

Practical Machine Learning Project: Classification of Exercises Quality

MARIANO MOLINA GARCIA

14/06/2020

Introduction

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.

In this project, the data from accelerometers on the belt, forearm, arm, and dumbbell of 6 participants will be used to quantify how well the exercise is done. The goal of your project is to predict the manner in which they did the exercise.

People were asked to perform barbell lifts correctly and incorrectly in 5 different ways, and the data previously commented have been stored in the file <https://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv>. The data for this project come from this source: <http://web.archive.org/web/20161224072740/http://groupware.les.inf.puc-rio.br/har>. More information about the data is available in that website, specifically in the section on the Weight Lifting Exercise Dataset.

Building the model. Data Analysis and selection of features.

The dataset contains 160 features, which can be used to predict if the exercise has been done properly or not. Six young health participants were asked to perform one set of 10 repetitions of the Unilateral Dumbbell Biceps Curl. The way the exercise has been done is labelled using the “classe” variable in the dataset. This “classe” variable can have 5 different values: A,B,C,D and E, which correspond to five different fashions: exactly according to the specification (Class A), throwing the elbows to the front (Class B), lifting the dumbbell only halfway (Class C), lowering the dumbbell only halfway (Class D) and throwing the hips to the front (Class E). Therefore, class A corresponds to the specified execution of the exercise, while the other 4 classes correspond to common mistakes.

First of all, an initial analysis of the data have been carried out. This analysis consists of:

1.- Load data. Checking the features and its range of values.

```
library(caret)
library(AppliedPredictiveModeling)
library(dplyr)
trainingData <- read.csv("pml-training.csv")
testingData<- read.csv("pml-testing.csv")
str(trainingData)
str(testingData)
```

The number of features is 160. The number of observations for the training dataset is 19622 and the number of observations for the test dataset is 20.

```
dim(trainingData)
```

```
## [1] 19622 160
```

```
dim(testingData)
```

```
## [1] 20 160
```

2.- Checking if there is feature with wrong NA values. If the percentage is very high, remove the column. If it is lower, analyze the possibilities to impute values. After this process, the number of features have been reduced to 93.

```
#Check if there are many rows of any variable with values NA
na_count <-sapply(trainingData, function(y) sum(length(which(is.na(y)))))
na_count <- data.frame(na_count)
perc_na <- data.frame(100*(na_count/nrow(trainingData)))
#Remove that cols with many NA from the analysis
trainingData<-trainingData[,perc_na<90]
testingData<-testingData[,perc_na<90]
dim(trainingData)
```

```
## [1] 19622 93
```

```
dim(testingData)
```

```
## [1] 20 93
```

3.- Checking if there is feature with empty values. If the percentage is very high, remove the column. If it is lower, analyze the possibilities to impute values. After this process, the number of features have been reduced to 60.

```
pat <- "^[[:space:]]*$"
matches <-sapply(trainingData, function(x) grepl(pat, x))
matches<- data.frame(matches)
val_count_empty <-sapply(matches, function(y) sum(length(which(y==TRUE))))
val_count_empty<- data.frame(val_count_empty)
perc_empty <- data.frame(100*(val_count_empty/nrow(trainingData)))
#Remove that cols with many empty values from the analysis
trainingData<-trainingData[,perc_empty<90]
testingData<-testingData[,perc_empty<90]
dim(trainingData)
```

```
## [1] 19622 60
```

```
dim(testingData)
```

```
## [1] 20 60
```

4.- Checking if there is features with a small variability of the values, which makes it useless for the prediction. There are not any feature with a single value, so the number of features for prediction is still 60.

```
val_count_unique <-sapply(trainingData, function(y) sum(length(unique(y))))
val_count_unique<- data.frame(val_count_unique)
trainingData<-trainingData[,val_count_unique>1]
testingData<-testingData[,val_count_unique>1]
dim(trainingData)
```

```
## [1] 19622 60
```

```
dim(testingData)
```

```
## [1] 20 60
```

5.- Checking if there are features useless for the prediction due to its meaning (time, names, etc).After removing these features, the number have been reduced to 53.

```
remove_cols = c(1:7)
trainingData<-trainingData[,-remove_cols]
testingData<-testingData[,-remove_cols]
dim(trainingData)
```

```
## [1] 19622    53
```

```
dim(testingData)
```

```
## [1] 20 53
```

6.- Remove near zero variables.

Near Zero variables: To identify these types of predictors, the following two metrics can be calculated: -the frequency of the most prevalent value over the second most frequent value (called the “frequency ratio”), which would be near one for well-behaved predictors and very large for highly-unbalanced data and -the “percent of unique values” is the number of unique values divided by the total number of samples (times 100) that approaches zero as the granularity of the data increases.

There are not any feature with near zero values, so the number of features for prediction is still 53.

```
nzv <- nearZeroVar(trainingData)
if (length(nzv) > 0) {
  trainingData <- trainingData[, -nzv]
  testingData <- testingData[, -nzv]
}
dim(trainingData)
```

```
## [1] 19622    53
```

```
dim(testingData)
```

```
## [1] 20 53
```

7.- Checking the correlation between the features to remove features with extremely high correlation, which makes the information redundant for the prediction. After analyzing the correlations, and removing the features with a correlation above 0.75, the number of features have been reduced to 32.

```
#Keep only numerical columns
numericCols <-supply(trainingData,is.numeric)
checkCorrTrainingData <- trainingData[,numericCols]
noCheckCorrTrainingData <- trainingData[,!numericCols]
#Calculate correlation
descrCor <- cor(checkCorrTrainingData)
summary(descrCor[upper.tri(descrCor)])
```

```
##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.
## -0.992008 -0.110080  0.002092  0.001790  0.092552  0.980924
```

```
#Remove columns with correlation > cutoff
highlyCorDescr <- findCorrelation(descrCor, cutoff = .75)
checkCorrTrainingData <- checkCorrTrainingData[, -highlyCorDescr]
descrCor2 <- cor(checkCorrTrainingData)
summary(descrCor2[upper.tri(descrCor2)])
```

```
##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.
## -0.606983 -0.103773  0.006527  0.003332  0.087527  0.736546
```

```
#Reconstruct dataframe with no numeric columns and numeric after correlation removal
trainingData <- data.frame(checkCorrTrainingData,"classe" = noCheckCorrTrainingData)
keepProblemId <- testingData$problem_id
testingData <- testingData[,intersect(colnames(testingData),colnames(trainingData))]
testingData <- data.frame(testingData,"problem_id" = keepProblemId)
dim(trainingData)
```

```
## [1] 19622    32
```

```
dim(testingData)
```

```
## [1] 20 32
```

8.- Check if there is outliers plotting the data with the violin shape and remove the outliers. The features gyros_dumbbell, gyros_forearm_x and gyros_forearm_z show outliers, and they have been removed. After analyzing the outliers, one observation of the training data have been removed, having 19621 to perform the analysis.

```
library(dplyr)
trainingData<-trainingData%>%filter(gyros_dumbbell_y<10)
trainingData<-trainingData%>%filter(gyros_forearm_x>-10)
trainingData<-trainingData%>%filter(gyros_forearm_z<10)
dim(trainingData)
```

```
## [1] 19621    32
```

```
dim(testingData)
```

```
## [1] 20 32
```

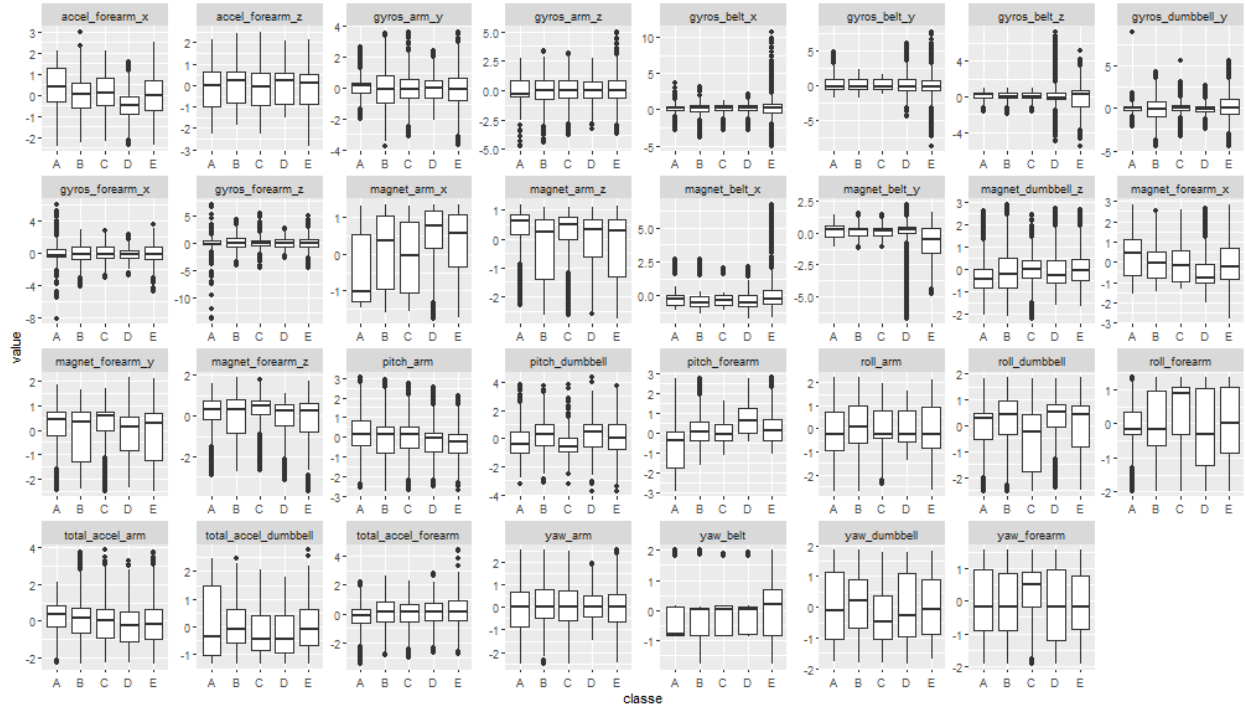
9.- Applying a preprocess to the data, to improve the performance or the prediction algorithms. From all the possible preprocess options (Possible values are “BoxCox”, “YeoJohnson”, “expoTrans”, “center”, “scale”, “range”, “knnImpute”, “bagImpute”, “medianImpute”, “pca”, “ica”, “spatialSign”, “corr”, “zv”, “nzv”, and “conditionalX”) centering and scaling have been selected. The correlation and nzv analysis have been carried out independently in previous steps.

```
#Centering and Scaling
preProcValues <- preprocess(trainingData, method = c("center", "scale"))
trainTransformed <- predict(preProcValues, trainingData)
testTransformed <- predict(preProcValues, testingData)
```

10.- Finally, some plots are performed to analyze the remaining features before proceeding to the use of multiclass classification models for prediction.

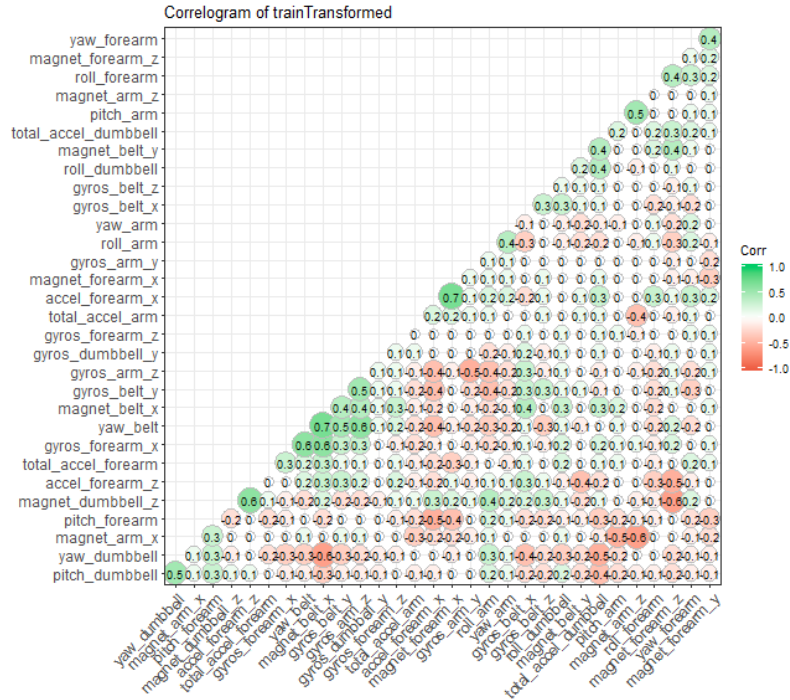
A boxplot analysis have been carried to, to visualize if there are some features which can be detected at first sight as good feature to distinguished between “classe” groups.

```
library(ggplot2)
library(tidyr)
trainTransformedlong <- gather(trainTransformed, key="measure", value="value"
                              ,c(names(trainTransformed[,1:31])))
p <- ggplot(data = trainTransformedlong, aes(x=classe,y=value)) +
  geom_boxplot()
p + facet_wrap( ~measure, scales="free",ncol=8)
```



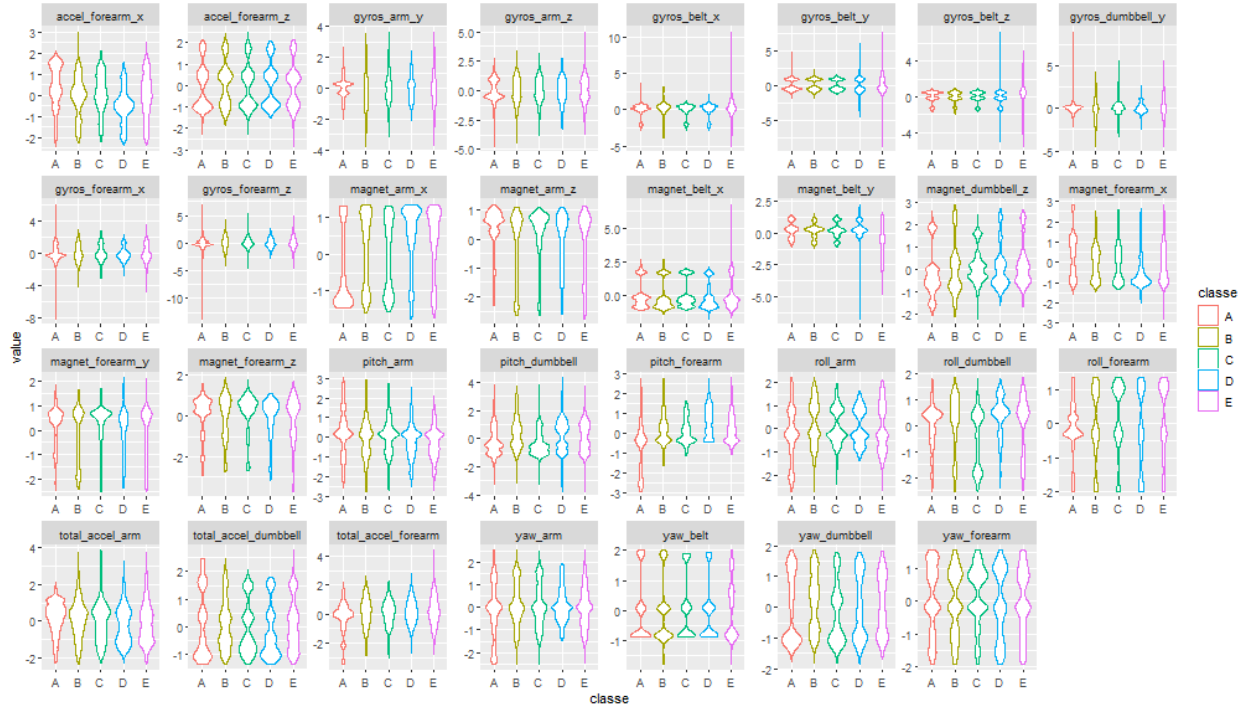
After that, a correlogram to show the correlation between the remaining features is presented as well.

```
#CorreloGram
library(ggplot2)
library(ggcorrplot)
corr <- round(cor(trainTransformed[,1:31]), 1)
ggcorrplot(corr, hc.order = TRUE,
            type = "lower",
            lab = TRUE,
            lab_size = 3,
            method="circle",
            colors = c("tomato2", "white", "springgreen3"),
            title="Correlogram of trainTransformed",
            ggtheme=theme_bw)
```



Finally, a violin plot to see the distribution of values per “classe” for each feature is shown as well.

```
library(ggplot2)
library(tidyr)
library(dplyr)
trainTransformedlong3 <- gather(trainTransformed, key="measure", value="value",
                                c(names(trainTransformed[,1:31])))
val=colnames(trainTransformedlong3)[1]
# plot
g <- ggplot(trainTransformedlong3, aes(x=classe, y=value,color=classe))
g <- g + geom_violin()
g + facet_wrap( ~measure,scales="free",ncol=8)
```



Building the model. Analysis of Multiclass classification algorithms.

The following 15 multiclass classification methods have been tested, and its accuracy compared: knn, rf, pda, parRF, sda, hddrda, LMT, slida, hdda, RRFglobal, C5.0, LogitBoost, pam, PART, rpart.

For all of them, a cv method has been selected for cross validation.

Other methods could have been selected. The available resampling methods are: The “boot”, “boot632”, “optimism_boot”, “boot_all”, “cv”, “repeatedcv”, “LOOCV”, “LGOCV” (for repeated training/test splits), “none” (only fits one model to the entire training set), “oob” (only for random forest, bagged trees, bagged earth, bagged flexible discriminant analysis, or conditional tree forest models), timeslice, “adaptive_cv”, “adaptive_boot” or “adaptive_LGOCV”

```
library(doParallel)
registerDoParallel(4)
getDoParWorkers()
```

- KNN

```
fitKNN = train(
  classe ~.,
  data = trainTransformed,
  method = "knn",
  trControl = trainControl(method = "cv", number = 10, savePredictions = "final",
    allowParallel = TRUE),
  tuneGrid = expand.grid(k = seq(1, 21, by = 2))
)
head(predict(fitKNN, type = "prob"))
predictedKNN <- predict(fitKNN, newdata = trainTransformed)
CM_fitKNN <- confusionMatrix(predictedKNN, trainTransformed$classe)
```

```
predictedKNN2<-predict(fitKNN,newdata=testTransformed)
varImp_fitKNN <- varImp(fitKNN)
```

- Random Forest

```
fitRF = train(
  classe ~.,
  data = trainTransformed,
  method = 'rf',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitRF, type = "prob"))
predictedRF<-predict(fitRF,newdata=trainTransformed)
CM_fitRF <- confusionMatrix(predictedRF, trainTransformed$classe)
predictedRF2<-predict(fitRF,newdata=testTransformed)
varImp_fitRF <- varImp(fitRF)
```

- pda

```
library(mda)
fitpda= train(
  classe ~.,
  data = trainTransformed,
  method = 'pda',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitpda, type = "prob"))
predictedpda<-predict(fitpda,newdata=trainTransformed)
CM_fitpda <- confusionMatrix(predictedpda, trainTransformed$classe)
predictedpda2<-predict(fitpda,newdata=testTransformed)
varImp_fitpda <- varImp(fitpda)
```

-Parallel Random Forest

```
library(e1071)
library(randomForest)
fitPRF = train(
  classe ~.,
  data = trainTransformed,
  method = 'parRF',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitPRF, type = "prob"))
predictedPRF<-predict(fitPRF,newdata=trainTransformed)
CM_fitPRF <- confusionMatrix(predictedPRF, trainTransformed$classe)
predictedPRF2<-predict(fitPRF,newdata=testTransformed)
varImp_fitPRF <- varImp(fitPRF)
```

-sda

```
library(sda)
fitsda= train(
  classe ~.,
```



```

data = trainTransformed,
method = 'sda',
trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitsda, type = "prob"))
predicted_sda<-predict(fitsda,newdata=trainTransformed)
CM_fitsda <- confusionMatrix(predicted_sda, trainTransformed$classe)
predicted_sda2<-predict(fitsda,newdata=testTransformed)
varImp_fitsda <- varImp(fitsda)

```

- hdrda

```

library(sparsediscrim)
fithdrda= train(
  classe ~.,
  data = trainTransformed,
  method = 'hdrda',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fithdrda, type = "prob"))
predicted_hdrda<-predict(fithdrda,newdata=trainTransformed)
CM_fithdrda <- confusionMatrix(predicted_hdrda, trainTransformed$classe)
predicted_hdrda2<-predict(fithdrda,newdata=testTransformed)
varImp_fithdrda <- varImp(fithdrda)

```

- LMT (Logistic Model Trees)

```

##Logistic Model Trees
library(RWeka)
fitLMT = train(
  classe ~.,
  data = trainTransformed,
  method = "LMT",
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitLMT, type = "prob"))
predicted_LMT<-predict(fitLMT,newdata=trainTransformed)
CM_fitLMT <- confusionMatrix(predicted_LMT, trainTransformed$classe)
predicted_LMT2<-predict(fitLMT,newdata=testTransformed)
varImp_fitLMT <- varImp(fitLMT)

```

- slda

```

library(ipred)
fitslda= train(
  classe ~.,
  data = trainTransformed,
  method = 'slda',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitslda, type = "prob"))
predicted_slda<-predict(fitslda,newdata=trainTransformed)

```

```
CM_fitslda <- confusionMatrix(predictedslda, trainTransformed$classe)
predictedslda2<-predict(fitslda,newdata=testTransformed)
varImp_fitslda <- varImp(fitslda)
```

- hdda

```
library(HDclassif)
fithdda= train(
  classe ~.,
  data = trainTransformed,
  method = 'hdda',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fithdda, type = "prob"))
predictedhdda<-predict(fithdda,newdata=trainTransformed)
CM_fithdda <- confusionMatrix(predictedhdda, trainTransformed$classe)
predictedhdda2<-predict(fithdda,newdata=testTransformed)
varImp_fithdda <- varImp(fithdda)
```

- RRFglobal

```
library(RRF)
fitRRFglobal= train(
  classe ~.,
  data = trainTransformed,
  method = 'RRFglobal',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitRRFglobal, type = "prob"))
predictedRRFglobal<-predict(fitRRFglobal,newdata=trainTransformed)
CM_fitRRFglobal <- confusionMatrix(predictedRRFglobal, trainTransformed$classe)
predictedRRFglobal2<-predict(fitRRFglobal,newdata=testTransformed)
varImp_fitRRFglobal <- varImp(fitRRFglobal)
```

- C5.0

```
fitC50 = train(
  classe ~.,
  data = trainTransformed,
  method = 'C5.0',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                           ,allowParallel = TRUE)
)
head(predict(fitC50, type = "prob"))
predictedC50<-predict(fitC50,newdata=trainTransformed)
CM_fitC50 <- confusionMatrix(predictedC50, trainTransformed$classe)
predictedC502<-predict(fitC50,newdata=testTransformed)
varImp_fitC50 <- varImp(fitC50)
```

- LogitBoost

```
library(caTools)
fitLogitBoost= train(
  classe ~.,
```

```

data = trainTransformed,
method = 'LogitBoost',
trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                          ,allowParallel = TRUE)
)
head(predict(fitLogitBoost, type = "prob"))
predictedLogitBoost<-predict(fitLogitBoost,newdata=trainTransformed)
CM_fitLogitBoost <- confusionMatrix(predictedLogitBoost, trainTransformed$classe)
predictedLogitBoost2<-predict(fitLogitBoost,newdata=testTransformed)
varImp_fitLogitBoost <- varImp(fitLogitBoost)

```

- pam

```

library(pamr)
fitpam= train(
  classe ~.,
  data = trainTransformed,
  method = 'pam',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                          ,allowParallel = TRUE)
)
head(predict(fitpam, type = "prob"))
predictedpam<-predict(fitpam,newdata=trainTransformed)
CM_fitpam <- confusionMatrix(predictedpam, trainTransformed$classe)
predictedpam2<-predict(fitpam,newdata=testTransformed)
varImp_fitpam <- varImp(fitpam)

```

- PART (Rule Based Classifier)

```

fitRBC = train(
  classe ~.,
  data = trainTransformed,
  method = 'PART',
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                          ,allowParallel = TRUE)
)
head(predict(fitRBC, type = "prob"))
predictedRBC<-predict(fitRBC,newdata=trainTransformed)
CM_fitRBC <- confusionMatrix(predictedRBC, trainTransformed$classe)
predictedRBC2<-predict(fitRBC,newdata=testTransformed)
varImp_fitRBC <- varImp(fitRBC)

```

- rpart (Tree with RPART)

```

fitRPART = train(
  classe ~.,
  data = trainTransformed,
  method = "rpart",
  trControl = trainControl(method = "cv", number = 10,savePredictions = "final"
                          ,allowParallel = TRUE)
)
head(predict(fitRPART, type = "prob"))
predictedRPART<-predict(fitRPART,newdata=trainTransformed)
CM_fitRPART <- confusionMatrix(predictedRPART, trainTransformed$classe)
predictedRPART2<-predict(fitRPART,newdata=testTransformed)
varImp_fitRPART <- varImp(fitRPART)

```

Results of the multiclass classification models. Importance of features.

In the following table, the results for accuracy for the algorithms evaluated are presented:

```
result_Analysis = data.frame(methods = methods<-c("C50","hdda","hdrda","KNN","LMT"
                                                    ,"LogitBoost","pam","pda","PRF"
                                                    ,"RBC","RF","RPART","RRFglobal"
                                                    ,"sda","slda"),
accuracy = c(CM_fitC50$overall["Accuracy"],CM_fithdda$overall["Accuracy"],
             CM_fithdrda$overall["Accuracy"],CM_fitC50$overall["Accuracy"],
             CM_fitKNN$overall["Accuracy"],CM_fitLogitBoost$overall["Accuracy"],
             CM_fitpam$overall["Accuracy"],CM_fitpda$overall["Accuracy"],
             CM_fitPRF$overall["Accuracy"],CM_fitRBC$overall["Accuracy"],
             CM_fitRF$overall["Accuracy"],CM_fitRPART$overall["Accuracy"],
             CM_fitRRFglobal$overall["Accuracy"],CM_fitsda$overall["Accuracy"],
             CM_fitslda$overall["Accuracy"]))
result_Analysis
```

##	methods	accuracy
## 1	C50	1.0000000
## 2	hdda	0.7601549
## 3	hdrda	0.8249834
## 4	KNN	1.0000000
## 5	LMT	1.0000000
## 6	LogitBoost	0.8700265
## 7	pam	0.4182254
## 8	pda	0.5834565
## 9	PRF	1.0000000
## 10	RBC	0.9992355
## 11	RF	1.0000000
## 12	RPART	0.5246929
## 13	RRFglobal	1.0000000
## 14	sda	0.5832017
## 15	slda	0.4235768

The results show that 6 methods have an in sample accuracy of 1, meaning that they are able to classify properly all the observations in the training data. With this in sample accuracy, it is expected to have a very high classification out of sample accuracy. To finalize the analysis, the importance of each predictor has been analyzed for those 6 methods with a 100% accuracy.

- Importance for C50

```
## C5.0 variable importance
##
## only 20 most important variables shown (out of 31)
##
## Overall
## gyros_belt_z 100.00
## magnet_forearm_x 100.00
## gyros_dumbbell_y 100.00
## magnet_forearm_y 100.00
## pitch_forearm 100.00
```

```
## magnet_belt_y      100.00
## yaw_belt           99.98
## yaw_arm            99.88
## magnet_arm_z       99.76
## roll_arm           98.78
## magnet_dumbbell_z  98.64
## gyros_belt_y       98.56
## roll_forearm       97.97
## gyros_arm_y        97.69
## magnet_belt_x      96.47
## magnet_forearm_z   94.15
## roll_dumbbell      93.48
## pitch_dumbbell     88.82
## gyros_belt_x       88.59
## yaw_dumbbell       86.33
```

- Importance for KNN

```
## ROC curve variable importance
##
##   variables are sorted by maximum importance across the classes
##   only 20 most important variables shown (out of 31)
##
##           A         B         C         D         E
## pitch_forearm    62.964 100.000 67.237 62.96 100.000
## accel_forearm_x  40.675  81.171 40.675 40.68  81.171
## magnet_arm_x     53.104  78.110 65.004 53.10  78.110
## pitch_dumbbell   52.842  52.842 52.842 71.31  44.049
## magnet_forearm_x 38.891  71.126 33.586 33.59  71.126
## magnet_belt_y    15.289  10.317 67.372 10.32  15.289
## roll_dumbbell    41.216  51.434 30.666 62.50  51.434
## magnet_dumbbell_z 56.342  36.753 53.959 23.75  56.342
## magnet_arm_z     52.443  52.443 52.443 52.44  40.079
## pitch_arm        26.743  42.163 49.028 26.74  42.163
## yaw_dumbbell     19.186  19.186 19.186 41.29  13.608
## total_accel_arm  30.686  40.517 32.030 15.97  40.517
## magnet_forearm_y 19.326  37.449 28.196 24.54  37.449
## roll_arm         34.545  34.545 34.545 34.54  24.087
## roll_forearm     34.196   6.325 13.790 23.73  34.196
## total_accel_forearm 23.437 27.044 31.981 23.44  27.044
## magnet_belt_x    25.177  25.177 25.177 25.26  18.092
## yaw_forearm      10.754  16.280 11.636 22.29  16.280
## total_accel_dumbbell 9.533 17.622  9.533 19.89  17.622
## yaw_belt         9.833   9.833 18.426 17.17   8.828
```

- Importance for LMT

```
## ROC curve variable importance
##
##   variables are sorted by maximum importance across the classes
##   only 20 most important variables shown (out of 31)
##
##           A         B         C         D         E
## pitch_forearm    62.964 100.000 67.237 62.96 100.000
## accel_forearm_x  40.675  81.171 40.675 40.68  81.171
## magnet_arm_x     53.104  78.110 65.004 53.10  78.110
```

## pitch_dumbbell	52.842	52.842	52.842	71.31	44.049
## magnet_forearm_x	38.891	71.126	33.586	33.59	71.126
## magnet_belt_y	15.289	10.317	67.372	10.32	15.289
## roll_dumbbell	41.216	51.434	30.666	62.50	51.434
## magnet_dumbbell_z	56.342	36.753	53.959	23.75	56.342
## magnet_arm_z	52.443	52.443	52.443	52.44	40.079
## pitch_arm	26.743	42.163	49.028	26.74	42.163
## yaw_dumbbell	19.186	19.186	19.186	41.29	13.608
## total_accel_arm	30.686	40.517	32.030	15.97	40.517
## magnet_forearm_y	19.326	37.449	28.196	24.54	37.449
## roll_arm	34.545	34.545	34.545	34.54	24.087
## roll_forearm	34.196	6.325	13.790	23.73	34.196
## total_accel_forearm	23.437	27.044	31.981	23.44	27.044
## magnet_belt_x	25.177	25.177	25.177	25.26	18.092
## yaw_forearm	10.754	16.280	11.636	22.29	16.280
## total_accel_dumbbell	9.533	17.622	9.533	19.89	17.622
## yaw_belt	9.833	9.833	18.426	17.17	8.828

- Importance for PRF

```
## parRF variable importance
##
##   only 20 most important variables shown (out of 31)
##
##           Overall
## yaw_belt           100.00
## magnet_dumbbell_z   80.55
## magnet_belt_y       67.07
## pitch_forearm       66.86
## roll_dumbbell       56.55
## roll_forearm        51.98
## gyros_belt_z        44.00
## roll_arm            38.62
## gyros_dumbbell_y    37.79
## total_accel_dumbbell 37.35
## yaw_dumbbell        37.15
## accel_forearm_x     32.61
## accel_forearm_z     31.97
## pitch_dumbbell      31.19
## magnet_forearm_z     30.62
## magnet_belt_x       30.46
## yaw_arm             29.72
## magnet_forearm_y    28.68
## magnet_forearm_x    27.56
## magnet_arm_x        27.42
```

- Importance for RF

```
## rf variable importance
##
##   only 20 most important variables shown (out of 31)
##
##           Overall
## yaw_belt           100.00
## magnet_dumbbell_z   74.64
## magnet_belt_y       67.66
```

```

## pitch_forearm      65.28
## roll_forearm       58.02
## roll_dumbbell      53.96
## gyros_belt_z       42.88
## roll_arm           40.64
## total_accel_dumbbell 39.35
## yaw_dumbbell       37.96
## gyros_dumbbell_y   37.50
## accel_forearm_x    34.37
## magnet_arm_x       32.23
## pitch_dumbbell     31.52
## magnet_forearm_z   30.72
## accel_forearm_z    29.36
## magnet_belt_x      29.25
## magnet_forearm_y   28.01
## magnet_forearm_x   27.51
## yaw_arm            27.46

```

- Importance for RRF Global

```

## RRFglobal variable importance
##
##   only 20 most important variables shown (out of 31)
##
##               Overall
## yaw_belt      100.000
## pitch_forearm  78.181
## magnet_belt_y  74.533
## magnet_dumbbell_z 55.788
## roll_dumbbell  47.250
## total_accel_dumbbell 40.856
## roll_forearm   38.187
## accel_forearm_z 30.353
## gyros_belt_z   26.907
## magnet_forearm_z 18.316
## accel_forearm_x 18.316
## yaw_dumbbell   17.891
## magnet_belt_x  16.838
## roll_arm       15.080
## yaw_arm        14.240
## pitch_arm      12.333
## magnet_arm_x   11.732
## gyros_arm_y    9.711
## pitch_dumbbell  9.587
## gyros_dumbbell_y 9.366

```