Machine Learning Project

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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.

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, a group of 6 people were asked to perform barbell lifts correctly and incorrectly in 5 different ways, and our goal will be to use the data gathered from accelerometers on the belt, forearm, arm, and dumbell, and predict the manner in which they did the exercise.

Importing and cleaning data

```
library (caret)
library (randomForest)
library (rattle)
library (rpart)
library (party)
library (rpart.plot)
training and validation <- read.csv("https://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv");
dim(training and validation)
## [1] 19622
              160
testing <- read.csv("https://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv"); dim(testing)
## [1] 20 160
set.seed(78863439)
inTrain \leftarrow sample(2, nrow(training\_and\_validation), replace = TRUE, prob = c(0.7,0.3))
training <- training and validation[inTrain==1,]</pre>
validation <- training_and_validation[inTrain==2,]</pre>
dim(training); dim(validation)
## [1] 13791 160
```

```
## [1] 5831 160
```

training <- subset(training, select=-c(X, user_name, raw_timestamp_part_1, raw_timestamp_part_2, cvtd_timest amp, new window, num window, kurtosis roll belt, kurtosis picth belt, kurtosis yaw belt, skewness roll belt, skewness_roll_belt.1, skewness_yaw_belt, max_yaw_belt, min_yaw_belt, amplitude_yaw_belt,kurtosis_roll_arm, k urtosis_picth_arm, kurtosis_yaw_arm, skewness_roll_arm, skewness_pitch_arm, skewness_yaw_arm, kurtosis_roll_ dumbbell, kurtosis picth_dumbbell, kurtosis_yaw_dumbbell, skewness_roll_dumbbell, skewness pitch_dumbbell, s kewness yaw dumbbell, max yaw dumbbell, min yaw dumbbell, amplitude yaw dumbbell, kurtosis roll forearm, kurt osis picth forearm, kurtosis yaw forearm, skewness roll forearm, skewness pitch forearm, skewness yaw forearm , max_yaw_forearm, min_yaw_forearm, amplitude_yaw_forearm, max_roll_belt, max_picth_belt, min_roll_belt, min_pitch_belt, amplitude_roll_belt, amplitude_pitch_belt, var_total_accel_belt , avg_roll_belt, stddev_roll_belt, var_roll_belt, avg_pitch_belt, stddev_pitch_belt,var_pitch_belt,avg_yaw_b elt,stddev_yaw_belt,var_yaw_belt,var_accel_arm,avg_roll_arm,stddev_roll_arm, var_roll_arm, avg_pitch_arm,std dev_pitch_arm, var_pitch_arm, avg_yaw_arm, stddev_yaw_arm, var_yaw_arm, max_roll_arm, max_picth_arm, max_yaw_ar m,min roll arm,min pitch arm,min yaw arm,amplitude roll arm,amplitude pitch arm,amplitude yaw arm,max roll d umbbell, max picth dumbbell, min roll dumbbell, min pitch dumbbell, amplitude roll dumbbell, amplitude pitch dumbbell, var accel dumbbell, avg roll dumbbell, stddev roll dumbbell, var roll dumbbell, avg pitch dumbbell, s tddev_pitch_dumbbell,stddev_pitch_dumbbell,var_pitch_dumbbell,avg_yaw_dumbbell, stddev_yaw_dumbbell, var_yaw _dumbbell, max_roll_forearm,max_picth_forearm, min_roll_forearm, min_pitch_forearm, amplitude_roll_forearm, amplitude pitch forearm, var_accel forearm, avg roll forearm, stddev_roll forearm, var_roll_forearm, avg pitch forearm, stddev_pitch fo rearm, var pitch forearm, avg yaw forearm, stddev yaw forearm, var yaw forearm))

We will build our predictive model using trees. For this purpose we will be comparing the R functions rpart (CART algorithm) and ctree (CHAID algorithm). As you can see above, the original training sample was split into training and validation, so that we can validate the performance of the models.

We have excluded any irrelevant predictors or predictors with too many missing values.

Model selection

Both, for rpart and ctree, we have decided to set \(minbucket=100\) - in order to have more stability out-of-sample).

We begin by using the ctree function. We calculate the accuracy both, on the training sample and on the validation sample:

```
## [1] 0.7159017
```

```
pred <- predict(modFit1,newdata=validation, type="response")
mean(pred==validation$classe)</pre>
```

```
## [1] 0.695078
```

Next we use the rpart function which has a built-in cross validation procedure:

```
modFit2 <- rpart(classe~.,data=training, control = rpart.control(minbucket = 100,cp=-1))
pred <- predict(modFit2,newdata=training, type="class")
mean(pred==training$classe)</pre>
```

```
## [1] 0.7988543
```

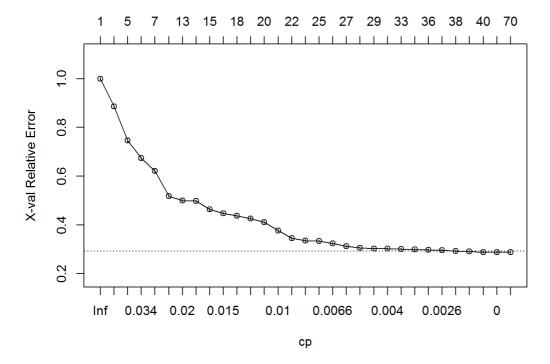
```
pred <- predict(modFit2,newdata=validation, type="class")
mean(pred==validation$classe)</pre>
```

```
## [1] 0.786143
```

The model modFit2 needs to be pruned, since we have selected \(cp=-1\). In order to decide where to prune, we visualize the relative cverrors:

```
plotcp(modFit2)
```





We choose to prune at \(cp=0.0052\) and hence reduce the number of nodes from 70 to 27. We obtain

```
modFit2_pruned <- prune.rpart(modFit2,cp=0.0052)
pred <- predict(modFit2_pruned,newdata=training, type="class")
mean(pred==training$classe)</pre>
```

[1] 0.7668044

pred <- predict(modFit2_pruned,newdata=validation, type="class")
mean(pred==validation\$classe)</pre>

[1] 0.7506431

plotcp(modFit2_pruned)

size of tree 2 5 6 7 12 14 16 19 21 24 26 0.1 X-val Relative Error 0.8 9.0 0.4 0.2 0.047 0.026 0.02 0.018 0.014 0.012 0.0072 0.0058 Inf

We choose modFit2_pruned to be our final model.

```
#fancyRpartPlot(modFit2_pruned, cex=0.15)
rpart.plot(modFit2_pruned,tweak=1.7)
               - C
- D
- E
                                                                                                                                                             .28 .19 .18 .16 .18
100%
                                                                                      A
.31 .21 .19 .18 .11
92%
                                                                                                                                                                   A .24 .23 .21 .20 .12 84%
                                                                                                                                                                                                                    .03 .51 .04 .23 .19
13%
                                                                                                                  .28 .18 .24 .19 .11
71%
                                                                                                                                                                     08 .18 .33 .23 .18
27%
                                                                                                                                                                                    .05 .31 .16 .14 .35
7%
                                                                                         D
.24 .12 .24 .29 .11
                                                                                                        D
20 .15 .12 .38 .14
15%
                                                                         .37 .02 .59 .01 .0
5%
                                                                                               .26 .10 .17 .43 .04
11%
                                                                                         .45 .12 .15 .23 .05
6%
                   81 08 06 19 05 08 08 18 88 03 03 11 00 30 58 82 07 00 04 07 03 14 76 03 03 0 58 85 5% 86 03 03 11 00 30 58 82 07 00 04 07 03 14 76 03 03 03 00 07 78 12 07 14 11 17 51 05 77 07 02 08 00 14 00 76 10
            99 01 00 00 00 02 26 63 00 09 01 152 01 151 13 02 86 80 03 02 09 03 86 02 00 09 03 86 02 00 09 03 86 02 00 00 22 63 35 30 06 03 74 01 06 14 01 04 88 03 04 11 80 14 00 00 00 03 07 52 14 25 02 54 23 10 11 06 13 04 88 08 00 00 00 00 00 100 01 00 00 09 99
```

The predictions on the testing sample are

```
testing$magnet_forearm_z <- as.numeric(testing$magnet_forearm_z)
predict(modFit2_pruned, newdata=testing, type="class")

## 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20
## B A B D A C D D A A C B C A E E A D D B
## Levels: A B C D E
```

The expected out-of-sample error should be close to the error of the validation set (ca. 0.25).