How Well Exercises Are Done

Hoi Leung

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The main goal for this report is to export how well exercises are being done. Model performance will be based on the accuracy statistics.

Initial Data Exploration

```
## [1] 19622 160
```

Because the number of columns and rows of the data, only half of the rows are used for the analysis to improve run time.

```
train_ind <- sample(1:dim(pml_train_full)[1], dim(pml_train_full)[1]/2, replace=F)
pml_train <- pml_train_full[train_ind, ]
pml_test <- pml_train_full[-train_ind, ]
dim(pml_train)</pre>
```

```
## [1] 9811 160
```

```
table(pml_train$classe)
```

```
##
## A B C D E
## 2772 1951 1697 1596 1795
```

```
#str(pml_train)
```

Upon first inspection, many variables can be converted from string to numbers. However, given the amount of time available to do this analysis, all the non-numeric variables are simply removed. Furthermore, missing values are replaced with the median values.

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```
numeric_var <- sapply(pml_train[, 1:160],is.numeric)
#sum(numeric_var)
numeric_var[1:7] <- FALSE
numeric_var[160] <- TRUE
pml_train <- pml_train[, numeric_var]
pml_test <- pml_test[, numeric_var]

mi <- preProcess(pml_train[, -120], method="medianImpute")
pml_train <- predict(mi, newdata=pml_train)
pml_test <- predict(mi, newdata=pml_test)</pre>
```

Model Training Data

Tree, random forest, and GBM were be used to predict classe. Afterward, another tree is built with the outputs of the previous 3 methods as inputs. Except for random forest, all the models are run with default options. For random forest, to shorten run time, only 100 trees were built.

```
#gbm_fit <- train(classe ~ ., data=pml_train, method="gbm")</pre>
#rf_fit <- train(classe ~ ., data=pml_train, method="rf", ntree = 100)</pre>
#tree_fit <- train(classe ~ ., data=pml_train, method="rpart")</pre>
#save(gbm_fit, file="gbm_fit.RData")
#save(rf_fit, file="rf_fit.RData")
#save(tree_fit, file="tree_fit.RData")
load("gbm_fit.RData"); load("rf_fit.RData"); load("tree_fit.RData")
pml_train_comb <- data.frame(gbm_pred=gbm_fit$train$.outcome,</pre>
                              rf_pred=rf_fit$train$.outcome,
                              tree_pred=tree_fit$train$.outcome,
                              classe=pml_train$classe)
comb_tree_fit <- train(classe ~ ., data=pml_train_comb, method="rpart")</pre>
acc <-
rbind(confusionMatrix(pml_train_comb$gbm_pred, pml_train_comb$classe)$overall[1],
      confusionMatrix(pml_train_comb$rf_pred, pml_train_comb$classe)$overall[1],
      confusionMatrix(pml_train_comb$tree_pred, pml_train_comb$classe)$overall[1],
      confusionMatrix(comb_tree_fit$train$.outcome, pml_train_comb$classe)$overall[1]
)
acc
```

```
## Accuracy
## [1,] 1
## [2,] 1
## [3,] 1
## [4,] 1
```

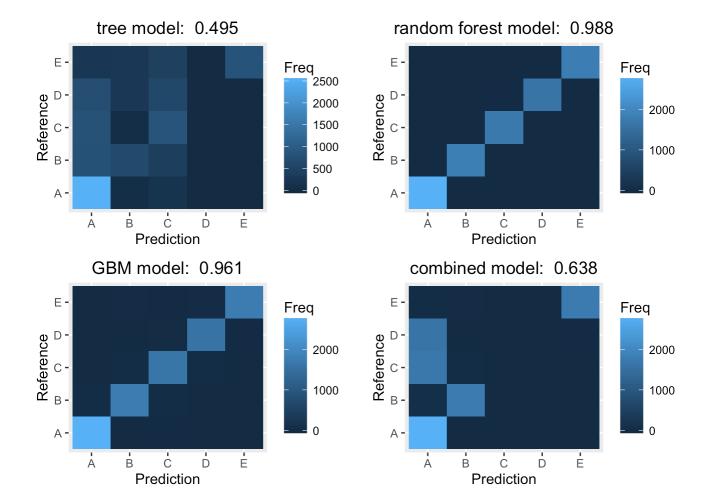
Examine Accuracy on Test Data

The test data were scored with the 4 models. Afterward, the heat maps and their accuracy statistics are plotted.

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```
gbm_pred <- predict(gbm_fit, newdata=pml_test)</pre>
rf_pred <- predict(rf_fit, newdata=pml_test)</pre>
tree_pred <- predict(tree_fit, newdata=pml_test)</pre>
pml_test_comb <- data.frame(gbm_pred=gbm_pred,</pre>
                              rf_pred=rf_pred,
                              tree_pred=tree_pred,
                              classe=pml_test$classe)
comb_tree_pred <- predict(comb_tree_fit, newdata=pml_test_comb)</pre>
gbm_test_accuracy <- confusionMatrix(gbm_pred, pml_test$classe)$overall[1]</pre>
rf test accuracy <- confusionMatrix(rf pred, pml test$classe)$overall[1]
tree_test_accuracy <- confusionMatrix(tree_pred, pml_test$classe)$overall[1]</pre>
combine_test_accuracy <- confusionMatrix(comb_tree_pred, pml_test$classe)$overall[1]</pre>
tree_table <- confusionMatrix(tree_pred, pml_test$classe)$table
rf_table <- confusionMatrix(rf_pred, pml_test$classe)$table</pre>
gbm_table <- confusionMatrix(gbm_pred, pml_test$classe)$table</pre>
comb_table <- confusionMatrix(comb_tree_pred, pml_test$classe)$table</pre>
library(grid)
library(gridExtra)
g1 <- ggplot(data.frame(tree_table))</pre>
g1 <- g1 + geom_tile(aes(x=Prediction, y=Reference, fill=Freq))</pre>
g1 <- g1 + ggtitle(paste("tree model: ", round(tree_test_accuracy, 3)))</pre>
g2 <- ggplot(data.frame(rf_table))</pre>
g2 <- g2 + geom_tile(aes(x=Prediction, y=Reference, fill=Freq))</pre>
g2 <- g2 + ggtitle(paste("random forest model: ", round(rf_test_accuracy, 3)))</pre>
g3 <- ggplot(data.frame(gbm_table))</pre>
g3 <- g3 + geom_tile(aes(x=Prediction, y=Reference, fill=Freq))
g3 <- g3 + ggtitle(paste("GBM model: ", round(gbm_test_accuracy, 3)))</pre>
g4 <- ggplot(data.frame(comb_table))</pre>
g4 <- g4 + geom_tile(aes(x=Prediction, y=Reference, fill=Freq))
g4 <- g4 + ggtitle(paste("combined model: ", round(combine_test_accuracy, 3)))
grid.arrange(g1, g2, g3, g4, ncol = 2, nrow=2)
```

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Conclusion

The random forest model happened to have the highest accuracy statistics in the test sample. Therefore, it will be selected as the final model.

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