Practical Machine Learning - Submission

Objective

The goal of this work is to predict from data collected with quantified self devices if people are performing barbell lifts correctly or incorrectly.

Librairies & seed

```
library(caret)

## Warning: package 'caret' was built under R version 3.1.2

## Loading required package: lattice
## Loading required package: ggplot2

set.seed(1234)
```

Collect data and build data sets

```
url_csv_train="http://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv"
url_csv_test="http://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv"
raw_data_train <- read.csv(url_csv_train)
raw_data_test <- read.csv(url_csv_test)

inTrain <- createDataPartition(raw_data_train$classe, p = 0.7, list = FALSE)
training <- raw_data_train[ inTrain,]
testing <- raw_data_train[-inTrain,]</pre>
```

Features

By looking at the data (using str), we can see that a majority of variables are filled with a large majority of NA, or are empty. We decide to eliminate: - these variables with a large majority of NA or empty values - the first columns (X, user_name, raw_timestamp_part_1, raw_timestamp_part_2, cvtd_timestamp, new_window, num_window) which may not give information on how well the exercise is performed

```
features=c("roll_belt","pitch_belt","yaw_belt","total_accel_belt","gyros_belt_x","gyros_belt_y","g
yros_belt_z","accel_belt_x","accel_belt_y","accel_belt_z","magnet_belt_x","magnet_belt_y","magnet_
belt_z","roll_arm","pitch_arm","yaw_arm","total_accel_arm","gyros_arm_x","gyros_arm_y","gyros_arm_
z","accel_arm_x","accel_arm_y","accel_arm_z","magnet_arm_x","magnet_arm_y","magnet_arm_z","roll_du
mbbell","pitch_dumbbell","yaw_dumbbell","total_accel_dumbbell","gyros_dumbbell_x","gyros_dumbbell_
y","gyros_dumbbell_z","accel_dumbbell_x","accel_dumbbell_y","accel_dumbbell_z","magnet_dumbbell_x"
,"magnet_dumbbell_y","magnet_dumbbell_z","roll_forearm","pitch_forearm","yaw_forearm","total_accel
_forearm","gyros_forearm_x","gyros_forearm_y","gyros_forearm_z","accel_forearm_x","accel_forearm_y
","accel_forearm_z","magnet_forearm_x","magnet_forearm_y","magnet_forearm_z")
```

At the end, we keep the variables which seems to be the measures: - roll - pitch - yaw - total accel - gyros x - gyros y - gyros z - accel x - accel y - accel z - magnet x - magnet y - magnet z

for different parts/captors: - belt - arm - dumbbell - forearm

Model

We choose "randomForest" algorithm and to perform 5-fold cross-validation.

```
control1 <- trainControl(method = "cv", number = 5, allowParallel = TRUE)
modelFit <- train(classe~.,data=training[,append(features,"classe")],trControl = control1,method="
rf")</pre>
```

```
## Loading required package: randomForest
```

```
## Warning: package 'randomForest' was built under R version 3.1.2
```

```
## randomForest 4.6-10
## Type rfNews() to see new features/changes/bug fixes.
```

Test - out of sample error

We use our own testing data set to determine the out of sample error.

```
predictions<-predict(modelFit,newdata=testing)
confusionMatrix(predictions,testing$classe)</pre>
```

```
## Confusion Matrix and Statistics
##
##
             Reference
                            C
##
  Prediction
                 Α
                      В
                                 D
                                      Ε
##
            A 1673
                      9
                            0
                                 0
                                      0
##
                 1 1129
                            7
            C
##
                      1 1018
                                10
##
            D
                 0
                      0
                            1 953
                                      1
##
            Ε
                      0
                            0
                                 1 1081
##
  Overall Statistics
##
##
##
                  Accuracy: 0.995
                     95% CI: (0.993, 0.996)
##
##
       No Information Rate: 0.284
##
       P-Value [Acc > NIR] : <2e-16
##
                     Kappa: 0.993
##
    Mcnemar's Test P-Value : NA
##
##
  Statistics by Class:
##
##
##
                         Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                            0.999
                                     0.991
                                              0.992
                                                        0.989
                                                                 0.999
## Specificity
                            0.998
                                     0.998
                                              0.998
                                                        1.000
                                                                 1.000
## Pos Pred Value
                            0.995
                                     0.993
                                              0.989
                                                        0.998
                                                                 0.999
## Neg Pred Value
                            1.000
                                     0.998
                                              0.998
                                                        0.998
                                                                 1.000
## Prevalence
                            0.284
                                     0.194
                                              0.174
                                                        0.164
                                                                 0.184
## Detection Rate
                            0.284
                                     0.192
                                                        0.162
                                                                  0.184
                                              0.173
## Detection Prevalence
                            0.286
                                     0.193
                                               0.175
                                                        0.162
                                                                  0.184
## Balanced Accuracy
                            0.999
                                     0.995
                                               0.995
                                                        0.994
                                                                  0.999
```

Predict the values for the test set

```
answers=predict(modelFit,newdata=raw_data_test)
answers
```

```
## [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
```

```
pml_write_files = function(x){
    n = length(x)
    for(i in 1:n){
        filename = paste0("problem_id_",i,".txt")
        write.table(x[i],file=filename,quote=FALSE,row.names=FALSE,col.names=FALSE)
    }
}

pml_write_files(answers)
```