STAT542 Statistical Learning Homework 2

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Question 1

a) [15 points]

Answer:

```
library(leaps)
# Read data set
data = read.csv('bitcoin dataset.csv')
# Get the index of train and test data
index1 = which(data[,1] == '2017-01-01 00:00:00')
index2 = which(data[,1] == '2017-09-12 00:00:00')
# Here we treat the variable Date as a continuous variable, say starting
# with value 0 at the first instances, then increase by 1 for each two days.
data$Date = 1:dim(data)[1]
# Split the data set into train and test data
# We remove Date information and treat the 2rd column 'btc_market_price'
# as the outcome variable
train data = data[1:index1-1,]
test data = data[index1:index2,]
train data x = train data[,-2]
train data y = train data[,2]
test_data_x = test_data[,-2]
test data y = test data[,2]
# Ignore the variable btc_trade_volume because it contains missing values
train_data_x = subset(train_data_x,select = -btc_trade_volume)
test_data_x = subset(test_data_x,select = -btc_trade_volume)
# Performs an exhaustive search over models, and gives back the best model
# (with low RSS) of each size.
RSSleaps=regsubsets(as.matrix(train data x), train data y,
                    nvmax = length(train data x))
best_subset= summary(RSSleaps, matrix=T)
```

You can find the best model of each size (1-22) from the output from summary(RSS1eaps, matrix=T). However, since the output maybe too wide (many columns) to fit a page, we will only report present a part of it. The following is the best model of size = 10 which means the best model with using only ten features(except Intercept).

```
#best subset
coef(RSSleaps, 10)
##
                 (Intercept)
                                   btc_total_bitcoins
                                                                  btc_market_cap
               9.204666e-01
                                        -8.102118e-07
##
                                                                    6.938252e-08
##
            btc blocks size
                                   btc avg block size
                                                          btc n orphaned blocks
##
              -7.289203e-03
                                          2.176334e+01
                                                                   -1.507009e+00
##
              btc hash rate
                                       btc difficulty
                                                             btc miners revenue
##
              -2.436673e-05
                                        -2.104494e-10
                                                                    1.859082e-05
## btc cost per transaction btc n transactions total
               5.870465e-01
                                          3.782071e-06
##
```

You can also get the the result of each size from the plot with their R square

```
plot(RSSleaps,scale='r2')
# The models are ordered by the specified model selection statistic. This plot is
# particularly useful when there are more than ten or so models and the simple
# table produced by summary.regsubsets is too big to read. Here we didn't plot here
# due the limitation of report page.
```

b) [15 points]

Answer:

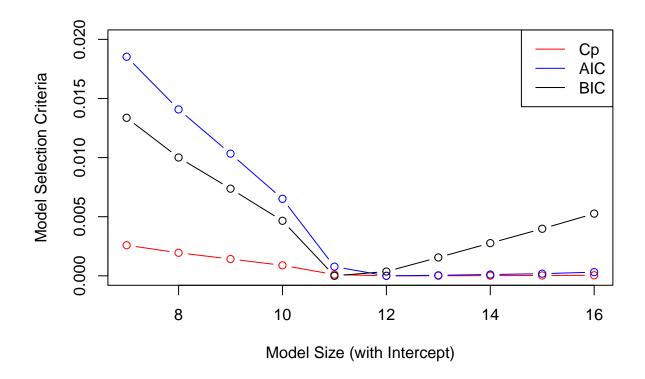
```
lmfit=lm(train data y~as.matrix(train data x))
msize=apply(best subset$which,1,sum)
n=dim(train data x)[1]
p=dim(train_data_x)[2]
# Calculate the Cp, AIC, BIC of the best model of each size
Cp = best subset$rss/(summary(lmfit)$sigma^2) + 2*msize - n
AIC = n*log(best subset$rss/n) + 2*msize
BIC = n*log(best subset$rss/n) + msize*log(n)
# Select the best model
result = data.frame(which.min(Cp), which.min(AIC), which.min(BIC),
                    row.names = "The best model")
colnames(result) = c("Cp","AIC","BIC")
result
##
                  Cp AIC BIC
## The best model 11 11
```

Thus, the best model using C_P and AIC criteria is that with size = 11. And the best model using BIC criteria is that with size = 10 The C_p , AIC and BIC of these two models are showed as below.

```
cbind(Cp,AIC,BIC)[c(10,11),]
##
                      AIC
                               BIC
             Ср
## 10 13.164159 5733.904 5792.052
## 11 9.591456 5730.287 5793.721
The features used in the best model with size = 10 and are showed as below.
names(train data x)[best subset$which[10,-1]]
    [1] "btc_total_bitcoins"
##
                                     "btc_market_cap"
##
    [3] "btc_blocks_size"
                                     "btc_avg_block_size"
##
    [5] "btc_n_orphaned_blocks"
                                     "btc_hash_rate"
##
    [7] "btc difficulty"
                                     "btc miners revenue"
##
    [9] "btc cost per transaction" "btc n transactions total"
The features used in the best model with size = 11 and are showed as below.
names(train data x)[best subset$which[11,-1]]
    [1] "Date"
##
                                     "btc total bitcoins"
##
   [3] "btc_market_cap"
                                     "btc_blocks_size"
    [5] "btc avg block size"
                                     "btc n orphaned blocks"
##
   [7] "btc hash rate"
                                     "btc difficulty"
##
##
    [9] "btc miners revenue"
                                     "btc_cost_per_transaction"
## [11] "btc_n_transactions_total"
Here we resacle C_p, AIC, BIC to (0,1) and made plots of these criterion of size from 6 to 15 (including
Intercept).
# Rescale Cp, AIC, BIC to (0,1).
inrange \leftarrow function(x) { (x - min(x)) / (max(x) - min(x)) }
Cp = best_subset$cp; Cp = inrange(Cp);
BIC = best subset$bic; BIC = inrange(BIC);
AIC = n*log(best_subset$rss/n) + 2*msize; AIC = inrange(AIC);
# Since we know when size = 10,11 we can get the minimun, here we
# plot the Model Selection Criteria(Cp, AIC, BIC) of size = 6-15
# zoom in
id=6:15:
plot(range(msize[id]), c(0, 0.02), type="n", xlab="Model Size (with Intercept)",
     ylab="Model Selection Criteria")
points(msize[id], Cp[id], col="red", type="b")
points(msize[id], AIC[id], col="blue", type="b")
points(msize[id], BIC[id], col="black", type="b")
```

legend("topright", lty=rep(1,3), col=c("red", "blue", "black"),

legend=c("Cp", "AIC", "BIC"))



Then we applied the fitted models to the testing dataset and report the prediction error of $n_{test}^{-1} \sum_{i \in test} (\hat{Y}_i - Y_i)^2$

[1] 46876.07

 mse_11

So, the prediction error of the best model with size = 10 which is selected by BIC criteria is 44235.49. And the prediction error of the best model with size = 11 which

is selected by C_p and AIC criteria is 46876.07.

c) [15 points]

Answer:

The best model of each size (1-22) can be found from the output from summary(RSS1eaps, matrix=T) Here we will only report present a part of it. The following is the best model of size = 10 which means the best model with using only ten features(except Intercept).

```
#best subset
coef(RSSleaps, 10)
##
                     (Intercept)
                                                          Date
##
                  -9.552736e-02
                                                 -5.264735e-03
             btc_total_bitcoins
##
                                                btc market cap
                   5.145858e-07
                                                 -8.651644e-11
##
                                           btc_avg_block_size
##
                btc blocks size
##
                  -7.743416e-04
                                                  1.628053e+00
## btc median confirmation time
                                                btc difficulty
##
                  -2.552420e-02
                                                 -1.296557e-11
##
       btc_cost_per_transaction
                                       btc_n_unique_addresses
                                                  3.466938e-06
##
                   4.430324e-02
##
       btc n transactions total
                   4.536846e-07
##
# You can also get the the result of each size from the plot with their R square
# plot(RSSleaps, scale='r2')
```

Then we use C_p , AIC and BIC criteria to select the best model.

```
# Calculate the Cp, AIC, BIC of the best model of each size
lmfit=lm(log(train_data_y+1)~as.matrix(train_data_x))
msize=apply(best_subset$which,1,sum)
n=dim(train_data_x)[1]
p=dim(train_data_x)[2]
Cp = best_subset$rss/(summary(lmfit)$sigma^2) + 2*msize - n
AIC = n*log(best_subset$rss/n) + 2*msize
BIC = n*log(best_subset$rss/n) + msize*log(n)

# Select the best model
```

```
result = data.frame(which.min(Cp), which.min(AIC), which.min(BIC),
                    row.names = "The best model")
colnames(result) = c("Cp", "AIC", "BIC")
result
##
                  Cp AIC BIC
## The best model 14
                     14
Thus, the best model using C_P and AIC criteria is that with size = 14. And the best
model using BIC criteria is that with size = 13 The C_p, AIC and BIC and the features
of these two models are showed as below.
cbind(Cp,AIC,BIC)[c(13,14),]
##
                     AIC
                                BIC
            Ср
## 13 17.23428 -3362.928 -3288.921
## 14 15.22976 -3364.968 -3285.675
# Selected features in the best model
names(train_data_x)[best_subset$which[13,-1]]
    [1] "Date"
##
##
    [2] "btc_total_bitcoins"
    [3] "btc_market_cap"
##
##
    [4] "btc_blocks_size"
    [5] "btc avg block size"
##
##
    [6] "btc median confirmation time"
##
    [7] "btc_difficulty"
##
    [8] "btc_transaction_fees"
##
    [9] "btc_cost_per_transaction"
## [10] "btc_n_unique_addresses"
## [11] "btc_n_transactions_total"
## [12] "btc_n_transactions_excluding_chains_longer_than_100"
  [13] "btc estimated transaction volume usd"
names(train data x)[best subset$which[14,-1]]
##
    [1] "Date"
##
    [2] "btc_total_bitcoins"
##
    [3] "btc_market_cap"
##
    [4] "btc blocks size"
##
    [5] "btc avg block size"
##
    [6] "btc_median_confirmation_time"
##
    [7] "btc difficulty"
##
    [8] "btc_transaction_fees"
##
   [9] "btc_cost_per_transaction"
## [10] "btc_n_unique_addresses"
## [11] "btc_n_transactions_total"
## [12] "btc n transactions excluding chains longer than 100"
## [13] "btc_estimated_transaction_volume"
```

```
## [14] "btc estimated transaction volume usd"
```

Then we applied the fitted models to the testing dataset and report the prediction error.

[1] 4426656

```
# The MSE of the best model with size = 14 (Cp, AIC)
mse_log_14
```

```
## [1] 4426691
```

So, the prediction error of the best model with size = 13 which is selected by BIC criteria is 4426656. And the prediction error of the best model with size = 14 which is selected by C_p and AIC criteria is 4426691.

Question 2

Part I [35 points]

Answer:

Firstly, we will not penalize the intercept term β_0 which means that we will center both your X and Y first and perform the algorithm in the question. Then let's derive the soft thresholding function/update rule of Lasoo using coordinate descent algorithm.

In the question, assume we have a scaled and centered dataset X with N instances, each of which consists of p features, and a centered outcome Y. We will use the objective function

$$f(\beta) = \frac{1}{2n} ||Y - X\beta||_2^2 + \lambda \sum_{j=1}^p |\beta_j| = \frac{1}{2n} \sum_{i=1}^N (y_i - \sum_{j=1}^p \beta_j x_{ij})^2 + \lambda \sum_{j=1}^p |\beta_j|$$

Consider the update rule of β_j , we take the derivative with respect to β_j of the objective function. Here we can split the objective function into two part: 1) Loss function term $g(\beta) = \frac{1}{2n} \sum_{i=0}^{N} (y_i - \sum_{j=1}^{p} \beta_j x_{ij})^2$ 2) Regularization function term $h(\beta) = \lambda \sum_{j=1}^{p} |\beta_j|$

Thus, the derivative with respect to β_j of the Loss function term is

$$\frac{\partial g(\beta)}{\partial \beta_j} = -\frac{1}{n} \sum_{i=1}^{N} (y_i - \sum_{j=1}^{p} \beta_j x_{ij}) x_{ij}$$

$$= -\frac{1}{n} \sum_{i=1}^{N} (y_i - \sum_{k \neq j}^{p} \beta_k x_{ik} - \beta_j x_{ij}) x_{ij}$$

$$= -\frac{1}{n} \sum_{i=1}^{N} (y_i - \sum_{k \neq j}^{p} \beta_k x_{ik}) x_{ij} + \frac{1}{n} \beta_j \sum_{i=1}^{N} x_{ij}^2$$

$$= -\frac{1}{n} p_j + \frac{1}{n} \beta_j z_j$$

where

$$p_{j} = \sum_{i=1}^{N} (y_{i} - \sum_{k \neq j}^{p} \beta_{k} x_{ik}) x_{ij}$$

$$= \sum_{i=1}^{N} (y_{i} - \sum_{k=1}^{p} \beta_{k} x_{ik} + \beta_{j} x_{ij}) x_{ij}$$

$$= \sum_{i=1}^{N} (r_{i} + \beta_{j} x_{ij}) x_{ij}$$

$$r_{i} = y_{i} - \sum_{k=1}^{p} \beta_{k} x_{ik}$$

$$z_{j} = \sum_{i=1}^{N} x_{ij}^{2}$$

For the regularization term, we can take the derivative with respect to β_j using subgradient method. So the subdifferential is

$$\partial_{\beta_j} \lambda \sum_{j=1}^p |\beta_j| = \partial_{\beta_j} \lambda |\beta_j| = \begin{cases} \lambda & \text{if } \beta_j > 0\\ [-\lambda, \lambda] & \text{if } \beta_j = 0\\ -\lambda & \text{if } \beta_j < 0 \end{cases}$$

So the have the derivative with respect to β_j of the objective function.

$$\frac{\partial f(\beta)}{\partial \beta_j} = \frac{\partial g(\beta)}{\partial \beta_j} + \frac{\partial h(\beta)}{\partial \beta_j} = \begin{cases} -\frac{1}{n}p_j + \frac{1}{n}\beta_j z_j + \lambda & \text{if } \beta_j > 0\\ [-\frac{1}{n}p_j + \frac{1}{n}\beta_j z_j - \lambda, -\frac{1}{n}p_j + \frac{1}{n}\beta_j z_j + \lambda] & \text{if } \beta_j = 0\\ -\frac{1}{n}p_j + \frac{1}{n}\beta_j z_j - \lambda & \text{if } \beta_j < 0 \end{cases}$$

Let the derivative to be zero $\partial_{\beta_j} f(\beta) = 0$ and solve for $\hat{\beta}_j$. We can get

$$\hat{\beta}_{j} = \begin{cases} \frac{p_{j} - n\lambda}{z_{j}} & \text{if } p_{j} > n\lambda \\ 0 & \text{if } p_{j} \in [-n\lambda, n\lambda] \\ \frac{p_{j} + n\lambda}{z_{j}} & \text{if } p_{j} < n\lambda \end{cases}$$
$$= \frac{1}{z_{j}} sign(p_{j})(|p_{j}| - n\lambda)_{+}$$

According to the soft thresholding function. we can write the Lasso fitting function using coordinate descent algorithm as below. The detail about the function can be found in the comments.

```
# Function LassoFit
# Usage:
# LassoFit(X, y, mybeta = rep(0, ncol(X)), lambda, tol = 1e-10, maxitr = 500)
# Arguments
# myX: input matrix, each row is an observation vector (don't need to be
       scaled and centered)
# myY: response variable (don't need to be centered)
# mybeta: the initialization of beta
# mylambda: tuning parameter (penalty level), which controls the amount of
       shrinkage. Usually, mylambda >= 0.
# tol: Convergence threshold for coordinate descent. Each inner coordinate-
       descent loop continues until the change in the objective function value
       after any update is less than tol (or run maxitr iterations). Defaults
       value is 1E-10.
# maxitr: The maximum number of iteration in coordinate-descent loop. Defaults
       value is 500.
# Value(Output)
# beta 0: The intercept term in our fitted model of the original scale of X.
# beta: The coefficient in our fitted model of the original scale of X.
LassoFit <- function(myX, myY, mybeta, mylambda, tol = 1e-10, maxitr = 500){
  # First we scale and center X, and record them. Also center y and record it.
  # dont scale it. Now since both y and X are centered at 0, we don't need to worry
  # about the intercept anymore.
  x center = colMeans(myX)
 x_scale = apply(myX, 2, sd)
 X = scale(myX)
 y mean = mean(myY)
 Y = myY - y mean
  # Calculate the number of instances
 n = nrow(X)
  # Calculate zj = \sum (x_i j)^2 in the formula
  z = apply(X, 2, function(x) sum(x^2))
  # Initia a matrix to record the objective function value
  f = rep(0, maxitr)
  # Start the iteration unless meet the stopping rule or run maxitr iterations
  for (k in 1:maxitr){
    # Compute the residual using current beta
    r = Y - X \% mybeta
    # Record the objective function value
    f[k] = sum(r*r) / (2*n) + mylambda * sum(abs(mybeta))
    # Calculate the stopping rule
```

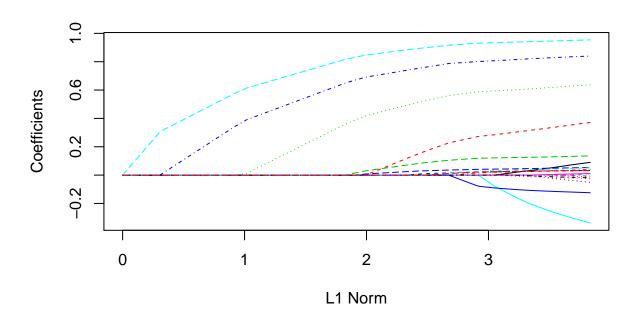
```
if (k > 10){
     if (abs(f[k] - f[k-1]) < tol) break;
   }
   # Use coordinate descent to update beta
   for (j in 1:ncol(X)){
     # Add the effect of jth variable back to r
       r = r + X[,j] * mybeta[j]
     # Calculate the pj in the formula
       p = sum(X[,j] * r)
     # Using the soft thresholding function
       # Remove the new effect of jth varaible out of r
       r = r - X[,j] * mybeta[j]
     # You can also write as the follow structure
     # Caluculate the pj in the formula
     \# p = sum(X[,j] * (Y - X[,-j] %*% mybeta[-j]))
     # Using the update rule / soft thresholding function
     # mybeta[j] = sign(p) * ifelse(abs(p)>n*mylambda,abs(p)-n*mylambda,0) / z[j]
   }
 }
 # Scale the beta back
 mybeta = mybeta / x_scale
 # Recalculte the intercept term in the original, uncentered and unscaled X
 beta 0 = mean(myY - myX %*% mybeta)
 # Return the beta and intercept
 return(list("beta" = mybeta, "beta 0" = beta 0))
}
```

Here, to demonstrate that the code is correct, we compare the results to glmnet. And compare the fitted beta values. First, we generate a matrix X and outcome Y.

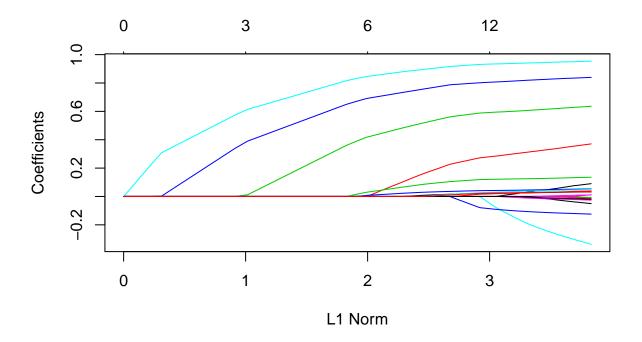
```
library(MASS)
library(glmnet)
set.seed(1)
N = 400
P = 20
Beta = c(1:5/5, rep(0, P-5))
Beta0 = 0.5
# Genrate X
V = matrix(0.5, P, P)
diag(V) = 1
X = as.matrix(mvrnorm(N, mu = 3*runif(P)-1, Sigma = V))
# Create artifical scale of X
X = sweep(X, 2, 1:10/5, "*")
# Genrate Y
y = Beta0 + X %*% Beta + rnorm(N)
```

Now we want to compare the result from our algorithm and glmnet

```
# Set the lambda sequence
lambda = exp(seq(log(max(abs(cov(scale(X), (y-mean(y)))))), log(0.001),
                 length.out = 100))
# Initiate a matrix that records the fitted beta for each lambda value
beta_all = matrix(NA, ncol(X), length(lambda))
# This vecter stores the intercept of each lambda value
beta0 all = rep(NA, length(lambda))
# Initialize beta
bhat = rep(0, ncol(X))
# Loop from the largest lambda value
for (i in 1:length(lambda)){
 lasso_beta = LassoFit(X, y, bhat, lambda[i])
 bhat = lasso beta$beta
 beta all[, i] = bhat
 beta0_all[i] = lasso_beta$beta_0
}
# Here we make a plot of the L1 Norm versus Coefficients
matplot(colSums(abs(beta all)), t(beta all), type="l",xlab = "L1 Norm",
       ylab = "Coefficients")
```



```
# We can also make a plot from the result of glmnet
plot(glmnet(X, y, lambda = lambda, alpha = 1))
```



We can find the two plots from our function and glmnet result are almost the same. In the other hand, we use the lambda sequence which is used in the glmnet result and rerun our algorithms. Then we compare the coefficients and intercept from our algorithm and glmnet to demonstrate that the code is correct.

```
# Set our lambda to the lambda value from glmnet and rerun your algorithm
lambda = glmnet(X, y)$lambda
beta_all = matrix(NA, ncol(X), length(lambda))
beta0 all = rep(NA, length(lambda))
# Initialize beta
bhat = rep(0, ncol(X))
# loop from the largest lambda value
for (i in 1:length(lambda)){
  lasso_beta = LassoFit(X, y, bhat, lambda[i])
  bhat = lasso_beta$beta
  beta all[, i] = lasso beta$beta
  beta0 all[i] = lasso beta$beta 0
}
# Then this distance should be pretty small which is no more than 0.01
glmnet result = glmnet(X, y, alpha = 1)
max(abs(beta all - glmnet result$beta))
## [1] 0.002095227
max(abs(beta0_all - glmnet_result$a0))
```

[1] 0.001404763

According to the result, we can say that the code is correct.

Part II [30 points]

Answer:

Here we use the lambda sequence from part of the result of glmnet, and use mean squared error as the test error to select the best model.

```
# Set our lambda to the lambda value from glmnet
result = glmnet(as.matrix(train_data_x),train_data_y,alpha = 1, nlambda = 35)
lambda = result$lambda[9:22]
# Initiate a matrix that records the fitted beta for each lambda value
beta all = matrix(NA, ncol(train data x), length(lambda))
# This vector stores the intercept of each lambda value
beta0 all = rep(NA, length(lambda))
# This vecter stores the test error of each lambda value
MSE = rep(NA, length(lambda))
# Initialize beta
bhat = rep(0, ncol(train data x))
# Loop from the largest lambda value
for (i in 1:length(lambda)){
  # Train the Lasso model
 lasso_beta = LassoFit(as.matrix(train_data_x), train_data_y, bhat, lambda[i])
  # Store the coefficients and intercepts
  beta_all[, i] = lasso_beta$beta
  beta0 all[i] = lasso beta$beta 0
  # Apply the fitted model on the test dataset
  test_y_hat = as.matrix(test_data_x) %*% lasso_beta$beta + lasso_beta$beta_0
  # Calculate the test error
 MSE[i] = mean((test y hat-test data y)^2)
}
```

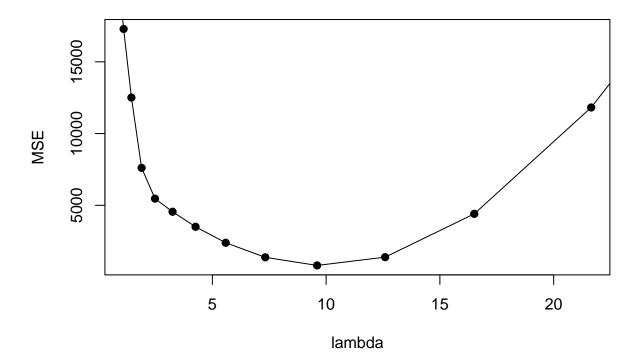
Then, the test error of each lambda is showed as below. We can also make plots of lambda versus MSE.

```
# The test error of each lambda cbind(lambda, MSE)
```

```
##
            lambda
                          MSE
##
   [1,] 28.3854431 25487.0375
   [2,] 21.6495373 11816.6553
##
   [3,] 16.5120715 4405.5510
   [4,] 12.5937336 1386.3619
##
   [5,] 9.6052228
##
                   810.1388
   [6,] 7.3258898 1374.9794
##
##
   [7,] 5.5874458 2390.0072
##
   [8,] 4.2615370 3504.0176
##
   [9,] 3.2502683 4551.3638
## [10,] 2.4789750 5465.1725
```

```
## [11,] 1.8907107 7610.5805
## [12,] 1.4420424 12508.4709
## [13,] 1.0998437 17288.9823
## [14,] 0.8388492 21551.4500

# Make plots of lambda versus MSE
sort_result = cbind(lambda,MSE)[order(cbind(lambda,MSE)[,2],decreasing=F),]
plot(MSE~lambda,data = sort_result[1:12,],pch=19)
lines(MSE~lambda)
```

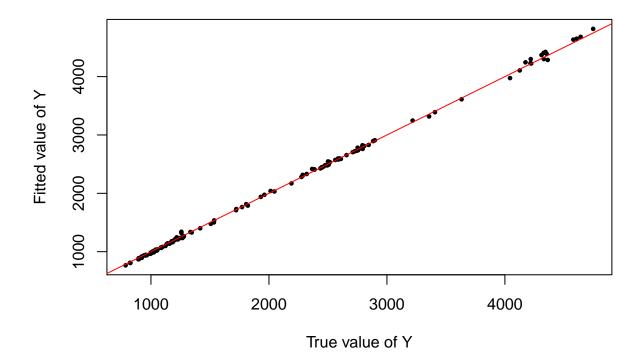


We select the model with minimum test error as the best model. The test error, lambda and coefficient are showed as below.

```
## Coefficient
## btc_market_cap 5.460582e-08
## btc_miners_revenue 4.265463e-05
## btc_cost_per_transaction 1.071608e+00
## Intercept 5.740412e+00
```

Thus, this is the coefficient and features in the best model. Now we can make a plot of the L1 Norm versus Coefficients (We don't show here due to the limitation of pages.)

Finally, we make a plot of fitted Y values versus the true value of y. We can find that lasso perform pretty well on the data set.



There will be a **More thought** part to analysis why the performance of lasso on the test data is much better than the best subset selection. The reason is about the distribution difference between train and test data, and it may be overfitting in Problem 1 since there is an enormous search space. However, we will not show the analysis here due to the limitation of report page. You can find more detail in the Rmd file.