Practical Machine Learning Course Proejct

RS

13 9 2020

R Markdown

Background

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. In this project, your goal will be to use data from accelerometers on the belt, forearm, arm, and dumbell of 6 participants. They were asked to perform barbell lifts correctly and incorrectly in 5 different ways. More information is available from the website here: http://web.archive.org/web/20161224072740/http:/groupware.les.inf.puc-rio.br/har (http://web.archive.org/web/20161224072740/http:/groupware.les.inf.puc-rio.br/har) (see the section on the Weight Lifting Exercise Dataset).

Data Preparation

Load the packages and the training and test data. Create partition with the training data set

```
## Loading required package: lattice

## Loading required package: ggplot2

library(corrplot)

## Warning: package 'corrplot' was built under R version 3.6.3

## corrplot 0.84 loaded

library(rpart)
library(rpart.plot)

## Warning: package 'rpart.plot' was built under R version 3.6.3
```

```
library(rattle)
## Warning: package 'rattle' was built under R version 3.6.3
## Loading required package: tibble
## Loading required package: bitops
## Rattle: A free graphical interface for data science with R.
## Version 5.4.0 Copyright (c) 2006-2020 Togaware Pty Ltd.
## Geben Sie 'rattle()' ein, um Ihre Daten mischen.
library(randomForest)
## randomForest 4.6-14
## 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
library (gbm)
## Warning: package 'gbm' was built under R version 3.6.3
## Loaded gbm 2.1.8
```

```
# Load training and test data
myTrain <- read.csv("pml-training.csv")
myTest <- read.csv("pml-testing.csv")

#Create partition with the training data set
inTrain <- createDataPartition(myTrain$classe, p=0.7, list=FALSE)
myTrainSet <- myTrain[inTrain, ]
myTestSet <- myTrain[-inTrain, ]
dim(myTrainSet)</pre>
```

```
## [1] 13737 160
```

```
dim(myTestSet)
```

```
## [1] 5885 160
```

```
#Remove variables with nearly zero variance AND ariables that are almost always NA

myNZV <- nearZeroVar(myTrainSet)
myTrainSet <- myTrainSet[, -myNZV]

myTestSet <- myTestSet[, -myNZV]

myMNA <- sapply(myTrainSet, function(x) mean(is.na(x))) > 0.95
myTrainSet <- myTrainSet[, myMNA==F]
myTestSet <- myTestSet[, myMNA==F]

# remove ID only variables: columns 1 - 5
myTrainSet <- myTrainSet[, -(1:5)]
myTestSet <- myTestSet[, -(1:5)]
dim(myTrainSet)</pre>
```

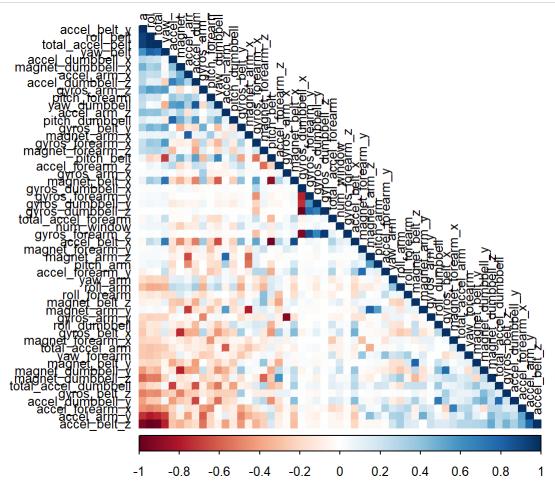
```
## [1] 13737 54
```

```
dim(myTestSet)
```

```
## [1] 5885 54
```

Correlation Analysis

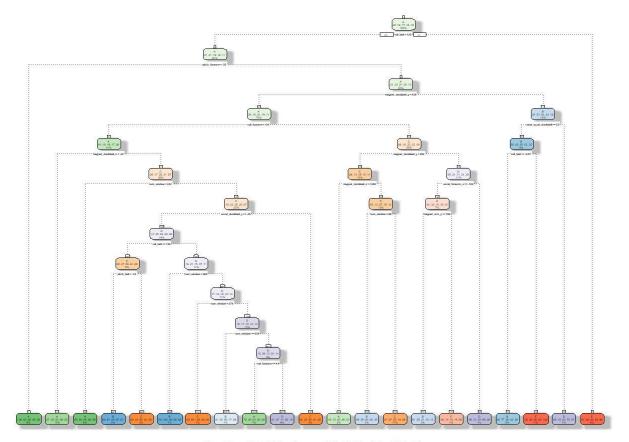
Perform a correlation analysis among variables before proceeding with the modeling procedure. Correlated variables/predictors are shown in dark colors



Model Evaluation / Building

I: Decision Trees

```
# Fit the model
set.seed(12345)
myDecTreeModel <- rpart(classe ~ ., data=myTrainSet, method="class")
fancyRpartPlot(myDecTreeModel)</pre>
```



Rattle 2020-Sep-13 22:18:55 Ronnz

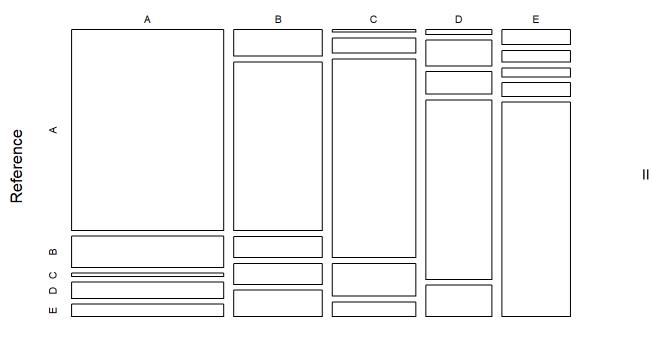
```
# Perform prediction on test data set
myDecTreePredict <- predict(myDecTreeModel, newdata=myTestSet, type="class")
myDecTreeConf <- confusionMatrix(myDecTreePredict, myTestSet$classe)
myDecTreeConf</pre>
```

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```
## Confusion Matrix and Statistics
##
##
           Reference
            A B
## Prediction
                     C D
                               Ε
         A 1485 230
                     25 119
##
                               93
##
         B 114 725 92
                         90 114
         C 10 62 807 133
##
                              60
         D 15 83
                     72 576 101
##
##
         E 50 39 30
                         46 714
##
## Overall Statistics
##
               Accuracy : 0.7319
##
                 95% CI: (0.7203, 0.7431)
##
    No Information Rate: 0.2845
##
     P-Value [Acc > NIR] : < 2.2e-16
##
##
                 Kappa: 0.6587
##
##
  Mcnemar's Test P-Value : < 2.2e-16
##
##
## Statistics by Class:
##
##
                    Class: A Class: B Class: C Class: D Class: E
                     0.8871 0.6365 0.7865 0.59751 0.6599
## Sensitivity
                     0.8891 0.9136 0.9455 0.94493 0.9656
## Specificity
## Pos Pred Value
                     ## Neg Pred Value
                     0.9519 0.9128 0.9545 0.92299 0.9265
## Prevalence
                     0.2845 0.1935 0.1743 0.16381 0.1839
## Detection Rate
                    0.2523 0.1232 0.1371 0.09788 0.1213
## Detection Prevalence 0.3317 0.1929 0.1822 0.14393 0.1494
## Balanced Accuracy
                    0.8881 0.7751 0.8660 0.77122 0.8128
```

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Decision Tree Accuracy = 0.732



Prediction

: Random Forest

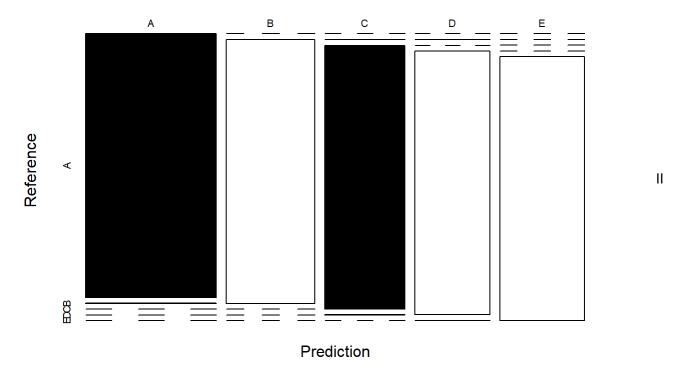
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```
##
## Call:
## randomForest(x = x, y = y, mtry = param$mtry)
##
              Type of random forest: classification
                    Number of trees: 500
##
## No. of variables tried at each split: 27
##
         OOB estimate of error rate: 0.23%
##
## Confusion matrix:
        B C D E class.error
##
      Α
         2 0 0 0.0005120328
## A 3904
     5 2651 2 0 0 0.0026335591
## B
## C
     0 5 2391 0 0 0.0020868114
     0 0 11 2240 1 0.0053285968
## D
    0 0 0 5 2520 0.0019801980
## E
```

```
# Perform prediction on test data set
myRFPredict <- predict(myRFModFit, newdata=myTestSet)
myRFConf <- confusionMatrix(myRFPredict, myTestSet$classe)
myRFConf</pre>
```

```
## Confusion Matrix and Statistics
##
##
           Reference
## Prediction
              A B
          A 1674
                   1
                        0
##
##
          В
               0 1136
                     0
                            0
               0
                  1 1026
##
          С
                          1
          D
               0
                   1
                        0 963
##
##
          E
               0
                  0
                        0 0 1081
##
## Overall Statistics
##
                Accuracy : 0.9992
##
                  95% CI: (0.998, 0.9997)
##
    No Information Rate: 0.2845
##
      P-Value [Acc > NIR] : < 2.2e-16
##
##
                  Kappa: 0.9989
##
##
##
  Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##
                     Class: A Class: B Class: C Class: D Class: E
                      1.0000 0.9974 1.0000 0.9990 0.9991
## Sensitivity
                      0.9998 1.0000 0.9996 0.9996 1.0000
## Specificity
## Pos Pred Value
                      0.9994 1.0000 0.9981 0.9979 1.0000
## Neg Pred Value
                      1.0000 0.9994 1.0000 0.9998 0.9998
## Prevalence
                      0.2845 0.1935 0.1743 0.1638 0.1839
## Detection Rate
                     0.2845 0.1930 0.1743 0.1636 0.1837
## Detection Prevalence 0.2846 0.1930 0.1747 0.1640 0.1837
## Balanced Accuracy
                     0.9999 0.9987 0.9998 0.9993 0.9995
```

Random Forest Accuracy = 0.999



: Generalized Boosted Model

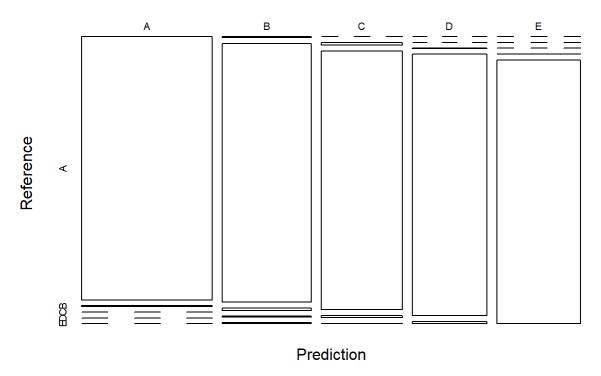
```
## A gradient boosted model with multinomial loss function.
## 150 iterations were performed.
## There were 53 predictors of which 53 had non-zero influence.
```

```
#Perform prediction on test data set
myGBMPrediction <- predict(myGBMmodfit, newdata=myTestSet)
myGBMconfm <- confusionMatrix(myGBMPrediction, myTestSet$classe)
myGBMconfm</pre>
```

```
## Confusion Matrix and Statistics
##
##
          Reference
## Prediction
            A B
                     С
                              E
         A 1670
                  5
                      0
##
##
          В
              4 1123 11
                11 1014
                         9
##
          С
              0
         D
              0
                   0
                       1 950
##
##
          E
              0
                 0
                       0 1 1069
##
## Overall Statistics
##
               Accuracy: 0.99
##
                 95% CI: (0.9871, 0.9924)
##
    No Information Rate: 0.2845
##
     P-Value [Acc > NIR] : < 2.2e-16
##
##
                  Kappa : 0.9873
##
##
##
  Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##
                    Class: A Class: B Class: C Class: D Class: E
                     0.9976 0.9860 0.9883 0.9855 0.9880
## Sensitivity
## Specificity
                     0.9988 0.9952 0.9957 0.9982 0.9998
## Pos Pred Value
                     0.9970 0.9799 0.9797 0.9906 0.9991
## Neg Pred Value
                     0.9990 0.9966 0.9975 0.9972 0.9973
## Prevalence
                     0.2845 0.1935 0.1743 0.1638 0.1839
## Detection Rate
                    ## Detection Prevalence 0.2846 0.1947 0.1759 0.1630 0.1818
## Balanced Accuracy
                    0.9982
                              0.9906 0.9920 0.9918 0.9939
```

```
# Plot the matrix results
plot(myGBMconfm$table, col = myGBMconfm$byClass,
    main = paste("GBM Accuracy =", round(myGBMconfm$overall['Accuracy'], 3)))
```

GBM Accuracy = 0.99



Applying the Selected Model to the Test Data. Since Random forest has best prediction use Random forest

#Random Forest model will be applied
myResults <- predict(myRFModFit, newdata=myTestSet)
myResults</pre>

##	[1]	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A
##	[38]	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	Α	А	Α	Α	Α	Α	А	Α	Α	Α	Α	Α	Α	Α
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##	[926]	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_
##	[963]	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_
##	[1000]	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_
##	[1037]	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_
##	[1074]	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_
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