

# Machine Learning Course Project: Predicting the manner of doing exercise based on parametric accelerometers data

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## Background

Using devices such as Jawbone Up, Nike FuelBand, and Fitbit it is now possible to collect a large amount of data about personal activity relatively inexpensively. These type of devices are part of the quantified self movement – a group of enthusiasts who take measurements about themselves regularly to improve their health, to find patterns in their behavior, or because they are tech geeks. One thing that people regularly do is quantify how much of a particular activity they do, but they rarely quantify how well they do it. In this project, your goal will be to use data from accelerometers on the belt, forearm, arm, and dumbbell of 6 participants. They were asked to perform barbell lifts correctly and incorrectly in 5 different ways. More information is available from the website here: <http://groupware.les.inf.puc-rio.br/har> (see the section on the Weight Lifting Exercise Dataset).

## Source Data

The training data for this project are available here:

<https://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv>

The test data are available here:

<https://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv>

The data for this project come from this source: <http://groupware.les.inf.puc-rio.br/har>. If you use the document you create for this class for any purpose please cite them as they have been very generous in allowing their data to be used for this kind of assignment.

## Purpose of the project / outcome

The goal of your project is to predict the manner in which they did the exercise. This is the "classe" variable in the training set. You may use any of the other variables to predict with. You should create a report describing how you built your model, how you used cross validation, what you think the expected out of sample error is, and why you made the choices you did. You will also use your prediction model to predict 20 different test cases.

## Loading the data

```
library(caret)

## Warning: package 'caret' was built under R version 3.2.4

## Loading required package: lattice

## Loading required package: ggplot2

## Warning: package 'ggplot2' was built under R version 3.2.4

library(rpart)
library(rpart.plot)

## Warning: package 'rpart.plot' was built under R version 3.2.4

library(RColorBrewer)
library(rattle)

## Warning: package 'rattle' was built under R version 3.2.4

## Rattle: A free graphical interface for data mining with R.
## Version 4.1.0 Copyright (c) 2006-2015 Togaware Pty Ltd.
## Type 'rattle()' to shake, rattle, and roll your data.

library(randomForest)

## Warning: package 'randomForest' was built under R version 3.2.4

## randomForest 4.6-12

## Type rfNews() to see new features/changes/bug fixes.

##
## Attaching package: 'randomForest'

## Следующий объект скрыт от 'package:ggplot2':
##
##      margin

set.seed(12345)
```

When we first take a look at the data (e.g. downloading from the link above and opening in XL spreadsheet) we observe that

The header contains the column names  
There are many missing values, which can be split into 3 types: NA, "", "#DIV/0!"  
The first column is not a variable, it counts the row number.

```
trainUrl <- "http://d396qusza40orc.cloudfront.net/predmachlearn/pml-
training.csv"
testUrl <- "http://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv"
training <- read.csv(url(trainUrl), na.strings=c("NA", "#DIV/0!", ""))
testing <- read.csv(url(testUrl), na.strings=c("NA", "#DIV/0!", ""))
```

Checking dim of 2 datasets, we conclude that both have 160 variables and different number of observations

```
dim(training); dim(testing)
## [1] 19622  160
## [1]  20 160
```

Looking at the variable CLASSE which we need to predict, we conclude that this is a "factor"-class variable which can take one of 5 meanings:

```
## [1] "factor"
##   Var1 Freq
## 1    A 5580
## 2    B 3797
## 3    C 3422
## 4    D 3216
## 5    E 3607
```

## Splitting Training dataset into 2 subsets

We want to split available Training dataset into 2 subsets: training1 (65%) and testing1 (35%)

```
train_partition <- createDataPartition(y=training$classe, p=0.65, list=FALSE)
training1 <- training[train_partition, ]
testing1 <- training[-train_partition, ]
dim(training1); dim(testing1)
## [1] 12757  160
## [1]  6865  160
```

We want to identify and remove all variables which have NearZeroVariance

```
nzv <- nearZeroVar(training1, saveMetrics=TRUE)
training1 <- training1[,nzv$nzv==FALSE]
nzv<- nearZeroVar(testing1,saveMetrics=TRUE)
testing1 <- testing1[,nzv$nzv==FALSE]
```

We want to remove the first column, which is simply the order of record but not a variable

```
training1 <- training1[,c(-1)]
```

Now we identify variables which have mostly NA. Those columns where NA are more than 60% will be removed

```
training1_temp <- training1
for(i in 1:length(training1)) {
  if( sum( is.na( training1[, i] ) ) /nrow(training1) >= .6 ) {
    for(j in 1:length(training1_temp)) {
      if( length( grep(names(training1[i]), names(training1_temp)[j]) )
==1) {
        training1_temp <- training1_temp[ , -j]
      }
    }
  }
}
```

```
}  
}
```

After checking how many columns / variables left after data cleaning, we conclude that their number reduced from 160 to 58

```
dim(training1_temp)  
## [1] 12757    58
```

resuming to our training subset

```
training1 <- training1_temp  
rm(training1_temp)
```

We need to repeat the same data cleaning for testing1 subset and for original testing dataset

```
clean1 <- colnames(training1)  
clean2 <- colnames(training1[, -58])  
testing1 <- testing1[clean1]  
testing <- testing[clean2]
```

Making sure the number of columns has reduced accordingly

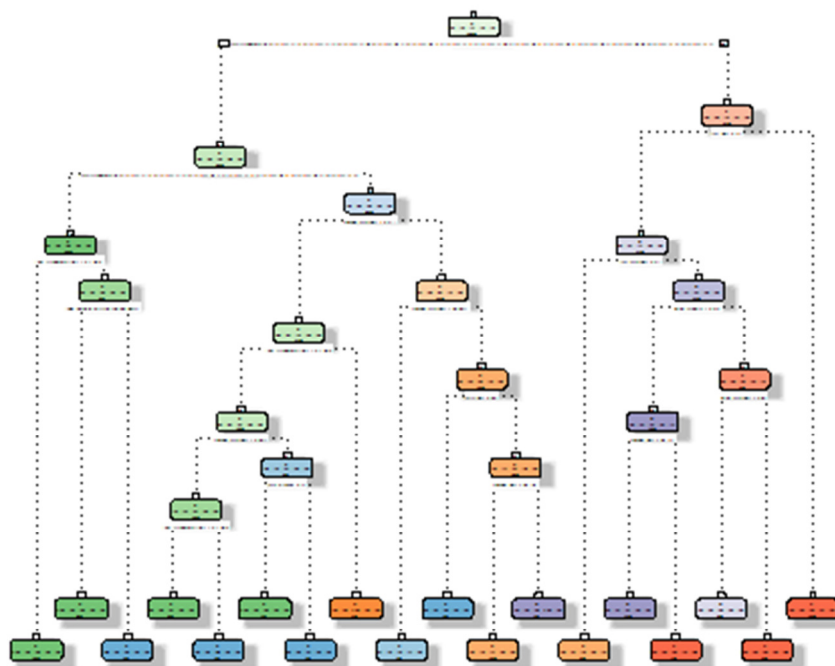
```
dim(testing1); dim(testing)  
## [1] 6865    58  
## [1] 20 57
```

We want to coerce the data into the same type

```
for (i in 1:length(testing) ) {  
  for(j in 1:length(training1)) {  
    if( length( grep(names(training1[i]), names(testing)[j]) ) ==1) {  
      class(testing[j]) <- class(training1[i])  
    }  
  }  
}  
  
testing <- rbind(training1[2, -58] , testing)  
testing <- testing[-1,]
```

## Prediction using Decision Trees

```
modFitA1 <- rpart(classe ~ ., data=training1, method="class")  
fancyRpartPlot(modFitA1)
```



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Predicting and checking accuracy using confusion matrix

```
predictionsA1 <- predict(modFitA1, testing1, type = "class")
confmatrix_tree <- confusionMatrix(predictionsA1, testing1$classe)
confmatrix_tree
```

## Confusion Matrix and Statistics

##

	Reference				
Prediction	A	B	C	D	E
A	1886	55	8	2	0
B	51	1124	71	57	0
C	16	141	1099	146	51
D	0	8	19	873	159
E	0	0	0	47	1052

##

## Overall Statistics

##

```
##           Accuracy : 0.879
##           95% CI   : (0.871, 0.8866)
##    No Information Rate : 0.2845
##    P-Value [Acc > NIR] : < 2.2e-16
```

##

```
##           Kappa : 0.8469
```

```
##  McNemar's Test P-Value : NA
```

##

## Statistics by Class:

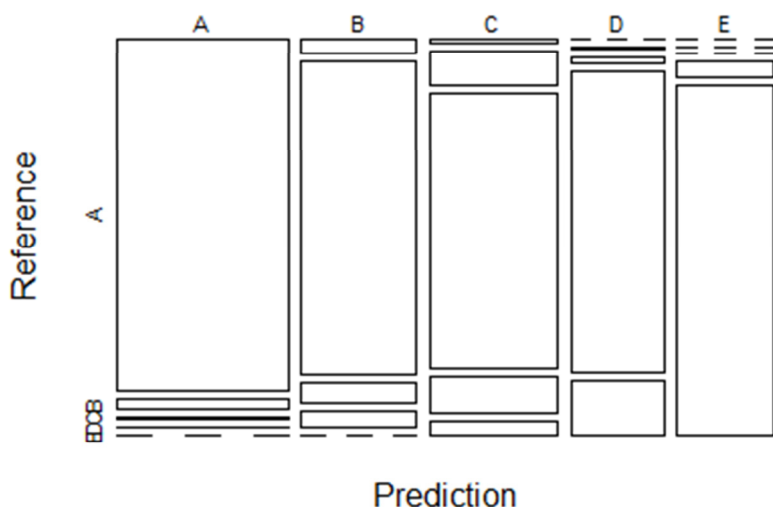
##

```
##           Class: A Class: B Class: C Class: D Class: E
```

```
## Sensitivity      0.9657  0.8464  0.9181  0.7760  0.8336
## Specificity     0.9868  0.9677  0.9375  0.9676  0.9916
## Pos Pred Value  0.9667  0.8626  0.7564  0.8244  0.9572
## Neg Pred Value  0.9864  0.9633  0.9819  0.9566  0.9636
## Prevalence      0.2845  0.1934  0.1744  0.1639  0.1838
## Detection Rate  0.2747  0.1637  0.1601  0.1272  0.1532
## Detection Prevalence 0.2842  0.1898  0.2117  0.1543  0.1601
## Balanced Accuracy 0.9762  0.9070  0.9278  0.8718  0.9126
```

```
plot(confmatrix_tree$table, col = confmatrix_tree$byClass, main =
paste("Decision Tree Confusion Matrix: Accuracy =",
round(confmatrix_tree$overall['Accuracy'], 4)))
```

## Decision Tree Confusion Matrix: Accuracy = 0.87



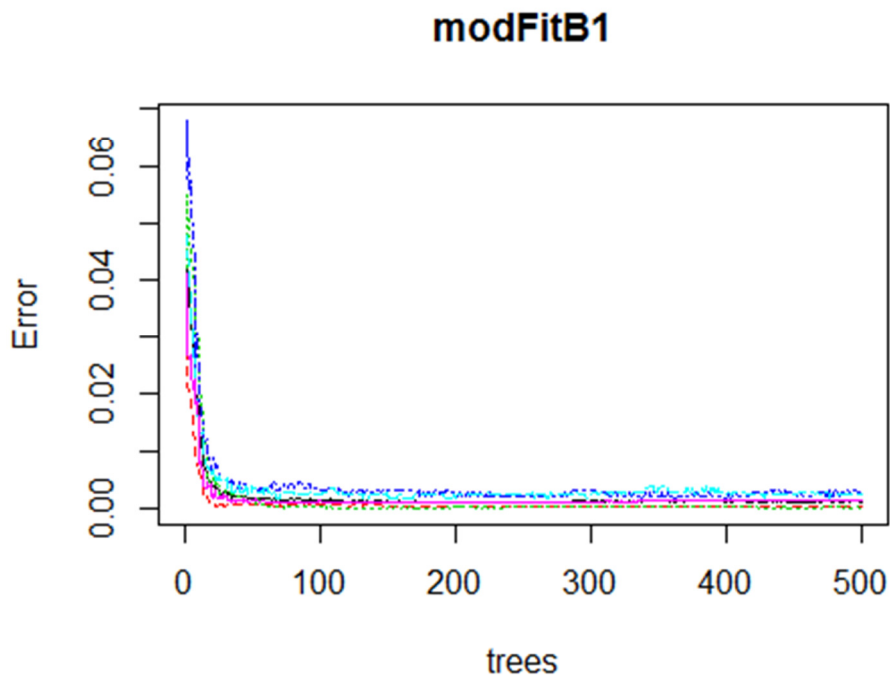
## Predictions using Random Forests

```
modFitB1 <- randomForest(classe ~. , data=training1)
predictionsB1 <- predict(modFitB1, testing1, type = "class")
confmatrix_randfor <- confusionMatrix(predictionsB1, testing1$classe)
confmatrix_randfor
```

```
## Confusion Matrix and Statistics
```

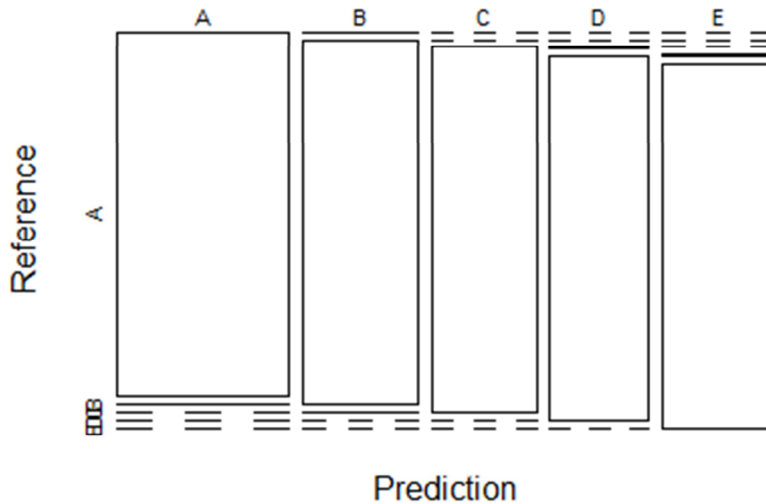
```
##
##           Reference
## Prediction    A    B    C    D    E
##           A 1952    2    0    0    0
##           B    1 1326    1    0    0
##           C    0    0 1194    0    0
##           D    0    0    2 1122    0
```

```
##           E      0      0      0      3 1262
##
## Overall Statistics
##
##           Accuracy : 0.9987
##           95% CI : (0.9975, 0.9994)
##           No Information Rate : 0.2845
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.9983
##           McNemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: A Class: B Class: C Class: D Class: E
## Sensitivity      0.9995  0.9985  0.9975  0.9973  1.0000
## Specificity      0.9996  0.9996  1.0000  0.9997  0.9995
## Pos Pred Value   0.9990  0.9985  1.0000  0.9982  0.9976
## Neg Pred Value   0.9998  0.9996  0.9995  0.9995  1.0000
## Prevalence       0.2845  0.1934  0.1744  0.1639  0.1838
## Detection Rate   0.2843  0.1932  0.1739  0.1634  0.1838
## Detection Prevalence 0.2846  0.1934  0.1739  0.1637  0.1843
## Balanced Accuracy 0.9995  0.9991  0.9987  0.9985  0.9997
plot(modFitB1)
```



```
plot(confmatrix_randfor$table, col = confmatrix_tree$byClass, main =
paste("Random Forest Confusion Matrix: Accuracy =",
round(confmatrix_randfor$overall['Accuracy'], 4)))
```

## Random Forest Confusion Matrix: Accuracy = 0.99



### Predicting CLASSE variable for the data in the Test dataset :

Random Forests gave a better accuracy on the testing1 dataset of 99.94% (compared to 86.6% for Decision Trees), so we'll use Random Forests outcome modfitB1 to predict CLASSE variable for the Test sample

```
predictionsB2 <- predict(modFitB1, testing, type = "class")
predictionsB2

##  2 31  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21
##  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
```

Generating Files for submission

```
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(predictionsB2)
```