Regression Diagnostics

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General Preparations

```
Go to the specified directory and do
data=read.table("CH09TA01.txt", header=T)
Attach data attach(data)
Fit the full model
mod = Im(y \sim x1 + x2 + x3 + x4 + x5 + x6 + x7 + x8)
Find the residuals:
e = resid(mod) OR e = y- predict(mod)
Calculate the X- Matrix
X = matrix(c(rep(1, nrow(data)), x1, x2, x3, x4, x5, x6, x7, x8),
nrow=nrow(data), ncol= 9)
Create the Hat-matrix: H = X\%^*\%solve(t(X)\%^*\%X)\%^*\%t(X)
```

Identifying Outlying Y variables

Semi-studentized Residuals:

$$e_i^* = rac{e_i}{\sqrt{(\mathit{MSE})}}$$

Studentized Residuals:

$$r_i = rac{e_i}{\sqrt{(MSE(1-h_{ii}))}}$$

Identifying Outlying Y variables

Semi-studentized Residuals:

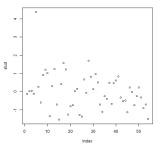
$$e_i^* = rac{e_i}{\sqrt{(\mathit{MSE})}}$$

Studentized Residuals:

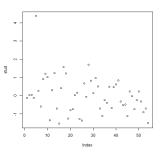
$$r_i = rac{e_i}{\sqrt{(extit{MSE}(1-h_{ii}))}}$$

```
MSE=tail( anova(mod)[, 3], 1)
h=diag(H)
semistud<-NULL
stud<- NULL
semistud[1:nrow(data)]= 0
stud[1:nrow(data)]=0
for (i in (1:nrow(data))){
semistud[i] = e[i]/(sqrt(MSE))
stud[i] = e[i]/(sqrt(MSE*(1-h[i])))}
```

png('stud.png') plot(stud) dev.off()

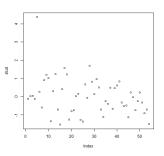


png('stud.png') plot(stud) dev.off()



-0.11 0.02 0.05 -0.13 4.38 0.26 -0.61 0.90 1.21 1.01 -1.36 0.30 1.24 -0.72 -1.54 0.41 1.58 1.23 -1.26 -0.80 -0.75 0.03 0.15 -1.29 -1.37 0.67 -0.06 1.69 0.82 0.13 0.96 0.51 -0.72 -1.12 -0.25 -0.41 0.49 -0.67 0.46 0.61 0.83 -0.33 -0.54 -0.49 -1.14 0.22 -0.04 -0.74 -0.26 0.23 -0.33 -0.92 -0.71 -1.50

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Rule of thumb:

Greater than 3*sd(stud)= 3.065305 OR

Bonferroni test-procedure: $t_{(1-\frac{\alpha}{2\alpha};n-p)} = \sim 3.43$



Identifying Outlying Y variables II

Deleted Residuals:

$$d_i = \frac{e_i}{1-h_{ii}}$$

Studentized deleted Residuals:

$$t_i = e_i \left\{ \frac{n-p-1}{(SSE(1-h_{ii})-e_i^2)} \right\}^{1/2}$$

Identifying Outlying Y variables II

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Studentized deleted Residuals:

$$t_i = e_i \left\{ \frac{n-p-1}{(SSE(1-h_{ii})-e_i^2)} \right\}^{1/2}$$

```
SSE=tail( anova(mod)[, 2], 1)

del<-NULL

studdel<- NULL

del[1:nrow(data)]= 0

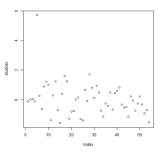
studdel[1:nrow(data)]=0

for (i in (1:nrow(data))){

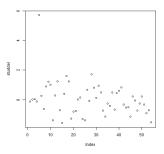
del[i] = e[i]/(1-h[i])

studdel[i] = e[i]*sqrt((nrow(data)-9-1)/(SSE*(1-h[i])-e[i]*e[i]))}
```

png('studdel.png') plot(studdel) dev.off()



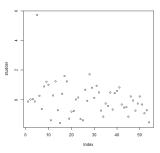
png('studdel.png') plot(studdel) dev.off()



round(studdel,2)

-0.11 0.02 0.05 -0.13 5.71 0.26 -0.60 0.90 1.21 1.01 -1.38 0.30 1.25 -0.72 -1.57 0.40 1.60 1.23 -1.27 -0.80 -0.75 0.03 0.15 -1.29 -1.38 0.67 -0.06 1.73 0.82 0.13 0.96 0.50 -0.72 -1.13 -0.24 -0.40 0.49 -0.66 0.45 0.61 0.83 -0.33 -0.53 -0.49 -1.15 0.22 -0.04 -0.74 -0.26 0.23 -0.32 -0.92 -0.70 -1.52

png('studdel.png') plot(studdel) dev.off()



round(studdel,2)

-0.11 0.02 0.05 -0.13 5.71 0.26 -0.60 0.90 1.21 1.01 -1.38 0.30 1.25 -0.72 -1.57 0.40 1.60 1.23 -1.27 -0.80 -0.75 0.03 0.15 -1.29 -1.38 0.67 -0.06 1.73 0.82 0.13 0.96 0.50 -0.72 -1.13 -0.24 -0.40 0.49 -0.66 0.45 0.61 0.83 -0.33 -0.53 -0.49 -1.15 0.22 -0.04 -0.74 -0.26 0.23 -0.32 -0.92 -0.70 -1.52

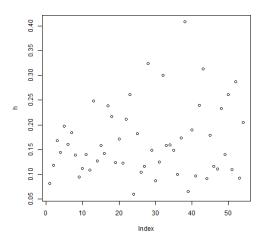
Rule of thumb:

Greater than 3*sd(studdel)= 3.428044 OR

Bonferroni test-procedure: $t_{(1-\frac{\alpha}{2n};n-p-1)}=\sim 3.42$

Identifying outlying *X* observations

Leverage values h_{ii} png('leverage.png') plot(h) dev.off()



Identifying outlying *X* observations

round(h,3)
0.082 0.119 0.168 0.144 0.197 0.160 0.184 0.139 0.095 0.112
0.141 0.109 0.248 0.127 0.159 0.142 0.238 0.217 0.124 0.172
0.123 0.212 0.261 0.060 0.182 0.105 0.116 0.323 0.148 0.088
0.125 0.300 0.159 0.160 0.149 0.100 0.174 0.408 0.066 0.190
0.097 0.239 0.313 0.091 0.179 0.116 0.111 0.233 0.140 0.261
0.110 0.287 0.093 0.205

Identifying outlying *X* observations

round(h,3)

0.082 0.119 0.168 0.144 0.197 0.160 0.184 0.139 0.095 0.112 0.141 0.109 0.248 0.127 0.159 0.142 0.238 0.217 0.124 0.172 0.123 0.212 0.261 0.060 0.182 0.105 0.116 0.323 0.148 0.088 0.125 0.300 0.159 0.160 0.149 0.100 0.174 0.408 0.066 0.190 0.097 0.239 0.313 0.091 0.179 0.116 0.111 0.233 0.140 0.261 0.110 0.287 0.093 0.205

Rule of thumb:

Greater than $\frac{2p}{n} = 0.337$

Point 28 and 38 needs to be seen further

Identifying Influential observations: DFFits

DFFits:

```
(\textit{DFFits})_i = rac{\hat{Y}_i - \hat{Y}_{i(i)}}{\sqrt{(\textit{MSE}_{(i)}h_{ii})}} = t_i ig\{rac{h_{ii}}{(1-h_{ii})}ig\}^{1/2}
```

```
dffits<-NULL
dffits[1:nrow(data)]=0
for (i in (1:nrow(data))){
  dffits[i] = studdel[i]*sqrt(h[i]/(1-h[i]))
}</pre>
```

Identifying Influential observations: DFFits

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```
dffits<-NULL
dffits[1:nrow(data)]=0
for (i in (1:nrow(data))){
  dffits[i] = studdel[i]*sqrt(h[i]/(1-h[i]))
}</pre>
```

Rule of thumb:

Influential if the value exceeds 1 (for small-medium datasets $(n \le 60)$ or if it exceeds $2\sqrt{\frac{p}{n}}$ (for large datasets)

Identifying Influential observations: DFBetas

DFBetas:

$$(DFBetas)_{k(i)} = \frac{b_k - b_{k(i)}}{\sqrt{(MSE_{(i)}c_{kk})}}$$

where c_{kk} is the kth diagonal of $(X^TX)^{-1}$.

```
 \begin{array}{l} c = diag(solve(t(X)\%^*\%X)) \\ mse = 0 \\ dfbetas1 < -NULL \\ dfbetas1[1:54] = 0 \\ for \ (i \ in \ (1:nrow(data))) \{ \\ fit < -lm(y[-i] \sim x1[-i] + x2[-i] + x3[-i] + x4[-i] + x5[-i] + x6[-i] + x7[-i] + x8[-i]) \\ mse < -tail(\ anova(fit)[,\ 3],\ 1) \\ dfbetas1[i] = (mod$coeff[2]-fit$coeff[2])/sqrt(mse* c[2]) \} \\ \end{array}
```

Identifying Influential observations: DFBetas

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$$(DFBetas)_{k(i)} = \frac{b_k - b_{k(i)}}{\sqrt{(MSE_{(i)}c_{kk})}}$$

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```
 \begin{array}{l} c = diag(solve(t(X)\%^*\%X)) \\ mse = 0 \\ dfbetas1 < -NULL \\ dfbetas1[1:54] = 0 \\ for (i in (1:nrow(data))) \{ \\ fit < -lm(y[-i] \sim x1[-i] + x2[-i] + x3[-i] + x4[-i] + x5[-i] + x6[-i] + x7[-i] + x8[-i]) \\ mse < -tail( anova(fit)[, 3], 1) \\ dfbetas1[i] = (mod$coeff[2]-fit$coeff[2])/sqrt(mse* c[2]) \} \\ ifelse(dfbetas(mod)[,] > 1,1,0) \\ \end{array}
```

Identifying Influential observations: DFBetas

DFBetas:

$$(DFBetas)_{k(i)} = \frac{b_k - b_{k(i)}}{\sqrt{(MSE_{(i)}c_{kk})}}$$

where c_{kk} is the kth diagonal of $(X^TX)^{-1}$.

```
c= diag(solve(t(X)%*%X))
mse=0
dfbetas1<-NULL
dfbetas1[1:54]=0
for (i in (1:nrow(data))){
fit < -lm(y[-i] \sim x1[-i] + x2[-i] + x3[-i] + x4[-i] + x5[-i] + x6[-i] + x7[-i] + x8[-i])
mse<-tail( anova(fit)[, 3], 1)
dfbetas1[i] = (mod$coeff[2]-fit$coeff[2])/sqrt(mse* c[2]) }
ifelse(dfbetas(mod)[,]>1,1,0)
Rule of thumb: Influential if the value exceeds 1 (for
small-medium datasets (n \leq 60) or if it exceeds 2\sqrt{\frac{p}{n}} (for large
datasets)
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```