Ensemble Methods

Importing Data

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```

```
library(randomForest)
library(adabag)
library(xgboost)
library(performanceEstimation)
library(mltools)

train <- read.csv("C:/Users/jah200003/Downloads/training.csv", header=TRUE, stringsAsFactors=TRUE)
test <- read.csv("C:/Users/jah200003/Downloads/testing.csv", header=TRUE, stringsAsFactors=TRUE)</pre>
```

Data Cleaning

Data is already prepared fairly well, but the classes are very imbalanced Applied synthetic oversampling and undersampling to improve balance before SVM (Copied from classification notebook)

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```
table(train$class)
```

farm	forest	grass imp	ervious	orchard	water
1441	7431	446	969	53	205

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```
train <- smote(class~ ., train, perc.over = 50, perc.under = 5)
table(train$class)</pre>
```

```
farm forest grass impervious orchard water
1777 9470 555 1173 2703 275
```

Random Forest

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```
rf <- randomForest(class~ ., data = train, importance = TRUE)

predrf <- predict(rf, newdata = test, type = "response")
accrf <- mean(predrf == test$class)
mccrf <- mcc(factor(predrf), test$class)

print(paste("Accuracy: ", accrf))</pre>
```

[1] "Accuracy: 0.6633333333333333"

```
print(paste("MCC: ", mccrf))
```

```
[1] "MCC: 0.604621407160539"
```

Random forest performed the best of all three ensemble techniques, likely due to the large amount of predictors and the relative lack of importance for most of them. For some types of land, like intact forests and open water, the vegetation index will not change much over time, causing the chronological indexes to potentially do more harm than good. None of the algorithms had an exceptionally high accuracy, but given the fact that there is a 1-in-6 chance of correctly guessing the type of land, hovering around 50% would be great in an even distribution, though the dominance of forest land in the training data threatens that idea. The runtime for random forest was about 45 seconds, being neither the slowest nor the fastest algorithm.

```
Ada Boost
                                                                                                               Hide
 adab <- boosting(class~ ., data = train, boos = TRUE, coeflearn = "Breiman")</pre>
 summary(adab)
             Length Class
                             Mode
 formula
                 3 formula call
 trees
               100
                    -none-
                            list
 weights
               100
                    -none-
                            numeric
 votes
             95718
                    -none-
                            numeric
 prob
             95718
                            numeric
                    -none-
 class
             15953
                    -none-
                            character
 importance
                    -none-
                            numeric
                28
                             call
 terms
                 3
                    terms
                    -none-
                            call
 call
                                                                                                               Hide
 predada <- predict(adab, newdata = test, type = "response")</pre>
 accada <- mean(predada$class == test$class)</pre>
 mccada <- mcc(factor(predada$class), test$class)</pre>
 print(paste("Accuracy: ", accada))
 [1] "Accuracy: 0.6"
                                                                                                               Hide
 print(paste("MCC: ", mccada))
```

On paper, this algorithm should have done very well because it works well with weak classifiers and nonbinary identification, but the algorithm was outdone by random forests. On top of that, this was the slowest algorithm, taking roughly a minute to go through the same number of iterations. The lower accuracy may be due to outliers or imbalance in the data set.

[1] "MCC: 0.520903676908143"

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```
train_labels <- sapply(as.numeric(train$class), function(x) x - 1)
train_matrix <- data.matrix(train[ , -29])
xgmodel <- xgboost(data = train_matrix, label = train_labels, verbose = 0, nrounds = 100, num_class = 6, ob
jective = "multi:softmax")

test_labels <- sapply(as.numeric(test$class), function(x) x - 1)
test_matrix <- data.matrix(test[ , -29])

probs <- predict(xgmodel, test_matrix)
predxg <- ifelse(probs > 0.5, 1, 0)

accxg <- mean(predxg == test_labels)
mccxg <- mcc(predxg, test_labels)

print(paste("Accuracy: ", accxg))</pre>
```

```
[1] "Accuracy: 0.4366666666667"
```

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```
print(paste("MCC: ", mccxg))
```

```
[1] "MCC: 0.391630713194483"
```

This algorithm performed the worst out of the three algorithms, which may be caused by trees that are too deep and are overfitting the data set. It is also certainly possible that I am doing something wrong, as the given documentation pertained to binary classification, but this land data set has six different classes to select from. On the bright side, this algorithm is blazingly fast and took fewer than 5 seconds.