

# project

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```
library(rJava, warn.conflicts = FALSE, quietly=TRUE)
library(xlsx, warn.conflicts = FALSE, quietly=TRUE)
library(stringr, warn.conflicts = FALSE, quietly=TRUE)
library(dplyr, warn.conflicts = FALSE, quietly=TRUE)
library(readr, warn.conflicts = FALSE, quietly=TRUE)
library(randomForestSRC, warn.conflicts = FALSE, quietly=TRUE)
library(ggplot2, warn.conflicts = FALSE, quietly=TRUE)
library(ggthemes, warn.conflicts = FALSE, quietly=TRUE)
library(caret, warn.conflicts = FALSE, quietly=TRUE)
library(tidyr, warn.conflicts = FALSE, quietly=TRUE)
library(scales, warn.conflicts = FALSE, quietly=TRUE)
library(data.table, warn.conflicts = FALSE, quietly=TRUE)
library(effects, warn.conflicts = FALSE, quietly=TRUE)
library(gridExtra, warn.conflicts = FALSE, quietly=TRUE)
library(ggRandomForests, warn.conflicts = FALSE, quietly=TRUE )
library(ROCR, warn.conflicts = FALSE, quietly=TRUE)
library(ggpubr, warn.conflicts = FALSE, quietly=TRUE)
library(grid, warn.conflicts = FALSE, quietly=TRUE)
```

## *#Functions*

*#AccuracyCutoffInfo, ConfusionMatrixInfo, ROCInfo function is completetly coded by github user ethen818*

*#The AccuracyCutoffInfo, ConfusionMatrixInfo, and ROCInfo functions in this Rmarkdown are a modified version*

*#The following function can be found at the following github page*

*#[https://github.com/ethen8181/machine-learning/blob/master/unbalanced/unbalanced\\_code/unbalanced\\_functions.R](https://github.com/ethen8181/machine-learning/blob/master/unbalanced/unbalanced_code/unbalanced_functions.R)*

*#Credit to user ethen8181 for creating these functions.*

```
# -----
# [AccuracyCutoffInfo] :
# Obtain the accuracy on the training and testing dataset.
# for cutoff value ranging from .4 to .8 ( with a .05 increase )
# @train   : your data.table or data.frame type training data ( assumes you have the predicted score in
# @test    : your data.table or data.frame type testing data
# @predict : prediction's column name (assumes the same for training and testing set)
# @actual  : actual results' column name
# returns  : 1. data : a data.table with three columns.
#            each row indicates the cutoff value and the accuracy for the
#            train and test set respectively.
#            2. plot : plot that visualizes the data.table
```

```
AccuracyCutoffInfo <- function( train, test, predict, actual )
{
  # change the cutoff value's range as you please
  cutoff <- seq( .05, 1, by = .025 )
```

```

accuracy <- lapply( cutoff, function(c)
{
  train_prediction <- as.factor(as.numeric( train[[predict]] > c ))
  test_prediction <- as.factor(as.numeric( test[[predict]] > c ))

  levels(train_prediction) <- c(levels(train[[actual]][1]),levels(train[[actual]][2]))
  levels(test_prediction) <- c(levels(test[[actual]][1]),levels(test[[actual]][2]))

  # use the confusionMatrix from the caret package
  cm_train <- confusionMatrix( train_prediction, train[[actual]] )
  cm_test  <- confusionMatrix( test_prediction, test[[actual]] )

  dt <- data.table( cutoff = c,
                    train  = cm_train$overall[["Accuracy"]],
                    test   = cm_test$overall[["Accuracy"]] )

  return(dt)
}) %>% rbindlist()

# visualize the accuracy of the train and test set for different cutoff value
# accuracy in percentage.
accuracy_long <- gather( accuracy, "data", "accuracy", -1 )

plot <- ggplot( accuracy_long, aes( cutoff, accuracy, group = data, color = data ) ) +
  geom_line( size = 1 ) + geom_point( size = 3 ) +
  scale_y_continuous( label = percent ) +
  ggtitle( "Train/Test Accuracy for Different Cutoff" ) +
  scale_x_continuous(breaks=seq(0, 1, 0.1)) +
  theme_bw()

return( list( data = accuracy, plot = plot ) )
}

#-----

# -----
# [ConfusionMatrixInfo] :
# Obtain the confusion matrix plot and data.table for a given
# dataset that already consists the predicted score and actual outcome.
# @data      : your data.table or data.frame type data that consists the column
#              of the predicted score and actual outcome
# @predict   : predicted score's column name
# @actual    : actual results' column name
# @cutoff    : cutoff value for the prediction score
# return     : 1. data : a data.table consisting of three column
#               the first two stores the original value of the prediction and actual outcome from
#               the passed in data frame, the third indicates the type, which is after choosing
#               cutoff value, will this row be a true/false positive/ negative
#             2. plot : plot that visualizes the data.table

ConfusionMatrixInfo <- function( data, predict, actual, cutoff )
{

```

```

# extract the column ;
# relevel making 1 appears on the more commonly seen position in
# a two by two confusion matrix
predict <- data[[predict]]
temp_data <- as.factor( as.numeric(data[[actual]]) )
levels(temp_data) <- c(0,1)
actual <- relevel(temp_data, "1")

result <- data.table( actual = actual, predict = predict )

# caculating each pred falls into which category for the confusion matrix
result[, type := ifelse( predict >= cutoff & actual == 1, "TP",
                        ifelse( predict >= cutoff & actual == 0, "FP",
                                ifelse( predict < cutoff & actual == 1, "FN", "TN" ) ) ) %>% as.factor()]

# jittering : can spread the points along the x axis
plot <- ggplot( result, aes( actual, predict, color = type ) ) +
  geom_violin( fill = "white", color = NA ) +
  geom_jitter( shape = 1 ) +
  geom_hline( yintercept = cutoff, color = "blue", alpha = 0.6 ) +
  scale_y_continuous( limits = c( 0, 1 ) ) +
  scale_color_discrete( breaks = c( "TP", "FN", "FP", "TN" ) ) + # ordering of the legend
  guides( col = guide_legend( nrow = 2 ) ) + # adjust the legend to have two rows
  ggtitle( sprintf( "Confusion Matrix with Cutoff at %.2f", cutoff ) )

return( list( data = result, plot = plot ) )
}

# -----
# [ROCInfo] :
# Pass in the data that already consists the predicted score and actual outcome.
# to obtain the ROC curve
# @data      : your data.table or data.frame type data that consists the column
#              of the predicted score and actual outcome
# @predict   : predicted score's column name
# @actual    : actual results' column name
# @cost.fp   : associated cost for a false positive
# @cost.fn   : associated cost for a false negative
# return     : a list containing
#              1. plot           : a side by side roc and cost plot, title showing optimal cutoff value
#                                title showing optimal cutoff, total cost, and area under the curve (auc)
#              2. cutoff        : optimal cutoff value according to the specified fp/fn cost
#              3. totalcost     : total cost according to the specified fp/fn cost
#              4. auc           : area under the curve
#              5. sensitivity    : TP / (TP + FN)
#              6. specificity    : TN / (FP + TN)

ROCInfo <- function( data, predict, actual, cost.fp, cost.fn )
{
  # calculate the values using the ROCR library
  # true positive, false postive

```

```

pred <- prediction( data[[predict]], data[[actual]] )
perf <- performance( pred, "tpr", "fpr" )
roc_dt <- data.frame( fpr = perf@x.values[[1]], tpr = perf@y.values[[1]] )

# cost with the specified false positive and false negative cost
# false positive rate * number of negative instances * false positive cost +
# false negative rate * number of positive instances * false negative cost
cost <- perf@x.values[[1]] * cost.fp * sum( data[[actual]] == 0 ) +
  ( 1 - perf@y.values[[1]] ) * cost.fn * sum( data[[actual]] == 1 )

cost_dt <- data.frame( cutoff = pred@cutoffs[[1]], cost = cost )

# optimal cutoff value, and the corresponding true positive and false positive rate
best_index <- which.min(cost)
best_cost <- cost_dt[ best_index, "cost" ]
best_tpr <- roc_dt[ best_index, "tpr" ]
best_fpr <- roc_dt[ best_index, "fpr" ]
best_cutoff <- pred@cutoffs[[1]][ best_index ]

# area under the curve
auc <- performance( pred, "auc" )@y.values[[1]]

# normalize the cost to assign colors to 1
normalize <- function(v) ( v - min(v) ) / diff( range(v) )

# create color from a palette to assign to the 100 generated threshold between 0 ~ 1
# then normalize each cost and assign colors to it, the higher the blacker
# don't times it by 100, there will be 0 in the vector
col_ramp <- colorRampPalette( c( "green", "orange", "red", "black" ) )(100)
col_by_cost <- col_ramp[ ceiling( normalize(cost) * 99 ) + 1 ]

roc_plot <- ggplot( roc_dt, aes( fpr, tpr ) ) +
  geom_line( color = rgb( 0, 0, 1, alpha = 0.3 ) ) +
  geom_point( color = col_by_cost, size = 4, alpha = 0.2 ) +
  geom_segment( aes( x = 0, y = 0, xend = 1, yend = 1 ), alpha = 0.8, color = "royalblue" ) +
  labs( title = "ROC", x = "False Positive Rate", y = "True Positive Rate" ) +
  geom_hline( yintercept = best_tpr, alpha = 0.8, linetype = "dashed", color = "steelblue4" ) +
  geom_vline( xintercept = best_fpr, alpha = 0.8, linetype = "dashed", color = "steelblue4" ) +
  theme_bw()

cost_plot <- ggplot( cost_dt, aes( cutoff, cost ) ) +
  geom_line( color = "blue", alpha = 0.5 ) +
  geom_point( color = col_by_cost, size = 4, alpha = 0.5 ) +
  ggtitle( "Cost" ) +
  scale_y_continuous( labels = comma ) +
  geom_vline( xintercept = best_cutoff, alpha = 0.8, linetype = "dashed", color = "steelblue4" ) +
  theme_bw()

# the main title for the two arranged plot
sub_title <- sprintf( "Cutoff at %.2f - Total Cost = %.2f, AUC = %.3f",
  best_cutoff, best_cost, auc )

# arranged into a side by side plot

```

```

plot <- arrangeGrob( roc_plot, cost_plot, ncol = 2,
                    top = textGrob( sub_title, gp = gpar( fontsize = 16, fontface = "bold" ) ) )

return( list( plot      = plot,
              cutoff     = best_cutoff,
              totalcost  = best_cost,
              auc         = auc,
              sensitivity = best_tpr,
              specificity = 1 - best_fpr ) )
}

#delete_dup

#Some variables are forced into the model regardless of variable selection result
#If the forced variable ended up being selected, this model will remove the duplicated variable.

delete_dup <- function(subset, data){
  remove <- c()
  for(i in 1:length(subset)){
    result <- str_detect(subset[i],names(data))
    for(j in 1:length(result)){
      if(result[j]){
        remove <- c(remove,i)
      }
    }
  }
  if(is.null(remove))
    return(subset)
  subset <- subset[-c(remove)]
  return(subset)
}

#data = data file
#Prediction: predicted result
#response: The name of response variable
#cut_off: probability cut off point

Classify <- function(data, prediction, response, cut_off ){
  for(i in 1:length(prediction)){
    if(prediction[i] < cut_off){
      prediction[i] <- levels(data[[response]])[1]
    } else{
      prediction[i] <- levels(data[[response]])[2]
    }
  }
}

prediction <- as.factor(prediction)
levels(prediction) <- c(levels(data[[response]])[1],levels(data[[response]])[2])
confusion_matrix <- table(data[[response]],prediction)
print(confusion_matrix)
Accuracy <- (confusion_matrix[1,1] + confusion_matrix[2,2])/sum(confusion_matrix)
TPR <- confusion_matrix[2,2] / (confusion_matrix[2,2] + confusion_matrix[2,1])

```

```

    return(cat(paste("The accuracy is", round(Accuracy*100,3), "%.\n", "The True positive rate is", round(TPR*
  })

#K fold K = 10

#data = data using for prediction
#response = name of the response variable
#cut off = probability cut off point
#interaction = you can type addition interaction term in text
#Example
#cv.error(CNP_logi_subset, "Subject_Type", "+Age*Auditory.global_eff", 0.8)

cv.error <- function(data, response, interaction = "", cut_off = 0.5){

  #generate random seeds
  r <- runif(1,0,9999)
  set.seed(r)
  folds <- createFolds(data[[response]],k = 10)
  Accuracy <- rep(NA,10)
  TPR <- rep(NA,10)

  for(i in 1:10){

    #training and testing
    train <- data[-folds[[i]],]
    test <- data[folds[[i]],]

    levels(test[[response]]) <- c(levels(data[[response]])[1],levels(data[[response]])[2])

    logi_cv <- glm(paste(response,"~.",interaction), data = train, family = "binomial")

    prediction <- predict(logi_cv, test, type = "response")
    for(j in 1:length(prediction)){
      if(prediction[j] < cut_off){
        prediction[j] <- levels(test[[response]])[1]
      } else{
        prediction[j] <- levels(test[[response]])[2]
      }
    }
    prediction <- as.factor(prediction)
    levels(prediction) <- c(levels(data[[response]])[1],levels(data[[response]])[2])

    confuseion_matrix <- table(test[[response]],prediction)
    Accuracy[i] <- (confuseion_matrix[1,1] + confuseion_matrix[2,2])/sum(confuseion_matrix)
    TPR[i] <- confuseion_matrix[2,2] / (confuseion_matrix[2,2] + confuseion_matrix[2,1])
  }
  return(list(Accuracy, TPR))
}

```

```
#Standardized variable
```

```
Standarize <- function(data){  
  for(i in 1:ncol(data)){  
    if(is.numeric(data[,i])){  
      data[,i] <- (data[,i] - mean(data[,i]))/sd(data[,i])  
    }  
  }  
  return(data)  
}
```

```
#Load data
```

```
setwd("A:/Winter 2018/Stats 141SL/project/")
```

```
#load CNP data
```

```
CNP_between <- read.table("CNP_between_nets.txt", header = TRUE)  
CNP_within <- read.table("CNP_within_nets.txt", header = TRUE)  
CNPDemographic <- read.xlsx("CNPDemographicMeasures.xlsx", sheetName = "SNF")
```

```
#load COBRE data
```

```
COBRE_between <- read.table("COBRE_between_nets.txt", header = TRUE)  
COBRE_within <- read.table("COBRE_within_nets.txt", header = TRUE)  
COBREDemographic <- read.xlsx("COBRE INDI Additional data.xls", sheetName = "NP")  
COBRE_phenotypic <- read_csv("COBRE_phenotypic_data.csv")
```

```
## Parsed with column specification:  
## cols(  
##   X1 = col_integer(),  
##   `Current Age` = col_character(),  
##   Gender = col_character(),  
##   Handedness = col_character(),  
##   `Subject Type` = col_character(),  
##   Diagnosis = col_character()  
## )
```

```
#Data cleaning process
```

```
#Removed character string
```

```
pattern <- "[a-z]*-"
```

```
CNP_within$Subject_ID <- as.numeric(str_replace_all(CNP_within$Subject_ID,  
  , pattern, ""))
```

```
CNP_between$Subject_ID <- as.numeric(str_replace_all(CNP_between$Subject_ID,  
  , pattern, ""))
```

```

#Merge data
CNP_within_merge <- left_join(CNP_within,CNPDemographic, by = c("Subject_ID" = "PTID"))

#summary(CNP_within_merge)

CNP_between_merge <- left_join(CNP_between,CNPDemographic, by = c("Subject_ID" = "PTID"))

#summary(CNP_between_merge)


#Reumove character string

COBRE_between$Subject_ID <- as.numeric(str_replace_all(COBRE_between$Subject_ID
, pattern,""))

COBRE_within$Subject_ID <- as.numeric(str_replace_all(COBRE_within$Subject_ID
, pattern,""))


#remove 00

pattern <- "^00"

COBREDemographic$ID <- as.numeric(str_replace_all(COBREDemographic$ID, pattern,""))

#Merge data

COBRE_within_merge <- left_join(COBRE_within,COBREDemographic, by = c("Subject_ID" = "ID"))

#summary(COBRE_within_merge)

COBRE_between_merge <- left_join(COBRE_between,COBREDemographic, by = c("Subject_ID" = "ID"))

#summary(COBRE_between_merge)

COBRE_phenotypic$Gender <- as.factor(COBRE_phenotypic$Gender)

COBRE_phenotypic <- COBRE_phenotypic %>%
  filter(!(COBRE_phenotypic$Gender == "Disenrolled"))

## Warning: package 'bindrcpp' was built under R version 3.4.2
COBRE_phenotypic$Gender <- droplevels(COBRE_phenotypic$Gender)

colnames(COBRE_phenotypic)[1:2] <- c("Subject_ID", "Age")

```



```
COBRE_between_merge <- merge(COBRE_between_merge, COBRE_phenotypic, all = TRUE)
COBRE_within_merge <- merge(COBRE_within_merge, COBRE_phenotypic, all = TRUE)
```

```
table(COBRE_between_merge$Diagnosis)
```

```
##
##           290.3           295.1           295.2
##           1           3           1
##           295.3           295.6           295.7
##           41           12           5
## 295.70 bipolar type 295.70 depressed type 295.9
##           1           1           5
##           295.92           296.26           296.4
##           1           1           1
##           311           None
##           1           72
```

```
table(COBRE_within_merge$Diagnosis)
```

```
##
##           290.3           295.1           295.2
##           1           3           1
##           295.3           295.6           295.7
##           41           12           5
## 295.70 bipolar type 295.70 depressed type 295.9
##           1           1           5
##           295.92           296.26           296.4
##           1           1           1
##           311           None
##           1           72
```

```
#CNP filter
```

```
CNP_within_merge <- CNP_within_merge %>%
  filter(Subject_Type == "Control" | Subject_Type == "Schizophrenia")
```

```
table(CNP_within_merge$Subject_Type)
```

```
##
##      ADHD      Bipolar      Control Schizophrenia
##      0         0         115         42
```

```
CNP_between_merge <- CNP_between_merge %>%
  filter(Subject_Type == "Control" | Subject_Type == "Schizophrenia")
```

```
table(CNP_between_merge$Subject_Type)
```

```
##
##      ADHD      Bipolar      Control Schizophrenia
##      0         0         115         42
```

```
#COBRE filter
```

```
COBRE_between_merge <- COBRE_between_merge %>%
```

```

filter(!(Diagnosis == 290.3 | Diagnosis == 296.26 | Diagnosis == 296.4 | Diagnosis == 311))

COBRE_within_merge <- COBRE_within_merge %>%
  filter(!(Diagnosis == 290.3 | Diagnosis == 296.26 | Diagnosis == 296.4 | Diagnosis == 311))

table(COBRE_between_merge$Diagnosis)

##
##           295.1           295.2           295.3
##           3           1           41
##           295.6           295.7   295.70 bipolar type
##           12           5           1
## 295.70 depressed type           295.9           295.92
##           1           5           1
##           None
##           72

table(COBRE_within_merge$Diagnosis)

##
##           295.1           295.2           295.3
##           3           1           41
##           295.6           295.7   295.70 bipolar type
##           12           5           1
## 295.70 depressed type           295.9           295.92
##           1           5           1
##           None
##           72

#Recoding Patients to Schizophrenia in COBRE

pattern <- "Patient"

COBRE_between_merge$Subject_Type <- str_replace_all(COBRE_between_merge$Subject_Type, pattern, "Schizophr")
COBRE_within_merge$Subject_Type <- str_replace_all(COBRE_within_merge$Subject_Type, pattern, "Schizophr")

table(COBRE_between_merge$Subject_Type)

##
##      Control Schizophrenia
##      72      70

table(COBRE_within_merge$Subject_Type)

##
##      Control Schizophrenia
##      72      70

CNP_between_merge$Subject_Type <- droplevels(CNP_between_merge$Subject_Type)
levels(CNP_between_merge$Subject_Type)

## [1] "Control"      "Schizophrenia"

CNP_within_merge$Subject_Type <- droplevels(CNP_within_merge$Subject_Type)
levels(CNP_within_merge$Subject_Type)

## [1] "Control"      "Schizophrenia"

```

```

#CNP between
#remove 96:98, 112
CNP_between_merge <- CNP_between_merge %>%
  select(-c(96:98,112))

#CNP within get rid of
#75 #76 #91
CNP_within_merge <- CNP_within_merge %>%
  select(-c(75:77,91))

#Merge both between and within data into CNP
CNP <- merge(CNP_between_merge,CNP_within_merge, all = TRUE)

CNP_RF_subset <- CNP %>%
  select(-c(1,5:41))

#Merge both between and within into COBRE
COBRE <- merge(COBRE_between_merge, COBRE_within_merge, all = TRUE)

#Use only the fMRI, MRI, and Age, keep global EFF
COBRE_RF_subset<- COBRE %>%
  select(-c(1,5:111))

COBRE_RF_subset$Subject_Type <- as.factor(COBRE_RF_subset$Subject_Type)

#CNP data modeling
set.seed(4321)

rfsrc_m1 <- rfsrc(as.factor(Subject_Type)~.,data = CNP_RF_subset, na.action = c("na.omit"), ntree= 1000)

max_var <- max.subtree(rfsrc_m1, conservative = TRUE)
max_var$topvars

## [1] "Ventral_Attention.Uncertain"
## [2] "Cingulo.opercular_Task_Control.mod"
#delete duplicate entity

#Logistic Regression Model
subset1 <- as.vector(max_var$topvars)

```

```

subset1 <- delete_dup(subset1,CNP_RF_subset[,c(1,137:150)])

CNP_logi_subset <- CNP_RF_subset[,c("Subject_Type",names(CNP_RF_subset[,c(1,137:150)])), subset1]]

#Using a previously grown forest, identify pairwise interactions for all pairs of variables from a spec

#method="maxsubtree"

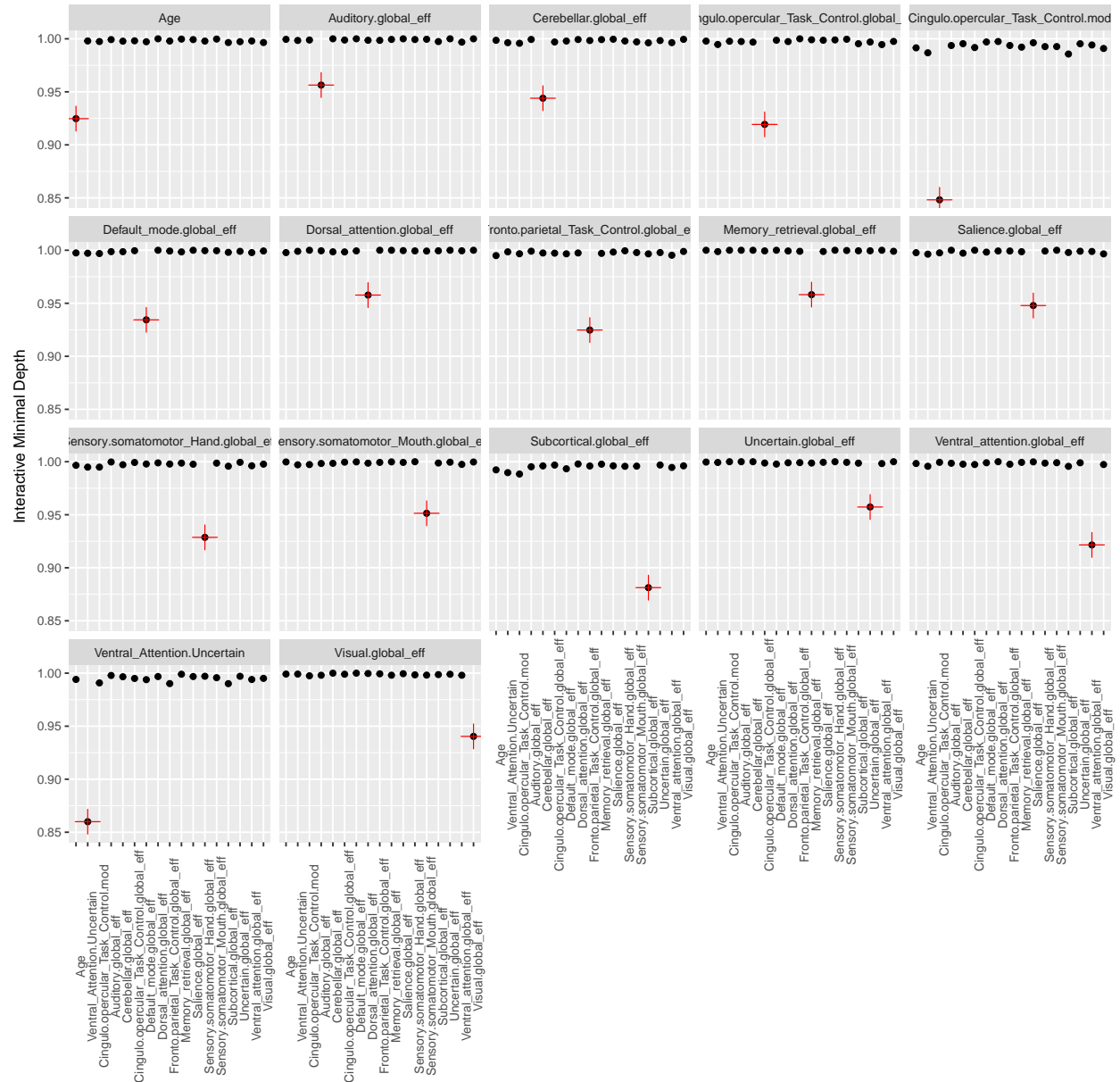
#This invokes a maximal subtree analysis.

CNP_logi_subset <- na.omit(CNP_logi_subset) %>%
  Standarize()

#Find interaction
gg_int <- gg_interaction(find.interaction(rfsrc_m1,
                                         xvar.names = names(CNP_logi_subset[, -c(1)]),
                                         sorted = FALSE,
                                         verbose = FALSE))

plot(gg_int)

```



*#Minimal depth variable interaction plot for all variables of interest.*

*#Higher values indicate lower interactivity with target variable marked in red.*

*#No interaction found based on the result, we don't have to add interaction term*

*#Correlation check*

```
high_cor <- findCorrelation(cor(CNP_logi_subset[, -c(1:2)]), cutoff = 0.75) + 2
```

*#No potential multicollinearity problem*

```
index <- sample(1:nrow(CNP_logi_subset), size = round(nrow(CNP_logi_subset)*0.7,0), replace = FALSE)
```

```
CNP_train <- CNP_logi_subset[index,]
```

```

CNP_test <- CNP_logi_subset[-index,]
logi_m1 <-glm(Subject_Type~. , data = CNP_train, family = "binomial")
summary(logi_m1)

##
## Call:
## glm(formula = Subject_Type ~ ., family = "binomial", data = CNP_train)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.6210  -0.6679  -0.3474   0.5809   3.0634
##
## Coefficients:
##              Estimate Std. Error z value
## (Intercept)    -1.73948    0.36956  -4.707
## Age              0.83260    0.34036   2.446
## Auditory.global_eff -0.09186    0.35094  -0.262
## Cerebellar.global_eff  0.15612    0.27872   0.560
## Cingulo.opercular_Task_Control.global_eff  0.03363    0.33229   0.101
## Default_mode.global_eff  0.87062    0.42690   2.039
## Dorsal_attention.global_eff  0.43827    0.31808   1.378
## Fronto.parietal_Task_Control.global_eff  0.38999    0.42524   0.917
## Memory_retrieval.global_eff -0.02610    0.29618  -0.088
## Salience.global_eff  0.36833    0.34606   1.064
## Sensory.somatomotor_Hand.global_eff  0.30430    0.50027   0.608
## Sensory.somatomotor_Mouth.global_eff -0.40188    0.37858  -1.062
## Subcortical.global_eff  0.56609    0.36193   1.564
## Uncertain.global_eff  0.78505    0.34559   2.272
## Ventral_attention.global_eff -0.20361    0.31489  -0.647
## Visual.global_eff  0.49797    0.38277   1.301
## Ventral_Attention.Uncertain -2.28226    0.67824  -3.365
## Cingulo.opercular_Task_Control.mod  0.88433    0.32735   2.701
##
##              Pr(>|z|)
## (Intercept)    2.51e-06 ***
## Age            0.014436 *
## Auditory.global_eff  0.793522
## Cerebellar.global_eff  0.575381
## Cingulo.opercular_Task_Control.global_eff  0.919375
## Default_mode.global_eff  0.041408 *
## Dorsal_attention.global_eff  0.168252
## Fronto.parietal_Task_Control.global_eff  0.359080
## Memory_retrieval.global_eff  0.929783
## Salience.global_eff  0.287175
## Sensory.somatomotor_Hand.global_eff  0.543008
## Sensory.somatomotor_Mouth.global_eff  0.288451
## Subcortical.global_eff  0.117797
## Uncertain.global_eff  0.023110 *
## Ventral_attention.global_eff  0.517884
## Visual.global_eff  0.193274
## Ventral_Attention.Uncertain  0.000766 ***
## Cingulo.opercular_Task_Control.mod  0.006903 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##

```

```
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 130.826 on 109 degrees of freedom
## Residual deviance: 91.494 on 92 degrees of freedom
## AIC: 127.49
##
## Number of Fisher Scoring iterations: 6
```

```
round(exp(coef(logi_m1)),3)
```

```
## (Intercept)
## 0.176
## Age
## 2.299
## Auditory.global_eff
## 0.912
## Cerebellar.global_eff
## 1.169
## Cingulo.opercular_Task_Control.global_eff
## 1.034
## Default_mode.global_eff
## 2.388
## Dorsal_attention.global_eff
## 1.550
## Fronto.parietal_Task_Control.global_eff
## 1.477
## Memory_retrieval.global_eff
## 0.974
## Salience.global_eff
## 1.445
## Sensory.somatomotor_Hand.global_eff
## 1.356
## Sensory.somatomotor_Mouth.global_eff
## 0.669
## Subcortical.global_eff
## 1.761
## Uncertain.global_eff
## 2.193
## Ventral_attention.global_eff
## 0.816
## Visual.global_eff
## 1.645
## Ventral_Attention.Uncertain
## 0.102
## Cingulo.opercular_Task_Control.mod
## 2.421
```

```
anova(logi_m1, test = "Chisq")
```

```
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: Subject_Type
##
```

```
## Terms added sequentially (first to last)
##
##
##
## Df Deviance Resid. Df Resid. Dev
## NULL 109 130.826
## Age 1 3.5821 108 127.244
## Auditory.global_eff 1 0.3403 107 126.903
## Cerebellar.global_eff 1 0.2195 106 126.684
## Cingulo.opercular_Task_Control.global_eff 1 0.0091 105 126.675
## Default_mode.global_eff 1 0.0117 104 126.663
## Dorsal_attention.global_eff 1 0.2303 103 126.433
## Fronto.parietal_Task_Control.global_eff 1 0.6998 102 125.733
## Memory_retrieval.global_eff 1 0.0689 101 125.664
## Salience.global_eff 1 0.0166 100 125.647
## Sensory.somatomotor_Hand.global_eff 1 1.5916 99 124.056
## Sensory.somatomotor_Mouth.global_eff 1 6.0241 98 118.032
## Subcortical.global_eff 1 0.0072 97 118.025
## Uncertain.global_eff 1 1.8716 96 116.153
## Ventral_attention.global_eff 1 1.1439 95 115.009
## Visual.global_eff 1 0.1207 94 114.888
## Ventral_Attention.Uncertain 1 13.8856 93 101.003
## Cingulo.opercular_Task_Control.mod 1 9.5089 92 91.494
## Pr(>Chi)
## NULL
## Age 0.0584057 .
## Auditory.global_eff 0.5596684
## Cerebellar.global_eff 0.6394285
## Cingulo.opercular_Task_Control.global_eff 0.9240192
## Default_mode.global_eff 0.9137170
## Dorsal_attention.global_eff 0.6313287
## Fronto.parietal_Task_Control.global_eff 0.4028610
## Memory_retrieval.global_eff 0.7929193
## Salience.global_eff 0.8975062
## Sensory.somatomotor_Hand.global_eff 0.2071033
## Sensory.somatomotor_Mouth.global_eff 0.0141116 *
## Subcortical.global_eff 0.9321680
## Uncertain.global_eff 0.1712969
## Ventral_attention.global_eff 0.2848198
## Visual.global_eff 0.7282840
## Ventral_Attention.Uncertain 0.0001943 ***
## Cingulo.opercular_Task_Control.mod 0.0020447 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
#R-squared
```

```
R_squared <- 1 - (summary(logi_m1)[[4]]/summary(logi_m1)[[8]])
R_squared
```

```
## [1] 0.3006434
```

```
#70/30 CV check
```

```
#Train
```

```
CNP_train$prediction <- predict(logi_m1, CNP_train, type = "response")
```



```

#Test
CNP_test$prediction <- predict(logi_m1, CNP_test, type = "response")

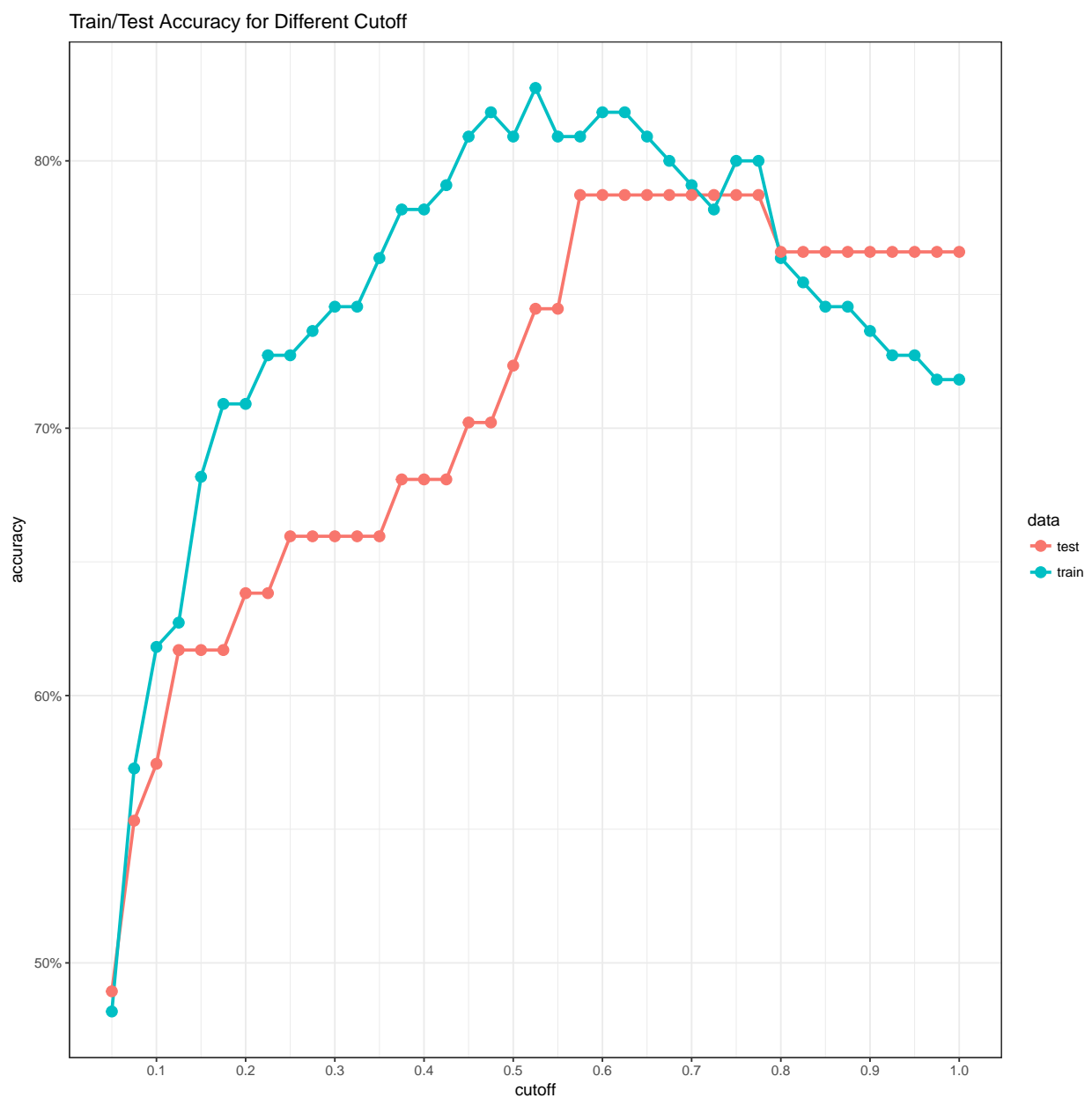
prop.table(table(CNP$Subject_Type))

##
##      Control Schizophrenia
##      0.7324841    0.2675159

accuracy_info <- AccuracyCutoffInfo( train = CNP_train, test = CNP_test,
                                     predict = "prediction", actual = "Subject_Type" )

accuracy_info$plot

```



```
Classify(CNP_train, CNP_train$prediction, "Subject_Type", 0.75 )
```

```
##           prediction
##           Control Schizophrenia
## Control           79           0
## Schizophrenia      22           9
## The accuracy is 80 %.
## The True positive rate is 29.032 %
```

```
Classify(CNP_test, CNP_test$prediction, "Subject_Type", 0.75 )
```

```
##           prediction
##           Control Schizophrenia
## Control           36           0
## Schizophrenia      10           1
## The accuracy is 78.723 %.
## The True positive rate is 9.091 %
```

```
set.seed(4321)
```

```
#CNP ROC search for better True positive rate.
```

```
#cutoff : Optimal cutoff value according to the specified FP and FN cost .
```

```
#totalcost : Total cost according to the specified FP and FN cost.
```

```
#auc : Area under the curve.
```

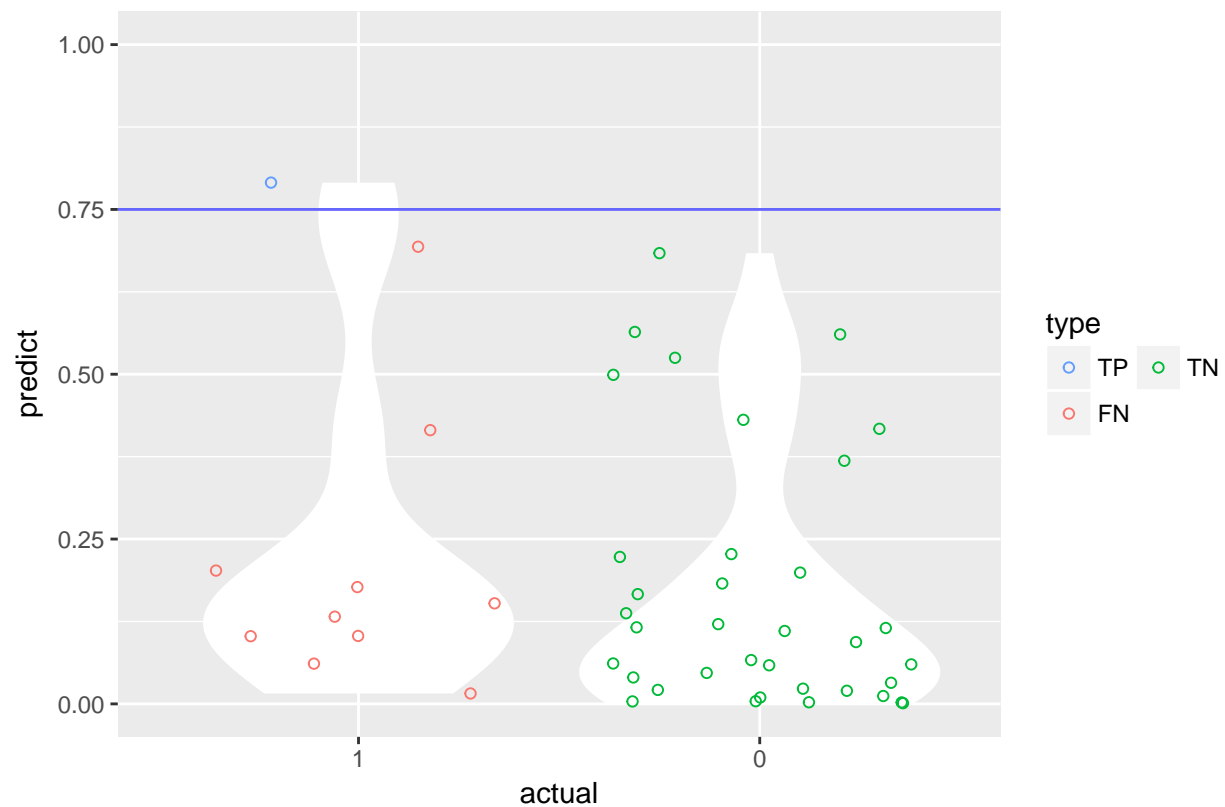
```
#sensitivity : TP / (TP + FN) for the optimal cutoff.
```

```
#specificity : TN / (FP + TN) for the optimal cutoff.
```

```
cm_info <- ConfusionMatrixInfo(data = CNP_test, predict = "prediction", actual = "Subject_Type", 0.75)
```

```
cm_info$plot
```

Confusion Matrix with Cutoff at 0.75



```
invisible(dev.off())

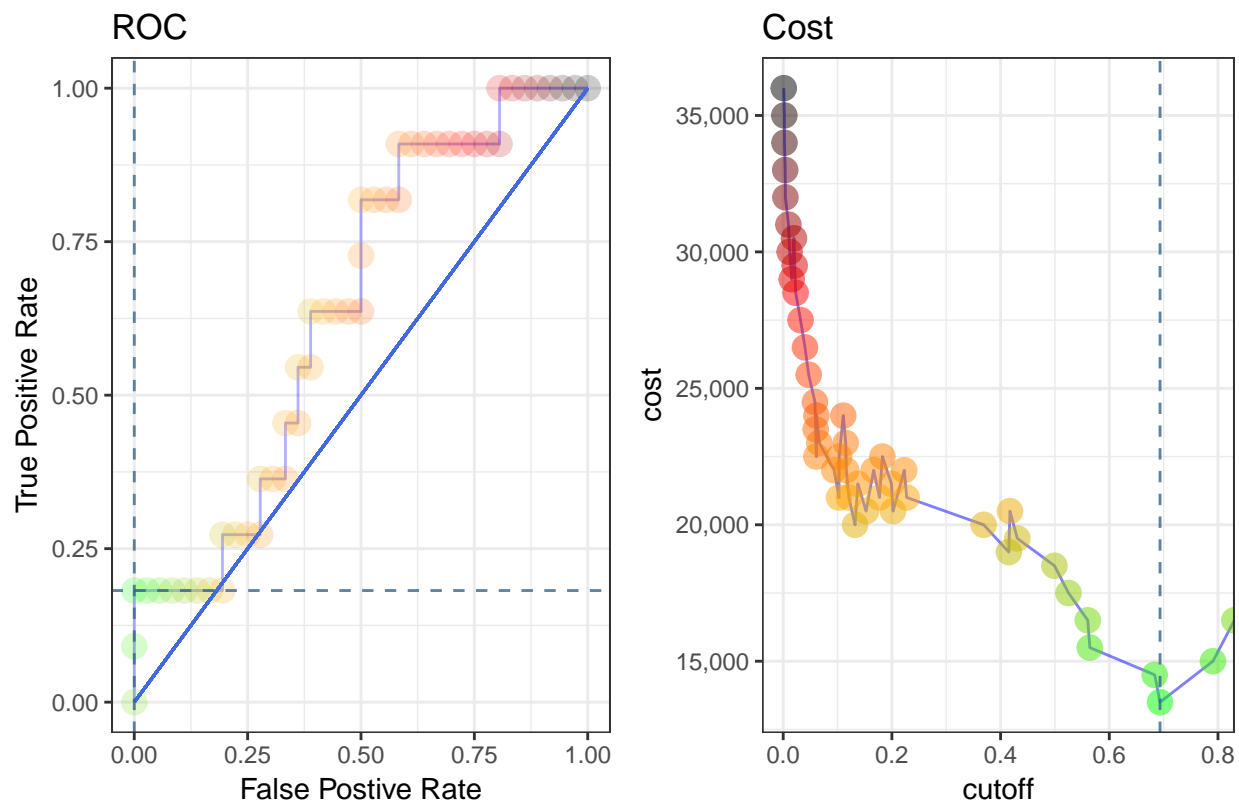
roc_info <- ROCInfo( data = cm_info$data, predict = "predict",
                    actual = "actual", cost.fp = 1000, cost.fn = 1500 )

#Optimal cutoff for True positive rate
roc_info$cutoff

## [1] 0.6933646

grid.draw(roc_info$plot)
```

**Cutoff at 0.69 – Total Cost = 13500.00, AUC = 0.641**



```
#CNP model k fold CV check
set.seed(4321)
```

```
#Optimal cutoff for Accuracy
result <- cv.error(CNP_logi_subset, "Subject_Type", cut_off = roc_info$cutoff)
Accuracy.k <- result[[1]]
mean(Accuracy.k)
```

```
## [1] 0.7263725
```

```
TTP.k <- result[[2]]
mean(TTP.k)
```

```
## [1] 0.15
```

```
#Optimal cutoff for True positive rate
result <- cv.error(CNP_logi_subset, "Subject_Type", cut_off = roc_info$cutoff)
Accuracy.k <- result[[1]]
mean(Accuracy.k)
```

```
## [1] 0.7401716
```

```
TTP.k <- result[[2]]
mean(TTP.k)
```

```
## [1] 0.195
```

```
set.seed(4321)
```

```

#Random Forest variable section
rfsrc_m2 <- rfsrc(Subject_Type~.,data = COBRE_RF_subset, na.action = c("na.omit"), ntree= 1000)

max_var <- max.subtree(rfsrc_m2, conservative = TRUE)
max_var$topvars

## [1] "Visual.Subcortical"
#delete duplicate entity

subset2 <- as.vector(max_var$topvars)

subset2 <- delete_dup(subset2,COBRE_RF_subset[,c(1,137:150)])

#Logistic Regression model

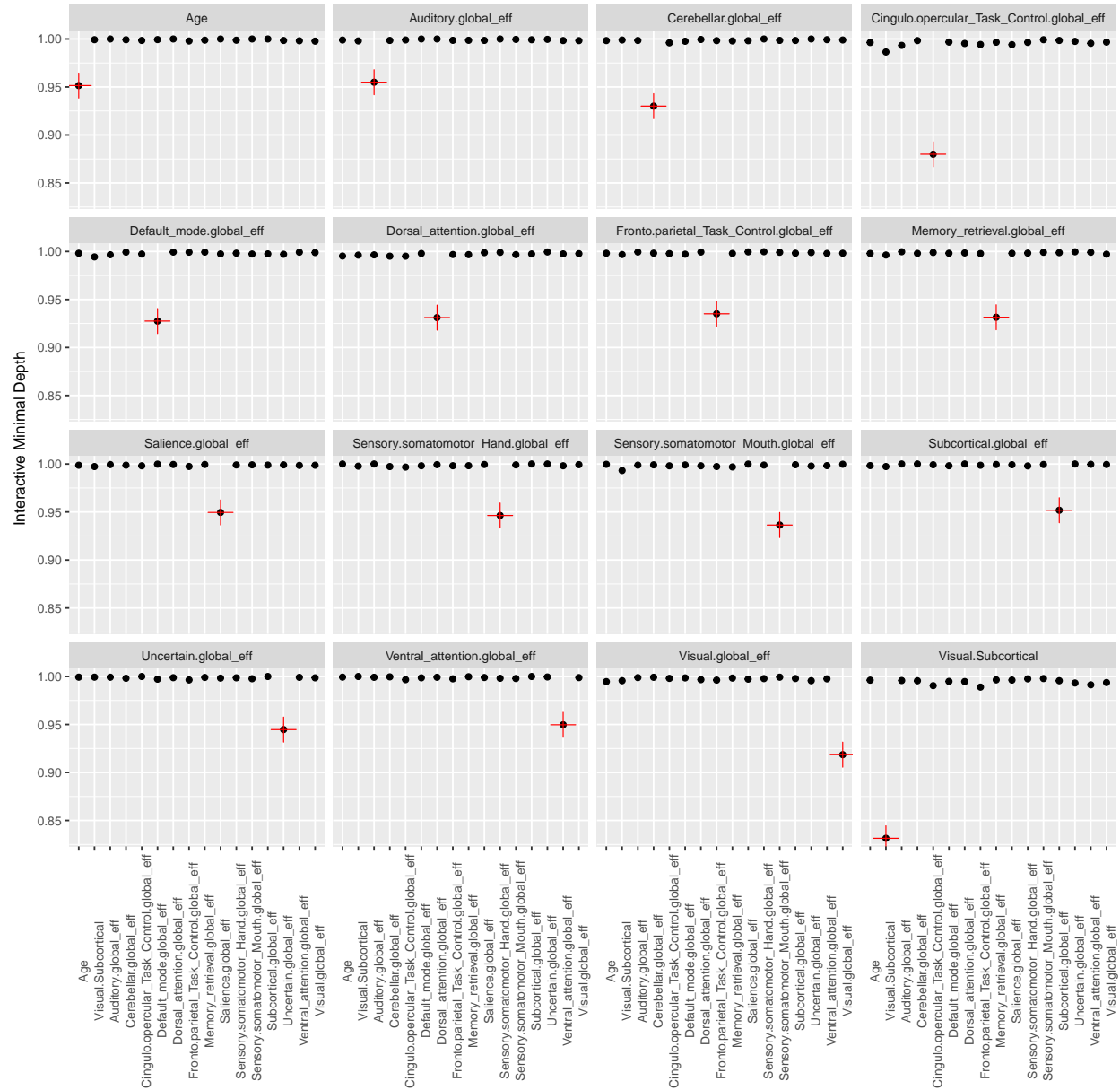
COBRE_logi_subset <- COBRE_RF_subset[,c("Subject_Type",names(COBRE_RF_subset[,c(1,137:150)]), subset2)]

COBRE_logi_subset <- na.omit(COBRE_logi_subset) %>%
  Standarize()

#Find interaction
gg_int <- gg_interaction(find.interaction(rfsrc_m2,
                                         xvar.names = names(COBRE_logi_subset[, -c(1)]),
                                         sorted = FALSE,
                                         verbose = FALSE))

plot(gg_int)

```



#No interaction in fund base on the result, we don't have to add interaction term

#Correlation check

```
high_cor <- findCorrelation(cor(COBRE_logi_subset[, -c(1:2)]), cutoff = 0.75) + 2
```

#No potential multicollinearity problem

```
index <- sample(1:nrow(COBRE_logi_subset), size = round(nrow(COBRE_logi_subset)*0.7, 0), replace = FALSE)
```

```
COBRE_train <- COBRE_logi_subset[index,]
```

```
COBRE_test <- COBRE_logi_subset[-index,]
```

```
logi_m2 <- glm(Subject_Type ~ ., data = COBRE_train, family = "binomial")
```

```
summary(logi_m2)
```

```
##
## Call:
## glm(formula = Subject_Type ~ ., family = "binomial", data = COBRE_train)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.7740  -0.8927  -0.3029   0.8723   2.7946
##
## Coefficients:
##                  Estimate Std. Error z value
## (Intercept)      -0.13280    0.24848  -0.534
## Age              -0.47861    0.31704  -1.510
## Auditory.global_eff    0.27499    0.28158   0.977
## Cerebellar.global_eff -0.02965    0.27095  -0.109
## Cingulo.opercular_Task_Control.global_eff -0.64555    0.33684  -1.916
## Default_mode.global_eff -0.41007    0.35894  -1.142
## Dorsal_attention.global_eff -0.16624    0.28377  -0.586
## Fronto.parietal_Task_Control.global_eff -0.59059    0.28493  -2.073
## Memory_retrieval.global_eff -0.13569    0.27979  -0.485
## Salience.global_eff -0.20282    0.31132  -0.652
## Sensory.somatomotor_Hand.global_eff    0.10884    0.36779   0.296
## Sensory.somatomotor_Mouth.global_eff -0.30390    0.31851  -0.954
## Subcortical.global_eff -0.27574    0.29112  -0.947
## Uncertain.global_eff    0.35474    0.28462   1.246
## Ventral_attention.global_eff    0.40859    0.29147   1.402
## Visual.global_eff    -0.74487    0.38911  -1.914
## Visual.Subcortical    1.09163    0.34568   3.158
##
##                  Pr(>|z|)
## (Intercept)      0.59303
## Age              0.13114
## Auditory.global_eff    0.32877
## Cerebellar.global_eff    0.91286
## Cingulo.opercular_Task_Control.global_eff 0.05530 .
## Default_mode.global_eff    0.25327
## Dorsal_attention.global_eff    0.55801
## Fronto.parietal_Task_Control.global_eff 0.03820 *
## Memory_retrieval.global_eff    0.62769
## Salience.global_eff    0.51472
## Sensory.somatomotor_Hand.global_eff    0.76728
## Sensory.somatomotor_Mouth.global_eff    0.34001
## Subcortical.global_eff    0.34356
## Uncertain.global_eff    0.21263
## Ventral_attention.global_eff    0.16097
## Visual.global_eff    0.05559 .
## Visual.Subcortical    0.00159 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 136.42  on 98  degrees of freedom
## Residual deviance: 105.62  on 82  degrees of freedom
```

```
## AIC: 139.62
##
## Number of Fisher Scoring iterations: 5
```

```
round(exp(coef(logi_m2)),3)
```

```
##                (Intercept)
##                0.876
##                Age
##                0.620
##          Auditory.global_eff
##                1.317
##          Cerebellar.global_eff
##                0.971
## Cingulo.opercular_Task_Control.global_eff
##                0.524
##          Default_mode.global_eff
##                0.664
##          Dorsal_attention.global_eff
##                0.847
## Fronto.parietal_Task_Control.global_eff
##                0.554
##          Memory_retrieval.global_eff
##                0.873
##          Salience.global_eff
##                0.816
##          Sensory.somatomotor_Hand.global_eff
##                1.115
##          Sensory.somatomotor_Mouth.global_eff
##                0.738
##          Subcortical.global_eff
##                0.759
##          Uncertain.global_eff
##                1.426
##          Ventral_attention.global_eff
##                1.505
##          Visual.global_eff
##                0.475
##          Visual.Subcortical
##                2.979
```

```
anova(logi_m2, test = "Chisq")
```

```
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: Subject_Type
##
## Terms added sequentially (first to last)
##
```

	Df	Deviance	Resid.	Df	Resid. Dev
## NULL			98		136.42
## Age	1	0.0417	97		136.38



```
## Auditory.global_eff      1  0.3108      96  136.07
## Cerebellar.global_eff    1  1.8761      95  134.19
## Cingulo.opercular_Task_Control.global_eff  1  3.9597      94  130.24
## Default_mode.global_eff  1  1.2625      93  128.97
## Dorsal_attention.global_eff  1  0.4660      92  128.51
## Fronto.parietal_Task_Control.global_eff  1  1.0467      91  127.46
## Memory_retrieval.global_eff  1  0.4759      90  126.98
## Salience.global_eff     1  0.0280      89  126.96
## Sensory.somatomotor_Hand.global_eff  1  0.0038      88  126.95
## Sensory.somatomotor_Mouth.global_eff  1  1.9567      87  125.00
## Subcortical.global_eff   1  0.0634      86  124.93
## Uncertain.global_eff     1  1.4932      85  123.44
## Ventral_attention.global_eff  1  2.6970      84  120.74
## Visual.global_eff        1  2.0923      83  118.65
## Visual.Subcortical       1 13.0328      82  105.62
##                           Pr(>Chi)
## NULL
## Age                       0.8382831
## Auditory.global_eff      0.5771834
## Cerebellar.global_eff    0.1707794
## Cingulo.opercular_Task_Control.global_eff 0.0466033 *
## Default_mode.global_eff  0.2611782
## Dorsal_attention.global_eff  0.4948425
## Fronto.parietal_Task_Control.global_eff  0.3062618
## Memory_retrieval.global_eff  0.4902644
## Salience.global_eff     0.8671524
## Sensory.somatomotor_Hand.global_eff  0.9505287
## Sensory.somatomotor_Mouth.global_eff  0.1618715
## Subcortical.global_eff   0.8011719
## Uncertain.global_eff     0.2217196
## Ventral_attention.global_eff  0.1005368
## Visual.global_eff        0.1480421
## Visual.Subcortical       0.0003061 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
#R-squared
```

```
R_squared <- 1 - (summary(logi_m2)[[4]]/summary(logi_m2)[[8]])
R_squared
```

```
## [1] 0.2258151
```

```
#70/30 CV check
```

```
#Train
```

```
COBRE_train$prediction <- predict(logi_m2, COBRE_train, type = "response")
```

```
#Test
```

```
COBRE_test$prediction <- predict(logi_m2, COBRE_test, type = "response")
```

```
prop.table(table(COBRE$Subject_Type))
```

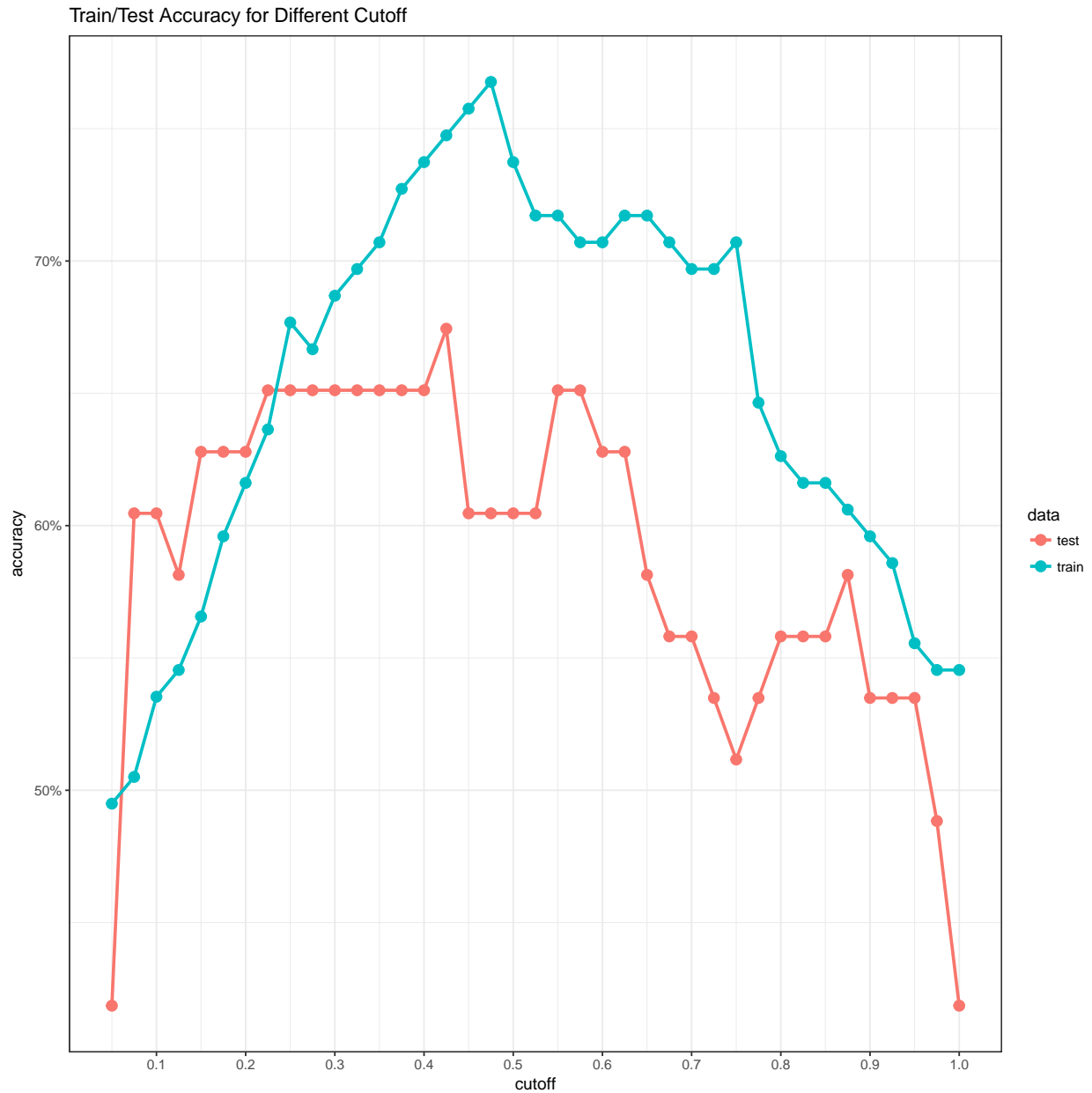
```
##
```

```
##      Control Schizophrenia
```

```
##      0.5070423      0.4929577
```

```
accuracy_info <- AccuracyCutoffInfo( train = COBRE_train, test = COBRE_test,
                                     predict = "prediction", actual = "Subject_Type" )
```

```
accuracy_info$plot
```



```
Classify(COBRE_train, COBRE_train$prediction, "Subject_Type", 0.425)
```

```
##           prediction
##           Control Schizophrenia
## Control           37           17
## Schizophrenia      8           37
## The accuracy is 74.747 %.
## The True positive rate is 82.222 %
```

```
Classify(COBRE_test, COBRE_test$prediction, "Subject_Type", 0.425)
```

```
##           prediction
##           Control Schizophrenia
## Control           9           9
## Schizophrenia      5          20
## The accuracy is 67.442 %.
## The True positive rate is 80 %
```

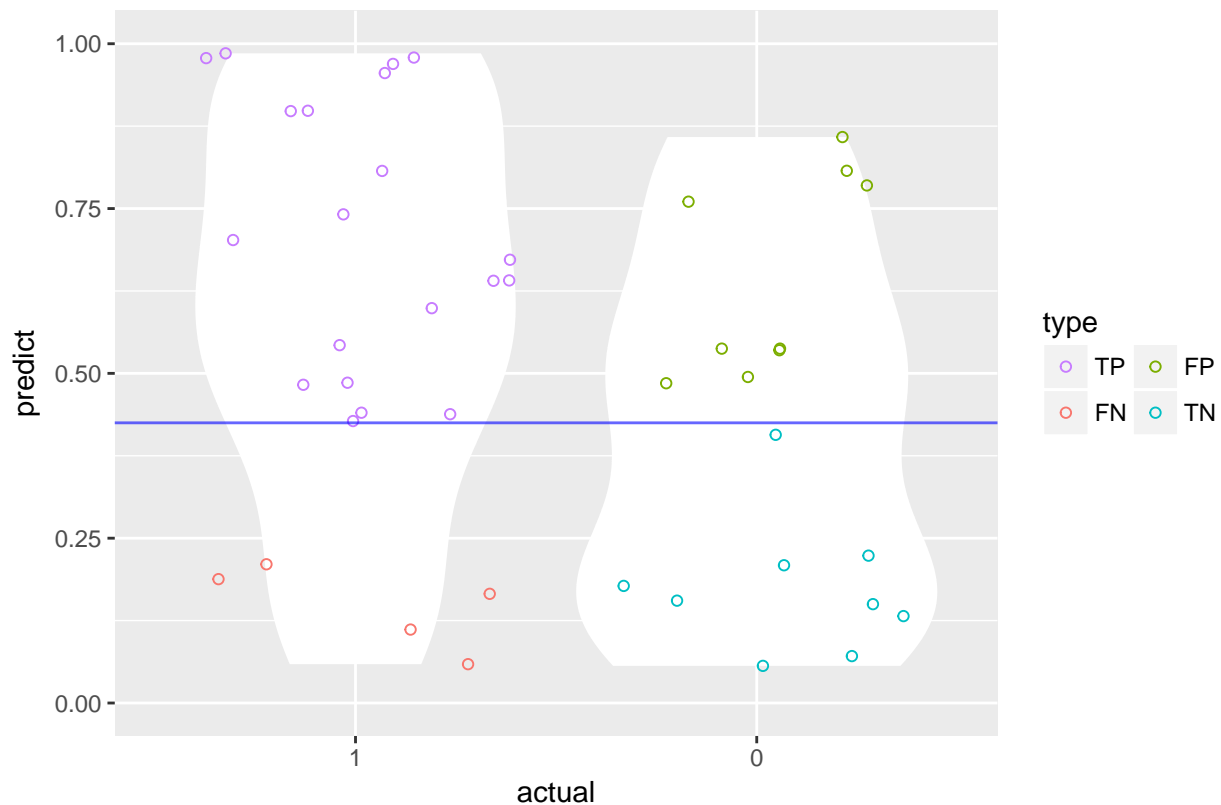
```
#COBRE ROC search for better True positive rate.
```

```
#cutoff : Optimal cutoff value according to the specified FP and FN cost .
#totalcost : Total cost according to the specified FP and FN cost.
#auc : Area under the curve.
#sensitivity : TP / (TP + FN) for the optimal cutoff.
#specificity : TN / (FP + TN) for the optimal cutoff.
```

```
cm_info <- ConfusionMatrixInfo(data = COBRE_test, predict = "prediction", actual = "Subject_Type", 0.425)
```

```
cm_info$plot
```

Confusion Matrix with Cutoff at 0.42



```
invisible(graphics.off())
roc_info <- ROCInfo( data = cm_info$data, predict = "predict",
                    actual = "actual", cost.fp = 1000, cost.fn = 1200 )
```

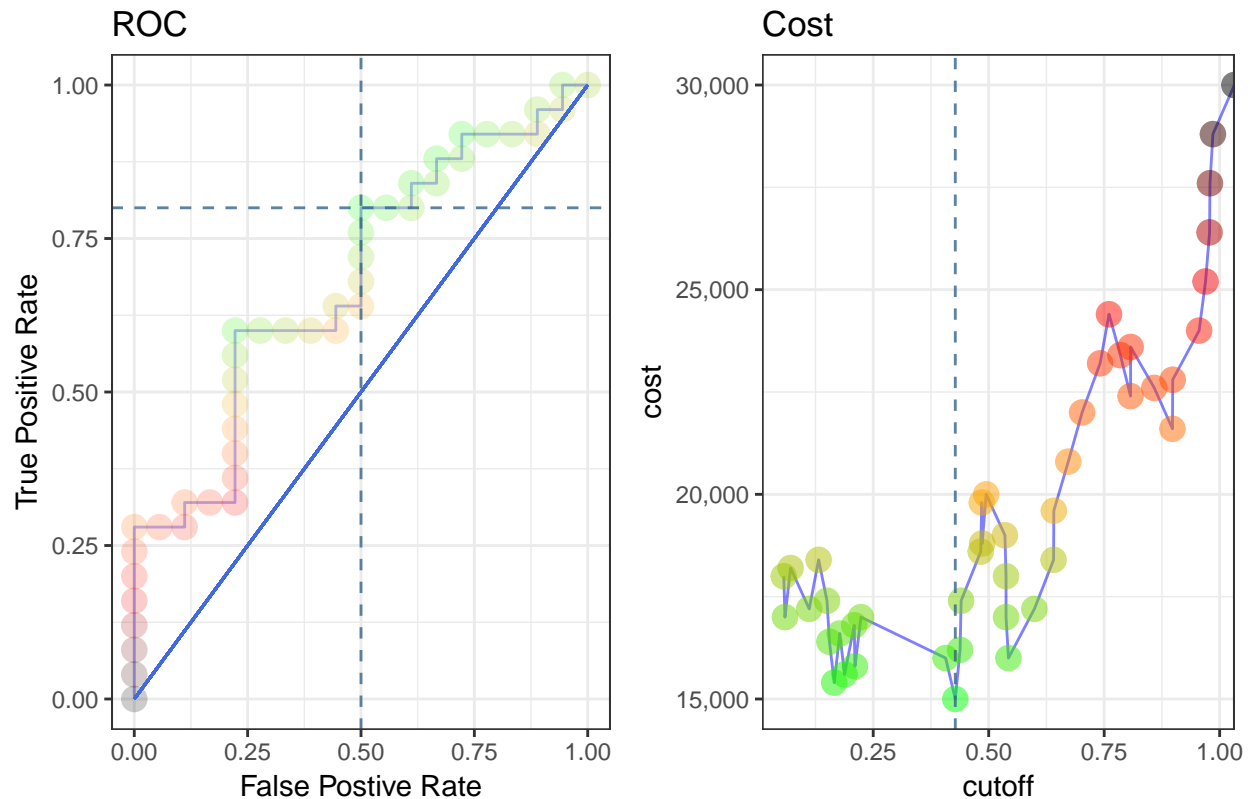
```
#Optimal cutoff for True positive rate
```

```
roc_info$cutoff
```

```
## [1] 0.4276777
```

```
grid.draw(roc_info$plot)
```

**Cutoff at 0.43 – Total Cost = 15000.00, AUC = 0.682**



```
#COBRE model k fold CV check
```

```
set.seed(4321)
```

```
#Optimal cutoff for Accuracy
```

```
result <- cv.error(COBRE_logi_subset, "Subject_Type", cut_off = 0.425)
```

```
Accuracy.k <- result[[1]]
```

```
mean(Accuracy.k)
```

```
## [1] 0.6185714
```

```
TTP <- result[[2]]
```

```
mean(TTP)
```

```
## [1] 0.7285714
```

```
#Optimal cutoff for True positive rate
```

```
result <- cv.error(COBRE_logi_subset, "Subject_Type", cut_off = roc_info$cutoff)
```

```
Accuracy.k <- result[[1]]
```

```
mean(Accuracy.k)
```

```
## [1] 0.6480952
```

```

TTP <- result[[2]]
mean(TTP)

## [1] 0.7285714

#When we want optimize True positive rate, we gave up about 10% of accuracy.

set.seed(4321)
#Fit Data into model build base on other study to test how it handles data from different study

#Fit COBRE data into CNP Model
Fit_COBRE_logi_subset <- COBRE_RF_subset[,c("Subject_Type",names(COBRE_RF_subset[,c(1,137:150)]), subset2)] %>%
  Standarize()

Fit_COBRE_test <- Fit_COBRE_logi_subset

invisible(rm(Fit_COBRE_logi_subset))

Fit_COBRE_test$prediction <- predict(logi_m1, Fit_COBRE_test, type = "response")
Classify(Fit_COBRE_test, Fit_COBRE_test$prediction,"Subject_Type", 0.17 )

##           prediction
##           Control Schizophrenia
## Control           39           33
## Schizophrenia      38           32
## The accuracy is 50 %.
## The True positive rate is 45.714 %

#Fit CNP data into COBRE model
Fit_CNP_logi_subset <- CNP_RF_subset[,c("Subject_Type",names(CNP_RF_subset[,c(1,137:150)]), subset2)] %>%
  Standarize()

Fit_CNP_test <- Fit_CNP_logi_subset

invisible(rm(Fit_CNP_logi_subset))

Fit_CNP_test$prediction <- predict(logi_m2, Fit_CNP_test, type = "response")
Classify(Fit_CNP_test, Fit_CNP_test$prediction,"Subject_Type", cut_off = 0.69 )

##           prediction
##           Control Schizophrenia
## Control           84           31
## Schizophrenia      29           13
## The accuracy is 61.783 %.
## The True positive rate is 30.952 %

#When we introduce data from the other study, the both model has a a low testing accuracy.
#This hint us that the two studys are different.

#Combine data

#Further data cleaning to merge CNP and COBRE data
Study <- rep("CNP",nrow(CNP))

CNP <- data.frame(CNP,Study)

```

```

CNP <- CNP %>%
  select(-c(7:41))

colnames(CNP)[5:6] <- c("Ethnicity", "Education")

levels(CNP$Gender) <- c("Female", "Male")

Study <- rep("COBRE", nrow(COBRE))
COBRE <- data.frame(COBRE, Study)

COBRE <- COBRE %>%
  select(-c(5, 8:111))

# CNP Ethnicity
#1=Hispanic origin
#2=Not of Hispanic origin

#COBRE Ethnicity
#Caucasian = 1
#African-American = 2
#Hispanic = 3

#Recoding required

table(COBRE$Ethnicity)

##
## 1 2 3
## 69 9 53

for(i in 1:length(COBRE$Ethnicity)){
  if(!is.na(COBRE$Ethnicity[i])){
    if(COBRE$Ethnicity[i] == 1 | COBRE$Ethnicity[i] == 2)
      COBRE$Ethnicity[i] <- 4
  }
}
COBRE$Ethnicity <- COBRE$Ethnicity - 2

table(COBRE$Ethnicity)

##
## 1 2
## 53 78

Data <- merge(CNP, COBRE, all = TRUE) %>%
  select(-c(1))
Data$Ethnicity <- as.factor(Data$Ethnicity)
levels(Data$Ethnicity) <- c("Hispanic", "non-Hispanic")

set.seed(4321)

# Combine Data modeling

#Random Forest variable selection

```

```

rfsrc_m3 <- rfsrc(Study~.,data = Data, na.action = c("na.omit"), ntree= 1000)

max_var <- max.subtree(rfsrc_m3, conservative = TRUE)

max_var$topvars

## [1] "Age"
## [2] "Education"
## [3] "Cingulo.opercular.Cerebellar"
## [4] "Fronto.parietal.Dorsal_Attention"
## [5] "Subcortical.Cerebellar"
## [6] "Visual.Fronto.parietal"
## [7] "Uncertain.char_path_length"
## [8] "Cingulo.opercular_Task_Control.mod"
## [9] "Uncertain.mod"
## [10] "Subcortical.global_eff"
## [11] "Uncertain.global_eff"
## [12] "Subcortical.clust_coef"

#delete duplicate entity

subset3 <- as.vector(max_var$topvars)

subset3 <- delete_dup(subset3,Data[,c(1:5,139:152)])

#Logistic Regression model

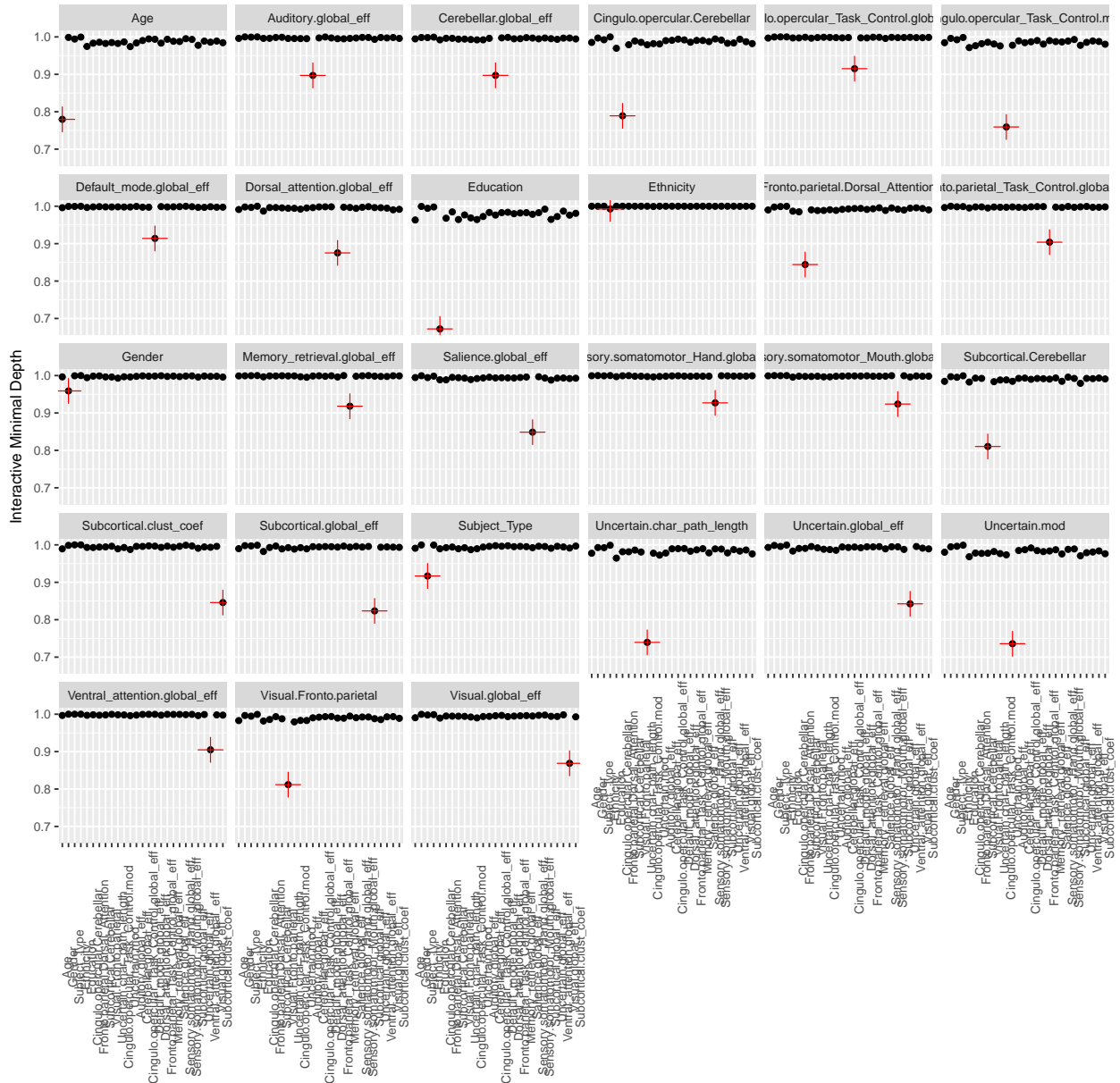
Data_logi <- Data[,c("Study",names(Data[,c(1:5,139:152)]), subset3)]

Data_logi <- na.omit(Data_logi) %>%
  Standarize()

#find interaction
gg_int <- gg_interaction(find.interaction(rfsrc_m3,
                                         xvar.names = names(Data_logi[, -c(1)]),
                                         sorted = FALSE,
                                         verbose = FALSE))

plot(gg_int)

```



*#No interaction in fund base on the result, we don't have to add interaction term*

*#check correlation*

```
high_cor <- findCorrelation(cor(Data_logi[, -c(1,3:5)]), cutoff = 0.75) + 4
```

*#Remove variables to prevent multicollinearity problem*

```
Data_logi <- Data_logi %>%
  select(-c(high_cor))
```

```
index <- sample(1:nrow(Data_logi), size = round(nrow(Data_logi)*0.7,0), replace = FALSE)
```

```
Data_train <- Data_logi[index,]
```

```
Data_test <- Data_logi[-index,]
```



```
logi_m3 <-glm(Study~. + Subject_Type*Age , data = Data_train, family = "binomial")
summary(logi_m3)
```

```
##
## Call:
## glm(formula = Study ~ . + Subject_Type * Age, family = "binomial",
##      data = Data_train)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7700  -0.6746  -0.3092   0.7643   2.4375
##
## Coefficients:
##                                Estimate Std. Error z value
## (Intercept)                   -0.53169    0.42039  -1.265
## Age                           0.36891    0.31853   1.158
## GenderMale                     0.46669    0.43638   1.069
## Subject_TypeSchizophrenia      0.38913    0.47163   0.825
## Ethnicitynon-Hispanic        -0.21121    0.44299  -0.477
## Education                     -0.48969    0.24136  -2.029
## Auditory.global_eff           0.22337    0.23506   0.950
## Cerebellar.global_eff        -0.29040    0.23386  -1.242
## Cingulo.opercular_Task_Control.global_eff 0.33690    0.26957   1.250
## Default_mode.global_eff      -0.09649    0.28185  -0.342
## Dorsal_attention.global_eff   0.25471    0.24005   1.061
## Fronto.parietal_Task_Control.global_eff  0.01053    0.23382   0.045
## Memory_retrieval.global_eff  -0.11565    0.21575  -0.536
## Salience.global_eff         0.33326    0.24352   1.368
## Sensory.somatomotor_Hand.global_eff      0.03733    0.30262   0.123
## Sensory.somatomotor_Mouth.global_eff    -0.12186    0.25212  -0.483
## Subcortical.global_eff       0.35924    0.26237   1.369
## Uncertain.global_eff        -0.33765    0.24043  -1.404
## Ventral_attention.global_eff -0.11511    0.22425  -0.513
## Visual.global_eff            -0.17071    0.26198  -0.652
## Cingulo.opercular.Cerebellar -0.59461    0.28980  -2.052
## Subcortical.Cerebellar       -0.31993    0.31572  -1.013
## Visual.Fronto.parietal       0.15127    0.31430   0.481
## Cingulo.opercular_Task_Control.mod      0.40354    0.23812   1.695
## Uncertain.mod                0.39313    0.24439   1.609
## Subcortical.clust_coef       0.50978    0.23507   2.169
## Age:Subject_TypeSchizophrenia -0.09416    0.42299  -0.223
##                                Pr(>|z|)
## (Intercept)                   0.2060
## Age                           0.2468
## GenderMale                     0.2849
## Subject_TypeSchizophrenia      0.4093
## Ethnicitynon-Hispanic         0.6335
## Education                     0.0425 *
## Auditory.global_eff           0.3420
## Cerebellar.global_eff         0.2143
## Cingulo.opercular_Task_Control.global_eff 0.2114
## Default_mode.global_eff       0.7321
## Dorsal_attention.global_eff   0.2887
## Fronto.parietal_Task_Control.global_eff  0.9641
```

```

## Memory_retrieval.global_eff      0.5919
## Salience.global_eff             0.1712
## Sensory.somatomotor_Hand.global_eff 0.9018
## Sensory.somatomotor_Mouth.global_eff 0.6289
## Subcortical.global_eff           0.1709
## Uncertain.global_eff             0.1602
## Ventral_attention.global_eff      0.6077
## Visual.global_eff                 0.5146
## Cingulo.opercular.Cerebellar      0.0402 *
## Subcortical.Cerebellar            0.3109
## Visual.Fronto.parietal            0.6303
## Cingulo.opercular_Task_Control.mod 0.0901 .
## Uncertain.mod                    0.1077
## Subcortical.clust_coef            0.0301 *
## Age:Subject_TypeSchizophrenia     0.8239
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 265.96  on 193  degrees of freedom
## Residual deviance: 178.00  on 167  degrees of freedom
## AIC: 232
##
## Number of Fisher Scoring iterations: 5

```

```
round(exp(coef(logi_m3)),3)
```

```

##                               (Intercept)
##                               0.588
##                               Age
##                               1.446
##                               GenderMale
##                               1.595
##      Subject_TypeSchizophrenia
##                               1.476
##      Ethnicitynon-Hispanic
##                               0.810
##      Education
##                               0.613
##      Auditory.global_eff
##                               1.250
##      Cerebellar.global_eff
##                               0.748
## Cingulo.opercular_Task_Control.global_eff
##                               1.401
##      Default_mode.global_eff
##                               0.908
##      Dorsal_attention.global_eff
##                               1.290
##      Fronto.parietal_Task_Control.global_eff
##                               1.011
##      Memory_retrieval.global_eff
##                               0.891
##      Salience.global_eff

```

```
## 1.396
## Sensory.somatomotor_Hand.global_eff
## 1.038
## Sensory.somatomotor_Mouth.global_eff
## 0.885
## Subcortical.global_eff
## 1.432
## Uncertain.global_eff
## 0.713
## Ventral_attention.global_eff
## 0.891
## Visual.global_eff
## 0.843
## Cingulo.opercular.Cerebellar
## 0.552
## Subcortical.Cerebellar
## 0.726
## Visual.Fronto.parietal
## 1.163
## Cingulo.opercular_Task_Control.mod
## 1.497
## Uncertain.mod
## 1.482
## Subcortical.clust_coef
## 1.665
## Age:Subject_TypeSchizophrenia
## 0.910
```

```
anova(logi_m3, test = "Chisq")
```

```
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: Study
##
## Terms added sequentially (first to last)
##
```

	Df	Deviance	Resid. Df	Resid. Dev
## NULL			193	265.96
## Age	1	3.3371	192	262.63
## Gender	1	6.2191	191	256.41
## Subject_Type	1	7.2228	190	249.19
## Ethnicity	1	0.5869	189	248.60
## Education	1	12.7223	188	235.88
## Auditory.global_eff	1	3.6053	187	232.27
## Cerebellar.global_eff	1	6.4624	186	225.81
## Cingulo.opercular_Task_Control.global_eff	1	0.2843	185	225.52
## Default_mode.global_eff	1	3.6685	184	221.86
## Dorsal_attention.global_eff	1	1.5829	183	220.27
## Fronto.parietal_Task_Control.global_eff	1	0.0726	182	220.20
## Memory_retrieval.global_eff	1	0.0668	181	220.13
## Salience.global_eff	1	3.7711	180	216.36
## Sensory.somatomotor_Hand.global_eff	1	0.0170	179	216.34

```
## Sensory.somatomotor_Mouth.global_eff      1  0.4466      178      215.90
## Subcortical.global_eff                    1  3.5802      177      212.32
## Uncertain.global_eff                     1  5.7028      176      206.62
## Ventral_attention.global_eff              1  0.4207      175      206.19
## Visual.global_eff                        1  2.0756      174      204.12
## Cingulo.opercular.Cerebellar              1 14.1765      173      189.94
## Subcortical.Cerebellar                   1  1.8643      172      188.08
## Visual.Fronto.parietal                   1  0.0379      171      188.04
## Cingulo.opercular_Task_Control.mod        1  2.3040      170      185.74
## Uncertain.mod                           1  2.3636      169      183.37
## Subcortical.clust_coef                   1  5.3243      168      178.05
## Age:Subject_Type                         1  0.0495      167      178.00
##                                           Pr(>Chi)
## NULL
## Age                                     0.0677343 .
## Gender                                0.0126379 *
## Subject_Type                          0.0071985 **
## Ethnicity                             0.4436360
## Education                             0.0003613 ***
## Auditory.global_eff                   0.0575963 .
## Cerebellar.global_eff                 0.0110179 *
## Cingulo.opercular_Task_Control.global_eff 0.5938826
## Default_mode.global_eff               0.0554491 .
## Dorsal_attention.global_eff            0.2083497
## Fronto.parietal_Task_Control.global_eff 0.7875658
## Memory_retrieval.global_eff            0.7959919
## Salience.global_eff                   0.0521457 .
## Sensory.somatomotor_Hand.global_eff     0.8961752
## Sensory.somatomotor_Mouth.global_eff    0.5039533
## Subcortical.global_eff                 0.0584732 .
## Uncertain.global_eff                   0.0169375 *
## Ventral_attention.global_eff            0.5165702
## Visual.global_eff                      0.1496697
## Cingulo.opercular.Cerebellar            0.0001664 ***
## Subcortical.Cerebellar                 0.1721237
## Visual.Fronto.parietal                 0.8457015
## Cingulo.opercular_Task_Control.mod      0.1290434
## Uncertain.mod                          0.1241928
## Subcortical.clust_coef                  0.0210304 *
## Age:Subject_Type                       0.8239025
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

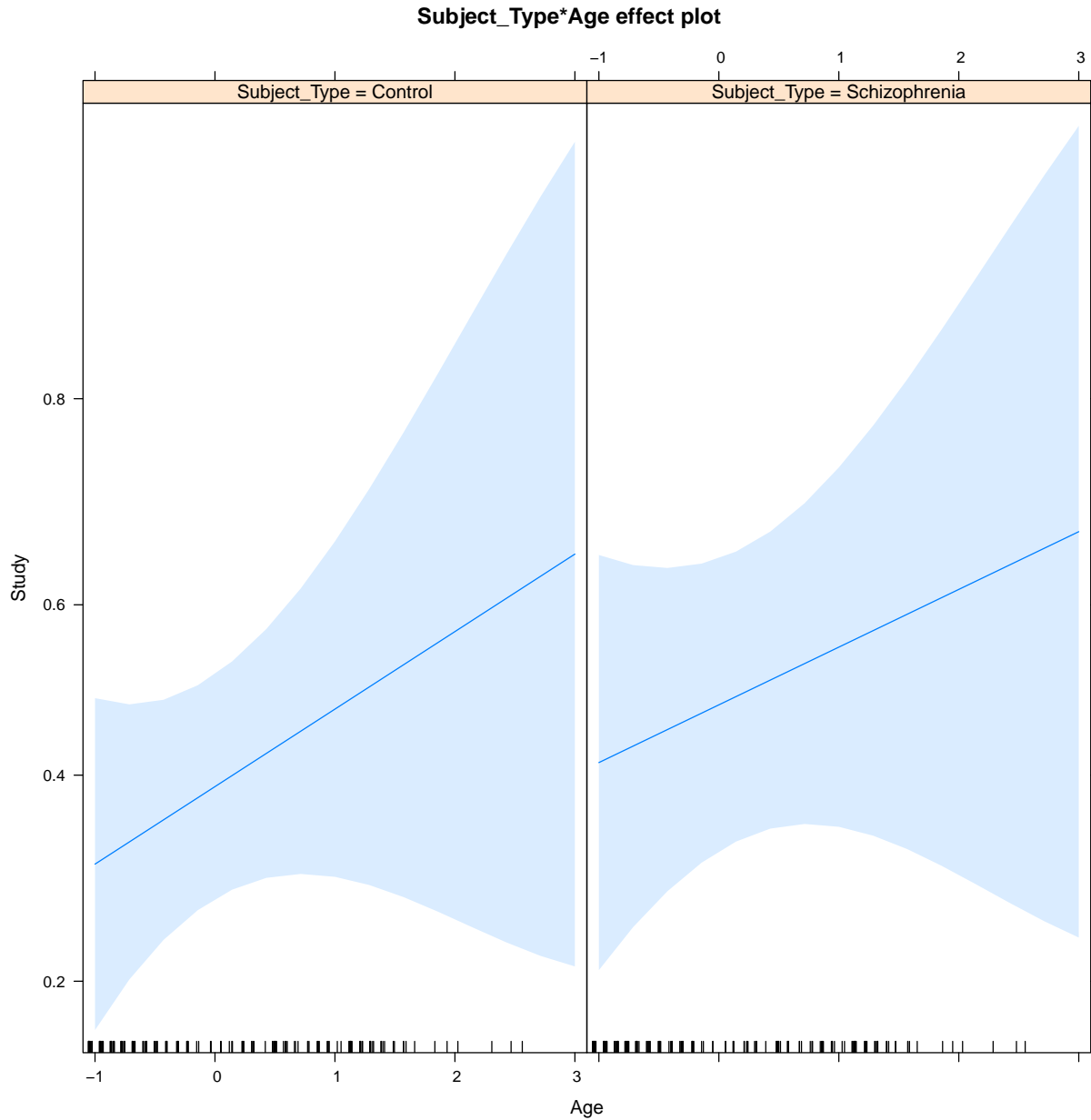
```
#R-squared
```

```
R_squared <- 1 - (summary(logi_m3)[[4]]/summary(logi_m3)[[8]])
R_squared
```

```
## [1] 0.3307403
```

```
#Effect plot
```

```
plot(Effect(c("Subject_Type", "Age"), logi_m3),ask = FALSE)
```



```
#70/30 CV check
```

```
#Train
```

```
Data_train$prediction <- predict(logi_m3, Data_train, type = "response")
```

```
#Test
```

```
Data_test$prediction <- predict(logi_m3, Data_test, type = "response")
```

```
prop.table(table(Data$Study))
```

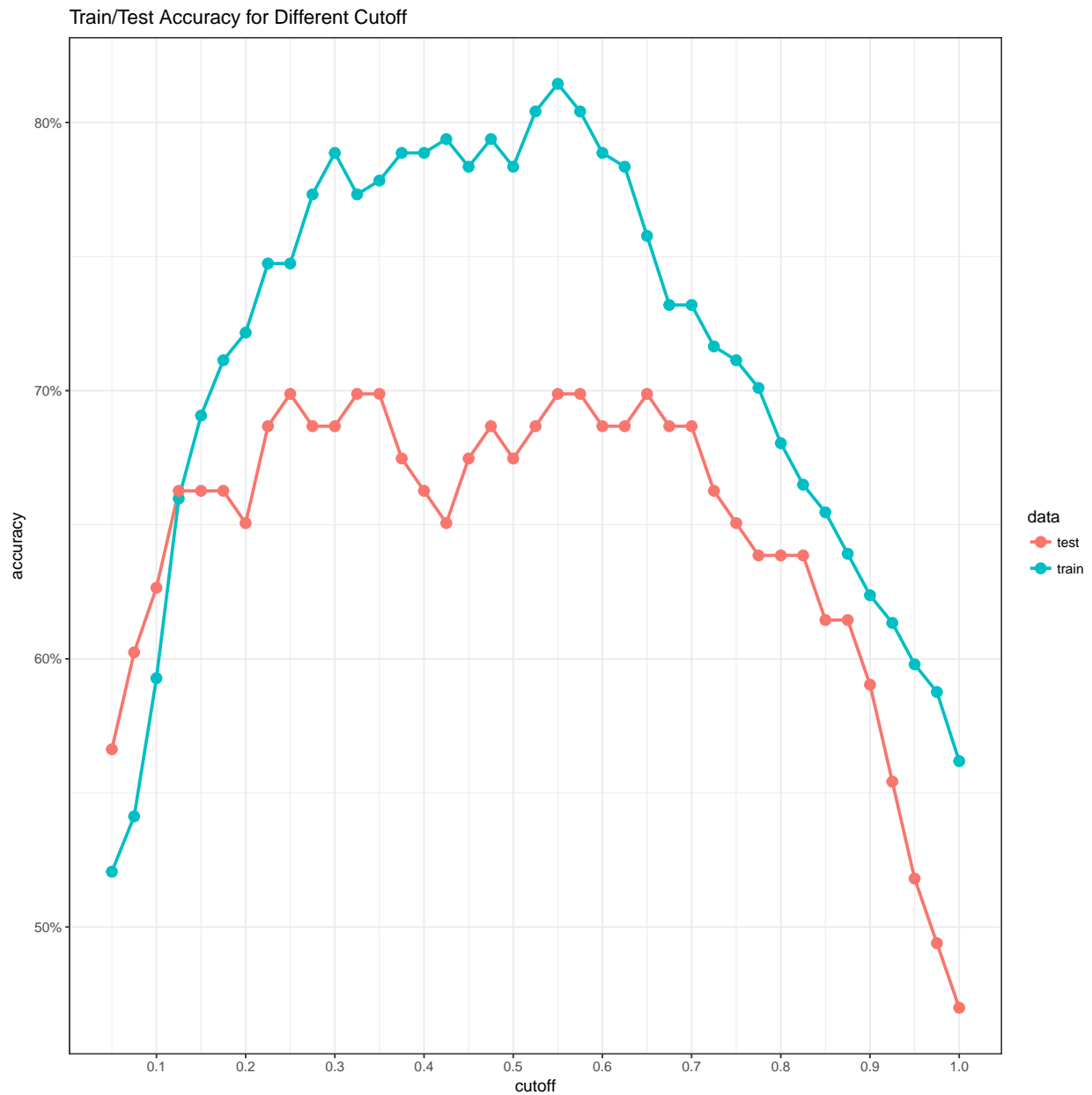
```
##
```

```
##      CNP      COBRE
```

```
## 0.5250836 0.4749164
```

```
accuracy_info <- AccuracyCutoffInfo( train = Data_train, test = Data_test,
                                     predict = "prediction", actual = "Study" )
```

```
accuracy_info$plot
```



```
Classify(Data_train, Data_train$prediction, "Study", 0.55)
```

```
##      prediction
##      CNP COBRE
## CNP      95   14
## COBRE    22   63
## The accuracy is 81.443 %.
## The True positive rate is 74.118 %
```

```
Classify(Data_test, Data_test$prediction, "Study", 0.55)
```

```
##      prediction
##      CNP COBRE
## CNP      30      9
## COBRE     16     28
## The accuracy is 69.88 %.
## The True positive rate is 63.636 %
```

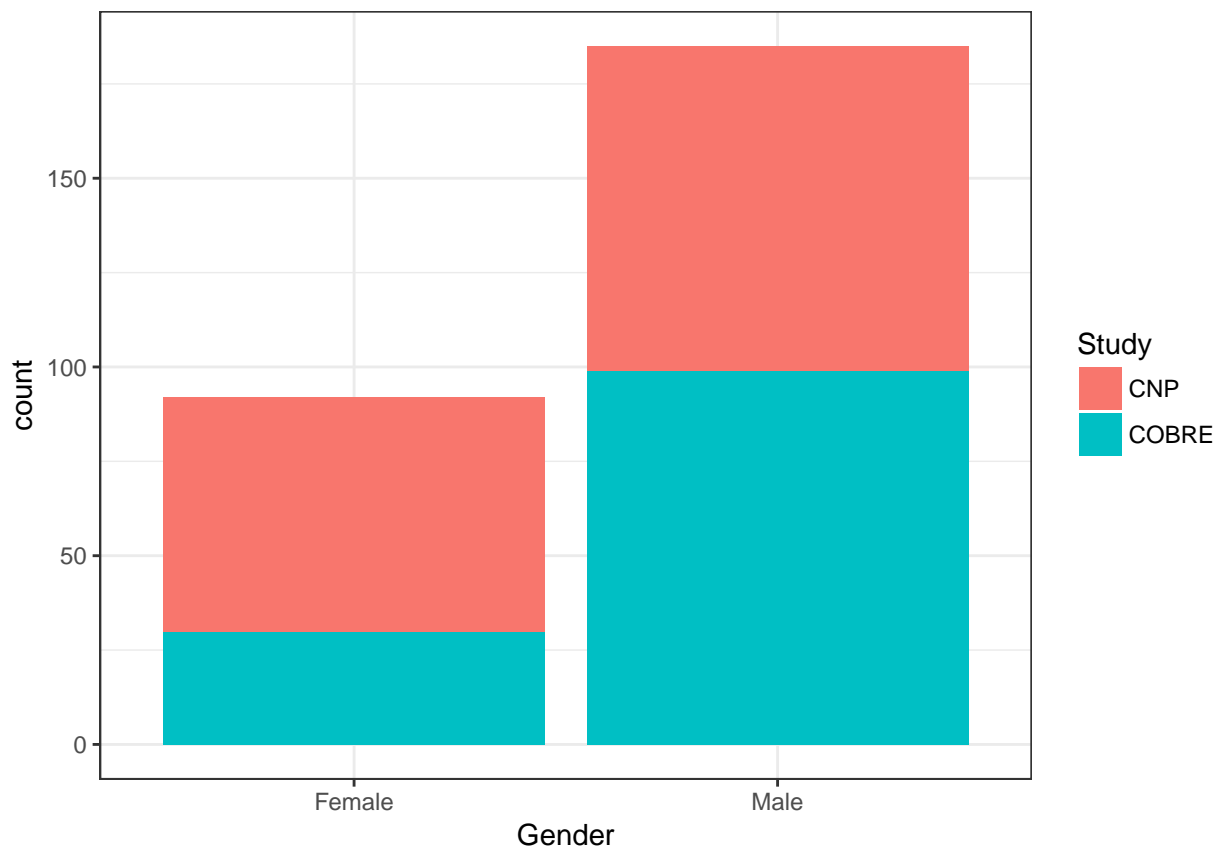
```
#Combine data model k fold CV check
#Here we are not interesting in looking at the True positive rate
set.seed(4321)
```

```
Accuracy.k <- cv.error(Data_logi, "Study", cut_off = 0.55)[[1]]
mean(Accuracy.k)
```

```
## [1] 0.7439153
```

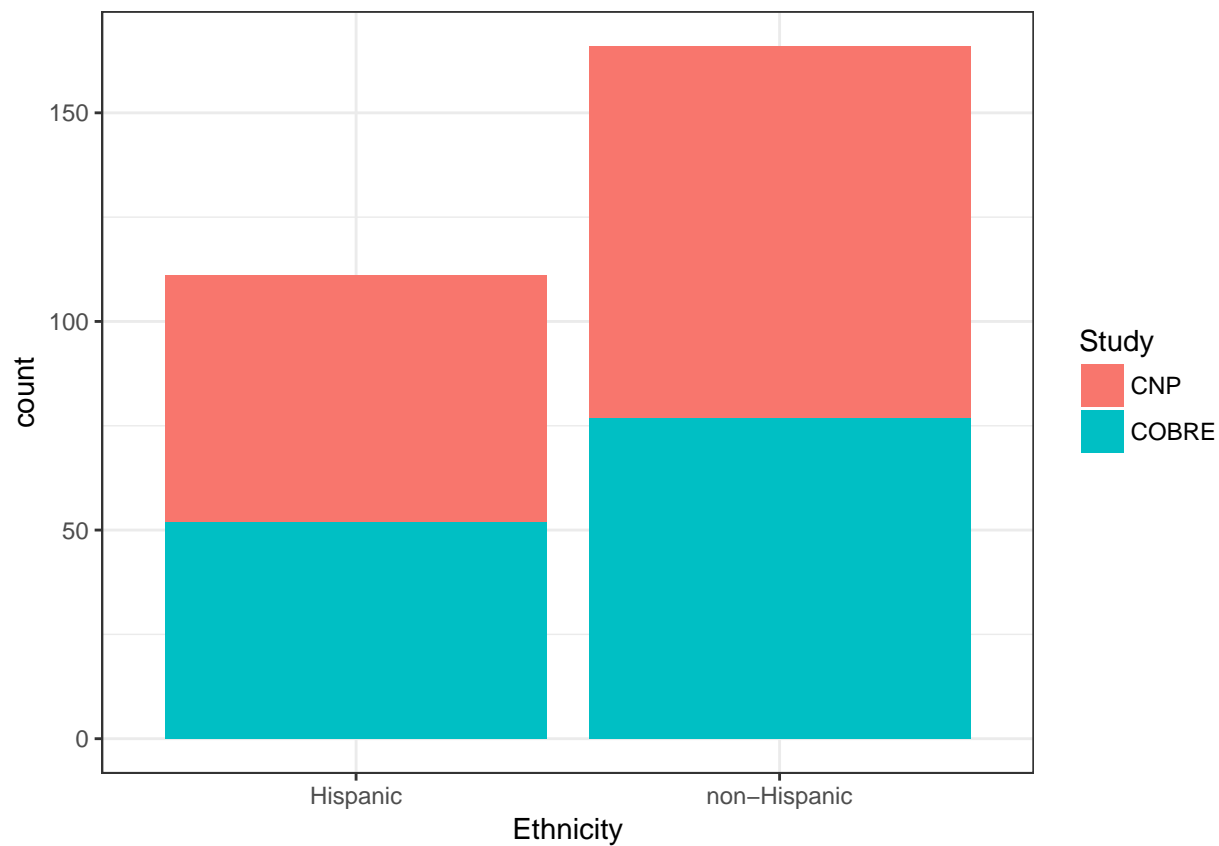
```
par(mfrow = c(2,2))
```

```
ggplot(data = na.omit(Data), aes(x = Gender, fill = Study)) +
  geom_bar() +
  theme_bw()
```



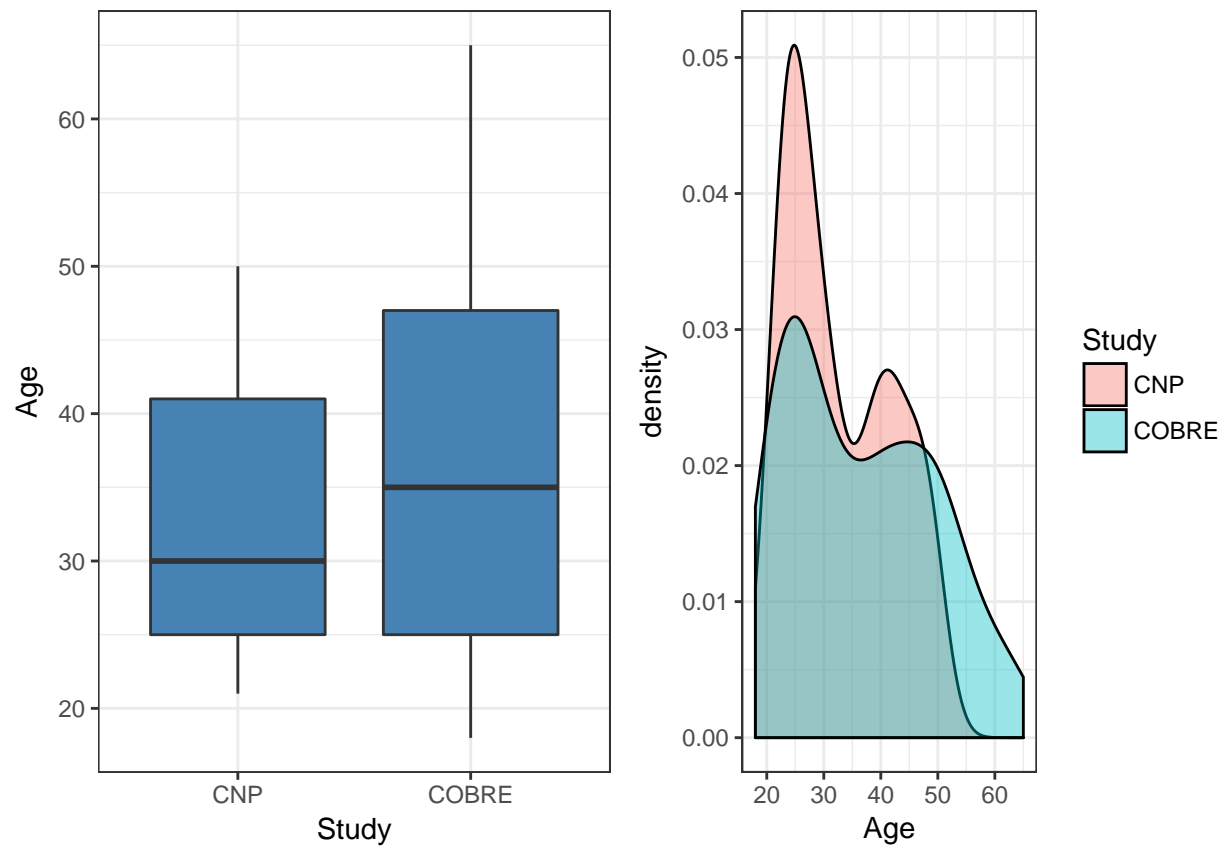
```
ggplot(data = na.omit(Data), aes(x = Ethnicity, fill = Study)) +
  geom_bar() +
```

```
theme_bw()
```



```
plot1 <- ggplot(data = na.omit(Data), aes(x = Study, y = Age)) +  
  geom_boxplot(fill = "steelblue") +  
  theme_bw()  
  
plot2 <- ggplot(data = na.omit(Data), aes(x = Age, fill = Study)) +  
  geom_density(alpha = 0.4) +  
  theme_bw()  
  
grid.arrange(plot1, plot2, nrow = 1, ncol = 2)
```

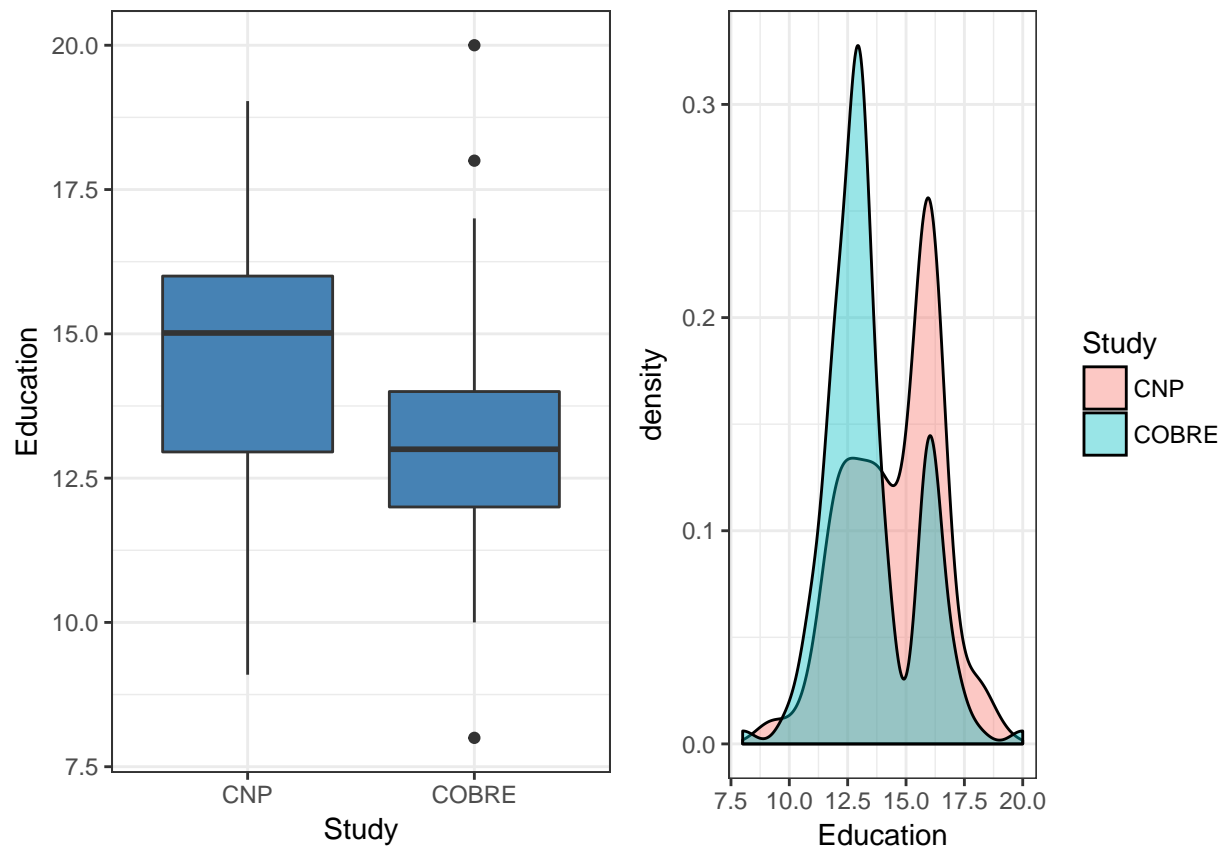




```
plot3 <- ggplot(data = na.omit(Data), aes(x = Study, y = Education)) +
  geom_boxplot(fill = "steelblue") +
  theme_bw()

plot4 <- ggplot(data = na.omit(Data), aes(x = Education, fill = Study)) +
  geom_density(alpha = 0.4) +
  theme_bw()

grid.arrange(plot3, plot4, nrow = 1, ncol = 2)
```



*#Recall the anova output for the combined data set logistic model*

```
anova(logi_m3, test = "Chisq")
```

```
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: Study
##
## Terms added sequentially (first to last)
##
##
```

	Df	Deviance	Resid. Df	Resid. Dev
## NULL			193	265.96
## Age	1	3.3371	192	262.63
## Gender	1	6.2191	191	256.41
## Subject_Type	1	7.2228	190	249.19
## Ethnicity	1	0.5869	189	248.60
## Education	1	12.7223	188	235.88
## Auditory.global_eff	1	3.6053	187	232.27
## Cerebellar.global_eff	1	6.4624	186	225.81
## Cingulo.opercular_Task_Control.global_eff	1	0.2843	185	225.52
## Default_mode.global_eff	1	3.6685	184	221.86
## Dorsal_attention.global_eff	1	1.5829	183	220.27
## Fronto.parietal_Task_Control.global_eff	1	0.0726	182	220.20

```
## Memory_retrieval.global_eff      1  0.0668      181    220.13
## Salience.global_eff              1  3.7711      180    216.36
## Sensory.somatomotor_Hand.global_eff 1  0.0170      179    216.34
## Sensory.somatomotor_Mouth.global_eff 1  0.4466      178    215.90
## Subcortical.global_eff           1  3.5802      177    212.32
## Uncertain.global_eff             1  5.7028      176    206.62
## Ventral_attention.global_eff      1  0.4207      175    206.19
## Visual.global_eff                1  2.0756      174    204.12
## Cingulo.opercular.Cerebellar      1 14.1765      173    189.94
## Subcortical.Cerebellar            1  1.8643      172    188.08
## Visual.Fronto.parietal            1  0.0379      171    188.04
## Cingulo.opercular_Task_Control.mod 1  2.3040      170    185.74
## Uncertain.mod                    1  2.3636      169    183.37
## Subcortical.clust_coef            1  5.3243      168    178.05
## Age:Subject_Type                 1  0.0495      167    178.00
##                                Pr(>Chi)
## NULL
## Age                             0.0677343 .
## Gender                          0.0126379 *
## Subject_Type                    0.0071985 **
## Ethnicity                       0.4436360
## Education                       0.0003613 ***
## Auditory.global_eff             0.0575963 .
## Cerebellar.global_eff           0.0110179 *
## Cingulo.opercular_Task_Control.global_eff 0.5938826
## Default_mode.global_eff         0.0554491 .
## Dorsal_attention.global_eff      0.2083497
## Fronto.parietal_Task_Control.global_eff 0.7875658
## Memory_retrieval.global_eff      0.7959919
## Salience.global_eff            0.0521457 .
## Sensory.somatomotor_Hand.global_eff 0.8961752
## Sensory.somatomotor_Mouth.global_eff 0.5039533
## Subcortical.global_eff          0.0584732 .
## Uncertain.global_eff            0.0169375 *
## Ventral_attention.global_eff     0.5165702
## Visual.global_eff               0.1496697
## Cingulo.opercular.Cerebellar     0.0001664 ***
## Subcortical.Cerebellar           0.1721237
## Visual.Fronto.parietal           0.8457015
## Cingulo.opercular_Task_Control.mod 0.1290434
## Uncertain.mod                   0.1241928
## Subcortical.clust_coef           0.0210304 *
## Age:Subject_Type                0.8239025
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

*#Hypothesis testing for demographic variables in the combined data set*

```
t.test(Age~Study, data = Data_logi)
```

```
##
## Welch Two Sample t-test
##
## data: Age by Study
```

```
## t = -2.8159, df = 227.22, p-value = 0.005292
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.5820745 -0.1028142
## sample estimates:
## mean in group CNP mean in group COBRE
## -0.1594777 0.1829667
```

```
t.test(Education~Study, data = Data_logi)
```

```
##
## Welch Two Sample t-test
##
## data: Education by Study
## t = 4.5959, df = 272.59, p-value = 6.597e-06
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.3046507 0.7612365
## sample estimates:
## mean in group CNP mean in group COBRE
## 0.2481939 -0.2847496
```

```
#Pearson's chi-squared test
```

```
#H_{0} = there is no difference between the distributions
#H_{1} = there is a difference between the distributions
```

```
chisq.test(table(Data_logi$Study, Data_logi$Gender))
```

```
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(Data_logi$Study, Data_logi$Gender)
## X-squared = 9.9677, df = 1, p-value = 0.001593
```

```
chisq.test(table(Data_logi$Study, Data_logi$Ethnicity))
```

```
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(Data_logi$Study, Data_logi$Ethnicity)
## X-squared = 1.2223e-30, df = 1, p-value = 1
```

```
chisq.test(table(Data_logi$Study, Data_logi$Subject_Type))
```

```
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: table(Data_logi$Study, Data_logi$Subject_Type)
## X-squared = 16.988, df = 1, p-value = 3.762e-05
```