

HW3

5/21/2021

```
##loading library
```

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.3      v purrr  0.3.4
```

```
## v tibble  3.1.1      v dplyr  1.0.5
```

```
## v tidyr   1.1.3      v stringr 1.4.0
```

```
## v readr   1.4.0      v forcats 0.5.1
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
```

```
## x dplyr::lag()     masks stats::lag()
```

```
library(ROCR)
```

```
library(tree)
```

```
## Registered S3 method overwritten by 'tree':
```

```
##   method      from
```

```
##   print.tree cli
```

```
library(maptree)
```

```
## Loading required package: cluster
```

```
## Loading required package: rpart
```

```
library(class)
```

```
library(lattice)
```

```
library(ggthemes)
```

```
library(superheat)
```

```
## loading the drug dataset and transform variables
```

```
drug_use <- read_csv('drug.csv',
```

```
col_names = c('ID', 'Age', 'Gender', 'Education', 'Country', 'Ethnicity', 'Nscore', 'Escore', 'Oscore', 'Ascore',  
'Choc', 'Coke', 'Crack', 'Ecstasy', 'Heroin', 'Ketamine', 'Legalh', 'LSD', 'Meth', 'Mushrooms', 'Nicotine', 'Seme
```

```
##
```

```
## -- Column specification -----
```

```
## cols(
```

```
##   .default = col_character(),
```

```

## ID = col_double(),
## Age = col_double(),
## Gender = col_double(),
## Education = col_double(),
## Country = col_double(),
## Ethnicity = col_double(),
## Nscore = col_double(),
## Escore = col_double(),
## Oscore = col_double(),
## Ascore = col_double(),
## Cscore = col_double(),
## Impulsive = col_double(),
## SS = col_double()
## )
## i Use 'spec()' for the full column specifications.

drug_use <- drug_use %>% mutate_at(as.ordered, .vars=vars(Alcohol:VSA))
drug_use <- drug_use %>%
mutate(Gender = factor(Gender, labels=c("Male", "Female"))) %>% mutate(Ethnicity = factor(Ethnicity, labels=c("White", "Black", "Hispanic", "Asian", "Other")))

##1. Logistic regression for drug use prediction ###a

## create a new factor response variable recent_cannabis_use

drug_use = drug_use %>% mutate(recent_cannabis_use=ifelse(Cannabis >= "CL3","Yes","No"))%>% mutate(recent_cannabis_use = factor(recent_cannabis_use, labels=c("Yes", "No")))

###b

##create a new tibble
drug_use_subset <- drug_use %>% select(Age:SS, recent_cannabis_use)
##Split drug_use_subset into a training data set and a test data set
set.seed(1)
training.indices = sample(1:nrow(drug_use_subset), 1500)
drug_use_train = drug_use_subset[training.indices,]
drug_use_test = drug_use_subset[-training.indices,]
dim(drug_use_train)

## [1] 1500 13

dim(drug_use_test)

## [1] 385 13

###c

##Fit a logistic regression
glm.fit = glm(recent_cannabis_use ~ ., data=drug_use_train, family=binomial)
# Summarize the logistic regression model
summary(glm.fit)

```

```
##
## Call:
## glm(formula = recent_cannabis_use ~ ., family = binomial, data = drug_use_train)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.9072  -0.5971   0.1416   0.5426   2.6600
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)      0.94949    0.64574   1.470 0.141457
## Age             -0.84406    0.09328  -9.049 < 2e-16 ***
## GenderFemale    -0.55929    0.15715  -3.559 0.000372 ***
## Education       -0.33389    0.07962  -4.193 2.75e-05 ***
## CountryCanada   13.10904   627.22755   0.021 0.983325
## CountryNew Zealand -1.16844    0.31848  -3.669 0.000244 ***
## CountryOther    -0.05676    0.46772  -0.121 0.903412
## CountryIreland  -0.28763    0.67573  -0.426 0.670354
## CountryUK       -0.43371    0.37043  -1.171 0.241674
## CountryUSA      -1.75636    0.19262  -9.118 < 2e-16 ***
## EthnicityAsian  -0.67025    0.96037  -0.698 0.485230
## EthnicityWhite   0.74053    0.63843   1.160 0.246081
## EthnicityMixed:White/Black -0.04713    1.09013  -0.043 0.965515
## EthnicityOther   1.07889    0.76823   1.404 0.160206
## EthnicityMixed:White/Asian  0.72525    1.01565   0.714 0.475178
## EthnicityMixed:Black/Asian 14.27149   766.28165   0.019 0.985141
## Nscore          -0.10143    0.09034  -1.123 0.261551
## Escore          -0.13375    0.09559  -1.399 0.161742
## Oscore           0.71000    0.09137   7.770 7.83e-15 ***
## Ascore           0.03058    0.08232   0.372 0.710251
## Cscore          -0.35855    0.09132  -3.926 8.63e-05 ***
## Impulsive       -0.09043    0.10093  -0.896 0.370290
## SS              0.58068    0.10836   5.359 8.39e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2077.2  on 1499  degrees of freedom
## Residual deviance: 1202.1  on 1477  degrees of freedom
## AIC: 1248.1
##
## Number of Fisher Scoring iterations: 14
```

##2. Decision tree models of drug use

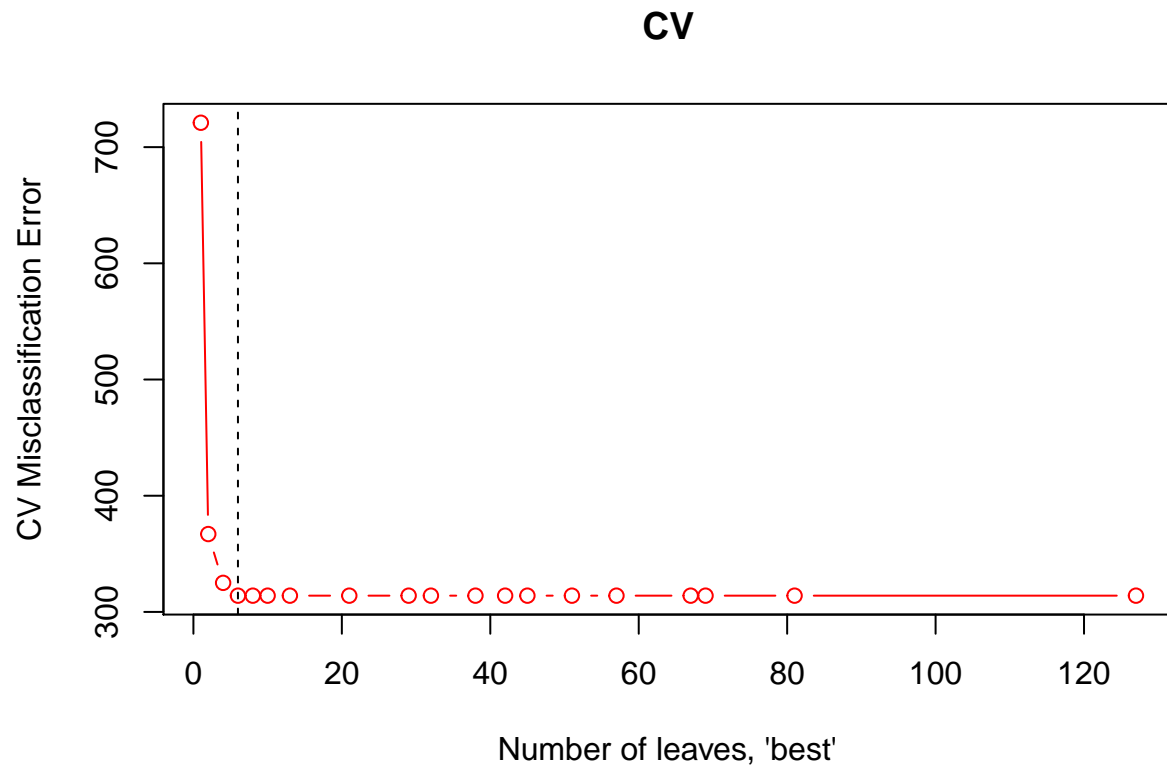
```
## Construct a decision tree
tree_parameters = tree.control(nobs=nrow(drug_use_train), minsize=10, mindev=1e-3)
drugtree = tree(recent_cannabis_use~.,control = tree_parameters,data = drug_use_train)
```

###a Use 10-fold CV to select the a tree

```

# Set random seed
set.seed(1)
cv = cv.tree(drugtree,FUN=prune.misclass, K=10)
best.size.cv = 6
plot(cv$size , cv$dev, type="b",
     xlab = "Number of leaves, \'best\'", ylab = "CV Misclassification Error",
     col = "red", main="CV")
abline(v=best.size.cv, lty=2)

```

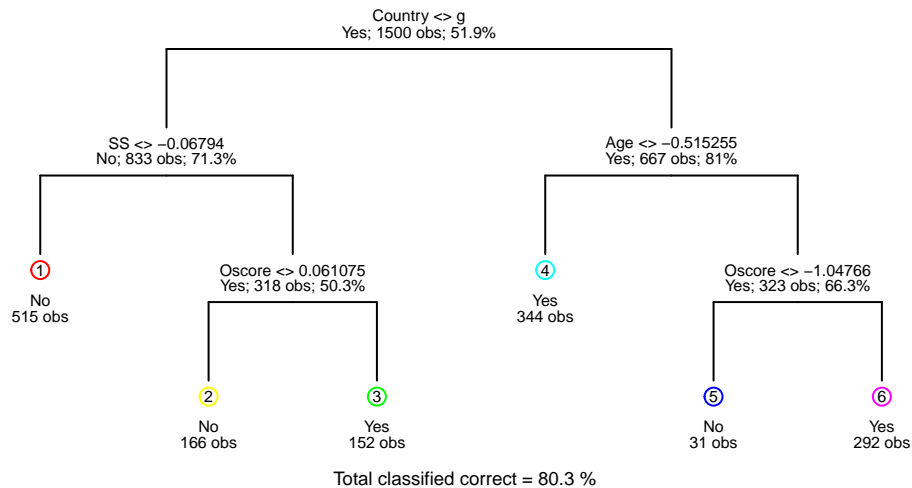


```
###b Prune the tree
```

```

# prune the original tree using the best size in a
drugtree.pruned = prune.misclass(drugtree, best=best.size.cv)
draw.tree(drugtree.pruned, nodeinfo=TRUE,cex = 0.5)

```



The variable 'Country' is split

first in this decision tree.

###c Compute and print the confusion matrix for the test data

```

# Predict on test set
predictions = predict(drugtree.pruned, drug_use_test, type="class")
# get the true response of the test data
truth = drug_use_test$recent_cannabis_use
# Obtain confusion matrix
confusion_matrix = table(truth, predictions)
confusion_matrix

```

```

##      predictions
## truth  No  Yes
##   No  125  40
##   Yes  45 175

```

```

# get true positive rate
true_positive_rate = confusion_matrix[2,2]/sum(confusion_matrix[2,])
true_positive_rate

```

```
## [1] 0.7954545
```

```

# get false positive rate
false_positive_rate = confusion_matrix[1,2]/sum(confusion_matrix[1,])
false_positive_rate

```

```
## [1] 0.2424242
```

###3.Model Comparison ###a

```

# get prediction from logistic regression model using test data
prob_log_testing = predict(glm.fit,drug_use_test,type="response")
pred_log = prediction(prob_log_testing, truth)
#calculate the True Positive Rate and False Positive Rate by performance()
perf_log = performance(pred_log, measure="tpr", x.measure="fpr")
# for the decision tree model

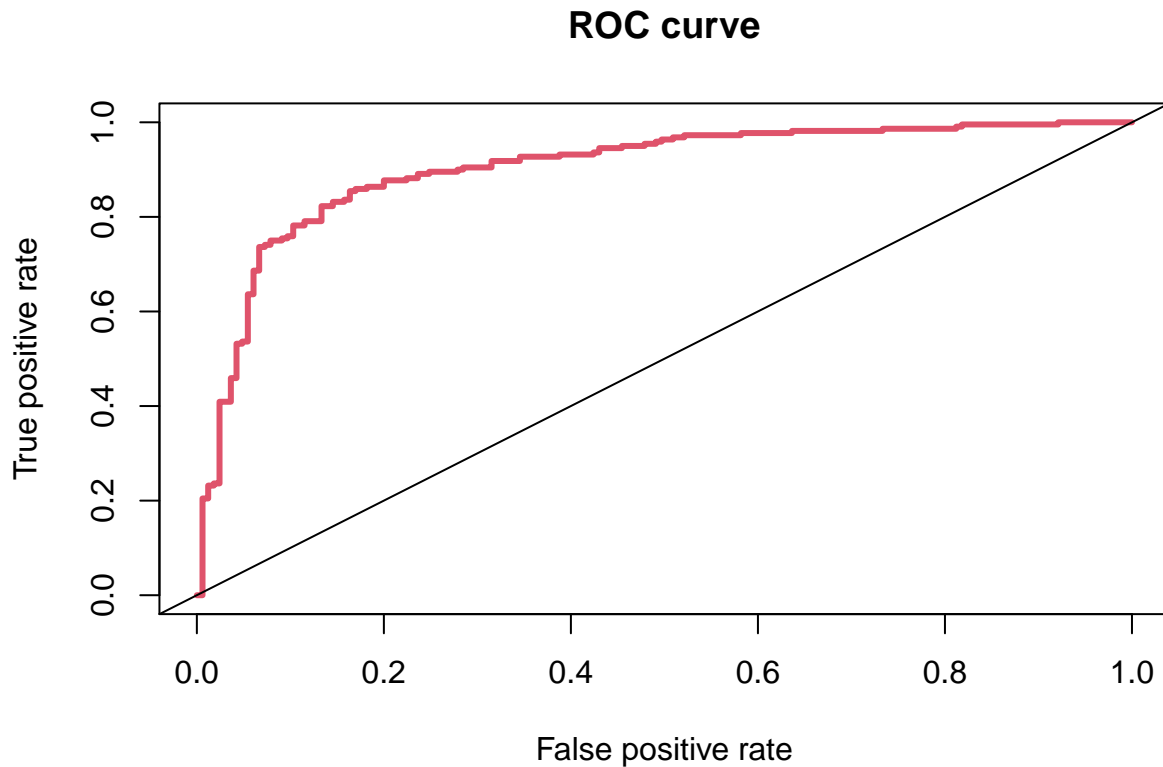
```

```

prob_tree_testing = predict(drugtree.pruned,newdata = drug_use_test)[,2]
pred_tree = prediction(prob_tree_testing,truth)
perf_tree = performance(pred_tree, measure="tpr", x.measure="fpr")

plot(perf_log, col=2, lwd=3, main="ROC curve")
abline(0,1)

```



```
###b
```

```

##Compute the AUC for logistic regression model
auc_log = performance(pred_log, "auc")@y.values
auc_log

```

```

## [[1]]
## [1] 0.902562

```

```

##Compute the AUC for decision tree model
auc_tree = performance(pred_tree, "auc")@y.values
auc_tree

```

```

## [[1]]
## [1] 0.8570523

```

```
# logistic is larger
```

```
##4. Clustering and dimension reduction for gene expression data
```

```
## read leukemia_data
leukemia_data <- read_csv("leukemia_data.csv")
```

```
## Warning: Duplicated column names deduplicated: 'FCGRT' => 'FCGRT_1' [3],
## 'TUBB4B' => 'TUBB4B_1' [49], 'SSR1' => 'SSR1_1' [67], 'HSP90AB1' =>
## 'HSP90AB1_1' [115], 'TMBIM6' => 'TMBIM6_1' [118], 'GAB1' => 'GAB1_1' [119],
## 'MPHOSPH9' => 'MPHOSPH9_1' [153], 'STK38' => 'STK38_1' [157], 'SFPQ' =>
## 'SFPQ_1' [159], 'RIPOR2' => 'RIPOR2_1' [181], 'HLA-F' => 'HLA-F_1' [188],
## 'PRPF40A' => 'PRPF40A_1' [198], 'SEPT6' => 'SEPT6_1' [205], 'CD22' =>
## 'CD22_1' [235], 'NCF4' => 'NCF4_1' [250], 'WAS' => 'WAS_1' [260], 'HLA-
## G' => 'HLA-G_1' [297], 'TRAF3IP3' => 'TRAF3IP3_1' [307], 'ZNF266' =>
## 'ZNF266_1' [364], 'CRYBG1' => 'CRYBG1_1' [441], 'BRD8' => 'BRD8_1' [460], 'MDC1'
## => 'MDC1_1' [464], 'RAC2' => 'RAC2_1' [478], 'IL10RB' => 'IL10RB_1' [483],
## 'AKAP17A' => 'AKAP17A_1' [542], 'N4BP2L1' => 'N4BP2L1_1' [547], 'ARPC4' =>
## 'ARPC4_1' [565], 'SRSF10' => 'SRSF10_1' [576], 'RAPGEF2' => 'RAPGEF2_1' [583],
## 'PARP2' => 'PARP2_1' [587], 'TRIM33' => 'TRIM33_1' [610], 'KAT8' =>
## 'KAT8_1' [665], 'ASMTL' => 'ASMTL_1' [715], 'LSM7' => 'LSM7_1' [727],
## 'HLA-DQB1' => 'HLA-DQB1_1' [732], 'FMR1' => 'FMR1_1' [826], 'RASGRP2' =>
## 'RASGRP2_1' [858], 'LIMK2' => 'LIMK2_1' [866], 'TMEM106C' => 'TMEM106C_1' [881],
## 'TGOLN2' => 'TGOLN2_1' [937], 'SLC25A1' => 'SLC25A1_1' [940], 'NMT1' =>
## 'NMT1_1' [942], 'ENSA' => 'ENSA_1' [947], 'ENSA' => 'ENSA_2' [948], 'UBR5'
## => 'UBR5_1' [963], 'UBE2J1' => 'UBE2J1_1' [966], 'ACTN1' => 'ACTN1_1' [994],
## 'TRA2A' => 'TRA2A_1' [1003], 'ATXN10' => 'ATXN10_1' [1057], 'CUL1' =>
## 'CUL1_1' [1077], 'XBP1' => 'XBP1_1' [1094], 'ATP2A2' => 'ATP2A2_1' [1110],
## 'LDLRAD4' => 'LDLRAD4_1' [1118], 'ARHGEF2' => 'ARHGEF2_1' [1134],
## 'IDH3B' => 'IDH3B_1' [1141], 'SERBP1' => 'SERBP1_1' [1188], 'TRIM44' =>
## 'TRIM44_1' [1205], 'TRIM44' => 'TRIM44_2' [1206], 'PTPRC' => 'PTPRC_1' [1219],
## 'PTPRC' => 'PTPRC_2' [1220], 'PPP2R5C' => 'PPP2R5C_1' [1235], 'PPP2R5C'
## => 'PPP2R5C_2' [1236], 'ADAM10' => 'ADAM10_1' [1241], 'NFATC3' =>
## 'NFATC3_1' [1252], 'ILF3' => 'ILF3_1' [1264], 'RBM6' => 'RBM6_1' [1274],
## 'CTNNA1' => 'CTNNA1_1' [1297], 'CTNNA1' => 'CTNNA1_2' [1298], 'IGHM' =>
## 'IGHM_1' [1302], 'IGHM' => 'IGHM_2' [1303], 'IGHM' => 'IGHM_3' [1304], 'SFPQ' =>
## 'SFPQ_2' [1321], 'RBCK1' => 'RBCK1_1' [1398], 'NFATC2IP' => 'NFATC2IP_1' [1408],
## 'ILF3' => 'ILF3_2' [1432], 'RAE1' => 'RAE1_1' [1436], 'ITPR1' =>
## 'ITPR1_1' [1443], 'NCBP2' => 'NCBP2_1' [1448], 'STAT1' => 'STAT1_1' [1486],
## 'AZIN1' => 'AZIN1_1' [1497], 'SEC13' => 'SEC13_1' [1517], 'ABI1' =>
## 'ABI1_1' [1565], 'CYB5B' => 'CYB5B_1' [1607], 'HUWE1' => 'HUWE1_1' [1624],
## 'RAB1A' => 'RAB1A_1' [1634], 'AHCYL1' => 'AHCYL1_1' [1652], 'EIF1AX' =>
## 'EIF1AX_1' [1661], 'MAGED2' => 'MAGED2_1' [1689], 'SCAF11' => 'SCAF11_1' [1709],
## 'BLCAP' => 'BLCAP_1' [1716], 'TROVE2' => 'TROVE2_1' [1729], 'CTCF' =>
## 'CTCF_1' [1745], 'RAB8A' => 'RAB8A_1' [1754], 'ACTR2' => 'ACTR2_1' [1768],
## 'HMGN4' => 'HMGN4_1' [1771], 'NDUFB7' => 'NDUFB7_1' [1793], 'VAMP3' =>
## 'VAMP3_1' [1796], 'SRSF6' => 'SRSF6_1' [1808], 'TNPO3' => 'TNPO3_1' [1811],
## 'SRSF1' => 'SRSF1_1' [1834], 'TMED10' => 'TMED10_1' [1847], 'AP3D1' =>
## 'AP3D1_1' [1872], 'MAPKAPK2' => 'MAPKAPK2_1' [1877], 'BRD2' => 'BRD2_1' [1891],
## 'BRD2' => 'BRD2_2' [1892], 'GARS' => 'GARS_1' [1901], 'SNX1' => 'SNX1_1' [1902],
## 'TSC22D3' => 'TSC22D3_1' [1927], 'AMD1' => 'AMD1_1' [1951], 'LITAF' =>
## 'LITAF_1' [2011], 'GLUD1' => 'GLUD1_1' [2059], 'KDELRL1' => 'KDELRL1_1' [2079],
## 'PGK1' => 'PGK1_1' [2099], 'VDAC2' => 'VDAC2_1' [2107], 'ADH5' =>
## 'ADH5_1' [2111], 'MEF2C' => 'MEF2C_1' [2113], 'MEF2C' => 'MEF2C_2' [2114],
## 'RCN2' => 'RCN2_1' [2125], 'PCMT1' => 'PCMT1_1' [2134], 'PCMT1' =>
## 'PCMT1_2' [2135], 'CD79A' => 'CD79A_1' [2149], 'MARCH6' => 'MARCH6_1' [2169],
## 'CBX3' => 'CBX3_1' [2180], 'LSM14A' => 'LSM14A_1' [2217], 'SORL1' =>
```

```

## 'SORL1_1' [2220], 'ICAM2' => 'ICAM2_1' [2244], 'SNRPB' => 'SNRPB_1' [2246],
## 'CYB5A' => 'CYB5A_1' [2248], 'BTN3A2' => 'BTN3A2_1' [2277], 'DICER1' =>
## 'DICER1_1' [2280], 'HADH' => 'HADH_1' [2281], 'HDGF' => 'HDGF_1' [2285], 'SEPT6'
## => 'SEPT6_2' [2306], 'SSBP1' => 'SSBP1_1' [2315], 'H2AFV' => 'H2AFV_1' [2318],
## 'PTPA' => 'PTPA_1' [2331], 'FBL' => 'FBL_1' [2354], 'OGT' => 'OGT_1' [2362],
## 'SLC25A1' => 'SLC25A1_2' [2377], 'FUBP1' => 'FUBP1_1' [2386], 'TUBGCP2' =>
## 'TUBGCP2_1' [2400], 'COX5B' => 'COX5B_1' [2402], 'VDAC1' => 'VDAC1_1' [2410],
## 'HNRNPDL' => 'HNRNPDL_1' [2431], 'THUMP1' => 'THUMP1_1' [2443], 'CDV3'
## => 'CDV3_1' [2444], 'UBE3B' => 'UBE3B_1' [2447], 'SFPQ' => 'SFPQ_3' [2451],
## 'STX16' => 'STX16_1' [2452], 'SMARCA2' => 'SMARCA2_1' [2471], 'CHD8' =>
## 'CHD8_1' [2475], 'TCF25' => 'TCF25_1' [2490], 'API5' => 'API5_1' [2491],
## 'SAP18' => 'SAP18_1' [2493], 'AHCYL1' => 'AHCYL1_2' [2501], 'CTBP1' =>
## 'CTBP1_1' [2503], 'AES' => 'AES_1' [2512], 'PURA' => 'PURA_1' [2514], 'BCL11A'
## => 'BCL11A_1' [2518], 'BUB3' => 'BUB3_1' [2534], 'RER1' => 'RER1_1' [2537],
## 'ATXN2L' => 'ATXN2L_1' [2541], 'JAK1' => 'JAK1_1' [2548], 'GUSBP11' =>
## 'GUSBP11_1' [2564], 'JTB' => 'JTB_1' [2568], 'BRD3' => 'BRD3_1' [2571], 'RSU1'
## => 'RSU1_1' [2584], 'ADD3' => 'ADD3_1' [2619], 'UBE2I' => 'UBE2I_1' [2627],
## 'MRPS12' => 'MRPS12_1' [2640], 'CTNNA1' => 'CTNNA1_3' [2641], 'XRCC5' =>
## 'XRCC5_1' [2642], 'ITGA4' => 'ITGA4_1' [2644], 'CTNNA1' => 'CTNNA1_4' [2647],
## 'FYN' => 'FYN_1' [2649], 'ERG' => 'ERG_1' [2652], 'RAC1' => 'RAC1_1' [2654],
## 'LCK' => 'LCK_1' [2657], 'PTK2B' => 'PTK2B_1' [2664], 'SKP1' =>
## 'SKP1_1' [2665], 'PRKDC' => 'PRKDC_1' [2666], 'MYC' => 'MYC_1' [2668], 'RBL2'
## => 'RBL2_1' [2673], 'AZIN1' => 'AZIN1_2' [2674], 'CCNA2' => 'CCNA2_1' [2681],
## 'FOS' => 'FOS_1' [2688], 'FOS' => 'FOS_2' [2689], 'RAF1' => 'RAF1_1' [2690],
## 'RAP1B' => 'RAP1B_1' [2692], 'ERCC1' => 'ERCC1_1' [2696], 'ERCC1' =>
## 'ERCC1_2' [2697], 'RAN' => 'RAN_1' [2702], 'TRIM27' => 'TRIM27_1' [2703],
## 'PMS2P3' => 'PMS2P3_1' [2708], 'TGFB2' => 'TGFB2_1' [2710], 'PCNA' =>
## 'PCNA_1' [2712], 'MYC' => 'MYC_2' [2714], 'CDK13' => 'CDK13_1' [2717],
## 'CCND3' => 'CCND3_1' [2719], 'FARSA' => 'FARSA_1' [2732], 'FARSA' =>
## 'FARSA_2' [2733], 'DAXX' => 'DAXX_1' [2734], 'UBE3A' => 'UBE3A_1' [2735],
## 'ARAF' => 'ARAF_1' [2739], 'UBE2N' => 'UBE2N_1' [2747], 'RASA1' =>
## 'RASA1_1' [2748], 'ABL1' => 'ABL1_1' [2749], 'ABL1' => 'ABL1_2' [2750], 'MTA1'
## => 'MTA1_1' [2753], 'EIF3I' => 'EIF3I_1' [2754], 'SYK' => 'SYK_1' [2761],
## 'TOP2A' => 'TOP2A_1' [2762], 'RB1' => 'RB1_1' [2764], 'TOP2B' =>
## 'TOP2B_1' [2765], 'TNFRSF1B' => 'TNFRSF1B_1' [2766], 'GRB2' => 'GRB2_1' [2769],
## 'RBM5' => 'RBM5_1' [2770], 'N4BP2L1' => 'N4BP2L1_2' [2773], 'N4BP2L2' =>
## 'N4BP2L2_1' [2774], 'NME1' => 'NME1_1' [2775], 'TYMS' => 'TYMS_1' [2776],
## 'DYRK1A' => 'DYRK1A_1' [2778], 'FEN1' => 'FEN1_1' [2779], 'FEN1' =>
## 'FEN1_2' [2780], 'ETS2' => 'ETS2_1' [2781], 'FNTA' => 'FNTA_1' [2783], 'JAK1'
## => 'JAK1_2' [2787], 'MYB' => 'MYB_1' [2792], 'MYB' => 'MYB_2' [2793], 'MYB' =>
## 'MYB_3' [2794], 'MYB' => 'MYB_4' [2795], 'MYB' => 'MYB_5' [2796], 'SMAD2' =>
## 'SMAD2_1' [2798], 'PTEN' => 'PTEN_1' [2799], 'MAPKAPK2' => 'MAPKAPK2_2' [2800],
## 'PSMD9' => 'PSMD9_1' [2801], 'PSMA4' => 'PSMA4_1' [2806], 'SRF' =>
## 'SRF_1' [2810], 'LYN' => 'LYN_1' [2815], 'IL7R' => 'IL7R_1' [2817], 'TCF3' =>
## 'TCF3_1' [2818], 'TCF3' => 'TCF3_2' [2819], 'NFKB1' => 'NFKB1_1' [2820], 'NFKB1'
## => 'NFKB1_2' [2821], 'RPA1' => 'RPA1_1' [2822], 'PPP2R2A' => 'PPP2R2A_1' [2823],
## 'TERF1' => 'TERF1_1' [2826], 'BCR' => 'BCR_1' [2828], 'RBBP4' =>
## 'RBBP4_1' [2830], 'TERF2' => 'TERF2_1' [2831], 'PSMB4' => 'PSMB4_1' [2834],
## 'PSMB7' => 'PSMB7_1' [2836], 'PARP1' => 'PARP1_1' [2838], 'RELA' =>
## 'RELA_1' [2840], 'RELA' => 'RELA_2' [2841], 'EIF2S3' => 'EIF2S3_1' [2842],
## 'YWHAZ' => 'YWHAZ_1' [2846], 'PTP4A2' => 'PTP4A2_1' [2847], 'POLR2H' =>
## 'POLR2H_1' [2850], 'GAB1' => 'GAB1_2' [2851], 'PRKDC' => 'PRKDC_2' [2852],
## 'PRKCB' => 'PRKCB_1' [2855], 'SAT1' => 'SAT1_1' [2862], 'PTPRE' =>
## 'PTPRE_1' [2865], 'RPL22' => 'RPL22_1' [2866], 'EIF2S1' => 'EIF2S1_1' [2867],

```



```
## 'CYC1' => 'CYC1_1' [2869], 'HSP90AB1' => 'HSP90AB1_2' [2870], 'CD44' =>
## 'CD44_1' [2873], 'MAP2K1' => 'MAP2K1_1' [2875], 'TNK2' => 'TNK2_1' [2877],
## 'GNA13' => 'GNA13_1' [2879], 'NR3C1' => 'NR3C1_1' [2882], 'RAB1A' =>
## 'RAB1A_2' [2888], 'ODC1' => 'ODC1_1' [2890], 'PLCG2' => 'PLCG2_1' [2891], 'RFC4'
## => 'RFC4_1' [2894], 'FLT3' => 'FLT3_1' [2895], 'EIF2AK2' => 'EIF2AK2_1' [2902],
## 'USP9X' => 'USP9X_1' [2913], 'PSMD7' => 'PSMD7_1' [2917], 'PPP1CA' =>
## 'PPP1CA_1' [2924], 'TUBB4B' => 'TUBB4B_2' [2926], 'ARRB2' => 'ARRB
```

```
##
## -- Column specification -----
## cols(
##   .default = col_double(),
##   Type = col_character()
## )
## i Use 'spec()' for the full column specifications.
```

```
####a
```

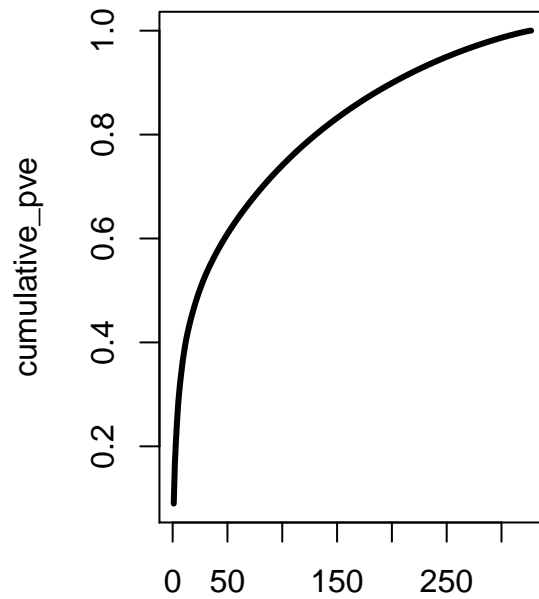
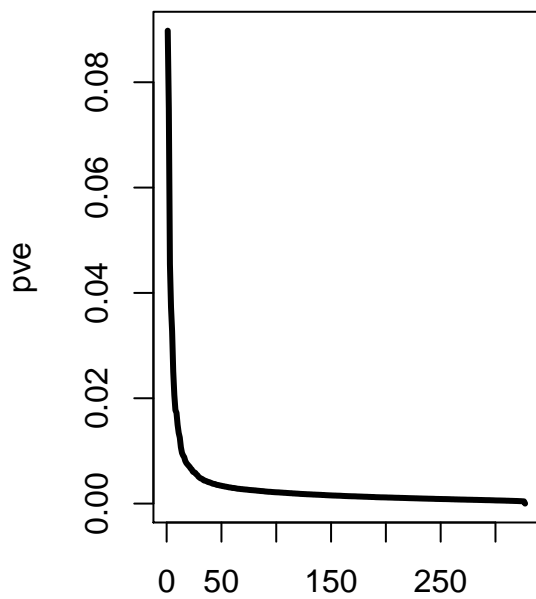
```
##Convert the Type column to factor
leukemia_data = leukemia_data %>% mutate(Type = factor(Type))
##the number of patients with each leukemia subtype
table(leukemia_data$Type)
```

```
##
##      BCR-ABL      E2A-PBX1 Hyperdip50      MLL      OTHERS      T-ALL      TEL-AML1
##           15           27           64           20           79           43           79
```

Subtype “BCR-ABL” occurs the least in this data

```
####b
```

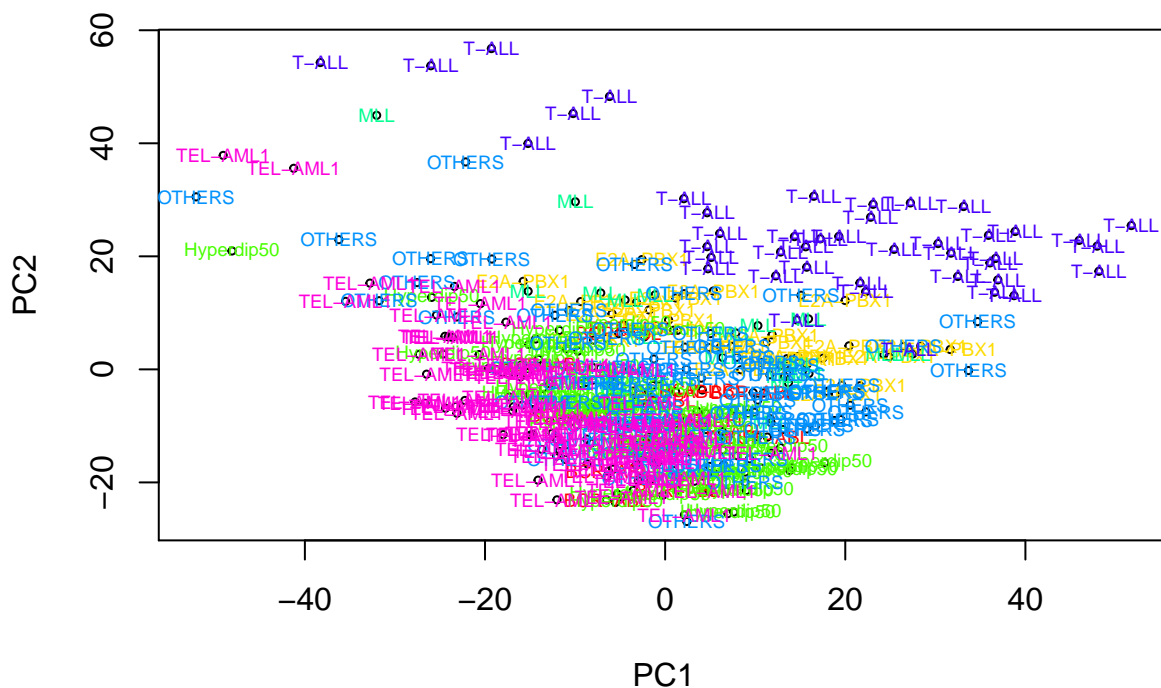
```
## exclude the Type column
leukemia_data_wt = leukemia_data %>% select(-Type)
pr.out=prcomp(leukemia_data_wt, scale=TRUE, center = TRUE)
## get the variance explained by each principal component
pr.var=pr.out$sdev ^2
pve=pr.var/sum(pr.var)
## get cumulative pve
cumulative_pve = cumsum(pve)
## plot PVE and cumulative pve
par(mfrow=c(1, 2))
plot(pve, type="l", lwd=3)
plot(cumulative_pve, type="l", lwd=3)
```



###c

```
## set color
rainbow_colors <- rainbow(7)

plot_colors <- rainbow_colors[leukemia_data$Type]
plot(pr.out$x[,1],pr.out$x[,2],xlab="PC1", ylab="PC2",cex=0.5)
text(pr.out$x[,1],pr.out$x[,2],labels = leukemia_data$Type,col=plot_colors,cex=0.6)
```



T-ALL

is most clearly separated from the others along the PC1 axis

```
## create a dataframe contain Type label and numeric PC1 value
sort(abs(pr.out$rotation[,1]),decreasing = T)
```

##	SEMA3F	CCT2	LDHB	COX6C	SNRPD2
##	4.517148e-02	4.323818e-02	4.231619e-02	4.183480e-02	4.179822e-02
##	ELK3	SNRPE	LDHA	ATP5G3	CD40
##	4.155821e-02	4.034466e-02	4.005093e-02	4.004077e-02	3.995309e-02
##	40134_at	UQCRQ	CCT3	ITPR1	MDH1
##	3.977691e-02	3.960993e-02	3.953677e-02	3.947052e-02	3.940276e-02
##	NME1	UBA2	TBX19	SNRPG	KARS
##	3.900101e-02	3.872838e-02	3.869340e-02	3.856511e-02	3.836629e-02
##	32980_f_at	TPTEP1	PSMA4_1	YWHAQ_1	PGAM1
##	3.805177e-02	3.804482e-02	3.804281e-02	3.798229e-02	3.794595e-02
##	ERVH-1	EPOR	TUBGCP2	C1QBP	ERVK-7
##	3.790849e-02	3.783244e-02	3.762640e-02	3.739828e-02	3.733545e-02
##	SNRPA	HLA-DMB	ATXN10	HSPD1	SNRPB
##	3.721083e-02	3.720706e-02	3.720134e-02	3.717927e-02	3.713661e-02
##	MIF	40189_at	ITPR1_1	UQCRC1	SDC3
##	3.702705e-02	3.694271e-02	3.680088e-02	3.673105e-02	3.672759e-02
##	COX8A	KHDRBS1	MFSD10	SCARB1	TRA2A
##	3.669869e-02	3.658927e-02	3.654335e-02	3.653583e-02	3.647413e-02
##	FUCA1	SNRPF	32815_at	ISLR	PSMB2
##	3.645552e-02	3.631937e-02	3.628221e-02	3.621483e-02	3.621209e-02
##	AKAP17A	STAT2	TGOLN2	RAN_1	ITPR1_2
##	3.619071e-02	3.612710e-02	3.604629e-02	3.598788e-02	3.597252e-02
##	1937_at	INSR	TAZ	PAICS	RAN
##	3.586579e-02	3.559505e-02	3.558546e-02	3.546331e-02	3.541676e-02
##	OR2B6	UTRN	PRRC2B	CYCS	ATXN10_1
##	3.524901e-02	3.521147e-02	3.520911e-02	3.513441e-02	3.509530e-02
##	NCOA3	HNRNPA2B1	CCT5	FOXO1	CALCOCO1
##	3.506708e-02	3.500773e-02	3.499333e-02	3.481261e-02	3.481129e-02
##	ENO1	PAX8	39224_at	TRAFD1	EFNA3
##	3.479382e-02	3.474603e-02	3.470130e-02	3.460836e-02	3.460753e-02
##	SAP18	SPTA1	GPRIN2	HTRA2	TERF2_1
##	3.459465e-02	3.454154e-02	3.450462e-02	3.444213e-02	3.442021e-02
##	CLNS1A	TBC1D9	1627_at	PARK7	HLA-DOB
##	3.440591e-02	3.437814e-02	3.436168e-02	3.435885e-02	3.435284e-02
##	CTGF	TERF2	MEF2C	MC2R	HNRNPA3
##	3.426306e-02	3.426204e-02	3.421616e-02	3.419521e-02	3.419210e-02
##	ALDOA	MAP3K11	HSP90AB1	NUBP1	NPIP15
##	3.417207e-02	3.416858e-02	3.416834e-02	3.412728e-02	3.402794e-02
##	ELOC	FAM214B	SSRP1	NME1_1	COX6B1
##	3.402533e-02	3.400191e-02	3.398180e-02	3.393633e-02	3.392899e-02
##	SNAP23	32872_at	MYBPC1	MAN2B2	HHLA1
##	3.388626e-02	3.388353e-02	3.386933e-02	3.381239e-02	3.377224e-02
##	CBX3_1	JRK	VAMP1	JMJD1C	AP1B1
##	3.363399e-02	3.362428e-02	3.362081e-02	3.358154e-02	3.345921e-02
##	SNRPB_1	DDB1	36104_at	UBE3B	GAB1_1
##	3.342569e-02	3.342495e-02	3.342244e-02	3.338361e-02	3.337236e-02
##	HIST1H2BE	NCL	ITGA2B	SERPINH1	SRP9
##	3.334487e-02	3.333263e-02	3.332936e-02	3.329747e-02	3.323575e-02
##	IL6ST	31522_f_at	SLC9A1	HNRNPAB	GTF3A
##	3.319240e-02	3.319006e-02	3.313786e-02	3.306500e-02	3.306426e-02

##	ELOB	HGS	MMP25	SAP18_1	PHB2
##	3.305441e-02	3.301701e-02	3.300088e-02	3.297428e-02	3.289758e-02
##	HNRNPC	34093_at	ERG_1	CBFB	JUP
##	3.284812e-02	3.284460e-02	3.284416e-02	3.280753e-02	3.280603e-02
##	HLA-DMA	ANKRD12	33415_at	LARGE1	CBX3
##	3.280560e-02	3.280357e-02	3.280334e-02	3.279051e-02	3.276945e-02
##	NDUFAB1	GRM4	33877_s_at	COX4I1	HIST1H2BN
##	3.276198e-02	3.271671e-02	3.270819e-02	3.268690e-02	3.262753e-02
##	AGPAT2	PHF1	1863_s_at	HSP90AA1	CANX
##	3.262535e-02	3.261911e-02	3.258375e-02	3.254858e-02	3.254530e-02
##	ANKS1A	GAB1_2	STX4	TRIM38	31524_f_at
##	3.244427e-02	3.240494e-02	3.239847e-02	3.237471e-02	3.231921e-02
##	PSMD8	ADAM15	SORBS3	H2AFZ	ENTPD4
##	3.225074e-02	3.223970e-02	3.218296e-02	3.218036e-02	3.216745e-02
##	TUBB2B	CD79A_1	CREBBP	TOMM20	PTPRE_1
##	3.211944e-02	3.211783e-02	3.210417e-02	3.207028e-02	3.206887e-02
##	TXN	HINT1	PML	PTP4A3	ZBTB22
##	3.201973e-02	3.200449e-02	3.189542e-02	3.187461e-02	3.183007e-02
##	MAPK8IP3	1980_s_at	1116_at	LSM12	SP100
##	3.180405e-02	3.174908e-02	3.174410e-02	3.171880e-02	3.169851e-02
##	MYL12B	CNOT4	YWHAQ	DYRK2	B4GALT3
##	3.169733e-02	3.168838e-02	3.162117e-02	3.158349e-02	3.157068e-02
##	NDUFB3	MEF2C_1	SLC6A7	ISG20	HIST2H2BE
##	3.156023e-02	3.153240e-02	3.151027e-02	3.150542e-02	3.147975e-02
##	COX5B	286_at	RAB5B	CEP135	DICER1_1
##	3.147555e-02	3.147422e-02	3.141388e-02	3.138793e-02	3.136186e-02
##	38056_at	UPF2	CAMK2G	CD79A	41584_at
##	3.135876e-02	3.133870e-02	3.127091e-02	3.124424e-02	3.122555e-02
##	MVK	RGL2	SZT2	RUVBL2	LILRA2
##	3.120694e-02	3.120089e-02	3.116577e-02	3.113261e-02	3.108732e-02
##	TP53I11	MAST3	HLA-DRB4	CECR7	DHX9
##	3.107584e-02	3.106266e-02	3.101751e-02	3.099712e-02	3.099314e-02
##	MMP11	40637_at	PTPRE	TMEM63A	EIF3K
##	3.097817e-02	3.096998e-02	3.095880e-02	3.095021e-02	3.093357e-02
##	SLC30A1	GDI1	PSMA4	IDH2	VCL
##	3.091483e-02	3.089683e-02	3.087614e-02	3.085391e-02	3.085283e-02
##	KAT7	BRD2_2	HLA-DQA1	SUMO1	40219_at
##	3.085107e-02	3.082461e-02	3.082064e-02	3.080533e-02	3.075465e-02
##	CD22	32007_at	KLHL18	EIF2AK2_1	SCAF11_1
##	3.075162e-02	3.074671e-02	3.070493e-02	3.070029e-02	3.067632e-02
##	41324_g_at	LIMK2	RAD21	ACAP2	HIST1H2BM
##	3.063258e-02	3.058023e-02	3.057923e-02	3.052959e-02	3.049033e-02
##	KDM3B	IFI35	DHRS1	TNFRSF25	HLA-DQB1
##	3.047770e-02	3.047589e-02	3.047339e-02	3.045203e-02	3.042830e-02
##	EML3	PIK3CG	CCT8	GSTO1	COX5A
##	3.041371e-02	3.041011e-02	3.040109e-02	3.039306e-02	3.037135e-02
##	LYPLA2	RPL36AL	MT4	STRAP	TRA2B
##	3.037060e-02	3.035987e-02	3.035944e-02	3.035802e-02	3.029614e-02
##	SIVA1	GABBR1	HSP90AB1_2	ERH	SNX1
##	3.026037e-02	3.025465e-02	3.024065e-02	3.023390e-02	3.020061e-02
##	PABPC4	ATP5C1	EIF3M	POU2AF1	TFF3
##	3.018692e-02	3.018406e-02	3.016916e-02	3.011169e-02	3.010201e-02
##	MGAT1	CDK13_1	SWAP70	HADHA	HIST1H2BD
##	3.002319e-02	3.002039e-02	2.999056e-02	2.995640e-02	2.993410e-02

##	NCDN	XRCC5_1	CLOCK	ZFPL1	ZKSCAN1
##	2.993268e-02	2.990962e-02	2.990836e-02	2.989667e-02	2.988100e-02
##	PSMA3	39073_at	NUP210	GOT2	AHDC1
##	2.981852e-02	2.977529e-02	2.976635e-02	2.970330e-02	2.967800e-02
##	ADD2	TMED2	FBL	CALM1	LAIR1
##	2.967128e-02	2.964562e-02	2.963796e-02	2.960429e-02	2.960253e-02
##	HNRNPC_1	SLC25A3	CLCN7	MAP4_1	NDUFB8
##	2.958161e-02	2.957001e-02	2.954128e-02	2.948158e-02	2.947318e-02
##	KHNYN	MAN1A1	SDHD	38372_at	32609_at
##	2.946988e-02	2.946224e-02	2.943356e-02	2.941767e-02	2.937711e-02
##	EIF4B	PSMA2	TRANK1	37796_at	32408_s_at
##	2.933043e-02	2.931375e-02	2.924136e-02	2.921429e-02	2.921045e-02
##	THEMIS2	34902_at	GNG7	40428_i_at	POU4F1
##	2.920335e-02	2.920067e-02	2.919628e-02	2.919175e-02	2.918585e-02
##	C6orf120	GRIK5	40888_f_at	PPP3CC	SMAD1
##	2.913761e-02	2.911913e-02	2.911601e-02	2.911436e-02	2.910618e-02
##	UQCR11	VIM	HIST1H2BL	ATP50	ZNF592
##	2.908412e-02	2.908403e-02	2.907560e-02	2.905689e-02	2.905233e-02
##	ZNF253	EHD1	HDGF	WWP1	XRCC6
##	2.904505e-02	2.902728e-02	2.900365e-02	2.894589e-02	2.894129e-02
##	PSMA1	33261_at	WSB1	PPM1G	GAB1
##	2.893038e-02	2.890891e-02	2.888559e-02	2.888266e-02	2.886167e-02
##	CD79B	STK19	RPP38	ARAF_1	HMCES
##	2.882244e-02	2.882230e-02	2.879154e-02	2.877867e-02	2.876046e-02
##	STK38_1	CGRRF1	TUFM	39731_at	EIF3H
##	2.875776e-02	2.870746e-02	2.870522e-02	2.869285e-02	2.867543e-02
##	NF2	POLR2L_1	ERG_2	PDE4B	THOC2
##	2.867018e-02	2.866709e-02	2.866085e-02	2.864795e-02	2.864544e-02
##	E2F1	DNAJC9	RPS6KB2	TMX4	37590_g_at
##	2.862848e-02	2.862719e-02	2.857131e-02	2.856679e-02	2.856578e-02
##	GRB10	HLA-DPB1	ZEB2	NHP2	MEF2C_2
##	2.855155e-02	2.855021e-02	2.853226e-02	2.851440e-02	2.850160e-02
##	ILF2	PSMB7_1	MR1	35748_at	FEN1
##	2.850113e-02	2.848936e-02	2.848060e-02	2.847257e-02	2.847133e-02
##	EIF3I	PSMC1	TCEA1	RNF103	UBE2L3
##	2.844696e-02	2.842423e-02	2.838678e-02	2.838239e-02	2.831617e-02
##	SSBP1	SETDB1	MTF1	NELFB	MAPK3
##	2.827725e-02	2.824072e-02	2.823578e-02	2.820575e-02	2.820042e-02
##	CCT7	TMEM41B	N4BP2L1_1	MARCH6	CDV3
##	2.819659e-02	2.819054e-02	2.816489e-02	2.814679e-02	2.813785e-02
##	CTBP1_1	CDK14	COX5B_1	THUMPDI	INPP5D
##	2.807658e-02	2.806863e-02	2.805710e-02	2.804008e-02	2.803757e-02
##	33458_r_at	GAK	LGALS9	ELOVL5	PIGB
##	2.803654e-02	2.801649e-02	2.800190e-02	2.799894e-02	2.799851e-02
##	MFN1	TBCA	PRDM2	BAZ2A	ATIC
##	2.799764e-02	2.799239e-02	2.798597e-02	2.794148e-02	2.793698e-02
##	IFNGR1	VAMP3_1	TNFRSF10B	UQCRFS1	41723_s_at
##	2.791136e-02	2.790906e-02	2.789719e-02	2.785014e-02	2.784152e-02
##	COPS5	ITSN2	ERG	HECTD4	CD55
##	2.783617e-02	2.783490e-02	2.783078e-02	2.776934e-02	2.774627e-02
##	HAUS5	BTF3	HIST1H2BK	GRINA	NAP1L1_1
##	2.773525e-02	2.769519e-02	2.768977e-02	2.767472e-02	2.767366e-02
##	FBL_1	KDM5C	36112_r_at	IRF2	PPM1D
##	2.765746e-02	2.763402e-02	2.760489e-02	2.757110e-02	2.756641e-02

##	CHST15	ENG	PRMT1	CEP68	LSM3
##	2.756245e-02	2.754350e-02	2.754125e-02	2.752379e-02	2.751029e-02
##	GFPT1	SERBP1_1	WDR37	YTHDF3	PCBP1
##	2.747630e-02	2.745969e-02	2.743865e-02	2.742745e-02	2.741211e-02
##	40819_at	UBE2I_1	LONRF1	SGSM2	VDAC3
##	2.739961e-02	2.736180e-02	2.734686e-02	2.734194e-02	2.733157e-02
##	GSTP1_1	HLA-F	31993_f_at	PSMD5	EIF3I_1
##	2.731901e-02	2.729233e-02	2.728488e-02	2.726113e-02	2.720465e-02
##	38257_at	36696_at	INTS3	CBLB	NPY
##	2.717578e-02	2.717471e-02	2.716060e-02	2.714758e-02	2.707235e-02
##	LSM1	34841_at	RXRB	37693_at	SP110
##	2.706019e-02	2.705143e-02	2.704096e-02	2.701924e-02	2.700697e-02
##	CCR7	USP11	TCFL5	UQCRB	PGD
##	2.699691e-02	2.698267e-02	2.697585e-02	2.694286e-02	2.694140e-02
##	FCMR	PIK3C3	SFSWAP	CIAPIN1	BRMS1
##	2.693558e-02	2.693502e-02	2.692848e-02	2.692721e-02	2.688753e-02
##	CENPC	RUSC1	HLA-DQB1_1	SRSF3_1	HERC2P2
##	2.686403e-02	2.685667e-02	2.683988e-02	2.680568e-02	2.678769e-02
##	PPP1CC	MLXIP	ARHGEF2_1	CD200	WDR45B
##	2.678568e-02	2.676572e-02	2.674855e-02	2.674032e-02	2.673659e-02
##	SMAGP	TRIM28	SLC2A3	BAX	ADA
##	2.669752e-02	2.668617e-02	2.668350e-02	2.666344e-02	2.663957e-02
##	35858_at	DCAF7	CSNK2B	PRKCB_1	LSM2
##	2.663449e-02	2.663120e-02	2.663028e-02	2.662507e-02	2.661834e-02
##	1199_at	ATF4	ZMYND8	SIPA1	PCGF3
##	2.657579e-02	2.657263e-02	2.656036e-02	2.654374e-02	2.651851e-02
##	PPIP5K2	SKP1_1	DNAJC16	PXDN	SNRNP27
##	2.651747e-02	2.650764e-02	2.648159e-02	2.647562e-02	2.647141e-02
##	CDK10	IGSF9B	BTN2A1	40130_at	ZNF271P
##	2.646819e-02	2.646735e-02	2.644912e-02	2.642062e-02	2.641456e-02
##	XRCC5_2	1179_at	TMEM187	NDUFA1	SUB1
##	2.640007e-02	2.639983e-02	2.636661e-02	2.634962e-02	2.630018e-02
##	BTN3A3	XRCC5_3	FHIT	P2RX1	PTPN18
##	2.629714e-02	2.625346e-02	2.624532e-02	2.624451e-02	2.617313e-02
##	PRKY	HMHB1	GPR35	PIM1_1	UQCRC2
##	2.613415e-02	2.612383e-02	2.610526e-02	2.610476e-02	2.610306e-02
##	HHEX	ZNF85	TNK2_1	ATP5H	1294_at
##	2.610156e-02	2.609944e-02	2.608975e-02	2.602682e-02	2.601751e-02
##	IMPDH2	CLIP2	SEC62	MCM7	NCOA2
##	2.598271e-02	2.598033e-02	2.596832e-02	2.593725e-02	2.592948e-02
##	35961_at	PCLAF	ZBTB17	ALDOC	ENSA
##	2.592854e-02	2.592794e-02	2.586750e-02	2.585044e-02	2.584788e-02
##	POLD4	PSMB4	ARRB2_1	MFN2	2075_s_at
##	2.584283e-02	2.583863e-02	2.583467e-02	2.583380e-02	2.583204e-02
##	34791_at	TRAK1	NBN	AMFR	CSF1
##	2.582262e-02	2.581055e-02	2.580794e-02	2.580600e-02	2.580439e-02
##	CD19	RIPK1	HMGXB3	NUDC	SERINC5
##	2.579214e-02	2.578772e-02	2.576304e-02	2.573652e-02	2.571918e-02
##	FABP5	LPGAT1	36111_s_at	PLCG2_1	SF3B1
##	2.571488e-02	2.571459e-02	2.569155e-02	2.568766e-02	2.566046e-02
##	LY9	IRF9	EIF1B	PSMB4_1	PRKCB
##	2.563119e-02	2.559176e-02	2.557737e-02	2.553532e-02	2.552056e-02
##	KLF10_1	FDPS	SLC25A1_2	PEX19	PRKCB_2
##	2.550105e-02	2.548686e-02	2.546127e-02	2.545595e-02	2.544616e-02

##	KDM1A	SRSF1	PTP4A2_1	UGCG	REEP5
##	2.541152e-02	2.540751e-02	2.540482e-02	2.540054e-02	2.539809e-02
##	PTPA	NAE1	JTB_1	GOLGA1	HIST1H2AL
##	2.536565e-02	2.536368e-02	2.536259e-02	2.534063e-02	2.533801e-02
##	FCHSD2	C11orf58	SMARCD2_1	34279_at	ISCA1
##	2.530778e-02	2.530052e-02	2.527844e-02	2.526375e-02	2.525227e-02
##	IARS	TUBGCP4	WDR47	SERP1	VDAC2_1
##	2.524650e-02	2.524405e-02	2.523432e-02	2.523200e-02	2.523010e-02
##	AGPAT1	ACTN4	JMJD6	BANF1	GSK3A
##	2.521805e-02	2.520967e-02	2.519238e-02	2.518258e-02	2.518028e-02
##	POLB	PSMB1	STX7	RANBP3	STK38_2
##	2.515553e-02	2.515486e-02	2.514758e-02	2.513928e-02	2.513456e-02
##	BLCAP	AUTS2	ESD	NCF4	AZIN1
##	2.512026e-02	2.509691e-02	2.509594e-02	2.509573e-02	2.506611e-02
##	SND1	LCP1	HSPE1	CD27	SRRM1
##	2.506462e-02	2.505697e-02	2.505344e-02	2.504183e-02	2.503523e-02
##	41215_s_at	PSMD14	SMARCC2	CUL2	ASCC2
##	2.503144e-02	2.501843e-02	2.496665e-02	2.495103e-02	2.493460e-02
##	PRKCSH	BTF3_1	PDCD6	CAMTA2	KLHDC10
##	2.492223e-02	2.491142e-02	2.490785e-02	2.490400e-02	2.488331e-02
##	SLF2	PNRC1	PREP	TSC22D3_1	TAF11
##	2.488157e-02	2.485823e-02	2.485749e-02	2.485524e-02	2.484350e-02
##	ZMYND11	TNFRSF1B_1	PPP2R5E	NDUFB1	NCKAP1L
##	2.482846e-02	2.482442e-02	2.480979e-02	2.480763e-02	2.477379e-02
##	PTPN12	NRDC	ABI1_1	PRPF4B	POLE3
##	2.476273e-02	2.471478e-02	2.470925e-02	2.470644e-02	2.470397e-02
##	MAPKAP1	CRIM1	ZNF117	WBP1L	AP2S1
##	2.469778e-02	2.468275e-02	2.467498e-02	2.464053e-02	2.463583e-02
##	LRP10	PSMB3	TUBB4B	HADH_1	IRF7
##	2.463496e-02	2.463317e-02	2.462850e-02	2.462307e-02	2.456765e-02
##	TBC1D22A	MAP3K7	ARHGAP25	SLC25A11	IFFO1
##	2.453299e-02	2.453087e-02	2.452718e-02	2.452293e-02	2.451443e-02
##	ADA_1	ZDHHC17	MSH2	PRPF3	FBXW11
##	2.449840e-02	2.449261e-02	2.447758e-02	2.444061e-02	2.443030e-02
##	INPP5E	CTR9	CKS2	WASHC4	CDK2AP2
##	2.442792e-02	2.442277e-02	2.441975e-02	2.441283e-02	2.439586e-02
##	MAPRE1	NUDT3	ZNF266	STARD7	CASP7
##	2.438438e-02	2.437601e-02	2.436271e-02	2.434864e-02	2.434634e-02
##	PRKX	GPI	NMT1_1	SSR2	RASSF1
##	2.433130e-02	2.433113e-02	2.432926e-02	2.432284e-02	2.428856e-02
##	MAN2C1	41131_f_at	RCN2_1	BBX	ATP5D
##	2.426438e-02	2.424426e-02	2.423953e-02	2.420675e-02	2.417791e-02
##	NDUFV2	39126_at	DCK	NPC1	FAM189B
##	2.414075e-02	2.410325e-02	2.409725e-02	2.409141e-02	2.409102e-02
##	MTHFD2	COIL	DPM2	ATP5G1	OPTN
##	2.406253e-02	2.402622e-02	2.400973e-02	2.398858e-02	2.398121e-02
##	EEF1D	YY1	CEP350	LTA4H	NFKBIA
##	2.397531e-02	2.397165e-02	2.396967e-02	2.396122e-02	2.394565e-02
##	EDEM1	TNFRSF1B	POLR2I	FMR1_1	PTP4A2
##	2.394183e-02	2.391985e-02	2.387617e-02	2.383462e-02	2.382721e-02
##	SLC2A5	DAXX_1	LIG4	SFPQ_3	CCND2
##	2.382458e-02	2.382284e-02	2.382260e-02	2.381704e-02	2.381198e-02
##	TRAF5	CDK13	CUL1_1	TSPOAP1	TLE1
##	2.378712e-02	2.376044e-02	2.374689e-02	2.373242e-02	2.372887e-02

##	GRK2	NDUFV1	APEX1	MYL12A	NDUFC1
##	2.372441e-02	2.371432e-02	2.369002e-02	2.368849e-02	2.368782e-02
##	FTSJ1	BUB3	DENND4B	GHITM	AMD1_3
##	2.367326e-02	2.366234e-02	2.366116e-02	2.365668e-02	2.362117e-02
##	TSTA3	CDYL	ABCC5	COX7C	AKR1A1
##	2.358765e-02	2.356921e-02	2.356864e-02	2.355262e-02	2.354188e-02
##	HNRNPUL1	SNRPB2	DPT	FOXK2	RPL22_1
##	2.353163e-02	2.352524e-02	2.350104e-02	2.347567e-02	2.346766e-02
##	38698_at	H1FO	TNRC6B	TSC2	PPIB
##	2.344353e-02	2.344008e-02	2.342387e-02	2.341866e-02	2.340635e-02
##	PHB	STOM	N4BP2L2	MCM6	PTK2B_1
##	2.339091e-02	2.337445e-02	2.335914e-02	2.335520e-02	2.335469e-02
##	RFTN1	KDM6B	UBXN7	32921_at	ZNF91_1
##	2.334857e-02	2.334850e-02	2.333751e-02	2.332925e-02	2.332412e-02
##	EIF4H	CIC	CNIH1	YBX3	DNM2
##	2.332200e-02	2.332093e-02	2.331986e-02	2.330409e-02	2.327594e-02
##	CDK8	SECISBP2L	POLR2L	STK38	COX7A2
##	2.325544e-02	2.323071e-02	2.322486e-02	2.322229e-02	2.321652e-02
##	CD24	TRAF3IP2	MAPKAPK2_2	SOCS2	TRIM33
##	2.321162e-02	2.321061e-02	2.318480e-02	2.315294e-02	2.313963e-02
##	ZBED1	DHRS7	SLC9A3R1	PCNA	40765_at
##	2.313162e-02	2.309884e-02	2.308353e-02	2.308132e-02	2.306545e-02
##	SCML2	PCBP3	SEC61B	PPIE	GTF2A2
##	2.305360e-02	2.304270e-02	2.302262e-02	2.300337e-02	2.299260e-02
##	ATP2A2	CSRP2	LTBP4	TUBB4B_2	HDAC5
##	2.297819e-02	2.297459e-02	2.295989e-02	2.295668e-02	2.294863e-02
##	LPXN	TSPAN31	TARDBP	SEC61G	FIBP
##	2.294304e-02	2.292633e-02	2.291715e-02	2.291542e-02	2.291308e-02
##	38350_f_at	GSTP1	VAPA	PAPOLA	SREBF1
##	2.289648e-02	2.288871e-02	2.288725e-02	2.288095e-02	2.287958e-02
##	CITED2	ZDHHHC3	FCGRT	DDX19A	38710_at
##	2.287533e-02	2.287024e-02	2.285960e-02	2.285519e-02	2.284739e-02
##	ABCB7	RNF114	LEPROT	MYC	EZH1
##	2.284410e-02	2.284063e-02	2.283013e-02	2.281011e-02	2.280197e-02
##	HLA-J	SOS2	TERF1	37250_at	PRPF6
##	2.279677e-02	2.279461e-02	2.277712e-02	2.276468e-02	2.273690e-02
##	RBM14	JTB	TOMM70	NIPBL	WWOX
##	2.271567e-02	2.270295e-02	2.269036e-02	2.268247e-02	2.266994e-02
##	PHF8	POLR2C	SETD3	CSHL1_1	PRPF40A
##	2.266034e-02	2.262112e-02	2.260091e-02	2.259646e-02	2.259423e-02
##	ZNF91	32573_at	RPS15A	ZZZ3	NRIP1
##	2.258979e-02	2.256464e-02	2.255552e-02	2.254896e-02	2.254548e-02
##	PSMC3	NGDN	RBM4B	EIF3G	HSPB11
##	2.253523e-02	2.252359e-02	2.251773e-02	2.249486e-02	2.248569e-02
##	DOK1	VAV1	HNRNPL	USP12	WBP4
##	2.247372e-02	2.245390e-02	2.245283e-02	2.242069e-02	2.241852e-02
##	SUN2	HSBP1	SNX2	YWHAB	PTEN_1
##	2.241391e-02	2.235433e-02	2.231210e-02	2.231139e-02	2.230615e-02
##	GTF2H1	RNPS1	HPCAL1	HCCS	S1PR4
##	2.229956e-02	2.229633e-02	2.227238e-02	2.225260e-02	2.222789e-02
##	SLC25A5	MSL3	ZNF451	PIBF1	CHD9
##	2.222020e-02	2.221775e-02	2.221568e-02	2.220311e-02	2.219169e-02
##	PLCL2	NOLC1	ANP32A	BTK	SLC9A6
##	2.218416e-02	2.217676e-02	2.216645e-02	2.213903e-02	2.211541e-02

##	LSM4	PIM1	RPA3	AP1G2	TOMM34
##	2.209858e-02	2.207523e-02	2.203405e-02	2.203380e-02	2.201408e-02
##	CTNNA1	NUCB2	AP2M1	EIF4A3	GNE
##	2.200451e-02	2.197990e-02	2.196674e-02	2.195386e-02	2.195021e-02
##	C9orf16	EIF5B	36940_at	MARCH6_1	ANAPC5
##	2.194453e-02	2.194439e-02	2.192903e-02	2.192836e-02	2.192279e-02
##	SEC23IP	INTS1	HNRNPM	WASF1	EFR3A
##	2.191580e-02	2.190657e-02	2.190480e-02	2.190350e-02	2.188216e-02
##	CTNNA1_4	CLK3	CSNK1D_1	KRIT1	UBE2D3
##	2.188135e-02	2.187286e-02	2.186604e-02	2.180403e-02	2.179231e-02
##	PSMD6	TALD01	RBBP6	SETD1B	PSMD1
##	2.173668e-02	2.171329e-02	2.170252e-02	2.166056e-02	2.165902e-02
##	RCN2	RAG1	34099_f_at	NDUFS3	DUSP7
##	2.165767e-02	2.165116e-02	2.163969e-02	2.159689e-02	2.158658e-02
##	SSB	ADGRE5	SEM1	ANKLE2	ZNF43
##	2.157618e-02	2.157537e-02	2.156397e-02	2.155127e-02	2.154151e-02
##	ETFA	CCNT2	RFX5	MME	CTNNA1_3
##	2.152137e-02	2.150008e-02	2.149952e-02	2.149761e-02	2.148227e-02
##	KHSRP	DCAF8	STAT6_1	SHOC2	RAB3GAP1
##	2.147723e-02	2.145747e-02	2.143197e-02	2.143100e-02	2.142918e-02
##	ARPC4_1	OAS2	P4HA1	NMT1	ATM
##	2.141508e-02	2.140919e-02	2.140919e-02	2.140722e-02	2.137672e-02
##	BLCAP_1	LINC00847	ZC3H14	PKD2	PBXIP1
##	2.137089e-02	2.135146e-02	2.134954e-02	2.134533e-02	2.132405e-02
##	ECH1	EIF3B	ADCY6	REV3L	SLC35A1
##	2.132211e-02	2.128631e-02	2.128618e-02	2.127540e-02	2.126691e-02
##	NCF4_1	BICRAL	TGIF2	HIST1H2AC	40593_at
##	2.124731e-02	2.122607e-02	2.122312e-02	2.122018e-02	2.122015e-02
##	CASP4	TMX1	LMNB2	POLR3C	PMS2P1_1
##	2.119070e-02	2.118173e-02	2.117878e-02	2.117770e-02	2.116533e-02
##	LCK_1	USP34	SPTBN1	CTNNA1_2	TMEM243
##	2.115204e-02	2.114926e-02	2.113866e-02	2.113385e-02	2.109276e-02
##	ACTL6A	ADD3_1	CYC1	PLCB2	RXRA
##	2.106489e-02	2.105934e-02	2.105804e-02	2.104834e-02	2.104508e-02
##	CLASP1	FEN1_2	PLCG2	GTF2F1	ATP5A1
##	2.102953e-02	2.102635e-02	2.102526e-02	2.098775e-02	2.098425e-02
##	C14orf2	UBR2	TUBB4B_1	NARS	HIF1A
##	2.092134e-02	2.091075e-02	2.090765e-02	2.089333e-02	2.088513e-02
##	CD52	GMPS	SERBP1	E2F4	HNRNPU
##	2.087523e-02	2.087088e-02	2.086111e-02	2.084975e-02	2.083425e-02
##	DUT	JADE3	LYN	33361_at	PHF3
##	2.081229e-02	2.080762e-02	2.080520e-02	2.080242e-02	2.079014e-02
##	COPB2	H2AFY	FOXN3	CSNK1D	PPP1R15A
##	2.078693e-02	2.077324e-02	2.076448e-02	2.076118e-02	2.076106e-02
##	DDX39A	DYNC1H1	LSM14A_1	PIGA	CBFA2T3
##	2.075818e-02	2.075215e-02	2.074921e-02	2.070894e-02	2.068071e-02
##	ANP32B	SYK_2	GAB2	ACTR3	HLX
##	2.067334e-02	2.064618e-02	2.063417e-02	2.062987e-02	2.062137e-02
##	CLIC1	BUB3_1	PPIL2	CUL1	TMEM11
##	2.061588e-02	2.057508e-02	2.057200e-02	2.055843e-02	2.055522e-02
##	TXNIP	PRPSAP1	GMFB	LPIN1	COASY
##	2.052620e-02	2.051523e-02	2.051310e-02	2.050656e-02	2.048366e-02
##	35304_at	SLC4A2	TBL1X	PFDN4	GGCT
##	2.047313e-02	2.044886e-02	2.043044e-02	2.040453e-02	2.040288e-02

##	PIGF	PRKDC_1	CLSTN1	TSP02	SUN1
##	2.036473e-02	2.035871e-02	2.035628e-02	2.035261e-02	2.035051e-02
##	CNP	BOLA2	PPP1R12A	35566_f_at	SSR1_1
##	2.031520e-02	2.031324e-02	2.028088e-02	2.027595e-02	2.026519e-02
##	DHRS3	DVL3	P4HB_1	RAB1A_2	COR01A
##	2.025837e-02	2.024430e-02	2.024065e-02	2.023721e-02	2.023497e-02
##	ZMIZ1	41174_at	RAPGEF2	TRIM22	MBNL1
##	2.023434e-02	2.022927e-02	2.022632e-02	2.020907e-02	2.018662e-02
##	FMR1	DUSP6	SYNPO	GSN	ETS2
##	2.018300e-02	2.018024e-02	2.016268e-02	2.014916e-02	2.010719e-02
##	TAF5	ZNF254	DYRK3	CTCF	NAP1L1
##	2.009268e-02	2.009063e-02	2.008503e-02	2.007986e-02	2.007522e-02
##	HAX1	RPGR	SRRM2	U2AF1	NMT2
##	2.005568e-02	2.004156e-02	2.003758e-02	2.003253e-02	2.002875e-02
##	MTMR9	KBTBD2	AMD1	PHLPP1	DMTF1
##	2.001897e-02	1.996905e-02	1.996854e-02	1.995424e-02	1.994540e-02
##	COPE	NT5C2	LCK	GLS	PLXNB2
##	1.994327e-02	1.994281e-02	1.994249e-02	1.993758e-02	1.993263e-02
##	USP14	VAT1	PNMT	HDAC2	SCAF11
##	1.992705e-02	1.992576e-02	1.991129e-02	1.990344e-02	1.989023e-02
##	XRCC5	RBBP4_1	MDK	WDR18	CAD
##	1.987135e-02	1.985505e-02	1.985389e-02	1.985261e-02	1.985104e-02
##	HNRNPA1	RAPGEF2_1	PGK1_1	C1D	ACIN1
##	1.983832e-02	1.983756e-02	1.983431e-02	1.981970e-02	1.979138e-02
##	BRD1	RP2	FUS	PITPNM1	GAPVD1
##	1.978918e-02	1.978672e-02	1.977054e-02	1.974870e-02	1.973718e-02
##	GTF3C1	41288_at	PRKDC	36234_at	CSNK2A1
##	1.971667e-02	1.970613e-02	1.969950e-02	1.968884e-02	1.968047e-02
##	SLC35D1	VEZF1_1	UBXN1	ATG14	FAM120A
##	1.967025e-02	1.964332e-02	1.963215e-02	1.963065e-02	1.960533e-02
##	MAML1	MRPL3	CD53	DENND3	USP1
##	1.960287e-02	1.958909e-02	1.958616e-02	1.956706e-02	1.956249e-02
##	MYC_1	SMURF2	BCR_1	CERS6	PRKD2
##	1.955597e-02	1.954022e-02	1.953914e-02	1.952374e-02	1.951936e-02
##	36625_at	HPRT1	VAMP5	JARID2	CD69
##	1.951816e-02	1.949841e-02	1.949765e-02	1.949302e-02	1.948645e-02
##	GARS	GLUL	MORF4L2	MAPKAPK2_1	WDR1
##	1.947120e-02	1.947019e-02	1.946115e-02	1.945961e-02	1.944574e-02
##	GL01	ARHGDIB	CCL5	ERCC1_2	DDOST
##	1.941958e-02	1.941086e-02	1.941008e-02	1.939829e-02	1.937696e-02
##	GTF2A2_1	IFI44	EEF1E1	TCF3	YARS
##	1.937477e-02	1.936393e-02	1.934469e-02	1.929805e-02	1.929734e-02
##	PSMB7	MTHFD1_1	TNK2	DCTD	PPP1R10
##	1.927423e-02	1.926171e-02	1.922296e-02	1.920706e-02	1.920341e-02
##	QKI	POLR2I_1	DYNLL1	SLC39A8	HLA-E
##	1.919053e-02	1.917773e-02	1.912341e-02	1.910993e-02	1.910493e-02
##	TXLNA	KDELRL1_1	SS18	TRIM44_2	VTI1B
##	1.908720e-02	1.908693e-02	1.907500e-02	1.906722e-02	1.904953e-02
##	RERE	NT5C2_1	33500_i_at	H2AFX	LTB
##	1.904495e-02	1.903987e-02	1.903629e-02	1.903247e-02	1.902170e-02
##	BCLAF1	SKAP2	LSM7	SIK3	POLD2
##	1.901693e-02	1.898704e-02	1.898529e-02	1.897995e-02	1.895288e-02
##	EEF2	PURA_1	FANCG	HMGN3	KAT8
##	1.894848e-02	1.894846e-02	1.894092e-02	1.893504e-02	1.892890e-02

##	SOD1	NEDD8	RBM3	36135_at	TMED9
##	1.891086e-02	1.890687e-02	1.888421e-02	1.887311e-02	1.887282e-02
##	DEK	UBE3B_1	AKAP9	TUBGCP3	LOC155060
##	1.886432e-02	1.886134e-02	1.884488e-02	1.883438e-02	1.882635e-02
##	MX1	GYPB	TNFRSF14	FARSA	ATP5I
##	1.879082e-02	1.876906e-02	1.876898e-02	1.876862e-02	1.875743e-02
##	PKM	MAP3K4	SYK_1	NDUFB5	IDI1
##	1.875137e-02	1.874916e-02	1.874538e-02	1.870371e-02	1.867952e-02
##	PTP4A1	CAPRIN1	DRG1	RHEB	ODC1_1
##	1.867242e-02	1.867043e-02	1.867024e-02	1.866757e-02	1.865663e-02
##	CIZ1	ASMTL_1	PPP1CA	VDAC1_1	HSP90AB1_1
##	1.865102e-02	1.862226e-02	1.861103e-02	1.860980e-02	1.859583e-02
##	NSL1	RPL22	TERF1_1	CYC1_1	ATP6V0E2
##	1.859345e-02	1.858411e-02	1.858223e-02	1.857577e-02	1.856701e-02
##	PTGES3	RBM5_1	FH	SNRK	ARAF
##	1.854406e-02	1.852879e-02	1.852576e-02	1.851685e-02	1.851637e-02
##	1007_s_at	CTBP2	FKBP15	MKNK1	IKZF1
##	1.849452e-02	1.849379e-02	1.847801e-02	1.847243e-02	1.842799e-02
##	ODC1	38835_at	CAPNS1	GNA13	EIF3D
##	1.842571e-02	1.839024e-02	1.837466e-02	1.837411e-02	1.836199e-02
##	LSM6	40567_at	RPA2	TPI1	GNA13_1
##	1.835728e-02	1.834968e-02	1.832443e-02	1.830123e-02	1.829342e-02
##	35615_at	NNT	CNOT8	UVRAG	MEF2D
##	1.829323e-02	1.829215e-02	1.828677e-02	1.828216e-02	1.826109e-02
##	COPS6	TCIRG1	CD22_1	TUBB	TSC22D4
##	1.826013e-02	1.825607e-02	1.825318e-02	1.824454e-02	1.824314e-02
##	SH2B3	NREP	APOBEC3C	39077_at	33300_at
##	1.823969e-02	1.823248e-02	1.823234e-02	1.818371e-02	1.816918e-02
##	NDUFAF3	RSBN1	FEN1_1	CSHL1	NDUFAF1
##	1.816726e-02	1.816269e-02	1.813416e-02	1.812673e-02	1.812666e-02
##	IP07	TOPORS	CNPY3	CEBPZ_1	TSC22D3
##	1.812389e-02	1.808114e-02	1.808097e-02	1.805764e-02	1.804314e-02
##	FCGRT_1	KIAA0232	CNBP	EPRS	GRB2
##	1.803831e-02	1.801708e-02	1.800409e-02	1.800394e-02	1.800096e-02
##	HAGH	ARPC2	COQ9	CREB3L2	RPN2
##	1.799111e-02	1.798824e-02	1.798684e-02	1.798048e-02	1.797972e-02
##	BPHL	RAF1_1	DARS	STRN3	MPC2
##	1.797542e-02	1.796855e-02	1.796728e-02	1.796675e-02	1.796537e-02
##	CSK	NDUFS5	NR2C1	CLK2	SAR1A
##	1.796393e-02	1.796391e-02	1.795359e-02	1.794566e-02	1.793860e-02
##	MDC1	FOSB	BCR_2	JAK1_1	EIF1AX_1
##	1.793680e-02	1.791442e-02	1.789705e-02	1.789637e-02	1.787756e-02
##	CHKB	41219_at	KCNA5	SDHB	OS9
##	1.785581e-02	1.785034e-02	1.782914e-02	1.781241e-02	1.780915e-02
##	ARPC5	LRIG1	ATAD2B	CENPB	FRYL
##	1.778820e-02	1.778268e-02	1.776834e-02	1.776814e-02	1.773528e-02
##	IRF3	OARD1	RAC1_1	ZCCHC11	MCM3
##	1.771623e-02	1.771153e-02	1.769879e-02	1.769403e-02	1.769170e-02
##	RBM10	TMEM184B	TAGLN2	SUGP1	MLEC
##	1.767177e-02	1.766177e-02	1.766172e-02	1.765892e-02	1.765119e-02
##	P4HB	PLSCR1	EIF3E	MAP4	RALY
##	1.763264e-02	1.762488e-02	1.762089e-02	1.760606e-02	1.760212e-02
##	COX7A2L	BICD2	TRAPPC3	ICE1	ETS2_1
##	1.758813e-02	1.758404e-02	1.758389e-02	1.758152e-02	1.757501e-02

##	MXI1	VDAC2	MPRIP	N4BP2L2_1	TUBB3
##	1.756732e-02	1.755957e-02	1.755719e-02	1.755234e-02	1.755184e-02
##	33501_r_at	ZNF273	PRRC2A	AIP	PARP2_1
##	1.754339e-02	1.753001e-02	1.751679e-02	1.751312e-02	1.751227e-02
##	UBE2N_1	33499_s_at	E2F5	ATXN2L_1	SIAH1
##	1.750536e-02	1.749504e-02	1.748613e-02	1.743857e-02	1.742669e-02
##	810_at	TRIM44_1	AATF	IFT27	CRYBG1_1
##	1.740819e-02	1.739747e-02	1.739723e-02	1.739199e-02	1.738760e-02
##	STXBP3	40877_s_at	SYK	SMARCA2_1	DYNLT3
##	1.737496e-02	1.734641e-02	1.731468e-02	1.730608e-02	1.730452e-02
##	GCDH	TAP1	CUX1	DCUN1D4	SMAD2
##	1.728390e-02	1.727448e-02	1.727437e-02	1.727424e-02	1.727408e-02
##	NUP62	TUBA4A	SEPT2	SLC25A1_1	SRSF4
##	1.725735e-02	1.725554e-02	1.725521e-02	1.724622e-02	1.724281e-02
##	CHMP2A	GNG5	IDH3B	ATP13A3	HADHB
##	1.723999e-02	1.722281e-02	1.718641e-02	1.717163e-02	1.716403e-02
##	NCK1	SFPQ_2	33893_r_at	DDAH2	RAB8A
##	1.714013e-02	1.712234e-02	1.712088e-02	1.711566e-02	1.711321e-02
##	FTO	KMT2A	WSB2	TYMS_1	OCRL
##	1.710828e-02	1.710799e-02	1.710571e-02	1.708235e-02	1.707159e-02
##	SMCHD1	SSSCA1	TMSB10	RPRD2	RSL1D1
##	1.706136e-02	1.703767e-02	1.702931e-02	1.702099e-02	1.701922e-02
##	PTPN11	CASC3	MDK_1	ARID5A	ZNF217
##	1.701570e-02	1.701559e-02	1.699672e-02	1.699291e-02	1.698718e-02
##	TGIF1	TROVE2_1	LMO4	PTDSS1	CPSF4
##	1.698702e-02	1.698094e-02	1.697455e-02	1.697240e-02	1.696913e-02
##	ATP6VOD1	TMEM147	SEC22B	FLNB	RBL2_1
##	1.696505e-02	1.696011e-02	1.693624e-02	1.693531e-02	1.690188e-02
##	CD72	BZW1	ARHGEF18	TUBB4A	HLA-G_1
##	1.690149e-02	1.687320e-02	1.686569e-02	1.686519e-02	1.686412e-02
##	XBP1	PSAP	RPA3_1	PIN1	SLC25A6
##	1.684796e-02	1.684446e-02	1.683587e-02	1.683411e-02	1.683210e-02
##	RIPOR2_1	C1QL1	CLN3	RASSF2	EXTL3
##	1.681658e-02	1.679929e-02	1.679360e-02	1.677072e-02	1.676961e-02
##	PCDH9	CLTA	PCNA_1	SIGMAR1	GTF2E2
##	1.676629e-02	1.675980e-02	1.671582e-02	1.667363e-02	1.666780e-02
##	SMS	TRIM27	ILF3_2	HBP1	CAPG
##	1.665647e-02	1.664744e-02	1.661801e-02	1.657727e-02	1.656229e-02
##	RAB22A	PARP2	NBAS	METAP1	MRPS12
##	1.655734e-02	1.654604e-02	1.653558e-02	1.651076e-02	1.649710e-02
##	JAK1	CBX1	PAN2	MAP2K2	PIM2
##	1.648406e-02	1.643897e-02	1.643470e-02	1.643320e-02	1.643155e-02
##	SF3A3	CTDNEP1	LGALS9_1	HIP1R	KDM5A
##	1.642949e-02	1.641456e-02	1.641338e-02	1.640982e-02	1.640649e-02
##	S100A13	HSPA4	ITGA6	TNFRSF1A	SREK1
##	1.639934e-02	1.637122e-02	1.636947e-02	1.634846e-02	1.634458e-02
##	ZHX2	XBP1_1	RTN4	TPX2	GIT2
##	1.634081e-02	1.634068e-02	1.633553e-02	1.632940e-02	1.631293e-02
##	GPS2	KDELR2	CSF2RB	ACTN1_1	MORC3
##	1.630877e-02	1.629177e-02	1.628386e-02	1.626465e-02	1.625925e-02
##	TCF3_1	SMAD4_2	SQSTM1	RIPOR2	MAN2B1
##	1.625646e-02	1.624937e-02	1.624509e-02	1.623803e-02	1.623593e-02
##	442_at	31895_at	SELENOF	TRIM27_1	SDHC
##	1.623430e-02	1.621604e-02	1.618971e-02	1.618938e-02	1.618510e-02

##	NPRL2	MRFAP1L1	TEX261	DKFZP586I1420	SRM
##	1.618006e-02	1.614354e-02	1.614274e-02	1.613828e-02	1.611054e-02
##	ELF4	GGPS1	SPTLC1	PMS1	FARSA_2
##	1.610512e-02	1.610308e-02	1.609640e-02	1.608437e-02	1.607208e-02
##	SOX4	TTC37	POLE	RAE1_1	UBE2S
##	1.606942e-02	1.606604e-02	1.605335e-02	1.604456e-02	1.602275e-02
##	IQSEC1	RAP1A	WBP2	RTCB	STOML2
##	1.600931e-02	1.599867e-02	1.598672e-02	1.596406e-02	1.595564e-02
##	H2AFV_1	PFDN5	39722_at	ATP5L	IKZF1_1
##	1.595459e-02	1.594278e-02	1.594131e-02	1.592912e-02	1.591612e-02
##	GUCY1A3	CLDND1	RTF1	MTMR3	CCSER2
##	1.591534e-02	1.590963e-02	1.589463e-02	1.588056e-02	1.587175e-02
##	SMARCD2	IMPDH1	AMD1_1	PSEN1	PFKP
##	1.587140e-02	1.587018e-02	1.584250e-02	1.583705e-02	1.581475e-02
##	TPST2	PPP1R2_1	COMMD4	ZBED4	CLK1
##	1.581224e-02	1.580546e-02	1.578994e-02	1.577380e-02	1.575890e-02
##	MTA1_1	NR3C1	RNMT	BASP1	HLA-F_1
##	1.566451e-02	1.564796e-02	1.564515e-02	1.559126e-02	1.556535e-02
##	TAF2	SLC4A1	GADD45A	MARF1	33301_g_at
##	1.556121e-02	1.555838e-02	1.555189e-02	1.553488e-02	1.551817e-02
##	ARHGAP45	ATRX	PPM1A	PPM1F	CMTR1
##	1.551453e-02	1.551448e-02	1.551251e-02	1.551056e-02	1.550349e-02
##	MRPL49	PCF11	CRYBG1	PMS2P3	ARHGDIA
##	1.548968e-02	1.548180e-02	1.547109e-02	1.544269e-02	1.543781e-02
##	MYL6B	RING1	VPS72	PSMC2	TBCB
##	1.543458e-02	1.541970e-02	1.541882e-02	1.541162e-02	1.540525e-02
##	STX8	39434_at	AHCYL1	NDUFA2	37209_g_at
##	1.539096e-02	1.539040e-02	1.537416e-02	1.537023e-02	1.534928e-02
##	SMC1A	IL1B	BLNK	USP9X_1	U2SURP
##	1.531902e-02	1.530668e-02	1.530648e-02	1.529826e-02	1.529629e-02
##	RANBP1	MLLT11	RASA1	RAP1GAP2	VBP1
##	1.528710e-02	1.527724e-02	1.526639e-02	1.526364e-02	1.525519e-02
##	IL2RG	RABGAP1	MTOR	GRB2_1	CD47
##	1.523476e-02	1.523199e-02	1.522847e-02	1.522812e-02	1.522677e-02
##	VGLL4	ZNF266_1	ARIH2	FRMD4B	TAPBP
##	1.522343e-02	1.521962e-02	1.521631e-02	1.518770e-02	1.518046e-02
##	TPP1	DPF2	AKAP10	DCAF11	MAK16
##	1.517449e-02	1.516365e-02	1.516232e-02	1.514983e-02	1.514014e-02
##	ELF2	ATP11B	IK_1	RCAN1	DBN1
##	1.513307e-02	1.511667e-02	1.511397e-02	1.510339e-02	1.508549e-02
##	RAB2A	GNB2	SYNCRIP	MAN2A2	CDV3_1
##	1.506411e-02	1.505903e-02	1.505825e-02	1.505467e-02	1.505353e-02
##	C10orf10	GLT8D1	MTHFD1	ZFP36L2	PCMT1
##	1.504043e-02	1.501891e-02	1.501532e-02	1.500768e-02	1.500744e-02
##	MRE11	MPI	SUPT4H1	PSMB9	ZNF292
##	1.499922e-02	1.498715e-02	1.493893e-02	1.493672e-02	1.493421e-02
##	WIPF2	MRPL23	PHF2	GALNT1	NCOR2
##	1.493329e-02	1.491852e-02	1.491332e-02	1.490654e-02	1.489206e-02
##	PRCC	PSMA5	JAK1_2	TBCC	EIF2AK2
##	1.488801e-02	1.486710e-02	1.484448e-02	1.482235e-02	1.481849e-02
##	MCFD2	PAK1	PECAM1_1	POLR2J	LYL1
##	1.481759e-02	1.481126e-02	1.480702e-02	1.480437e-02	1.480258e-02
##	SPECC1L	ILK	MSL1	MCM3AP	38160_at
##	1.480071e-02	1.479718e-02	1.479295e-02	1.479102e-02	1.478818e-02

##	INPP5K	PDCD10	THUMPD1_1	AASDHPPT	CD34_1
##	1.477167e-02	1.476574e-02	1.474167e-02	1.472875e-02	1.471877e-02
##	ZC3H15	UGP2	IRF1	STX16	LPCAT1
##	1.470795e-02	1.469599e-02	1.469411e-02	1.468431e-02	1.467840e-02
##	ZAP70	LAMP2	WNK1	PTK2B	FOXO3
##	1.466166e-02	1.466074e-02	1.464694e-02	1.462763e-02	1.462203e-02
##	ATP5J	TAB2	MAEA	TOB1	UBE2C
##	1.457971e-02	1.457011e-02	1.455044e-02	1.454880e-02	1.452225e-02
##	FLT3	DCTD_1	IL10RB	SERINC3	CTCF_1
##	1.452085e-02	1.450482e-02	1.448834e-02	1.446704e-02	1.445265e-02
##	JOSD1	TYK2	ZFP36L1	BUD31	34532_at
##	1.444453e-02	1.442939e-02	1.440408e-02	1.436655e-02	1.435600e-02
##	PFDN1	TLE4	31526_f_at	EIF4G3	41224_at
##	1.434022e-02	1.433316e-02	1.431494e-02	1.429739e-02	1.429179e-02
##	LBR	ARHGAP1	TARS	PSIP1	HELZ
##	1.428703e-02	1.428077e-02	1.427981e-02	1.426145e-02	1.425118e-02
##	PAPD7	SLC39A14	SDCBP	769_s_at	TCOF1
##	1.424734e-02	1.423154e-02	1.422128e-02	1.420382e-02	1.419339e-02
##	NISCH	ENSA_2	DEGS1	LMO2	POLR2G
##	1.418243e-02	1.416907e-02	1.416380e-02	1.414583e-02	1.414250e-02
##	LAMTOR5	PTPN18_1	HDAC1	RNGTT	PITPNB
##	1.413791e-02	1.413165e-02	1.412353e-02	1.412001e-02	1.408540e-02
##	WIPI2	SYNGR2	MSH3	TRIM13	SELENOW
##	1.406054e-02	1.404913e-02	1.404637e-02	1.401437e-02	1.399695e-02
##	HMGNA_1	KAT6A	37339_at	MZF1	DNAJC8
##	1.399663e-02	1.394666e-02	1.394405e-02	1.394335e-02	1.393638e-02
##	HSPA9	VDAC1	PDHB	TMED10_1	MNT
##	1.393081e-02	1.392859e-02	1.391672e-02	1.391284e-02	1.390803e-02
##	39025_at	DDX1	MAP2K1	PPID	BAZ2B
##	1.389912e-02	1.389895e-02	1.388978e-02	1.388190e-02	1.388090e-02
##	PSME4	TMEM109	MTF2	CAPZA1	RPS4Y1
##	1.387172e-02	1.386935e-02	1.386507e-02	1.386037e-02	1.385833e-02
##	KXD1	PTPN6	KDM3A	BTBD2	DHX15
##	1.385807e-02	1.385611e-02	1.385414e-02	1.384376e-02	1.383923e-02
##	SFPQ	37864_s_at	MKRN1	ATP6AP2	SEC24B
##	1.383464e-02	1.382709e-02	1.382704e-02	1.382680e-02	1.381949e-02
##	DYNC1I2	SEPT7	SART1	FAM89B	IFI16
##	1.380005e-02	1.379695e-02	1.379668e-02	1.377831e-02	1.376366e-02
##	39969_at	MAP2K3	TRAPPC8	PGK1	B4GALT1
##	1.374313e-02	1.372256e-02	1.371909e-02	1.371458e-02	1.371292e-02
##	RAB1A	CREB1	PEBP1	ADSL	TOP2A
##	1.370539e-02	1.370422e-02	1.368666e-02	1.368361e-02	1.368037e-02
##	HCLS1	KAT6B	CYBA	DCP2	BTG2
##	1.366704e-02	1.365809e-02	1.364216e-02	1.363999e-02	1.363352e-02
##	TXNDC9	PMAIP1	SMARCE1	SAFB	ZFP36
##	1.361680e-02	1.361508e-02	1.358864e-02	1.357573e-02	1.356982e-02
##	HMGNA1	ANAPC13	UBQLN2	UBE2J1_1	ADD3
##	1.356795e-02	1.355673e-02	1.354671e-02	1.353000e-02	1.352168e-02
##	HMGB1	COX7B	SPCS2	SIK1	SRP19
##	1.351172e-02	1.349743e-02	1.349620e-02	1.349385e-02	1.349058e-02
##	PPP1R2	TNFAIP3	UPF3A	HMGB2	PPP2R5A
##	1.347003e-02	1.345413e-02	1.345332e-02	1.345291e-02	1.343806e-02
##	WAS	UBE2E3	MAP4_2	ANKRD46	CRLF3
##	1.342947e-02	1.341904e-02	1.340389e-02	1.336735e-02	1.335819e-02

##	PTPRA	ACADM	EEF1A2	BCL11A_1	TNFAIP8
##	1.335696e-02	1.335687e-02	1.333364e-02	1.332300e-02	1.331710e-02
##	FOXJ2	39879_s_at	STAT1_1	THAP11	WTAP
##	1.331433e-02	1.331038e-02	1.330729e-02	1.330470e-02	1.329666e-02
##	DGUOK	GANAB	TMEM123	TSR3	ARF4
##	1.328517e-02	1.328037e-02	1.327973e-02	1.326915e-02	1.325798e-02
##	MECP2	TOP2B_1	FLT3_1	LANCL1	BCL2
##	1.325569e-02	1.324225e-02	1.323425e-02	1.317843e-02	1.317813e-02
##	RPL15	N4BP2L1	PSMB6	TMED10	ARID5B
##	1.317225e-02	1.314585e-02	1.313571e-02	1.312321e-02	1.311466e-02
##	RSU1_1	ADAR	PMS1_1	TMEM131	MRPS12_1
##	1.309639e-02	1.307398e-02	1.306497e-02	1.306282e-02	1.303935e-02
##	PEX3	PRKCI	ZNF32	ADGRG1	37313_at
##	1.303926e-02	1.303880e-02	1.302759e-02	1.302313e-02	1.302261e-02
##	UBXN4	CTBP1	JUNB	HLA-G	HNRNPR
##	1.298985e-02	1.298953e-02	1.297657e-02	1.297079e-02	1.295145e-02
##	MEN1	RPS6KA1	DPYSL2	SLC25A1	RAB1A_1
##	1.294772e-02	1.294062e-02	1.293445e-02	1.293382e-02	1.292187e-02
##	TRIM26	33817_at	AKR1B1	DESI2	NONO
##	1.291865e-02	1.291559e-02	1.291419e-02	1.290125e-02	1.290036e-02
##	APRT	CLK1_1	RASGRP2_1	AHSA1	DFFA
##	1.289709e-02	1.288721e-02	1.288618e-02	1.288031e-02	1.286668e-02
##	UBE2M	DUSP1	ZWINT	DNTT	TRIM33_1
##	1.286622e-02	1.284587e-02	1.284260e-02	1.283501e-02	1.283420e-02
##	CHD8_1	39421_at	KPNA2	ACLY	CDK9
##	1.283076e-02	1.281739e-02	1.281351e-02	1.281295e-02	1.279966e-02
##	NUMA1_1	PTEN	LIMK2_1	39342_at	COPS3
##	1.279284e-02	1.277517e-02	1.275527e-02	1.274295e-02	1.273927e-02
##	EIF2S3_1	DECR1	MED13L	CEBPG	TSG101
##	1.273479e-02	1.273478e-02	1.271515e-02	1.268611e-02	1.267200e-02
##	35450_s_at	CD9	RFC4_1	YWHAH	JCHAIN
##	1.266272e-02	1.264867e-02	1.264584e-02	1.263381e-02	1.262749e-02
##	HCFC1	XPO1	PRPSAP2	36122_at	EIF1AX
##	1.262634e-02	1.262542e-02	1.260713e-02	1.257793e-02	1.257393e-02
##	GUSBP11	TGOLN2_1	CD58	PCMT1_1	LST1
##	1.256660e-02	1.255739e-02	1.254971e-02	1.254169e-02	1.254026e-02
##	ARPP19	RYBP	UBN1	MICU2	PUM2
##	1.253840e-02	1.253245e-02	1.252783e-02	1.250129e-02	1.249732e-02
##	CD37	ARL2	TCF25_1	SKP1	IGHD
##	1.248267e-02	1.246242e-02	1.246139e-02	1.245713e-02	1.244968e-02
##	SSR4	KIFAP3	CENPF	PARG	CSTB
##	1.244610e-02	1.243221e-02	1.243099e-02	1.242891e-02	1.242429e-02
##	UBE2E1	ALAS2	NOP2	GGA3	ZMYM2
##	1.242086e-02	1.239234e-02	1.238598e-02	1.237432e-02	1.237080e-02
##	IDH1	RRP1B	35351_at	UBAC1	40951_at
##	1.236816e-02	1.234852e-02	1.233504e-02	1.233274e-02	1.232823e-02
##	C2CD5	RBM5	SSBP1_1	RAB9A	MBD2
##	1.228093e-02	1.226852e-02	1.226151e-02	1.225631e-02	1.225514e-02
##	HEBP2	FADS3	41407_at	NFRKB	ZNF337
##	1.225445e-02	1.224853e-02	1.224259e-02	1.223430e-02	1.222836e-02
##	TGFB1	38194_s_at	CDK1	DDB2	TRBC1
##	1.221808e-02	1.221532e-02	1.219751e-02	1.218279e-02	1.215715e-02
##	RASGRP2	G3BP1	FOS	FAM8A1	CKS1B
##	1.215273e-02	1.215156e-02	1.214655e-02	1.214181e-02	1.213662e-02

##	CYFIP2	DRAP1	POLR2A	SRP14	MGRN1
##	1.213533e-02	1.211141e-02	1.207797e-02	1.204205e-02	1.203420e-02
##	CLEC11A	IL23A	PTPN2	MGAT5	LSM14A
##	1.203373e-02	1.201382e-02	1.200973e-02	1.199807e-02	1.199031e-02
##	PTPRC	TECR	38709_at	ALG3	PSMD12
##	1.198676e-02	1.198481e-02	1.196671e-02	1.196555e-02	1.196399e-02
##	TNP03	STIP1	RABEP1	GLUD1_1	HMGXB4
##	1.196302e-02	1.195846e-02	1.195586e-02	1.194716e-02	1.194650e-02
##	PUF60	BIN1	ANXA7	DNAJB1	DEXI
##	1.193643e-02	1.187837e-02	1.187454e-02	1.186852e-02	1.185828e-02
##	TOR1A	PSMC5	TCF7	BRD3	CHD1
##	1.185112e-02	1.184961e-02	1.184804e-02	1.183316e-02	1.180104e-02
##	GNB1	VAT1_1	GUSB	ATRX_1	CDC37
##	1.179062e-02	1.178993e-02	1.178908e-02	1.178460e-02	1.178022e-02
##	31800_at	SNRPC	KLC1	RHOA	GABARAPL2
##	1.177450e-02	1.177257e-02	1.176257e-02	1.176160e-02	1.176018e-02
##	PSMD4	OXSRI	PRDX2	LTN1	PIGC
##	1.175081e-02	1.173565e-02	1.173522e-02	1.172652e-02	1.172513e-02
##	TM9SF2	CHD8	34677_f_at	MED13	CLTB
##	1.170933e-02	1.170552e-02	1.170290e-02	1.169341e-02	1.167881e-02
##	PCNT	PRPS2	TMC6	TFDP2	955_at
##	1.167081e-02	1.166908e-02	1.166128e-02	1.164559e-02	1.164361e-02
##	35736_at	ARPC4	EHMT2	NR3C1_1	MEF2A
##	1.163068e-02	1.162267e-02	1.159886e-02	1.157755e-02	1.157368e-02
##	FOS_2	NCAPD3	NSF	RUBCN	ZNF146
##	1.156696e-02	1.156131e-02	1.154662e-02	1.154313e-02	1.153431e-02
##	SEC13	37264_at	CNOT1	CTSC	KIF11
##	1.152910e-02	1.152434e-02	1.152203e-02	1.151673e-02	1.151213e-02
##	CSNK1A1	GNL2	RELA_2	GPSM3	TCEAL1
##	1.149784e-02	1.148094e-02	1.147246e-02	1.146645e-02	1.145863e-02
##	ALAS1	38019_at	NAB1	TAF11_1	IGF2R
##	1.144114e-02	1.142232e-02	1.141947e-02	1.140908e-02	1.140429e-02
##	SUSD6	H2AFV	FAM193A	EIF4EBP2	LYPLA1
##	1.134544e-02	1.133082e-02	1.131948e-02	1.129414e-02	1.127638e-02
##	C6orf62	CCT6A	SEC23A	TOP2A_2	NCSTN
##	1.126631e-02	1.126628e-02	1.125103e-02	1.122992e-02	1.121914e-02
##	TADA3	FAM208A	ITGA4	GABARAP	NDUFS2
##	1.121674e-02	1.121479e-02	1.119851e-02	1.118342e-02	1.118292e-02
##	RBCK1_1	TAF10	CRYZ	LYN_1	943_at
##	1.115402e-02	1.114437e-02	1.114427e-02	1.113442e-02	1.113059e-02
##	CDC123	FAF2	NFYC	NEK9	POLR2F
##	1.112906e-02	1.112015e-02	1.108997e-02	1.108780e-02	1.106675e-02
##	RAB11A	SSR1	MARK3	SRSF6	NR3C1_2
##	1.106654e-02	1.106592e-02	1.106417e-02	1.105359e-02	1.104096e-02
##	CFDP1	STAG2	VEZF1	ENTPD6	PHF21A
##	1.102215e-02	1.100879e-02	1.099615e-02	1.098838e-02	1.095439e-02
##	IP05	PPP1CA_1	EAPP	OGFR	STK10
##	1.094662e-02	1.094582e-02	1.094518e-02	1.093104e-02	1.091367e-02
##	SAT1	PLIN3	ARHGAP1_1	AKAP11	RSRC2
##	1.091228e-02	1.090813e-02	1.090353e-02	1.089161e-02	1.088292e-02
##	PECAM1	MIA2	SETX	FHL1	SEC31A
##	1.087369e-02	1.087046e-02	1.086593e-02	1.086381e-02	1.086281e-02
##	ADAM10_1	CELF1	NXF1	SLC25A44	PRKDC_2
##	1.084676e-02	1.084163e-02	1.082624e-02	1.081748e-02	1.081633e-02

##	NKG7	NFYA	IL27RA	MMS19	HMGA1
##	1.081349e-02	1.081193e-02	1.080609e-02	1.080264e-02	1.079312e-02
##	OFD1	RABGGB	DNAJC7	TOP2B	RAC2
##	1.078938e-02	1.077030e-02	1.076187e-02	1.074836e-02	1.073974e-02
##	AMOT	SERPINB1	193_at	DAP3	ATXN2
##	1.073682e-02	1.072406e-02	1.072150e-02	1.071220e-02	1.071102e-02
##	NCOA1	CDK2AP1	SARS	FYN_1	TRA2A_1
##	1.070635e-02	1.070516e-02	1.070478e-02	1.070037e-02	1.069544e-02
##	DDX21	SMARCC1	FOS_1	TROVE2	USP33
##	1.069115e-02	1.069085e-02	1.068831e-02	1.068119e-02	1.067692e-02
##	ASAH1	TCEAL4	ATP6V1G1	RBM6	BRD2_1
##	1.066595e-02	1.066544e-02	1.066160e-02	1.064780e-02	1.062303e-02
##	CS	38445_at	RBMS1_1	CYTH2	NFATC2IP
##	1.061123e-02	1.059880e-02	1.059685e-02	1.059176e-02	1.058656e-02
##	MX2	HNRNPK	TSFM	E2F3	AURKB
##	1.056497e-02	1.056361e-02	1.056143e-02	1.056047e-02	1.055767e-02
##	RECQL	YWHAZ	MTMR1	ME2	N4BP2L1_2
##	1.050982e-02	1.049954e-02	1.049342e-02	1.046679e-02	1.046564e-02
##	KIAA0922	APOBEC3G	RPA1	USP5	NASP
##	1.045077e-02	1.043187e-02	1.043118e-02	1.041567e-02	1.041498e-02
##	GRAMD4	PLPBP	PDHX	ID3	BLVRB
##	1.041251e-02	1.041197e-02	1.041121e-02	1.040787e-02	1.040109e-02
##	MRPS18B	ERCC3	PIP4K2A	MPHOSPH9_1	AKAP17A_1
##	1.038981e-02	1.038229e-02	1.036215e-02	1.035023e-02	1.032592e-02
##	KRT8	UTP3	YTHDC1	EIF2S3	TOPBP1
##	1.031396e-02	1.030320e-02	1.028343e-02	1.025345e-02	1.024458e-02
##	1158_s_at	RCOR1	CD44_1	PMS2P3_1	MDC1_1
##	1.023428e-02	1.023392e-02	1.021753e-02	1.021688e-02	1.016872e-02
##	PNISR	LY86	IST1	ZFC3H1	VPS52
##	1.015866e-02	1.015076e-02	1.013340e-02	1.013245e-02	1.013240e-02
##	AP3D1_1	FSCN1	PRPF8	TAF6L	BRE
##	1.012876e-02	1.012804e-02	1.012346e-02	1.011548e-02	1.011483e-02
##	FLNA	ARMCX6	CTNNA1_1	IGHM	STAM
##	1.010821e-02	1.009885e-02	1.009791e-02	1.009133e-02	1.009030e-02
##	UBE3A	RUVBL1	PSMD9	40552_s_at	BTN3A1
##	1.007746e-02	1.007688e-02	1.006630e-02	1.006557e-02	1.005379e-02
##	GPX1	NPAT	KDM2A	RIF1	GCN1
##	1.005053e-02	1.004753e-02	1.004618e-02	1.004031e-02	1.003407e-02
##	KDM5D	ZMYM4	STK39	VAMP2	CAT
##	1.003188e-02	1.000229e-02	9.999490e-03	9.996970e-03	9.995180e-03
##	TP53BP1	UNG	BIN1_1	ZMYM3	TOP1P2
##	9.994479e-03	9.986613e-03	9.962996e-03	9.962339e-03	9.945497e-03
##	SMAD2_1	UBE2S_2	SLBP	NIPSNAP1	WEE1
##	9.932878e-03	9.925123e-03	9.905697e-03	9.902963e-03	9.902306e-03
##	DAAM1	CALU	KCNAB2	ROCK1	NDUFB7_1
##	9.900186e-03	9.898942e-03	9.886441e-03	9.885384e-03	9.867490e-03
##	UBR5_1	MOB1A	IGHV5-78	CHTOP	32004_s_at
##	9.845516e-03	9.834891e-03	9.822859e-03	9.819821e-03	9.808490e-03
##	UGDH	PIK3C2B	PTPRC_2	GPAA1	SPOP
##	9.798553e-03	9.795352e-03	9.770973e-03	9.757316e-03	9.727563e-03
##	TYMS	SEC24C	IER2	BST2	SHC1
##	9.727450e-03	9.726091e-03	9.713487e-03	9.692563e-03	9.686242e-03
##	KIAA0100	MOAP1	NME3	SFN	RPS6KA3_1
##	9.686237e-03	9.682366e-03	9.678139e-03	9.677161e-03	9.676169e-03

##	AHCYL1_1	LDLRAD4	GPN1	DNPEP	PUM1
##	9.665084e-03	9.648507e-03	9.646888e-03	9.642146e-03	9.635598e-03
##	FMNL1	ASXL1	BPTF	MBTPS1	HSPB1
##	9.619781e-03	9.615926e-03	9.609441e-03	9.600967e-03	9.600653e-03
##	ARL6IP1	DHFR	MUTYH	CRY1	AES
##	9.585380e-03	9.582466e-03	9.556019e-03	9.555748e-03	9.553298e-03
##	LITAF_1	BRD8_1	MDN1	GLRX	CABIN1
##	9.547203e-03	9.542317e-03	9.535444e-03	9.530769e-03	9.512272e-03
##	ST6GAL1	UFD1	RBM6_1	GNL1	CD34
##	9.487854e-03	9.463228e-03	9.463052e-03	9.456389e-03	9.452583e-03
##	PDE6D	ARFGEF1	HNRNPDL_1	TBCE	PHIP
##	9.449683e-03	9.447858e-03	9.446177e-03	9.432176e-03	9.426914e-03
##	SRF_1	RNF167	32547_at	EID1	GM2A
##	9.414007e-03	9.411801e-03	9.381769e-03	9.367784e-03	9.361196e-03
##	API5_1	CNPPD1	UBE2S_1	DPM1	LARP4
##	9.358840e-03	9.354192e-03	9.329798e-03	9.318301e-03	9.316380e-03
##	CTNNB1	MAPKAPK2	TSC22D1	33414_at	LAPTM4A
##	9.312856e-03	9.312366e-03	9.293634e-03	9.286361e-03	9.282198e-03
##	TOP2A_1	SERTAD2	PRKAR1A_2	DDX3X_1	SBF1
##	9.269513e-03	9.256706e-03	9.253387e-03	9.251435e-03	9.251325e-03
##	CYB5A	CAMKK2	DSCR3	SENP3	CYFIP1
##	9.245213e-03	9.231951e-03	9.225792e-03	9.222666e-03	9.219886e-03
##	NDUFS4	CYB5B	SLC7A1	PRKCH	FUBP1
##	9.219173e-03	9.214994e-03	9.207465e-03	9.203659e-03	9.196553e-03
##	UPF1	LSM7_1	RAC2_1	GNA15	SCP2
##	9.187446e-03	9.176945e-03	9.167610e-03	9.159811e-03	9.156806e-03
##	SPEN	MED12	BCAP31	CD63	NDC80
##	9.153363e-03	9.147649e-03	9.141152e-03	9.140615e-03	9.134094e-03
##	CDK4	HNRNPH3	AFF1	STX16_1	SARAF
##	9.126460e-03	9.110108e-03	9.099745e-03	9.079862e-03	9.061261e-03
##	RALBP1	PBX3	TAF15	DEAF1	KRAS
##	9.060219e-03	9.027777e-03	9.020802e-03	9.019189e-03	9.008227e-03
##	ARIH1	ARHGAP4	URI1	ARRB2	LCP2
##	9.000411e-03	8.992381e-03	8.987060e-03	8.971077e-03	8.967054e-03
##	TRAF3IP3_1	SLC25A46	SATB1	MAPK6	MRPL9
##	8.966625e-03	8.956032e-03	8.952910e-03	8.935332e-03	8.929069e-03
##	RABAC1	ADAM10	SUM01_1	MSH2_1	TACC1
##	8.923979e-03	8.919201e-03	8.915481e-03	8.912618e-03	8.906580e-03
##	31444_s_at	RBCK1	NCOA4	SRSF5	TCF3_2
##	8.901288e-03	8.884275e-03	8.865711e-03	8.857914e-03	8.849417e-03
##	MANF	METTL3	PRRC2C	DHX38	UCKL1
##	8.842211e-03	8.804450e-03	8.804378e-03	8.797676e-03	8.796545e-03
##	USP24	ZNF428	36158_at	ZNF22	SUPT5H
##	8.791375e-03	8.787748e-03	8.784896e-03	8.754319e-03	8.751007e-03
##	SLC1A5	GNS	TRIM37	ERP29	CD44
##	8.749522e-03	8.747527e-03	8.741930e-03	8.741033e-03	8.738893e-03
##	RER1_1	CLEC16A	SDR39U1	TUSC2	SERINC1
##	8.732706e-03	8.731221e-03	8.727938e-03	8.720534e-03	8.711608e-03
##	NDRG1	SEC63	EIF1	RBPJ	EIF2B2
##	8.702577e-03	8.697542e-03	8.694713e-03	8.690925e-03	8.686339e-03
##	SEPT6	VEGFB	HNRNPA0	EIF2S1	HIVEP2
##	8.680785e-03	8.680314e-03	8.677014e-03	8.675326e-03	8.616208e-03
##	35292_at	DDHD2	NFKB1_1	LYRM1	TGFBR2
##	8.613047e-03	8.597997e-03	8.595840e-03	8.581814e-03	8.580699e-03

##	CCDC28A	TRIB2	AP3S1	MGA	PRKAG1
##	8.571304e-03	8.568333e-03	8.556435e-03	8.553714e-03	8.540963e-03
##	CAP1	SDHA	UBE2G2	PPP2R1A	PLXND1
##	8.523436e-03	8.512014e-03	8.504199e-03	8.498144e-03	8.485586e-03
##	CHD4	CD164	NUP93	CNPY2	BAG5
##	8.479081e-03	8.478104e-03	8.475796e-03	8.471537e-03	8.462504e-03
##	STAT1	CLIC4	EVI2A	UTP18	RGS19
##	8.461452e-03	8.452686e-03	8.451115e-03	8.449697e-03	8.436599e-03
##	PRKAR1A	SACS	TNFAIP2	MEGF9	ACBD3
##	8.433227e-03	8.427863e-03	8.425462e-03	8.422338e-03	8.415847e-03
##	NUP188	33987_at	SF3B2	GDI2	IGF2BP3
##	8.407784e-03	8.403602e-03	8.402602e-03	8.395878e-03	8.390248e-03
##	HAT1	TMEM59	SPN	MTMR4	CEP57
##	8.378495e-03	8.371860e-03	8.364372e-03	8.361053e-03	8.358256e-03
##	DIAPH1	MAPRE2	YWHAE	31519_f_at	FOXM1
##	8.346396e-03	8.331205e-03	8.326622e-03	8.301094e-03	8.294305e-03
##	SEPT6_2	PICALM	RIOK3	FNTA	TRRAP
##	8.289832e-03	8.286817e-03	8.246990e-03	8.235005e-03	8.232219e-03
##	MPP1	ATP6VOE1	SART3	TCF25	ELAC2
##	8.230598e-03	8.226979e-03	8.224240e-03	8.215025e-03	8.173970e-03
##	PTTG1IP	PTPRCAP	COA1	PFKL	DAD1
##	8.172686e-03	8.171272e-03	8.169806e-03	8.167156e-03	8.157663e-03
##	MYD88	SMAD4	TMEM106C	STAMPB	IQGAP2
##	8.152430e-03	8.134745e-03	8.127522e-03	8.119933e-03	8.115892e-03
##	RFC5	HACD2	TRAM1	MGST2	DBI
##	8.114506e-03	8.113385e-03	8.108294e-03	8.107213e-03	8.103145e-03
##	BCR	SASH3	EIF2S2	CCNH	MYB
##	8.090003e-03	8.087959e-03	8.083113e-03	8.080267e-03	8.059864e-03
##	SMG7	ADNP	PPP4R1	SLC7A5	KIAA1109
##	8.056391e-03	8.054022e-03	8.050226e-03	8.045123e-03	8.027160e-03
##	CDC7	VCP	DHX9_1	IARS2	MPHOSPH6
##	8.019138e-03	8.012448e-03	8.003192e-03	8.001011e-03	7.994259e-03
##	TRAF3IP3	SRSF6_1	CD46	HUWE1	LARP1
##	7.992645e-03	7.987862e-03	7.971625e-03	7.963704e-03	7.947263e-03
##	SNW1	DHX16	AZIN1_1	TDG	MPHOSPH9
##	7.945491e-03	7.942332e-03	7.917652e-03	7.911616e-03	7.908819e-03
##	FLI1	FNTA_1	GLOD4	32878_f_at	FKBP1A_1
##	7.892524e-03	7.873933e-03	7.862207e-03	7.849024e-03	7.846110e-03
##	PARN	PRPF40A_1	MPST	TK1_1	GLUD1
##	7.846024e-03	7.841301e-03	7.828926e-03	7.794959e-03	7.785451e-03
##	AP3B1	CHMP1A	SDF2	CELF2	GNPAT
##	7.784183e-03	7.784065e-03	7.774139e-03	7.768400e-03	7.765025e-03
##	NUP205	GOSR1	TIMM17A	NPC2	NME4
##	7.733177e-03	7.709275e-03	7.703709e-03	7.701782e-03	7.684842e-03
##	BLMH	MTMR6	KLF10	PKMYT1	PKN1
##	7.682892e-03	7.662870e-03	7.662508e-03	7.656176e-03	7.654501e-03
##	KLHL9	BTG1	POLR2B	CDC16	ERCC1
##	7.641062e-03	7.629914e-03	7.625629e-03	7.619415e-03	7.617931e-03
##	TSNAX	POP5	41143_at	IGHM_3	PSMD9_1
##	7.616700e-03	7.609957e-03	7.608577e-03	7.606283e-03	7.589332e-03
##	STIM1	TRIP12	TTK	ARHGEF6	EIF5
##	7.587709e-03	7.582021e-03	7.573785e-03	7.557458e-03	7.552120e-03
##	PCBP2	GUK1	PRMT2	NFKB1	1106_s_at
##	7.548660e-03	7.546468e-03	7.535913e-03	7.533628e-03	7.529241e-03

##	ELM01	PIK3R3	AP3D1	KIF2A	EDC4
##	7.515728e-03	7.514207e-03	7.502749e-03	7.502334e-03	7.496816e-03
##	SAMM50	ISG15	MLX	RAB8A_1	DDX3Y
##	7.478293e-03	7.474476e-03	7.474472e-03	7.471803e-03	7.468825e-03
##	SEPT9	SMAD4_1	DDX11	S100A10	MGEA5
##	7.460737e-03	7.458650e-03	7.454037e-03	7.452619e-03	7.451022e-03
##	MAP1LC3B	VAMP7	ELF1	KTN1	SYPL1
##	7.449272e-03	7.446413e-03	7.443726e-03	7.441885e-03	7.438950e-03
##	TSPAN3	DOCK2	FKBP4	ACOT13	NAP1L4
##	7.437368e-03	7.435164e-03	7.429769e-03	7.424601e-03	7.415829e-03
##	TNP03_1	41258_at	IFITM2	TFDP1	ARL2BP
##	7.402528e-03	7.397381e-03	7.395616e-03	7.385873e-03	7.381824e-03
##	ATP6V1B2	AMD1_2	VOPP1	XPC	SF3B4
##	7.374032e-03	7.355555e-03	7.338052e-03	7.333067e-03	7.311525e-03
##	TPD52L2	TRIP13	AIF1	AK2	RSU1
##	7.311161e-03	7.300195e-03	7.281337e-03	7.267725e-03	7.255289e-03
##	SRSF1_1	CCNG2	CHERP	SRGN	RPA1_1
##	7.250753e-03	7.250375e-03	7.245321e-03	7.242639e-03	7.218968e-03
##	CRADD	AKR7A2	DNAJA1_1	CAPZA2	TMEM106C_1
##	7.215463e-03	7.212157e-03	7.204607e-03	7.200938e-03	7.189144e-03
##	ZNHIT3	GPX4	NCOA6	FUBP1_1	IFITM1
##	7.184750e-03	7.184660e-03	7.180742e-03	7.157558e-03	7.149868e-03
##	COPS2	C5orf15	USP7	NUP153	RB1
##	7.145441e-03	7.134440e-03	7.130062e-03	7.112829e-03	7.076529e-03
##	KAT2A	HSF1	PTPRC_1	NADK	HERPUD1
##	7.074207e-03	7.060796e-03	7.060578e-03	7.058966e-03	7.028136e-03
##	CALCOCO2	34647_at	EIF3F	KIAA0907	NPEPPS
##	7.014570e-03	7.004160e-03	7.000974e-03	6.991801e-03	6.987802e-03
##	RMND5A	XP07	MYH10	CDKN1B	CIRBP
##	6.982398e-03	6.934736e-03	6.930399e-03	6.915050e-03	6.910793e-03
##	NFE2L2	DUSP11	MAT2A	CD3E	POLR2H_1
##	6.906716e-03	6.896567e-03	6.895086e-03	6.881783e-03	6.858806e-03
##	EZH2	IL10RB_1	PRKRA	STMN1	RAP1B_1
##	6.852470e-03	6.851515e-03	6.841970e-03	6.837188e-03	6.832545e-03
##	HNRNPH1	UBE2D3_1	PRDX6	TCF12	AKT1
##	6.801118e-03	6.790878e-03	6.787321e-03	6.785324e-03	6.780680e-03
##	MYC_2	SCAF8	MYBL2	NFATC3_2	41733_at
##	6.774701e-03	6.771999e-03	6.771895e-03	6.769481e-03	6.762789e-03
##	41243_at	ATP6AP1	MAD2L1	MAGED1	CD81
##	6.755067e-03	6.754532e-03	6.743998e-03	6.740799e-03	6.714017e-03
##	TLN1	TBCD	TKT	BRD3_1	TFRC
##	6.705881e-03	6.695087e-03	6.673184e-03	6.671704e-03	6.669872e-03
##	POLDIP3	PSMD11	33836_at	RBBP8	ETHE1
##	6.649040e-03	6.630869e-03	6.629059e-03	6.627731e-03	6.626915e-03
##	KDELRL1	CAPZB	RPN1	CASP8	ABL1_2
##	6.618350e-03	6.615944e-03	6.594308e-03	6.593020e-03	6.585298e-03
##	DNAJA1	ENSA_1	ALDH9A1	SORL1_1	ACADVL
##	6.585143e-03	6.575073e-03	6.561170e-03	6.552972e-03	6.549170e-03
##	41718_g_at	FGFR1	ITM2B	2064_g_at	RNF11
##	6.547092e-03	6.544976e-03	6.517320e-03	6.515733e-03	6.513707e-03
##	DDX18	39728_at	MXRA7	AP1S2	31673_s_at
##	6.512542e-03	6.508779e-03	6.507427e-03	6.499305e-03	6.496461e-03
##	IGHM_1	KAT8_1	CSNK1G2	FAM53B	BTN3A2
##	6.484783e-03	6.475562e-03	6.474051e-03	6.469870e-03	6.466415e-03

##	SLC30A9	1184_at	TDP2	ARCN1	ALOX5AP
##	6.445967e-03	6.393714e-03	6.383781e-03	6.378096e-03	6.362933e-03
##	HUWE1_1	SULT1A1	PAK2	DDX3X	OXA1L
##	6.349231e-03	6.328225e-03	6.311979e-03	6.309708e-03	6.303433e-03
##	RAB14	TIA1	41562_at	INSIG1	LDLRAD4_1
##	6.301641e-03	6.299230e-03	6.289257e-03	6.282416e-03	6.278038e-03
##	ZBTB18	PRNP	MYO9B	CAD_1	RAF1
##	6.274028e-03	6.261353e-03	6.254759e-03	6.246752e-03	6.229780e-03
##	SKIV2L2	VPS13B	CCND3_1	374_f_at	HDLBP
##	6.219733e-03	6.194135e-03	6.184472e-03	6.177384e-03	6.175480e-03
##	RARRES3	CCNF	CARS	ALDH5A1	CDC20
##	6.167576e-03	6.136248e-03	6.134463e-03	6.131647e-03	6.114121e-03
##	PSMD7_1	FNBP1	SRP54	38585_at	IMP4
##	6.109705e-03	6.109121e-03	6.094617e-03	6.092622e-03	6.087379e-03
##	TXNL4A	SEPHS2	SACM1L	466_at	TSPAN7
##	6.081574e-03	6.078301e-03	6.051976e-03	6.038137e-03	6.022637e-03
##	MAGED2_1	RNF4	TUG1	PNP	ANXA11
##	6.019246e-03	6.018995e-03	5.993896e-03	5.993700e-03	5.986343e-03
##	KLF6	DYNLT1	39065_s_at	TMBIM6_1	RFC4
##	5.984831e-03	5.976607e-03	5.970538e-03	5.954487e-03	5.942143e-03
##	SMARCA2	VPS26A	MAZ	SAFB2	IVNS1ABP
##	5.939798e-03	5.934716e-03	5.927599e-03	5.922402e-03	5.912355e-03
##	RASA1_1	GRHPR	UFL1	MPPE1	IGFBP7
##	5.896599e-03	5.896318e-03	5.894596e-03	5.877180e-03	5.874688e-03
##	PLEKHB2	EMP3	CLINT1	EIF4G1	PRPS1
##	5.871294e-03	5.856192e-03	5.846925e-03	5.846668e-03	5.843494e-03
##	NUDT21	MAPK1IP1L	FNBP4	YME1L1	PSMC6
##	5.832788e-03	5.818444e-03	5.816188e-03	5.809639e-03	5.791745e-03
##	SCAMP3	AHCY	PPP2CA	TFAP4	32317_s_at
##	5.756856e-03	5.754388e-03	5.748390e-03	5.740686e-03	5.736688e-03
##	HNRNPDL	TARBP1	969_s_at	AEBP1	CIB1
##	5.734238e-03	5.734081e-03	5.730227e-03	5.715821e-03	5.695890e-03
##	PURA	SPTAN1	PTOV1	BRD8	STAT5A
##	5.692284e-03	5.686162e-03	5.669223e-03	5.651899e-03	5.638224e-03
##	FKBP1A	LAGE3	MARCH7	NSA2	PRKAR1A_1
##	5.633990e-03	5.631130e-03	5.629926e-03	5.627965e-03	5.626854e-03
##	CNOT2	LRMP	SEPT6_1	FTH1	HDDC2
##	5.624300e-03	5.624140e-03	5.620928e-03	5.599879e-03	5.599206e-03
##	1728_at	XPO6	PSMD3	BARD1	POM121
##	5.574632e-03	5.573395e-03	5.553169e-03	5.544031e-03	5.536496e-03
##	RHOH	COPA	SURF1	NUP214	TK1
##	5.533050e-03	5.528843e-03	5.504576e-03	5.503578e-03	5.499426e-03
##	BCL7A	33234_at	PTTG1	PPP1R7	SUM03
##	5.494039e-03	5.491644e-03	5.490616e-03	5.488374e-03	5.484991e-03
##	GARS_1	YWHAZ_1	OTUD4	GYPC	VAMP3
##	5.484845e-03	5.483971e-03	5.481382e-03	5.475814e-03	5.474297e-03
##	39416_at	RANBP2	RBMS1	MKI67	R3HCC1
##	5.473255e-03	5.471861e-03	5.463717e-03	5.454384e-03	5.453238e-03
##	TMED5	IGHM_2	NIPSNAP2	ADD1	PATZ1
##	5.445274e-03	5.430095e-03	5.418247e-03	5.407951e-03	5.403799e-03
##	ERGIC3	SMC4	UBE4A	37907_at	40427_at
##	5.399575e-03	5.398079e-03	5.391514e-03	5.375143e-03	5.364442e-03
##	EIF2S1_1	RAD23A	SORL1	SGTA	40783_s_at
##	5.355241e-03	5.351682e-03	5.321039e-03	5.314839e-03	5.313164e-03

##	TP53	39372_at	ADH5_1	SSBP2	NFKB1_2
##	5.311293e-03	5.301947e-03	5.298284e-03	5.290645e-03	5.285816e-03
##	41132_r_at	ATP6V1A	IGFBP4	BCL11A	PMS2P1
##	5.283529e-03	5.283293e-03	5.282215e-03	5.278081e-03	5.262931e-03
##	CRIP1	OSBPL8	PSMD10	LRRFIP1	SLC35A3
##	5.245757e-03	5.230035e-03	5.203902e-03	5.161772e-03	5.147186e-03
##	ASMTL	BYSL	CCT4	SEC14L1	BAZ1B
##	5.113100e-03	5.064270e-03	5.063276e-03	5.062568e-03	5.055012e-03
##	SMC5	HDGF_1	RBM42	ANXA6	TRIM14
##	5.031861e-03	5.019554e-03	5.014094e-03	5.010865e-03	5.002251e-03
##	ZFR	CYB5B_1	CKAP5	TYROBP	ICAM2_2
##	4.999275e-03	4.995920e-03	4.983345e-03	4.981240e-03	4.963767e-03
##	33689_s_at	CHSY1	CHAF1A	CALR	ISCU
##	4.937184e-03	4.927999e-03	4.924514e-03	4.918605e-03	4.915327e-03
##	SMARCC1_1	CCNB1	UROD	STAT5B	PSMD7
##	4.908579e-03	4.902557e-03	4.895483e-03	4.891485e-03	4.860104e-03
##	PDIA6	SNU13	DDIT4	TGFB2_1	PPP6R1
##	4.858979e-03	4.857616e-03	4.838424e-03	4.832644e-03	4.829177e-03
##	925_at	FEZ2	BEX4	XPOT	GRSF1
##	4.826320e-03	4.818852e-03	4.813719e-03	4.813232e-03	4.807502e-03
##	SH3GLB1	LSP1	CBX6	UBE2I	SRF
##	4.797379e-03	4.796706e-03	4.783136e-03	4.773695e-03	4.772297e-03
##	PGRMC1	ETF1	FKBP2	HIPK1	CYTH1
##	4.770114e-03	4.764506e-03	4.737350e-03	4.734316e-03	4.731427e-03
##	PRDX3	FBXO21	TIAL1	NCBP2_1	ATR
##	4.717913e-03	4.700016e-03	4.672382e-03	4.659048e-03	4.646461e-03
##	TTC1	41128_at	TPR	ATP2B4	SCAP
##	4.623966e-03	4.621249e-03	4.604415e-03	4.586544e-03	4.584446e-03
##	32877_i_at	FKBP5	HAUS7	SNRNP70	RAP1GDS1
##	4.581611e-03	4.575337e-03	4.551646e-03	4.538053e-03	4.532915e-03
##	TUBG1	SRSF11	ADH5	LEPROTL1	NFATC2IP_1
##	4.517003e-03	4.507539e-03	4.495129e-03	4.484180e-03	4.482905e-03
##	TOP1	SH3BP5	GNAI2	RNH1	EED
##	4.475063e-03	4.468193e-03	4.461926e-03	4.455764e-03	4.454429e-03
##	SMARCB1	IGLC1	PA2G4	HADH	RNPEP
##	4.454113e-03	4.452920e-03	4.447290e-03	4.445901e-03	4.439463e-03
##	S100A4	ADCY7	PSMD2	DAZAP2	PPP2R1A_1
##	4.437799e-03	4.436219e-03	4.429477e-03	4.419747e-03	4.419137e-03
##	AHNAK	MYB_1	NUMA1	SP3	G3BP2
##	4.412720e-03	4.398237e-03	4.390990e-03	4.363777e-03	4.347712e-03
##	37742_at	38923_at	IMMT	PPT1	EXOSC8
##	4.323671e-03	4.315768e-03	4.309706e-03	4.291592e-03	4.290540e-03
##	PSD4	PTPA_1	UBR4	31510_s_at	PLP2
##	4.277007e-03	4.274757e-03	4.266979e-03	4.256371e-03	4.240069e-03
##	ATP6V1E1	MARCKSL1	RETREG3	GMFG	NDUFB7
##	4.228658e-03	4.225972e-03	4.223912e-03	4.218314e-03	4.202332e-03
##	UBA3	HBD	PITRM1	33274_f_at	CAPN2
##	4.201755e-03	4.192187e-03	4.186277e-03	4.183978e-03	4.176217e-03
##	SUZ12	EPS15	SRRT	MT1E	CLEC2B
##	4.166840e-03	4.164058e-03	4.136685e-03	4.129721e-03	4.115444e-03
##	39129_at	ACAA2	HLTF	TXNL1	FAM168B
##	4.112958e-03	4.112534e-03	4.104949e-03	4.104640e-03	4.103361e-03
##	DGCR2	APLP2	MBOAT7	35688_g_at	TERF2IP
##	4.101617e-03	4.100482e-03	4.100013e-03	4.093947e-03	4.073527e-03

##	TBC1D1	GDI2_1	ABL1_1	INTS10	SMC3
##	4.063374e-03	4.055309e-03	4.044238e-03	4.031053e-03	4.017949e-03
##	EFCAB14	PPP6C	BNIP3L	IMPA1	SLA
##	4.015608e-03	4.015569e-03	4.012829e-03	3.995349e-03	3.981656e-03
##	WAPL	RBBP4	CSTF3	CRKL	DHPS
##	3.970988e-03	3.969240e-03	3.964929e-03	3.964641e-03	3.962583e-03
##	TOR1AIP1	ACAP1	LEF1	RBL2	BRD2
##	3.946715e-03	3.943021e-03	3.925919e-03	3.903129e-03	3.897744e-03
##	FBX09	GBE1	DICER1	ULK1	31503_at
##	3.892184e-03	3.885410e-03	3.864582e-03	3.863224e-03	3.850973e-03
##	40132_g_at	LASP1	FOXJ3	NEK7	R3HDM1
##	3.804319e-03	3.802494e-03	3.793163e-03	3.790164e-03	3.774655e-03
##	SMIM10L1	ZNF384	ITGB3BP	ITGA4_1	ATOX1
##	3.759364e-03	3.758210e-03	3.725181e-03	3.715843e-03	3.709427e-03
##	IL7R_1	FDFT1	MFAP1	SLC39A6	CUL4B
##	3.698420e-03	3.687271e-03	3.661532e-03	3.657913e-03	3.646350e-03
##	MTA1	729_i_at	RELA_1	SRSF10	SUM04
##	3.628754e-03	3.627134e-03	3.626914e-03	3.619393e-03	3.612181e-03
##	NDUFS7	NAA10	POLA1	DR1	TAF7
##	3.591422e-03	3.588204e-03	3.585921e-03	3.585187e-03	3.559269e-03
##	DDX42	ARF3	RRAGA	ATP6VOC	41233_at
##	3.548930e-03	3.546768e-03	3.525441e-03	3.523495e-03	3.512876e-03
##	YIF1A	CYB5R3	CNOT9	IDH3B_1	CDKN2C
##	3.512670e-03	3.493291e-03	3.483199e-03	3.455139e-03	3.423441e-03
##	CDK19	MYCBP2	PPP2R5C_1	ANKRD17	37703_at
##	3.419378e-03	3.394900e-03	3.391221e-03	3.372155e-03	3.369426e-03
##	RB1_1	SF3A2	SF1	TRAP1	SIAH2
##	3.365086e-03	3.360735e-03	3.357114e-03	3.343028e-03	3.342683e-03
##	CYB5A_1	TAX1BP1_1	RAB5A	SPG11	LUC7L3
##	3.342426e-03	3.340607e-03	3.332299e-03	3.312254e-03	3.308713e-03
##	SFPQ_1	UBE2I_2	CHD3	IQGAP1	CCNG1
##	3.307518e-03	3.295087e-03	3.285706e-03	3.274677e-03	3.272721e-03
##	USP22	RAG2	UBE2A	HTATSF1	ACO2
##	3.270883e-03	3.258323e-03	3.242597e-03	3.231900e-03	3.223816e-03
##	PDAP1	MRPS27	MAP2K1_1	CETN3	GTSE1
##	3.213771e-03	3.201839e-03	3.193755e-03	3.191873e-03	3.182359e-03
##	DDX19B	H1FX	GTF2B	AHCYL1_2	EMG1
##	3.176989e-03	3.118901e-03	3.108822e-03	3.107396e-03	3.066832e-03
##	VAR5	TRAPPC6A	MACF1	UBTF	PAPSS1
##	3.041889e-03	3.037404e-03	3.033142e-03	3.032942e-03	3.031688e-03
##	ABI1	CEBPB	40036_at	MAGED2	SEC13_1
##	3.025061e-03	3.023665e-03	2.994032e-03	2.990141e-03	2.989861e-03
##	ATP2A3	ECHS1	RAE1	COPB1	IL7R
##	2.978850e-03	2.977322e-03	2.969024e-03	2.948804e-03	2.945326e-03
##	GOLGA3	TLK1	USF2	SPAG7	PPWD1
##	2.938382e-03	2.929258e-03	2.928943e-03	2.924472e-03	2.921290e-03
##	FBX07	ICAM2	MBD4	FYN	FLII
##	2.920046e-03	2.917144e-03	2.888169e-03	2.884362e-03	2.883417e-03
##	LMNB1	EI24	MYB_5	RANGAP1	PPP2R5C_2
##	2.873310e-03	2.868161e-03	2.864917e-03	2.861856e-03	2.857809e-03
##	CCND3	IFITM3	PIEZ01	SAT1_1	35926_s_at
##	2.855130e-03	2.849162e-03	2.839203e-03	2.815434e-03	2.813057e-03
##	SNRNP40	DNAJC11	HMGN4	LARP4B	MAPK1
##	2.810393e-03	2.809527e-03	2.800051e-03	2.798585e-03	2.788156e-03

##	EWSR1	KIDINS220	ABCD3	ATXN2L	PRDX1
##	2.770070e-03	2.762269e-03	2.762008e-03	2.747230e-03	2.734838e-03
##	PARP1_1	ITGB1	API5	LITAF	TLK2
##	2.721206e-03	2.720419e-03	2.697873e-03	2.682845e-03	2.682505e-03
##	CTDSP2	ZC3HAV1	AARS	PSME1	SLC20A1
##	2.679800e-03	2.674402e-03	2.664926e-03	2.660519e-03	2.660342e-03
##	SRSF10_1	34882_at	UBE2D2_1	SLC7A6	EBP
##	2.656749e-03	2.645139e-03	2.619799e-03	2.606154e-03	2.560844e-03
##	ATXN7L3B	PRPF4	GPX7	MGMT	NCBP2
##	2.560700e-03	2.554736e-03	2.533066e-03	2.509244e-03	2.500862e-03
##	AFG3L2	MAPK14	ACTR2	PNPLA6	PJA2
##	2.489990e-03	2.487604e-03	2.485183e-03	2.484261e-03	2.478266e-03
##	WDR7	ATF2	P2RX5	PAFAH1B1	PPP2CB
##	2.471492e-03	2.470531e-03	2.467980e-03	2.455969e-03	2.454814e-03
##	PDS5A	MICB	ITGB2	DENND5A	DGKA
##	2.449811e-03	2.431217e-03	2.412050e-03	2.405029e-03	2.401529e-03
##	RCBTB2	WAS_1	UBE2L6	SUGP2	OAT
##	2.399539e-03	2.390650e-03	2.382886e-03	2.381090e-03	2.357141e-03
##	JADE2	AMPD2	MSH6	HNRNPD	ACYPI
##	2.356571e-03	2.352651e-03	2.351462e-03	2.342236e-03	2.338171e-03
##	SON	PSMC4	STK24	PEX11B	UCP2
##	2.330068e-03	2.323355e-03	2.318786e-03	2.315126e-03	2.290432e-03
##	SF3A1	TMBIM6	ARPC3	34829_at	IDH3G
##	2.287525e-03	2.284310e-03	2.275717e-03	2.272614e-03	2.232990e-03
##	PSMB5	GINS1	PSMB10	SRSF3	TCERG1
##	2.230924e-03	2.230002e-03	2.202775e-03	2.190250e-03	2.176700e-03
##	MYDGF	ATP2A2_1	BECN1	LRPPRC	MAP4K1
##	2.173507e-03	2.170688e-03	2.135152e-03	2.124057e-03	2.109551e-03
##	PPP4C	PHC2	CD2AP	MYB_2	UBE2D2
##	2.108603e-03	2.106852e-03	2.104457e-03	2.100032e-03	2.093571e-03
##	S100A8	TRIM44	ZNF207	NFYB	IGBP1
##	2.092395e-03	2.071522e-03	2.071001e-03	2.050889e-03	2.045199e-03
##	CBX5	CCNA2_1	GCFC2	NMI	TSPYL4
##	2.044607e-03	2.036996e-03	2.026882e-03	2.023071e-03	2.001860e-03
##	ATP6V0B	HSD17B10	PGM1	EIF6	SPOCK2
##	1.994985e-03	1.963764e-03	1.960744e-03	1.957998e-03	1.956138e-03
##	BLM	NDUFA7	PRIM1	ICAM2_1	ACP1
##	1.954181e-03	1.950269e-03	1.933575e-03	1.926758e-03	1.926719e-03
##	SNX3	SLC3A2	ILF3	SNX4	PWP1
##	1.923252e-03	1.915501e-03	1.913932e-03	1.901348e-03	1.897823e-03
##	UBE2N	39050_at	FUBP3	FARSA_1	SELL
##	1.887766e-03	1.885704e-03	1.874283e-03	1.860710e-03	1.854690e-03
##	ARF5	PIP4K2B	SLC5A3	PSMB8	ATP5F1
##	1.853600e-03	1.849137e-03	1.844738e-03	1.838556e-03	1.836344e-03
##	MCM5	31600_s_at	MCL1_1	ASF1A	OGT
##	1.833538e-03	1.830185e-03	1.796610e-03	1.762893e-03	1.757275e-03
##	LAMP1	BAG6	CD247	USP4	UBR5
##	1.724267e-03	1.709562e-03	1.704877e-03	1.685346e-03	1.674547e-03
##	GNAQ	PPP2R2A_1	ILF3_1	OGT_1	MYB_4
##	1.674147e-03	1.666526e-03	1.655367e-03	1.648951e-03	1.644786e-03
##	ARF6	VAMP8	DNMT1	MPV17	STAT3
##	1.635393e-03	1.629162e-03	1.592096e-03	1.589295e-03	1.581819e-03
##	MPHOSPH10	FAM32A	RNF44	IK	CHUK
##	1.570860e-03	1.560153e-03	1.550090e-03	1.549054e-03	1.545485e-03

##	ITPA	DCTN2	PCMT1_2	41597_s_at	PRKACB
##	1.539724e-03	1.535626e-03	1.532380e-03	1.529428e-03	1.528614e-03
##	RPS6KA3	CDIPT	MBD3	VRK1	POGZ
##	1.528415e-03	1.526445e-03	1.524515e-03	1.515685e-03	1.493180e-03
##	TUBGCP2_1	ATP6V1F	R3HDM2	ALG8	MT3
##	1.476129e-03	1.458978e-03	1.442291e-03	1.440742e-03	1.440635e-03
##	SNRPA1	ABL1	PARP1	SCCPDH	AIMP2
##	1.438148e-03	1.424408e-03	1.405295e-03	1.387031e-03	1.383872e-03
##	IRAK1	SPINT2	CTSH	MT1B	38072_at
##	1.382998e-03	1.372070e-03	1.368710e-03	1.366674e-03	1.356369e-03
##	SRP72	LETMD1	GSPT1	RHOG	CDKN2D
##	1.339519e-03	1.337087e-03	1.300125e-03	1.273071e-03	1.264146e-03
##	BAZ1A	UBE3A_1	NUP98	TSP0	EIF1AX_2
##	1.253329e-03	1.251788e-03	1.249759e-03	1.248350e-03	1.243761e-03
##	MRPL33	CAPN1	MCL1	34842_at	CD48
##	1.230854e-03	1.203481e-03	1.200879e-03	1.198623e-03	1.195972e-03
##	GUSBP11_1	ZNHIT1	41185_f_at	ESYT1	PPP2R5C_3
##	1.172795e-03	1.172049e-03	1.167148e-03	1.161702e-03	1.158217e-03
##	NPTN	DAXX	PPP1R11	32674_at	RAC1
##	1.157567e-03	1.153130e-03	1.152033e-03	1.144137e-03	1.143769e-03
##	SH3BGRL	MYB_3	TPGS2	SREBF2	33132_at
##	1.138716e-03	1.134079e-03	1.131171e-03	1.121541e-03	1.115529e-03
##	IGLL1	ERCC1_1	ACTR2_1	RNF6	ADIPOR2
##	1.114425e-03	1.103698e-03	1.102930e-03	1.084751e-03	1.082464e-03
##	PDLIM1	EIF4E2	NUDT1	TMF1	PPP2R5C
##	1.080275e-03	1.073748e-03	1.072276e-03	1.061440e-03	1.048266e-03
##	CCNA2	DDX46	DGKD	BSG	SNX1_1
##	1.035799e-03	1.026049e-03	9.968248e-04	9.903010e-04	9.886669e-04
##	BCL7B	UBAP2L	DYRK1A_1	41170_at	ARHGEF2
##	9.771515e-04	9.674280e-04	9.651185e-04	9.522824e-04	9.157203e-04
##	MYH9	DCTN3	SMARCA5	ACTN1	SSNA1
##	9.134535e-04	8.890275e-04	8.745456e-04	8.573573e-04	8.499408e-04
##	PPP2R2A	MTX1	39133_at	NFATC3	POLG
##	8.405383e-04	8.268736e-04	8.054509e-04	7.988562e-04	7.910434e-04
##	RRM1	PAIP1	ANXA5	39341_at	RBM39
##	7.850166e-04	7.833753e-04	7.811430e-04	7.748676e-04	7.719494e-04
##	PRDX4	UBE2J1	CDK2	DYRK1A	SLC25A36
##	7.653927e-04	7.596108e-04	7.550153e-04	7.488844e-04	7.488368e-04
##	POLR2H	NDUFA5	EXOSC7	PHKB	CLASP2
##	7.464619e-04	7.271278e-04	7.239739e-04	7.237445e-04	7.200520e-04
##	ST13	RPIA	RNASEH2B	HDAC1_1	ATP1B3
##	7.120455e-04	7.008217e-04	6.935114e-04	6.837224e-04	6.741265e-04
##	ZNF24	DGKZ	USP9X	MCM2	STAT6
##	6.548647e-04	6.490966e-04	6.414715e-04	6.231849e-04	6.185794e-04
##	CPNE1	TAX1BP1	METTL7A	ABCF1	MDM1
##	6.134568e-04	6.118785e-04	6.057066e-04	5.758713e-04	5.742352e-04
##	USP10	CEBPZ	CSE1L	EIF5_1	BNIP2
##	5.681019e-04	5.517715e-04	5.461995e-04	5.343836e-04	5.338330e-04
##	ARFGAP2	SMARCA4	41171_at	PLIN2	PIKFYVE
##	5.267058e-04	5.190541e-04	5.033864e-04	5.012445e-04	4.746192e-04
##	ARPC1B	HYOU1	COMT	SRSF7	NUP88
##	4.724262e-04	4.713211e-04	4.603091e-04	4.561579e-04	4.554182e-04
##	MLH1	CD99	ZC3H4	393_s_at	RER1
##	4.531458e-04	4.419164e-04	4.379201e-04	4.302755e-04	4.214908e-04

```
##          KRT10          NUP50          DAP          TPM4          TXNRD1
## 4.134017e-04 4.061259e-04 3.948040e-04 3.907445e-04 3.873447e-04
##      432_s_at      LGALS3BP      C21orf33      ITPKB      AES_1
## 3.848690e-04 3.785471e-04 3.561622e-04 3.545129e-04 3.504927e-04
##      SRPK1      GORASP2      RAP1B      PNN      ATP8A1
## 3.415470e-04 3.398254e-04 3.281394e-04 3.264299e-04 3.088563e-04
##      NFATC3_1      RAD23B      RARS      RELA      1420_s_at
## 2.986639e-04 2.982330e-04 2.943287e-04 2.801652e-04 2.781550e-04
##      AZIN1_2      STAU1      ARHGEF7      BTN3A2_1      ZPR1
## 2.550922e-04 2.526246e-04 2.477764e-04 2.380802e-04 2.299243e-04
##      NSD2      MEA1      NDUFS6      37735_at      DDX17
## 2.255866e-04 2.091612e-04 2.069900e-04 1.956215e-04 1.607080e-04
##      UBA1      PQBP1      MKI67_1      EZR      CCDC69
## 1.576586e-04 1.493142e-04 1.281797e-04 1.277985e-04 1.186925e-04
##      BIRC2      ICAM3      IFT20      GTF3C2      HSPA5
## 1.160605e-04 1.065337e-04 5.701387e-05 5.199234e-05 3.773451e-05
##      ZFAND5      EVI2B      35985_at      SEC11A      BUB1B
## 3.513191e-05 3.282533e-05 3.193166e-05 2.400636e-05 3.499181e-06
##      SRSF8
## 7.950999e-07
```

```
pc_1 = data.frame(leukemia_data$Type,pr.out$x[,1])
colnames(pc_1) <- c('Tpye','PC1')
## sort the PC1 value
sorted <- pc_1[order(-abs(pc_1$PC1)),]
head(sorted,6)
```

```
##          Tpye          PC1
## 161      OTHERS -52.06836
## 247      T-ALL  51.75384
## 303      TEL-AML1 -49.06527
## 231      T-ALL  48.16035
## 56      Hyperdip50 -48.09339
## 214      T-ALL  47.96769
```

```
###f
```

```
## load dendextend library
library(dendextend)
```

```
##
## -----
## Welcome to dendextend version 1.15.1
## Type citation('dendextend') for how to cite the package.
##
## Type browseVignettes(package = 'dendextend') for the package vignette.
## The github page is: https://github.com/talgalili/dendextend/
##
## Suggestions and bug-reports can be submitted at: https://github.com/talgalili/dendextend/issues
## Or contact: <tal.galili@gmail.com>
##
## To suppress this message use: suppressPackageStartupMessages(library(dendextend))
## -----
```

```
##
## Attaching package: 'dendextend'

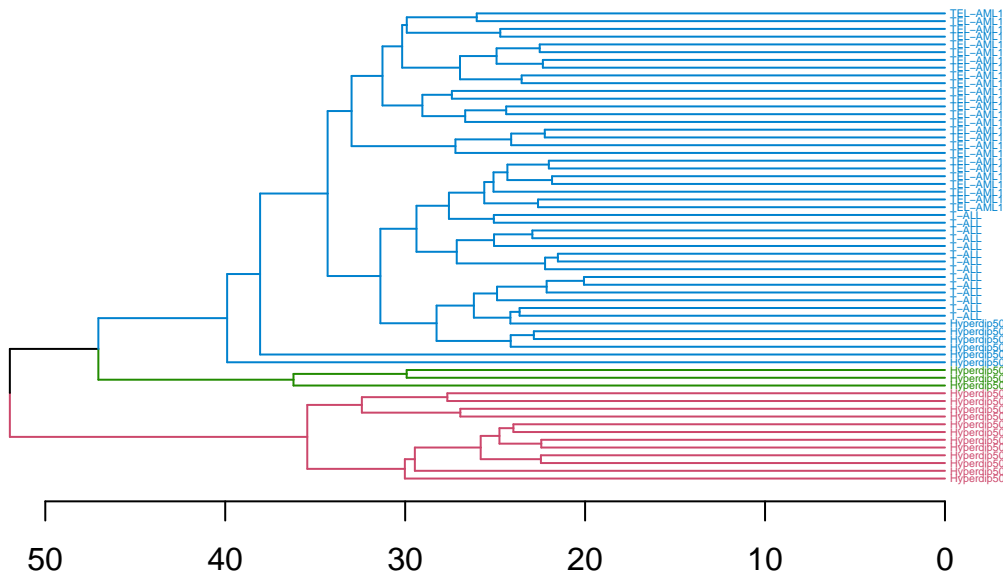
## The following object is masked from 'package:rpart':
##
##      prune

## The following object is masked from 'package:stats':
##
##      cutree

##subsetting to include only rows for which Type is either T-ALL, TEL-AML1, or Hyperdip50
leukemia_subset_1 <- filter(leukemia_data, Type == c("T-ALL", "TEL-AML1", "Hyperdip50"))
## exclude the first column Type
leukemia_subset = leukemia_subset_1 %>% select(-Type)
## calculate the distance matrix
subset_dist = dist(leukemia_subset)
set.seed(1)
## Hierarchical Clustering using complete linkage
drug.hclust = hclust(subset_dist)

## first plot
x = as.dendrogram(drug.hclust)
x %>% set_labels(leukemia_subset_1$Type) %>%
  set("labels_col", k=3) %>%
  set("branches_k_color", k = 3) %>%
  set("labels_cex", 0.3) %>%
  plot(main='Three Groups for hclust', horiz = TRUE)
```

Three Groups for hclust



```
## second plot
y = as.dendrogram(drug.hclust)
y %>% set_labels(leukemia_subset_1$Type) %>% set("labels_col",k=5) %>% set("branches_k_color", k = 5) %>%
  set("labels_cex", 0.3) %>% plot(main='Five Groups for hclust',horiz = TRUE)
```

