MATH-650 Assignment 9

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Chapter 12: 14

```
require(leaps)
## Loading required package: leaps
data <- read.csv('case1102.csv')</pre>
data$logY = log(data$Brain/data$Liver)
Y <- data$logY
X <- data[,c('Days', 'Sex', 'Weight', 'Loss', 'Tumor')]</pre>
We use the leaps package to perform subset selection.
rsubsets <- regsubsets(logY ~ Days+Sex+Weight+Loss+Tumor, data=data)</pre>
s <- summary(rsubsets, matrix.logical=TRUE)</pre>
s$cp
## [1] 9.457598 1.430200 2.006538 4.000835 6.000000
Part (a): C_p
plot(rsubsets, scale='Cp')
     1.4 -
       2 -
S
       4 -
       6
     9.5 -
                                                     Weight
                                         SexMale
```

The way to interpet this plot is to look at first the smallest C_p values, which happens to be around 1.4 and see the black dots which in this case are given by Days, SexMale So if we were to choose the covarates based only on C_p values, we select: Days and Sex Here p=5 and in principle any model with $C_p < p$ is better than the full model, so we can also select these:

```
• Days, Sex: C_p=1.43
• Days, Sex, Weight: C_p=2.006
• Days, Sex, Weight, Tumor: C_p=4.00008
```

Part (b): Forward Selection

```
rsubsets <- regsubsets(logY ~ Days+Sex+Weight+Loss+Tumor,
                       data=data,
                       method='forward')
sforward <- summary(rsubsets, matrix.logical=TRUE)</pre>
sforward
## Subset selection object
## Call: regsubsets.formula(logY ~ Days + Sex + Weight + Loss + Tumor,
      data = data, method = "forward")
## 5 Variables (and intercept)
##
          Forced in Forced out
## Days
              FALSE
                         FALSE
              FALSE
## SexMale
                         FALSE
## Weight
              FALSE
                         FALSE
## Loss
              FALSE
                         FALSE
                         FALSE
## Tumor
              FALSE
## 1 subsets of each size up to 5
## Selection Algorithm: forward
            Days SexMale Weight Loss Tumor
                    TRUE FALSE FALSE FALSE
## 1 ( 1 ) FALSE
## 2 (1) TRUE
                     TRUE FALSE FALSE FALSE
## 3 (1) TRUE
                     TRUE
                           TRUE FALSE FALSE
## 4 ( 1 ) TRUE
                     TRUE
                           TRUE FALSE
## 5 (1) TRUE
                     TRUE
                           TRUE TRUE TRUE
```

Part (c): Backward Selection

```
## Days
              FALSE
                         FALSE
## SexMale
              FALSE
                         FALSE
## Weight
              FALSE
                         FALSE
                         FALSE
## Loss
              FALSE
## Tumor
              FALSE
                         FALSE
## 1 subsets of each size up to 5
## Selection Algorithm: backward
            Days SexMale Weight Loss Tumor
## 1
     (1) FALSE
                    TRUE FALSE FALSE FALSE
## 2 ( 1 ) TRUE
                    TRUE
                         FALSE FALSE FALSE
## 3 (1) TRUE
                    TRUE
                           TRUE FALSE FALSE
## 4 (1)
            TRUE
                    TRUE
                           TRUE FALSE
                                      TRUE
## 5 (1) TRUE
                    TRUE
                           TRUE TRUE
                                      TRUE
```

Part(d): Stepwise Regression

```
rsubsets <- regsubsets(logY ~ Days+Sex+Weight+Loss+Tumor,</pre>
                       data=data,
                       method="segrep")
sboth <- summary(rsubsets, matrix.logical=TRUE)</pre>
sboth
## Subset selection object
## Call: regsubsets.formula(logY ~ Days + Sex + Weight + Loss + Tumor,
       data = data, method = "segrep")
## 5 Variables (and intercept)
##
           Forced in Forced out
## Days
               FALSE
                          FALSE
## SexMale
               FALSE
                          FALSE
                          FALSE
## Weight
               FALSE
               FALSE
                          FALSE
## Loss
## Tumor
               FALSE
                          FALSE
## 1 subsets of each size up to 5
## Selection Algorithm: 'sequential replacement'
##
             Days SexMale Weight Loss Tumor
## 1 ( 1 ) FALSE
                     TRUE FALSE FALSE FALSE
## 2 (1) TRUE
                     TRUE FALSE FALSE FALSE
## 3 (1) TRUE
                     TRUE
                            TRUE FALSE FALSE
## 4 ( 1 ) TRUE
                     TRUE
                            TRUE FALSE
                                        TRUE
## 5 (1) TRUE
                     TRUE
                            TRUE TRUE
                                        TRUE
```

Conclusion

From the above, we conclude that the variable selection in this case gives us the same set for all four methods.