Ozone_Assignment.R

SAMSON

2023-02-14

```
#### Assignment
getwd()
```

```
## [1] "C:/Users/SAMSON"
```

```
setwd("C:/Users/SAMSON")
ozone <- read.table("ozone (1).txt", header=TR
UE)
head(ozone)</pre>
```

```
## rad temp wind ozone
## 1 190 67 7.4 41
## 2 118 72 8.0 36
## 3 149 74 12.6 12
## 4 313 62 11.5 18
## 5 299 65 8.6 23
## 6 99 59 13.8 19
```

```
Y = ozone$ozone
X1 = ozone$rad
X2 = ozone$temp
X3 = ozone$wind
model.identity = glm(Y \sim X1 + X2 + X3), family
= gaussian(link = "identity"))
model.inverse = glm(Y \sim X1 + X2 + X3, family =
gaussian(link = "inverse"))
model.log = glm(Y \sim X1 + X2 + X3, family = ga
ussian(link = "log"))
model.exponential = glm(Y \sim log(X1) + log(X2))
+ log(X3), family = gaussian(link = "log"))
#1a
summary(model.identity)$aic
```

[1] 998.6276

summary(model.inverse)\$aic

[1] 999.0104

summary(model.log)\$aic # iii Best Model AIC =
972.5169

```
## [1] 972.5169
```

summary(model.exponential)\$aic

[1] 974.9654

```
# 1b
var.yi = summary(model.log)$dispersion
mu = predict(model.log,type="response")
pearson = (Y - mu)/sqrt(var.yi)
model.pearson.normal = glm(pearson^2 ~ mu,family = gaussian(link ="identity"))
model.pearson.gamma = glm(pearson^2 ~ mu,family = Gamma(link ="identity"))
model.pearson.inv = glm(pearson^2 ~ mu,family = inverse.gaussian(link ="identity"))
summary(model.pearson.normal)#i insignificant
at 1%
```

```
##
## Call:
## glm(formula = pearson^2 ~ mu, family = gaus
sian(link = "identity"))
##
## Deviance Residuals:
                1Q Median
      Min
                                  30
##
                                          Max
## -2.4306 -0.7060 -0.2830 0.0626 19.4220
##
## Coefficients:
               Estimate Std. Error t value Pr
##
(>|t|)
## (Intercept) -0.147945 0.405849 -0.365
0.71617
               0.026197 0.008072 3.246
## mu
0.00156 **
## ---
## Signif. codes: 0 '***' 0.001 '**'
                                     0.01
0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family t
aken to be 5.255567)
##
      Null deviance: 628.22 on 110 degrees
##
of freedom
## Residual deviance: 572.86 on 109
                                     degrees
```

```
of freedom

## AIC: 503.17

##

## Number of Fisher Scoring iterations: 2
```

summary(model.pearson.gamma)

```
##
## Call:
## glm(formula = pearson^2 ~ mu, family = Gamm
a(link = "identity"))
##
## Deviance Residuals:
       Min
                 10 Median
                                          Max
                                   30
##
## -3.8385 -1.5794 -0.6042 0.1484
                                       4.0541
##
## Coefficients:
                Estimate Std. Error t value Pr
##
(>|t|)
## (Intercept) -0.092842 0.046616 -1.992
0.0489 *
## mu
                0.023788 0.004176 5.697 1.
05e-07 ***
## ---
## Signif. codes: 0 '***' 0.001 '**'
                                     0.01
0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for Gamma family take
n to be 2.63651)
##
   Null deviance: 328.61 on 110 degrees
##
of freedom
## Residual deviance: 256.46 on 109
                                     degrees
```

```
of freedom

## AIC: 115.83

##

## Number of Fisher Scoring iterations: 7
```

summary(model.pearson.inv)

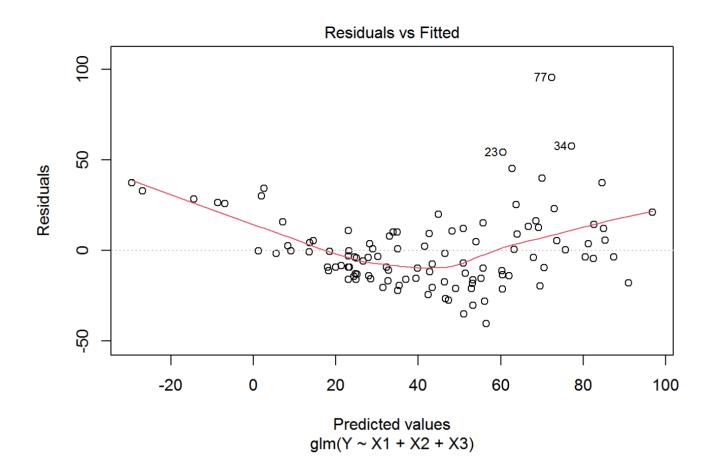
```
##
## Call:
## glm(formula = pearson^2 ~ mu, family = inve
rse.gaussian(link = "identity"))
##
## Deviance Residuals:
        Min
                         Median
                                       30
                   10
##
Max
## -136.619 -2.754
                         -0.782 0.177
4.780
##
## Coefficients:
                Estimate Std. Error t value Pr
##
(>|t|)
## (Intercept) -0.103365 0.021468 -4.815 4.
78e-06 ***
                0.025080 0.003892 6.445 3.
## mu
24e-09 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01
0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for inverse.gaussian
family taken to be 3.935746)
##
       Null deviance: 36009 on 110 degrees o
##
```

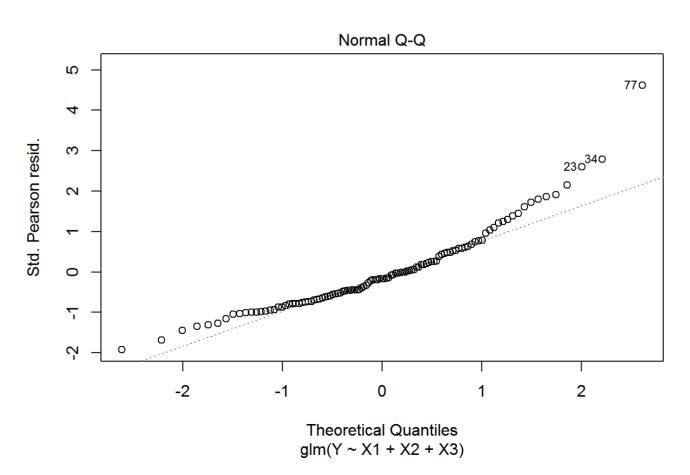
```
f freedom
## Residual deviance: 35867 on 109 degrees o
f freedom
## AIC: 457.23
##
## Number of Fisher Scoring iterations: 6
```

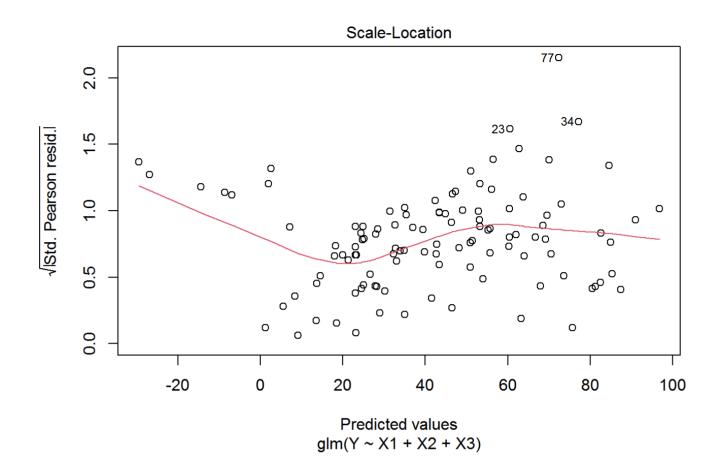
```
# 1c
betahat = summary(model.log)$coef[1:4]
betahat = matrix(betahat)
K = matrix(c(0,0,1,1),nrow = 4,ncol = 1)
W = (t(t(K)%*%betahat))*solve(t(K)%*%vcov(model.log)%*%K)*(t(K)%*%betahat)
W # 10.66784
```

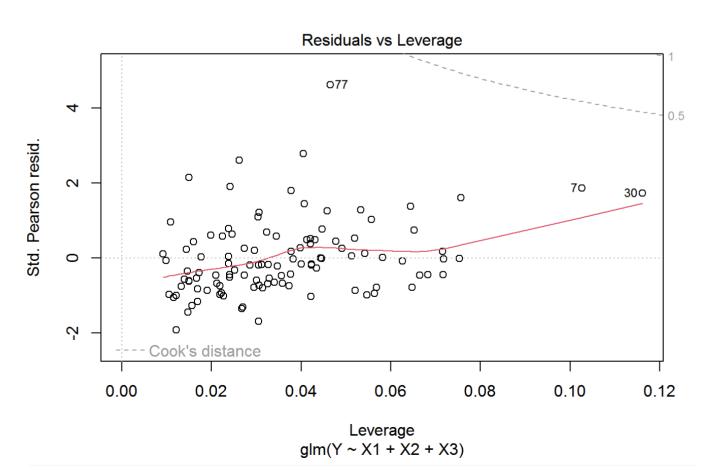
```
## [,1]
## [1,] 10.66784
```

```
plot(model.identity)
```

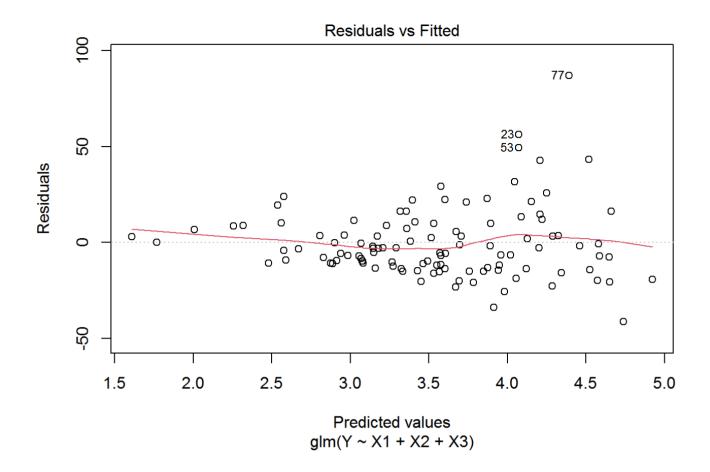


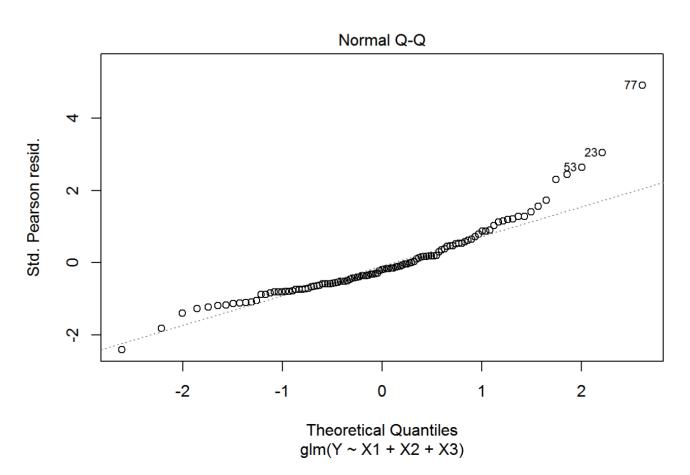


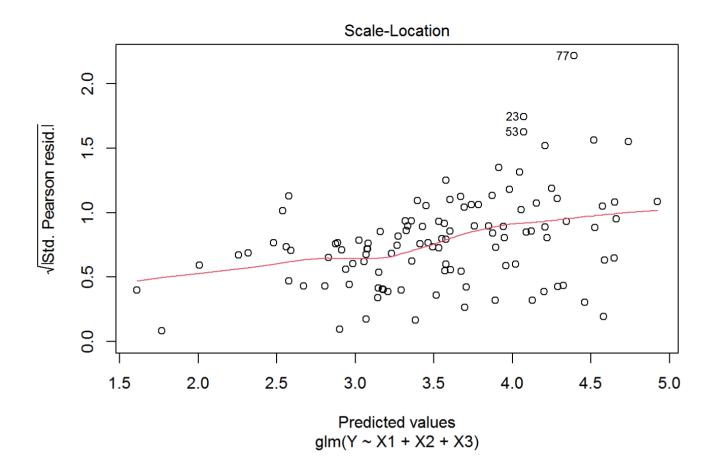


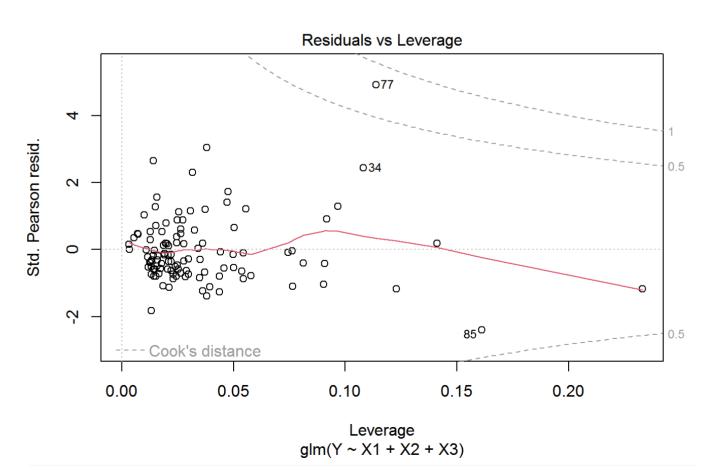


plot(model.log)

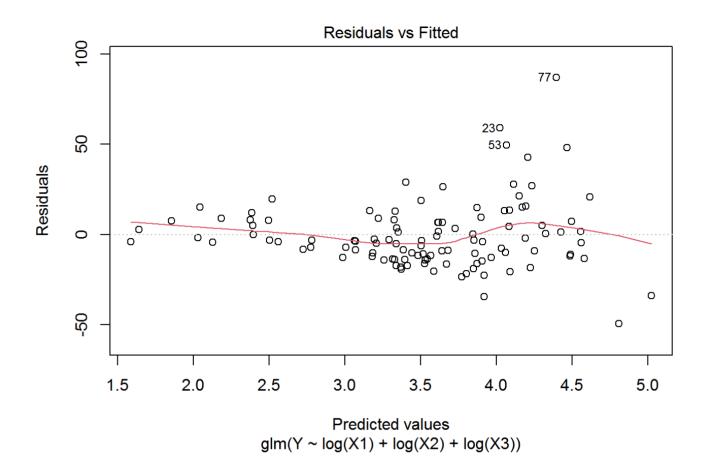


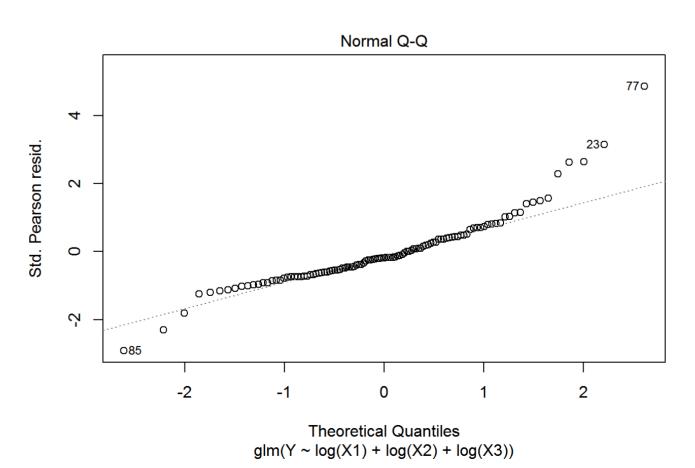


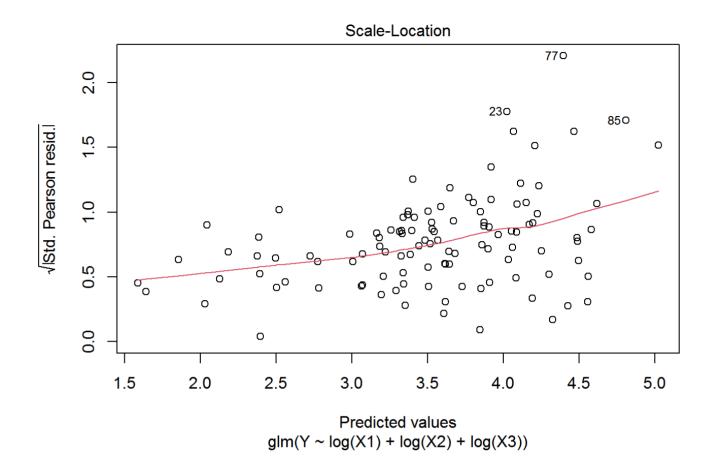


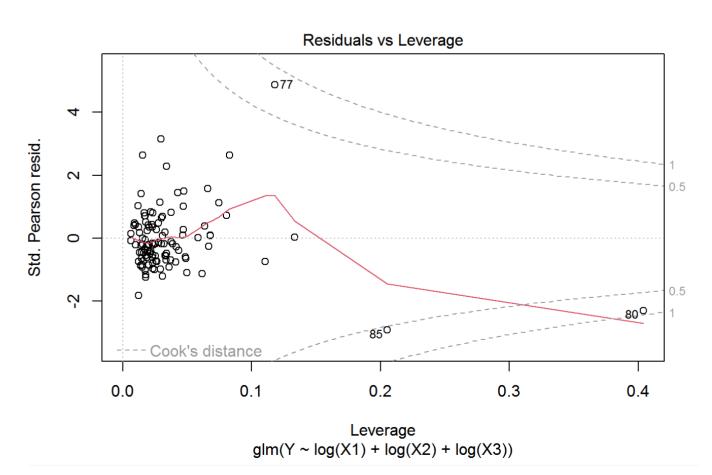


plot(model.exponential)









plot(model.inverse)

