

# 573 Final Paper RMD HT

2022-11-08

```
knitr::opts_chunk$set(echo = TRUE)
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.2 --
## v ggplot2 3.3.6      v purrr  0.3.4
## v tibble  3.1.8      v dplyr  1.0.9
## v tidyr   1.2.0      v stringr 1.4.0
## v readr   2.1.2      v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

```
library(dplyr)
library(gbm)
```

```
## Loaded gbm 2.1.8.1
```

```
library(ggplot2)
library(caret)
```

```
## Loading required package: lattice
##
## Attaching package: 'caret'
##
## The following object is masked from 'package:purrr':
##
##     lift
```

```
library(glmnet)
```

```
## Loading required package: Matrix
##
## Attaching package: 'Matrix'
##
## The following objects are masked from 'package:tidyr':
##
##     expand, pack, unpack
##
## Loaded glmnet 4.1-4
```

```
library(pROC)
```

```
## Type 'citation("pROC")' for a citation.
##
## Attaching package: 'pROC'
##
## The following objects are masked from 'package:stats':
##
##     cov, smooth, var
```

```
library(class)
library(MASS)
```

```
##
## Attaching package: 'MASS'
##
## The following object is masked from 'package:dplyr':
##
##     select
```

```
library(randomForest)
```

```
## randomForest 4.7-1.1
## Type rfNews() to see new features/changes/bug fixes.
##
## Attaching package: 'randomForest'
##
## The following object is masked from 'package:dplyr':
##
##     combine
##
## The following object is masked from 'package:ggplot2':
##
##     margin
```

```
library(tree)
library(e1071)
```

Read and clean data

```
set.seed(12345)
data <- read.csv("Drug_Consumption.csv")
head(data)
```

|      | ID | Age   | Gender | Education                         | Country | Ethnicity | Nscore   |
|------|----|-------|--------|-----------------------------------|---------|-----------|----------|
| ## 1 | 2  | 25-34 | M      | Doctorate degree                  | UK      | White     | -0.67825 |
| ## 2 | 3  | 35-44 | M      | Professional certificate/ diploma | UK      | White     | -0.46725 |
| ## 3 | 4  | 18-24 | F      | Masters degree                    | UK      | White     | -0.14882 |
| ## 4 | 5  | 35-44 | F      | Doctorate degree                  | UK      | White     | 0.73545  |
| ## 5 | 6  | 65    | F      | Left school at 18 years           | Canada  | White     | -0.67825 |

```
## 6 7 45-54 M Masters degree USA White -0.46725
## EScore OScore AScore Cscore Impulsive SS Alcohol Amphet Amyl
## 1 1.93886 1.43533 0.76096 -0.14277 -0.71126 -0.21575 CL5 CL2 CL2
## 2 0.80523 -0.84732 -1.62090 -1.01450 -1.37983 0.40148 CL6 CL0 CL0
## 3 -0.80615 -0.01928 0.59042 0.58489 -1.37983 -1.18084 CL4 CL0 CL0
## 4 -1.63340 -0.45174 -0.30172 1.30612 -0.21712 -0.21575 CL4 CL1 CL1
## 5 -0.30033 -1.55521 2.03972 1.63088 -1.37983 -1.54858 CL2 CL0 CL0
## 6 -1.09207 -0.45174 -0.30172 0.93949 -0.21712 0.07987 CL6 CL0 CL0
## Benzos Caff Cannabis Choc Coke Crack Ecstasy Heroin Ketamine Legalh LSD Meth
## 1 CL0 CL6 CL4 CL6 CL3 CL0 CL4 CL0 CL2 CL0 CL2 CL3
## 2 CL0 CL6 CL3 CL4 CL0 CL0 CL0 CL0 CL0 CL0 CL0 CL0
## 3 CL3 CL5 CL2 CL4 CL2 CL0 CL0 CL0 CL2 CL0 CL0 CL0
## 4 CL0 CL6 CL3 CL6 CL0 CL0 CL1 CL0 CL0 CL1 CL0 CL0
## 5 CL0 CL6 CL0 CL4 CL0 CL0 CL0 CL0 CL0 CL0 CL0 CL0
## 6 CL0 CL6 CL1 CL5 CL0 CL0 CL0 CL0 CL0 CL0 CL0 CL0
## Mushrooms Nicotine Semer VSA
## 1 CL0 CL4 CL0 CL0
## 2 CL1 CL0 CL0 CL0
## 3 CL0 CL2 CL0 CL0
## 4 CL2 CL2 CL0 CL0
## 5 CL0 CL6 CL0 CL0
## 6 CL0 CL6 CL0 CL0
```

```
table(data$Age)
```

```
##
## 18-24 25-34 35-44 45-54 55-64 65
## 643 481 355 294 93 18
```

```
table(data$Gender)
```

```
##
## F M
## 941 943
```

```
table(data$Education)
```

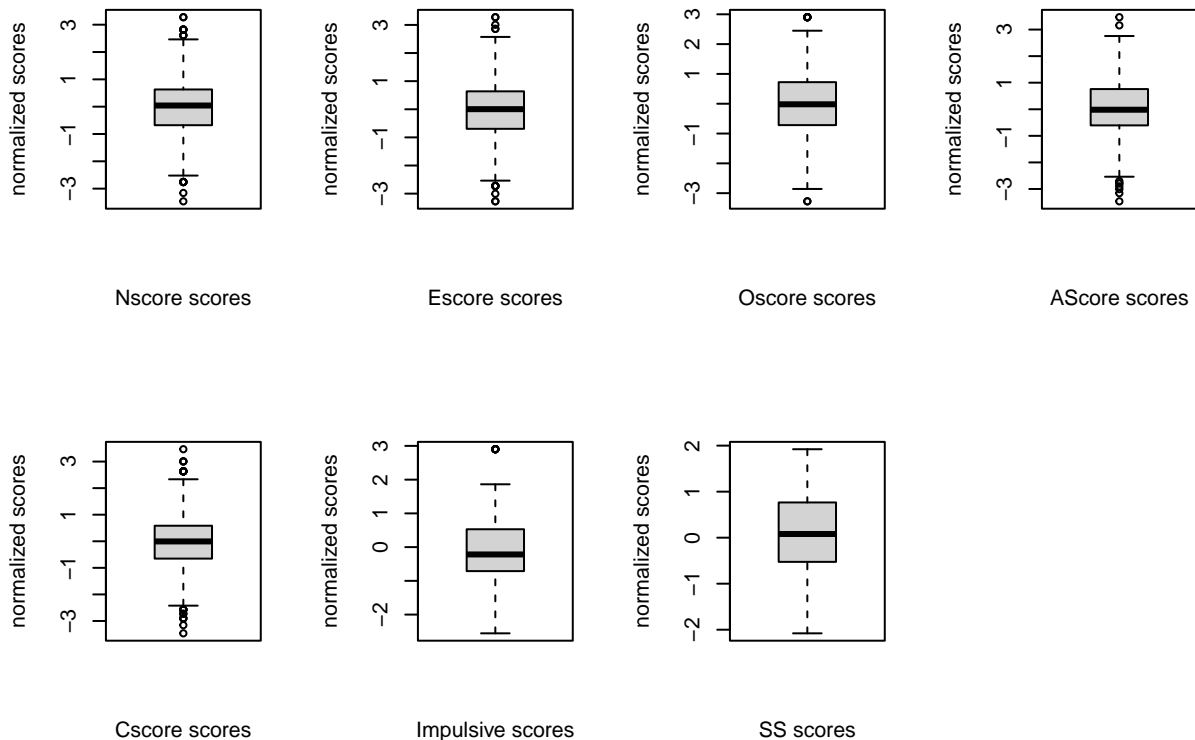
```
##
## Doctorate degree
## 89
## Left school at 16 years
## 99
## Left school at 17 years
## 30
## Left school at 18 years
## 100
## Left school before 16 years
## 28
## Masters degree
## 283
## Professional certificate/ diploma
## 269
```

```
## Some college or university, no certificate or degree
##                                     506
##                                     University degree
##                                     480
```

```
table(data$Country)
```

```
##
##      Australia      Canada      New Zealand      Other
##           54           87           5           118
## Republic of Ireland      UK      USA
##           20           1043          557
```

```
par(mfrow = c(2,4))
for(i in 7:13){
  boxplot(data[,i], xlab = paste(colnames(data)[i], "scores"), ylab = "normalized scores")
}
```



Read and clean data

```
set.seed(12345)
data <- read.csv("Drug_Consumption.csv")

# Remove the over-claimers using the control drug "Semer"
data <- subset(data, data$Semer == "CL0")
```

```

for(i in 14:ncol(data)){
  data[,i] <- as.numeric(data[, i] == "CL4" | data[, i] == "CL5" | data[, i] == "CL6")
}

# Drop 65+
data <- data %>% mutate(dummy=1) %>%
spread(key=Age,value=dummy,fill=0)

# Drop Doctorate
data <- data %>% mutate(dummy=1) %>%
spread(key=Education,value=dummy,fill=0)

# Drop other
data <- data %>% mutate(dummy=1) %>%
spread(key=Country,value=dummy,fill=0)

# Drop other
data <- data %>% mutate(dummy=1) %>%
spread(key=Ethnicity,value=dummy,fill=0)

# Drop 'F' variable and rename to gender
data <- data %>% mutate(dummy=1) %>%
spread(key=Gender,value=dummy, fill=0)

# Drop variables that we aren't using.
drop <- c("ID", "65+", "Doctorate degree", "Other", "F", "Amphet", "Amyl", "Benzos", "Choc", "Crack", "Coke", "E")
data <- data[,!(names(data) %in% drop)]

names(data)[names(data) == "M"] <- "Gender"

# Split into test and train data
test.i <- sample(1:nrow(data), .3*nrow(data))
test.data <- data[test.i,]
train.data <- data[-test.i,]

```

Generate Tables for Data

```
head(data)
```

| ##   | Nscore   | Escore   | Oscore   | AScore   | Cscore   | Impulsive | SS       | Alcohol | Caff                    |
|------|----------|----------|----------|----------|----------|-----------|----------|---------|-------------------------|
| ## 1 | -0.67825 | 1.93886  | 1.43533  | 0.76096  | -0.14277 | -0.71126  | -0.21575 | 1       | 1                       |
| ## 2 | -0.46725 | 0.80523  | -0.84732 | -1.62090 | -1.01450 | -1.37983  | 0.40148  | 1       | 1                       |
| ## 3 | -0.14882 | -0.80615 | -0.01928 | 0.59042  | 0.58489  | -1.37983  | -1.18084 | 1       | 1                       |
| ## 4 | 0.73545  | -1.63340 | -0.45174 | -0.30172 | 1.30612  | -0.21712  | -0.21575 | 1       | 1                       |
| ## 5 | -0.67825 | -0.30033 | -1.55521 | 2.03972  | 1.63088  | -1.37983  | -1.54858 | 0       | 1                       |
| ## 6 | -0.46725 | -1.09207 | -0.45174 | -0.30172 | 0.93949  | -0.21712  | 0.07987  | 1       | 1                       |
| ##   | Cannabis | Nicotine | 18-24    | 25-34    | 35-44    | 45-54     | 55-64    | 65      | Left school at 16 years |
| ## 1 | 1        | 1        | 0        | 1        | 0        | 0         | 0        | 0       |                         |
| ## 2 | 0        | 0        | 0        | 0        | 1        | 0         | 0        | 0       |                         |
| ## 3 | 0        | 0        | 1        | 0        | 0        | 0         | 0        | 0       |                         |
| ## 4 | 0        | 0        | 0        | 0        | 1        | 0         | 0        | 0       |                         |
| ## 5 | 0        | 1        | 0        | 0        | 0        | 0         | 0        | 1       |                         |
| ## 6 | 0        | 1        | 0        | 0        | 0        | 1         | 0        | 0       |                         |

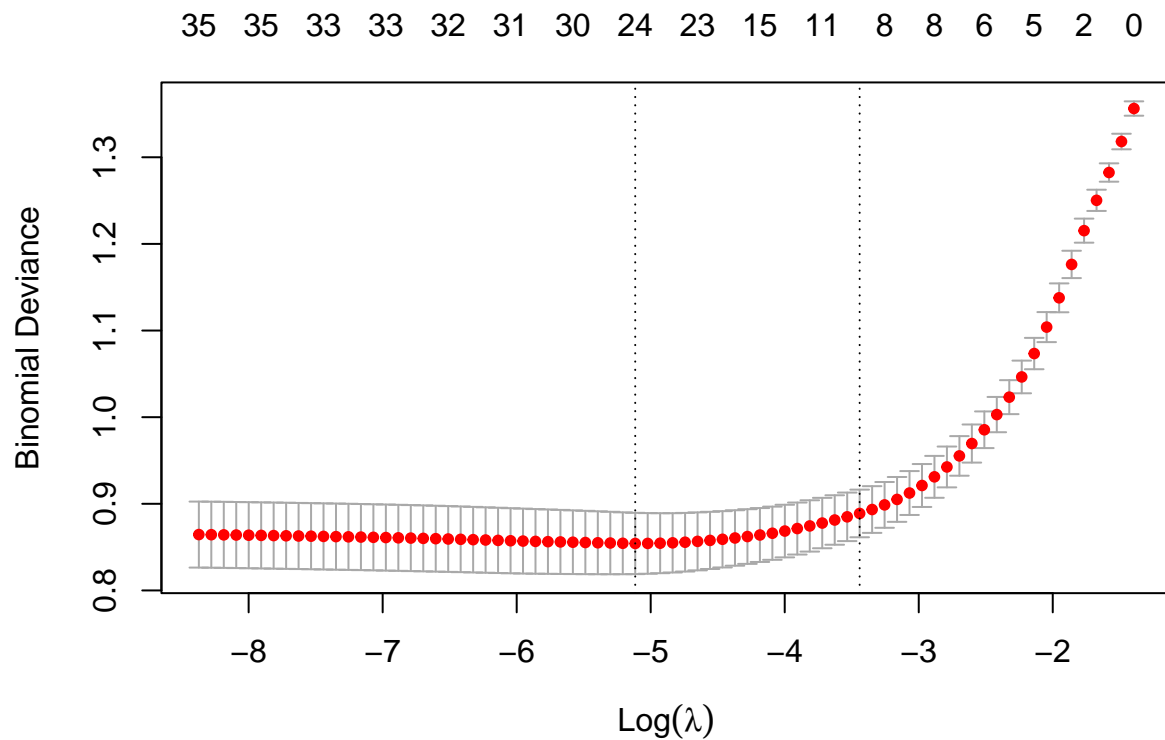
```
## Left school at 17 years Left school at 18 years Left school before 16 years
## 1 0 0 0
## 2 0 0 0
## 3 0 0 0
## 4 0 0 0
## 5 0 1 0
## 6 0 0 0
## Masters degree Professional certificate/ diploma
## 1 0 0
## 2 0 1
## 3 1 0
## 4 0 0
## 5 0 0
## 6 1 0
## Some college or university, no certificate or degree University degree
## 1 0 0
## 2 0 0
## 3 0 0
## 4 0 0
## 5 0 0
## 6 0 0
## Australia Canada New Zealand Republic of Ireland UK USA Asian Black
## 1 0 0 0 0 1 0 0 0
## 2 0 0 0 0 1 0 0 0
## 3 0 0 0 0 1 0 0 0
## 4 0 0 0 0 1 0 0 0
## 5 0 1 0 0 0 0 0 0
## 6 0 0 0 0 0 1 0 0
## Mixed-Black/Asian Mixed-White/Asian Mixed-White/Black White Gender
## 1 0 0 0 1 1
## 2 0 0 0 1 1
## 3 0 0 0 1 0
## 4 0 0 0 1 0
## 5 0 0 0 1 0
## 6 0 0 0 1 1
```

```
par(mfrow = c(1,4))
Alc_table <- table(data$Alcohol)
Caff_table <- table(data$Caff)
Cann_table <- table(data$Cannabis)
Nic_table <- table(data$Nicotine)
```

## LASSO Exploration

```
set.seed(123)
#Setting up matrices for lasso
x <- model.matrix(Cannabis~., data = data)[, -1]
y <- data$Cannabis
x.test <- as.matrix(test.data[, -10])
y.test <- test.data$Cannabis

#CV for Optimal Lambda
cv.out <- cv.glmnet(x, y, alpha = 1, family = 'binomial')
plot(cv.out)
```



```
lambda.opt <- cv.out$lambda.min
lambda.opt # 0.006588544
```

```
## [1] 0.006003236
```

```
# Lasso
lasso <- glmnet(x, y, alpha = 1, lambda = lambda.opt, family = "binomial")

#Lasso Regression
lasso.pred <- predict(lasso, s = lambda.opt, newx = x.test, type = "response")

# Assign a class to predictions based on boundary optimization found by this
# code.

cutoffs <- seq(.05, .95, by = .025)
preds <- rep(0, length(lasso.pred))
error.lasso <- rep(0, length(lasso.pred))
lasso.test.err <- rep(NA, length(cutoffs))

for(i in 1:length(cutoffs)){
  preds <- ifelse(lasso.pred < cutoffs[i], 0, 1)

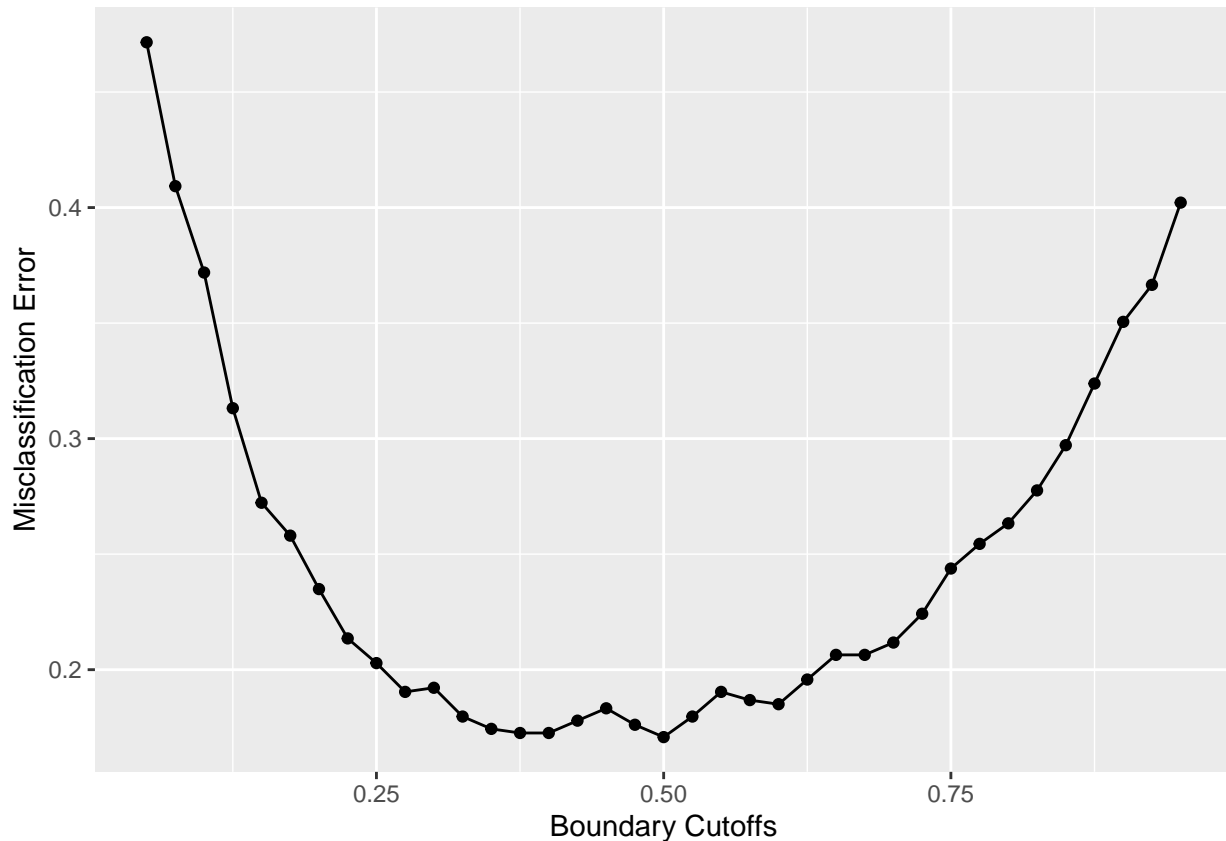
  for(e in 1:length(preds)){
    error.lasso[e] <- (preds[e] == y.test[e])
  }
}
```

```

  lasso.test.err[i] = (length(error.lasso)-sum(error.lasso))/length(error.lasso)
}

df <- data.frame(cutoffs, lasso.test.err)
ggplot(data = df, aes(x = cutoffs, y = lasso.test.err)) +
  geom_point() +
  geom_line() +
  xlab("Boundary Cutoffs") +
  ylab("Misclassification Error")

```



```
min(lasso.test.err) # 0.1725979
```

```
## [1] 0.1708185
```

```
cutoffs[which.min(lasso.test.err)] # 0.5
```

```
## [1] 0.5
```

```

# This process verified that 0.5 is the optimal cutoff to minimize test error
# using this lasso regression. We reached a test error rate of 0.1725979 or
# a success rate of 82.74%

```

```

#Predictor Coefficients after Lasso
coef(lasso)

```



```
## 38 x 1 sparse Matrix of class "dgCMatrix"
##
## (Intercept) -1.03621469
## Nscore -0.08503095
## Escore .
## Oscore 0.45119455
## AScore .
## Cscore -0.08470724
## Impulsive .
## SS 0.33743838
## Alcohol 0.12387857
## Caff .
## Nicotine 1.07268340
## '18-24' 0.91697043
## '25-34' .
## '35-44' .
## '45-54' -0.23307344
## '55-64' -0.13107620
## '65' -1.73335963
## 'Left school at 16 years' 0.36636759
## 'Left school at 17 years' .
## 'Left school at 18 years' 0.36093765
## 'Left school before 16 years' 0.39234000
## 'Masters degree' -0.42727518
## 'Professional certificate/ diploma' .
## 'Some college or university, no certificate or degree' 0.16860336
## 'University degree' -0.31771326
## Australia 0.04905410
## Canada .
## 'New Zealand' 0.92819062
## 'Republic of Ireland' .
## UK -1.14315070
## USA 0.40420861
## Asian -1.15089309
## Black .
## 'Mixed-Black/Asian' 0.67707959
## 'Mixed-White/Asian' 0.37697095
## 'Mixed-White/Black' .
## White .
## Gender 0.49309850
```

```
# Make a new data set removing the variables considered insignificant by the
# lasso regression.
```

```
data.lasso <- subset(data, select = -c(Escore, AScore, Impulsive, Caff, `35-44`, `Left school at 17 years`))
```

```
# Test and training sets for lasso
```

```
test.lasso <- data.lasso[test.i,]
```

```
train.lasso <- data.lasso[-test.i,]
```

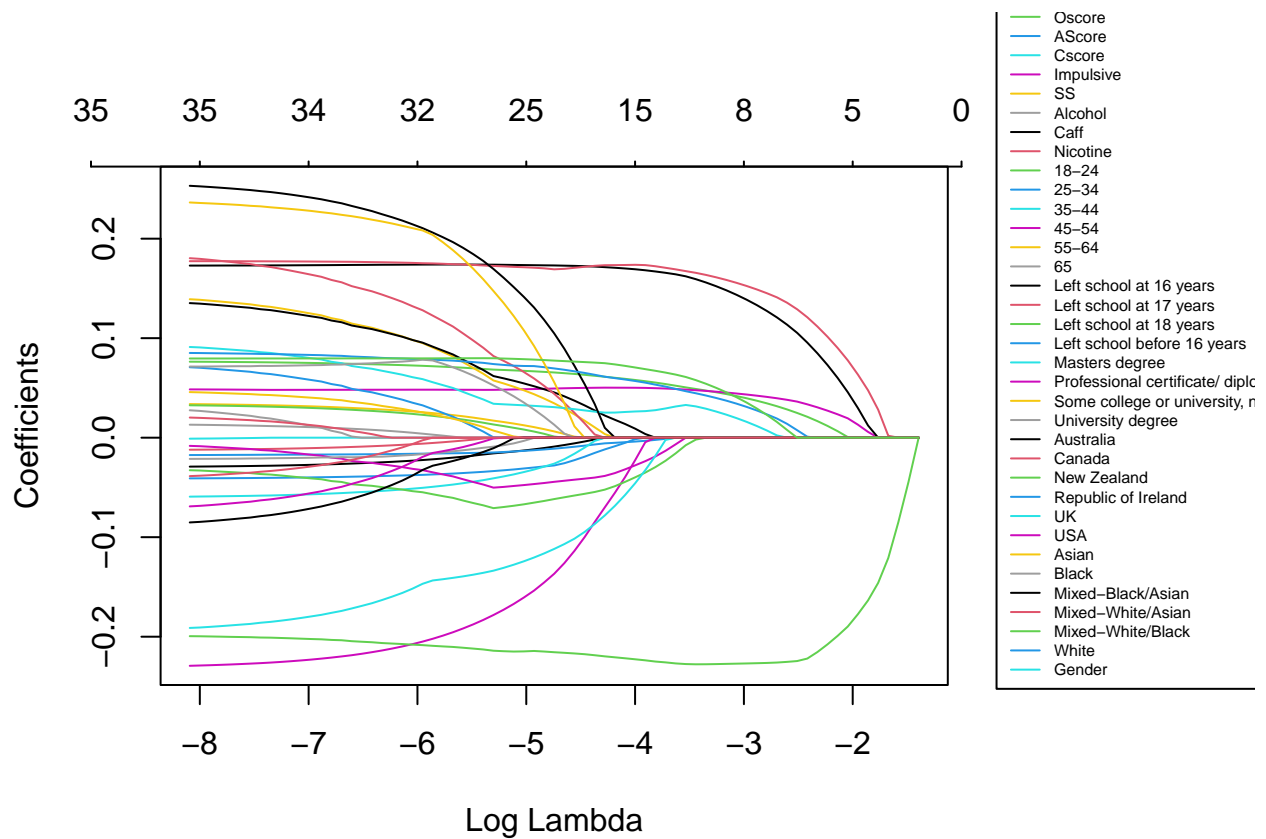
```
#Lasso Plot
```

```
par(mar=c(5, 4, 4, 8), xpd=TRUE)
```

```
lasso.plot <- glmnet(x, y, alpha = 1)
```

```
plot(lasso.plot, "lambda", col = 1:36)
```

```
legend("topright", inset=c(-0.6, -.4), lwd = 1, col= 1:37, legend = colnames(data[, -10]), cex = 0.5)
```

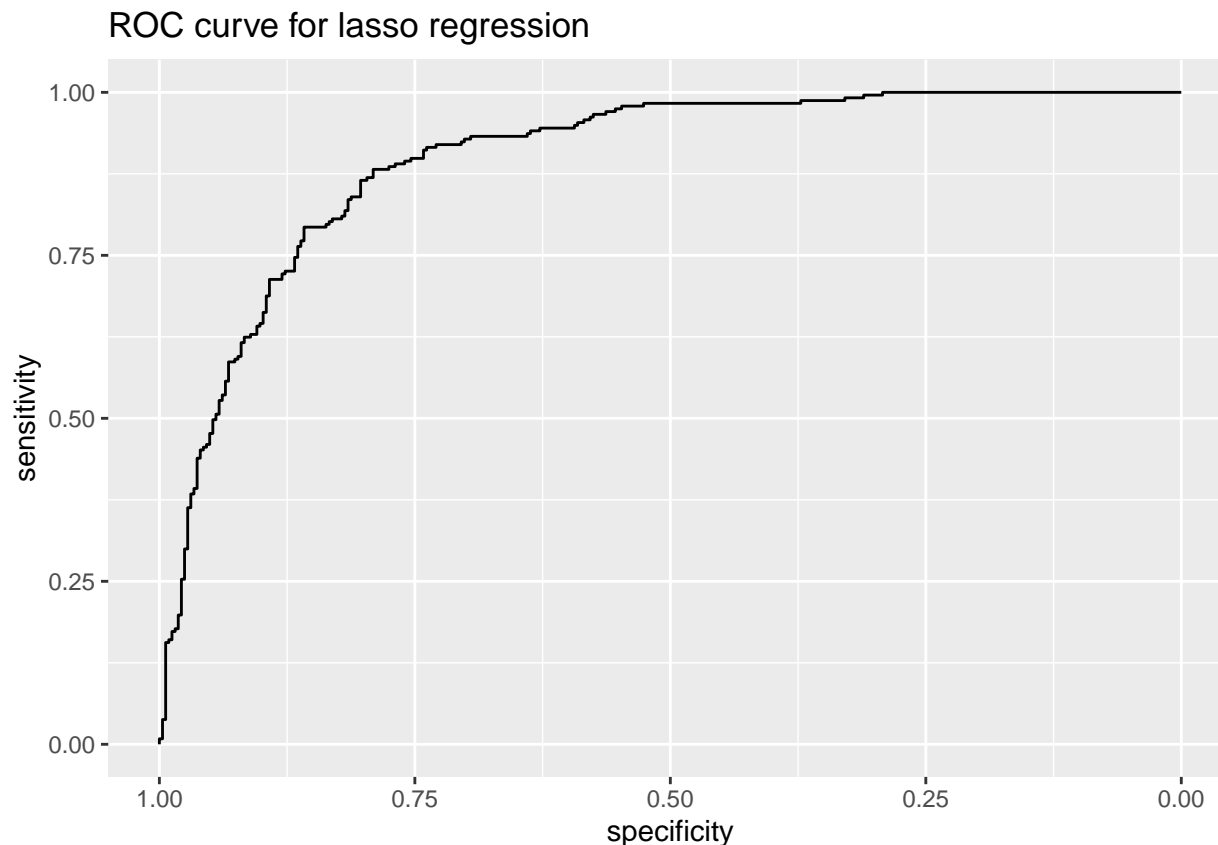


```
lasso.pred <- as.numeric(lasso.pred)
ROC.score.lasso <- roc(test.data$Cannabis, lasso.pred)
```

```
## Setting levels: control = 0, case = 1
```

```
## Setting direction: controls < cases
```

```
ggroc(ROC.score.lasso, legacy.axes = FALSE) +
  ggtitle("ROC curve for lasso regression")
```



Boosting - Finding the optimal shrinkage parameter

```
# Cycle through the shrinkage parameters to find the ideal value based on
# test MSE. Plot test error along different shrinkage values to find the
# ideal value.

# We will find the optimal cutoff for the prediction boundary using the
# optimal shrinkage coefficient found through this process. We will use
# 0.5 as the cutoff for this process and we will optimize the decision
# boundary based on the optimal shrinkage value to compensate for the unequal
# distribution of class 0 (not used Cannabis within the last month) and 1 (used
# Cannabis within the last month) in the data set. We also aim to reduce
# test error by optimizing the decision boundary.
set.seed(12345)
shrinkage <- seq(from = 0.01, to = .5, by = .0049)
boost.test.err <- rep(0, length(shrinkage))
error <- rep(0, nrow(test.data))

for(i in 1:length(shrinkage)){
  boost <- gbm(Cannabis ~ ., data = train.data,
               distribution = 'bernoulli',
               n.trees = 200,
               shrinkage = shrinkage[i])

  pred.boost <- predict(boost,
                        n.trees=100,
```

```

      newdata = test.data,
      type = 'response')

pclass.boost <- rep(NA, length(pred.boost))

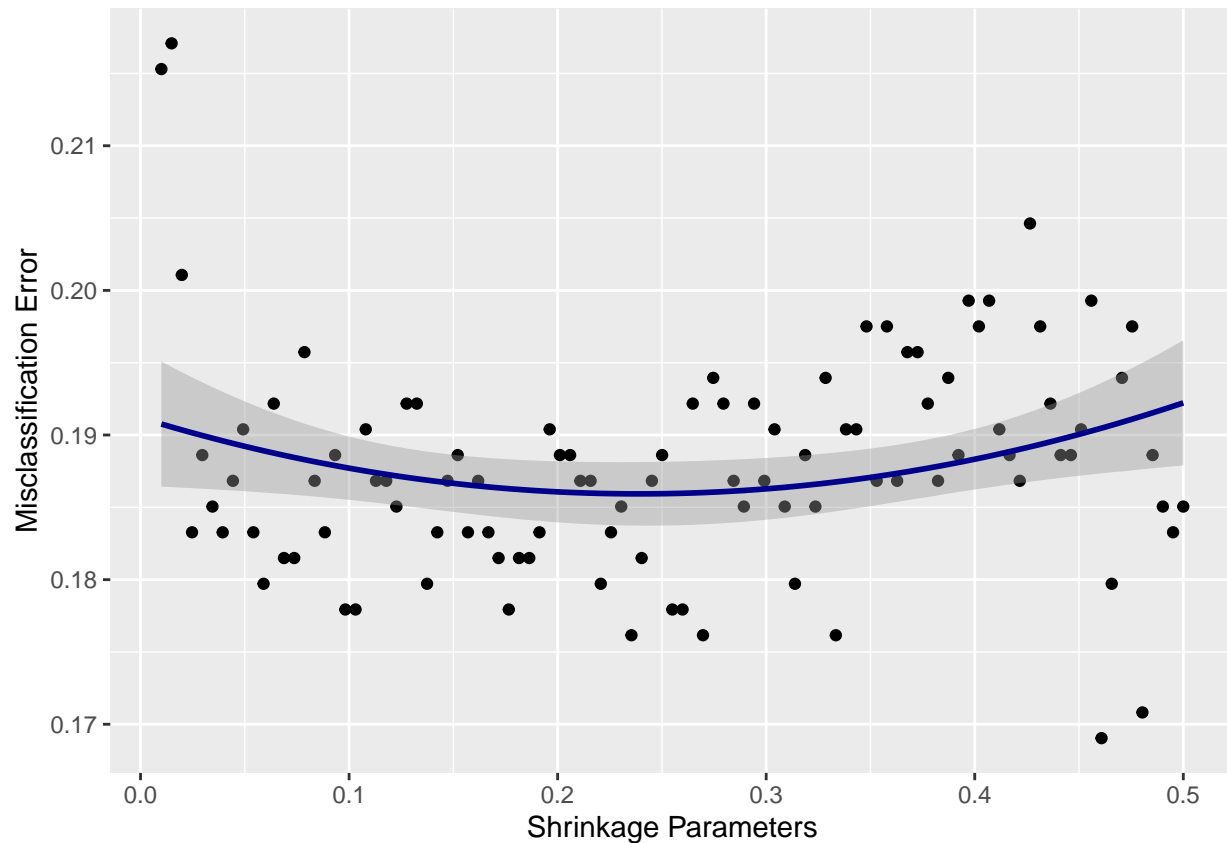
for(n in 1:length(pred.boost)){
  if(pred.boost[n] < 0.5){
    pclass.boost[n] = 0
  }else{
    pclass.boost[n] = 1
  }
}

for(e in 1:length(pclass.boost)){
  error[e] <- ((pclass.boost[e]) == test.data$Cannabis[e])
}

boost.test.err[i] = (length(error)-sum(error))/length(error)
}

df <- data.frame(shrinkage, boost.test.err)
ggplot(data = df, aes(x = shrinkage, y = boost.test.err)) +
  geom_point() +
  stat_smooth(method = "glm", formula = y ~ x + I(x^2), size = 1, col = "dark blue") +
  xlab("Shrinkage Parameters") +
  ylab("Misclassification Error")

```



```
shrinkage[which.min(boost.test.err)] # Use .23
```

```
## [1] 0.4608
```

```
min(boost.test.err) # 0.186
```

```
## [1] 0.1690391
```

*# From this chart, we see that shrinkage coefficients between .01 and .5 are the ideal values. I will not use the shrinkage value with the lowest test error (0.4804) because it appears to be an outlier. I will stick within the ideal range and use the shrinkage value of 0.186 as it had a low test error and it is the approximate bottom of the regression line of test errors.*

Boosting - Finding the optimal decision boundary

```
# Pick the ideal boundary cutoff using the ideal shrinkage value
# Plot the test MSE along different cutoff values of class 0/1
cutoffs <- seq(.05, .95, by = .025)
set.seed(12345)
boost.test.err <- rep(0, length(cutoffs))
error <- rep(0, nrow(test.data))

boost.2 <- gbm(Cannabis ~ ., data = train.data,
```

```

      distribution = 'bernoulli',
      n.trees = 200,
      shrinkage = .23)

pred.boost.2 <- predict(boost.2,
                        n.trees=100,
                        newdata = test.data,
                        type = 'response')

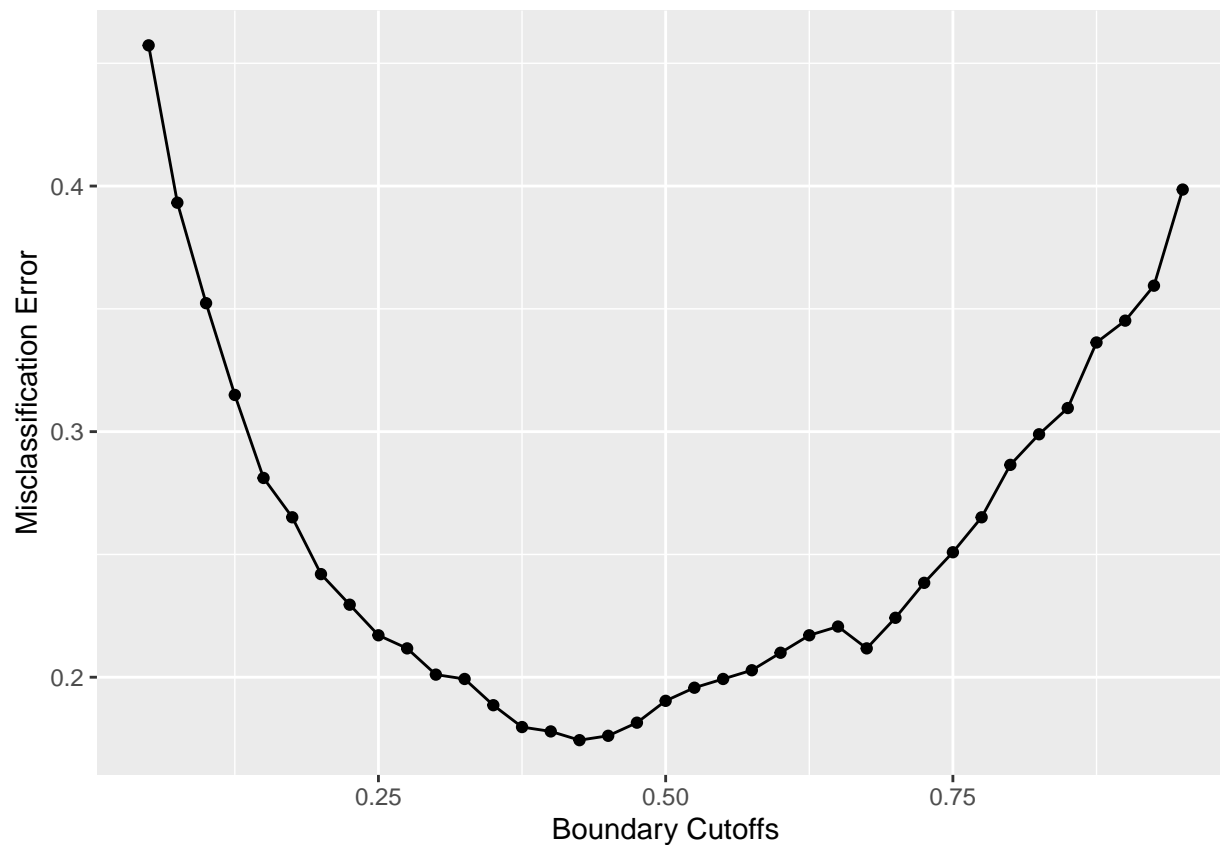
pclass.boost.2 <- rep(NA, length(pred.boost.2))

for(i in 1:length(cutoffs)){
  pclass.boost.2 <- ifelse(pred.boost.2 < cutoffs[i], 0, 1)

  for(e in 1:length(pclass.boost.2)){
    error[e] <- (pclass.boost.2[e] == test.data$Cannabis[e])
  }
  boost.test.err[i] = (length(error)-sum(error))/length(error)
}

df <- data.frame(cutoffs, boost.test.err)
ggplot(data = df, aes(x = cutoffs, y = boost.test.err)) +
  geom_point() +
  geom_line() +
  xlab("Boundary Cutoffs") +
  ylab("Misclassification Error")

```



```
cutoffs[which.min(boost.test.err)] # 0.425
```

```
## [1] 0.425
```

```
min(boost.test.err) # 0.1743772
```

```
## [1] 0.1743772
```

Boosting - Combine ideal shrinkage coefficient and ideal cutoff value

```
set.seed(12345)
error <- rep(0, nrow(test.data))

boost <- gbm(Cannabis ~ ., data = train.data,
             distribution = 'bernoulli',
             n.trees = 500,
             shrinkage = 0.23)

pred.boost <- predict(boost,
                     n.trees=100,
                     newdata = test.data,
                     type = 'response')

pclass.boost <- ifelse(pred.boost < .425, 0, 1)
```

```
for(e in 1:length(pclass.boost)){
  error[e] <- (pclass.boost[e] == test.data$Cannabis[e])
}
```

```
boost.test.err = (length(error)-sum(error))/length(error)
boost.test.err # 0.1743772
```

```
## [1] 0.1743772
```

```
boost.success.rate <- 1 - boost.test.err
boost.success.rate # 0.8256228
```

```
## [1] 0.8256228
```

```
# This code runs the model using the optimized shrinkage parameter and boundary
# cutoff. We reached an error rate of 17.43%, or a success rate of 82.56%.
```

## Logistic Regression

```
# In this code, we use logistic regression to generate a binary prediction
# model to predict if an individual has used Cannabis within the last month.
```

```
# We will cycle through decision boundaries from 5% to 95% and calculate test
# error at each cutoff. This will be used to find the error-minimizing decision
# boundary of our model.
```

```
set.seed(12345)
log.fit <- glm(Cannabis ~ ., data = train.data, family = "binomial")
```

```
## Warning: glm.fit: algorithm did not converge
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
cutoffs <- seq(.05, .95, by = .025)
probs <- predict(log.fit, test.data, type = "response")
preds <- rep(0, length(probs))
error.log <- rep(0, length(probs))
log.test.err <- rep(NA, length(cutoffs))
```

```
for(i in 1:length(cutoffs)){
  preds <- ifelse(probs < cutoffs[i], 0, 1)

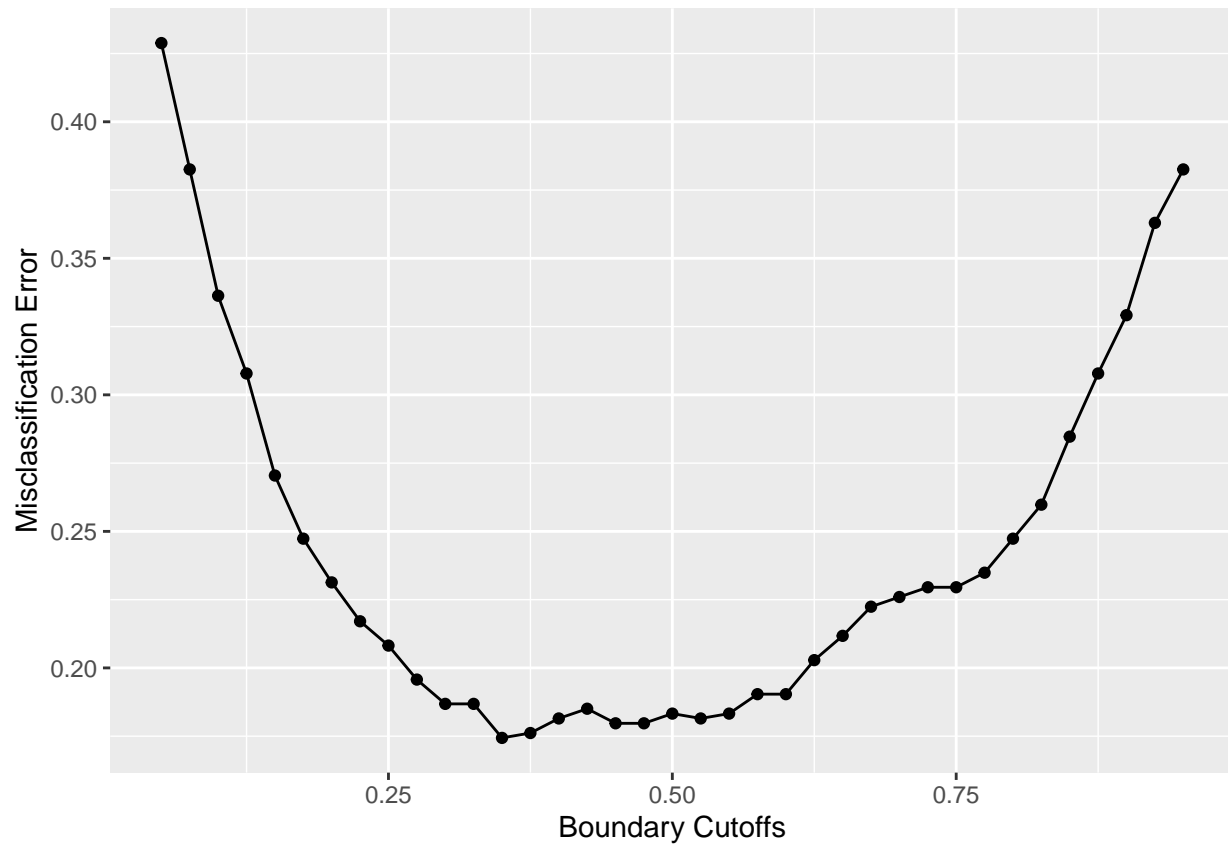
  for(e in 1:length(preds)){
    error.log[e] <- (preds[e] == test.data$Cannabis[e])
  }

  log.test.err[i] = (length(error.log)-sum(error.log))/length(error.log)
}
```

```
df <- data.frame(cutoffs, log.test.err)
```



```
ggplot(data = df, aes(x = cutoffs, y = log.test.err)) +
  geom_point() +
  geom_line() +
  xlab("Boundary Cutoffs") +
  ylab("Misclassification Error")
```



```
min(log.test.err) # 0.1761566
```

```
## [1] 0.1743772
```

```
cutoffs[which.min(log.test.err)] # 0.35
```

```
## [1] 0.35
```

```
# =====
# Identified ideal cutoff at 0.325 Rerun logistic regression using the ideal
# cutoff and calculate the confusion matrix to see the false positive rate,
# false negative rate, and model accuracy.
```

```
log.fit <- glm(Cannabis ~ ., data = train.data, family = "binomial")
```

```
## Warning: glm.fit: algorithm did not converge
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
summary(log.fit)
```

```
##
## Call:
## glm(formula = Cannabis ~ ., family = "binomial", data = train.data)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7728  -0.5792  -0.2688   0.5949   2.4302
##
## Coefficients:
##                                     Estimate Std. Error
## (Intercept)                    -6.797e+13  9.079e+13
## Nscore                        -1.995e-01  9.307e-02
## Escore                       -9.156e-02  9.697e-02
## Oscore                        5.252e-01  9.564e-02
## AScore                       1.701e-03  8.564e-02
## Cscore                       -8.587e-02  9.667e-02
## Impulsive                     -1.218e-02  1.087e-01
## SS                           3.742e-01  1.151e-01
## Alcohol                      5.020e-01  2.080e-01
## Caff                         2.393e-01  3.447e-01
## Nicotine                     1.175e+00  1.601e-01
## '18-24'                      6.797e+13  9.079e+13
## '25-34'                      6.797e+13  9.079e+13
## '35-44'                      6.797e+13  9.079e+13
## '45-54'                      6.797e+13  9.079e+13
## '55-64'                      6.797e+13  9.079e+13
## '65'                        -4.436e+15  9.079e+13
## 'Left school at 16 years'     5.997e-01  5.131e-01
## 'Left school at 17 years'     1.333e+00  7.342e-01
## 'Left school at 18 years'     8.872e-01  5.047e-01
## 'Left school before 16 years' 1.585e+00  7.303e-01
## 'Masters degree'             -1.500e-01  4.113e-01
## 'Professional certificate/ diploma' 5.638e-01  4.174e-01
## 'Some college or university, no certificate or degree' 4.859e-01  4.087e-01
## 'University degree'          -1.335e-01  3.925e-01
## Australia                    9.315e-01  5.236e-01
## Canada                      3.252e-02  4.267e-01
## 'New Zealand'                1.863e+00  1.414e+00
## 'Republic of Ireland'        -2.967e-01  6.786e-01
## UK                           -9.892e-01  2.979e-01
## USA                          6.479e-01  3.030e-01
## Asian                       -1.483e+00  9.593e-01
## Black                       -1.018e+00  8.492e-01
## 'Mixed-Black/Asian'          2.555e+01  2.161e+05
## 'Mixed-White/Asian'          5.387e-01  8.629e-01
## 'Mixed-White/Black'         -5.882e-01  9.162e-01
## White                       -4.016e-01  4.088e-01
## Gender                      6.471e-01  1.661e-01
##
## z value Pr(>|z|)
## (Intercept)                -0.749 0.454066
## Nscore                      -2.144 0.032047 *
```

```

## Escore -0.944 0.345052
## Oscore 5.492 3.98e-08 ***
## AScore 0.020 0.984158
## Cscore -0.888 0.374375
## Impulsive -0.112 0.910797
## SS 3.253 0.001143 **
## Alcohol 2.414 0.015792 *
## Caff 0.694 0.487549
## Nicotine 7.337 2.19e-13 ***
## '18-24' 0.749 0.454066
## '25-34' 0.749 0.454066
## '35-44' 0.749 0.454066
## '45-54' 0.749 0.454066
## '55-64' 0.749 0.454066
## '65' -48.856 < 2e-16 ***
## 'Left school at 16 years' 1.169 0.242562
## 'Left school at 17 years' 1.816 0.069438 .
## 'Left school at 18 years' 1.758 0.078784 .
## 'Left school before 16 years' 2.171 0.029949 *
## 'Masters degree' -0.365 0.715286
## 'Professional certificate/ diploma' 1.351 0.176837
## 'Some college or university, no certificate or degree' 1.189 0.234541
## 'University degree' -0.340 0.733867
## Australia 1.779 0.075241 .
## Canada 0.076 0.939248
## 'New Zealand' 1.318 0.187582
## 'Republic of Ireland' -0.437 0.661984
## UK -3.321 0.000898 ***
## USA 2.138 0.032505 *
## Asian -1.546 0.122194
## Black -1.199 0.230713
## 'Mixed-Black/Asian' 0.000 0.999906
## 'Mixed-White/Asian' 0.624 0.532427
## 'Mixed-White/Black' -0.642 0.520857
## White -0.982 0.325891
## Gender 3.896 9.78e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 1782.5 on 1313 degrees of freedom
## Residual deviance: 1069.9 on 1276 degrees of freedom
## AIC: 1145.9
##
## Number of Fisher Scoring iterations: 25

```

```

probs <- predict(log.fit, test.data, type = "response")
preds <- rep(0, length(probs))
preds[probs > 0.35] = 1

preds <- as.factor(preds)
test.data$Cannabis <- as.factor(test.data$Cannabis)
confusionMatrix(test.data$Cannabis, preds) # 82.38%

```

```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction  0    1
##           0 259  66
##           1  32 205
##
##           Accuracy : 0.8256
##           95% CI : (0.7917, 0.8561)
##       No Information Rate : 0.5178
##       P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.6493
##
##  McNemar's Test P-Value : 0.0008576
##
##           Sensitivity : 0.8900
##           Specificity : 0.7565
##       Pos Pred Value : 0.7969
##       Neg Pred Value : 0.8650
##           Prevalence : 0.5178
##       Detection Rate : 0.4609
##       Detection Prevalence : 0.5783
##       Balanced Accuracy : 0.8232
##
##       'Positive' Class : 0
##
```

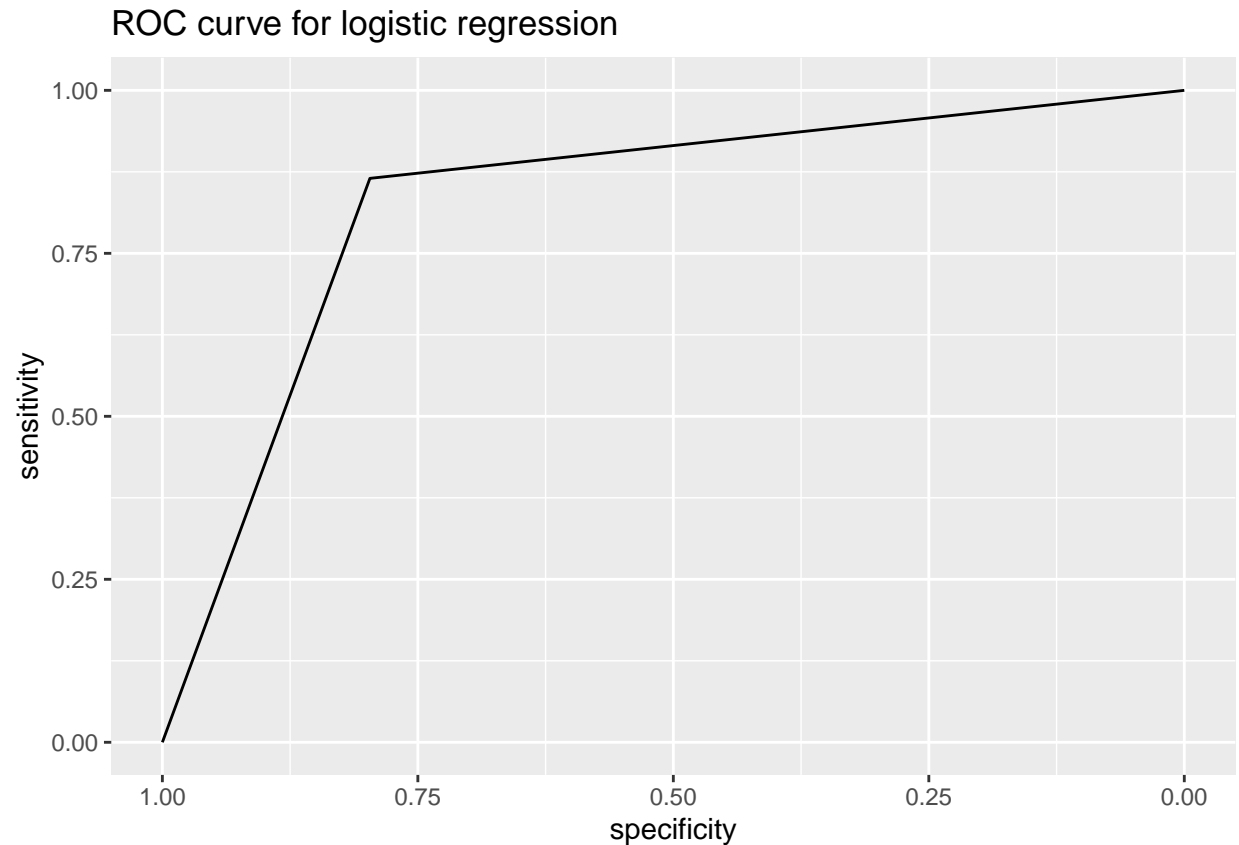
```
# Accuracy of 82.38%
# FPR = 5.69395%
# FNR = 11.92171%
```

```
#ROC-curve using pROC library
test.data$Cannabis <- as.numeric(test.data$Cannabis)
preds <- as.numeric(preds)
ROC.score.log <- roc(test.data$Cannabis, preds)
```

```
## Setting levels: control = 1, case = 2
```

```
## Setting direction: controls < cases
```

```
ggroc(ROC.score.log, legacy.axes = FALSE) +
  ggtitle("ROC curve for logistic regression")
```



#### RECLEAN DATA

```
set.seed(12345)
data <- read.csv("Drug_Consumption.csv")

# Remove the over-claimers using the control drug "Semer"
data <- subset(data, data$Semer == "CL0")

for(i in 14:ncol(data)){
  data[,i] <- as.numeric(data[, i] == "CL4" | data[, i] == "CL5" | data[, i] == "CL6")
}

# Drop 65+
data <- data %>% mutate(dummy=1) %>%
spread(key=Age,value=dummy,fill=0)

# Drop Doctorate
data <- data %>% mutate(dummy=1) %>%
spread(key=Education,value=dummy,fill=0)

# Drop other
data <- data %>% mutate(dummy=1) %>%
spread(key=Country,value=dummy,fill=0)

# Drop other
data <- data %>% mutate(dummy=1) %>%
```

```

spread(key=Ethnicity,value=dummy,fill=0)

# Drop 'F' variable and rename to gender
data <- data %>% mutate(dummy=1) %>%
spread(key=Gender,value=dummy, fill=0)

# Drop variables that we aren't using.
drop <- c("ID", "65+", "Doctorate degree", "Other", "F", "Amphet", "Amyl", "Benzos", "Choc", "Crack", "Coke", "E")
data <- data[,!(names(data) %in% drop)]

names(data)[names(data) == "M"] <- "Gender"

# Split into test and train data
test.i <- sample(1:nrow(data), .3*nrow(data))
test.data <- data[test.i,]
train.data <- data[-test.i,]

```

kNN

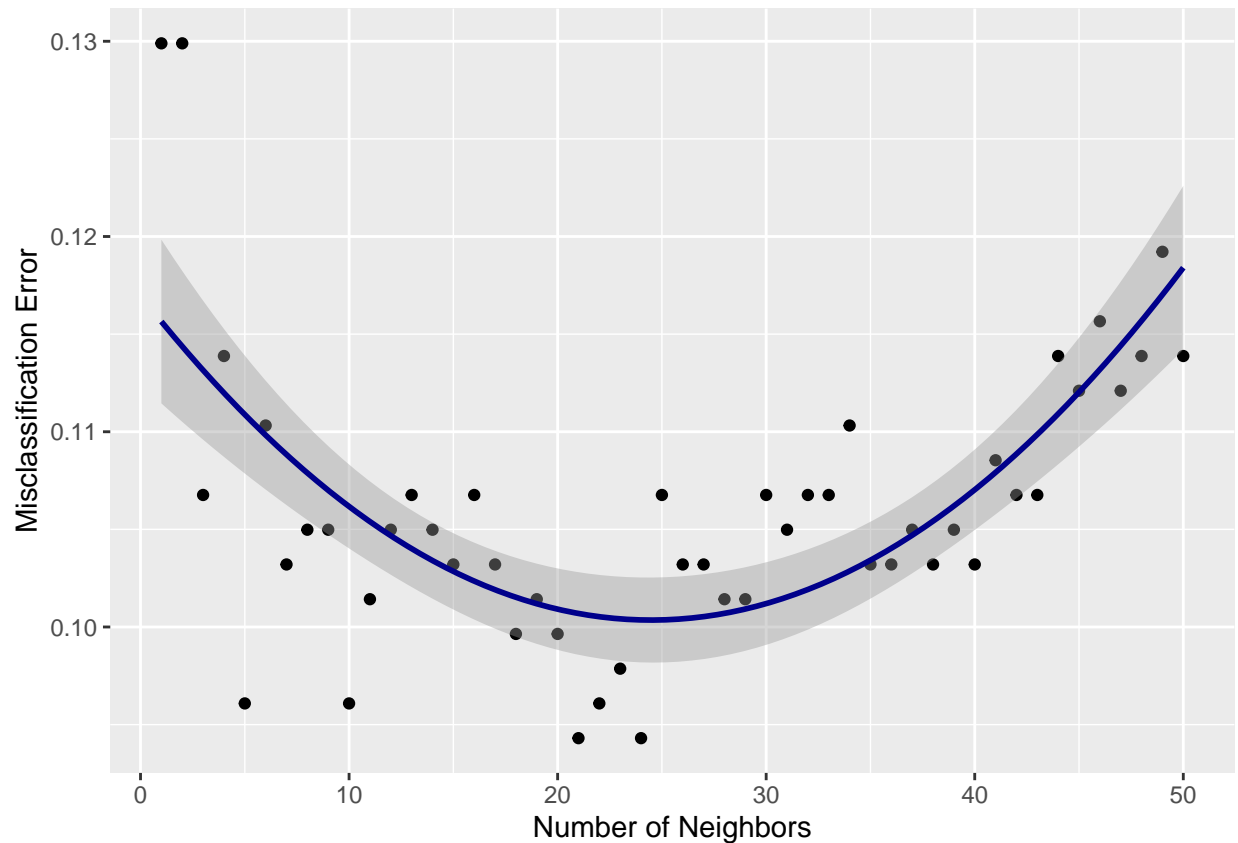
```

set.seed(12345)
ks <- 1:50
knn.error <- rep(0, length(ks))

for(i in 1:length(ks)){
  pred.knn <- knn(train.data, test.data, train.data$Cannabis, k = ks[i])
  table.knn <- table(pred.knn, test.data$Cannabis)
  knn.error[i] <- (table.knn[1,2] + table.knn[2,1])/(table.knn[1,2] + table.knn[2,1] + table.knn[2,2] +
}

df.knn = data.frame(ks, knn.error)
ggplot(data = df.knn, aes(x = ks, y = knn.error)) +
  geom_point() +
  stat_smooth(method = "glm", formula = y ~ x + I(x^2), size = 1, col = "dark blue") +
  xlab("Number of Neighbors") +
  ylab("Misclassification Error")

```



```
which.min(knn.error) # k = 22 results in the minimum error
```

```
## [1] 21
```

```
min(knn.error) # 0.09252669, or a success rate of 90.74733%
```

```
## [1] 0.09430605
```

SVM

```
x_SVM <- train.data[,-10]
y_SVM <- train.data[,10]
SVM_data <- data.frame(x = x_SVM, y = as.factor(y_SVM))
SVM_model <- svm(y~., data = SVM_data, kernel = "linear", scale = FALSE, cost = 10)
SVM_model
```

```
##
## Call:
## svm(formula = y ~ ., data = SVM_data, kernel = "linear", cost = 10,
##      scale = FALSE)
##
##
## Parameters:
##   SVM-Type:  C-classification
```

```
## SVM-Kernel: linear
## cost: 10
##
## Number of Support Vectors: 588
```

```
SVM_predict <- predict(SVM_model, data.frame(x = test.data[, -10], y = test.data[, 10]))
# Ideal cost is 1.92875e-22
# minimum error is 0.192923

table.SVM <- table(SVM_predict, test.data$Cannabis)
table.SVM
```

```
##
## SVM_predict 0 1
##           0 276 51
##           1 49 186
```

```
(table.SVM[1,2] + table.SVM[2,1]) / (table.SVM[1,2] + table.SVM[2,1] + table.SVM[1,1] + table.SVM[2,2])
```

```
## [1] 0.1779359
```

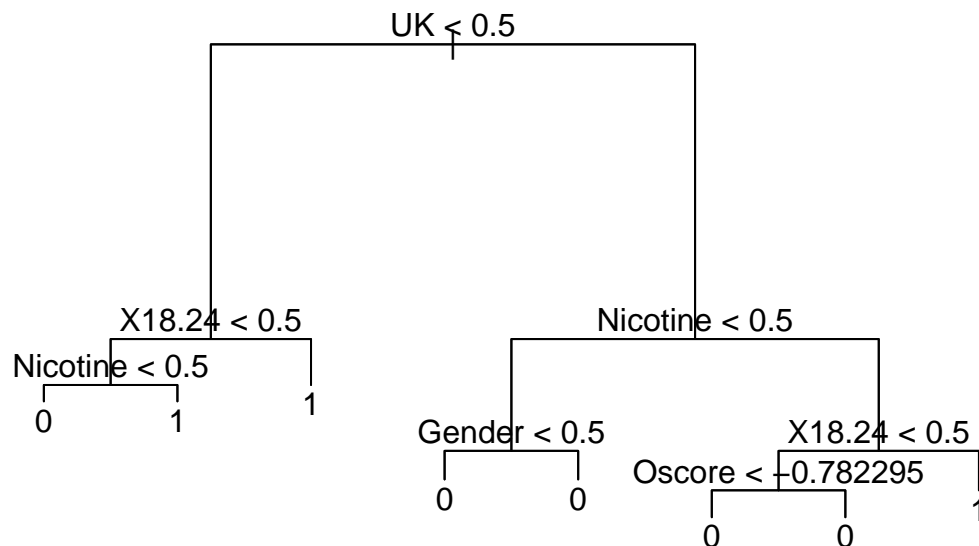
Decision Trees

```
set.seed(12345)
tree_train <- data.frame(train.data)
tree_test <- data.frame(test.data)
treefit <- tree(as.factor(Cannabis) ~ ., data = tree_train)
summary(treefit)
```

```
##
## Classification tree:
## tree(formula = as.factor(Cannabis) ~ ., data = tree_train)
## Variables actually used in tree construction:
## [1] "UK" "X18.24" "Nicotine" "Gender" "Oscore"
## Number of terminal nodes: 8
## Residual mean deviance: 0.8736 = 1141 / 1306
## Misclassification error rate: 0.2032 = 267 / 1314
```

```
# variables used : UK, 18-24, Oscore, Nicotine, gender, and SS
plot(treefit)
text(treefit)
```





```

tree.predict <- predict(treefit, tree_test, type = "class")
tree.table <- table(tree.predict, tree_test$Cannabis)
tree.error <- (tree.table[1,2] + tree.table[2,1]) / (tree.table[1,2] + tree.table[2,1] + tree.table[1,1] + tree.table[2,2])
tree.error # 0.1992883

```

```
## [1] 0.1992883
```

Random Forest

```

set.seed(12345)
rF <- randomForest(as.factor(Cannabis)~., data = tree_train, importance = TRUE)
rf.predict <- predict(rF, tree_test)
rf.table <- table(rf.predict, tree_test$Cannabis) # .2009 error rate
rf.table

```

```

##
## rf.predict    0    1
##              0 272  47
##              1  53 190

```

```

rf.error <- (rf.table[1,2] + rf.table[2,1]) / (rf.table[1,2] + rf.table[2,1] + rf.table[1,1] + rf.table[2,2])
rf.error # 0.1814947

```

```
## [1] 0.1779359
```

LDA

```
set.seed(12345)
lda.fit <- lda(as.factor(Cannabis)~., data = train.data)
```

```
## Warning in lda.default(x, grouping, ...): variables are collinear
```

```
lda.fit
```

```
## Call:
## lda(as.factor(Cannabis) ~ ., data = train.data)
##
## Prior probabilities of groups:
##      0      1
## 0.585997 0.414003
##
## Group means:
##      Nscore      Escore      Oscore      AScore      Cscore      Impulsive
## 0 -0.05089368 -0.0002935974 -0.2980838  0.07746401  0.1988430 -0.2055659
## 1  0.02314386  0.0304835294  0.4436558 -0.12047708 -0.2295484  0.3013005
##      SS      Alcohol      Caff      Nicotine      '18-24'      '25-34'      '35-44'
## 0 -0.3303218  0.8155844  0.9259740  0.2974026  0.1701299  0.2844156  0.2428571
## 1  0.4434787  0.8400735  0.9522059  0.6875000  0.5588235  0.2150735  0.1213235
##      '45-54'      '55-64'      '65'      'Left school at 16 years'
## 0 0.22597403 0.05844156 0.01818182 0.06363636
## 1 0.07904412 0.02573529 0.00000000 0.03492647
##      'Left school at 17 years'      'Left school at 18 years'
## 0      0.01038961      0.03246753
## 1      0.02389706      0.07536765
##      'Left school before 16 years'      'Masters degree'
## 0      0.01168831      0.20259740
## 1      0.01654412      0.08639706
##      'Professional certificate/ diploma'
## 0      0.1649351
## 1      0.1286765
##      'Some college or university, no certificate or degree'      'University degree'
## 0      0.1441558      0.3090909
## 1      0.4283088      0.1764706
##      Australia      Canada      'New Zealand'      'Republic of Ireland'      UK      USA
## 0 0.01558442 0.03896104 0.001298701 0.009090909 0.7597403 0.1298701
## 1 0.04411765 0.05147059 0.003676471 0.016544118 0.2591912 0.5275735
##      Asian      Black      'Mixed-Black/Asian'      'Mixed-White/Asian'
## 0 0.023376623 0.02597403 0.000000000 0.009090909
## 1 0.003676471 0.01102941 0.005514706 0.016544118
##      'Mixed-White/Black'      White      Gender
## 0      0.007792208 0.9142857 0.3805195
## 1      0.011029412 0.8897059 0.6801471
##
## Coefficients of linear discriminants:
##
## LD1
## Nscore      -0.128719708
## Escore      -0.063894941
## Oscore      0.309557993
```

```
## AScore -0.009474028
## Cscore -0.044445079
## Impulsive -0.006703214
## SS 0.205778013
## Alcohol 0.273200620
## Caff 0.099991319
## Nicotine 0.747972175
## '18-24' 0.528292204
## '25-34' -0.030754587
## '35-44' -0.171371758
## '45-54' -0.343321680
## '55-64' -0.197907859
## '65' -1.129097364
## 'Left school at 16 years' 0.388558815
## 'Left school at 17 years' 0.814517525
## 'Left school at 18 years' 0.522752868
## 'Left school before 16 years' 0.944261957
## 'Masters degree' -0.092648764
## 'Professional certificate/ diploma' 0.347386663
## 'Some college or university, no certificate or degree' 0.344774309
## 'University degree' -0.038493892
## Australia 0.476439450
## Canada -0.080914690
## 'New Zealand' 1.228977238
## 'Republic of Ireland' -0.174970800
## UK -0.756964941
## USA 0.402876112
## Asian -0.649234781
## Black -0.407009859
## 'Mixed-Black/Asian' 0.989634489
## 'Mixed-White/Asian' 0.109803221
## 'Mixed-White/Black' -0.393579887
## White -0.282101194
## Gender 0.380486085
```

```
lda.pred <- predict(lda.fit, test.data)$class
table.lda <- table(lda.pred, test.data$Cannabis)

lda.error <- (table.lda[1,2] + table.lda[2,1])/(table.lda[1,2] + table.lda[2,1] + table.lda[1,1] + table.lda[2,2])
lda.error # 0.1886121
```

```
## [1] 0.1886121
```