# Machine Learning Project\_3

suszanna

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#### BACKGROUND

Various devices equipped with embedded accelerometers can be worn on the body during physical exercise to collect data for subsequent analysis of the quality of the exercise after the fact. These devices are used to measure quality of performance of the movements of arms, legs etc. Resulting data is trained by resampling with cross validation to come up with reproducible metrics for accuracy and error rates to classify each performance of the exercise. The data we use was collected from 6 participants who performed one set of 10 repetitions of the Unilateral Dumbbell Biceps Curl in five different variations of the specified exercise: \* Class A - exactly according to the specification \* Class B - throwing the elbows to the front \* Class C - lifting the dumbbell only halfway \* Class D - lowering the dumbbell only halfway \* Class E - throwing the hips to the front.

#### **GOALS**

The goal of this study is to make a reasonable prediction. To do this, we run models that produce predictions typical of each model. We compare the metrics of each (accuracy and error rates) & select the best prediction as our outcome. We focus on the 'classe' variable in the above described training set: A,B,C,D,E.

#### DATA

Load, clean and split

library(rattle)

## 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.
 ## Type 'rattle()' to shake, rattle, and roll your data.
 library(caret)
 ## Loading required package: lattice
 ## Loading required package: ggplot2
 library(rpart)
 library(rpart.plot)
 library(corrplot)
 ## corrplot 0.84 loaded
 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:ggplot2':
 ##
 ## margin
 ## The following object is masked from 'package:rattle':
 ##
 ## importance
 library(RColorBrewer)
Load and Read CSV Data
 trainFile <- "./data/pml-training.csv"
 testFile <- "./data/pml-testing.csv"
 trainRaw <- read.csv(trainFile)
```

testing <- testing[, cond]

#### Clean

```
NZV <- nearZeroVar(trainRaw, saveMetrics = TRUE)
head(NZV, 20)
```

```
freqRatio percentUnique zeroVar nzv
               1.000000 100.00000000 FALSE FALSE
## user name 1.100679 0.03057792 FALSE FALSE
## raw timestamp part 1 1.000000 4.26562022 FALSE FALSE
## raw timestamp part 2 1.000000 85.53154622 FALSE FALSE
## cvtd_timestamp
                   1.000668 0.10192641 FALSE FALSE
## new_window
                    47.330049 0.01019264 FALSE TRUE
## num_window
                    1.000000 4.37264295 FALSE FALSE
## roll_belt
                1.101904 6.77810621 FALSE FALSE
## pitch belt
                  1.036082 9.37722964 FALSE FALSE
## yaw belt
                  1.058480 9.97349913 FALSE FALSE
## total accel belt 1.063160 0.14779329 FALSE FALSE
## kurtosis roll belt 1921.600000 2.02323922 FALSE TRUE
## kurtosis picth belt 600.500000 1.61553358 FALSE TRUE
## kurtosis_yaw_belt 47.330049 0.01019264 FALSE TRUE
## skewness_roll_belt 2135.111111 2.01304658 FALSE TRUE
## skewness_roll_belt.1 600.500000 1.72255631 FALSE TRUE
## skewness_yaw_belt 47.330049 0.01019264 FALSE TRUE
## max_roll_belt
                   1.000000 0.99378249 FALSE FALSE
## max picth belt
                    1.538462 0.11211905 FALSE FALSE
## max yaw belt
                   training01 <- trainRaw[, !NZV$nzv]
testing01 <- testRaw[, !NZV$nzv]
regex <- grepl("^X|timestamp|user_name", names(training01))
training <- training01[, !regex]
testing <- testing01[, !regex]
cond <- (colSums(is.na(training)) == 0)
training <- training[, cond]
```

```
dim(training)

## [1] 19622 54

dim(testing)

## [1] 20 54

Split

set.seed(9573644)
inTrain <- createDataPartition(training$classe, p = 0.70, list = FALSE)
validation <- training[-inTrain, ]
training <- training[inTrain, ]

dim(training)

## [1] 13737 54

dim(validation)

## [1] 5885 54
```

#### **APPROACH**

TWO PREDICTIVE MODELS ARE RESAMPLED WITH CROSS VALIDATION TO PROVIDE ACCURACY & OUT OF SAMPLE ERROR RATES

PREDICTION MODEL 1: Recursive Partitioning and Regression Tree (rpart)

0.0538 accuracy/ 0.462 out-of-sample error

Train the model: for the Training data set, find accuracy for rpart with Model 1.

The resampling method 'cv' (cross validation) is used to train with the trainControl function. A re-sampling method involves repeatedly drawing samples from a training data set and refitting a model to obtain additional information about that model.

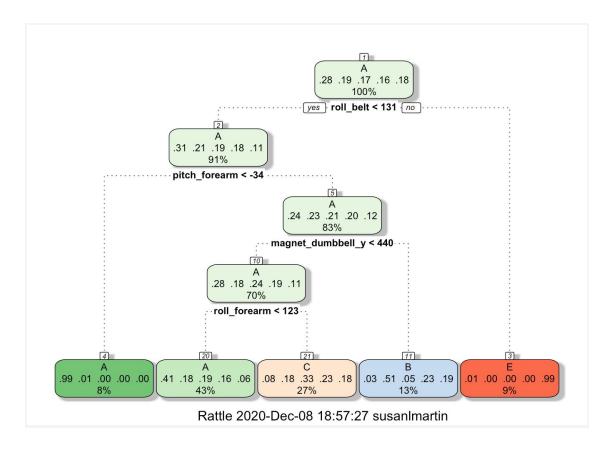
```
con_trol <- trainControl(method = "cv", number = 5)
model1_rpart <- train(classe ~ ., data = training, method = "rpart", trControl = con_trol)
print(model1_rpart, digits = 4)</pre>
```

```
## CART
##
## 13737 samples
    53 predictor
    5 classes: 'A', 'B', 'C', 'D', 'E'
##
## No pre-processing
## Resampling: Cross-Validated (5 fold)
## Summary of sample sizes: 10990, 10989, 10990, 10989, 10990
## Resampling results across tuning parameters:
##
## cp Accuracy Kappa
## 0.03901 0.5381 0.40609
## 0.05978 0.3928 0.16952
## 0.11830 0.3167 0.04917
##
```

## Accuracy was used to select the optimal model using the largest value.

## The final value used for the model was cp = 0.03901.

fancyRpartPlot(model1\_rpart\$finalModel)



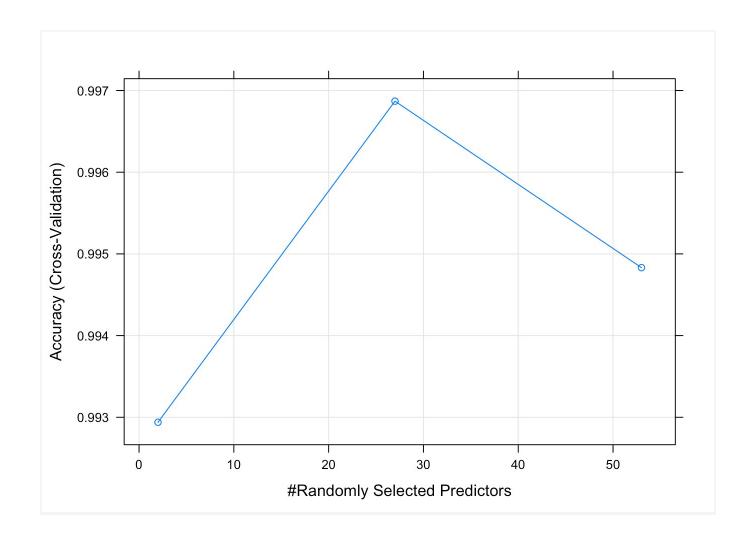
## PREDICTION MODEL 2: Random Forest

0.993 accuracy/ 0.007 out-of-sample error

Train the model: for the Training data set, find accuracy of the Random Forest model 2.

The resampling method 'cv' (cross validation) is used again here to train with the trainControl function. As a resampling procedure, cross validation is used to evaluate machine learning models on a limited data sample. It uses a limited sample in order to estimate how the model is expected to perform in general. It is then used to make predictions on data not used during the training of the model.

```
modelRF <- train(classe ~ ., data = training, method = "rf", trControl = trainControl(method = "cv", 5), ntree =
251)
modelRF
## Random Forest
##
## 13737 samples
## 53 predictor
## 5 classes: 'A', 'B', 'C', 'D', 'E'
##
## No pre-processing
## Resampling: Cross-Validated (5 fold)
## Summary of sample sizes: 10990, 10988, 10991, 10989, 10990
## Resampling results across tuning parameters:
##
## mtry Accuracy Kappa
## 2 0.9929393 0.9910682
## 27 0.9968699 0.9960406
## 53 0.9948318 0.9934618
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was mtry = 27.
plot(modelRF, log="y")
```



## CONCLUSION

As these models indicate, the performance of the Random Forest (RF) method is superior to that of the Recursive Partitioning and Regression Trees (rpart) method. This is useful information, as recursion is generally known to be slow yet accurate. Our outcome shows that the accuracy of the RF model was 0.993 compared to 0.538 of the rpart Model. The expected out-of-sample error for the RF model is estimated at 0.007, or 0.7%. In contrast, and the recursive rpart Model shows an out-of-sample error rate of 0.462 or 46%. Our outcome is not conclusive, but interesting.

# **PREDICTION**

The following prediction is our research outcome. It is based on the Prediction Model 2 (Random Forest) and is applied against our test data.

```
x <- testing

answers <- predict(modelRF, newdata=x)

answers

## [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
```

Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.