Homework 2

Liting Hu

November 15, 2016

Problem 1

1. Logistic regression

sum(lda.pred\$posterior[, 1] > .9)

lda.pred <- rep("No", dimDP[1])</pre>

DefaultPredict\$lda.pred <- lda.pred</pre>

```
# Logistic Regression
glm.fit <- glm(default~balance+student+income, data=Default, family=binomial)</pre>
summary(glm.fit)
summary(glm.fit$coefficients)
glm.probs <- predict(glm.fit, data=Default, type ="response")</pre>
glm.pred=rep("No", 10000)
glm.pred[glm.probs >.5]="Yes"
table(glm.pred, default)
mean(glm.pred == default)
# 0.9732
glm.probs <- predict(glm.fit, DefaultPredict, type="response")</pre>
glm.pred <- rep("No", dimDP[1])</pre>
glm.pred[glm.probs >.5] <- "Yes"</pre>
DefaultPredict$glm.pred <- glm.pred</pre>
2. Linear discriminant analysis
lda.fit <- lda(default~balance+student+income, data=Default)</pre>
lda.fit
plot(lda.fit)
lda.pred <- predict(lda.fit, Default)</pre>
lda.class <- lda.pred$class</pre>
table(lda.class, default)
mean(lda.class == default)
# 0.9724
sum(lda.pred$posterior[, 1] >= .5)
sum(lda.pred$posterior[, 1] < .5)</pre>
```

lda.probs <- predict(lda.fit, DefaultPredict, type="response")</pre>

lda.pred[lda.probs\$posterior[, 2] > .5] <- "Yes"</pre>

3. quadratic discriminant analysis

```
qda.fit <- qda(default~balance+student+income, data=Default)
qda.fit

qda.probs <- predict(qda.fit, Default)
qda.class <- qda.pred$class
table(qda.class, default)
mean(qda.class == default)
# 0.973

qda.probs <- predict(qda.fit, DefaultPredict, type="response")
DefaultPredict$qda.pred <- qda.probs$class</pre>
```

4. K-nearest neighbor classification

```
# K-Nearest Neighbors
Default.X <- cbind(Default$student, Default$balance, Default$income)
DefaultPredict.X <- cbind(DefaultPredict$student, DefaultPredict$balance, DefaultPredict$income)
knn.pred1 <- knn(Default.X, DefaultPredict.X, default, k = 1)
knn.pred3 <- knn(Default.X, DefaultPredict.X, default, k = 3)
knn.pred6 <- knn(Default.X, DefaultPredict.X, default, k = 6)
knn.pred10 <- knn(Default.X, DefaultPredict.X, default, k = 10)
DefaultPredict$knn.pred1 <- knn.pred1
DefaultPredict$knn.pred3 <- knn.pred3
DefaultPredict$knn.pred6 <- knn.pred6
DefaultPredict$knn.pred10 <- knn.pred10
detach(Default)</pre>
```

All results are stored in dataframe DefaultPredict:

>	Defaul:	tPredict									
	index	student	balance	income	alm.pred	lda.pred	ada.pred	knn.pred1	knn.pred3	knn.pred6	knn.pred10
1	1	Yes			No	No	No	No	No	No	No
	2	No			No	No		No	No	No	No
2	3	Yes	771.7039646		No	No	No	No	No	No	No
4	4	No	754.6104381		No	No	No	No	No	No	No
5	5	No			No	No	No	No	No	No	No
5 6	6	No	510.0996253	49837.10	No	No	No	No	No	No	No
7	7	No	886.2620474	52852.43	No	No	No	No	No	No	No
8	8	No	831.1602780	39944.24	No	No	No	No	No	No	No
9	9	Yes	400.9717764	15317.38	No	No	No	No	No	No	No
10	10	No	1092.2819530	45468.86	No	No	No	No	No	No	No
11	11	No	0.1797604	41724.94	No	No	No	No	No	No	No
12	12	Yes	996.9292528	20057.01	No	No	No	No	No	No	No
13	13	No	376.3586612	25420.12	No	No	No	No	No	No	No
14	14	No	654.0181025	54844.04	No	No	No	No	No	No	No
15	15	No	1109.1008990	45525.52	No	No	No	No	No	No	No
16	16	No	937.2591203	56696.03	No	No	No	No	No	No	No
17	17	Yes	184.3792088	14995.35	No	No	No	No	No	No	No
18	18	No	710.9217520	53003.81	No	No	No	No	No	No	No
19	19	No	747.1608518	19649.92	No	No	No	No	No	No	No
20	20	No	852.5276677		No	No	No	No	No	No	No
21	21	No	1567.9187140	36668.33	No	No	No	No	No	No	No
22	22	Yes	187.4985488	16885.93	No	No	No	No	No	No	No
23	23	Yes	1490.8859400	17899.42	No	No	No	No	No	No	No
24	24	Yes	2204.5416940	14309.70	Yes	Yes	Yes	Yes	Yes	No	No
25	25	Yes	1777.7084840	20363.12	No	No	No	Yes	No	No	No
26	26	No	1877.1215410	48977.54	Yes	No	No	Yes	Yes	Yes	No
27	27	Yes	1902.5514700	20575.27	No	No	No	Yes	No	No	No
28	28		1571.8059100		No	No	No	Yes	No	No	No
29	29	No	1968.9988070	39084.55	Yes	Yes	Yes	Yes	Yes	No	Yes
30	30		1526.1299570		No	No	No	Yes	No	No	No
31	31		1641.5593160		No	No	No	Yes	No	No	No
32	32	No			Yes	Yes	Yes	Yes	Yes	Yes	Yes
33	33	No			No	No	No	Yes	No	No	No
34	34				No	No	No	Yes	No	No	No
35	35	No	1700.2979030	30458.13	No	No	No	No	No	No	No
36	36		1118.1097410		No	No	No	Yes	No	No	No
37	37		1132.5249430		No	No	No	Yes	No	No	No
38	38	No			Yes	Yes	No	Yes	Yes	Yes	No
39	39	No			No	No	No	Yes	Yes	Yes	No
40	40	No	1457.0124430		No	No	No	Yes	No	No	No
41	41	No			No	No	No	Yes	No	No	No
42	42	Yes	1759.1077950	15980.12	No	No	No	Yes	No	No	No

Figure 1: Default predict under different measures

In K-nearest neighbor classification, the results under k=1 and k=10 seem unreasonable compared to other classifications. Choosing a k value between 3 and 6 may produce a better result.

Problem 2

1. Read the Email Messages

There are 9353 messages in the 5 directories combined.

2. Find the Words in a Message

Firstly, define a function to split the message into header and body. The body of a message is separated from the header by a single empty line.

```
splitMessage = function(msg) {
    splitPoint = match("", msg)
    header = msg[1:(splitPoint-1)]
    body = msg[ -(1:splitPoint) ]
    return(list(header = header, body = body))
}
```

Then we should remove attachments from the message. Based on the anatomy of email messages, If an attachment is added to a message, the MIME type is multipart and the Content-Type field provides a boundary string that can be used to locate the attachments. To get the boundary, define function:

```
getBoundary = function(header) {
    boundaryIdx = grep("boundary=", header)
    boundary = gsub('"', "", header[boundaryIdx])
    gsub(".*boundary= *([^;]*);?.*", "\\1", boundary)
}
Then we can remove attachments from messages by
dropAttach = function(body, boundary){
    bString = paste("--", boundary, sep = "")
    bStringLocs = which(bString == body)
    # Search for the boundary string
    if (length(bStringLocs) <= 1) return(body)</pre>
    eString = paste("--", boundary, "--", sep = "")
    eStringLoc = which(eString == body)
    # The closing boundary
    if (length(eStringLoc) == 0)
        return(body[ (bStringLocs[1] + 1) : (bStringLocs[2] - 1)])
    n = length(body)
    if (eStringLoc < n)</pre>
        return( body[ c( (bStringLocs[1] + 1) : (bStringLocs[2] - 1),
                          ( (eStringLoc + 1) : n )) ] )
    return( body[ (bStringLocs[1] + 1) : (bStringLocs[2] - 1) ])
}
```

After all these steps, words can be extracted from the messages.

```
cleanText =
    function(msg)
        tolower(gsub("[[:punct:]0-9[:space:][:blank:]]+", " ", msg))
findMsgWords =
    function(msg) {
        if(is.null(msg))
            return(character())
        words = unique(unlist(strsplit(cleanText(msg), "[[:blank:]\t]+")))
        # drop empty and 1 letter words
        words = words[ nchar(words) > 1]
        invisible(words)
    }
processAllWords = function(dirName)
    # read all files in the directory
    fileNames = list.files(dirName, full.names = TRUE)
    # drop files that are not email, i.e., cmds
    notEmail = grep("cmds$", fileNames)
    if ( length(notEmail) > 0) fileNames = fileNames[ - notEmail ]
    messages = lapply(fileNames, readLines, encoding = "latin1")
    # split header and body
    emailSplit = lapply(messages, splitMessage)
    # put body and header in own lists
    bodyList = lapply(emailSplit, function(msg) msg$body)
    headerList = lapply(emailSplit, function(msg) msg$header)
    rm(emailSplit)
    # determine which messages have attachments
    hasAttach = sapply(headerList, function(header) {
        CTloc = grep("Content-Type", header)
        if (length(CTloc) == 0) return(0)
        multi = grep("multi", tolower(header[CTloc]))
        if (length(multi) == 0) return(0)
        multi
    })
    hasAttach = which(hasAttach > 0)
    # find boundary strings for messages with attachments
    boundaries = sapply(headerList[hasAttach], getBoundary)
    # drop attachments from message body
    bodyList[hasAttach] = mapply(dropAttach, bodyList[hasAttach],
                                 boundaries, SIMPLIFY = FALSE)
    # extract words from body
    msgWordsList = lapply(bodyList, findMsgWords)
```

```
invisible(msgWordsList)
}
msgWordsList = lapply(fullDirNames, processAllWords)
```

For now we have collected all the necessary words from all the emails. Create a logical vector to define whether the messages are spam or not based on the number of elements in each list and flatten all five lists into one list:

```
numMsgs = sapply(msgWordsList, length)
numMsgs
# [1] 5051 1400 500 1000 1397

isSpam = rep(c(FALSE, FALSE, FALSE, TRUE, TRUE), numMsgs)
msgWordsList = unlist(msgWordsList, recursive = FALSE)
```

3. The Naive Bayes Classifier

For spam messages (similar to ham messages), we should estimate the following probabilities (1/2 used to avoid zero probabilities):

$$P(\text{awordispresent}|\text{spam}) \approx \frac{\#\text{ofspammessageswiththisword} + 1/2}{\#\text{ofspammessages} + 1/2}$$

$$P(\text{awordisabsent}|\text{spam}) \approx \frac{\#\text{ofspammessageswithoutthisword} + 1/2}{\#\text{ofspammessages} + 1/2}$$

We use log of ratios to avoid products and it tends to have better statistical properties. So for a new message we compute the log likelihood ratio as

```
\sum_{\substack{\text{wordsinmessage}\\ \text{wordsnotinmessage}}} (\log P(\text{wordpresent}|\text{spam}) - \log P(\text{wordpresent}|\text{ham})) + \\ \sum_{\substack{\text{wordsnotinmessage}\\ \text{wordsnotinmessage}}} (\log P(\text{wordabsent}|\text{spam}) - \log P(\text{wordabsent}|\text{ham})) + \log P(\text{spam}) - \log P(\text{ham})
    where \log P(\text{spam}) - \log P(\text{ham}) should be constant.
To collect these probabilities, construct the function that returns all log likelihood ratios as a
matrix:
computeFreqs =
     function(wordsList, spam, bow = unique(unlist(wordsList)))
      {
           # create a matrix for spam, ham, and log odds
           wordTable = matrix(0.5, nrow = 4, ncol = length(bow),
                                      dimnames = list(c("spam", "ham",
                                                               "presentLogOdds",
                                                               "absentLogOdds"), bow))
           # For each spam message, add 1 to counts for words in message
           counts.spam = table(unlist(lapply(wordsList[spam], unique)))
           wordTable["spam", names(counts.spam)] = counts.spam + .5
           # Similarly for ham messages
           counts.ham = table(unlist(lapply(wordsList[!spam], unique)))
           wordTable["ham", names(counts.ham)] = counts.ham + .5
```

```
# Find the total number of spam and ham
numSpam = sum(spam)
numHam = length(spam) - numSpam

# Prob(word|spam) and Prob(word | ham)
wordTable["spam", ] = wordTable["spam", ]/(numSpam + .5)
wordTable["ham", ] = wordTable["ham", ]/(numHam + .5)

# log odds
wordTable["presentLogOdds", ] =
    log(wordTable["spam",]) - log(wordTable["ham", ])
wordTable["absentLogOdds", ] =
    log((1 - wordTable["spam", ])) - log((1 -wordTable["ham", ]))
invisible(wordTable)
}
```

trainTable = computeFreqs(trainMsgWords, trainIsSpam)

The trainTable can be used to compute the log likelihood ratio for a new message. This value is used to classify the message as spam or ham. Besides, we apply function computeMsgLLR to each of the messages in our test set

```
computeMsgLLR = function(words, freqTable)
{
    # Discards words not in training data.
    words = words[!is.na(match(words, colnames(freqTable)))]

# Find which words are present
    present = colnames(freqTable) %in% words

sum(freqTable["presentLogOdds", present]) +
        sum(freqTable["absentLogOdds", !present])
}
```

To compare the summary statistics of the LLR values for the ham and spam in the test data with

```
tapply(testLLR, testIsSpam, summary)
$`FALSE`
    Min.
                     Median
                                                    Max.
          1st Qu.
                                 Mean
                                       3rd Qu.
         -132.50
-1367.00
                   -104.60
                             -120.50
                                                  697.80
$`TRUE`
     Min.
            1st Qu.
                        Median
                                     Mean
                                            3rd Qu.
                                                          Max.
  -63.650
                        56.330
                                            138.300 23600.000
              9.347
                                  142.500
```

and the boxplots for these data:

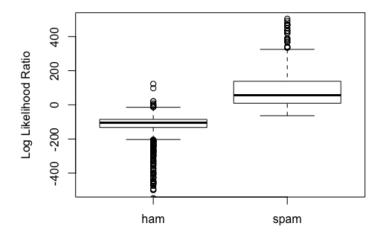


Figure 2: Boxplot of log likelyhood ratio for spam and ham

which shows that most ham data have log likelihood ratio below zero and spam data above zero. So naive Bayes approximation is a good way to group spam or ham messages. And we need to decide a cut-off τ which used to judge a message is spam or not.

Before that, define functions to calculate type I and type II errors. Then draw the plot to display these two errors in different τ .

```
typeIErrorRates =
    function(llrVals, isSpam)
    {
        o = order(llrVals)
        llrVals = llrVals[o]
        isSpam = isSpam[o]
        idx = which(!isSpam)
        N = length(idx)
        list(error = (N:1)/N, values = llrVals[idx])
    }
typeIIErrorRates = function(llrVals, isSpam) {
    o = order(llrVals)
    llrVals = llrVals[o]
    isSpam = isSpam[o]
    idx = which(isSpam)
    N = length(idx)
    list(error = (1:(N))/N, values = llrVals[idx])
}
```

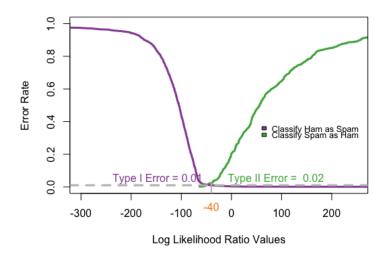


Figure 3: Type I and II Error Rates

with a threshold of $\tau = -40$, messages with an LLR value above -40 will be classified as spam messages while others as ham. Since in this case, type I error is 0.01 and type II error is 0.023, 1% of ham is misclassified as spam and 2.3% of spam is misclassified as ham.

4. Result by removing the stop words

Stop words also can be found in "tm" package and they are easily extracted as a vector.

```
library(tm)
stopwords <- stopwords()</pre>
```

Then add a function after processAllWords to remove stop words as:

```
removeStopWords =
  function(x, stopWords)
  {
     stopWords <- stopwords()
     if(is.character(x))
        setdiff(x, stopWords)
     else if(is.list(x))
        lapply(x, removeStopWords, stopWords)
     else x
}</pre>
```

msgWordsList = lapply(msgWordsList, removeStopWords, stopWords)

Other steps are similar. Then we have the result under the condition that the stop words are removed. The summary statistics of the LLR values for the ham and spam in the test data are

```
> tapply(testLLR, testIsSpam, summary)
$`FALSE`
                                  Mean
    Min.
           1st Qu.
                      Median
                                        3rd Qu.
                                                      Max.
-1363.00
           -127.90
                     -101.60
                              -116.90
                                          -81.32
                                                   700.00
$`TRUE`
    Min.
           1st Qu.
                      Median
                                  Mean
                                        3rd Qu.
                                                      Max.
  -60.33
                                          131.70 23600.00
              7.15
                               138.50
                       50.01
```

And the boxplot:

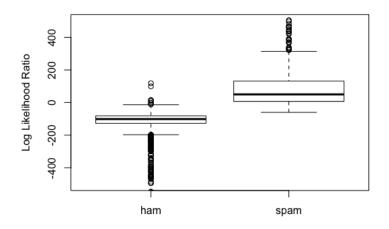


Figure 4: Boxplot of log likelyhood ratio for spam and ham when stop words are removed

Type I and type II Error Rates:

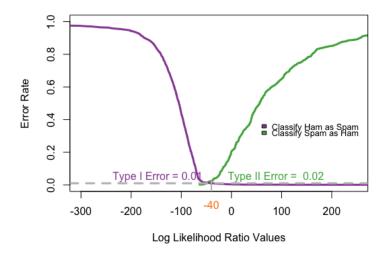


Figure 5: Type I and II Error Rates when stop words are removed

That is just a little different with the situation that remains stop words. Setting $\tau = -43$, messages with an LLR value above -43 will be classified as spam messages while others as ham. Since in this case, type I error is 0.01 and type II error is 0.02, 1% of ham is misclassified as spam and 2% of spam is misclassified as ham.

Problem 3

1. Get and Read the Data

Firstly, download the historical stock data from Yahoo Finance and save these contents to local files.

Then read the data and make sure that the date values remain as strings and not interpreted as a factor but converted into date format.

```
for(fmt in dateFormat) {
        tmp = as.Date(data$Date, fmt)
        if(all(!is.na(tmp))) {
            data$Date = tmp
            break
        }
    }
    data[ order(data$Date), ]
}

pep <- readData("PEP.csv")
ko <- readData("KO.csv")
dps <- readData("DPS.csv")</pre>
```

2. Proceed the Data

For now we use stock "PEP" and "KO" to do the pairs trading.

Define a function to compute the subsets with common dates and return a data frame with records for each day with the adjusted closing prices for the two stocks. After combining stock "PEP" and "KO", we can create the ratio of the adjusted closing price.

```
combine2Stocks =
    function(a, b, stockNames = c(deparse(substitute(a)),
                                   deparse(substitute(b))))
    {
        rr = intersect(a$Date, b$Date)
        a.sub=a[which(a$Date %in% rr),]
        b.sub=b[which(b$Date %in% rr),]
        structure(data.frame(as.Date(a.sub$Date),
                             a.sub$Adj.Close,
                             b.sub$Adj.Close),
                  names = c("Date", stockNames))
overlap = combine2Stocks(pep, ko)
r = overlap$pep/overlap$ko
Visualize the ratio (k = 1):
plotRatio =
    function(r, k = 1, date = seq(along = r), ...)
        plot(date, r, type = "l", ...)
        abline(h = c(mean(r),
                     mean(r) + k * sd(r),
                     mean(r) - k * sd(r)),
               col = c("darkgreen", rep("red", 2*length(k))),
               lty = "dashed")
    }
plotRatio(r, k = 1, overlap$Date, col = "lightgray", xlab = "Date", ylab = "Ratio")
length(r)
```

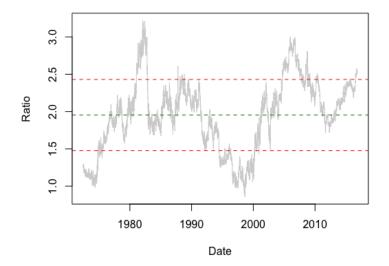


Figure 6: The ratio of stock prices for PEP and KO

3. Finding positions

In our case, k = 1, the upper bound is m + ks and the lower bound is m - ks. Compute two opening positions (green dots) and two closing positions (red dots). The plot was drawn below.

```
findNextPosition =
    # Check they are increasing and correctly offset
    function(ratio, startDay = 1, k = 1,
             m = mean(ratio), s = sd(ratio))
    {
        up = m + k *s
        down = m - k *s
        if(startDay > 1)
            ratio = ratio[ - (1:(startDay-1)) ]
        isExtreme = ratio >= up | ratio <= down
        if(!any(isExtreme))
            return(integer())
        start = which(isExtreme)[1]
        backToNormal = if(ratio[start] > up)
            ratio[ - (1:start) ] <= m
        else
            ratio[ - (1:start) ] >= m
        # return either the end of the position or the index
        # of the end of the vector.
        # Could return NA for not ended, i.e. which(backToNormal)[1]
        # for both cases. But then the caller has to interpret that.
```

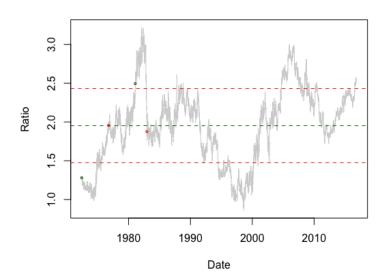


Figure 7: Position finding function test

In order to find all positions, define the function "getPositions":

```
getPositions =
  function(ratio, k = 1, m = mean(ratio), s = sd(ratio))
{
    when = list()
    cur = 1

  while(cur < length(ratio)) {
      tmp = findNextPosition(ratio, cur, k, m, s)
      if(length(tmp) == 0) # done
            break</pre>
```

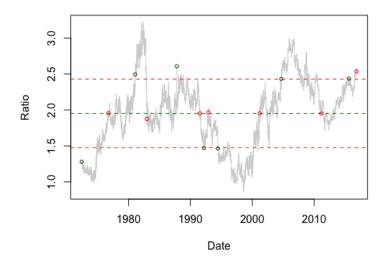


Figure 8: All positions when k=1

Set k = 0.5, the plot is showed below. There are much more positions than that in k = 1.

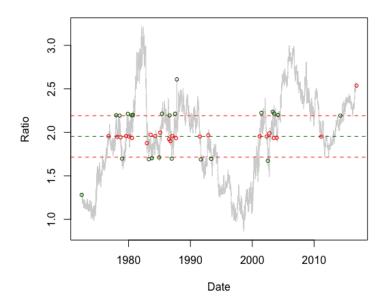


Figure 9: All positions when k=0.5

More positions existing means that we may earn more but cost more in transaction fees. To pursue the maximum earnings, we should find appropriate value k based on the historical data for our pair of stocks.

4. Computing the Profit and Find the Optimal Value for k

Calculate the position profits:

```
positionProfit =
    # r = overlap$pep/overlap$ko
    # pos = getPositions(r, k)
    # positionProfit(pos[[1]], overlap$pep, overlap$ko)
    function(pos, stockPriceA, stockPriceB,
             ratioMean = mean(stockPriceA/stockPriceB),
             p = .0002, byStock = FALSE)
    {
        if(is.list(pos)) {
            ans = sapply(pos, positionProfit,
                         stockPriceA, stockPriceB, ratioMean, p, byStock)
            if(byStock)
                rownames(ans) = c("A", "B", "commission")
            return(ans)
        # prices at the start and end of the positions
        priceA = stockPriceA[pos]
        priceB = stockPriceB[pos]
        # how many units can we by of A and B with $1
        unitsOfA = 1/priceA[1]
```

```
unitsOfB = 1/priceB[1]
        # The dollar amount of how many units we would buy of A and B
        # at the cost at the end of the position of each.
        amt = c(unitsOfA * priceA[2], unitsOfB * priceB[2])
        # Which stock are we selling
        sellWhat = if(priceA[1]/priceB[1] > ratioMean) "A" else "B"
        profit = if(sellWhat == "A")
            c((1 - amt[1]), (amt[2] - 1), -p * sum(amt))
        else
            c((1 - amt[2]), (amt[1] - 1), - p * sum(amt))
        if (byStock)
            profit
        else
            sum(profit)
    }
When calculating for PEP/KO ratios, the result is:
[1] 0.37333954 0.42307159 0.99590876 0.34030945 0.84320973 0.35930336 -0.04535196
which seems reasonable.
Then to find the optimal value for k, divide the data into a training period and a test period.
we just use the training period to determine optimal value.
i = 1:floor(nrow(overlap)/2)
train = overlap[i, ]
test = overlap[ - i, ]
# Divide the data half to a training period and half to a test period
r.train = train$pep/train$ko
r.test = test$pep/test$ko
k.max = max((r.train - mean(r.train))/sd(r.train))
k.min = min((abs(r.train - mean(r.train))/sd(r.train)))
ks = seq(k.min, k.max, length = 1000)
m = mean(r.train)
profits =
    sapply(ks,
           function(k) {
               pos = getPositions(r.train, k)
               sum(positionProfit(pos, train$pep, train$ko,
                                  mean(r.train)))
           })
plot(ks, profits, type = "l", xlab = "k", ylab = "Profit")
tmp.k = ks[profits == max(profits)]
# [1] 0.03661572
pos = getPositions(r.test, tmp.k, mean(r.train), sd(r.train))
testProfit = sum(positionProfit(pos, test$pep, test$ko))
testProfit
# [1] 1.548414
```

Loop over 1000 values of k and compute the 1000 corresponding profits. We can see the relationship between k and profits as below

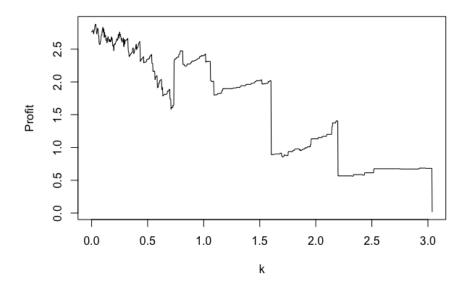


Figure 10: Relationship between profits and k

The profit reach its maximum 1.548414 (about 155%) at k = 0.03661572.

5. Pairs trading on PEP and DPS

All steps are similar to former trading.

```
# PEP vs DPS
overlap_pd <- combine2Stocks(pep, dps)</pre>
r_pd <- overlap_pd$ko/overlap_pd$dps</pre>
i = 1:floor(nrow(overlap_pd)/2)
train = overlap_pd[i, ]
test = overlap_pd[ - i, ]
# Divide the data half to a training period and half to a test period
r.train = train$pep/train$dps
r.test = test$pep/test$dps
sd(r.train)
k.max = min(max((r.train - mean(r.train))/sd(r.train)),
            (mean(r.train)-min(r.train))/sd(r.train))
#k.max = max((r.train - mean(r.train))/sd(r.train))
k.min = max(0.1, min((abs(r.train - mean(r.train))/sd(r.train))))
ks = seq(k.min, k.max, length = 1000)
  = mean(r.train)
profits =
    sapply(ks,
```

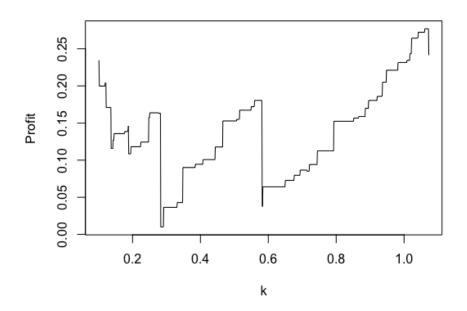


Figure 11: Relationship between profits and k in pairs trading PEP vs DPS

The profit reach its maximum 0.3586618 (about 35.9%) at k = 1.066602.



6. Pairs trading on KO and DPS

```
# KO vs DPS
overlap_kd <- combine2Stocks(ko, dps)</pre>
r_kd <- overlap_kd$ko/overlap_kd$dps
i = 1:floor(nrow(overlap_kd)/2)
train = overlap_kd[i, ]
test = overlap_kd[ - i, ]
# Divide the data half to a training period and and half to a test period
r.train = train$ko/train$dps
r.test = test$ko/test$dps
k.max = max((r.train - mean(r.train))/sd(r.train))
k.min = max(0.1, min((abs(r.train - mean(r.train))/sd(r.train))))
ks = seq(k.min, k.max, length = 1000)
m = mean(r.train)
profits =
    sapply(ks,
           function(k) {
               pos = getPositions(r.train, k)
               sum(positionProfit(pos, train$ko, train$dps,
                                  mean(r.train)))
           })
plot(ks, profits, type = "l", xlab = "k", ylab = "Profit")
tmp.k = ks[profits == max(profits)]
# [1] 0.4365644 0.4402227
```

```
k.star = mean(ks[profits == max(profits)])
# [1] 0.4383935

pos = getPositions(r.test, k.star)
testProfit = sum(positionProfit(pos, test$ko, test$dps))
testProfit
```

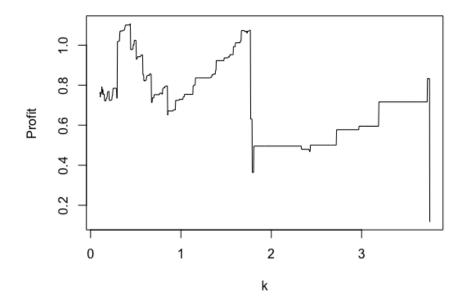


Figure 12: Relationship between profits and k in pairs trading KO vs DPS

The profit reach its maximum 0.1493933 (about 14.9%) at k=0.4383935.

