

Human Activity Recognition

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Load raw data

```
library(caret)
```

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

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

```
library(rattle)
```

```
## 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(e1071)
```

```
library(ggplot2)
```

```
library(C50)
```

```
trainData <- read.csv("pml-training.csv",na.strings=c("NA","#DIV/0!",""))
```

```
testData <- read.csv("pml-testing.csv",na.strings=c("NA","#DIV/0!",""))
```

Clean the raw data

Remove first column that is not required - running number

```
cleanTrainData <- trainData[,c(-1)]
```

Find NA values and exclude

```
notNA <- sapply(cleanTrainData,function(i){sum(is.na(i))/length(i)})<0.9
```

```
cleanTrainData <- cleanTrainData[, notNA]
```

Remove near zero variance columns/variables

```
nzv <- nearZeroVar(cleanTrainData, saveMetrics=TRUE)
```

```
cleanTrainData <- cleanTrainData[,nzv$nzv==FALSE]
```

Drop unnecessary columns from the testing dataset

```
columnNames <- colnames(cleanTrainData[,-58]) # remove class column.
cleanTestData <- testData[columnNames]
```

Split the training set into a training and validation set.

```
intrain<-createDataPartition(y=cleanTrainData$classe,p=0.7,list=FALSE)
training<-cleanTrainData[intrain,]
testing<-cleanTrainData[-intrain,]
```

Cross Validation to select the best model performance

```
fitControl <- trainControl(method = "cv", number = 10, repeats = 0,savePred=T, classProb=T)
modelHAR_1 <- train(classe~., data = training, method = "rpart", trControl= fitControl)
```

Loading required package: rpart

```
modelHAR_2 <- train(classe~., data = training, method = "rpart1SE", trControl= fitControl)
modelHAR_3 <- train(classe~., data = training, method = "C5.0Tree", trControl= fitControl)
```

Confusion martix

```
activity_1 <- predict(modelHAR_1, newdata = testing[,-58])
conMatrix_1 <- confusionMatrix(activity_1, testing$classe)
conMatrix_1
```

Confusion Matrix and Statistics

##

Reference

## Prediction		A	B	C	D	E
## A	1260	284	21	31	13	
## B	107	493	17	0	0	
## C	83	178	540	28	48	
## D	218	184	448	905	512	
## E	6	0	0	0	509	

##

Overall Statistics

##

Accuracy : 0.6299

95% CI : (0.6174, 0.6423)

No Information Rate : 0.2845

P-Value [Acc > NIR] : < 2.2e-16

##

Kappa : 0.5355

McNemar's Test P-Value : NA

##

Statistics by Class:

##

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

```
## Sensitivity      0.7527  0.43284  0.52632  0.9388  0.47043
## Specificity      0.9171  0.97387  0.93064  0.7232  0.99875
## Pos Pred Value   0.7831  0.79903  0.61574  0.3992  0.98835
## Neg Pred Value   0.9032  0.87737  0.90296  0.9837  0.89330
## Prevalence       0.2845  0.19354  0.17434  0.1638  0.18386
## Detection Rate   0.2141  0.08377  0.09176  0.1538  0.08649
## Detection Prevalence 0.2734  0.10484  0.14902  0.3852  0.08751
## Balanced Accuracy 0.8349  0.70335  0.72848  0.8310  0.73459
```

```
activity_2 <- predict(modelHAR_1, newdata = training[,-58])
conMatrix_2 <- confusionMatrix(activity_2, training$classe)
conMatrix_2
```

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

```
##
##           Reference
## Prediction   A    B    C    D    E
##           A 3021  665   68   79   31
##           B  208 1183   28    0    0
##           C  186  368 1270   59  150
##           D  483  442 1030 2114 1222
##           E    8    0    0    0 1122
##
```

```
## Overall Statistics
```

```
##
##           Accuracy : 0.6341
##           95% CI : (0.6259, 0.6421)
##           No Information Rate : 0.2843
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.5402
##           McNemar's Test P-Value : NA
##
```

```
## Statistics by Class:
```

```
##
##           Class: A Class: B Class: C Class: D Class: E
## Sensitivity      0.7734  0.44507  0.53005  0.9387  0.44436
## Specificity      0.9143  0.97870  0.93272  0.7234  0.99929
## Pos Pred Value   0.7818  0.83369  0.62469  0.3995  0.99292
## Neg Pred Value   0.9104  0.88026  0.90379  0.9837  0.88871
## Prevalence       0.2843  0.19349  0.17442  0.1639  0.18381
## Detection Rate   0.2199  0.08612  0.09245  0.1539  0.08168
## Detection Prevalence 0.2813  0.10330  0.14799  0.3852  0.08226
## Balanced Accuracy 0.8438  0.71188  0.73139  0.8310  0.72182
```

```
activity_3 <- predict(modelHAR_3, newdata = training[,-58])
conMatrix_3 <- confusionMatrix(activity_3, training$classe)
conMatrix_3
```

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

```
##
##           Reference
## Prediction   A    B    C    D    E
```

```
##           A 3900    12    0    0    0
##           B    5 2636    1    0    0
##           C    0    9 2395    2    2
##           D    1    1    0 2249    2
##           E    0    0    0    1 2521
##
## Overall Statistics
##
##           Accuracy : 0.9974
##           95% CI : (0.9964, 0.9982)
##           No Information Rate : 0.2843
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.9967
##           McNemar's Test P-Value : NA
##
## Statistics by Class:
##
##           Class: A Class: B Class: C Class: D Class: E
## Sensitivity           0.9985  0.9917  0.9996  0.9987  0.9984
## Specificity           0.9988  0.9995  0.9989  0.9997  0.9999
## Pos Pred Value        0.9969  0.9977  0.9946  0.9982  0.9996
## Neg Pred Value        0.9994  0.9980  0.9999  0.9997  0.9996
## Prevalence            0.2843  0.1935  0.1744  0.1639  0.1838
## Detection Rate        0.2839  0.1919  0.1743  0.1637  0.1835
## Detection Prevalence  0.2848  0.1923  0.1753  0.1640  0.1836
## Balanced Accuracy      0.9986  0.9956  0.9992  0.9992  0.9992
```

As the “C5.0Tree” has the highest accuracy (0.9972) when predicting the test data. we select this model to build our final model.

Build final model

Final model is build with the “C5.0Tree” algorithm using the entire training data set.

```
fitControl <- trainControl(method = "cv", number = 10, repeats = 0, savePred=T, classProb=T)
modelHAR_F <- train(classe~., data = cleanTrainData, method = "C5.0Tree", trControl= fitControl)
```

Perform prediction with final model

```
act_prediction <- predict(modelHAR_F, newdata = cleanTestData)
```

The predistions obtained are:

```
act_prediction

## [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
```