Predicting Exercise Method

Friday, March 20, 2015

Overview

The goal of this project is to predict the manner in which the people have done the exercise after training a model using the training data.

Read in the data

We first download and read in the data.

```
library(caret)

train_url = "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.c
sv"

#download.file(train_url,destfile=".//pml-training.csv")

test_url = "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.cs
v"

#download.file(test_url,destfile=".//pml-testing.csv")

train = read.csv(".//pml-training.csv")

test = read.csv(".//pml-testing.csv")
```

Data Cleansing

To start with there are a possible 159 predictor variables in the dataset. On exploring the data it was found that there are a lot of variables with NA values and also lot of variables with mostly spaces. The following code identifies such variables and removes them from the dataset thus reducing the dimensions. After removing the sparse columns we get 52 predictors.

```
#Check nr. of predictors
colnames = names(train)

#Detect columns with NA's
x=c()
n = length(colnames)
for (i in 1:n)
{
   t = summary(train[,i])
   m = t["NA's"]
   if (!is.na(m)) {print(paste(i,"-", colnames[i], " NAs =", m)); x=c(x,i)}
}
```

```
## [1] "18 - max roll belt NAs = 19216"
## [1] "19 - max picth belt NAs = 19216"
## [1] "21 - min roll belt NAs = 19216"
## [1] "22 - min pitch belt NAs = 19216"
## [1] "24 - amplitude roll belt NAs = 19216"
## [1] "25 - amplitude pitch belt NAs = 19216"
## [1] "27 - var total accel belt NAs = 19216"
## [1] "28 - avg roll belt NAs = 19216"
## [1] "29 - stddev roll belt NAs = 19216"
## [1] "30 - var roll belt NAs = 19216"
## [1] "31 - avg pitch belt NAs = 19216"
## [1] "32 - stddev pitch belt NAs = 19216"
## [1] "33 - var pitch belt NAs = 19216"
## [1] "34 - avg yaw belt NAs = 19216"
## [1] "35 - stddev yaw belt NAs = 19216"
## [1] "36 - var yaw belt NAs = 19216"
## [1] "50 - var accel arm NAs = 19216"
## [1] "51 - avg roll arm NAs = 19216"
## [1] "52 - stddev roll arm NAs = 19216"
## [1] "53 - var roll arm NAs = 19216"
## [1] "54 - avg_pitch_arm NAs = 19216"
## [1] "55 - stddev pitch arm NAs = 19216"
## [1] "56 - var pitch arm NAs = 19216"
## [1] "57 - avg yaw arm NAs = 19216"
## [1] "58 - stddev yaw arm NAs = 19216"
## [1] "59 - var yaw arm NAs = 19216"
\#\# [1] "75 - max roll arm NAs = 19216"
\#\# [1] "76 - max picth arm NAs = 19216"
## [1] "77 - max yaw arm NAs = 19216"
## [1] "78 - min roll arm NAs = 19216"
## [1] "79 - min pitch arm NAs = 19216"
## [1] "80 - min yaw arm NAs = 19216"
## [1] "81 - amplitude roll arm NAs = 19216"
## [1] "82 - amplitude pitch arm NAs = 19216"
## [1] "83 - amplitude yaw arm NAs = 19216"
\#\# [1] "93 - max roll dumbbell NAs = 19216"
## [1] "94 - max_picth dumbbell NAs = 19216"
## [1] "96 - min roll dumbbell NAs = 19216"
## [1] "97 - min pitch dumbbell NAs = 19216"
## [1] "99 - amplitude roll dumbbell NAs = 19216"
\#\# [1] "100 - amplitude pitch dumbbell NAs = 19216"
## [1] "103 - var accel dumbbell NAs = 19216"
## [1] "104 - avg_roll dumbbell NAs = 19216"
## [1] "105 - stddev roll dumbbell NAs = 19216"
\#\# [1] "106 - var roll dumbbell NAs = 19216"
## [1] "107 - avg pitch dumbbell NAs = 19216"
## [1] "108 - stddev pitch dumbbell NAs = 19216"
## [1] "109 - var pitch dumbbell NAs = 19216"
```

```
## [1] "110 - avg yaw dumbbell NAs = 19216"
## [1] "111 - stddev yaw dumbbell NAs = 19216"
## [1] "112 - var yaw dumbbell NAs = 19216"
## [1] "131 - max roll forearm NAs = 19216"
\#\# [1] "132 - max picth forearm NAs = 19216"
## [1] "134 - min roll forearm NAs = 19216"
## [1] "135 - min pitch forearm NAs = 19216"
## [1] "137 - amplitude roll forearm NAs = 19216"
## [1] "138 - amplitude pitch forearm NAs = 19216"
## [1] "141 - var accel forearm NAs = 19216"
## [1] "142 - avg roll forearm NAs = 19216"
\#\# [1] "143 - stddev roll forearm NAs = 19216"
\#\# [1] "144 - var roll forearm NAs = 19216"
\#\# [1] "145 - avg pitch forearm NAs = 19216"
## [1] "146 - stddev pitch forearm NAs = 19216"
\#\# [1] "147 - var pitch forearm NAs = 19216"
## [1] "148 - avg yaw forearm NAs = 19216"
## [1] "149 - stddev yaw forearm NAs = 19216"
\#\# [1] "150 - var yaw forearm NAs = 19216"
```

```
#Since all of the columns containing NA's are sparce(19000+ records in each con
tain NA), remove them from the dataset
train=train[,-c(1:7,x)]

test=test[,-c(1:7,x)]

#Detect columns with spaces
colnames=names(train)
x=c()
n = length(colnames)
for (i in 1:n)
{
    t = as.character(train[,i])
    m = which(t == "")
    if (length(m)!=0) { x=c(x,i); print(paste(i,"-", colnames[i], " Spaces =", le
ngth(m)))}
}
```

```
## [1] "5 - kurtosis roll belt Spaces = 19216"
## [1] "6 - kurtosis picth belt Spaces = 19216"
## [1] "7 - kurtosis yaw belt Spaces = 19216"
## [1] "8 - skewness roll belt Spaces = 19216"
## [1] "9 - skewness roll belt.1 Spaces = 19216"
## [1] "10 - skewness yaw belt Spaces = 19216"
## [1] "11 - max yaw belt Spaces = 19216"
## [1] "12 - min yaw belt Spaces = 19216"
## [1] "13 - amplitude yaw belt Spaces = 19216"
## [1] "36 - kurtosis roll arm Spaces = 19216"
## [1] "37 - kurtosis picth arm Spaces = 19216"
## [1] "38 - kurtosis yaw arm Spaces = 19216"
## [1] "39 - skewness roll arm Spaces = 19216"
## [1] "40 - skewness_pitch_arm Spaces = 19216"
## [1] "41 - skewness yaw arm Spaces = 19216"
## [1] "45 - kurtosis roll dumbbell Spaces = 19216"
## [1] "46 - kurtosis picth dumbbell Spaces = 19216"
## [1] "47 - kurtosis yaw dumbbell Spaces = 19216"
## [1] "48 - skewness roll dumbbell Spaces = 19216"
## [1] "49 - skewness pitch dumbbell Spaces = 19216"
## [1] "50 - skewness yaw dumbbell Spaces = 19216"
## [1] "51 - max yaw dumbbell Spaces = 19216"
## [1] "52 - min yaw dumbbell Spaces = 19216"
## [1] "53 - amplitude yaw dumbbell Spaces = 19216"
## [1] "67 - kurtosis roll forearm Spaces = 19216"
## [1] "68 - kurtosis picth forearm Spaces = 19216"
## [1] "69 - kurtosis yaw forearm Spaces = 19216"
## [1] "70 - skewness roll forearm Spaces = 19216"
## [1] "71 - skewness pitch forearm Spaces = 19216"
## [1] "72 - skewness yaw forearm Spaces = 19216"
## [1] "73 - max yaw forearm Spaces = 19216"
## [1] "74 - min yaw forearm Spaces = 19216"
## [1] "75 - amplitude yaw forearm Spaces = 19216"
```

```
#Since all of the columns containing spaces are sparce(19000+ records in each c ontain spaces), remove them from the dataset train=train[,-c(x)] test=test[,-c(x)]
```

Model Creation

The following models were tried and their findings are explained in the sections below:

Cartesian Tree

Cartesian Tree on the entire dataset without any pre-processing gave poor predictive accuracy. Hence was rejected.

```
modFit = train(classe~., method="rpart", data=train)
## Loading required package: rpart
confusionMatrix(train$classe,predict(modFit,newdata=train[,-53]))
## Confusion Matrix and Statistics
##
##
           Reference
## Prediction A B C D
                                Ε
##
         A 5080 81 405 0 14
##
         В 1581 1286 930 0 0
##
          C 1587 108 1727
                            0
                                0
##
         D 1449 568 1199 0 0
         E 524 486 966 0 1631
##
##
## Overall Statistics
##
##
                Accuracy: 0.4956
                  95% CI: (0.4885, 0.5026)
##
##
   No Information Rate: 0.5209
     P-Value [Acc > NIR] : 1
##
##
##
                   Kappa : 0.3407
```

```
## Sensitivity 0.4970 0.50850 0.33040 NA 0.99149
## Specificity 0.9468 0.85310 0.88225 0.8361 0.89008
## Pos Pred Value 0.9104 0.33869 0.50468 NA 0.45218
## Neg Pred Value 0.6339 0.92145 0.78395 NA 0.99913
## Prevalence 0.5209 0.12889 0.26638 0.0000 0.08383
## Detection Rate 0.2589 0.06554 0.08801 0.0000 0.08312
## Detection Prevalence 0.2844 0.19351 0.17440 0.1639 0.18382
## Balanced Accuracy 0.7219 0.68080 0.60633 NA 0.94079
```

Pre-processing using PCA

Statistics by Class:

Mcnemar's Test P-Value : NA

##

##

##

Principal Component Analysis identified Principal components, but a tree model created using the PCA predictors had very poor predictive accuracy. PCA was tried at 100%, 99% and 95% threshold levels. But all gave poor predictive accuracy and hence this effort was rejected.

Class: A Class: B Class: C Class: D Class: E

```
prComp = prcomp(train[,-53])
preProc = preProcess(train[,-53],method="pca", pcaComp=2)
preProc
```

```
##
## Call:
## preProcess.default(x = train[, -53], method = "pca", pcaComp = 2)
##
## Created from 19622 samples and 52 variables
## Pre-processing: principal component signal extraction, scaled, centered
##
## PCA used 2 components as specified.
```

```
trainPC = predict(preProc, train[,-53])
trainPC = cbind(trainPC, classe=train[,53])
modFit = train(classe~., method="rpart", data=trainPC)
confusionMatrix(trainPC$classe, predict(modFit, newdata=trainPC[,-6]))
```

```
## Confusion Matrix and Statistics
##
##
              Reference
## Prediction A B
                           С
                                   D
            A 3405 464 291 646 774
##
             B 1295 822 289 475 916
             C 1349 453 471 418 731
##
##
            D 514 577 76 1230 819
##
            E 706 382 190 695 1634
##
## Overall Statistics
##
##
                    Accuracy: 0.3854
##
                      95% CI: (0.3786, 0.3922)
      No Information Rate: 0.3705
##
##
      P-Value [Acc > NIR] : 8.053e-06
##
##
                        Kappa: 0.2138
  Mcnemar's Test P-Value : < 2.2e-16
##
##
## Statistics by Class:
##
##
                           Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                           0.4684 0.30467 0.35763 0.35508 0.33525
## Specificity
                           0.8239 0.82421 0.83879 0.87709 0.86622

      0.6102
      0.21649
      0.13764
      0.38246
      0.45301

      0.7248
      0.88145
      0.94778
      0.86383
      0.79769

      0.3705
      0.13750
      0.06712
      0.17654
      0.24839

## Pos Pred Value
## Neg Pred Value
## Prevalence
## Detection Rate 0.1735 0.04189 0.02400 0.06268 0.08327
## Detection Prevalence 0.2844 0.19351 0.17440 0.16390 0.18382
## Balanced Accuracy 0.6462 0.56444 0.59821 0.61608 0.60073
```

Random Forest

Random Forest was finally selected since it gave a predictive accurancy of 97%. Random Forest could not run successfully on my laptop for the entire data. It would error out after a while citing memory allocation issues. It ran successfully for 60% of the training data. The remainder of the training data was used to test the model. Models were created using ntree=5 and 10 since anything above 10 would take more than 1 hr and the laptop would get scaringly hot. With ntree=10, the model gave an accuracy rate around 97% and was accepted since I could not try for better values due to laptop limitations.

```
#Splitting the training set into training and testing
inTrain = createDataPartition(y=train$classe, p=0.6, list=FALSE)
training = train[inTrain,]
testing = train[-inTrain,]

#Creating a random forest. Not running this again while report generation sinc
e it takes a long time. Just executing the final model in the section below
#modFit = train(classe~., method="rf", data=training, ntree=10, prox=T)
#Checking the confusion matrix
#confusionMatrix(training$classe,predict(modFit,newdata=training[,-53]))
```

Crossvalidation

Default Bootstrap reSampling with 25 repeats provided a good accuracy rate of 97.7%.

5 fold crossvalidation with 5 repeats provided and accurance of 98.56%.

10 fold cross-validation with 10 repeats provided and accurance of 98.64%. Submitted the predictions using this model because it provided the highest accuracy.

Although the 10 fold cross-validation model did not provide a significant improvement in accuracy over the 5 fold model compared to the time it took to get generated, it gave a better accuracy compared to the bootstrap resampling cross-validation.

```
#During report generation am running a 5 fold crossvalidation model since 10-fo
ld model takes toooo long. The actual prediction was done with 10 fold cross-va
lidation model.
fitControl <- trainControl(## 5-fold CV
  method = "repeatedcv",
  number = 5,
  ## repeated 5 times
  repeats = 5)
modFit = train(classe~., method="rf", data=training, trControl = fitControl, nt
ree=10, prox=T)</pre>
```

```
confusionMatrix(testing$classe,predict(modFit,newdata=testing[,-53]))
```

```
## Confusion Matrix and Statistics
##
##
            Reference
## Prediction A B
                          С
          A 2221
                    6
           B 21 1482 14
                              1
##
           C 0 18 1344 6
##
                 0 1 19 1264 2
##
          D
##
                 0 5 8 4 1425
##
## Overall Statistics
##
##
                  Accuracy: 0.986
##
                    95% CI: (0.9831, 0.9885)
     No Information Rate: 0.2858
##
##
      P-Value [Acc > NIR] : < 2.2e-16
##
##
                     Kappa: 0.9823
## Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##
                        Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                        0.9906 0.9802 0.9704 0.9890 0.9972
## Specificity
                        0.9980 0.9943 0.9963 0.9967 0.9974

      0.9951
      0.9763
      0.9825
      0.9829
      0.9882

      0.9963
      0.9953
      0.9937
      0.9979
      0.9994

## Pos Pred Value
## Neg Pred Value
## Prevalence
                        0.2858 0.1927 0.1765 0.1629 0.1821
## Detection Rate 0.2831 0.1889 0.1713 0.1611 0.1816
## Detection Prevalence 0.2845 0.1935 0.1744 0.1639 0.1838
## Balanced Accuracy 0.9943 0.9872 0.9833 0.9928 0.9973
```

Expected Out of Sample Error

I expect the out of sample error to be less than 3% i.e on a sample of 20, I expect less than 1 to be wrong.FYI:I got 1 wrong when I predicted in first run. But in the 2nd run without any change of parameters it predicted correctly for the one that it had got wrong earlier.

Predicted Output

The predicted output is as follows:

```
predict(modFit, newdata=test[,-53])
```

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