## **Example report of OUKS**

This R Markdown document was provided as an example to reproduce the *OUKS* code script. Fig. 4 from article can be reproduced with the following code. Preliminarily download "xcms after IPO MVI QC-XGB filter repeats annot+filtr LMM adj KEGG.csv" and "8 peaks.csv" files into your working directory (for example: ("D:/OUKS/").

### **Prepare environment**

First, set the folder for the working directory and load the packages.

```
setwd("D:/OUKS/")
library(data.table)
## Warning: package 'data.table' was built under R version 4.0.2
library(factoextra)
## Loading required package: ggplot2
## Registered S3 methods overwritten by 'tibble':
##
    method
               from
##
    format.tbl pillar
     print.tbl pillar
##
## Welcome! Want to learn more? See two factoextra-related books at https://goo.
gl/ve3WBa
library(FactoMineR)
library(dendextend)
##
## -----
## Welcome to dendextend version 1.13.4
## 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(dendex
tend))
## -----
## Attaching package: 'dendextend'
```

```
## The following object is masked from 'package:data.table':
##
##
       set
## The following object is masked from 'package:stats':
##
##
       cutree
library(rafalib)
library(pROC)
## Type 'citation("pROC")' for a citation.
##
## Attaching package: 'pROC'
## The following objects are masked from 'package:stats':
##
##
       cov, smooth, var
library(parallel)
library(doParallel)
## Loading required package: foreach
## Warning: package 'foreach' was built under R version 4.0.3
## Loading required package: iterators
## Warning: package 'iterators' was built under R version 4.0.3
library(grid)
library(caret)
## Loading required package: lattice
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:data.table':
##
##
       between, first, last
## The following objects are masked from 'package:stats':
##
       filter, lag
##
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
library(MKinfer)
## Warning: package 'MKinfer' was built under R version 4.0.2
```

### **Load datasets**

```
ds <- as.data.frame(fread(input = "8 peaks.csv", header=T))
rownames(ds) <- ds[,1]
ds <- ds[,-1]
colnames(ds)[1] <-"Label"
ds[,-1] <- sapply(ds[,-1], as.numeric)
ds$Label <- as.factor(ds$Label)

ds2 <- as.data.frame(fread(input = "xcms after IPO MVI QC-XGB filter repeats ann
ot+filtr LMM adj KEGG.csv", header=T))
ds2 <- ds2[-c(1:12),]
rownames(ds2) <- ds2[,5]
ds2 <- ds2[,-c(1,3:5)]
ds2[,-1] <- sapply(ds2[,-1], as.numeric)
ds2$Label <- as.factor(ds2$Label)</pre>
```

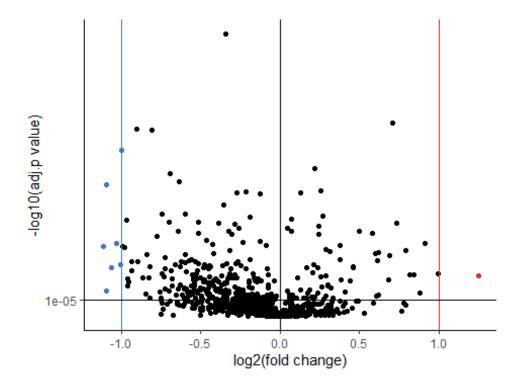
# Plot volcano plot

```
ds_log <- as.data.frame(log2(ds2[,-1]))
ds_log <- cbind(Label = ds2[,1], ds_log)

FOLD.CHANGE <- function(data) {
    ds_log_subsets <- lapply(1:length(unique(data[,1])), function(y) filter(data[,-1], data$Label == unique(data[,1])[y]))
    mean_r_l <- lapply(1:length(ds_log_subsets), function(y) apply(ds_log_subsets[[y]], 2, mean, na.rm = T))
    foldchange <- mean_r_l[[1]] - mean_r_l[[2]]
    fc_res <- as.data.frame(foldchange)
    return(fc_res)
}

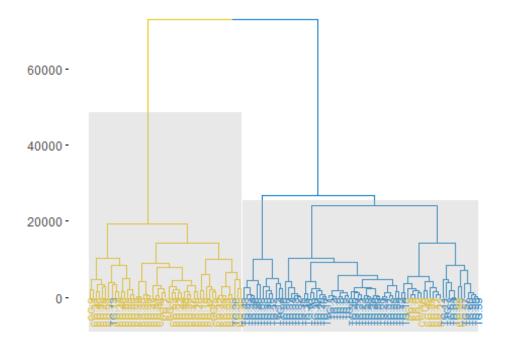
fc_res <- FOLD.CHANGE(ds_log)
foldchange <- as.numeric(fc_res$foldchange)

mdl_mtrx <- model.matrix(~Label, ds2)
lmf <- lmFit(t(ds2[,-1]), method = "robust", design = mdl_mtrx, maxit = 1000)</pre>
```

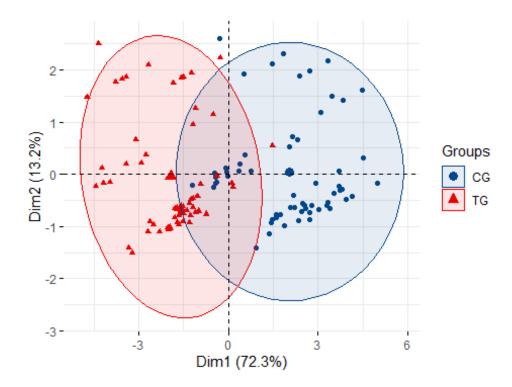


# **HCA plot**

```
color_labels_by_k = F,
label_cols = Cols(grp1,res.hc1$order),
rect = T,
rect_fill = T,
horiz = F,
lwd = 0.7,
show_labels = T,
main = "",
ylab = "")
```

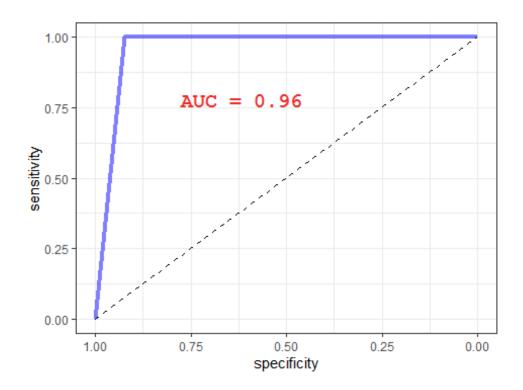


# **PCA plot**



### **ROC** curve

```
# start parallel processing
fc <- as.numeric(detectCores(logical = F))</pre>
cl <- makePSOCKcluster(fc+1)</pre>
registerDoParallel(cl)
# cross-validation
set.seed(1234)
trainIndex <- createDataPartition(ds$Label, p = 0.8, list = F, times = 1)</pre>
dsTrain <- ds[ trainIndex,]</pre>
dsValid <- ds[-trainIndex,]</pre>
trainControl <- trainControl(method="repeatedcv", number=10, repeats=10, classPr</pre>
obs = F)
metric <- "Accuracy"</pre>
# machine learning
set.seed(1234)
fit.cl <- train(Label~., data=dsTrain, method="svmRadial", metric=metric, trCont</pre>
rol=trainControl, tuneLength = 10)
predicted.classes <- predict(fit.cl, newdata=dsValid)</pre>
# ROC curve
res.roc <- roc(as.numeric(as.factor(predicted.classes)), as.numeric(dsValid$Labe
1))
## Setting levels: control = 1, case = 2
## Setting direction: controls < cases</pre>
```



## bootstrap histogram

```
# start parallel processing
fc <- as.numeric(detectCores(logical = F))
cl <- makePSOCKcluster(fc+1)
registerDoParallel(cl)

# bootstrap
set.seed(1234)
trainControl <-trainControl(method="boot", number=1000)
metric <- "Accuracy"

# machine Learning
set.seed(1234)
fit.cl <- train(Label~., data=ds, method="svmRadial", metric=metric, trControl=trainControl)
results <- resamples(list(svm=fit.cl, svm1=fit.cl), trControl = trainControl, metric=metric)

# histogram</pre>
```

```
d <-ggplot(results$values, aes(x=results$values[,2])) +
   geom_histogram(colour="blue", fill="white") + theme_bw() + xlab("Accuracy") +</pre>
ylab("Frequency")
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

