

Machine Learning | Homework 5

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

```
## Loading required package: lattice
```

```
library(ISLR)
library(MASS)
library(ROCR)
```

```
## Loading required package: gplots
```

```
##
## Attaching package: 'gplots'
```

```
## The following object is masked from 'package:stats':
##
##      lowess
```

```
library(rpart.plot)
```

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

Getting the data.

```
german_credit <-
  read.table("http://archive.ics.uci.edu/ml/machine-learning-databases/statlog/german/german.data")

colnames(german_credit) <- c("chk_acct", "duration", "credit_his", "purpose",
  "amount", "saving_acct", "present_emp",
  "installment_rate", "sex", "other_debtor",
  "present_resid", "property", "age",
  "other_install", "housing", "n_credits",
  "job", "n_people", "telephone",
  "foreign", "response")

ind <- (german_credit$response == 2)
german_credit$response <- rep("Negative_class", length(german_credit$response))
german_credit$response[ind] <- "Positive_class"
german_credit$response <- as.factor(german_credit$response)
```

Divide dataset into training (80%) and test sets (20%).

```
set.seed(1)
train_index <- createDataPartition(german_credit$response, p = 0.8, list = F)
train_data <- german_credit[train_index,]
test_data <- german_credit[-train_index,]
```

1. Apply classification trees (method = “rpart” in the Caret package) on the training set with 10-fold cross-validation. Show the confusion matrices and the ROC curves for the training and test sets. Draw the corresponding decision tree (score = 30).

```
grid = expand.grid(cp = seq(0, 0.4, by = 0.01))

train_control_1 = trainControl(method = "cv",
                               number = 10,
                               classProbs = T)

model_tree_1 = train(response ~ .,
                      data = train_data,
                      method = "rpart",
                      tuneLength = 30,
                      trControl = train_control_1,
                      tuneGrid = grid)

model_tree_1
```

```
## CART
##
## 800 samples
## 20 predictor
## 2 classes: 'Negative_class', 'Positive_class'
##
## No pre-processing
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 720, 720, 720, 720, 720, 720, ...
## Resampling results across tuning parameters:
##
##   cp    Accuracy  Kappa
##   0.00  0.70000   0.25005688
##   0.01  0.69625   0.20343988
##   0.02  0.70250   0.19655228
##   0.03  0.70500   0.09199016
##   0.04  0.69250   0.01739927
##   0.05  0.70000   0.00000000
##   0.06  0.70000   0.00000000
##   0.07  0.70000   0.00000000
##   0.08  0.70000   0.00000000
##   0.09  0.70000   0.00000000
##   0.10  0.70000   0.00000000
##   0.11  0.70000   0.00000000
##   0.12  0.70000   0.00000000
##   0.13  0.70000   0.00000000
##   0.14  0.70000   0.00000000
```

```
## 0.15 0.70000 0.00000000
## 0.16 0.70000 0.00000000
## 0.17 0.70000 0.00000000
## 0.18 0.70000 0.00000000
## 0.19 0.70000 0.00000000
## 0.20 0.70000 0.00000000
## 0.21 0.70000 0.00000000
## 0.22 0.70000 0.00000000
## 0.23 0.70000 0.00000000
## 0.24 0.70000 0.00000000
## 0.25 0.70000 0.00000000
## 0.26 0.70000 0.00000000
## 0.27 0.70000 0.00000000
## 0.28 0.70000 0.00000000
## 0.29 0.70000 0.00000000
## 0.30 0.70000 0.00000000
## 0.31 0.70000 0.00000000
## 0.32 0.70000 0.00000000
## 0.33 0.70000 0.00000000
## 0.34 0.70000 0.00000000
## 0.35 0.70000 0.00000000
## 0.36 0.70000 0.00000000
## 0.37 0.70000 0.00000000
## 0.38 0.70000 0.00000000
## 0.39 0.70000 0.00000000
## 0.40 0.70000 0.00000000
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was cp = 0.03.
```

Now to see the confusion matrices and ROC curves for both train and test datasets, we need to make predictions with the help of the model constructed above:

```
pred_tree_train_1 = predict(model_tree_1, newdata = train_data, type = "raw")
pred_tree_test_1 = predict(model_tree_1, newdata = test_data, type = "raw")
```

So the confusion matrix for the Training dataset is as follows:

```
cm_train_1 = confusionMatrix(pred_tree_train_1, data = train_data$response, positive = "Positive_class")
cm_train_1
```

```
## Confusion Matrix and Statistics
##
##              Reference
## Prediction   Negative_class Positive_class
## Negative_class      497             63
## Positive_class      135            105
##
##              Accuracy : 0.7525
##              95% CI : (0.7211, 0.7821)
##      No Information Rate : 0.79
##      P-Value [Acc > NIR] : 0.9954
##
```

```
##           Kappa : 0.3555
## McNemar's Test P-Value : 4.517e-07
##
##           Sensitivity : 0.6250
##           Specificity : 0.7864
##           Pos Pred Value : 0.4375
##           Neg Pred Value : 0.8875
##           Prevalence : 0.2100
##           Detection Rate : 0.1313
##           Detection Prevalence : 0.3000
##           Balanced Accuracy : 0.7057
##
##           'Positive' Class : Positive_class
##
```

And the one for Test dataset is:

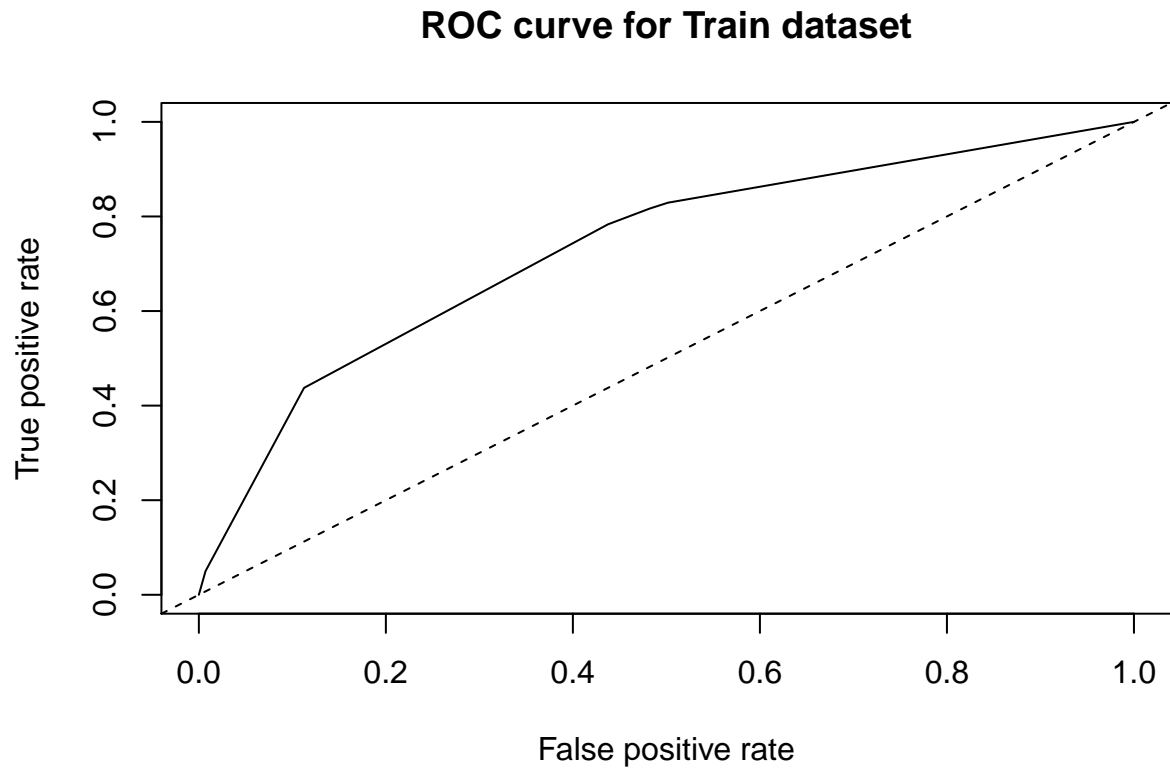
```
cm_test_1 = confusionMatrix(pred_tree_test_1, data = test_data$response, positive = "Positive_class")
cm_test_1
```

```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction   Negative_class Positive_class
## Negative_class      120          20
## Positive_class       28          32
##
##           Accuracy : 0.76
##           95% CI : (0.6947, 0.8174)
##           No Information Rate : 0.74
##           P-Value [Acc > NIR] : 0.2894
##
##           Kappa : 0.4059
## McNemar's Test P-Value : 0.3123
##
##           Sensitivity : 0.6154
##           Specificity : 0.8108
##           Pos Pred Value : 0.5333
##           Neg Pred Value : 0.8571
##           Prevalence : 0.2600
##           Detection Rate : 0.1600
##           Detection Prevalence : 0.3000
##           Balanced Accuracy : 0.7131
##
##           'Positive' Class : Positive_class
##
```

In addition, here are the ROC curves for the training and testing data respectively:

```
pred_tree_train_prob_1 = predict(model_tree_1, newdata = train_data, type = "prob")
prediction_obj_train_1 = prediction(pred_tree_train_prob_1[,2], train_data$response)
```

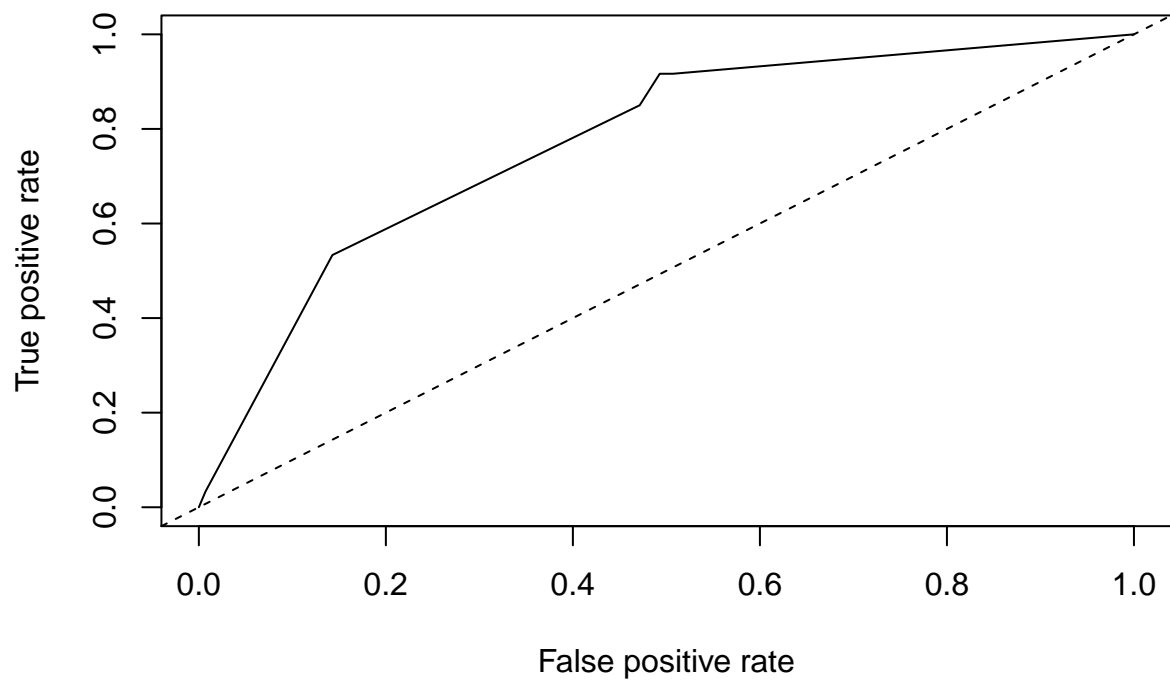
```
ROC_train_1 = performance(prediction_obj_train_1, "tpr", "fpr")
plot(ROC_train_1, main = "ROC curve for Train dataset")
abline(0, 1, lty = 2)
```



```
pred_tree_test_prob_1 = predict(model_tree_1, newdata = test_data, type = "prob")
prediction_obj_test_1 = prediction(pred_tree_test_prob_1[,2], test_data$response)

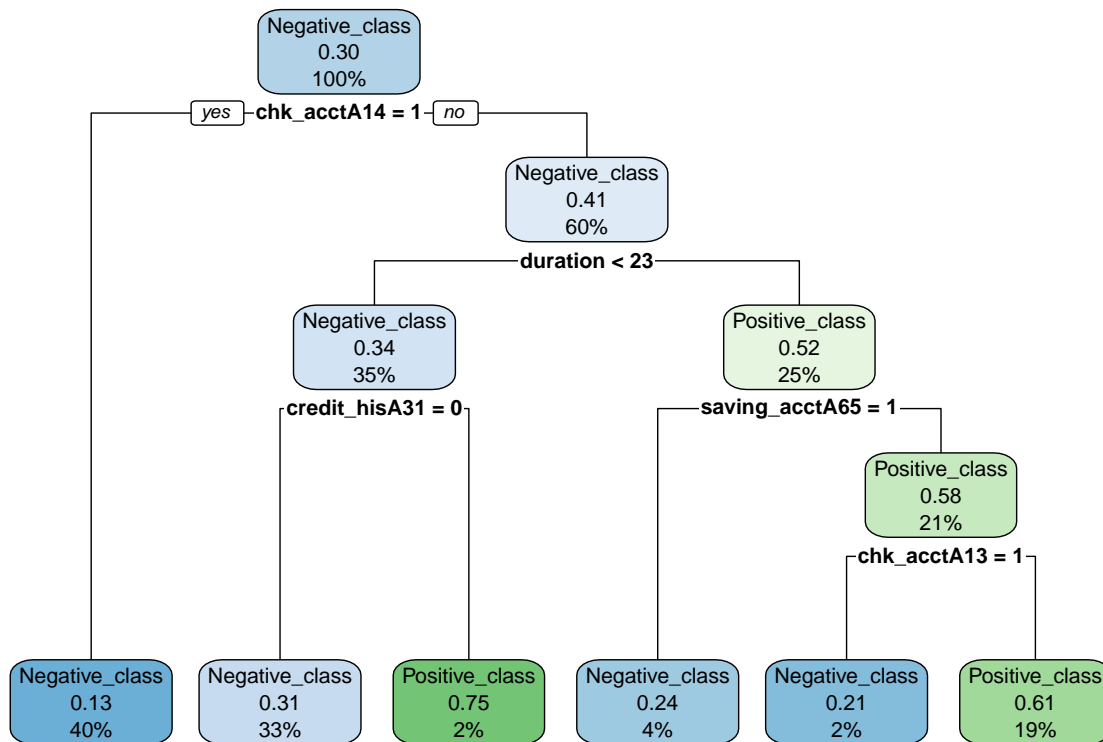
ROC_test_1 = performance(prediction_obj_test_1, "tpr", "fpr")
plot(ROC_test_1, main = "ROC curve for Test dataset")
abline(0, 1, lty = 2)
```

ROC curve for Test dataset



And here is the decision tree:

```
rpart.plot(model_tree_1$finalModel)
```



2. Apply random forests (method = "rf") on the training set. Show the confusion matrices and the ROC curves for the training and test sets. (score = 30).

```

grid2 = expand.grid(mtry = seq(2, ncol(train_data)-1, by = 1))

train_control_2 = trainControl(method = "repeatedcv",
                               number = 10,
                               repeats = 3,
                               allowParallel = TRUE)

model_tree_2 = train(response ~ .,
                      data = train_data,
                      method = "rf",
                      trControl = train_control_2,
                      tuneGrid = grid2,
                      num.threads = 3,
                      ntree = 500)

```

The confusion matrices are for training and testing sets respectively:

```

pred_tree_train_2 = predict(model_tree_2, newdata = train_data, type = "raw")
pred_tree_test_2 = predict(model_tree_2, newdata = test_data, type = "raw")

```

```
cm_train_2 = confusionMatrix(pred_tree_train_2, data = train_data$response, positive = "Positive_class")
cm_test_2 = confusionMatrix(pred_tree_test_2, data = test_data$response, positive = "Positive_class")
```

```
cm_train_2
```

```
## Confusion Matrix and Statistics
##
##              Reference
## Prediction  Negative_class Positive_class
## Negative_class      560           0
## Positive_class       0          240
##
##              Accuracy : 1
##              95% CI : (0.9954, 1)
##      No Information Rate : 0.7
##      P-Value [Acc > NIR] : < 2.2e-16
##
##              Kappa : 1
##  McNemar's Test P-Value : NA
##
##      Sensitivity : 1.0
##      Specificity : 1.0
##      Pos Pred Value : 1.0
##      Neg Pred Value : 1.0
##      Prevalence : 0.3
##      Detection Rate : 0.3
##      Detection Prevalence : 0.3
##      Balanced Accuracy : 1.0
##
##      'Positive' Class : Positive_class
##
```

```
cm_test_2
```

```
## Confusion Matrix and Statistics
##
##              Reference
## Prediction  Negative_class Positive_class
## Negative_class      129          11
## Positive_class       35          25
##
##              Accuracy : 0.77
##              95% CI : (0.7054, 0.8264)
##      No Information Rate : 0.82
##      P-Value [Acc > NIR] : 0.970362
##
##              Kappa : 0.3817
##  McNemar's Test P-Value : 0.000696
##
##      Sensitivity : 0.6944
##      Specificity : 0.7866
##      Pos Pred Value : 0.4167
```



```
##          Neg Pred Value : 0.9214
##          Prevalence : 0.1800
##          Detection Rate : 0.1250
##          Detection Prevalence : 0.3000
##          Balanced Accuracy : 0.7405
##
##          'Positive' Class : Positive_class
##
```

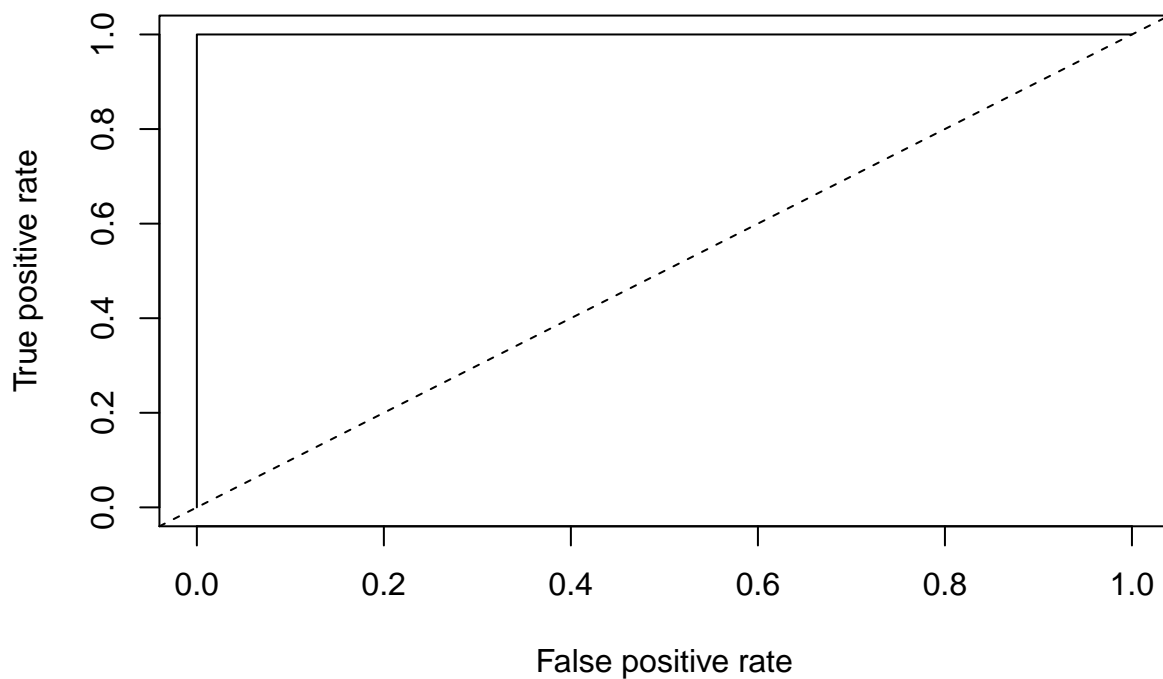
And the ROC curves for training and testing sets respectively:

```
pred_tree_train_prob_2 = predict(model_tree_2, newdata = train_data, type = "prob")

prediction_obj_train_2 = prediction(pred_tree_train_prob_2[,2], train_data$response)

ROC_train_2 = performance(prediction_obj_train_2, "tpr", "fpr")
plot(ROC_train_2, main = "ROC curve for Train dataset")
abline(0, 1, lty = 2)
```

ROC curve for Train dataset

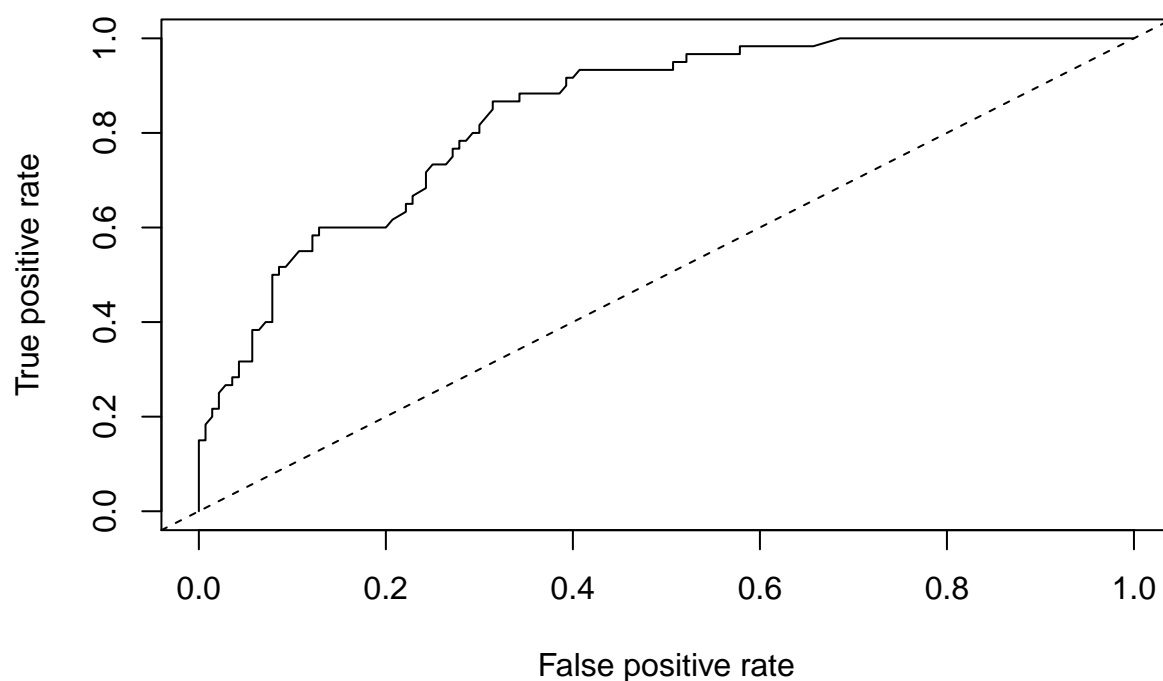


```
pred_tree_test_prob_2 = predict(model_tree_2, newdata = test_data, type = "prob")

prediction_obj_test_2 = prediction(pred_tree_test_prob_2[,2], test_data$response)

ROC_test_2 = performance(prediction_obj_test_2, "tpr", "fpr")
plot(ROC_test_2, main = "ROC curve for Test dataset")
abline(0, 1, lty = 2)
```

ROC curve for Test dataset



3. Apply boosting (method = “adaboost”) of decision trees. Show the confusion matrices and the ROC curves for training and test sets (score = 30).

```
set.seed(1)
train_control_3 = trainControl(method = "cv",
                               number = 5,
                               verbose = T)

model_tree_3 = train(response ~ .,
                      data = train_data,
                      method = "adaboost",
                      trControl = train_control_3,
                      tuneLength = 10)
```

```
## + Fold1: nIter= 50, method=Adaboost.M1
## - Fold1: nIter= 50, method=Adaboost.M1
## + Fold1: nIter=100, method=Adaboost.M1
## - Fold1: nIter=100, method=Adaboost.M1
## + Fold1: nIter=150, method=Adaboost.M1
## - Fold1: nIter=150, method=Adaboost.M1
## + Fold1: nIter=200, method=Adaboost.M1
## - Fold1: nIter=200, method=Adaboost.M1
## + Fold1: nIter=250, method=Adaboost.M1
## - Fold1: nIter=250, method=Adaboost.M1
```

```

## + Fold1: nIter=300, method=Adaboost.M1
## - Fold1: nIter=300, method=Adaboost.M1
## + Fold1: nIter=350, method=Adaboost.M1
## - Fold1: nIter=350, method=Adaboost.M1
## + Fold1: nIter=400, method=Adaboost.M1
## - Fold1: nIter=400, method=Adaboost.M1
## + Fold1: nIter=450, method=Adaboost.M1
## - Fold1: nIter=450, method=Adaboost.M1
## + Fold1: nIter=500, method=Adaboost.M1
## - Fold1: nIter=500, method=Adaboost.M1
## + Fold1: nIter= 50, method=Real adaboost
## - Fold1: nIter= 50, method=Real adaboost
## + Fold1: nIter=100, method=Real adaboost
## - Fold1: nIter=100, method=Real adaboost
## + Fold1: nIter=150, method=Real adaboost
## - Fold1: nIter=150, method=Real adaboost
## + Fold1: nIter=200, method=Real adaboost
## - Fold1: nIter=200, method=Real adaboost
## + Fold1: nIter=250, method=Real adaboost
## - Fold1: nIter=250, method=Real adaboost
## + Fold1: nIter=300, method=Real adaboost
## - Fold1: nIter=300, method=Real adaboost
## + Fold1: nIter=350, method=Real adaboost
## - Fold1: nIter=350, method=Real adaboost
## + Fold1: nIter=400, method=Real adaboost
## - Fold1: nIter=400, method=Real adaboost
## + Fold1: nIter=450, method=Real adaboost
## - Fold1: nIter=450, method=Real adaboost
## + Fold1: nIter=500, method=Real adaboost
## - Fold1: nIter=500, method=Real adaboost
## + Fold2: nIter= 50, method=Adaboost.M1
## - Fold2: nIter= 50, method=Adaboost.M1
## + Fold2: nIter=100, method=Adaboost.M1
## - Fold2: nIter=100, method=Adaboost.M1
## + Fold2: nIter=150, method=Adaboost.M1
## - Fold2: nIter=150, method=Adaboost.M1
## + Fold2: nIter=200, method=Adaboost.M1
## - Fold2: nIter=200, method=Adaboost.M1
## + Fold2: nIter=250, method=Adaboost.M1
## - Fold2: nIter=250, method=Adaboost.M1
## + Fold2: nIter=300, method=Adaboost.M1
## - Fold2: nIter=300, method=Adaboost.M1
## + Fold2: nIter=350, method=Adaboost.M1
## - Fold2: nIter=350, method=Adaboost.M1
## + Fold2: nIter=400, method=Adaboost.M1
## - Fold2: nIter=400, method=Adaboost.M1
## + Fold2: nIter=450, method=Adaboost.M1
## - Fold2: nIter=450, method=Adaboost.M1
## + Fold2: nIter=500, method=Adaboost.M1
## - Fold2: nIter=500, method=Adaboost.M1
## + Fold2: nIter= 50, method=Real adaboost
## - Fold2: nIter= 50, method=Real adaboost
## + Fold2: nIter=100, method=Real adaboost
## - Fold2: nIter=100, method=Real adaboost

```

```

## + Fold2: nIter=150, method=Real adaboost
## - Fold2: nIter=150, method=Real adaboost
## + Fold2: nIter=200, method=Real adaboost
## - Fold2: nIter=200, method=Real adaboost
## + Fold2: nIter=250, method=Real adaboost
## - Fold2: nIter=250, method=Real adaboost
## + Fold2: nIter=300, method=Real adaboost
## - Fold2: nIter=300, method=Real adaboost
## + Fold2: nIter=350, method=Real adaboost
## - Fold2: nIter=350, method=Real adaboost
## + Fold2: nIter=400, method=Real adaboost
## - Fold2: nIter=400, method=Real adaboost
## + Fold2: nIter=450, method=Real adaboost
## - Fold2: nIter=450, method=Real adaboost
## + Fold2: nIter=500, method=Real adaboost
## - Fold2: nIter=500, method=Real adaboost
## + Fold3: nIter= 50, method=Adaboost.M1
## - Fold3: nIter= 50, method=Adaboost.M1
## + Fold3: nIter=100, method=Adaboost.M1
## - Fold3: nIter=100, method=Adaboost.M1
## + Fold3: nIter=150, method=Adaboost.M1
## - Fold3: nIter=150, method=Adaboost.M1
## + Fold3: nIter=200, method=Adaboost.M1
## - Fold3: nIter=200, method=Adaboost.M1
## + Fold3: nIter=250, method=Adaboost.M1
## - Fold3: nIter=250, method=Adaboost.M1
## + Fold3: nIter=300, method=Adaboost.M1
## - Fold3: nIter=300, method=Adaboost.M1
## + Fold3: nIter=350, method=Adaboost.M1
## - Fold3: nIter=350, method=Adaboost.M1
## + Fold3: nIter=400, method=Adaboost.M1
## - Fold3: nIter=400, method=Adaboost.M1
## + Fold3: nIter=450, method=Adaboost.M1
## - Fold3: nIter=450, method=Adaboost.M1
## + Fold3: nIter=500, method=Adaboost.M1
## - Fold3: nIter=500, method=Adaboost.M1
## + Fold3: nIter= 50, method=Real adaboost
## - Fold3: nIter= 50, method=Real adaboost
## + Fold3: nIter=100, method=Real adaboost
## - Fold3: nIter=100, method=Real adaboost
## + Fold3: nIter=150, method=Real adaboost
## - Fold3: nIter=150, method=Real adaboost
## + Fold3: nIter=200, method=Real adaboost
## - Fold3: nIter=200, method=Real adaboost
## + Fold3: nIter=250, method=Real adaboost
## - Fold3: nIter=250, method=Real adaboost
## + Fold3: nIter=300, method=Real adaboost
## - Fold3: nIter=300, method=Real adaboost
## + Fold3: nIter=350, method=Real adaboost
## - Fold3: nIter=350, method=Real adaboost
## + Fold3: nIter=400, method=Real adaboost
## - Fold3: nIter=400, method=Real adaboost
## + Fold3: nIter=450, method=Real adaboost
## - Fold3: nIter=450, method=Real adaboost

```

```

## + Fold3: nIter=500, method=Real adaboost
## - Fold3: nIter=500, method=Real adaboost
## + Fold4: nIter= 50, method=Adaboost.M1
## - Fold4: nIter= 50, method=Adaboost.M1
## + Fold4: nIter=100, method=Adaboost.M1
## - Fold4: nIter=100, method=Adaboost.M1
## + Fold4: nIter=150, method=Adaboost.M1
## - Fold4: nIter=150, method=Adaboost.M1
## + Fold4: nIter=200, method=Adaboost.M1
## - Fold4: nIter=200, method=Adaboost.M1
## + Fold4: nIter=250, method=Adaboost.M1
## - Fold4: nIter=250, method=Adaboost.M1
## + Fold4: nIter=300, method=Adaboost.M1
## - Fold4: nIter=300, method=Adaboost.M1
## + Fold4: nIter=350, method=Adaboost.M1
## - Fold4: nIter=350, method=Adaboost.M1
## + Fold4: nIter=400, method=Adaboost.M1
## - Fold4: nIter=400, method=Adaboost.M1
## + Fold4: nIter=450, method=Adaboost.M1
## - Fold4: nIter=450, method=Adaboost.M1
## + Fold4: nIter=500, method=Adaboost.M1
## - Fold4: nIter=500, method=Adaboost.M1
## + Fold4: nIter= 50, method=Real adaboost
## - Fold4: nIter= 50, method=Real adaboost
## + Fold4: nIter=100, method=Real adaboost
## - Fold4: nIter=100, method=Real adaboost
## + Fold4: nIter=150, method=Real adaboost
## - Fold4: nIter=150, method=Real adaboost
## + Fold4: nIter=200, method=Real adaboost
## - Fold4: nIter=200, method=Real adaboost
## + Fold4: nIter=250, method=Real adaboost
## - Fold4: nIter=250, method=Real adaboost
## + Fold4: nIter=300, method=Real adaboost
## - Fold4: nIter=300, method=Real adaboost
## + Fold4: nIter=350, method=Real adaboost
## - Fold4: nIter=350, method=Real adaboost
## + Fold4: nIter=400, method=Real adaboost
## - Fold4: nIter=400, method=Real adaboost
## + Fold4: nIter=450, method=Real adaboost
## - Fold4: nIter=450, method=Real adaboost
## + Fold4: nIter=500, method=Real adaboost
## - Fold4: nIter=500, method=Real adaboost
## + Fold5: nIter= 50, method=Adaboost.M1
## - Fold5: nIter= 50, method=Adaboost.M1
## + Fold5: nIter=100, method=Adaboost.M1
## - Fold5: nIter=100, method=Adaboost.M1
## + Fold5: nIter=150, method=Adaboost.M1
## - Fold5: nIter=150, method=Adaboost.M1
## + Fold5: nIter=200, method=Adaboost.M1
## - Fold5: nIter=200, method=Adaboost.M1
## + Fold5: nIter=250, method=Adaboost.M1
## - Fold5: nIter=250, method=Adaboost.M1
## + Fold5: nIter=300, method=Adaboost.M1
## - Fold5: nIter=300, method=Adaboost.M1

```

```

## + Fold5: nIter=350, method=Adaboost.M1
## - Fold5: nIter=350, method=Adaboost.M1
## + Fold5: nIter=400, method=Adaboost.M1
## - Fold5: nIter=400, method=Adaboost.M1
## + Fold5: nIter=450, method=Adaboost.M1
## - Fold5: nIter=450, method=Adaboost.M1
## + Fold5: nIter=500, method=Adaboost.M1
## - Fold5: nIter=500, method=Adaboost.M1
## + Fold5: nIter= 50, method=Real adaboost
## - Fold5: nIter= 50, method=Real adaboost
## + Fold5: nIter=100, method=Real adaboost
## - Fold5: nIter=100, method=Real adaboost
## + Fold5: nIter=150, method=Real adaboost
## - Fold5: nIter=150, method=Real adaboost
## + Fold5: nIter=200, method=Real adaboost
## - Fold5: nIter=200, method=Real adaboost
## + Fold5: nIter=250, method=Real adaboost
## - Fold5: nIter=250, method=Real adaboost
## + Fold5: nIter=300, method=Real adaboost
## - Fold5: nIter=300, method=Real adaboost
## + Fold5: nIter=350, method=Real adaboost
## - Fold5: nIter=350, method=Real adaboost
## + Fold5: nIter=400, method=Real adaboost
## - Fold5: nIter=400, method=Real adaboost
## + Fold5: nIter=450, method=Real adaboost
## - Fold5: nIter=450, method=Real adaboost
## + Fold5: nIter=500, method=Real adaboost
## - Fold5: nIter=500, method=Real adaboost
## Aggregating results
## Selecting tuning parameters
## Fitting nIter = 500, method = Real adaboost on full training set

```

The Confusion matrices for Training and Test sets respectively:

```

pred_tree_train_3 = predict(model_tree_3, newdata = train_data, type = "raw")
cm_train_3 = confusionMatrix(pred_tree_train_3, data = train_data$response, positive = "Positive_class")

cm_train_3

```

```

## Confusion Matrix and Statistics
##
##              Reference
## Prediction      Negative_class Positive_class
##   Negative_class           560             0
##   Positive_class            0           240
##
##              Accuracy : 1
##              95% CI : (0.9954, 1)
##   No Information Rate : 0.7
##   P-Value [Acc > NIR] : < 2.2e-16
##
##              Kappa : 1
##   Mcnemar's Test P-Value : NA

```

```
##
##          Sensitivity : 1.0
##          Specificity : 1.0
##          Pos Pred Value : 1.0
##          Neg Pred Value : 1.0
##          Prevalence : 0.3
##          Detection Rate : 0.3
##          Detection Prevalence : 0.3
##          Balanced Accuracy : 1.0
##
##          'Positive' Class : Positive_class
##

pred_tree_test_3 = predict(model_tree_3, newdata = test_data, type = "raw")
cm_test_3 = confusionMatrix(pred_tree_test_3, data = test_data$response, positive = "Positive_class")

cm_test_3

## Confusion Matrix and Statistics
##
##              Reference
## Prediction   Negative_class Positive_class
## Negative_class      128           12
## Positive_class       38           22
##
##              Accuracy : 0.75
##              95% CI : (0.684, 0.8084)
##          No Information Rate : 0.83
##          P-Value [Acc > NIR] : 0.998463
##
##              Kappa : 0.3207
## Mcnemar's Test P-Value : 0.000407
##
##              Sensitivity : 0.6471
##              Specificity : 0.7711
##              Pos Pred Value : 0.3667
##              Neg Pred Value : 0.9143
##              Prevalence : 0.1700
##              Detection Rate : 0.1100
##          Detection Prevalence : 0.3000
##              Balanced Accuracy : 0.7091
##
##          'Positive' Class : Positive_class
##
```

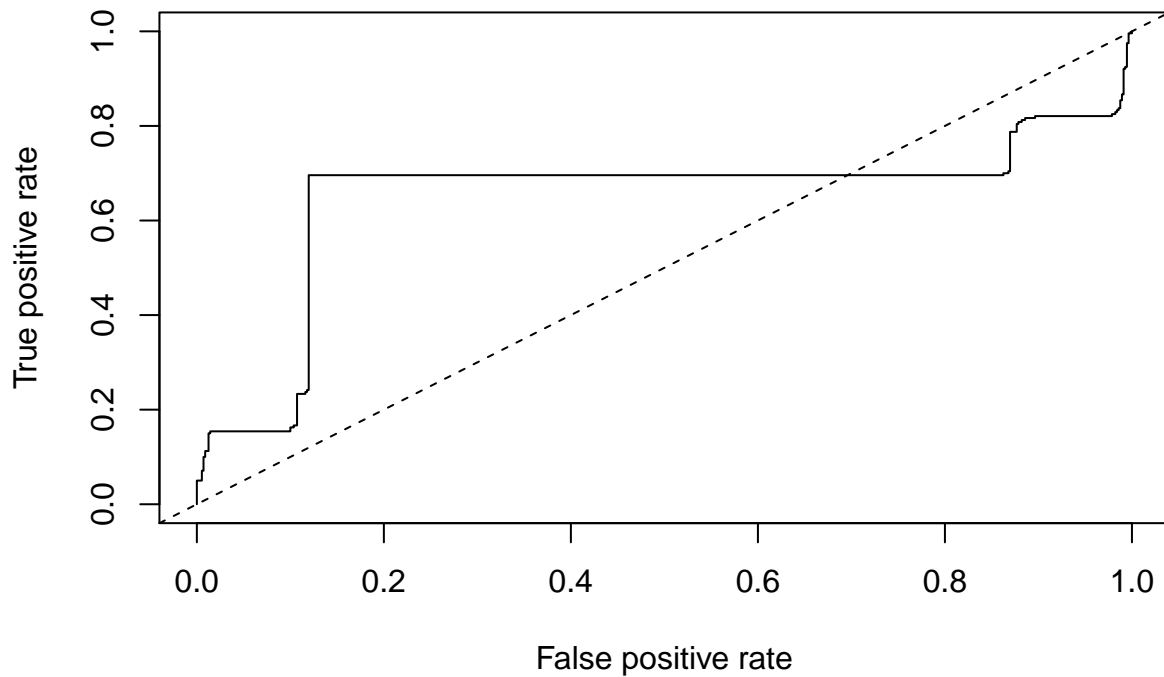
And finally, here are the ROC curves for Training and Test set for Boosting:

```
pred_tree_train_prob_3 = predict(model_tree_3, newdata = train_data, type = "prob")

prediction_obj_train_3 = prediction(pred_tree_train_prob_3[,2], train_data$response)

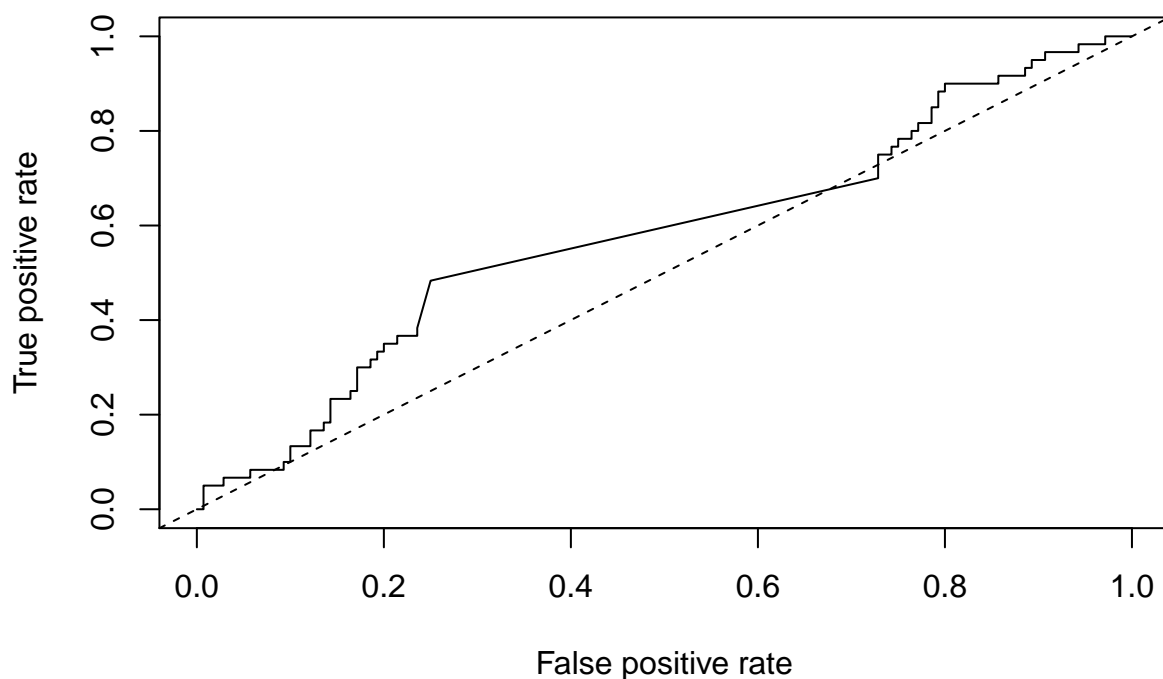
ROC_train_3 = performance(prediction_obj_train_3, "tpr", "fpr")
plot(ROC_train_3, main = "ROC curve for Train dataset")
abline(0, 1, lty = 2)
```

ROC curve for Train dataset



```
pred_tree_test_prob_3 = predict(model_tree_3, newdata = test_data, type = "prob")  
prediction_obj_test_3 = prediction(pred_tree_test_prob_3[,2], test_data$response)  
  
ROC_test_3 = performance(prediction_obj_test_3, "tpr", "fpr")  
plot(ROC_test_3, main = "ROC curve for Test dataset")  
abline(0, 1, lty = 2)
```


ROC curve for Test dataset



4. Compare the precisions on the test set and pick the best model for classification (score = 10).

```
prec_1 <- cm_test_1$byClass["Pos Pred Value"]
prec_2 <- cm_test_2$byClass["Pos Pred Value"]
prec_3 <- cm_test_3$byClass["Pos Pred Value"]
precisions <- c(prec_1, prec_2, prec_3)
names(precisions) <- c("Classification Trees", "Random Forests", "Boosting of Dec. Trees")
precisions
```

##	Classification Trees	Random Forests	Boosting of Dec. Trees
##	0.5333333	0.4166667	0.3666667

So by comparing the precisions, the best model for classification is Classification Trees.