reduced.lmfit)

#Cook's distance values

ols_plot_cooksd_chart(reduced.lmfit)

Transformation

library(EnvStats)

boxcox.summary <- boxcox(reduced.lmfit, optimize=TRUE)

lambda <- boxcox.summary\$lambda

lambda

trans.Y <- streamflow\$Y^lambda

streamflow <- cbind(streamflow,trans.Y)

streamflow

Re-fitting a model using the transformed response variable.

boxcox.lmfit <- lm(trans.Y ~ X2 + X3 + X5, data=streamflow)

```
summary(boxcox.lmfit)
```

```
anova(boxcox.lmfit)
```

```
boxcox.res <- rstudent(boxcox.lmfit)
```

```
boxcox.fitted.y <- fitted(boxcox.lmfit)
```

```
##### Residual Plots #####
```

```
par(mfrow=c(2,2))
```

```
plot(boxcox.res ~ streamflow$X2, xlab="X2", ylab="Residual", main="Residuals vs.  
X2")
```

```
abline(h=0)
```

```
plot(boxcox.res ~ streamflow$X3, xlab="X3", ylab="Residual", main="Residuals vs.  
X3")
```

```
abline(h=0)
```

```
plot(boxcox.res ~ streamflow$X5, xlab="X5", ylab="Residual", main="Residuals vs.  
X5")
```

```
abline(h=0)
```

```
plot(boxcox.res ~ fitted.y, xlab="Fitted value", ylab="Residual", main="Residuals vs.  
Fitted Values")
```

```
abline(h=0)
```

```
##### Multicollinearity #####
```

```
library(HH)
```

```
vif(boxcox.lmfit)
```

```
##### Constancy of Error Variances #####
```

```
bptest(boxcox.lmfit)
```

```
dwtest(boxcox.lmfit, alternative="two.sided")
```

Normality

```
qqnorm(boxcox.res);qqline(boxcox.res)
shapiro.test(boxcox.res)
```

Final Model

```
final.lmfit <- boxcox.lmfit
summary(final.lmfit)
```

```
##Obtain DFFITS, DFBETAS, and Cook's distance values
library(olsrr)
```

```
#DFFITS values
ols_plot_dffits(final.lmfit)
```

```
#DFBETAS values
ols_plot_dfbetas(final.lmfit)
```

```
#Cook's distance values
ols_plot_cooksd_chart(final.lmfit)
```

Fit a regression model with interaction terms

```
streamflow.lmfit <- lm(Y ~ X2 + X3 + X5 + X2*X3 + X2*X5 + X3*X5, data=streamflow)
summary(streamflow.lmfit)
anova(streamflow.lmfit)
```

Fit a regression model with no interaction terms

```
streamflow.reduced <- lm(Y ~ X2 + X3 + X5, data=streamflow)
```

Test for significance of the interaction terms

anova(streamflow.reduced,streamflow.lmfit)