## HW3

Keyi Zhong

5/7/2020

### 1.1

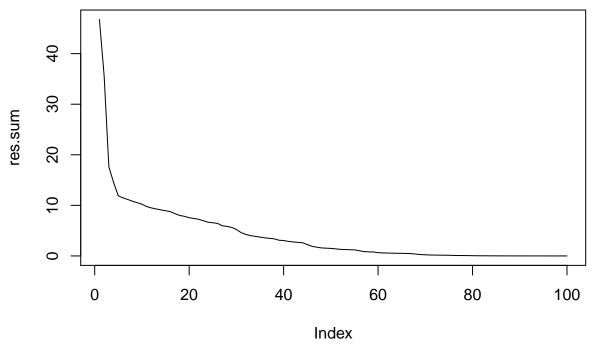
```
source("test-data.R")
library("leaps")
truncated.power.design.matrix <- function(x) {
    result <- matrix(0,nrow=length(x),ncol=length(x))
    for (i in 1:length(x)) {
        for (j in 1:length(x)) {
            result[i,j] <- max(0,x[i]-x[j])
        }
    }
    for (j in 1:length(x)) {
        result[j,length(x)] <- 1
    }
    return(result)
}</pre>
```

### 1.2

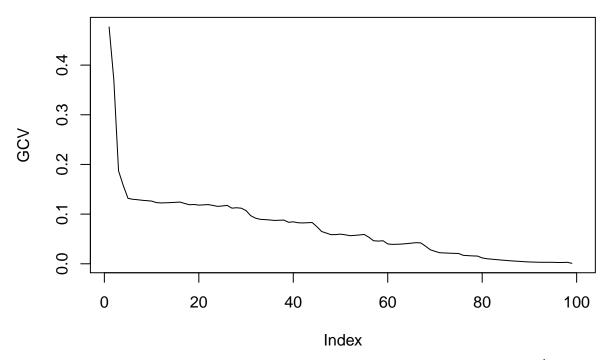
```
regsubsets.fitted.values <- function(X, regsubsets.out, nterm) {
  out.summary <- summary(regsubsets.out)
  vorder <- regsubsets.out$vorder[1:nterm][order(regsubsets.out$vorder[1:nterm])]
  coeff <- coef(regsubsets.out,nterm)
  coefficient <- rep(0,100)
  for (i in 1:nterm) {
    coefficient[vorder[i]] <- coeff[i]
  }
  return(X%*%coefficient)
}</pre>
```

```
truncated <- truncated.power.design.matrix(x)
regsubsets.out <- regsubsets(truncated,y,method="forward",nvmax=100,intercept = FALSE)
out.summary <- summary(regsubsets.out)
res.sum <- rep(0,100)
for(k in 1:100) {
  fit.values <- regsubsets.fitted.values(truncated,regsubsets.out,k)</pre>
```

```
res.sum[k] <- sum((y-fit.values)^2)
}
plot(res.sum,type='l')</pre>
```

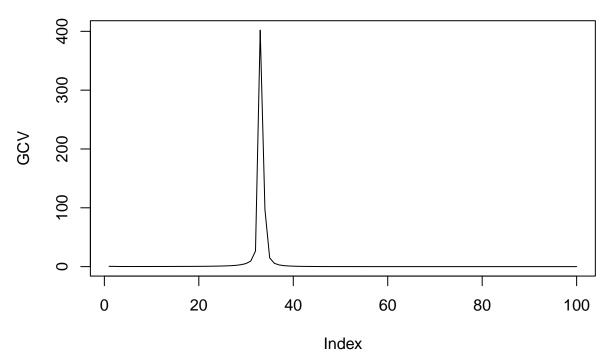


```
GCV <- rep(0,100)
for(i in 1:100) {
  fit.values <- regsubsets.fitted.values(truncated,regsubsets.out,i)
  GCV[i] <- mean((y-fit.values)^2/(1-i/100)^2)
}
plot(GCV,type='l')</pre>
```



Yes, because GCV score is high with small k, because GCV is basically residual sum,  $1 - \frac{k}{n}$  is decreasing

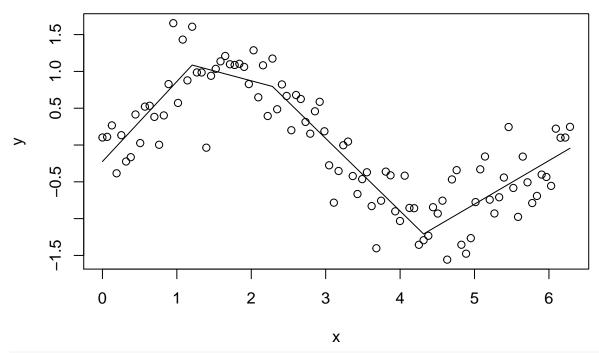
```
GCV <- rep(0,100)
for(i in 1:100) {
  fit.values <- regsubsets.fitted.values(truncated,regsubsets.out,i)
  GCV[i] <- mean((y-fit.values)^2/(1-3*i/100)^2)
}
plot(GCV,type='l')</pre>
```



No, 3\*k close to 100 when k is 33 so it gives highest score. With different degree of freedom, it gives different highest score

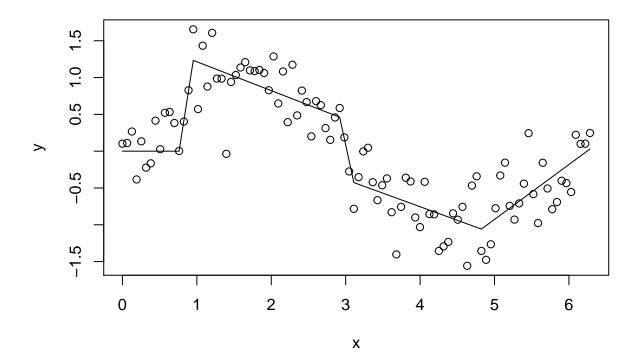
```
k <- which(GCV==min(GCV[1:30]))
fit.values <- regsubsets.fitted.values(truncated,regsubsets.out,k)
plot(x,y,main='forward')
lines(x,fit.values)</pre>
```

# forward



regsubsets.out <- regsubsets(truncated,y,method="backward",nvmax=100,intercept = FALSE)
fit.values <- regsubsets.fitted.values(truncated,regsubsets.out,k)
plot(x,y,main='backward')
lines(x,fit.values)</pre>

# backward



### 2.1

```
\begin{split} \hat{a} &= argmin||y - Xa||^2 + \lambda a^T \Omega a \ \frac{d}{da}||y - Xa||^2 + \lambda a^T \Omega a = 0 = \frac{d}{da} a^T X^T a - a^T X^T y - y^T X a + y^T y + \lambda a^T \Omega a \\ &= 2X^T X a - 2X^T y + 2\lambda \Omega a = X^T X a - X^T y + \lambda \Omega a \ a = (X^T X + \lambda \Omega)^{-1} X^T y \end{split} so \hat{y} &= X(X^T X + \lambda \Omega)^{-1} X^T y = S_{\lambda} y
```

### 2.2

```
library('glmnet')

## Loading required package: Matrix

## Loaded glmnet 3.0-2

lambda <- c(0,1,10,1000000)

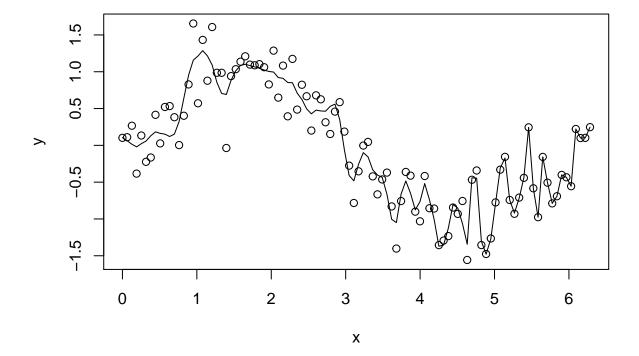
ridge.mod <- glmnet(truncated,y,alpha=0,lambda = c(0,1,10,1000000)))

coeff <- coef(ridge.mod)

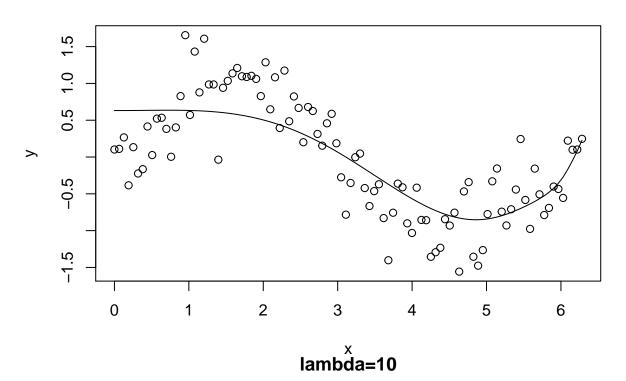
coeff[101,]=coeff[1,]

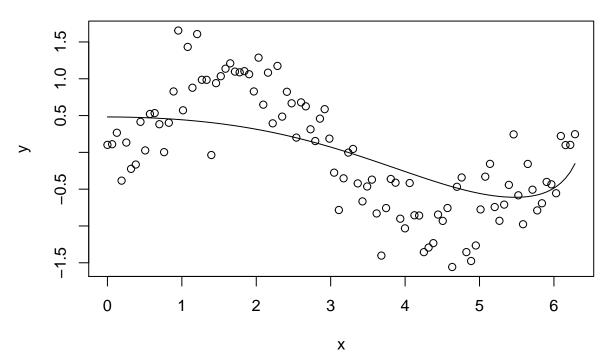
for (i in 1:4) {
   plot(x,y,main = paste0("lambda=",lambda[i]))
    lines(x,truncated%*%coeff[,5-i][2:101])
}</pre>
```

# lambda=0

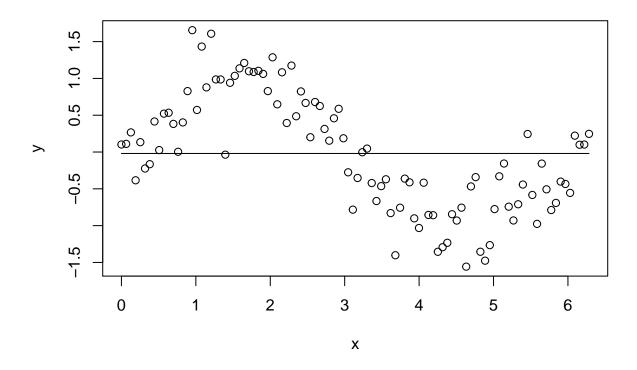


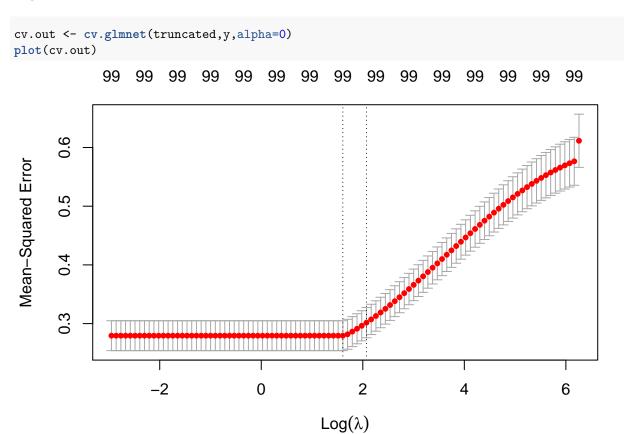
# lambda=1



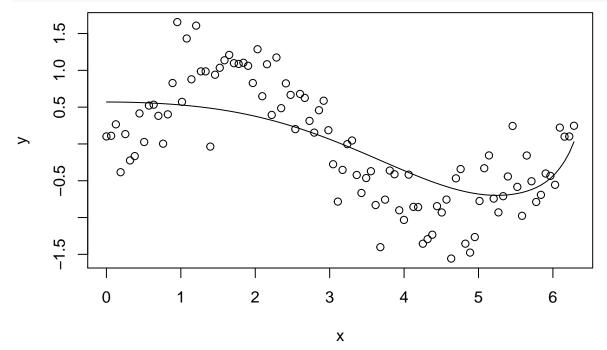


# lambda=1e+06





```
optlam <- cv.out$lambda.min
ridge.mod <- glmnet(truncated,y,alpha=0,lambda = optlam)
coeff <- coef(ridge.mod)
coeff[101,]=coeff[1,]
y_hat <- truncated%*%coeff[2:101]
plot(x,y)
lines(x,y_hat)</pre>
```



3

 $\mathbf{a}$ 

Best subset gives the smallest training RSS, because it chooses the best k among all predictors as backward and forward both choose k from top to bottom or vice versa.

### b

Also best subset, because we don't have control on test set, in general it gives the best prediction.

 $\mathbf{c}$ 

1

True, because forward choose k from beginnig to the end, so a k+1 model must include.

 $\mathbf{2}$ 

True, same reason as last question.

|  | ٠, |  |
|--|----|--|
|  |    |  |

False, forward and backward choose from different direction.

### 4

False, same as last question.

### 5

False, the additional variable could take one of the predictor's place.

### 4

 $\mathbf{a}$ 

Increase, as  $\lambda$  increase, the model is more off, because of the  $-\lambda \sum \beta^2$  part.

### $\mathbf{b}$

Decrease then increase, first, increasing  $\lambda$  can make the model close to the true model, but it will eventually off the model.

### $\mathbf{c}$

Decrease, it lessen the variance of  $x_{ij}$ 

### $\mathbf{d}$

Increase, it makes the model off the predict model.

#### $\mathbf{e}$

Not change, irreducible error does not depend on the model