

Machine Learning Project

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```
library(knitr)
library(caret)

## Loading required package: lattice
## Loading required package: ggplot2
library(rpart)
library(rpart.plot)
library(rattle)

## Rattle: A free graphical interface for data science with R.
## Version 5.1.0 Copyright (c) 2006-2017 Togaware Pty Ltd.
## Type 'rattle()' to shake, rattle, and roll your data.
library(randomForest)

## randomForest 4.6-12
## Type rfNews() to see new features/changes/bug fixes.
##
## Attaching package: 'randomForest'
## The following object is masked from 'package:rattle':
##
##     importance
## The following object is masked from 'package:ggplot2':
##
##     margin
set.seed(12345)
UrlTrain <- "http://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv"
UrlTest  <- "http://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv"

# Dataset
training <- read.csv(url(UrlTrain))
testing  <- read.csv(url(UrlTest))

# Training and Testing
inTrain <- createDataPartition(training$classe, p=0.7, list=FALSE)
TrainSet <- training[inTrain, ]
TestSet  <- training[-inTrain, ]
dim(TrainSet)

## [1] 13737  160
dim(TestSet)

## [1] 5885  160
```

```

NZV <- nearZeroVar(TrainSet)
TrainSet <- TrainSet[, -NZV]
TestSet <- TestSet[, -NZV]
dim(TrainSet)

## [1] 13737 106

dim(TestSet)

## [1] 5885 106

AllNA <- sapply(TrainSet, function(x) mean(is.na(x))) > 0.95
TrainSet <- TrainSet[, AllNA==FALSE]
TestSet <- TestSet[, AllNA==FALSE]
dim(TrainSet)

## [1] 13737 59

dim(TestSet)

## [1] 5885 59

TrainSet <- TrainSet[, -(1:5)]
TestSet <- TestSet[, -(1:5)]
dim(TrainSet)

## [1] 13737 54

dim(TestSet)

## [1] 5885 54

# RANDOM FOREST
set.seed(12345)
controlRF <- trainControl(method="cv", number=3, verboseIter=FALSE)
modFitRandForest <- train(classe ~ ., data=TrainSet, method="rf",
                          trControl=controlRF)
modFitRandForest$finalModel

##
## Call:
## randomForest(x = x, y = y, mtry = param$mtry)
##              Type of random forest: classification
##              Number of trees: 500
## No. of variables tried at each split: 27
##
##              OOB estimate of  error rate: 0.19%
## Confusion matrix:
##      A    B    C    D    E  class.error
## A 3904     1     0     0     1 0.0005120328
## B   6 2651     1     0     0 0.0026335591
## C    0     6 2390     0     0 0.0025041736
## D    0     0   8 2244     0 0.0035523979
## E    0     0     0    3 2522 0.0011881188

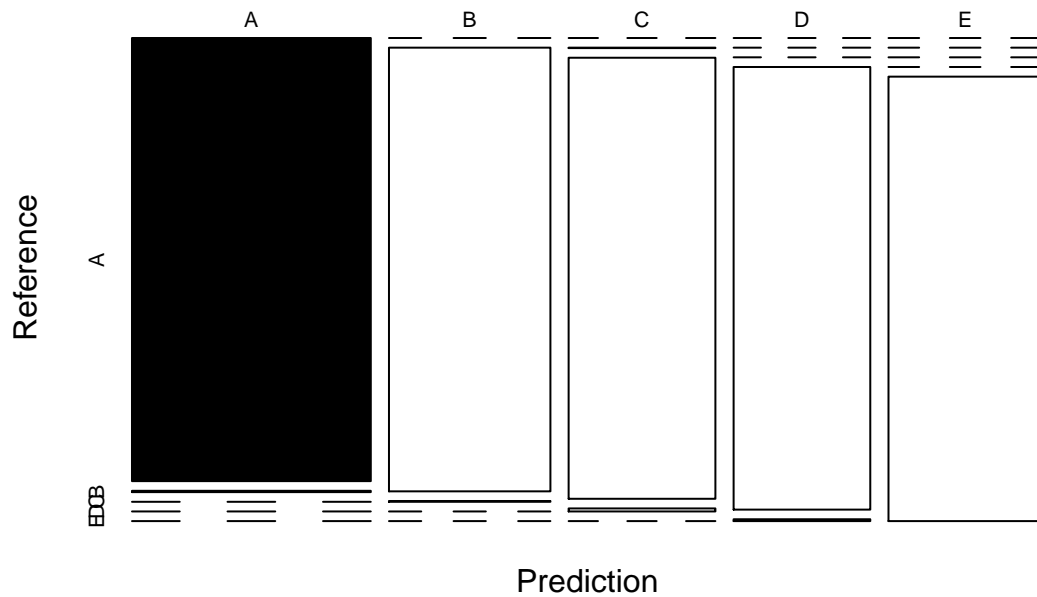
predictRandForest <- predict(modFitRandForest, newdata=TestSet)
confMatRandForest <- confusionMatrix(predictRandForest, TestSet$classe)
confMatRandForest

## Confusion Matrix and Statistics

```

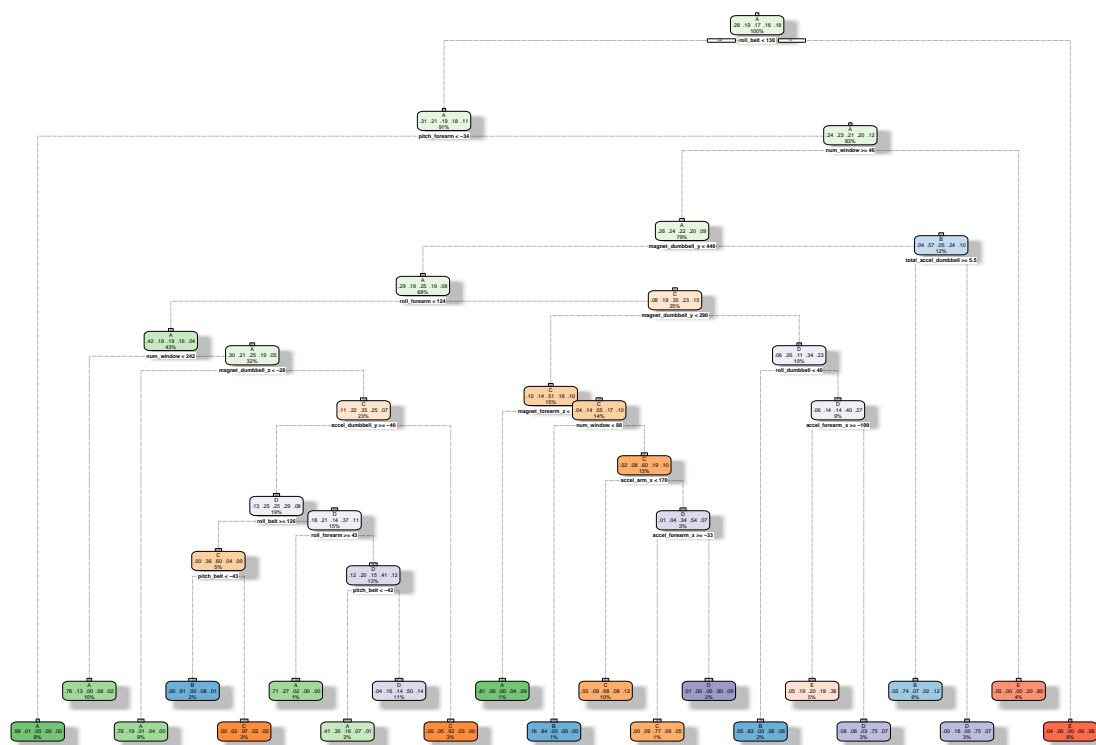
```
##
##           Reference
## Prediction    A    B    C    D    E
##           A 1674    5    0    0    0
##           B    0 1133    2    0    0
##           C    0    1 1024    7    0
##           D    0    0    0  957    4
##           E    0    0    0    0 1078
##
## Overall Statistics
##
##           Accuracy : 0.9968
##           95% CI : (0.995, 0.9981)
##           No Information Rate : 0.2845
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.9959
##           McNemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: A Class: B Class: C Class: D Class: E
## Sensitivity      1.0000  0.9947  0.9981  0.9927  0.9963
## Specificity      0.9988  0.9996  0.9984  0.9992  1.0000
## Pos Pred Value   0.9970  0.9982  0.9922  0.9958  1.0000
## Neg Pred Value    1.0000  0.9987  0.9996  0.9986  0.9992
## Prevalence       0.2845  0.1935  0.1743  0.1638  0.1839
## Detection Rate   0.2845  0.1925  0.1740  0.1626  0.1832
## Detection Prevalence 0.2853  0.1929  0.1754  0.1633  0.1832
## Balanced Accuracy 0.9994  0.9972  0.9982  0.9960  0.9982
plot(confMatRandForest$table, col = confMatRandForest$byClass,
     main = paste("Random Forest - Accuracy =",
                  round(confMatRandForest$overall['Accuracy'], 4)))
```

Random Forest – Accuracy = 0.9968



```
#Decision Tree
set.seed(12345)
modFitDecTree <- rpart(classe ~ ., data=TrainSet, method="class")
fancyRpartPlot(modFitDecTree)
```

```
## Warning: labs do not fit even at cex 0.15, there may be some overplotting
```



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```
predictDecTree <- predict(modFitDecTree, newdata=TestSet, type="class")
confMatDecTree <- confusionMatrix(predictDecTree, TestSet$classe)
confMatDecTree
```

Confusion Matrix and Statistics

##

Reference

## Prediction		A	B	C	D	E
## A	1530	269	51	79	16	
## B	35	575	31	25	68	
## C	17	73	743	68	84	
## D	39	146	130	702	128	
## E	53	76	71	90	786	

##

Overall Statistics

##

Accuracy : 0.7368
 ## 95% CI : (0.7253, 0.748)
 ## No Information Rate : 0.2845
 ## P-Value [Acc > NIR] : < 2.2e-16

##

Kappa : 0.6656
 ## McNemar's Test P-Value : < 2.2e-16

##

Statistics by Class:

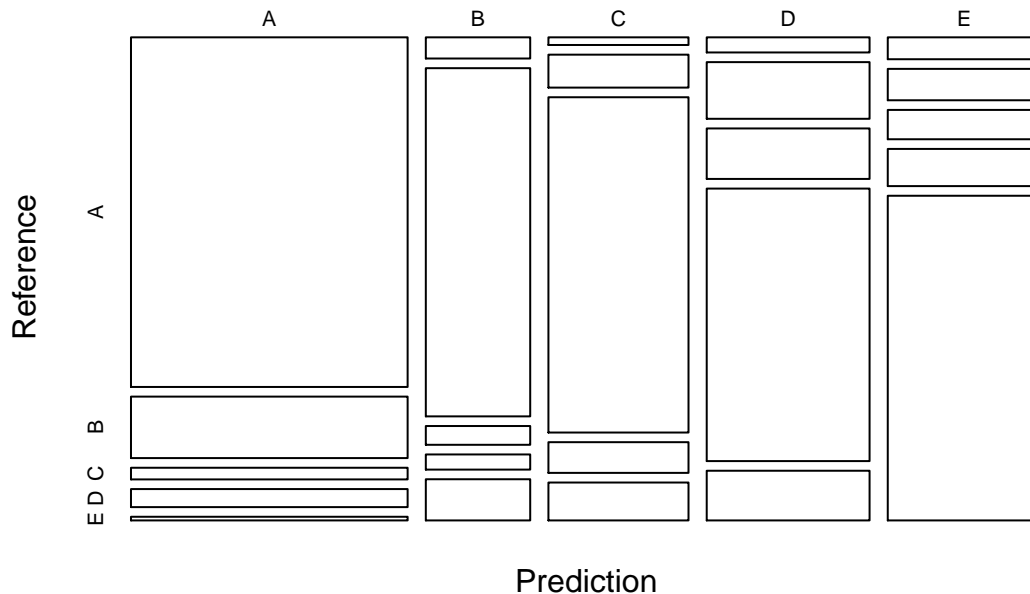
##

Class: A Class: B Class: C Class: D Class: E

```
## Sensitivity      0.9140  0.50483  0.7242  0.7282  0.7264
## Specificity     0.9014  0.96650  0.9502  0.9100  0.9396
## Pos Pred Value  0.7866  0.78338  0.7543  0.6131  0.7305
## Neg Pred Value  0.9635  0.89051  0.9422  0.9447  0.9384
## Prevalence      0.2845  0.19354  0.1743  0.1638  0.1839
## Detection Rate  0.2600  0.09771  0.1263  0.1193  0.1336
## Detection Prevalence 0.3305 0.12472 0.1674 0.1946 0.1828
## Balanced Accuracy 0.9077 0.73566 0.8372 0.8191 0.8330
```

```
plot(confMatDecTree$table, col = confMatDecTree$byClass,
     main = paste("Decision Tree - Accuracy =",
                  round(confMatDecTree$overall['Accuracy'], 4)))
```

Decision Tree – Accuracy = 0.7368



```
#GBM
set.seed(12345)
controlGBM <- trainControl(method = "repeatedcv", number = 5, repeats = 1)
modFitGBM <- train(classe ~ ., data=TrainSet, method = "gbm",
                  trControl = controlGBM, verbose = FALSE)
```

```
## Loading required package: survival
##
## Attaching package: 'survival'
## The following object is masked from 'package:caret':
##
##   cluster
## Loading required package: splines
```

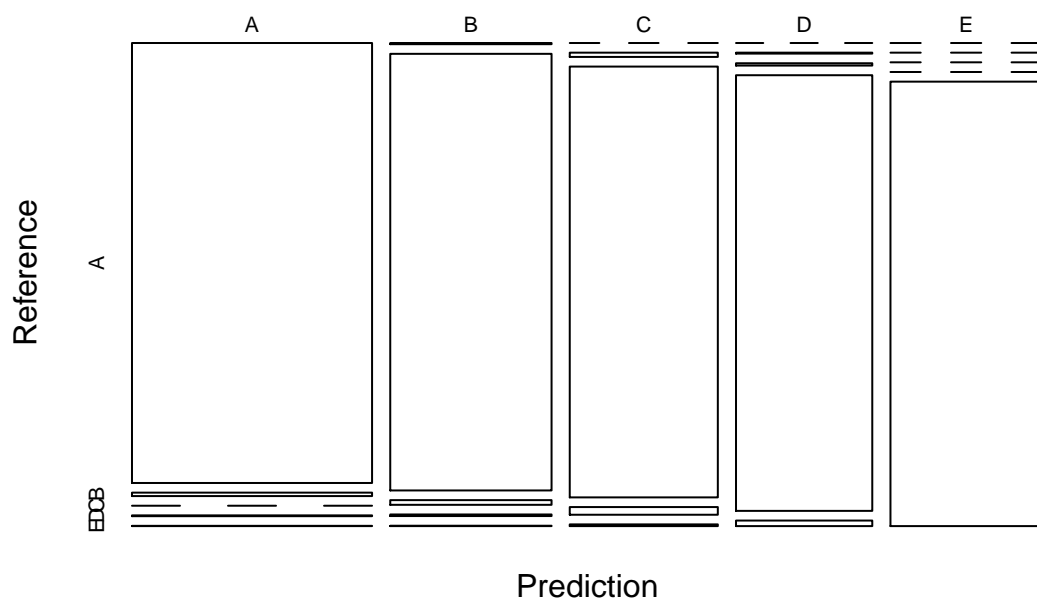
```
## Loading required package: parallel
## Loaded gbm 2.1.3
modFitGBM$finalModel

## A gradient boosted model with multinomial loss function.
## 150 iterations were performed.
## There were 53 predictors of which 43 had non-zero influence.
predictGBM <- predict(modFitGBM, newdata=TestSet)
confMatGBM <- confusionMatrix(predictGBM, TestSet$classe)
confMatGBM

## Confusion Matrix and Statistics
##
##           Reference
## Prediction    A    B    C    D    E
##           A 1671   13    0    3    1
##           B    3 1114   12    4    1
##           C    0   10 1009   18    4
##           D    0    2    5  939   12
##           E    0    0    0    0 1064
##
## Overall Statistics
##
##           Accuracy : 0.985
##           95% CI : (0.9816, 0.988)
##           No Information Rate : 0.2845
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.9811
##           McNemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: A Class: B Class: C Class: D Class: E
## Sensitivity      0.9982  0.9781  0.9834  0.9741  0.9834
## Specificity      0.9960  0.9958  0.9934  0.9961  1.0000
## Pos Pred Value   0.9899  0.9824  0.9693  0.9802  1.0000
## Neg Pred Value   0.9993  0.9947  0.9965  0.9949  0.9963
## Prevalence       0.2845  0.1935  0.1743  0.1638  0.1839
## Detection Rate   0.2839  0.1893  0.1715  0.1596  0.1808
## Detection Prevalence 0.2868  0.1927  0.1769  0.1628  0.1808
## Balanced Accuracy 0.9971  0.9869  0.9884  0.9851  0.9917

plot(confMatGBM$table, col = confMatGBM$byClass,
     main = paste("GBM - Accuracy =", round(confMatGBM$overall['Accuracy'], 4)))
```

GBM – Accuracy = 0.985



```
predictTEST <- predict(modFitRandForest, newdata=testing)
predictTEST
```

```
## [1] B A B A A E D B A A B C B A E E A B B B
## Levels: A B C D E
```