Human Activity Recognition: Accuracy of Activities in Weight Lifting

Introduction

Considerable amount of research have been conducted by Universities and businesses on human wear devices for monitoring human's physical activities. A branch of the HAR research is focusing on accuracy of the intended activities, for example, daily routine activities of elderly people and athletics body movement during their regular exercise.

We will use the data set in http://groupware.les.inf.puc-rio.br/har to predict the accuracy by which the weight lifters are using dumbbell during their exercise. Six subjects are monitored while carrying out the unilateral dumbbell biceps curl in five different ways: exactly according to the specification (Class A), throwing the elbows to the front (Class B), lifting the dumbbell only halfway (Class C), lowering the dumbbell only halfway (Class D), and throwing the hips to the front (Class E). Class A corresponds to proper execution of the activity, whereas other four classes correspond to common mistakes. During these activities accelerometer attached to the subject arm, forearm, belt, and dumbbell are used to record the movement. We will build a prediction learning model from the data in the training set. The prediction model will be used to classify the observations in the test set.

Following packages and libraries are required: caret, Random Forest, AppliedPredictiveModeling.

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

library(randomForest)

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

library(AppliedPredictiveModeling)
```

Data preparation

The data set are partitioned into two sets: Training and Testing, which are transferred from the above website to the working directory and then imported to the appropriate data frame.

```
train <- read.csv2("pml-training.csv", header = TRUE, sep = ",")
test <- read.csv2("pml-testing.csv", header = TRUE, sep = ",")
for (i in ncol(train):1)
   if (sum(is.na(train[,i])) > 19000) {
        train[,i] <- NULL
        test[,i] <- NULL
   }
for (i in ncol(test):1) {
   s <- sum(is.na(test[,i]))
   if (s == nrow(test)) {
        test[,i] <- NULL
   }
</pre>
```

```
##
##
                  Α
                        В
                             С
                                   D
                                        Ε
##
                           750
                                      686
               1165
                      776
                                 515
     adelmo
                           493
##
     carlitos
                834
                      690
                                 486
                                      609
##
     charles
                899
                      745
                           539
                                 642
                                      711
##
     eurico
                865
                      592
                           489
                                 582
                                      542
##
     jeremy
               1177
                      489
                           652
                                 522
                                      562
##
     pedro
                640
                     505
                           499
                                 469
                                     497
```

```
train <- train[,-c(1:7)]
test <- test[,-c(1:7)]</pre>
```

Predictors with large number of NA (> 19000) are removed from both data sets and the remaining ones except classe in train and problem_id in test, are covnerted to numeric. The cross-table between user_name and classe in train set shows a uniform distribution of observations for each pair of class (classe and user_name) values. This is an indication that the model should be able to predict an outcome for every combination of class pairs. Highly correlated predictors are also removed from the data sets. There were no duplicate observation in the training set.

```
Corr <- cor(train[,-53])
highCorr <- findCorrelation(Corr, 0.90)
strain <- train[, -highCorr]
stest <- test[, -highCorr]</pre>
```

The Model

Considering the number of predictors and the inherent dependencies among them, we used Random Forest to build the prediction model, which exhaustively generates multiple trees and selects the best tree model from the permutation of predictors, averaged over accuracy on the out-of-bag portion of the data.

```
set.seed(123)
rfmodel <- randomForest(classe~., data=strain, importance=T, ntree=500)
print(rfmodel)
##</pre>
```

```
С
##
                         D
                              E class.error
        Α
## A 5576
              4
                   0
                         0
                              0
                                   0.0007168
        9 3780
## B
                   8
                                   0.0044772
## C
        0
             15 3406
                              0
                                   0.0046756
                         1
## D
        1
              0
                  23 3189
                              3
                                   0.0083955
## E
              0
                         8 3598
                                   0.0024951
                   1
```

```
predict(rfmodel, newdata=stest)
```

```
## 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 ## B A A A E E E A A A E A A E A A D B B ## Levels: A B C D E
```

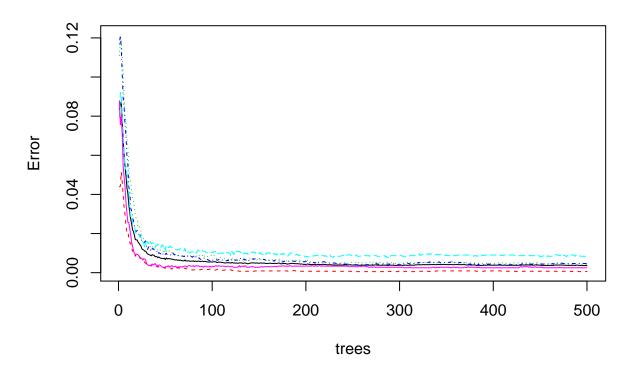
The estimate of error rate of 0.37% and near zero classification errors for each classe is an indication of a good fit, perhaps somewhat an overfit. The prediction from the test data is also displayed. We will look at the output of few statistics of this model.

```
head(varImp(rfmodel, sort=T, scale=F))
```

```
##
                          Α
                                   В
                                           С
                                                             Ε
                    0.10386 0.123599 0.14901 0.16466 0.238025
## roll_belt
## pitch_belt
                    0.09857 0.079791 0.16154 0.15131 0.047394
## yaw_belt
                    0.09735 0.107848 0.12593 0.13824 0.090971
## total_accel_belt 0.02514 0.029686 0.03575 0.03261 0.035702
## gyros_belt_x
                    0.01335 0.008935 0.01522 0.01208 0.003760
                    0.02241 0.020379 0.02637 0.01877 0.008315
## gyros_belt_y
```

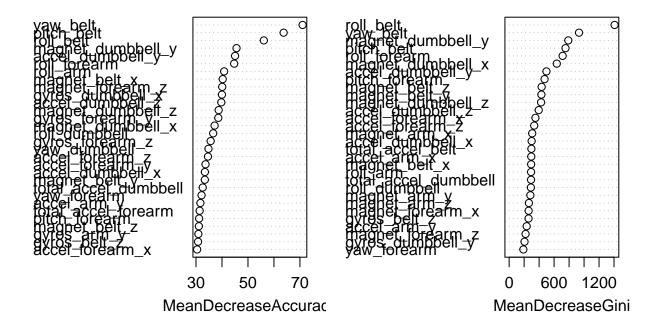
```
plot(rfmodel, main="Estimate of error rate")
```

Estimate of error rate



varImpPlot(rfmodel, sort=T)

rfmodel



```
cv <- rfcv(stest[,-1],stest[,1],cv.fold=10)
cv$error.cv

## 50 25 12 6 3 1
## 24.07 25.37 28.09 28.56 28.12 27.01</pre>
```

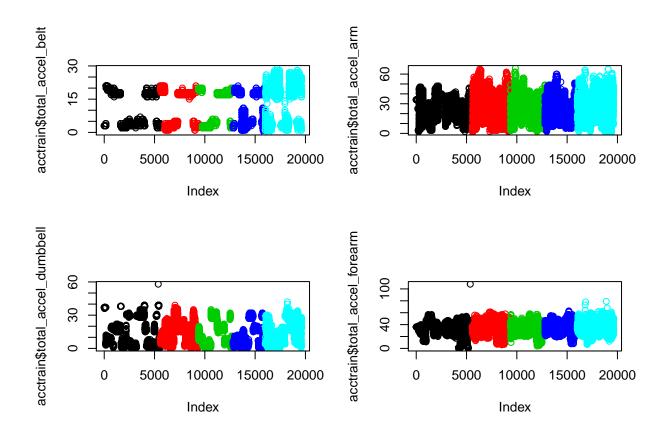
The plot of estimate error shows the errors for each class, trending down as the number of trees grow and the plot of varImp displays the contribution of the predictors to the model. Results of cross-validation shows the error for the number of variables used in each step of building the model.

Refining the Model

The above random forest model, though has a small miss-classification error did not reflect a uniform prediction of all classe (e.g. no prediction for Class C and more than 50% of the test observation are of Class A, correct execution of dumbbell exercise) of test data. To explore for better model, we used the list of important predictors and divided to different groups of accelerator, Gyroscope, and Mag. and for each group or combination of them generated a random forest model and measured the estimated error. It appeared that the predictors of accelerometers has the minimal error rate and fully predicted all classes in the test data.

```
acctrain <- train[,c(53,4,8:10,17,21:23,30,34:36,43,47:49)]
acctest <- test[,c(53,4,8:10,17,21:23,30,34:36,43,47:49)]
par(mfrow=c(2,2))
plot(acctrain$total_accel_belt,col=acctrain$classe)
plot(acctrain$total_accel_arm,col=acctrain$classe)</pre>
```

```
plot(acctrain$total_accel_dumbbell,col=acctrain$classe)
plot(acctrain$total_accel_forearm,col=acctrain$classe)
```

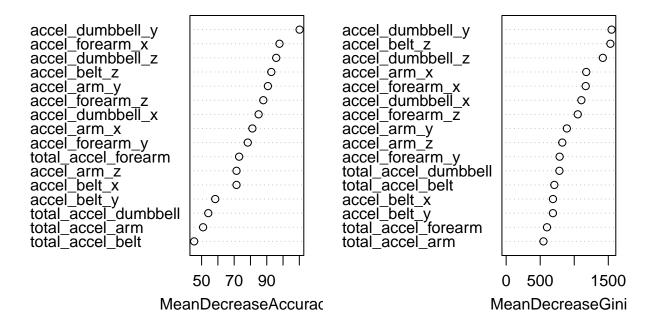


Plot of four predictors shown above depicts a clear separation of the data for total acceleration.

```
rfmod <- randomForest(classe~., data=acctrain, importance=T, ntree=500)
print(rfmod)</pre>
```

```
##
## Call:
    randomForest(formula = classe ~ ., data = acctrain, importance = T,
##
                                                                                  ntree = 500)
##
                   Type of random forest: classification
##
                         Number of trees: 500
## No. of variables tried at each split: 4
##
##
           OOB estimate of
                             error rate: 4.31%
##
  Confusion matrix:
##
        Α
             В
                   С
                        D
                              E class.error
## A 5414
            29
                  63
                       72
                              2
                                    0.02975
## B
      106 3558
                  86
                       21
                            26
                                    0.06294
                       30
## C
       51
            70 3266
                                    0.04559
                              5
## D
       58
            16
                 110 3017
                            15
                                    0.06188
## E
                       26 3522
                                    0.02357
        5
            37
                  17
```

rfmod



```
predict(rfmod,newdata=acctest, type="class")
```

```
## 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20
## B A C A A E D B A A B C B A E E A B B B
## Levels: A B C D E
```

Summary

As it appears to build the best random forest model with high accuracy we need all the quantitative predictors, giving $\sim 100\%$ accuracy (miss-classifications error of 0.37%). To predict the data in the test set we need a model that can predict for all classes (A, B, C, D, E), for which the initial random forest model did not accomplish. However, this model divided the training data into two groups with proper use of dumbbell (28%) and without proper use of the dumbbell (78%). we felt that the model has overfit the training data and is not providing a good prediction for the test data.

Using the importance() variables we narrowed down the model to the set of acceleration predictors and produced a random forest model for validation of the test data. This model exhibited a 4% miss-classifications error but gave an accurate prediction for the test data.