<!DOCTYPE html>

```
<html xmlns="http://www.w3.org/1999/xhtml">
<head>
<meta charset="utf-8">
<meta http-equiv="Content-Type" content="text/html; charset=utf-8" />
<meta name="generator" content="pandoc" />
<meta name="author" content="Onur Akpolat" />
<title>Predict activity quality from activity monitors</title>
<script src="data:application/x-javascript,%2F%2A%21%20jQuery%20v1%2E11%2E0%20%7C%20%28c%29%202005%2C%202014%20jQuery%20Foundation%2C%20Inc%2E%20%7C</pre>
<meta name="viewport" content="width=device-width, initial-scale=1.0" />
k href="data:text/css,%2F%2A%21%0A%20%2A%20Bootstrap%20v2%2E3%2E2%0A%20%2A%20M20%2A%20Copyright%202013%20Twitter%2C%20Inc%0A%20%2A%20Licensed%20
<link href="data:text/css,%2F%2A%21%0A%20%2A%20Bootstrap%20Responsive%20v2%2E3%2E2%0A%20%2A%0A%20%2A%20Copyright%202013%20Twitter%2C%20Inc%0A%20%2A%</pre>
<script src="data:application/x-javascript,%2F%2A%21%0A%2A%20Bootstrap%2Ejs%20by%20%40fat%20%26%20%40mdo%0A%2A%20Copyright%202013%20Twitter%2C%20Inc</pre>
<style type="text/css">code{white-space: pre;}</style>
k href="data:text/css,pre%20%2Eoperator%2C%0Apre%20%2Eparen%20%7B%0A%20color%3A%20rgb%28104%2C%20118%2C%20135%29%0A%7D%0A%0Apre%20%2Eliteral%20%
<script src="data:application/x-javascript,%0Avar%20hljs%3Dnew%20function%28%29%7Bfunction%20m%28p%29%7Breturn%20p%2Ereplace%28%2F%26%2Fgm%2C%22%26a</pre>
<style type="text/css">
 pre:not([class]) {
   background-color: white;
 }
</style>
<script type="text/javascript">
if (window.hljs && document.readyState && document.readyState === "complete") {
  window.setTimeout(function() {
     hljs.initHighlighting();
  }, 0);
</script>
</head>
<body>
<style type="text/css">
.main-container {
 max-width: 940px;
 margin-left: auto;
 margin-right: auto;
<div class="container-fluid main-container">
<div id="header">
<h1 class="title">Predict activity quality from activity monitors</h1>
<h4 class="author"><em>Onur Akpolat</em></h4>
<h4 class="date"><em>
January 2015
</em></h4>
</div>
<div id="synopsis" class="section level2">
Using devices such as Jawbone Up, Nike FuelBand, and Fitbit it is now possible to collect a large amount of data about personal activity relative
The goal of this project is to predict the manner in which they did the exercise. This is the <code>classe</code> variable in the training set.
</div>
<div id="data-description" class="section level2">
The outcome variable is <code>classe</code>, a factor variable with 5 levels. For this data set, participants were asked to perform one set of 10
exactly according to the specification (Class A)
throwing the elbows to the front (Class B)
```

```
lifting the dumbbell only halfway (Class C)
lowering the dumbbell only halfway (Class D)
throwing the hips to the front (Class E)
</div>
<div id="initial-configuration" class="section level2">
<h2>Initial configuration</h2>
The initial configuration consists of loading some required packages and initializing some variables.
<code>#Data variables
training.file < - './data/pml-training.csv'
test.cases.file < - './data/pml-testing.csv'
training.url <- 'http://d396qusza40orc.cloudfront.net/predmachlearn/pml-training.csv'
test.cases.url &lt:- 'http://d396gusza40orc.cloudfront.net/predmachlearn/pml-testing.csv'
if (!file.exists("data")){
 dir.create("data")
if (!file.exists("data/submission")){
 dir.create("data/submission")
}
#R-Packages
IscaretInstalled <- require(&quot;caret&quot;)</code>
<code>## Loading required package: caret
## Loading required package: lattice
## Loading required package: ggplot2</code>
<code>if(!IscaretInstalled){
   install.packages("caret")
   library("caret")
IsrandomForestInstalled <- require(&quot;randomForest&quot;)</code>
<code>## Loading required package: randomForest
## randomForest 4.6-10
## Type rfNews() to see new features/changes/bug fixes.</code>
<code>if(!IsrandomForestInstalled){
   install.packages("randomForest")
   library("randomForest")
   }
IsRpartInstalled <- require(&quot;rpart&quot;)</code>
<code>## Loading required package: rpart</code>
<code>if(!IsRpartInstalled){
   install.packages("rpart")
   library(&quot:rpart&quot:)
IsRpartPlotInstalled <- require(&quot;rpart.plot&quot;)</code>
<code>## Loading required package: rpart.plot</code>
<code>if(!IsRpartPlotInstalled){
   install.packages("rpart.plot")
   library("rpart.plot")
# Set seed for reproducability
set.seed(9999)</code>
</div>
<div id="data-processing" class="section level2">
<h2>Data processing</h2>
In this section the data is downloaded and processed. Some basic transformations and cleanup will be performed, so that <code>NA</code> values at
The <code>pml-training.csv</code> data is used to devise training and testing sets. The <code>pml-test.csv</code> data is used to predict and ans
<code># Download data
download.file(training.url, training.file)
download.file(test.cases.url,test.cases.file )
# Clean data
training \\ \< -read.csv(training.file, na.strings = c(\&quot; NA\&quot;, \&quot; \#DIV/0!\&quot;, \&quot; \&quot;))
testing \ \< -read.csv(test.cases.file \ , \ na.strings = c(\&quot; NA\&quot;, \&quot; \#DIV/0!\&quot;, \&quot; \&quot;))
training< -training[,colSums(is.na(training)) == 0]
testing <-testing[,colSums(is.na(testing)) == 0]
# Subset data
```

```
training <-training[,-c(1:7)]
testing <-testing[,-c(1:7)]</code>
</div>
<div id="cross-validation" class="section level2">
In this section cross-validation will be performed by splitting the training data in training (75%) and testing (25%) data.
<code>subSamples &lt;- createDataPartition(y=training$classe, p=0.75, list=FALSE)
subTraining <- training[subSamples, ]
subTesting < - training[-subSamples, ]</code>
<div id="expected-out-of-sample-error" class="section level2">
<h2>Expected out-of-sample error</h2>
<div id="exploratory-analysis" class="section level2">
<h2>Exploratory analysis</h2>
<code>plot(subTraining$classe, col=&quot;orange&quot;, main=&quot;Levels of the variable classe&quot;, xlab=&quot;classe levels&quot;
<img src=">
The plot above shows that Level A is the most frequent classe. D appears to be the least frequent one.
</div>
<div id="prediction-models" class="section level2">
<h2>Prediction models</h2>
In this section a decision tree and random forest will be applied to the data.
<div id="decision-tree" class="section level3">
<h3>Decision tree</h3>
<code># Fit model
modFitDT <- rpart(classe ~ ., data=subTraining, method=&quot;class&quot;)
# Perform prediction
predictDT <- predict(modFitDT, subTesting, type = &quot;class&quot;)
rpart.plot(modFitDT, main="Classification Tree", extra=102, under=TRUE, faclen=0)</code>
Following confusion matrix shows the errors of the prediction algorithm.
<code>confusionMatrix(predictDT, subTesting$classe)</code>
<code>## Confusion Matrix and Statistics
##
##
         Reference
## Prediction A B
                   C D
                            E
##
        A 1266 208 25 91
         B 33 535 71 30
##
        C 28 90 676 130
                           94
##
        D 45 72 59 501 43
##
        E 23 44 24 52 668
##
## Overall Statistics
##
##
              Accuracy: 0.7435
##
               95% CI: (0.731, 0.7557)
##
    No Information Rate : 0.2845
##
    P-Value [Acc > NIR] : < 2.2e-16
##
                Kappa : 0.6738
##
## Mcnemar's Test P-Value : < 2.2e-16
## Statistics by Class:
##
##
                 Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                   0.9075 0.5638 0.7906 0.6231 0.7414
## Specificity
                   0.8994 0.9492 0.9155 0.9466 0.9643
## Pos Pred Value
                   0.7820 0.7269 0.6640 0.6958 0.8237
                  0.9607 0.9007 0.9539 0.9276 0.9431
## Neg Pred Value
## Prevalence
                  0.2845 0.1935 0.1743 0.1639 0.1837
## Detection Rate
                  0.2582 0.1091 0.1378 0.1022 0.1362
## Detection Prevalence 0.3301 0.1501 0.2076 0.1468 0.1654
## Balanced Accuracy 0.9035 0.7565 0.8531 0.7849 0.8528</code>
</div>
<div id="random-forest" class="section level3">
<h3>Random forest</h3>
<code># Fit model
modFitRF <- randomForest(classe ~ ., data=subTraining, method=&quot;class&quot;)
```

```
# Perform prediction
predictRF <- predict(modFitRF, subTesting, type = &quot;class&quot;)</code>
Following confusion matrix shows the errors of the prediction algorithm.
<code>confusionMatrix(predictRF, subTesting$classe)</code>
<code>## Confusion Matrix and Statistics
##
##
           Reference
## Prediction A B C D
          A 1394 2 0 0
          B 1 946 8 0
          C 0 1 846 6
##
                  0 1 796
##
          D
             0
                               1
##
             0 0
                      0 2 900
##
## Overall Statistics
##
               Accuracy : 0.9955
##
                 95% CI : (0.9932, 0.9972)
##
    No Information Rate : 0.2845
##
     P-Value [Acc > NIR] : < 2.2e-16
##
##
                  Kappa: 0.9943
## Mcnemar's Test P-Value : NA
##
## Statistics by Class:
##
##
                   Class: A Class: B Class: C Class: D Class: E
## Sensitivity
                    0.9993 0.9968 0.9895 0.9900 0.9989
## Specificity
                     0.9994 0.9977 0.9983 0.9995 0.9995
## Pos Pred Value
                     0.9986 0.9906 0.9918 0.9975 0.9978
## Neg Pred Value
                      0.9997 0.9992 0.9978 0.9981 0.9998
## Prevalence
                      0.2845 0.1935 0.1743 0.1639
                                                     0.1835
                      0.2843 0.1929 0.1725 0.1623
## Detection Rate
## Detection Prevalence 0.2847 0.1947 0.1739 0.1627 0.1839
## Balanced Accuracy 0.9994 0.9973 0.9939 0.9948 0.9992</code>
</div>
<div id="conclusion" class="section level2">
<h2>Conclusion</h2>
<div id="result" class="section level3">
<h3>Result</h3>
The confusion matrices show, that the Random Forest algorithm performens better than decision trees. The accuracy for the Random Forest model was
<div id="expected-out-of-sample-error-1" class="section level3">
<h3>Expected out-of-sample error</h3>
The expected out-of-sample error is estimated at 0.005, or 0.5%. The expected out-of-sample error is calculated as 1 - accuracy for predictions n
</div>
</div>
<div id="submission" class="section level2">
<h2>Submission</h2>
In this section the files for the project submission are generated using the random forest algorithm on the testing data.
<code># Perform prediction
predictSubmission <- predict(modFitRF, testing, type=&quot;class&quot;)
predictSubmission</code>
<code>## 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20
## 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</code>
<code># Write files for submission
pml_write_files = function(x){
 n = length(x)
 for(i in 1:n){
   filename = paste0("./data/submission/problem_id_",i,".txt")
   write.table(x[i],file=filename,quote=FALSE,row.names=FALSE,col.names=FALSE)
 }
}
pml_write_files(predictSubmission)</code>
</div>
</div>
```

```
<script>
// add bootstrap table styles to pandoc tables
$(document).ready(function () {
 $('tr.header').parent('thead').parent('table').addClass('table table-condensed');
});
</script>
<!-- dynamically load mathjax for compatibility with self-contained -->
<script>
 (function () {
   var script = document.createElement("script");
    script.type = "text/javascript";
    script.src = "https://cdn.mathjax.org/mathjax/latest/MathJax.js?config=TeX-AMS-MML_HTMLorMML";
   document.getElementsByTagName("head")[0].appendChild(script);
 })();
</script>
</body>
</html>
```