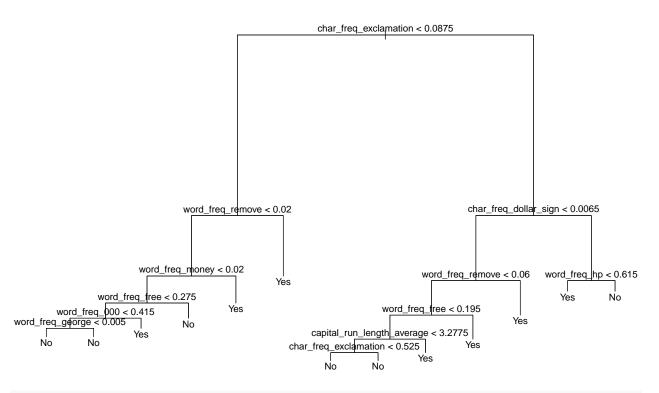
## STATS 415 - Homework 8 - Classification Trees

Marian L. Schmidt | April 1, 2016

(a) Fit a classification tree using only the training set. Find the percentage of emails in the test set that were misclassified by your optimal tree. Of all the spam emails of the test set what percentage was misclassified and of all the non-spam emails of the test set what percentage was misclassified?

```
attach(train_data)
spam_status <- ifelse(spam_yes == 1,"Yes","No")
train_data <- data.frame(train_data, spam_status)
# Fit the classification tree
tree_spam <- tree(spam_status~.-spam_yes, data = train_data)
plot(tree_spam) # Plot the tree
text(tree_spam, pretty = 0) # add the predictor names and cutoffs</pre>
```



summary(tree\_spam) # The training error rate is 8.77%

```
## Residual mean deviance: 0.4927 = 1505 / 3054
## Misclassification error rate: 0.08771 = 269 / 3067

set.seed(111)
spam_prediction <- predict(tree_spam, test_data, type="class")
yes_test <- ifelse(spam_yes == 1,"Yes","No")
table(spam_prediction,yes_test)

## yes_test
## spam_prediction No Yes
## No 854 78
## Yes 62 540</pre>
(854+540)/1534
```

## [1] 0.9087353

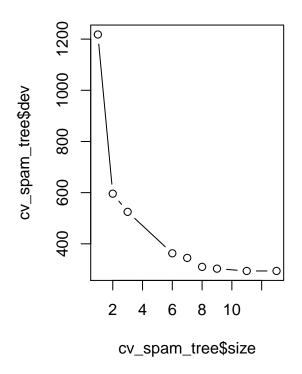
90.8735332% of the data was classified correctly, therefore the misclassification error is 9.1264668%. There were 540 spam e-mails classified by the above classification tree model, however, in reality 618 of the test emails were spam. Therefore, the misclassification of spam e-mails is 87.3786408%. On the other hand, there were 854 e-mails classified as non-spam when in reality, there were 916 non-spam e-mails in the test data. Thus, the misclassification of the non-spam e-mails is 93.231441%

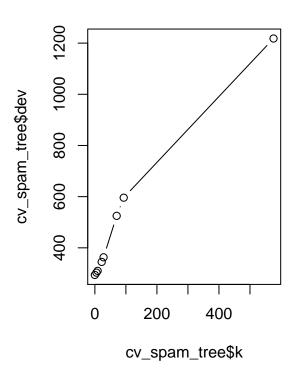
(b) Plot a subtree of the optimal tree that has at most 8 terminal nodes. What are some of the variables that were used in tree construction?

```
cv_spam_tree <- cv.tree(tree_spam, FUN = prune.misclass)
names(cv_spam_tree)

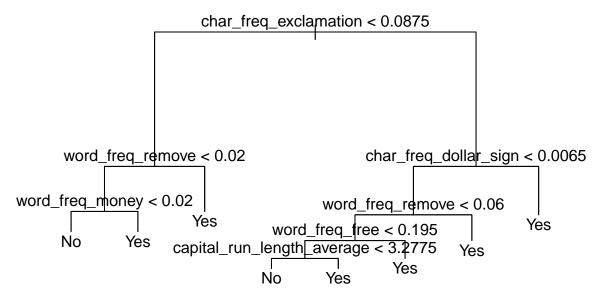
## [1] "size" "dev" "k" "method"

par(mfrow = c(1,2)) # plot it!
plot(cv_spam_tree$size, cv_spam_tree$dev, type = "b")
plot(cv_spam_tree$k, cv_spam_tree$dev, type = "b")</pre>
```





```
# Time to prune the tree to 8 nodes
prune_spam_tree <- prune.misclass(tree_spam, best = 8)
par(mfrow = c(1,1))
plot(prune_spam_tree)
text(prune_spam_tree, pretty = 0)</pre>
```



```
pred_spam_tree <- predict(prune_spam_tree, test_data, type="class")
table(pred_spam_tree, yes_test)</pre>
```

```
## yes_test
## pred_spam_tree No Yes
## No 854 87
## Yes 62 531
```

```
(854+531)/1534
```

## ## [1] 0.9028683

The variables that were used in the pruned tree with 8 nodes are char\_freq\_exclamation, word\_freq\_remove, char\_freq\_dollar\_sign, word\_freq\_remove, word\_freq\_money, word\_freq\_free, and capitol\_run\_length\_average. These are many of the variables that were used in the larger tree.

(c) Try (a) again using Random Forest. Use the "importance()" function to determine which variables are most important. Describe the effect of m, the number of variables considered at each split, on the error rate obtained.

```
## Including 8 variables
rf_spam_8 <- randomForest(spam_status~.-spam_yes, data=train_data, mtry=8, importance = TRUE)
yhat_rf_8 <- predict(rf_spam_8, newdata = test_data)
yhat_rf_8_num <- as.numeric(ifelse(yhat_rf_8 == "Yes","1","0"))
mean((yhat_rf_8_num-test_data$spam_yes)^2)</pre>
```

## [1] 0.02281617

```
head(importance(rf_spam_8), n = 5)
```

```
##
                                    Yes MeanDecreaseAccuracy
                           No
## word_freq_make
                    10.910402 11.076553
                                                   14.327894
## word_freq_address 11.435203 9.745039
                                                   12.837664
## word freq all 14.661538 16.281953
                                                   18.621406
## word_freq_3d
                    6.235549 6.902190
                                                    8.413608
                    22.305054 24.657604
## word_freq_our
                                                   26.005484
                    MeanDecreaseGini
##
## word_freq_make
                            4.862249
## word_freq_address
                            6.110087
## word_freq_all
                          14.303099
## word_freq_3d
                            2.061795
## word_freq_our
                           55.104642
```

## # Including 12 variables

```
rf_spam_12 <- randomForest(spam_status~.-spam_yes, data=train_data, mtry=12, importance = TRUE)
yhat_rf <- predict(rf_spam_12, newdata = test_data)
yhat_rf_12_num <- as.numeric(ifelse(yhat_rf == "Yes","1","0"))
mean((yhat_rf_12_num-test_data$spam_yes)^2)</pre>
```

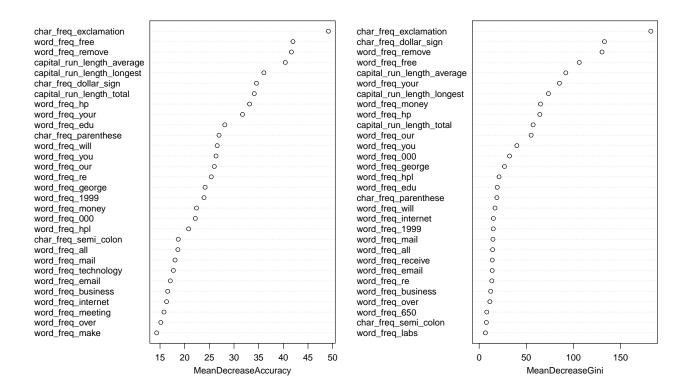
## [1] 0.02477184

```
head(importance(rf_spam_12), n = 5)
```

```
## No Yes MeanDecreaseAccuracy
## word_freq_make 10.470837 10.976760 14.31353
## word_freq_address 12.502835 9.258132 13.93724
## word_freq_all 12.809631 14.483414 15.96055
## word freq 3d 8.527507 6.475422 9.28751
```

```
## word freq our
                     27.373639 25.112105
                                                      30.54497
##
                     MeanDecreaseGini
                             4.175492
## word freq make
## word_freq_address
                             5.393501
## word freq all
                            12.362576
## word freq 3d
                             2.921882
## word freq our
                            46.691482
#Plot most important variables for each model
varImpPlot(rf_spam_8)
```

rf spam 8



For the 8 variable model the top 5 predictors, in order, are char\_freq\_exclamation, capital\_run\_length\_average, word\_freq\_remove, word\_freq\_free and capital\_run\_length\_longest while for the 12 variable model the top 5 predictors are char\_freq\_exclamation, word\_freq\_remove, word\_freq\_free, capital\_run\_length\_average, and char\_freq\_dollar\_sign. The model with fewer variables has a lower MSE as reported in the code above.