Machine Learning Project

Thursday, June 18, 2015

## Introduction

Using devices such as Jawbone Up, Nike FuelBand, and Fitbit it is now possible to collect a large amount of data about personal activity relatively inexpensively. These type of devices are part of the quantified self movement - a group of enthusiasts who take measurements about themselves regularly to improve their health, to find patterns in their behavior, or because they are tech geeks. One thing that people regularly do is quantify how much of a particular activity they do, but they rarely quantify how well they do it.

The goal of project is to predict the manner in which device users did the exercise.

## Reading, Cleaning and Preprocessing Data

# Load required libraries  
library(caret)

## Loading required package: lattice  
## Loading required package: ggplot2

library(randomForest)

## Warning: package 'randomForest' was built under R version 3.2.1

## randomForest 4.6-10  
## Type rfNews() to see new features/changes/bug fixes.

library(e1071)

## Warning: package 'e1071' was built under R version 3.2.1

set.seed(9999)  
  
# Training Data - Identifying "NA", "" and "#DIV/0!" as NA strings while reading the data  
training <- read.csv("pml-training.csv",na.strings=c("NA","","#DIV/0!"))  
  
# Testing Data - Identifying "NA", "" and "#DIV/0!" as NA strings while reading the data  
testing <- read.csv("pml-testing.csv",na.strings=c("NA","","#DIV/0!"))  
  
# Delete columns with all missing values  
training <-training[,colSums(is.na(training))== 0]  
testing <-testing[,colSums(is.na(testing)) == 0]  
  
# Some variables are not required for the current project - user\_name, raw\_timestamp\_part\_1, raw\_timestamp\_part\_,2 cvtd\_timestamp, new\_window, and num\_window (columns 1 to 7) so deleting these variables.  
  
training <- training[,-c(1:7)]  
testing <- testing[,-c(1:7)]  
  
# Partitioning the original training set into a training set and a validation set to validate the model. Splitting on the 'classe' variable (variable of interest) with a 70-30 split  
  
trainSet <- createDataPartition(y = training$classe, p = 0.7, list = FALSE)  
trainData <- training[trainSet,]  
validData <- training[-trainSet,]

## Training the Model

We will apply 3 different learning methods Classification tree, Gradient boosting (gbm) and Random forest models.

# Classification Tree Model  
  
CTmodel <- train(classe ~ ., method="rpart", data = trainData)

## Loading required package: rpart

CTpredict <- predict(CTmodel, validData)  
  
confusionMatrix(CTpredict, validData$classe)

## Confusion Matrix and Statistics  
##   
## Reference  
## Prediction A B C D E  
## A 1008 176 27 55 15  
## B 4 198 20 3 7  
## C 509 498 609 301 404  
## D 148 267 370 605 150  
## E 5 0 0 0 506  
##   
## Overall Statistics  
##   
## Accuracy : 0.4972   
## 95% CI : (0.4843, 0.5101)  
## No Information Rate : 0.2845   
## P-Value [Acc > NIR] : < 2.2e-16   
##   
## Kappa : 0.3737   
## Mcnemar's Test P-Value : < 2.2e-16   
##   
## Statistics by Class:  
##   
## Class: A Class: B Class: C Class: D Class: E  
## Sensitivity 0.6022 0.17384 0.5936 0.6276 0.46765  
## Specificity 0.9352 0.99284 0.6477 0.8100 0.99896  
## Pos Pred Value 0.7869 0.85345 0.2624 0.3929 0.99022  
## Neg Pred Value 0.8553 0.83354 0.8830 0.9174 0.89282  
## Prevalence 0.2845 0.19354 0.1743 0.1638 0.18386  
## Detection Rate 0.1713 0.03364 0.1035 0.1028 0.08598  
## Detection Prevalence 0.2177 0.03942 0.3944 0.2617 0.08683  
## Balanced Accuracy 0.7687 0.58334 0.6206 0.7188 0.73331

# Gradient Boosting (GBM)  
  
GBmodel <- train(classe ~ ., method="gbm", data = trainData, verbose=F)

## Loading required package: gbm

## Warning: package 'gbm' was built under R version 3.2.1

## 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 2.1.1  
## Loading required package: plyr

GBpredict <- predict(GBmodel, validData)  
  
confusionMatrix(GBpredict, validData$classe)

## Confusion Matrix and Statistics  
##   
## Reference  
## Prediction A B C D E  
## A 1647 45 0 1 3  
## B 18 1063 31 1 13  
## C 3 27 981 37 10  
## D 4 3 11 918 18  
## E 2 1 3 7 1038  
##   
## Overall Statistics  
##   
## Accuracy : 0.9596   
## 95% CI : (0.9542, 0.9644)  
## No Information Rate : 0.2845   
## P-Value [Acc > NIR] : < 2.2e-16   
##   
## Kappa : 0.9488   
## Mcnemar's Test P-Value : 1.881e-07   
##   
## Statistics by Class:  
##   
## Class: A Class: B Class: C Class: D Class: E  
## Sensitivity 0.9839 0.9333 0.9561 0.9523 0.9593  
## Specificity 0.9884 0.9867 0.9842 0.9927 0.9973  
## Pos Pred Value 0.9711 0.9440 0.9272 0.9623 0.9876  
## Neg Pred Value 0.9936 0.9840 0.9907 0.9907 0.9909  
## Prevalence 0.2845 0.1935 0.1743 0.1638 0.1839  
## Detection Rate 0.2799 0.1806 0.1667 0.1560 0.1764  
## Detection Prevalence 0.2882 0.1913 0.1798 0.1621 0.1786  
## Balanced Accuracy 0.9861 0.9600 0.9701 0.9725 0.9783

# Random Forest Model  
RFmodel <- randomForest(classe ~. , method="class", data=trainData)  
  
RFpredict <- predict(RFmodel, validData, type = "class")  
  
confusionMatrix(RFpredict, validData$classe)

## Confusion Matrix and Statistics  
##   
## Reference  
## Prediction A B C D E  
## A 1674 8 0 0 0  
## B 0 1128 10 0 0  
## C 0 3 1013 11 2  
## D 0 0 3 951 0  
## E 0 0 0 2 1080  
##   
## Overall Statistics  
##   
## Accuracy : 0.9934   
## 95% CI : (0.991, 0.9953)  
## No Information Rate : 0.2845   
## P-Value [Acc > NIR] : < 2.2e-16   
##   
## Kappa : 0.9916   
## Mcnemar's Test P-Value : NA   
##   
## Statistics by Class:  
##   
## Class: A Class: B Class: C Class: D Class: E  
## Sensitivity 1.0000 0.9903 0.9873 0.9865 0.9982  
## Specificity 0.9981 0.9979 0.9967 0.9994 0.9996  
## Pos Pred Value 0.9952 0.9912 0.9845 0.9969 0.9982  
## Neg Pred Value 1.0000 0.9977 0.9973 0.9974 0.9996  
## Prevalence 0.2845 0.1935 0.1743 0.1638 0.1839  
## Detection Rate 0.2845 0.1917 0.1721 0.1616 0.1835  
## Detection Prevalence 0.2858 0.1934 0.1749 0.1621 0.1839  
## Balanced Accuracy 0.9991 0.9941 0.9920 0.9930 0.9989

## Interpreting Model Results

1. Random Forest model performed better than classification Trees and Gradient Boosting (GBM).
2. Random Forest model has accuracy of 99.3% compared to 95.9% for GBM and classification Tree for 49.7%.
3. Expected out-of-sample error (1 - accuracy) is estimated at 0.6% for Random Forest.

## Predicting 20 test cases

Finally, the Random forest model (RFmodel) tuned with cross-validation set will be used to predict 20 test cases available in the test data loaded earlier.

# Predicting on the testing data set using Random Forest Model  
  
prediction <- predict(RFmodel, testing)  
print(prediction)

## 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20   
## 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

answers <- as.vector(prediction)  
pml\_write\_files(answers)