Practical Machine Learning Course Project

Summary

Two modeling methods were utilized for this project, Classification Tree and Random Forest. For both methods, 5-fold cross validation is used.

Load Packages

```
library (caret)
## Warning: package 'caret' was built under R version 3.4.4
## Loading required package: lattice
## Loading required package: ggplot2
## Warning: package 'ggplot2' was built under R version 3.4.4
library (randomForest)
## Warning: package 'randomForest' was built under R version 3.4.4
## randomForest 4.6-14
## Type rfNews() to see new features/changes/bug fixes.
## Attaching package: 'randomForest'
## The following object is masked from 'package:ggplot2':
##
##
     margin
library (rattle)
## Warning: package 'rattle' was built under R version 3.4.4
## Rattle: A free graphical interface for data science with R.
## Version 5.1.0 Copyright (c) 2006-2017 Togaware Pty Ltd.
## Type 'rattle()' to shake, rattle, and roll your data.
##
## Attaching package: 'rattle'
```

```
## The following object is masked from 'package:randomForest':
##
## importance

library(rpart.plot)

## Warning: package 'rpart.plot' was built under R version 3.4.4

## Loading required package: rpart

library(AppliedPredictiveModeling)

## Warning: package 'AppliedPredictiveModeling' was built under R version
## 3.4.4
```

Import Data

It is assumed that data files are located in working directory

```
training <- read.csv("pml-training.csv", na.strings=c("NA",""), header=TRUE)
colnames_train <- colnames(training)
testing <- read.csv("pml-testing.csv", na.strings=c("NA",""), header=TRUE)
colnames_test <- colnames(testing)</pre>
```

Remove columns that aren't needed. NA data is cleaned, as well.

```
training <- training[, colSums(is.na(training)) == 0]
testing <- testing[, colSums(is.na(testing)) == 0]
training <- training[, -c(1:7)]
testing <- testing[, -c(1:7)]</pre>
```

Training data is separated into training and test sets

```
set.seed(1111)
ids_small <- createDataPartition(y=training$classe, p=0.25, list=FALSE)
small <- training[ids_small,]
remainder <- training[-ids_small,]

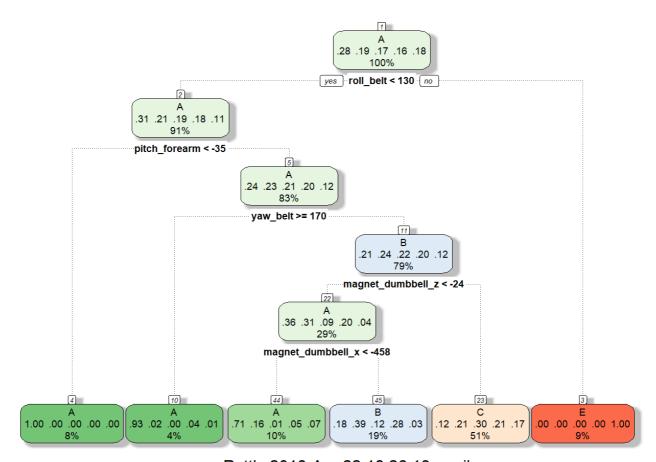
set.seed(1111)
Train <- createDataPartition(y=small$classe, p=0.6, list=FALSE)
small_training <- small[Train,]
small_testing <- small[-Train,]</pre>
```

Classification Tree

```
set.seed(1111)
control <- trainControl(method = "cv", number = 5)
fit_rpart <- train(classe ~ ., data = small_training, method = "rpart", trControl = control)
print(fit_rpart, digits=3)</pre>
```

```
## CART
##
## 2946 samples
## 52 predictor
    5 classes: 'A', 'B', 'C', 'D', 'E'
##
##
## No pre-processing
## Resampling: Cross-Validated (5 fold)
## Summary of sample sizes: 2356, 2358, 2357, 2356, 2357
## Resampling results across tuning parameters:
##
##
   ср
          Accuracy Kappa
## 0.0365 0.537 0.4130
   0.0408 0.516
                     0.3873
##
  0.1214 0.334
                    0.0753
##
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was cp = 0.0365.
```

```
fancyRpartPlot(fit rpart$finalModel)
```



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Classification Tree Prediction

```
predict_rpart <- predict(fit_rpart, newdata=small_testing)
print(confusionMatrix(predict_rpart, small_testing$classe), digits=4)</pre>
```

```
## Confusion Matrix and Statistics
##
##
          Reference
## Prediction A B C D
         A 357 47 3 12 17
          в 76 108 47 116 17
##
         C 125 225 292 193 148
##
         D 0 0 0 0 0
##
##
         E 0 0 0 0 178
##
## Overall Statistics
##
##
               Accuracy: 0.4768
##
                 95% CI: (0.4545, 0.4992)
##
    No Information Rate: 0.2845
##
    P-Value [Acc > NIR] : < 2.2e-16
##
##
                  Kappa: 0.3433
## Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##
                    Class: A Class: B Class: C Class: D Class: E
                      0.6398 0.28421 0.8538 0.0000 0.49444
## Sensitivity
                      0.9437 0.83808 0.5732 1.0000 1.00000
## Specificity
## Pos Pred Value
                     0.8188 0.29670 0.2970 NaN 1.00000
## Neg Pred Value
                     0.8682 0.82968 0.9489 0.8363 0.89792
## Prevalence
                     0.2845 0.19378 0.1744 0.1637 0.18358
## Detection Rate 0.1820 0.05507 0.1489 0.0000 0.09077
## Detection Prevalence 0.2223 0.18562 0.5013 0.0000 0.09077
## Balanced Accuracy 0.7917 0.56114 0.7135 0.5000 0.74722
```

CT Accuracy 0.4768

Random Forest

```
set.seed(1111)
fit_rf <- train(classe ~ ., data = small_training, method = "rf", trControl = contr
ol)
print(fit_rf, digits = 4)</pre>
```

```
## Random Forest
##
## 2946 samples
   52 predictor
##
    5 classes: 'A', 'B', 'C', 'D', 'E'
##
##
## No pre-processing
## Resampling: Cross-Validated (5 fold)
## Summary of sample sizes: 2356, 2358, 2357, 2356, 2357
## Resampling results across tuning parameters:
##
##
  mtry Accuracy Kappa
## 2 0.9569 0.9454
## 27 0.9603 0.9497
  52 0.9525 0.9399
##
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was mtry = 27.
```

Random Forest Prediction

```
predict_rf <- predict(fit_rf, newdata=small_testing)
print(confusionMatrix(predict_rf, small_testing$classe), digits=4)</pre>
```

```
## Confusion Matrix and Statistics
##
##
         Reference
## Prediction A B C D E
##
     A 556 22 0 1 0
         в 0 351 5 0 1
##
         C 0 7 334 9 1
##
##
         D 2 0 3 311 7
         E 0 0 0 0 351
##
##
## Overall Statistics
##
##
               Accuracy: 0.9704
                95% CI: (0.9619, 0.9775)
##
##
    No Information Rate: 0.2845
##
    P-Value [Acc > NIR] : < 2.2e-16
##
##
                 Kappa: 0.9625
## Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##
                   Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                    0.9964 0.9237 0.9766 0.9688 0.9750
## Specificity
                     0.9836 0.9962 0.9895 0.9927 1.0000
                    0.9603 0.9832 0.9516 0.9628 1.0000
## Pos Pred Value
## Neg Pred Value
                    0.9986 0.9819 0.9950 0.9939 0.9944
                    0.2845 0.1938 0.1744 0.1637 0.1836
## Prevalence
## Detection Rate 0.2835 0.1790 0.1703 0.1586 0.1790
## Detection Prevalence 0.2953 0.1820 0.1790 0.1647 0.1790
## Balanced Accuracy 0.9900 0.9599 0.9831 0.9808 0.9875
```

RF Accuracy 0.9704

After a head to head comparison, Random Forest method (0.9704) proved to be more accurate than Classification Tree method (0.4768).

Test set Prediction

```
prediction <- predict(fit_rf, newdata=testing)
prediction</pre>
```

```
## [1] BABAAEDBAABCBAEEABBB
## Levels: ABCDE
```