Machine Learning Prediction Assignment

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```
Load relevant libraries, set seed
```

```
library(caret)
library(dplyr)
library(DataExplorer)
set.seed(1000)

Read in data

training <- read.csv("pml-training.csv")
testing <- read.csv("pml-testing.csv")</pre>
```

```
Simplify dataset during coding work, set to false for final run
```

```
simple = FALSE
if (simple == TRUE) {
   training <- sample_n(training, 100)
}</pre>
```

Explore data

```
plot_missing(training)
plot_histogram(training)
```

Clean data

```
# First 6 columns identify record, not predictive of classe so remove
training_clean <- training[, -c(1:7)]</pre>
# Identify and remove variables where records are primarily blank - no value
# in imputing variables that are mostly blank
bad <- intersect(which(apply(training_clean, 2, function(x) length(which(x ==
    "#DIV/0!")) > 0)), which(apply(training_clean, 2, function(x) length(unique(x)) <
    5)))
training_clean <- training_clean[-bad]</pre>
# All remaining variables are continuous, but many have been misidentified
# as factor variables. Reclass as numeric
classe <- training_clean$classe</pre>
training_clean <- lapply(training_clean[-79], function(x) as.numeric(as.character(x)))</pre>
training_clean <- as.data.frame(training_clean)</pre>
training_clean$classe = classe
# Some remaining columns are primarily NAs; remove these
completeCols <- which(colSums(is.na(training_clean)) == 0)</pre>
training_clean <- training_clean[completeCols]</pre>
```

Fit models, balancing parameter tuning to maximize resampling without taking prohibatively long

```
# Fit simple tree model with 2 cross-validation iterations
fitControl <- trainControl(method = "cv", number = 2)</pre>
treeFit <- train(classe ~ ., data = training_clean, method = "rpart", trControl = fitControl)</pre>
# Fit bagging models with 10 iterations
bagFit <- train(classe ~ ., data = training_clean, method = "bagEarth", B = 10)</pre>
# Fit xqboost tree model with 50 iterations
parametersGrid <- expand.grid(eta = 0.3, colsample_bytree = 0.6, max_depth = 3,
    nrounds = 50, gamma = 0.5, min_child_weight = 2, subsample = 0.75)
trainX <- training_clean[-53]</pre>
trainY <- training_clean[53]</pre>
xgfit <- train(x = trainX, y = trainY$classe, method = "xgbTree", trControl = fitControl,</pre>
    tuneGrid = parametersGrid)
# Fit random forest model with k-fold cross-validation with k = 10 and 3
fitControl <- trainControl(method = "repeatedcv", number = 10, repeats = 3)</pre>
rffit <- train(x = trainX, y = trainY$classe, method = "rf", trControl = fitControl,
    prox = TRUE)
Print accuracy of each model. Out of sample accuracy is expected to be slightly lower than accuracy of
testing on training sets
treepredicted <- predict(treeFit, training_clean)</pre>
treeMatrix <- confusionMatrix(treepredicted, training_clean$classe)</pre>
bagpredicted <- predict(bagFit, training_clean)</pre>
bagMatrix <- confusionMatrix(bagpredicted, training_clean$classe)</pre>
xgbpredicted <- predict(xgfit, training_clean)</pre>
xgbMatrix <- confusionMatrix(xgbpredicted, training_clean$classe)</pre>
rfpredicted <- predict(rffit, training_clean)</pre>
rfMatrix <- confusionMatrix(rfpredicted, training_clean$classe)</pre>
treeMatrix$overall[1]
## Accuracy
## 0.4955662
bagMatrix$overall[1]
## Accuracy
## 0.7364693
xgbMatrix$overall[1]
## Accuracy
## 0.9726837
rfMatrix$overall[1]
```

```
## Accuracy
## 1
```

Random forest model appears to be the most accurate, so select this one as final model and run on test data set

```
testPredict <- predict(rffit, testing)
print(testPredict)</pre>
```

[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