Prediction Assignment Project

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Project Goal

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 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, etc. The goal of your project is to predict the manner in which they did the exercise. You should create a report describing how you built your model, how you used cross validation, what you think the expected out of sample error is, and why you made the choices you did. You will also use your prediction model to predict 20 different test cases.

Acquire Datasets and Prepare for Analysis

```
#load the data libraries that may be needed to support analysis
library(caret, warn.conflicts = FALSE, quietly = TRUE)
library(rpart, warn.conflicts = FALSE, quietly = TRUE)
library(rpart.plot, warn.conflicts = FALSE, quietly = TRUE)
library(RColorBrewer, warn.conflicts = FALSE, quietly = TRUE)
library(rattle, warn.conflicts = FALSE, quietly = TRUE)
## Rattle: A free graphical interface for data science with R.
## Version 5.2.0 Copyright (c) 2006-2018 Togaware Pty Ltd.
## Type 'rattle()' to shake, rattle, and roll your data.
library(randomForest, warn.conflicts = FALSE, quietly = TRUE)
## randomForest 4.6-14
## Type rfNews() to see new features/changes/bug fixes.
library(gbm, warn.conflicts = FALSE, quietly = TRUE)
## Loaded gbm 2.1.4
library(plyr, warn.conflicts = FALSE, quietly = TRUE)
#download data from the internet
download.file(url = "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-
training.csv",
              destfile = "./pml-train.csv", method = "curl")
```

Clean the Data

Remove columns with missing values, "NA" values, etc. Time dependence values will be removed.

```
features <- names(dfTest[,colSums(is.na(dfTest)) == 0])[8:59]

# Only use features used in testing cases
dfTrain <- dfTrain[,c(features,"classe")]
dfTest <- dfTest[,c(features,"problem_id")]

#View data file dimensions for each file post cleansing
dim(dfTrain); dim(dfTest);
## [1] 19622 53
## [1] 20 53</pre>
```

Partition the Dataset

Use a 60/40 split (train/test) for partitioning the dataset as was alluded to in the training materials to increase performance and accuracy of the model. Therefore, p is set = to 0.6

```
set.seed(10)

inTrain <- createDataPartition(dfTrain$classe, p=0.6, list=FALSE)

train <- dfTrain[inTrain,]

test <- dfTrain[-inTrain,]</pre>
```

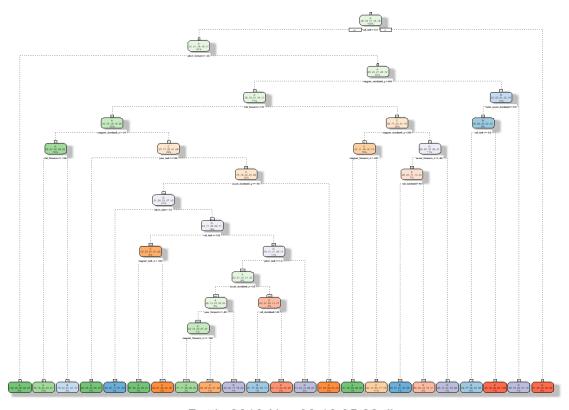
```
dim(train); dim(test);
## [1] 11776 53
## [1] 7846 53
```

Build the Decision Tree

Although easy to interpret, results may be variable which will affect accuracy. Will set cross validation to "cv" and 10 for resampling. Will set method to "class" to get a factor of classifications based on the responses.

```
modFDT <- rpart(classe ~ ., data = train, method="class", control = rpart.con
trol(method = "cv", number = 10))

fancyRpartPlot(modFDT)</pre>
```



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Predict with the Decision Tree Model

If we can get an accuracy of 70% or above, then we'll consider it to be acceptable.

```
pred <- predict(modFDT, test, type = "class")</pre>
confusionMatrix(pred, test$classe)
## Confusion Matrix and Statistics
##
            Reference
##
## Prediction
                A
                     В
                         С
                              D
                                   Ε
           A 1994 264
                         49
                             85
                                  15
##
               65 836
                         72 108
                                 121
##
           В
##
           С
               69 194 1095 174
                                186
           D
               67 112
                        78 805
                                 78
##
               37 112
                        74 114 1042
##
           Ε
##
## Overall Statistics
##
                 Accuracy: 0.7357
##
                   95% CI : (0.7258, 0.7454)
##
     No Information Rate: 0.2845
##
      P-Value [Acc > NIR] : < 2.2e-16
##
##
##
                    Kappa: 0.6649
   Mcnemar's Test P-Value : < 2.2e-16
## Statistics by Class:
##
##
                       Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                                0.5507 0.8004
                        0.8934
                                                   0.6260
                                                           0.7226
## Specificity
                        0.9264
                                0.9422 0.9038
                                                   0.9489
                                                           0.9474
## Pos Pred Value
                        0.8284
                                0.6955 0.6374
                                                   0.7061 0.7556
  Neg Pred Value
                        0.9562 0.8974 0.9555
                                                   0.9283
                                                           0.9381
## Prevalence
                        0.2845
                                0.1935 0.1744
                                                   0.1639
                                                           0.1838
## Detection Rate
                        0.2541 0.1066 0.1396
                                                   0.1026
                                                           0.1328
## Detection Prevalence 0.3068 0.1532 0.2190 0.1453
                                                           0.1758
```

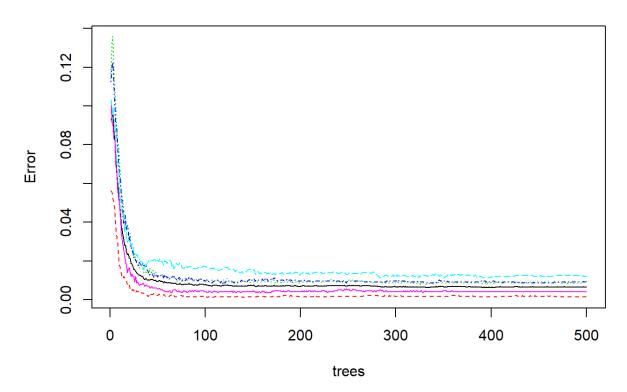
Accuracy is 72.6%, thereby, we consider to be acceptable.

Build the Random Forest (rf) Model

One of the proces of the Random Forest Model is accuracy. But Overfitting can be a problem. As stated previously, we will use a 40% test sample. The error estimate is expected to be less than 5%.

```
modFRF <- randomForest(classe ~ ., data = train, method = "rf", importance =
T, trControl = trainControl(method = "cv", classProbs=TRUE, savePredictions=TR
UE, allowParallel=TRUE, number = 10))
plot(modFRF)</pre>
```



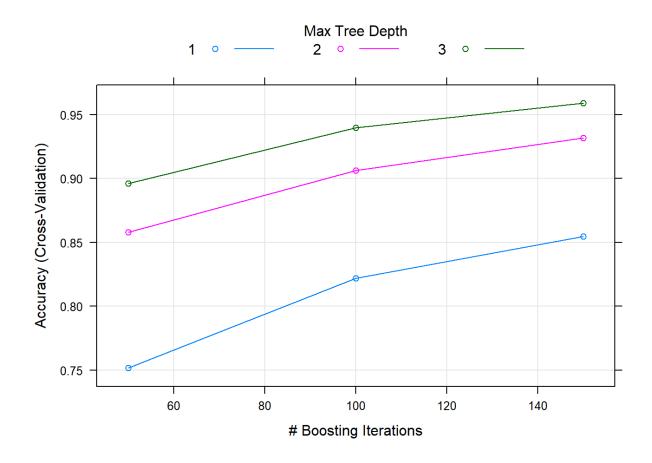


Build the Generalized Boosted Regression Model (gbm)

The goal is to minimize error on the training set. We will use gbm (boosting with trees). Will set cross validation to "cv" and 10 for resampling. Will set Verbose to False to avoid the extensive info and error logs being printed.

```
modFB <- train(classe ~ ., method = "gbm", data = train,</pre>
                    verbose = F,
                     trControl = trainControl(method = "cv", number = 10))
modFB
## Stochastic Gradient Boosting
##
## 11776 samples
##
      52 predictor
       5 classes: 'A', 'B', 'C', 'D', 'E'
## No pre-processing
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 10598, 10599, 10600, 10597, 10598, 10599, ...
## Resampling results across tuning parameters:
##
##
     interaction.depth n.trees Accuracy
                                             Kappa
##
                          50
                                  0.7516973 0.6850818
                         100
                                  0.8220076 0.7747438
##
     1
                                  0.8546175 0.8159894
                         150
##
     1
     2
                         50
                                  0.8577619 0.8197622
##
                                  0.9064200 0.8815455
     2
                        100
##
     2
                         150
                                  0.9318102 0.9136974
##
     3
                         50
                                  0.8960575 0.8683848
##
                                  0.9397924 0.9238092
##
     3
                         100
     3
                         150
                                  0.9588157 0.9478923
##
##
## Tuning parameter 'shrinkage' was held constant at a value of 0.1
##
## Tuning parameter 'n.minobsinnode' was held constant at a value of 10
## Accuracy was used to select the optimal model using the largest value.
```

```
## The final values used for the model were n.trees = 150,
## interaction.depth = 3, shrinkage = 0.1 and n.minobsinnode = 10.
plot(modFB)
```



Predict with the rf Model

```
pred <- predict(modFRF, test, type = "class")</pre>
confusionMatrix(pred, test$classe)
## Confusion Matrix and Statistics
##
              Reference
## Prediction
                  Α
                                   D
                                         Ε
             A 2232
                        3
                              0
                                   0
                                         0
                             7
##
             В
                  0 1513
                                   1
                                         0
                        2 1361
##
                                  13
                                         0
                             0 1271
```

```
E 0 0 0 1 1436
##
##
## Overall Statistics
##
##
               Accuracy: 0.9958
                 95% CI: (0.9941, 0.9971)
##
     No Information Rate: 0.2845
     P-Value [Acc > NIR] : < 2.2e-16
##
                  Kappa: 0.9947
##
  Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
                   Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                     1.0000 0.9967 0.9949 0.9883 0.9958
## Specificity
                     0.9995 0.9987 0.9977 0.9991 0.9998
## Pos Pred Value 0.9987 0.9947 0.9891 0.9953 0.9993
                1.0000 0.9992 0.9989 0.9977 0.9991
## Neg Pred Value
## Prevalence
                     0.2845 0.1935 0.1744 0.1639 0.1838
                     0.2845 0.1928 0.1735 0.1620 0.1830
## Detection Rate
## Detection Prevalence 0.2849 0.1939 0.1754 0.1628 0.1832
## Balanced Accuracy 0.9997 0.9977 0.9963 0.9937 0.9978
```

The rf model achieved 99.4% accuracy.

Predict with gbm

```
pred <- predict(modFB, test)
confusionMatrix(pred, test$classe)
## Confusion Matrix and Statistics
##
## Reference
## Prediction A B C D E
## A 2189 48 0 1 3</pre>
```

```
29 1423 42 10
##
##
                   44 1308
                            36
                                 18
                                 20
##
          D
                    1
                       15 1227
                    2
                        3
##
                            12 1383
##
  Overall Statistics
                Accuracy: 0.9597
##
                  95% CI: (0.9551, 0.964)
##
     No Information Rate: 0.2845
##
      P-Value [Acc > NIR] : < 2.2e-16
##
##
##
                   Kappa: 0.9491
   Mcnemar's Test P-Value: 8.925e-09
## Statistics by Class:
##
##
                     Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                       0.9807 0.9374
                                       0.9561
                                                 0.9541
                                                         0.9591
 Specificity
                       0.9907 0.9844 0.9836
                                                 0.9938 0.9972
                       0.9768 0.9350 0.9250
## Pos Pred Value
                                                 0.9677 0.9872
## Neg Pred Value
                      0.9923 0.9850 0.9907 0.9910 0.9908
## Prevalence
                       0.2845 0.1935 0.1744 0.1639 0.1838
## Detection Rate
                       0.2790 0.1814 0.1667 0.1564 0.1763
## Detection Prevalence 0.2856 0.1940 0.1802 0.1616 0.1786
## Balanced Accuracy
                       0.9857 0.9609 0.9699
                                                 0.9739 0.9781
```

The gbm achieved 95.9% accuracy.

Predict with the Test Dataset

```
predDT <- predict(modFDT, dfTest)
predDT
## A B C D E
## 1 0.03041363 0.119221411 0.53163017 0.16788321 0.150851582</pre>
```

```
0.75634518 0.182741117 0.01840102 0.03362944 0.008883249
     0.04885993 0.203583062 0.17752443 0.15960912 0.410423453
     0.13056836 0.033794163 0.11981567 0.67895545 0.036866359
     0.70662461 0.110410095 0.06309148 0.09463722 0.025236593
     0.03041363 0.119221411 0.53163017 0.16788321 0.150851582
     0.05823293 0.116465863 0.04016064 0.70080321 0.084337349
     0.70662461 0.110410095 0.06309148 0.09463722 0.025236593
     0.99676724 0.003232759 0.00000000 0.00000000 0.000000000
  10 0.75634518 0.182741117 0.01840102 0.03362944 0.008883249
  11 0.03041363 0.119221411 0.53163017 0.16788321 0.150851582
  12 0.04885993 0.203583062 0.17752443 0.15960912 0.410423453
  13 0.03041363 0.119221411 0.53163017 0.16788321 0.150851582
## 14 0.99676724 0.003232759 0.00000000 0.00000000 0.000000000
  15 0.04885993 0.203583062 0.17752443 0.15960912 0.410423453
  16 0.03816794 0.106870229 0.00000000 0.20229008 0.652671756
  17 0.96932515 0.000000000 0.01840491 0.00000000 0.012269939
  18 0.09830508 0.416949153 0.01355932 0.33559322 0.135593220
## 19 0.09830508 0.416949153 0.01355932 0.33559322 0.135593220
## 20 0.04968944 0.813664596 0.01242236 0.03105590 0.093167702
```

Apply the rf Prediction

```
predRF <- predict(modFRF, dfTest)
predRF
## 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20
## B A B A E D B A A B C B A E E A B B B
## Levels: A B C D E</pre>
```

Apply the gbm Prediction

```
predgbm <- predict(modFB, dfTest)
predgbm
## [1] B A B A E D B A A B C B A E E A B B B
## Levels: A B C D E</pre>
```

File to be Submitted

The rf model appears to have a high level of accuracy at 99.5%. With a level of accuracy this high, we can feel confident that any test cases that are submitted for analysis will be accurate.

```
project_files = function(x) {
    n = length(x)
    for(i in 1:n) {
        filename = paste0("problem_id_",i,".txt")
        write.table(x[i], file=filename, quote=FALSE, row.names=FALSE, col.names=FALSE)
    }
}
project_files(predRF)
```