

```
In [1]: library(caret)
```

```
Warning message:  
"package 'caret' was built under R version 3.6.3"  
Loading required package: lattice  
Loading required package: ggplot2  
Warning message:  
"package 'ggplot2' was built under R version 3.6.3"
```

```
In [2]: library(rpart)
```

```
In [3]: library(rpart.plot)
```

```
Error in library(rpart.plot): there is no package called 'rpart.plot'  
Traceback:
```

```
1. library(rpart.plot)
```

```
In [4]: install.packages("rpart.plot", repos='http://cran.us.r-project.org')
```

```
package 'rpart.plot' successfully unpacked and MD5 sums checked
```

```
The downloaded binary packages are in  
C:\Users\Acer\AppData\Local\Temp\RtmpgFPMX\downloaded_packages
```

```
In [5]: library(rpart.plot)
```

```
Warning message:  
"package 'rpart.plot' was built under R version 3.6.3"
```

```
In [6]: library(rpart.plot)
```

```
Warning message:  
"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
```

```
In [7]: library(corrplot)
```

```
Error in library(corrplot): there is no package called 'corrplot'  
Traceback:
```

```
1. library(corrplot)
```

```
In [8]: install.packages("corrplot", repos='http://cran.us.r-project.org')
```

```
package 'corrplot' successfully unpacked and MD5 sums checked
```

```
The downloaded binary packages are in  
C:\Users\Acer\AppData\Local\Temp\RtmpgFPMMX\downloaded_packages
```

```
In [9]: library(corrplot)
```

```
Warning message:  
"package 'corrplot' was built under R version 3.6.3"corrplot 0.84 loaded
```

```
In [10]: trainingURL <- "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-  
training.csv"  
testingURL <- "https://d396qusza40orc.cloudfront.net/predmachlearn/pml-  
testing.csv"  
trainingFile <- "./data/pml-training.csv"  
testingFile <- "./data/pml-testing.csv"  
if (!file.exists("./data")) {  
  dir.create("./data")  
}
```

```
if (!file.exists(trainingFile)) {  
  download.file(trainingURL, destfile=trainingFile, method="curl")  
}  
if (!file.exists(testingFile)) {  
  download.file(testingURL, destfile=testingFile, method="curl")  
}
```

```
In [11]: trainRaw <- read.csv("./data/pml-training.csv")  
testRaw <- read.csv("./data/pml-testing.csv")  
dim(trainRaw)
```

19622 160

```
In [12]: dim(testRaw)
```

20 160

```
In [13]: sum(complete.cases(trainRaw))
```

406

```
In [14]: trainRaw <- trainRaw[, colSums(is.na(trainRaw)) == 0]
```

```
In [15]: testRaw <- testRaw[, colSums(is.na(testRaw)) == 0]
```

```
In [16]: classe <- trainRaw$classe  
trainRemove <- grepl("^X|timestamp|window", names(trainRaw))  
trainRaw <- trainRaw[, !trainRemove]  
trainCleaned <- trainRaw[, sapply(trainRaw, is.numeric)]  
trainCleaned$classe <- classe  
testRemove <- grepl("^X|timestamp|window", names(testRaw))  
testRaw <- testRaw[, !testRemove]  
testCleaned <- testRaw[, sapply(testRaw, is.numeric)]
```

```
In [17]: set.seed(2108) # For reproducible purpose  
inTrain <- createDataPartition(trainCleaned$classe, p=0.70, list=F)
```

```
trainData <- trainCleaned[inTrain, ]
testData <- trainCleaned[-inTrain, ]
```

```
In [18]: controlRf <- trainControl(method="cv", 5)
modelRf <- train(classe ~ ., data=trainData, method="rf", trControl=controlRf, ntree=250)
modelRf
```

Error: package e1071 is required
Traceback:

```
1. train(classe ~ ., data = trainData, method = "rf", trControl = controlRf,
.      ntree = 250)
2. train.formula(classe ~ ., data = trainData, method = "rf", trControl = controlRf,
.      ntree = 250)
3. train(x, y, weights = w, ...)
4. train.default(x, y, weights = w, ...)
5. evalSummaryFunction(y, wts = weights, ctrl = trControl, lev = classLevels,
.      metric = metric, method = method)
6. ctrl$summaryFunction(testOutput, lev, method)
7. postResample(data[, "pred"], data[, "obs"])
8. requireNamespaceQuietStop("e1071")
9. stop(paste("package", package, "is required"), call. = FALSE)
```

```
In [19]: install.packages("e1071", repos='http://cran.us.r-project.org')
```

package 'e1071' successfully unpacked and MD5 sums checked

The downloaded binary packages are in
C:\Users\Acer\AppData\Local\Temp\RtmpgFPMX\downloaded_packages

```
In [20]: controlRf <- trainControl(method="cv", 5)
modelRf <- train(classe ~ ., data=trainData, method="rf", trControl=controlRf, ntree=250)
modelRf
```

Random Forest

```
13737 samples
 52 predictor
 5 classes: 'A', 'B', 'C', 'D', 'E'
```

No pre-processing

Resampling: Cross-Validated (5 fold)

Summary of sample sizes: 10989, 10989, 10990, 10990, 10990

Resampling results across tuning parameters:

mtry	Accuracy	Kappa
2	0.9892259	0.9863692
27	0.9913372	0.9890413
52	0.9826747	0.9780829

Accuracy was used to select the optimal model using the largest value.
The final value used for the model was mtry = 27.

```
In [21]: predictRf <- predict(modelRf, testData)
confusionMatrix(testData$classe, predictRf)
```

Confusion Matrix and Statistics

		Reference				
Prediction		A	B	C	D	E
A	1671	1	1	0	1	
B	13 1124		2	0	0	
C	0 3 1022			1	0	
D	0 0 7 957				0	
E	0 0 4 3 1075					

Overall Statistics

```
Accuracy : 0.9939
95% CI : (0.9915, 0.9957)
No Information Rate : 0.2862
P-Value [Acc > NIR] : < 2.2e-16
```

Kappa : 0.9923

McNemar's Test P-Value : NA

Statistics by Class:

	Class: A	Class: B	Class: C	Class: D	Class: E
Sensitivity	0.9923	0.9965	0.9865	0.9958	0.9991
Specificity	0.9993	0.9968	0.9992	0.9986	0.9985
Pos Pred Value	0.9982	0.9868	0.9961	0.9927	0.9935
Neg Pred Value	0.9969	0.9992	0.9971	0.9992	0.9998
Prevalence	0.2862	0.1917	0.1760	0.1633	0.1828
Detection Rate	0.2839	0.1910	0.1737	0.1626	0.1827
Detection Prevalence	0.2845	0.1935	0.1743	0.1638	0.1839
Balanced Accuracy	0.9958	0.9967	0.9928	0.9972	0.9988

```
In [22]: accuracy <- postResample(predictRf, testData$classe)
accuracy
```

Accuracy 0.993882752761257
Kappa 0.992261060800132

```
In [23]: oose <- 1 - as.numeric(confusionMatrix(testData$classe, predictRf)$overall[1])
oose
```

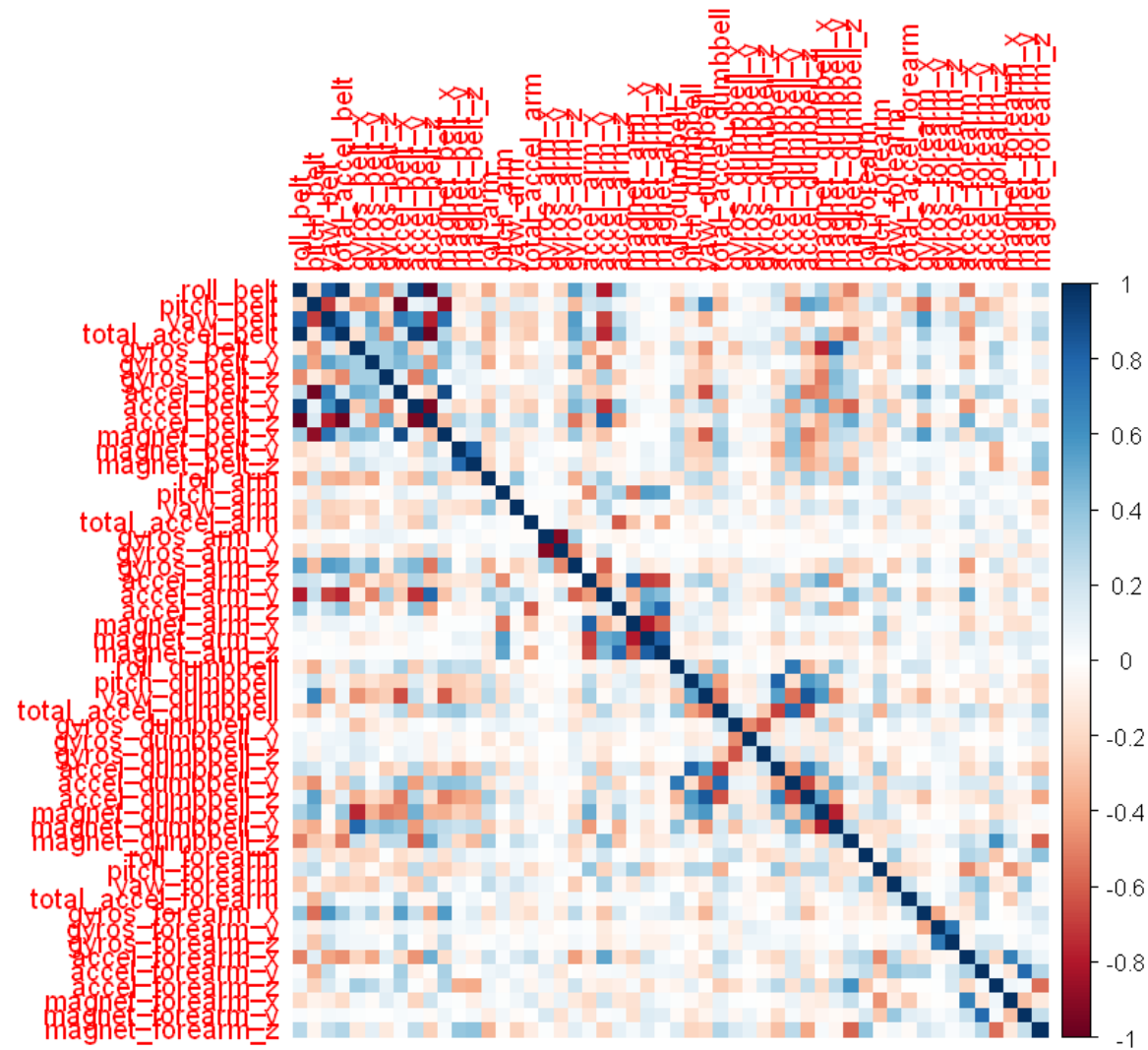
0.00611724723874252

```
In [24]: result <- predict(modelRf, testCleaned[, -length(names(testCleaned))])
result
```

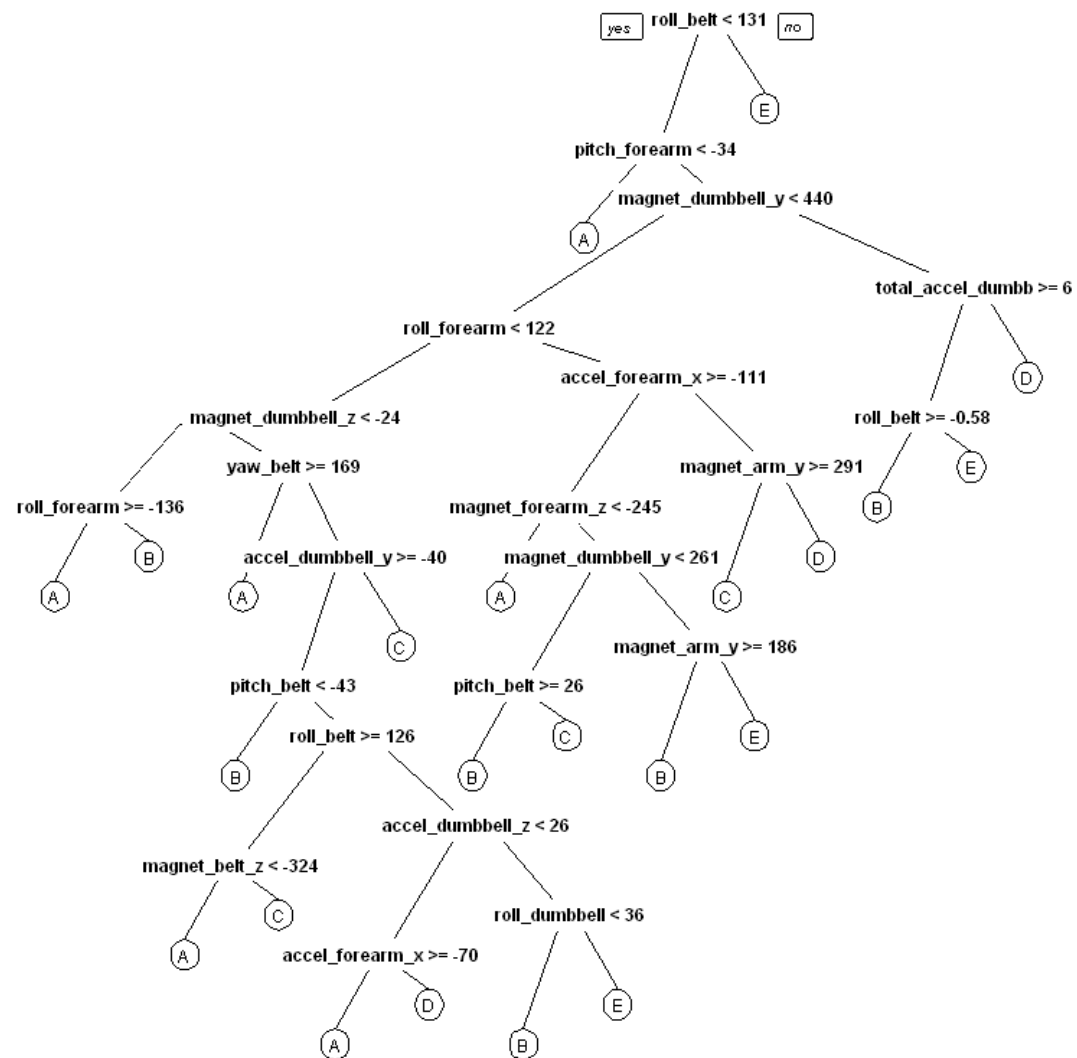
B A B A A E D B A A B C B A E E A B B B

► **Levels:**

```
In [25]: corrPlot <- cor(trainData[, -length(names(trainData))])
corrplot(corrPlot, method="color")
```



```
In [26]: treeModel <- rpart(classe ~ ., data=trainData, method="class")
prp(treeModel)
```



In []: