

# Machine Learning PML Project

Tanya Nijhawan

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```
install.packages("e1071")
```

## Overview

Predicting the manner, in which participants, performed the exercise with the help of the *Weight Lifting Exercise Dataset* from the website :: <http://web.archive.org/web/20161224072740/http://groupware.les.inf.puc-rio.br/har> (<http://web.archive.org/web/20161224072740/http://groupware.les.inf.puc-rio.br/har>)

The data is collected from the arm, forearm, belt and dumbbells of 6 participants.

## Importing of the Libraries and Loading the Data-

```
library(caret)
```

```
## Warning: package 'caret' was built under R version 3.6.3
```

```
## Loading required package: lattice
```

```
## Loading required package: ggplot2
```

```
## Warning: package 'ggplot2' was built under R version 3.6.3
```

```
library(randomForest)
```

```
## Warning: package 'randomForest' was built under R version 3.6.3
```

```
## 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(Metrics)
```

```
## Warning: package 'Metrics' was built under R version 3.6.3
```

```
##  
## Attaching package: 'Metrics'
```

```
## The following objects are masked from 'package:caret':  
##  
##      precision, recall
```

```
trainData <- read.csv("pml-training.csv",header=TRUE,na.strings=c("NA",""))  
testData <- read.csv("pml-testing.csv",header=TRUE,na.strings=c("NA",""))  
#replacing all strings such as NA and empty strings to NA value of R
```

Dimensions of training and testing data::

```
dim(trainData)
```

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

```
dim(testData)
```

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

## Preprocessing or the Cleaning of Data-

1. Eliminate variables - that have a variance equal to or close to zero.
2. Drop the missing value...
3. Drop all the unnecessary columns...

```
# Remove near zero covariates
NSV <- nearZeroVar(trainData,saveMetrics=TRUE)
trainData <- trainData[,!NSV$nzv]
testData <- testData[,!NSV$nzv]

# Drop missing values
train_filt_na <- trainData[, (colSums(is.na(trainData)) == 0)]
test_filt_na <- testData[, (colSums(is.na(testData)) == 0)]

# Drop unnecessary columns
rmCol_train <- c("user_name","raw_timestamp_part_1","raw_timestamp_part_2","cvtd_timestamp","num_
_window")
rmCol_test <- c("user_name","raw_timestamp_part_1","raw_timestamp_part_2","cvtd_timestamp","num_
window","problem_id")
trainData_rmCol <- train_filt_na[,!(names(train_filt_na) %in% rmCol_train)]
testData_rmCol <- test_filt_na[,!(names(test_filt_na) %in% rmCol_test)]
```

The dimensions are:

```
dim(trainData_rmCol)
```

```
## [1] 19622    54
```

```
dim(testData_rmCol)
```

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

## Partitioning of the Dataset

We will create the training and validation dataset.

```
inTrain <- createDataPartition(y=trainData$classe, p=0.7, list=FALSE)
train_clean <- trainData_rmCol[inTrain,]
valid_clean <- trainData_rmCol[-inTrain,]
```

```
cor <- abs(sapply(colnames(train_clean[, -ncol(trainData)]), function(x) cor(as.numeric(train_cl
ean[, x]), as.numeric(train_clean$classe), method = "spearman")))
```

No predictors seem to be strongly correlated with the outcome. Linear regression may not be a good option. Therefore we select random forest model.

## Random Forest Model

We attempt to fit a random forest model and test the model performance on the validation set.

```
set.seed(71)
```

```
# Fit randomforest model
```

```
model <- train(classe ~ ., method = "rf", data = train_clean, importance = TRUE, trControl = tra  
inControl(method = "cv", number = 4))  
model
```

```
## Random Forest
```

```
##
```

```
## 13737 samples
```

```
## 53 predictor
```

```
## 5 classes: 'A', 'B', 'C', 'D', 'E'
```

```
##
```

```
## No pre-processing
```

```
## Resampling: Cross-Validated (4 fold)
```

```
## Summary of sample sizes: 10302, 10303, 10304, 10302
```

```
## Resampling results across tuning parameters:
```

```
##
```

```
## mtry Accuracy Kappa
```

```
## 2 0.9958506 0.9947512
```

```
## 27 0.9997089 0.9996318
```

```
## 53 0.9996361 0.9995397
```

```
##
```

```
## Accuracy was used to select the optimal model using the largest value.
```

```
## The final value used for the model was mtry = 27.
```

```
valid_pred <- predict(model, newdata=valid_clean)
```

```
# To check the performance of the model
```

```
confusionMatrix(valid_pred,valid_clean$classe)
```

```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction   A    B    C    D    E
##           A 1674    0    0    0    0
##           B    0 1139    0    0    0
##           C    0    0 1026    0    0
##           D    0    0    0  964    0
##           E    0    0    0    0 1082
##
## Overall Statistics
##
##           Accuracy : 1
##           95% CI : (0.9994, 1)
##           No Information Rate : 0.2845
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 1
##
##           Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: A Class: B Class: C Class: D Class: E
## Sensitivity           1.0000   1.0000   1.0000   1.0000   1.0000
## Specificity           1.0000   1.0000   1.0000   1.0000   1.0000
## Pos Pred Value        1.0000   1.0000   1.0000   1.0000   1.0000
## Neg Pred Value        1.0000   1.0000   1.0000   1.0000   1.0000
## Prevalence            0.2845   0.1935   0.1743   0.1638   0.1839
## Detection Rate        0.2845   0.1935   0.1743   0.1638   0.1839
## Detection Prevalence  0.2845   0.1935   0.1743   0.1638   0.1839
## Balanced Accuracy      1.0000   1.0000   1.0000   1.0000   1.0000
```

## Prediction

We now use this model to predict on the testing data.

```
test_pred <- predict(model, newdata=testData_rmCol)
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)
  }
}
write_files(test_pred)
```

## Results

We used 52 variables to build the random forest model with 100 trees using 4 fold cross validation.