Activity Recognition Using Predictive Analytics

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Overview

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. The aim of this project is to predict the manner in which participants perform a barbell lift. The data comes from http://groupware.les.inf.puc-rio.br/har wherein 6 participants were asked to perform the same set of exercises correctly and incorrectly with accelerometers placed on the belt, forearm, arm, and dumbell.

For the purpose of this project, the following steps would be followed:

```
1. Data Preprocessing
```

2. Exploratory Analysis 3. Prediction Model Selection

4. Predicting Test Set Output

Data Preprocessing First, we load the training and testing set from the online sources and then split the training set further into training and test sets.

```
library(caret)
 setwd("~/Projects/R/Coursera-Practical-Machine-Learning-Assignment-1/")
 trainURL <- "http://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv"
 testURL <- "http://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv"
 training <- read.csv(url(trainURL))</pre>
 testing <- read.csv(url(testURL))</pre>
 label <- createDataPartition(training$classe, p = 0.7, list = FALSE)</pre>
 train <- training[label, ]</pre>
 test <- training[-label, ]</pre>
From among 160 variables present in the dataset, some variables have nearly zero variance whereas some contain a lot of NA terms which need
```

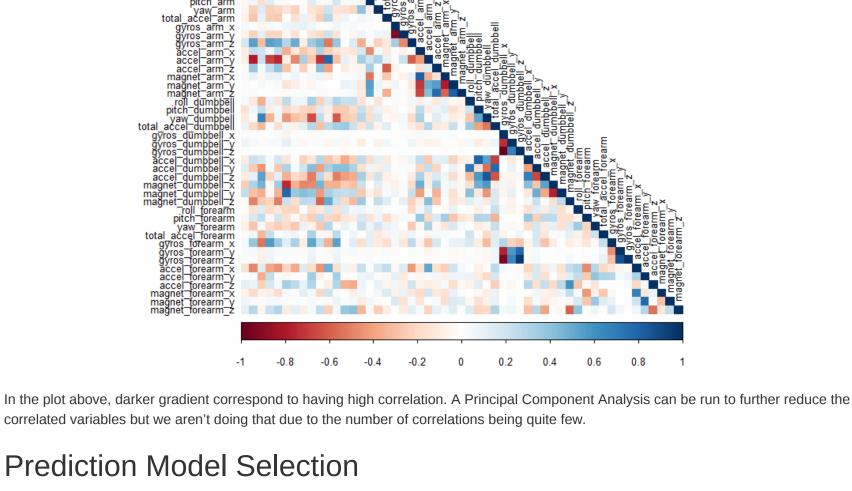
to be excluded from the dataset. Moreover, other 5 variables used for identification can also be removed. NZV <- nearZeroVar(train)

```
train <- train[ ,-NZV]</pre>
 test <- test[ ,-NZV]</pre>
 label <- apply(train, 2, function(x) mean(is.na(x))) > 0.95
 train <- train[, -which(label, label == FALSE)]</pre>
 test <- test[, -which(label, label == FALSE)]</pre>
 train <- train[ , -(1:5)]</pre>
 test <- test[ , -(1:5)]
As a result of the preprocessing steps, we were able to reduce 160 variables to 54.
```

Exploratory Analysis Now that we have cleaned the dataset off absolutely useless varibles, we shall look at the dependence of these variables on each other through a

correlation plot.

library(corrplot) corrMat <- cor(train[,-54])</pre> corrplot(corrMat, method = "color", type = "lower", tl.cex = 0.8, tl.col = rgb(0,0,0))



We will use 3 methods to model the training set and thereby choose the one having the best accuracy to predict the outcome variable in the testing set. The methods are Decision Tree, Random Forest and Generalized Boosted Model. A confusion matrix plotted at the end of each model will help visualize the analysis better.

Decision Tree

49

8

С

D

##

##

627

34 32

control <- trainControl(method = "cv", number = 3, verboseIter=FALSE)</pre>

Type of random forest: classification Number of trees: 500

randomForest(x = x, y = y, mtry = param\$mtry)

No. of variables tried at each split: 27

A 3904 1 0 0 1 0.0005120328

No Information Rate: 0.2845 P-Value [Acc > NIR] : < 2.2e-16

Mcnemar's Test P-Value : NA

Confusion Matrix and Statistics

Reference

Α A 1670 10

0

D 1

E 0

Detection Prevalence 0.2856

predictRF <- predict(modelRF, testing)</pre>

Balanced Accuracy

predictRF

С

Overall Statistics

3 1117

1

0

0

11

11 1009 10

Accuracy : 0.9888

3 949

3 0 1074

1

3

2

Statistics by Class:

Kappa: 0.9968

modelRF <- train(classe ~ ., data = train, method = "rf", trControl = control)</pre>

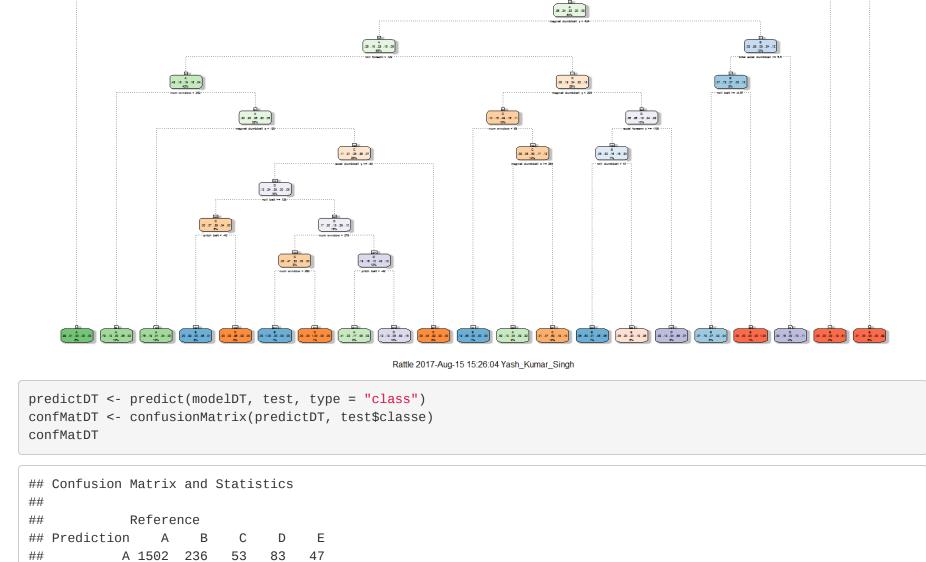
67 829 139

95 152 51 609 116

library(rpart) library(rpart.plot) library(rattle)

set.seed(13908) modelDT <- rpart(classe ~ ., data = train, method = "class")</pre>

fancyRpartPlot(modelDT)



```
E 20
                  57
                         59 101 826
 ## Overall Statistics
 ##
                 Accuracy: 0.7465
 ##
 ##
                   95% CI: (0.7352, 0.7575)
 ##
       No Information Rate: 0.2845
       P-Value [Acc > NIR] : < 2.2e-16
 ##
 ##
 ##
                    Kappa: 0.678
 ##
    Mcnemar's Test P-Value : < 2.2e-16
 ##
 ## Statistics by Class:
 ##
 ##
                       Class: A Class: B Class: C Class: D Class: E
 ## Sensitivity
                       0.8973 0.5505 0.8080 0.6317 0.7634
                      0.9005 0.9707 0.9418 0.9159 0.9507
 ## Specificity
 ## Pos Pred Value
                      0.7819 0.8185 0.7455 0.5953 0.7770
                      0.9566 0.9000 0.9587 0.9270
 ## Neg Pred Value
                                                          0.9469
 ## Prevalence
                        0.2845 0.1935 0.1743 0.1638 0.1839
 ## Prevalence 0.2845 0.1935 0.1743 0.1638 0.1839  
## Detection Rate 0.2552 0.1065 0.1409 0.1035 0.1404
 ## Detection Prevalence 0.3264 0.1302 0.1890 0.1738 0.1806
 ## Balanced Accuracy
                         0.8989 0.7606 0.8749 0.7738 0.8570
Random Forest
```

00B estimate of error rate: 0.21% ## Confusion matrix: ## A B C D E class.error

library(caret) set.seed(13908)

##

##

Call:

confMatRF

##

##

##

##

##

##

##

##

Prediction

modelRF\$finalModel

```
## B 4 2651 3 0 0 0.0026335591
    0 2 2394 0 0 0.0008347245
## C
## D 0 0 13 2238 1 0.0062166963
## E 0 1 0 3 2521 0.0015841584
predictRF <- predict(modelRF, test)</pre>
confMatRF <- confusionMatrix(predictRF, test$classe)</pre>
```

```
## Confusion Matrix and Statistics
         Reference
## Prediction A B C
     A 1674 2 0 0
      B 0 1136 4 0
       C 0 0 1022 7 0
       D 0 1 0 957 1
##
##
       E 0 0 0 01081
##
## Overall Statistics
            Accuracy: 0.9975
##
##
              95% CI: (0.9958, 0.9986)
```

```
## Detection Prevalence 0.2848 0.1937 0.1749 0.1630 0.1837
 ## Balanced Accuracy 0.9998 0.9983 0.9973 0.9962 0.9995
Generalized Boosted Model
 library(caret)
 set.seed(13908)
 control <- trainControl(method = "repeatedcv", number = 5, repeats = 1, verboseIter = FALSE)</pre>
 modelGBM <- train(classe ~ ., data = train, trControl = control, method = "gbm", verbose = FALSE)</pre>
 modelGBM$finalModel
 ## A gradient boosted model with multinomial loss function.
 ## 150 iterations were performed.
 ## There were 53 predictors of which 41 had non-zero influence.
 predictGBM <- predict(modelGBM, test)</pre>
 confMatGBM <- confusionMatrix(predictGBM, test$classe)</pre>
 confMatGBM
```

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

```
95% CI: (0.9858, 0.9913)
##
##
       No Information Rate: 0.2845
##
       P-Value [Acc > NIR] : < 2.2e-16
##
                       Kappa: 0.9858
##
##
    Mcnemar's Test P-Value : NA
## Statistics by Class:
                          Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                         0.9976 0.9807 0.9834 0.9844
                                                                   0.9926
## Sensitivity 0.9974 0.9956 0.9951 0.9986
## Pos Pred Value 0.9935 0.9815 0.9768 0.9927
## Neg Pred Value 0.9990 0.9954 0.9965 0.9970
                                                                   0.9994
                                                                   0.9972
                                                                   0.9983
## Prevalence
                         0.2845 0.1935 0.1743 0.1638
                                                                   0.1839
## Detection Rate
                            0.2838 0.1898 0.1715 0.1613
                                                                   0.1825
```

0.9915

0.1830

0.9960

As Random Forest offers the maximum accuracy of 99.75%, we will go with Random Forest Model to predict our test data class variable. **Predicting Test Set Output**

0.1934 0.1755 0.1624

0.9975 0.9881 0.9892

```
## [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
```