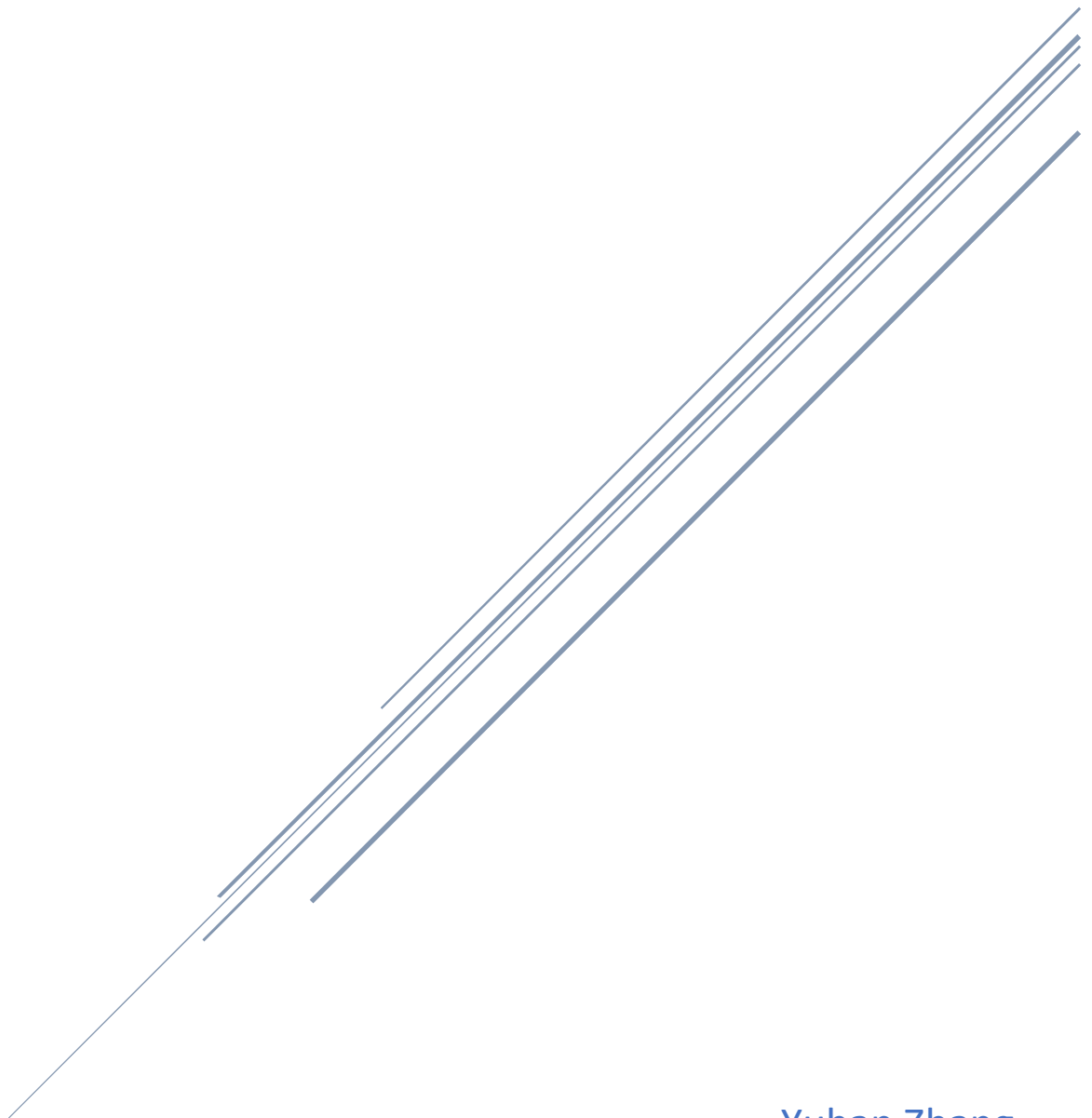


# STAT 350

## Assignment4



Yuhan Zhang  
301345627

Assignment 4.

$$1. D_i = \frac{(\hat{\beta}_i - \bar{\beta})'(X'X)(\hat{\beta}_i - \bar{\beta})}{p \cdot MS_{Res}}$$

$$[\hat{\beta}_i - \bar{\beta}] = \frac{(X'X)^{-1} X_i e_i}{1 - h_{ii}} \text{ from (81)}$$

$$= \frac{[(X'X)^{-1} X_i e_i]' (X'X) [(X'X)^{-1} X_i e_i]}{(1 - h_{ii})^2 p \cdot MS_{Res}}$$

$$= \frac{e_i' X_i' (X'X)^{-1} (X'X) (X'X)^{-1} X_i e_i}{(1 - h_{ii})^2 p \cdot MS_{Res}}$$

$$= \frac{e_i' X_i' (X'X)^{-1} X_i e_i}{(1 - h_{ii})^2 p \cdot MS_{Res}}$$

$$= \frac{X_i' (X'X)^{-1} e_i^2 X_i}{(1 - h_{ii})^2 p \cdot MS_{Res}}$$

$$= \frac{h_{ii} e_i^2}{(1 - h_{ii})^2 p \cdot MS_{Res}}$$

$$= \frac{e_i^2}{(1 - h_{ii}) MS_{Res}} \cdot \frac{h_{ii}}{(1 - h_{ii}) \cdot p} \quad \left[ r_i = \frac{e_i}{\sqrt{MS_{Res} (1 - h_{ii})}} \right]$$

$$= \frac{r_i^2 \cdot h_{ii}}{p \cdot (1 - h_{ii})}$$

Q2 a

Coefficients:

(Intercept)	z1	z2
-5.959e-17	9.137e-01	3.068e-01

Q2 b

```
##b.
```{r}
round(vcov(fit1),5)
```

	(Intercept)	z1	z2
(Intercept)	0.00132	0.00000	0.00000
z1	0.00000	0.01185	0.00000
z2	0.00000	0.00000	0.01185

As the covariance are all zero, They are independent.

Q2 c

```
##c
```{r}
vif(fit1)|
```

z1	z2
1	1

Q2 d

Coefficients:

(Intercept)	z1	z2
-5.508e-17	8.941e-01	-1.155e-02

Q2 e

```
## e
```{r}
round(vcov(fit2),5)
```

	(Intercept)	z1	z2
(Intercept)	0.00395	0.00000	0.00000
z1	0.00000	0.05908	-0.03727
z2	0.00000	-0.03727	0.05908

They are not independent because  $\text{var}(\text{Beta1}(\text{hat}), \text{Beta2}(\text{hat}))$  is not equal to 0.

Q2 f

```
##f
```{r}
vif(fit2)
```

      z1      z2
1.660661 1.660661
```

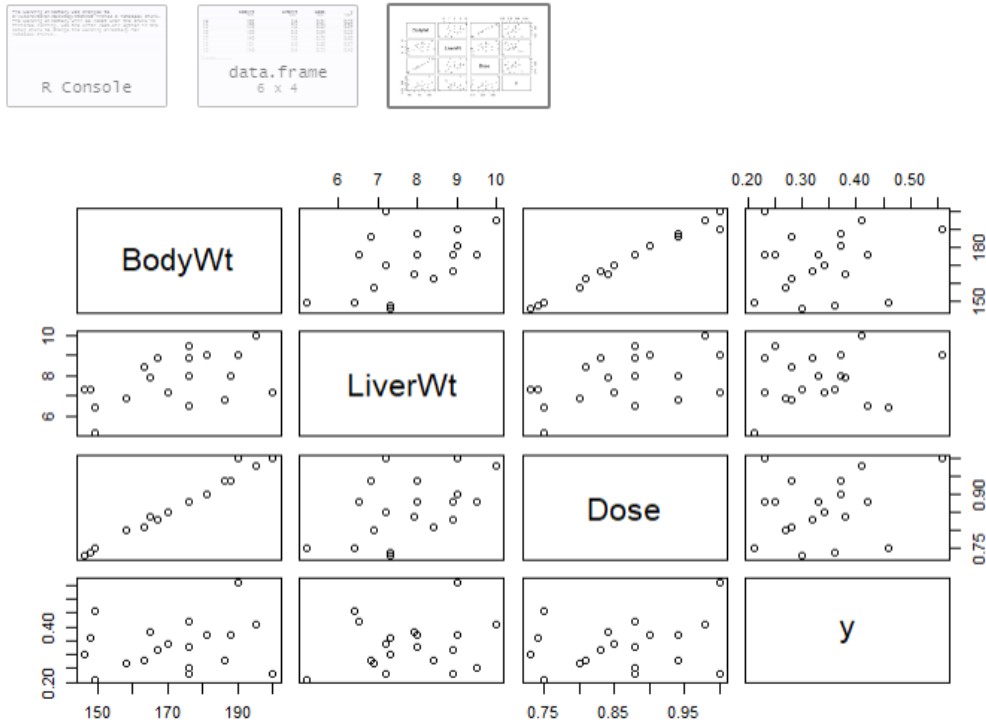
Q2 g

## g.  
The first is preferable as it does not have multicollinearity.  
Besides, the variance the  $\beta(\hat{\beta})$  is smaller.

Q4 a

```
## a.
```

```
##{r}  
setwd("C:/Users/carol/Desktop/stat350")  
df3 = read.csv("guinea_pig.csv")[, -1]  
tail(df3)  
pairs(df3)
```



```
##i
```

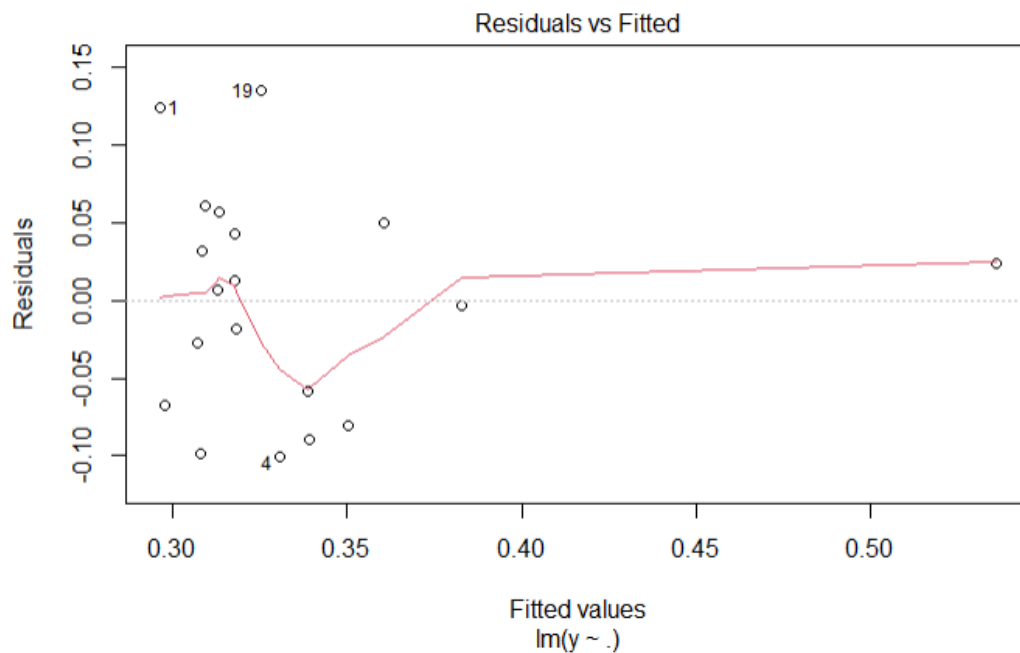
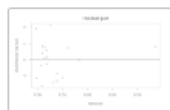
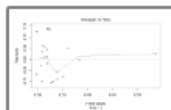
There is a positive linear relation between body weight and dose. There is a weak positive linear relationship between body weight and liver weight, and a weak positive linear relationship between liverweight and dose.

```
##ii
```

There is no obviould relationship between predictors and response variables.

Q4 bc

```
## b.  
body weight and dose  
`{r}`  
fit3 = lm(y~., data=df3)  
summary(fit3)  
  
##body and dose, there is a significant linear relationship.  
  
## c.  
plot(fit3, 1)  
plot(fitted(fit3), rstudent(fit3), main="residual plot", ylab="studentized residual")  
abline(h=0)  
`{r}`
```



```
##c  
There is no linear relationship.
```

#### Q4 d

```
## d.
```

```
```{r}  
nrow(df3)  
x<-cbind(rep(1,nrow(df3)), df3[,1],df3[,2],df3[,3])  
  
hii<-diag(X%%solve(t(X)%%X)%%t(X))  
hii  
# Identify points of high Leverage  
p<-ncol(X) # number of betas in the model (beta0,beta1,beta2)  
n<-nrow(X) # number of observations  
which(hii>2*p/n) #points with high leverage  
```
```

```
[1] 19  
[1] 0.17798270 0.17934099 0.85091457 0.10761585 0.39153825 0.16115958 0.13688107 0.25367448  
[9] 0.06701578 0.11968672 0.11950583 0.17239599 0.31618336 0.13140699 0.07617481 0.21661460  
[17] 0.19522441 0.14872221 0.17796183  
[1] 3
```

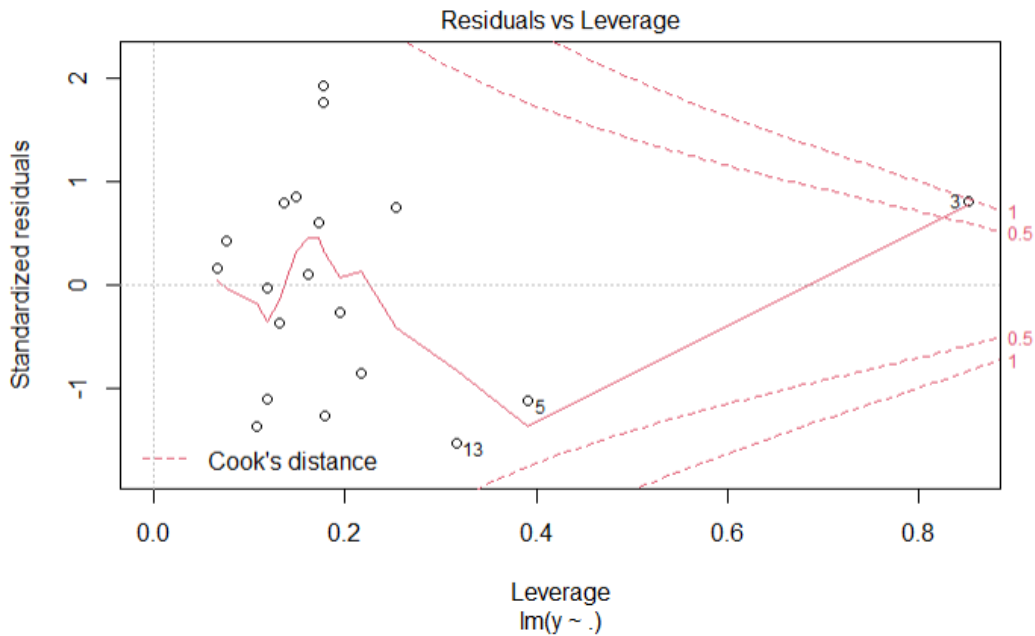
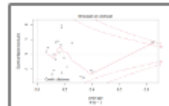
yes.The third one has large hii.

Q4 e

```
## e.  
{r}  
influence.measures(fit3)  
plot(fit3, 5)
```

R Console

data.frame  
19 x 9



There is no cook's distance larger than one.

The 3rd obs has large leverage and cook's distance, it has relative large impact on the regression coef.



#### Q4 f

```
## f.
```

```
```{r}
vif(fit3)
```
```

```
      Bodywt      Liverwt      Dose
52.101917    1.335679    51.427154
```

Yes there is evidence of collinearity, as the vif of bodywt and dose is 52.1 and 51.43 which are larger than 10.

#### Q4 g

```
## g.
```

```
```{r}
fit4 = lm(y~., data=df3[-3,])
summary(fit4)
vif(fit4)
```
```

```
Call:
lm(formula = y ~ ., data = df3[-3, ])

Residuals:
    Min       1Q   Median       3Q      Max
-0.102154 -0.056486  0.002838  0.046519  0.137059

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  0.311427   0.205094   1.518   0.151
Bodywt      -0.007783   0.018717  -0.416   0.684
Liverwt      0.008989   0.018659   0.482   0.637
Dose         1.484877   3.713064   0.400   0.695

Residual standard error: 0.07825 on 14 degrees of freedom
Multiple R-squared:  0.02106, Adjusted R-squared:  -0.1887
F-statistic: 0.1004 on 3 and 14 DF, p-value: 0.9585

      Bodywt      Liverwt      Dose
259.449422    1.445674    253.199751
```

As all the p-value is larger than 0.05, no independent variable have significant linear relationship with the response variable, which is as a result of the multicollinearity increase the standard error of the beata hats. Therefore reduce the t-test and increase the p-value leading to non-significant result.