## Prediction Assignment

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## **Executive Summary**

The goal in this work was to develop models to estimate the Classe parameter from the "Human Active Recognition" dataset, which collects vast data on human movement to infer on the quality of exercise, among other variables. Initially we refined the data, removing parameters with >10% blank values, autocorrelated values and near zero variances values. We also removed parameters related to the data provider ID, since these would not contribute to datasets other than the training. We developed a Random Forest model with the remaining parameters to estimate the Classe parameter on the training data, and estimated these values for the test data provided. Our results were uploaded on GitHub on the following link: http://graphics8.nytimes.com/images/2008/04/16/us/0416-nat-subOBAMA.jpg

## Preparing the dataset

```
pml_training <- read.csv('pml-training.csv')
pml_testing <- read.csv('pml-testing.csv')</pre>
```

After loading the huge dataset provided. I removed collums that contained >10% of blanck (NA) values:

```
pml_training[pml_training==""]<-NA

pml_training <- pml_training[, colSums(is.na(pml_training)) < (nrow(pml_training)*0.1)]</pre>
```

After that, I looked for correlated variables

```
pml.cor <- cor(pml_training[,-which(sapply(pml_training, class) == "factor")])
#removing the lower layer
for (i in 2:56) {
    pml.cor[i,1:(i-1)] <- 0
}
#Keeping only those which correlation is <0.8 or >0.8
pml.cor.loc <- which(pml.cor > 0.8 | pml.cor < -0.8,arr.ind = TRUE )

pml.cor.loc <- as.data.frame(pml.cor.loc)</pre>
```

```
#Removing variables that are correlated with themselves (row = collum
pml.cor.loc1 <- data.frame(row=as.integer(), col=as.integer())</pre>
for(i in 1:nrow(pml.cor.loc)){
if (pml.cor.loc$row[i] != pml.cor.loc$col[i]){
  pml.cor.loc1[i,] <- pml.cor.loc[i,]</pre>
} else {
  pml.cor.loc1[i,] <- c('NA', 'NA')</pre>
}
}
#removing rows that are NA
pml.cor.loc1 <- as.data.frame(pml.cor.loc1)</pre>
pml.cor.loc2 <- pml.cor.loc1[pml.cor.loc1 != 'NA', ]</pre>
pml.cor.loc2 <- na.omit(pml.cor.loc2)</pre>
pml.cor.loc2$row <- as.integer(pml.cor.loc2$row)</pre>
pml.cor.loc2$col <- as.integer(pml.cor.loc2$col)</pre>
#Finding the correlations
correlations <- data.frame()</pre>
for (i in 1:nrow(pml.cor.loc2)) {
  correlations[i,1] <- rownames(pml.cor)[pml.cor.loc2$row[i]]</pre>
  correlations[i,2] <- colnames(pml.cor)[pml.cor.loc2$col[i]]</pre>
names.corr <- unique(correlations)</pre>
temp1 <- unique(names.corr$V1)</pre>
temp2 <- unique(names.corr$V2)</pre>
names.corr <- unique(correlations)</pre>
library(dplyr)
pml_training1 <- select(pml_training, -c(temp1, temp2))</pre>
```

This made us remove the following parameters (additional to those removed due to NAs)

```
V2
                 ۷1
          roll_belt
1
                            yaw_belt
2
          roll_belt total_accel_belt
3
         pitch_belt
                        accel_belt_x
4
          roll_belt
                        accel_belt_y
5 total_accel_belt
                        accel_belt_y
6
          roll_belt
                        accel_belt_z
7
  total_accel_belt
                        accel_belt_z
8
       accel_belt_y
                        accel_belt_z
9
         pitch_belt
                       magnet_belt_x
10
       accel_belt_x
                       magnet_belt_x
```

```
11
       gyros_arm_x
                      gyros_arm_y
12
       accel_arm_x
                       magnet_arm_x
13
      magnet_arm_y
                       magnet_arm_z
14 gyros_dumbbell_x gyros_dumbbell_z
15
    pitch_dumbbell accel_dumbbell_x
      yaw dumbbell accel dumbbell z
16
17 gyros_dumbbell_x gyros_forearm_z
18 gyros_dumbbell_z gyros_forearm_z
19 gyros_forearm_y gyros_forearm_z
```

I removed the following variables related to identification of samples, since they will not be useful for testing data

```
[1] "X" "user_name" "raw_timestamp_part_1" "raw_timestamp_part_2" [5] "cvtd_timestamp" "new_window" "num_window"
```

We looked for and removed parameters with near zero variance.

```
library('caret')
zero.var <- nearZeroVar(pml_training, saveMetrics = TRUE, names = TRUE)</pre>
```

However, there was no Near Zero Variance parameters in the dataset with the remnant parameters.

## Generating the model to estimate Classe

We used Random Forets in the package 'ranger' to estimate the Classe values based on the remnant 30 parameters after data processing (of the original 159 parameters).

```
library('ranger')
modelfit <- train(classe ~ ., method = 'ranger', data = pml_training2)
trainingpredict <- predict(modelfit, pml_training2)
table(trainingpredict, pml_training2$classe)</pre>
```

Finally, we included our estimated values for "Classe" parameter in the testing dataset

```
pred_test <- predict(modelfit, pml_testing)
pml_testing$classe <- pred_test
pred_test</pre>
```

```
[1] "B" "A" "B" "A" "A" "A" "E" "D" "B" "A" "A" "B" "C" "B" "A" "E" "E" "A" "B" "B" "B"
```