

CSE – 4020 – Machine Learning

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Slot: L59 + L60

Lab Assignment – 5

1. Implement kNN(k-nearest neighbors) in R for classification(Consider binary class of predictors of any data sets of your choice)

Soln.

```
> # Read data
> # load library
> library(caret)
> library(e1071)
> # Transforming the dependent variable to a factor
> data1$win.Loss = as.factor(data1$win.Loss)
> #Partitioning the data into training and validation data
> set.seed(101)
> index = createDataPartition(data1$win.Loss, p = 0.7, list = F )
> train = data1[index,]
> validation = data1[-index,]
> # Explore data
> dim(train)
[1] 1068  14
> dim(validation)
[1] 456  14
> names(train)
[1] "win.Loss"      "Optimism"      "Pessimism"     "PastUsed"      "FutureUsed"
"PresentUsed"    "OwnPartyCount" "OppPartyCount"
[9] "NumericContent" "Extra"         "Emoti"         "Agree"         "Consc"
"Openn"
> head(train)
  win.Loss Optimism Pessimism PastUsed FutureUsed PresentUsed OwnPartyCount
OppPartyCount NumericContent Extra Emoti Agree Consc Openn
1          1 0.10450450 0.05045045 0.4381443  0.4948454  0.06701031          2
2          0 0.001877543 4.041 4.049 3.469 2.450 2.548
3          1 0.11257190 0.04930156 0.4159664  0.5168067  0.06722689          1
1          0 0.002131163 3.463 4.039 3.284 2.159 2.465
5          1 0.10582640 0.05172414 0.3342618  0.5821727  0.08356546          3
4          0 0.002229220 4.658 4.023 3.283 2.415 2.836
7          1 0.09838275 0.06401617 0.3240741  0.6018519  0.07407407          6
4          0 0.002251985 3.727 4.108 3.357 2.128 2.231
9          1 0.10610734 0.04688464 0.3633540  0.5372671  0.09937888          2
5          0 0.002446440 4.119 4.396 3.661 2.572 2.599
10         1 0.10066128 0.05951506 0.3554817  0.5382060  0.10631229          1
2          0 0.002107436 3.800 4.501 3.624 2.117 2.154
```

```

> head(validation)
  Win.Loss Optimism Pessimism PastUsed FutureUsed PresentUsed OwnPartyCount
OppPartyCount NumericContent Extra Emoti Agree Consc Openn
2      1 0.11457521 0.05923617 0.2912621 0.6213592 0.08737864      1
4      0 0.001418909 3.446 3.633 3.528 2.402 2.831
4      1 0.10723350 0.04631980 0.4634921 0.4666667 0.06984127      1
3      0 0.001871715 4.195 4.661 4.007 2.801 3.067
6      1 0.07586207 0.03448276 0.2800000 0.5200000 0.20000000      0
0      0 0.003290827 2.843 3.563 3.075 1.769 1.479
8      1 0.10377924 0.05638872 0.3692722 0.5498652 0.08086253      2
4      0 0.002215028 4.027 4.631 3.920 2.417 2.291
17     1 0.11289199 0.05505227 0.3891051 0.5214008 0.08949416      2
7      0 0.001165647 4.086 4.173 3.368 2.348 2.412
21     1 0.11466373 0.03858875 0.2736842 0.6210526 0.10526316      1
7      0 0.003105161 3.770 3.858 2.874 1.949 2.006
> # Setting levels for both training and validation data
> levels(train$Win.Loss) <- make.names(levels(factor(train$Win.Loss)))
> levels(validation$Win.Loss) <- make.names(levels(factor(validation$Win.Loss)))
> # Setting levels for both training and validation data
> levels(train$Win.Loss) <- make.names(levels(factor(train$Win.Loss)))
> levels(validation$Win.Loss) <- make.names(levels(factor(validation$Win.Loss)))
> # Setting up train controls
> repeats = 3
> numbers = 10
> tune1 = 10
> set.seed(1234)
> x = trainControl(method = "repeatedcv",
+                   number = numbers,
+                   repeats = repeats,
+                   classProbs = TRUE,
+                   summaryFunction = twoClassSummary)
> model1 <- train(Win.Loss~. , data = train, method = "knn",
+                 preProcess = c("center","scale"),
+                 trControl = x,
+                 metric = "ROC",
+                 tuneLength = tune1)
> # Summary of model
> model1
k-Nearest Neighbors

1068 samples
 13 predictor
 2 classes: 'x0', 'x1'

```

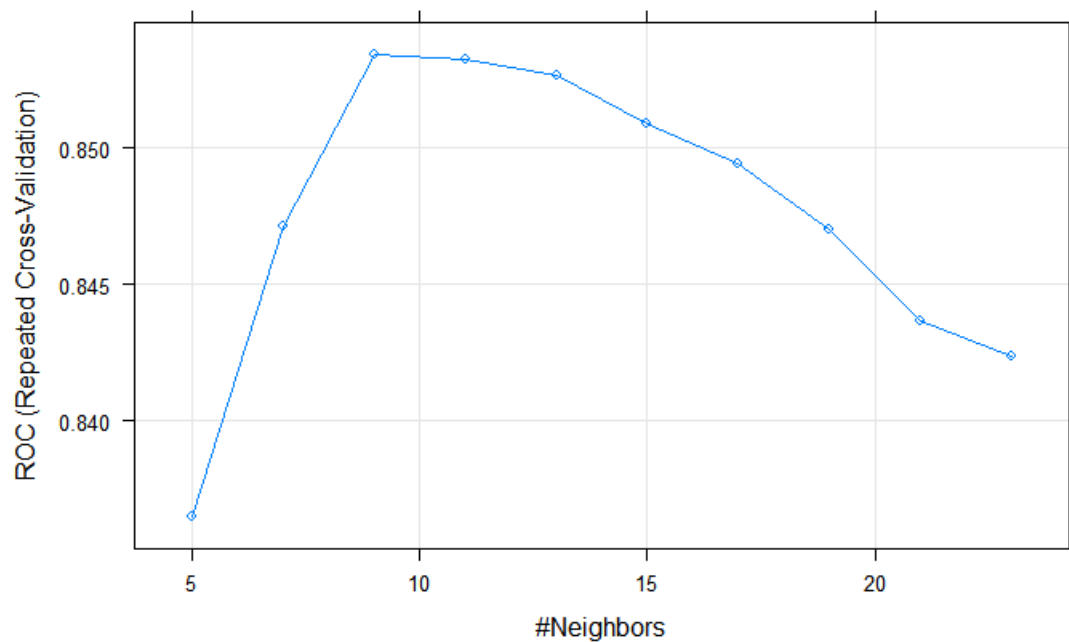
Pre-processing: centered (13), scaled (13)
 Resampling: Cross-Validated (10 fold, repeated 3 times)
 Summary of sample sizes: 962, 961, 961, 961, 961, 961, ...
 Resampling results across tuning parameters:

k	ROC	Sens	Spec
5	0.8364872	0.6900890	0.8412665
7	0.8471507	0.6684475	0.8494250
9	0.8534144	0.6587689	0.8525019
11	0.8532324	0.6540457	0.8602020
13	0.8526851	0.6531940	0.8683994
15	0.8509041	0.6491096	0.8607071
17	0.8494333	0.6411537	0.8560995
19	0.8470142	0.6267325	0.8612432
21	0.8436754	0.6146922	0.8637529
23	0.8423458	0.6042973	0.8714375

ROC was used to select the optimal model using the largest value.

The final value used for the model was $k = 9$.

```
> plot(model1)
```



```
> # Validation
> valid_pred <- predict(model1, validation, type = "prob")
> # Storing Model Performance Scores
> library(ROCR)
> pred_val <- prediction(valid_pred[,2], validation$win.Loss)
> # Calculating Area under Curve (AUC)
> perf_val <- performance(pred_val, "auc")
> perf_val
```

An object of class "performance"

Slot "x.name":

[1] "None"

Slot "y.name":

[1] "Area under the ROC curve"

Slot "alpha.name":

[1] "none"

Slot "x.values":

list()

Slot "y.values":

[[1]]

[1] 0.8670378

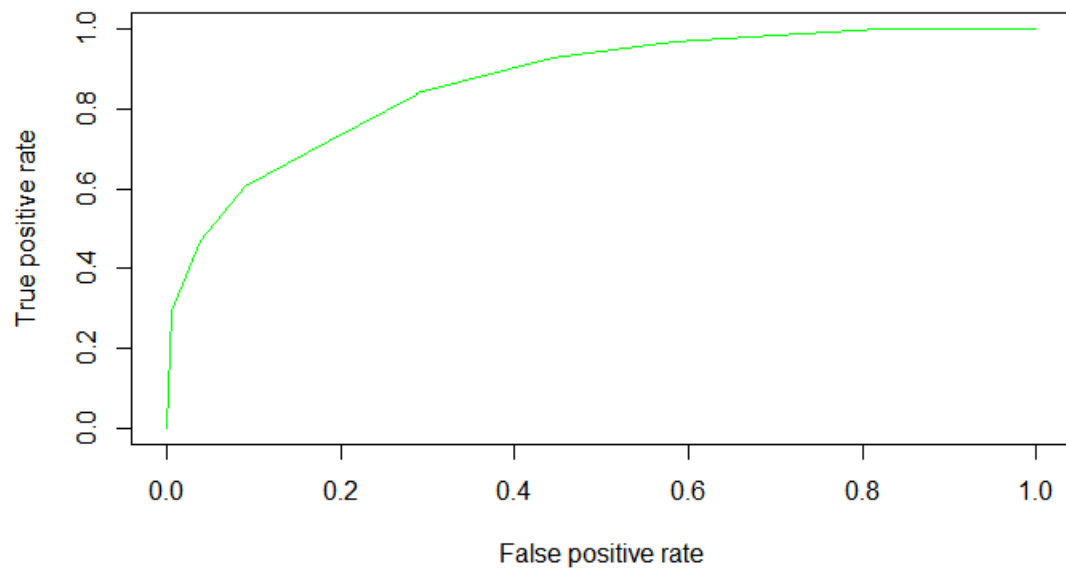
Slot "alpha.values":

list()

```
> # Plot AUC
```

```
> perf_val <- performance(pred_val, "tpr", "fpr")
```

```
> plot(perf_val, col = "green", lwd = 1.5)
```



3. Implement K-means clustering? (Consider any clustering data set from internet except iris data)

Soln:

Dataset: Wholesale Customers - UCI

```
> summary(data)
  Channel      Region      Fresh      Milk      Grocery      Frozen
Detergents_Paper Delicassen
Min.   :1.000  Min.   :1.000  Min.   :    3  Min.   :   55  Min.   :    3  Min.   :
25.0  Min.   :    3.0  Min.   :    3.0
1st Qu.:1.000  1st Qu.:2.000  1st Qu.: 3128  1st Qu.: 1533  1st Qu.: 2153  1st Qu.:
742.2  1st Qu.: 256.8  1st Qu.: 408.2
Median :1.000  Median :3.000  Median : 8504  Median : 3627  Median : 4756  Median :
1526.0  Median : 816.5  Median : 965.5
Mean   :1.323  Mean   :2.543  Mean   : 12000  Mean   : 5796  Mean   : 7951  Mean   :
3071.9  Mean   : 2881.5  Mean   : 1524.9
3rd Qu.:2.000  3rd Qu.:3.000  3rd Qu.: 16934  3rd Qu.: 7190  3rd Qu.:10656  3rd Qu.:
3554.2  3rd Qu.: 3922.0  3rd Qu.: 1820.2
Max.   :2.000  Max.   :3.000  Max.   :112151  Max.   :73498  Max.   :92780  Max.   :
60869.0  Max.   :40827.0  Max.   :47943.0

> top.n.custs <- function (data,cols,n=5) { #Requires some data frame and the top N to remove
+   idx.to.remove <-integer(0) #Initialize a vector to hold customers being removed
+   for (c in cols){ # For every column in the data we passed to this function
+     col.order <-order(data[,c],decreasing=T) #Sort column "c" in descending order (bigger
on top)
+     #Order returns the sorted index (e.g. row 15, 3, 7, 1, ...) rather than the actual
values sorted.
+     idx <-head(col.order, n) #Take the first n of the sorted column C to
+     idx.to.remove <-union(idx.to.remove,idx) #Combine and de-duplicate the row ids that
need to be removed
+   }
}
```

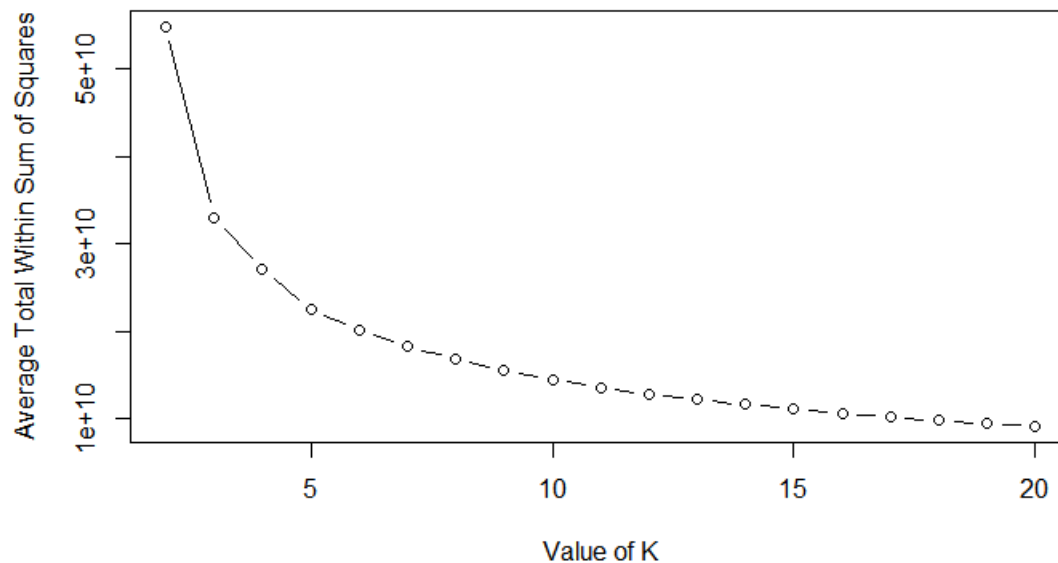
```

+   return(idx.to.remove) #Return the indexes of customers to be removed
+ }
> top.custs <-top.n.custs(data,cols=3:8,n=5)
> length(top.custs) #How Many Customers to be Removed?
[1] 19
> data[top.custs,] #Examine the customers
  Channel Region Fresh Milk Grocery Frozen Detergents_Paper Delicassen
182      1      3 112151 29627  18148  16745           4948      8550
126      1      3  76237  3473   7102  16538           778       918
285      1      3  68951  4411  12609  8692           751      2406
40       1      3  56159   555    902  10002           212      2916
259      1      1  56083  4563   2124  6422           730      3321
87       2      3  22925 73498  32114   987          20070       903
48       2      3  44466 54259  55571  7782          24171      6465
86       2      3  16117 46197  92780  1026          40827      2944
184      1      3  36847 43950  20170 36534           239     47943
62       2      3  35942 38369  59598  3254          26701      2017
334      2      2   8565  4980   67298   131          38102      1215
66       2      3     85 20959  45828    36          24231      1423
326      1      2  32717 16784  13626 60869          1272      5609
94       1      3  11314  3090   2062 35009           71      2698
197      1      1  30624  7209   4897 18711           763      2876
104      1      3  56082  3504   8906 18028          1480      2498
24       2      3  26373 36423  22019  5154          4337     16523
72       1      3  18291  1266   21042  5373          4173     14472
88       1      3  43265  5025   8117  6312          1579     14351
> data.rm.top <-data[-c(top.custs),] #Remove the Customers
> set.seed(76964057) #Set the seed for reproducibility
> k <-kmeans(data.rm.top[, -c(1,2)], centers=5) #Create 5 clusters, Remove columns 1 and 2
> k$centers #Display cluster centers
  Fresh      Milk  Grocery  Frozen Detergents_Paper Delicassen
1 4189.747 7645.639 11015.277 1335.145      4750.4819 1387.1205
2 16470.870 3026.491  4264.741 3217.306      996.5556 1319.7593
3 33120.163 4896.977  5579.860 3823.372      945.4651 1620.1860
4  5830.214 15295.048 23449.167 1936.452     10361.6429 1912.7381
5  5043.434  2329.683  2786.138 2689.814       652.8276  849.8414
> table(k$cluster) #Give a count of data points in each cluster

 1    2    3    4    5
83 108  43  42 145
> rng<-2:20 #K from 2 to 20
> tries<-100 #Run the K Means algorithm 100 times
> avg.totw.ss<-integer(length(rng)) #Set up an empty vector to hold all of points
> for(v in rng){ # For each value of the range variable
+   v.totw.ss<-integer(tries) #Set up an empty vector to hold the 100 tries
+   for(i in 1:tries){
+     k.temp<-kmeans(data.rm.top,centers=v) #Run kmeans
+     v.totw.ss[i]<-k.temp$tot.withinss#Store the total withinss
+   }
+   avg.totw.ss[v-1]<-mean(v.totw.ss) #Average the 100 total withinss
+ }
> plot(rng,avg.totw.ss,type="b", main="Total within SS by Various K",
+       ylab="Average Total within Sum of Squares",
+       xlab="Value of K")

```

Total Within SS by Various K



4. Spam classification using any Ensemble classifier? Find AUC, ROC, Confusion Matrix, and accuracy?
 Data set link : <https://archive.ics.uci.edu/ml/machine-learning-databases/spambase/spambase.data>
 The last attribute is the predictor

Soln:

```
> names <- read.csv("C:/Users/chait/Desktop/VIT/Machine_Learning/Lab/Assesment - 5/names.csv",header=FALSE,sep=";")
> names(dataset) <- sapply((1:nrow(names)),function(i) toString(names[i,1]))
> dataset$y <- as.factor(dataset$y)
> sample <- dataset[sample(nrow(dataset), 300),]
> smp_size = floor(nrow(sample)*0.7)
> ind = sample(seq_len(nrow(sample)),size = smp_size)
> df_train = sample[ind,]
> df_test = sample[-ind,]
> resample.spam.train <- function()
+ {
+   indices <- sample(1:nrow(df_train),nrow(df_train),replace=TRUE)
+   df <- df_train[indices,]
+   return(df)
+ }
> bag.trees <- function(B) # B is the number of bootstrap samples
+ {
+   bootstrap.samples <- list()
+   pred.mat <- matrix(NA, nrow = nrow(df_test), ncol = B)
+   for(i in 1:B)
+   {
+     spam.sample <- resample.spam.train() # gets a bootstrap sample
+     tree <- rpart(y ~ .,
+                   data = spam.sample) # fits a tree
+     pruned.tree <- prune(tree,
+                           cp= tree$cptable[which.min(tree$cptable[, "xerror"]),
+                           "cp"]) #prunes tree
```

```

+
+   # predictions
+   pred.mat[,i] <- predict(pruned.tree,
+                           newdata = df_test[, -ncol(df_test)],
+                           type = "class")
+
+   pred.mat[,i] <- pred.mat[,i] - 1 # convert to (0/1)
+ }
+ return(pred.mat)
+ }
> library(rpart)
> set.seed(11)
> bag <- bag.trees(50)
> # the 0/1 output is the one that the majority of the trees select
> bagged.preds <- (rowMeans(bag) >= 0.5)+0
> # final classification accuracy
> round(mean(bagged.preds == df_test$spam),2)
0.8

```

5. Implement polynomial regression and find all the necessary errors (Take any regression data from UCI machine learning repository) (if possible, in MS EXCEL; R2 and RMSE are expected to be calculated as I have demonstrated in both F1 and F2 slot classes)

Soln:

```

> indexes = sample(1:nrow(df), size=0.2*nrow(df))
> train = df[-indexes,]
> test = df[indexes,]
> model = lm(formula = ERP ~ MYCT+MMIN+MMAX+CACH+CHMIN+CHMAX+PRP,
data=train)
> confint(model)

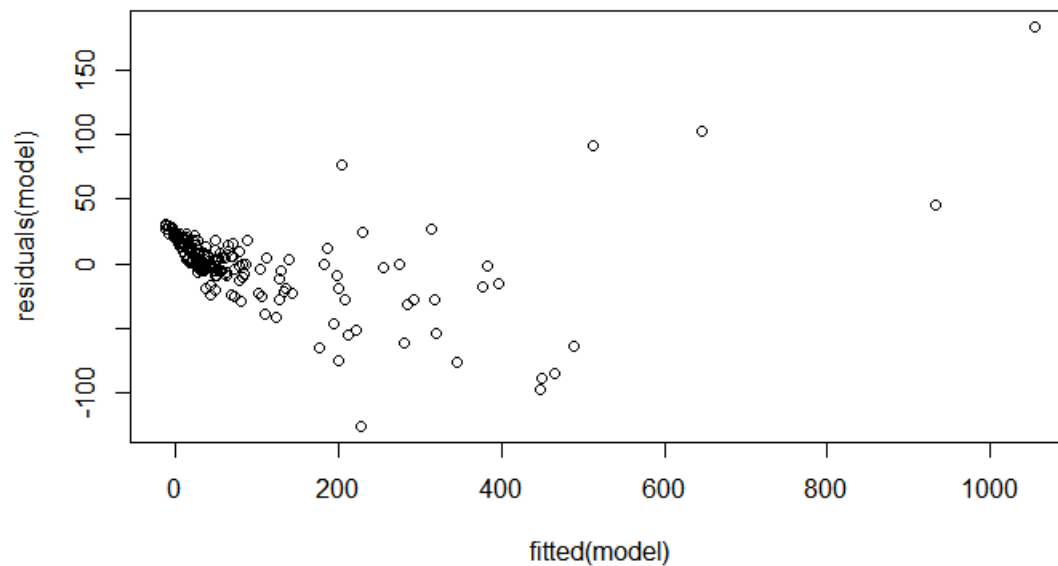
```

	2.5 %	97.5 %
(Intercept)	-42.403438945	-21.223996296
MYCT	0.015685479	0.056745731
MMIN	0.003530103	0.008609015
MMAX	0.002112134	0.004028666
CACH	0.023046146	0.353958726
CHMIN	-0.705511161	1.310470281
CHMAX	-0.067176460	0.513565018
PRP	0.481381893	0.650260325

```

> plot(fitted(model),residuals(model))

```



```
> predicted.intervals = predict(model,df,interval='confidence',level=0.99)
> actual = df[,"ERP"]
> predicted = predicted.intervals[, "fit"]
> rss = sum((actual-predicted)^2)
> tss = sum((actual-mean(actual))^2)
> rsq = 1 - rss/tss
> rmse = (rss/nrow(df))^(.5)
> cat("RMSE value is",rmse)
RMSE value is 31.44807
> cat("R-squared value is",rsq)
R-squared value is 0.9585075
```

6. Implement PCA with high dimension data set.

7. Hierarchical clustering with any data set of your choice.

Soln of 6 & 7:

```
> library(dplyr)
> library(tibble)
> library(ggplot2)
> library(readr)
> proteoms <- read.csv("C:/Users/chaith/Desktop/VIT/Machine_Learning/Lab/Assesment - 5/data.csv")
> colnames(proteoms)
[1] "RefSeq_accession_number" "gene_symbol" "gene_name"
"AO.A12D.01TCGA" "C8.A131.01TCGA" "C8.A130.02TCGA"
[6] "AO.A12B.01TCGA" "BH.A18Q.02TCGA" "C8.A138.03TCGA"
"C8.A138.03TCGA" "E2.A154.03TCGA" "C8.A12L.04TCGA"
[11] "C8.A12L.04TCGA" "A2.A0EX.04TCGA" "AN.A04A.05TCGA"
"AN.A04A.05TCGA" "BH.A0AV.05TCGA" "C8.A12T.06TCGA"
[16] "C8.A12T.06TCGA" "A8.A06Z.07TCGA" "BH.A18U.08TCGA"
"BH.A18U.08TCGA" "A2.A0EQ.08TCGA"
```


[21] "AR.A0U4.09TCGA"	"AO.A0J9.10TCGA"	"AR.A1AP.11TCGA"
"AN.A0FK.11TCGA"	"AO.A0J6.11TCGA"	
[26] "A7.A13F.12TCGA"	"BH.A0E1.12TCGA"	"A7.A0CE.13TCGA"
"A2.A0YC.13TCGA"	"AO.A0JC.14TCGA"	
[31] "A8.A08Z.14TCGA"	"AR.A0TX.14TCGA"	"A8.A076.15TCGA"
"AO.A126.15TCGA"	"BH.A0C1.16TCGA"	
[36] "A2.A0EY.16TCGA"	"AR.A1AW.17TCGA"	"AR.A1AV.17TCGA"
"C8.A135.17TCGA"	"A2.A0EV.18TCGA"	
[41] "AN.A0AM.18TCGA"	"D8.A142.18TCGA"	"AN.A0FL.19TCGA"
"BH.A0DG.19TCGA"	"AR.A0TV.20TCGA"	
[46] "C8.A12Z.20TCGA"	"AO.A0JJ.20TCGA"	"AO.A0JE.21TCGA"
"AN.A0AJ.21TCGA"	"A7.A0CJ.22TCGA"	
[51] "AO.A12F.22TCGA"	"A8.A079.23TCGA"	"A2.A0T3.24TCGA"
"A2.A0YD.24TCGA"	"AR.A0TR.25TCGA"	
[56] "AO.A030.25TCGA"	"AO.A12E.26TCGA"	"A8.A06N.26TCGA"
"A2.A0YG.27TCGA"	"BH.A18N.27TCGA"	
[61] "AN.A0AL.28TCGA"	"A2.A0T6.29TCGA"	"E2.A158.29TCGA"
"E2.A15A.29TCGA"	"AO.A0JM.30TCGA"	
[66] "C8.A12V.30TCGA"	"A2.A0D2.31TCGA"	"C8.A12U.31TCGA"
"AR.A1AS.31TCGA"	"A8.A09G.32TCGA"	
[71] "C8.A131.32TCGA"	"C8.A134.32TCGA"	"A2.A0YF.33TCGA"
"BH.A0DD.33TCGA"	"BH.A0E9.33TCGA"	
[76] "AR.A0TT.34TCGA"	"AO.A12B.34TCGA"	"A2.A0SW.35TCGA"
"AO.A0JL.35TCGA"	"BH.A0BV.35TCGA"	
[81] "A2.A0YM.36TCGA"	"BH.A0C7.36TCGA"	"A2.A0SX.36TCGA"
"X263d3f.I.CPTAC"	"b1cdb9.I.CPTAC"	
[86] "c4155b.C.CPTAC"		

```
> clean.proteoms <- na.omit(proteoms)
```

```
> head(as.matrix(clean.proteoms[,4:length(colnames(clean.proteoms))]))
```

```
AO.A12D.01TCGA C8.A131.01TCGA AO.A12B.01TCGA BH.A18Q.02TCGA C8.A130.02TCGA C8.A138.03TCGA
E2.A154.03TCGA C8.A12L.04TCGA A2.A0EX.04TCGA
```

1	1.0961312	2.60994298	-0.6598280	0.1953407	-0.4940596	2.7650807
0.8626593	1.407570262	1.185108				
3	1.1113704	2.65042179	-0.6542851	0.2154129	-0.5006193	2.7797092
0.8701860	1.410311827	1.188860				
6	1.1075606	2.64637391	-0.6542851	0.2154129	-0.5038992	2.7797092
0.8701860	1.407570262	1.188860				
7	1.1113704	2.65042179	-0.6487422	0.2154129	-0.5006193	2.7833664
0.8701860	1.410311827	1.188860				
10	0.4827537	-1.04529350	1.2220027	-0.5172257	-0.4055031	0.7499970
2.3491966	-0.007077155	2.138081				
11	0.2617854	-0.03737115	1.0196851	-0.7246394	-0.7039712	-0.1569735
1.5814659	-0.023526544	1.732880				

```
AO.A12D.05TCGA AN.A04A.05TCGA BH.A0AV.05TCGA C8.A12T.06TCGA A8.A06Z.07TCGA A2.A0CM.07TCGA
BH.A18U.08TCGA A2.A0EQ.08TCGA AR.A0U4.09TCGA
```

1	1.1006881	0.38458773	0.35053566	-0.2049179	-0.4964091	0.6834035
-0.2650304	-0.9126703	-0.03322133				
3	1.1006881	0.37139283	0.36740533	-0.1666684	-0.4964091	0.6980976
-0.2516423	-0.9279787	-0.02721152				
6	1.0970232	0.37799028	0.36740533	-0.1666684	-0.4964091	0.6980976
-0.2516423	-0.9279787	-0.03021642				
7	1.0970232	0.37469156	0.36065746	-0.1666684	-0.4964091	0.6980976
-0.2516423	-0.9279787	-0.03021642				
10	0.5436299	0.05141659	0.40114465	0.4708229	1.3493193	-0.9843733
-0.6465901	-2.2789429	-2.10059548				
11	0.8404833	-1.62433531	-0.05433625	1.5163087	-1.9641727	-0.5472247
0.2336749	-2.2368449	-0.78745230				

```
AO.A0J9.10TCGA AR.A1AP.11TCGA AN.A0FK.11TCGA AO.A0J6.11TCGA A7.A13F.12TCGA BH.A0E1.12TCGA
A7.A0CE.13TCGA A2.A0YC.13TCGA AO.A0JC.14TCGA
```

1	0.02000705	0.4610875	0.9735642	0.8311317	1.2791847	0.7620444
-1.123173	0.8188241	-0.3072668				

3	0.01195532	0.4610875	0.9774761	0.8565398	1.2751671	0.7663844
-1.116861	0.8148772	-0.3072668				
6	0.01195532	0.4610875	0.9774761	0.8565398	1.2791847	0.7620444
-1.120017	0.8148772	-0.3072668				
7	0.01195532	0.4610875	0.9774761	0.8508936	1.2791847	0.7620444
-1.123173	0.8148772	-0.3072668				
10	0.32597284	-0.2836251	3.2620206	2.8383705	0.9617945	2.1855460
-2.575065	1.0516860	1.1702139				
11	-0.53153654	-0.4656659	2.4248758	3.4058177	0.2145214	1.8079709
-2.742348	0.8661858	0.9551376				
A8.A08Z.14TCGA AR.A0TX.14TCGA A8.A076.15TCGA AO.A126.15TCGA BH.A0C1.16TCGA A2.A0EY.16TCGA						
AR.A1AW.17TCGA AR.A1AV.17TCGA C8.A135.17TCGA						
1	0.5688946	-0.5834286	1.873982	0.1958767	-0.5183665	1.1748810
0.5783087	-0.7598231	1.120502				
3	0.5688946	-0.5671090	1.870383	0.1958767	-0.5072138	1.1832088
0.5783087	-0.7491137	1.137618				
6	0.5688946	-0.5779888	1.870383	0.1997197	-0.5072138	1.1832088
0.5783087	-0.7437590	1.127348				
7	0.5688946	-0.5779888	1.870383	0.1997197	-0.5100020	1.1832088
0.5822129	-0.7544684	1.137618				
10	0.8877477	1.6387646	1.377356	0.6531924	-2.1745373	0.8251135
-1.6978260	0.7261015	2.257001				
11	1.2173487	1.7883614	0.981495	0.6378204	-2.5342107	0.3087899
-1.7602928	-1.4130944	2.154306				
A2.A0EV.18TCGA AN.A0AM.18TCGA D8.A142.18TCGA AN.A0FL.19TCGA BH.A0DG.19TCGA AR.A0TV.20TCGA						
C8.A12Z.20TCGA AO.A0JJ.20TCGA AO.A0JE.21TCGA						
1	0.4529859	1.501967	0.5385958	2.455138	-0.2056375	-1.514278
-0.78719498	0.7571881	0.5597770				
3	0.4725901	1.501967	0.5422105	2.480137	-0.2056375	-1.528285
-0.75594056	0.7741042	0.5597770				
6	0.4725901	1.510348	0.5422105	2.471046	-0.2103218	-1.525484
-0.77156777	0.7774874	0.5597770				
7	0.4725901	1.506158	0.5422105	2.480137	-0.2056375	-1.525484
-0.77156777	0.7774874	0.5597770				
10	1.8112774	1.552255	0.2674902	-1.858279	0.2721667	-1.738392
0.02542003	1.9277820	1.7540156				
11	1.9569086	1.149947	0.1590480	-1.117407	-1.0253996	-2.231442
0.96305274	1.1428751	0.7303825				
AN.A0AJ.21TCGA A7.A0CJ.22TCGA AO.A12F.22TCGA A8.A079.23TCGA A2.A0T3.24TCGA A2.A0YD.24TCGA						
AR.A0TR.25TCGA AO.A03O.25TCGA AO.A12E.26TCGA						
1	-0.4281815	-1.0012398	-1.947792	1.048959	0.5837133	0.06377853
-1.1016752	1.053225	0.26485911				
3	-0.4063780	-1.0046198	-1.955180	1.052257	0.5806231	0.08446902
-1.1087826	1.055948	0.27571131				
6	-0.4063780	-1.0012398	-1.955180	1.052257	0.5868034	0.09333637
-1.1064135	1.055948	0.27299826				
7	-0.4063780	-1.0012398	-1.955180	1.052257	0.5868034	0.08446902
-1.1087826	1.058671	0.27571131				
10	0.9720842	0.5636905	-2.551133	3.436487	1.5014792	1.30520788
1.0755482	-1.250614	0.13191966				
11	0.9018286	1.2227864	-1.915778	3.489250	1.0534116	1.38501405
-0.0000333	-1.375882	0.07765866				
A8.A06N.26TCGA A2.A0YG.27TCGA BH.A18N.27TCGA AN.A0AL.28TCGA A2.A0T6.29TCGA E2.A158.29TCGA						
E2.A15A.29TCGA AO.A0JM.30TCGA C8.A12V.30TCGA						
1	0.2385471	-0.07820182	1.101261	0.3236627	0.7939756	-1.086529
2.1801233	1.395247	0.6739047				
3	0.2441826	-0.07143937	1.097767	0.3269726	0.8147235	-1.095492
2.1801233	1.412341	0.6887176				
6	0.2498182	-0.06805814	1.101261	0.3269726	0.8112655	-1.093252
2.1801233	1.412341	0.6887176				
7	0.2441826	-0.07143937	1.101261	0.3269726	0.8112655	-1.093252
2.1801233	1.412341	0.6887176				

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10      -0.2292004      -0.28783799      2.103995      -1.9105423      1.5651080      -1.149272
0.8962164      1.802077      -2.1331321
11      0.5344115      -1.56594237      1.146681      -1.2551755      1.5097801      -1.270277
0.8112191      1.655071      -1.6628238
      A2.A0D2.31TCGA C8.A12U.31TCGA AR.A1AS.31TCGA A8.A09G.32TCGA C8.A131.32TCGA C8.A134.32TCGA
A2.A0YF.33TCGA BH.A0DD.33TCGA BH.A0E9.33TCGA
1      0.1074909      -0.4815502      1.222507      -1.5233435      2.7072502      0.1401818
0.3113192      -0.6923158      1.466665
3      0.1074909      -0.4815502      1.222507      -1.5099719      2.7376293      0.1331178
0.2961771      -0.6641611      1.474474
6      0.1041645      -0.4852105      1.218974      -1.5126462      2.7376293      0.1260538
0.2961771      -0.6618149      1.474474
7      0.1041645      -0.4815502      1.222507      -1.5153205      2.7376293      0.1154577
0.2961771      -0.6641611      1.474474
10     -1.5124729      -0.9976551      1.741961      0.1347327      -1.1205244      -2.3604810
2.3517137      0.9805402      2.813743
11     -1.4060276      -1.3527060      1.038755      -0.1139788      -0.6800265      -1.8589356
2.4955634      1.6914453      2.341290
      AR.A0TT.34TCGA AO.A12B.34TCGA A2.A0SW.35TCGA AO.A0JL.35TCGA BH.A0BV.35TCGA A2.A0YM.36TCGA
BH.A0C7.36TCGA A2.A0SX.36TCGA X263d3f.I.CPTAC
1      -0.51142119      -0.9639039      -0.4877725      -0.106680      -0.06583842      0.6558497
-0.5522120      -0.3985598      0.5985845
3      -0.52606668      -0.9439194      -0.4877725      -0.106680      -0.06583842      0.6558497
-0.5522120      -0.3926014      0.6039931
6      -0.52972805      -0.9382095      -0.4877725      -0.106680      -0.05589267      0.6581426
-0.5477494      -0.3926014      0.6066975
7      -0.52972805      -0.9439194      -0.4877725      -0.106680      -0.06252317      0.6558497
-0.5522120      -0.3926014      0.6039931
10     -0.52606668      1.3257521      0.7311482      -1.177327      0.70993057      1.3070365
0.4875738      0.6948104      2.7782634
11     -0.06107243      1.6455045      0.3580749      -3.466189      -0.50345170      1.3001577
0.7218603      0.5726628      3.8464683
      b1cdb9.I.CPTAC c4155b.C.CPTAC
1      -0.1912845      0.5669753
3      -0.1860225      0.5767473
6      -0.1839177      0.5787017
7      -0.1860225      0.5767473
10     1.3673298      3.2151898
11     1.0884420      3.4809884
> proteoms.pca <- prcomp(clean.proteoms[,4:length(colnames(clean.proteoms))], center = T)
> names(proteoms.pca)
[1] "sdev" "rotation" "center" "scale" "x"
> head(proteoms.pca$x)
      PC1      PC2      PC3      PC4      PC5      PC6      PC7      PC8
PC9      PC10     PC11     PC12     PC13
1 -3.599764 -1.566944 0.5272164 -3.2158079 3.177636 -0.7555303 1.236568 -1.2388591
-0.7678642 -1.2713068 0.1458712 -1.6907957 0.5198961
3 -3.663029 -1.561971 0.5348120 -3.1743285 3.207102 -0.7749749 1.242069 -1.2660165
-0.7820288 -1.2971958 0.1380706 -1.6795248 0.5232510
6 -3.659489 -1.562818 0.5234919 -3.1827055 3.196925 -0.7673091 1.240862 -1.2665938
-0.7776568 -1.2972613 0.1396268 -1.6770480 0.5201774
7 -3.653587 -1.567978 0.5246178 -3.1841037 3.201360 -0.7729774 1.241310 -1.2640772
-0.7821890 -1.2963795 0.1389090 -1.6797551 0.5243077
10 -6.987977 3.848287 -8.3213847 -1.8855901 3.140966 -0.9524179 -1.123623 -0.8112765
0.2357237 0.6794838 0.1724692 1.1846840 -0.7580815
11 -6.221785 5.179066 -6.4654576 0.2749185 3.310336 -0.7082599 -1.232022 -1.8427013
-0.2702671 -1.3154296 -1.2513509 -0.2886961 -1.3710253
      PC14      PC15      PC16      PC17      PC18      PC19      PC20      PC21
PC22      PC23      PC24      PC25
1 0.1278848 -0.3091430 0.05983299 0.2519446 -0.7335441 -0.6369246 -2.0793540 1.630920
0.3721621 -0.1727348 -1.17669988 1.1744078

```

0.	0.1187915	-0.3098289	0.05973205	0.2378868	-0.7372402	-0.6419921	-2.0874183	1.649980	
0.	0.3701661	-0.1682483	-1.19661315	1.1759788					
6	0.1144474	-0.3055530	0.05963252	0.2414510	-0.7345915	-0.6479520	-2.0839287	1.648393	
0.	0.3849554	-0.1689400	-1.18751383	1.1778023					
7	0.1230106	-0.3094607	0.06432937	0.2444940	-0.7423673	-0.6436291	-2.0861637	1.653336	
0.	0.3773675	-0.1672175	-1.19194202	1.1769329					
10	-0.7586473	-0.2943068	0.01816792	1.1820989	-0.1288812	-0.3481304	0.4277494	1.253868	
1.	4022907	-0.2483602	-0.04800123	-0.1380246					
11	1.2258601	-1.0441680	-1.57195407	0.8718249	-1.4045816	-0.9149134	0.7500041	2.066840	
0.	4275349	-1.6789903	0.09785348	-0.4863625					
	PC26	PC27	PC28	PC29	PC30	PC31	PC32	PC33	
PC34	PC35	PC36	PC37						
1	1.2730789	0.1341272	-0.7448445	-0.7943628	1.3235885	-0.4043017	-1.841656	0.9764388	
-0.	005675926	-0.31359739	0.011202981	-1.043821					
3	1.2903815	0.1275161	-0.7459513	-0.8061584	1.3376292	-0.4174232	-1.854814	0.9651947	
-0.	011545822	-0.31674621	0.007301704	-1.055907					
6	1.2946446	0.1220445	-0.7505712	-0.8057893	1.3398976	-0.4177089	-1.853340	0.9623030	
-0.	014780420	-0.32223840	0.004752804	-1.056947					
7	1.2950546	0.1262595	-0.7518949	-0.8091630	1.3413802	-0.4184353	-1.855226	0.9640567	
-0.	012931139	-0.32380739	0.006513727	-1.061367					
10	-0.2838253	-1.7303282	-2.2830296	-1.4509413	0.1545269	0.5648301	-2.599145	1.5567802	
1.	095657011	-0.03378417	1.110011541	-1.045809					
11	-0.2806181	-1.1364446	-3.3429159	-2.0558134	0.1427084	0.9110828	-2.865932	1.7171899	
1.	734126207	-0.43004480	1.076600608	-1.313402					
	PC38	PC39	PC40	PC41	PC42	PC43	PC44	PC45	
PC46	PC47	PC48	PC49						
1	0.4677184	-0.5213558	0.5530872	1.4508069	0.1519437	-0.5177118	0.1571119	0.3512621	
-0.	6542297	-0.21949651	0.8885389	0.6484622					
3	0.4707762	-0.5327662	0.5713006	1.4725120	0.1616131	-0.5057932	0.1645801	0.3607477	
-0.	6630465	-0.21168309	0.8934769	0.6415484					
6	0.4661709	-0.5350788	0.5665955	1.4757970	0.1662754	-0.5058287	0.1650478	0.3631333	
-0.	6674184	-0.21289532	0.8896392	0.6441700					
7	0.4677679	-0.5323697	0.5670929	1.4795896	0.1629602	-0.5067911	0.1672954	0.3605918	
-0.	6706043	-0.21082389	0.8894013	0.6380101					
10	-0.3650018	0.3574800	0.6458406	0.1879803	0.8767433	-1.7227723	-0.8167122	-0.7023169	
0.	6607983	-0.90787179	-0.5494380	0.6938012					
11	0.5048332	0.3143009	0.5880441	0.2633403	-0.1787429	-1.1603788	-0.9667014	-0.8833431	
0.	5741339	0.07906612	-1.2556128	1.5393121					
	PC50	PC51	PC52	PC53	PC54	PC55	PC56	PC57	
PC58	PC59	PC60	PC61						
1	0.7389762	0.5054221	1.3039426	-0.5745544	-0.4280911				

```

10  0.1201587 -0.4771048 0.6366893  1.6612986 -0.19757528 -0.7090803 -0.6617768 -0.5074338
0.09124969  0.500687867 -0.11114594  0.3997009
11  0.1137704  0.6415849 0.3727557  1.3547096  0.03217751 -0.7852475 -0.6970829 -0.6019859
0.47869164 -0.263715277 -0.38793632  0.1247979

```

```

          PC74      PC75      PC76      PC77      PC78      PC79      PC80
PC81      PC82      PC83
1  -0.22110227 -0.07298477 -0.09755746  0.5551588  0.1641169 -0.1179238  0.1555411
0.03029241 -0.1525013 -0.017847283
3  -0.22229465 -0.07052686 -0.10640685  0.5404295  0.1648348 -0.1177008  0.1487354
0.02516098 -0.1655298 -0.011432206
6  -0.22163271 -0.07040523 -0.10422538  0.5396141  0.1684001 -0.1177256  0.1535451
0.02395941 -0.1671409 -0.008769818
7  -0.22310874 -0.07021702 -0.10415823  0.5431360  0.1649801 -0.1163722  0.1468095
0.02538999 -0.1617406 -0.009363245
10  1.02721192  0.04138428  0.07813679  0.6814881 -0.4870878  0.4792118 -0.0723777
-0.64051379  0.1210410  0.156150165
11  0.09586813  0.23616488 -0.28745803 -0.5772096  0.3530266 -0.6093605  0.0755825
-0.94027481 -0.2943216 -0.267363756

```

```

> pca.var <- proteoms.pca$sdev^2
> pca.var.per <- round(pca.var/sum(pca.var)*100,1)
> barplot(pca.var.per, main="Scree Plot", xlab="Principal Component", ylab="Percent
Variation")
> (x <- cbind(1:length(pca.var.per),cumsum(pca.var.per)))[x[,1] == 8,]
[1] 8.0 52.6
> hclust.out <- hclust(dist(as.matrix(proteoms.pca$x[,1:8])), method = "complete")
> plot(hclust.out)

```

Scree Plot

