# Predict exercise manners based on data from the quantified self context

'Practical Machine Learning' course project of Hendrik L., September 2015

## Download and read the training and test data

```
# download and prepare the subTrainData data
if (!file.exists("pmlTraining.csv")) {
    fileURL <- "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-subTrainData.csv"
    download.file(fileURL, "pmlTraining.csv", mode = "wb")
}
trainData <- read.csv("pmlTraining.csv",
    na.strings = c("NA", "NAs", "NULL", "#DIV/0!"," ", ""))

# download and prepare the test data
if (!file.exists("pmlTest.csv")) {
    fileURL <- "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-subTestData.csv"
    download.file(fileURL, "pmlTest.csv", mode = "wb")
}
testData <- read.csv("pmlTest.csv",
    na.strings = c("NA", "NAs", "NULL", "#DIV/0!"," ", ""))</pre>
```

# Drop columns with NA values and unnecessary data

## Remove zero and near-zero variance predictors

```
# zero variance predictors: they have one unique value across samples
# near-zero variance predictors: they have both few unique values relative to the
# number of samples and a large ratio of the frequency of the most common value
# to the frequency of the second most common value
nearZV <- nearZeroVar(trainData[sapply(trainData, is.numeric)], saveMetrics = TRUE)
trainData <- trainData[, nearZV[, 'nzv'] == 0]</pre>
```

```
corMatrix <- cor(na.omit(trainData[sapply(trainData, is.numeric)]))
# correlationmatrixdegreesoffreedom <- expand.grid(row = 1:52, col = 1:52)
# correlationmatrixdegreesoffreedom$correlation <- as.vector(corMatrix)
corIndex <- findCorrelation(corMatrix, cutoff = 0.7, verbose = TRUE)</pre>
```

```
## Compare row 10 and column 1 with corr 0.992
    Means: 0.27 vs 0.168 so flagging column 10
##
## Compare row 1 and column 9 with corr 0.925
    Means: 0.25 vs 0.164 so flagging column 1
##
## Compare row 9 and column 22 with corr 0.722
    Means: 0.233 vs 0.161 so flagging column 9
##
## Compare row 22 and column 4 with corr 0.759
##
    Means: 0.224 vs 0.158 so flagging column 22
## Compare row 4 and column 3 with corr 0.762
##
    Means: 0.2 vs 0.155 so flagging column 4
## Compare row 3 and column 8 with corr 0.708
##
    Means: 0.2 vs 0.153 so flagging column 3
## Compare row 36 and column 29 with corr 0.849
    Means: 0.257 vs 0.151 so flagging column 36
##
## Compare row 8 and column 2 with corr 0.966
##
    Means: 0.229 vs 0.146 so flagging column 8
## Compare row 2 and column 11 with corr 0.884
##
    Means: 0.212 vs 0.143 so flagging column 2
## Compare row 37 and column 38 with corr 0.769
    Means: 0.198 vs 0.139 so flagging column 37
## Compare row 35 and column 30 with corr 0.773
    Means: 0.195 vs 0.137 so flagging column 35
## Compare row 38 and column 5 with corr 0.781
    Means: 0.177 vs 0.134 so flagging column 38
## Compare row 21 and column 24 with corr 0.814
    Means: 0.176 vs 0.133 so flagging column 21
##
## Compare row 34 and column 28 with corr 0.808
    Means: 0.176 vs 0.13 so flagging column 34
## Compare row 23 and column 26 with corr 0.779
##
    Means: 0.137 vs 0.129 so flagging column 23
## Compare row 25 and column 24 with corr 0.792
##
    Means: 0.145 vs 0.128 so flagging column 25
## Compare row 12 and column 13 with corr 0.779
##
    Means: 0.122 vs 0.127 so flagging column 13
  Compare row 48 and column 51 with corr 0.772
    Means: 0.145 vs 0.127 so flagging column 48
##
## Compare row 19 and column 18 with corr 0.918
    Means: 0.095 vs 0.127 so flagging column 18
##
## Compare row 46 and column 45 with corr 0.846
##
    Means: 0.131 vs 0.129 so flagging column 46
## Compare row 45 and column 31 with corr 0.71
    Means: 0.098 vs 0.129 so flagging column 31
##
## Compare row 45 and column 33 with corr 0.716
    Means: 0.078 vs 0.132 so flagging column 33
## All correlations <= 0.7
```

```
trainData <- trainData[, -corIndex]</pre>
```

Remove blank columns from training and test data

```
for (i in c(8:ncol(trainData) - 1)) {
    trainData[,i] <- as.numeric(as.character(trainData[,i]))
}
for (i in c(8:ncol(testData) - 1)) {
    testData[,i] <- as.numeric(as.character(testData[,i]))
}</pre>
```

Create the final feature set and model data

```
featureset <- colnames(trainData[colSums(is.na(trainData)) == 0])[-(1:7)]
modeldata <- trainData[featureset]</pre>
```

Subset the training data (60% for training, 40% for testing)

```
trainIndex <- createDataPartition(modeldata$classe, p = 0.6, list = FALSE )
subTrainData <- modeldata[trainIndex,]
subTestData <- modeldata[-trainIndex,]</pre>
```

Fit a random forest model with 5-fold cross validation

Evaluate the model performance

```
predict <- predict(model, subTestData)
confusionMatrix(subTestData$classe, predict)</pre>
```

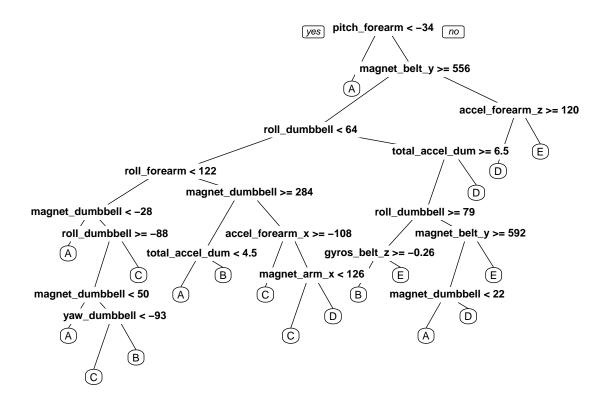
```
## Confusion Matrix and Statistics
##
            Reference
##
              Α
                                   Ε
## Prediction
                     В
                          С
                               D
##
           A 2223
                     4
                          1
                               2
##
           В
               34 1468
                          9
                               6
                                    1
##
           С
                1
                    23 1329
                              11
                2
##
           D
                     0
                         52 1222
                                   10
##
           Ε
                3
                     9
                          2
                              11 1417
##
## Overall Statistics
##
                 Accuracy : 0.9762
##
##
                   95% CI: (0.9725, 0.9794)
##
      No Information Rate: 0.2884
      P-Value [Acc > NIR] : < 2.2e-16
##
##
##
                    Kappa: 0.9698
  Mcnemar's Test P-Value : 4.833e-11
##
##
## Statistics by Class:
##
                       Class: A Class: B Class: C Class: D Class: E
##
## Sensitivity
                         0.9823 0.9761 0.9541
                                                   0.9760
                                                            0.9881
## Specificity
                         0.9984 0.9921 0.9940 0.9903
                                                           0.9961
## Pos Pred Value
                         0.9960 0.9671 0.9715 0.9502
                                                           0.9827
## Neg Pred Value
                         0.9929 0.9943
                                         0.9901
                                                   0.9954
                                                            0.9973
## Prevalence
                         0.2884 0.1917
                                          0.1775
                                                   0.1596
                                                            0.1828
## Detection Rate
                                                  0.1557
                         0.2833 0.1871
                                          0.1694
                                                            0.1806
## Detection Prevalence
                         0.2845 0.1935
                                          0.1744
                                                   0.1639
                                                            0.1838
                         0.9904 0.9841
                                          0.9740
## Balanced Accuracy
                                                   0.9832
                                                            0.9921
accuracy <- postResample(predict, subTestData$classe)</pre>
```

## **Estimated results:**

The estimated model accuracy is 97.62% and the estimated out of sample error is 2.38%.

## Show the tree model

```
treeModel <- rpart(classe ~ ., data = trainData, method = "class")
prp(treeModel)</pre>
```



## Part 2 of the course project:

Predict the results for the test data and write them to files for submission

```
testData <- testData[featureset[featureset != 'classe']]
predictedResults <- predict(model, newdata = testData)
for (i in 1:length(predictedResults)) {
    filename = paste0("problem_id_", i, ".txt")
    write.table(predictedResults[i], file = filename, quote = FALSE,
        row.names = FALSE, col.names = FALSE)
}</pre>
```