

ParticalMachineLeaning_couse4

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Overview

This reports describes that

- 1.Preprocess Remove the colums which includes NA valume more than 80%. Also, check the zero convariates. 2.Create prediction model.Be aware of multi-corrleation. The methods of dicision tree,random fores, and boosting will be used.
- 3.Cross validation with training set. In sample versus out of sample error, prevent from overfitting.
- 4.The resons of the prediction model choice
- 5.Test the data with the prediction model

Method

The data was obtained from following Websites. The training data for this project are available here: <https://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv> (<https://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv>) The test data are available here: <https://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv> (<https://d396qusza40orc.cloudfront.net/predmachlearn/pml-testing.csv>)

The data for this project come from this source:

<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>).

Results

install library

```
library(caret)
```

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

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

```
library(ISLR)
library(rattle)
```

```
## Rattle: A free graphical interface for data science with R.
## バージョン 5.3.0 Copyright (c) 2006-2018 Togaware Pty Ltd.
## 'rattle()' と入力して、データを多角的に分析します。
```

```
library(doParallel)
```

```
## Loading required package: foreach
```

```
## Loading required package: iterators
```

```
## Loading required package: parallel
```

```
library(tictoc)
library(e1071)
```

set the training/test data.

```
train<-read.csv("/Users/yuki/Documents/R/Cousera_PML/pml-training.
csv",header = T,na.strings = c("#DIV/0!","", "NA"))
test<-read.csv("/Users/yuki/Documents/R/Cousera_PML/pml-testing.cs
v",header = T,na.strings= c("#DIV/0!","", "NA"))
```

remove the columns which includes NA volume more than 80%.

```
# Following column names are important for later analysis.
y1<-names(train[c(1:7,160)])

na.ratio<-function(x){sum(is.na(x)=="TRUE")/length(x)}
y<-apply(train[,-c(1:7,160)],2,na.ratio)
y2<-names(subset(y,y<0.8))

train2<-data.frame(train[160],train[y2])
```

removing zero convariates

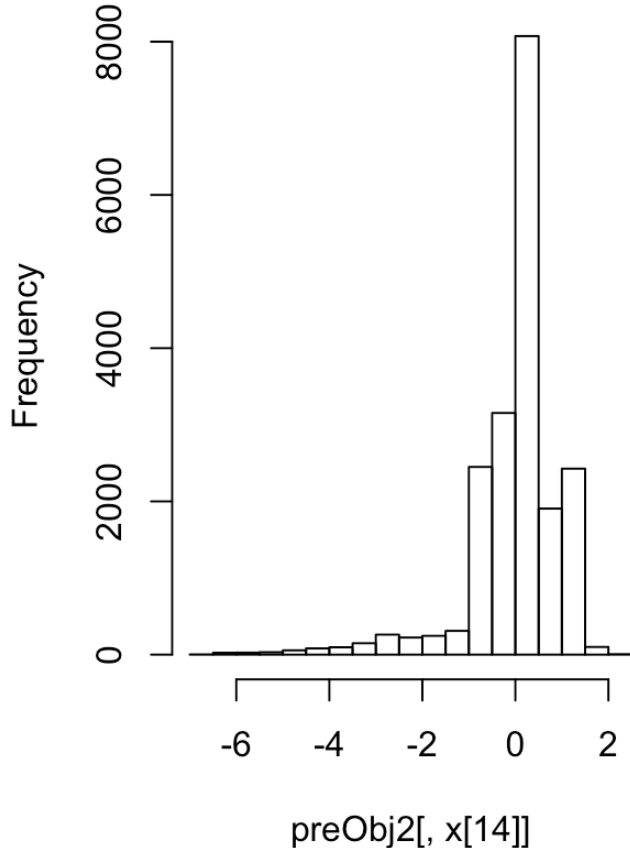
```
preObj<-preProcess(train2,method=c("center","scale"))
predict(preObj,train2)->preObj2

nsv<-nearZeroVar(train2,saveMetrics = T)
nsv2<-subset(nsv,nzv=="FALSE")
train3<-(train[,rownames(nsv2)])

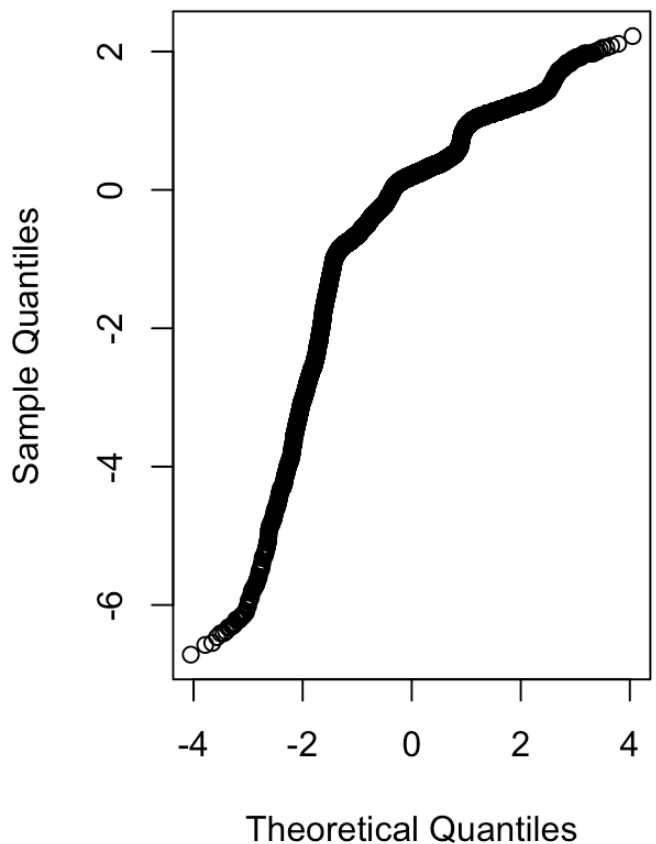
x<-0
for (i in 1:nrow(nsv2)){
  x<-append(x,grep(rownames(nsv2)[i],colnames(train3)))
}

# check the histogram
par(mfrow=c(1,2))
hist(preObj2[,x[14]])
qqnorm(preObj2[,x[14]])
```

Histogram of preObj2[, x[14]]



Normal Q-Q Plot



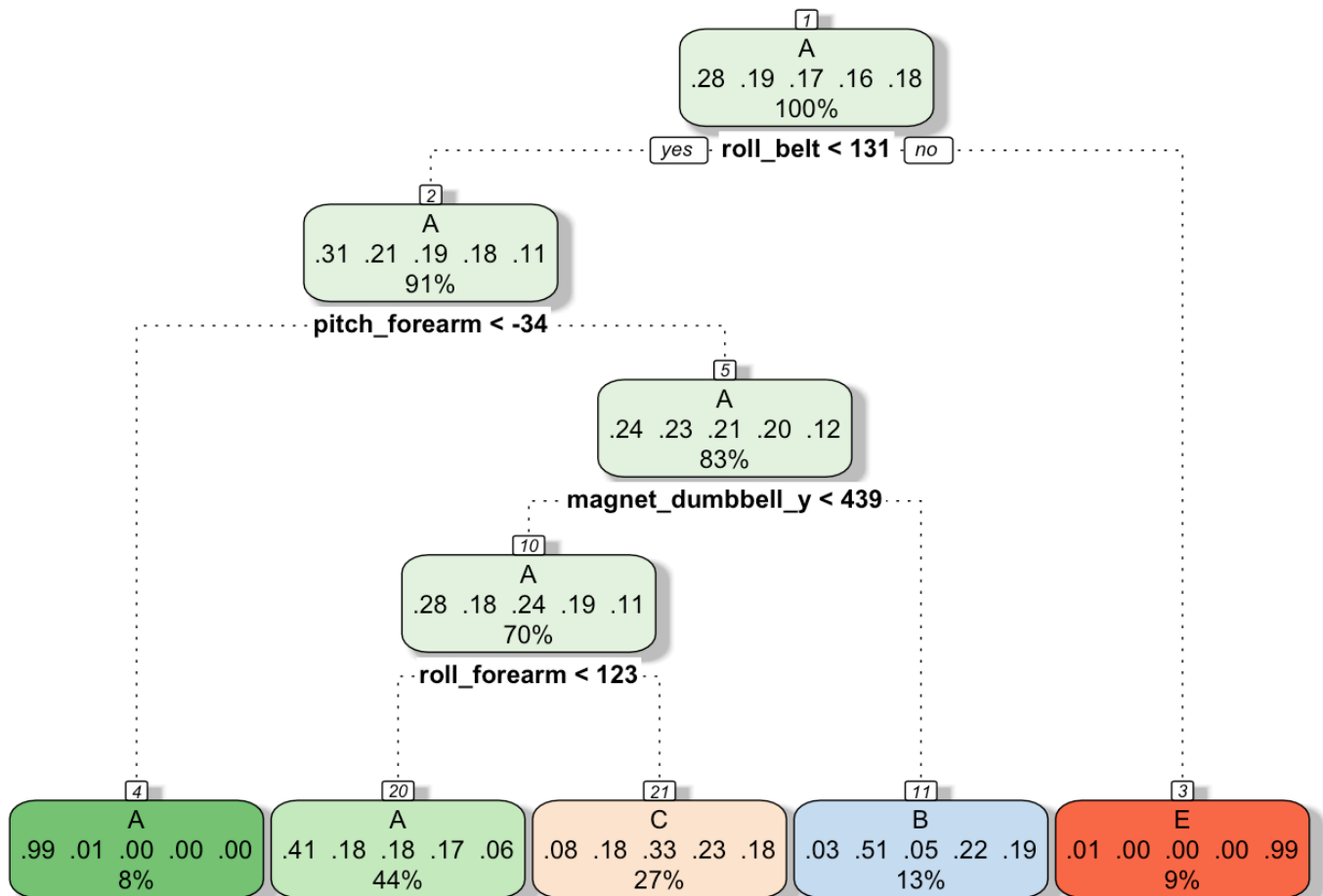
```
test2<-data.frame(test[y2])
```

Create the model “Predicting with trees”

```
cl <- makePSOCKcluster(4)
registerDoParallel(cl)
tic()
```

```
#folds<-createFolds(y=train2$classe,k=10,list=T,returnTrain =F)
trainPart <- createDataPartition(train3$classe, p=0.70, list=F)
trainSubset <- train3[trainPart, ]
validSubset <- train3[-trainPart, ]
```

```
modFit<-train(classe~.,data=trainSubset,method="rpart")
fancyRpartPlot(modFit$finalModel)
```



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```
Pred<-predict(modFit,validSubset)
table(Pred,validSubset$classe)
```

```
##  
## Pred      A      B      C      D      E  
##      A 1527   469   481   428   147  
##      B   29   384    25   168   149  
##      C  117   286   520   368   313  
##      D    0     0     0     0     0  
##      E    1     0     0     0   473
```

```
toc()
```

```
## 13.049 sec elapsed
```

```
stopCluster(cl)
```

Create the model “Random Forest”

```
cl <- makePSOCKcluster(4)  
registerDoParallel(cl)  
tic()  
  modFit2<-train(classe~.,data=trainSubset,method="rf",prox=T)  
  modFit2
```

```
## Random Forest
##
## 13737 samples
##    52 predictor
##    5 classes: 'A', 'B', 'C', 'D', 'E'
##
## No pre-processing
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 13737, 13737, 13737, 13737, 13737, 13737
## , ...
## Resampling results across tuning parameters:
##
##   mtry  Accuracy   Kappa
##    2    0.9885105  0.9854658
##   27    0.9886352  0.9856246
##   52    0.9796360  0.9742395
##
## Accuracy was used to select the optimal model using the largest v
alue.
## The final value used for the model was mtry = 27.
```

```
Pred2<-predict(modFit2,validSubset)
table(Pred2,validSubset$classe)
```

```
##
## Pred2      A      B      C      D      E
##   A 1673    12      0      0      0
##   B   0 1125      5      0      0
##   C   1   2 1018     11      3
##   D   0   0   3  953      8
##   E   0   0   0   0 1071
```

```
toc()
```

```
## 4197.716 sec elapsed
```

```
stopCluster(cl)
```

Create the model “Boosting”

```

cl <- makePSOCKcluster(4)
registerDoParallel(cl)
tic()
  modFit3<-train(classe~.,data=trainSubset,method="gbm",verbose=F)
  modFit3

```

```

## Stochastic Gradient Boosting
##
## 13737 samples
##    52 predictor
##    5 classes: 'A', 'B', 'C', 'D', 'E'
##
## No pre-processing
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 13737, 13737, 13737, 13737, 13737, 13737
, ...
## Resampling results across tuning parameters:
##
##  interaction.depth  n.trees  Accuracy  Kappa
##  1                  50      0.7492410  0.6820228
##  1                  100      0.8164125  0.7675400
##  1                  150      0.8498122  0.8098757
##  2                   50      0.8525329  0.8130941
##  2                  100      0.9039753  0.8784375
##  2                  150      0.9280041  0.9088712
##  3                   50      0.8944516  0.8663303
##  3                  100      0.9382539  0.9218437
##  3                  150      0.9571930  0.9458308
##
## Tuning parameter 'shrinkage' was held constant at a value of 0.1
##
## Tuning parameter 'n.minobsinnode' was held constant at a value of
10
## Accuracy was used to select the optimal model using the largest v
alue.
## The final values used for the model were n.trees = 150, interacti
on.depth =
##  3, shrinkage = 0.1 and n.minobsinnode = 10.

```

```

Pred3<-predict(modFit3,validSubset)
table(Pred3,validSubset$classe)

```

```
##
## Pred3      A      B      C      D      E
##      A 1646      45      1      1      3
##      B   18 1051      24      3      8
##      C    6   40  989     33      9
##      D    3    3   11  920     20
##      E    1    0    1    7 1042
```

```
toc()
```

```
## 314.567 sec elapsed
```

```
stopCluster(cl)
```

Conclusion

Compared the decision tree, random forest, and boosting methods, the random forest shows the best accuracy. The 52 predictors which has less than 50 % NA value in each column was adapted for these models. Therefore, it is concluded that the random forest as used this dataset, and the prediction with test set shows below.

```
predict(modFit2, test2)
```

```
##  [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
```

Supplymental data

No data.