# Introduction to Machine Learning

### Lab 2 Block 2

### Rasmus Holm

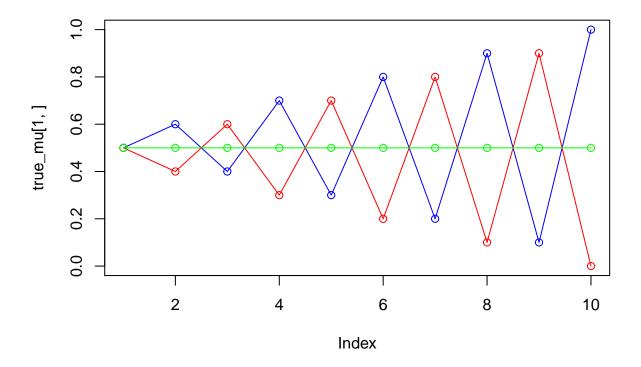
#### 2016-12-01

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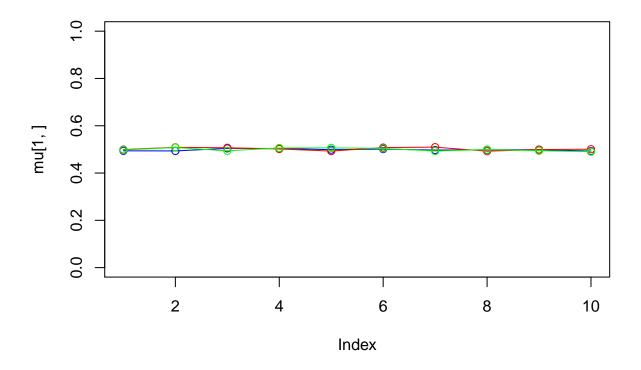
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## Assignment 1a

### Assignment 1b



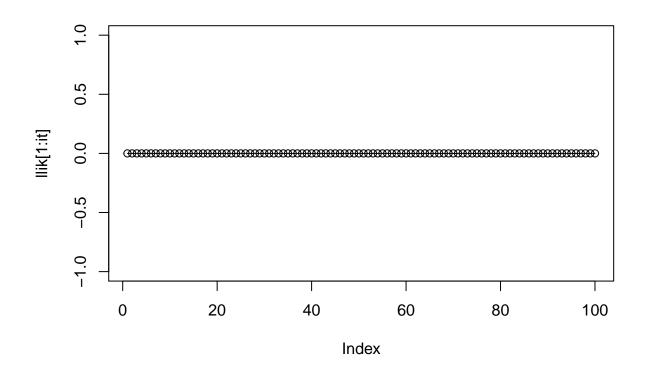
```
#> [1] 0.3326090 0.3336558 0.3337352
             [,1]
                       [,2]
                                  [,3]
                                            [,4]
                                                      [,5]
                                                                [,6]
#> [1,] 0.4939877 0.4935375 0.5042511 0.5040286 0.4987810 0.5012754 0.4971036
#> [2,] 0.4993719 0.5088453 0.5068730 0.5016720 0.4929275 0.5077146 0.5095075
#> [3,] 0.4975302 0.5077926 0.4939841 0.5059821 0.5063490 0.5041462 0.4929400
#>
             [,8]
                       [,9]
                                 [,10]
#> [1,] 0.4982144 0.4987654 0.4929075
#> [2,] 0.4924574 0.4992470 0.5008651
#> [3,] 0.4992362 0.4943482 0.4903974
```



```
#> iteration:
               1 log likelihood:
                                   FALSE
               2 log likelihood:
#> iteration:
                                   FALSE
#> iteration:
               3 log likelihood:
                                   FALSE
#> iteration:
               4 log likelihood:
                                   FALSE
#> iteration:
               5 log likelihood:
                                   FALSE
#> iteration:
               6 log likelihood:
                                   FALSE
#> iteration:
               7 log likelihood:
                                   FALSE
#> iteration:
               8 log likelihood:
                                   FALSE
#> iteration:
               9 log likelihood:
                                   FALSE
               10 log likelihood:
#> iteration:
                                   FALSE
#> iteration:
               11 log likelihood:
                                    FALSE
               12 log likelihood:
#> iteration:
                                    FALSE
#> iteration:
               13 log likelihood:
                                    FALSE
#> iteration:
               14 log likelihood:
                                    FALSE
#> iteration:
               15 log likelihood:
                                    FALSE
               16 log likelihood:
                                    FALSE
#> iteration:
#> iteration:
               17 log likelihood:
                                    FALSE
#> iteration:
               18 log likelihood:
                                    FALSE
#> iteration:
               19 log likelihood:
                                    FALSE
               20 log likelihood:
#> iteration:
                                    FALSE
               21 log likelihood:
#> iteration:
                                    FALSE
#> iteration:
               22 log likelihood:
                                    FALSE
#> iteration:
               23 log likelihood:
                                    FALSE
#> iteration:
               24 log likelihood:
                                    FALSE
#> iteration:
               25 log likelihood:
                                    FALSE
#> iteration: 26 log likelihood:
```

```
#> iteration:
              27 log likelihood: FALSE
              28 log likelihood: FALSE
#> iteration:
#> iteration:
              29 log likelihood:
              30 log likelihood:
                                  FALSE
#> iteration:
#> iteration:
              31 log likelihood:
                                  FALSE
#> iteration:
              32 log likelihood:
                                  FALSE
#> iteration:
              33 log likelihood:
                                  FALSE
#> iteration:
              34 log likelihood:
                                  FALSE
#> iteration:
               35 log likelihood: FALSE
#> iteration:
               36 log likelihood:
                                  FALSE
#> iteration:
              37 log likelihood:
                                  FALSE
#> iteration:
               38 log likelihood:
                                  FALSE
#> iteration:
               39 log likelihood: FALSE
               40 log likelihood: FALSE
#> iteration:
#> iteration:
               41 log likelihood: FALSE
#> iteration:
               42 log likelihood: FALSE
#> iteration:
               43 log likelihood: FALSE
#> iteration:
               44 log likelihood:
                                  FALSE
#> iteration:
              45 log likelihood:
                                  FALSE
#> iteration:
              46 log likelihood:
                                  FALSE
#> iteration:
              47 log likelihood:
                                  FALSE
               48 log likelihood:
#> iteration:
               49 log likelihood:
#> iteration:
                                  FALSE
              50 log likelihood:
#> iteration:
                                  FALSE
#> iteration:
              51 log likelihood:
                                  FALSE
#> iteration:
              52 log likelihood:
                                  FALSE
              53 log likelihood:
                                  FALSE
#> iteration:
#> iteration:
              54 log likelihood: FALSE
#> iteration:
               55 log likelihood: FALSE
#> iteration:
               56 log likelihood: FALSE
#> iteration:
               57 log likelihood: FALSE
#> iteration:
               58 log likelihood: FALSE
#> iteration:
               59 log likelihood: FALSE
#> iteration:
              60 log likelihood: FALSE
#> iteration:
               61 log likelihood:
                                  FALSE
#> iteration:
              62 log likelihood: FALSE
#> iteration:
               63 log likelihood:
#> iteration:
              64 log likelihood:
                                  FALSE
#> iteration:
               65 log likelihood:
                                  FALSE
               66 log likelihood:
#> iteration:
                                  FALSE
              67 log likelihood:
#> iteration:
                                   FALSE
#> iteration:
              68 log likelihood: FALSE
#> iteration:
              69 log likelihood: FALSE
              70 log likelihood: FALSE
#> iteration:
#> iteration:
               71 log likelihood: FALSE
               72 log likelihood: FALSE
#> iteration:
#> iteration:
               73 log likelihood: FALSE
#> iteration:
               74 log likelihood: FALSE
#> iteration:
              75 log likelihood: FALSE
#> iteration:
              76 log likelihood: FALSE
#> iteration:
              77 log likelihood: FALSE
#> iteration:
              78 log likelihood: FALSE
#> iteration:
              79 log likelihood: FALSE
#> iteration: 80 log likelihood: FALSE
```

```
#> iteration: 81 log likelihood: FALSE
#> iteration: 82 log likelihood: FALSE
#> iteration: 83 log likelihood: FALSE
#> iteration: 84 log likelihood: FALSE
#> iteration: 85 log likelihood: FALSE
#> iteration: 86 log likelihood: FALSE
#> iteration: 87 log likelihood: FALSE
#> iteration: 88 log likelihood: FALSE
#> iteration: 89 log likelihood: FALSE
#> iteration: 90 log likelihood: FALSE
#> iteration: 91 log likelihood: FALSE
#> iteration: 92 log likelihood: FALSE
#> iteration: 93 log likelihood: FALSE
#> iteration: 94 log likelihood: FALSE
#> iteration: 95 log likelihood: FALSE
#> iteration: 96 log likelihood: FALSE
#> iteration: 97 log likelihood: FALSE
#> iteration: 98 log likelihood: FALSE
#> iteration: 99 log likelihood: FALSE
#> iteration: 100 log likelihood: FALSE
#> [1] 0.3326090 0.3336558 0.3337352
             [,1]
                      [,2]
                                [,3]
                                          [,4]
                                                    [,5]
#> [1,] 0.4939877 0.4935375 0.5042511 0.5040286 0.4987810 0.5012754 0.4971036
#> [2,] 0.4993719 0.5088453 0.5068730 0.5016720 0.4929275 0.5077146 0.5095075
#> [3,] 0.4975302 0.5077926 0.4939841 0.5059821 0.5063490 0.5041462 0.4929400
             [,8]
                      [,9]
                               [,10]
#> [1,] 0.4982144 0.4987654 0.4929075
#> [2,] 0.4924574 0.4992470 0.5008651
#> [3,] 0.4992362 0.4943482 0.4903974
```



### Assignment 2a

### 1

**#>** [1] 37.10301

#### 2

**#>** [1] 30.8038

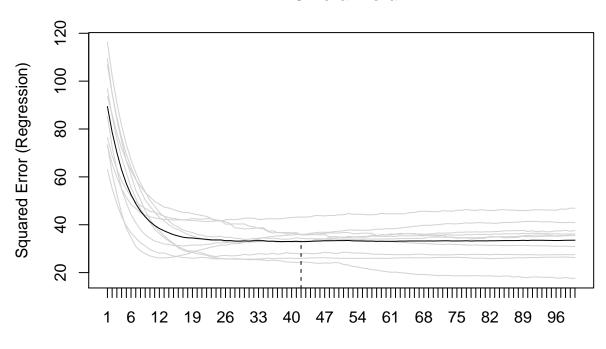
3

## Assignment 2b

### Assignment 3a

1





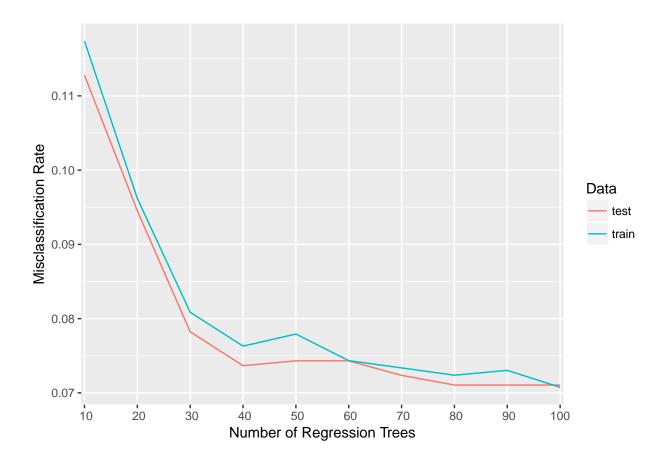
Number of boosting iterations

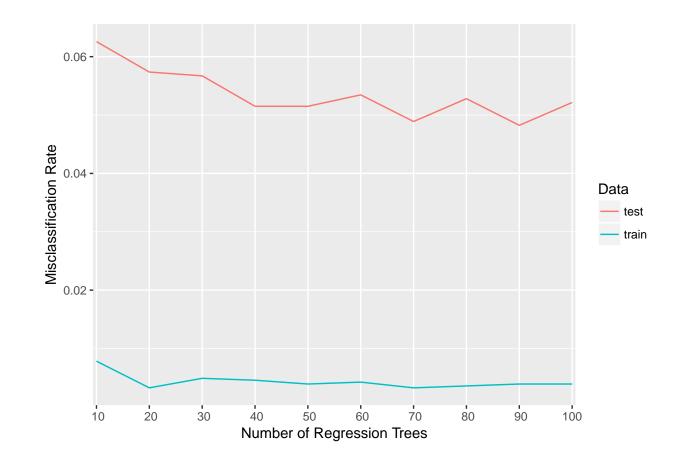
 $\mathbf{2}$ 

**#>** [1] 988.8233

**#>** [1] 1433.941

## Assignment 4a





#### **Appendix**

Code for Assignment 1a

Code for Assignment 1b

Code for Assignment 2a

```
library(tree)
data <- read.csv2("../data/bodyfatregression.csv")</pre>
names(data) <- c("Waist", "Weight", "Bodyfat")</pre>
set.seed(1234567890)
train_idx <- sample(nrow(data), floor(nrow(data) * (2 / 3)))</pre>
train <- data[train_idx,]</pre>
test <- data[-train_idx,]</pre>
set.seed(1234567890)
tree_count <- 100
test_errors <- rep(0, tree_count)</pre>
for (i in 1:tree_count) {
    newdata <- train[sample(nrow(train), replace=TRUE),]</pre>
    fit <- tree(Bodyfat ~ ., data=newdata, split="deviance")</pre>
    test_error <- mean((predict(fit, test) - test$Bodyfat)^2)</pre>
    test_errors[i] <- test_error</pre>
}
mean(test errors)
set.seed(1234567890)
tree_count <- 100
k <- 3
errors <- rep(0, tree_count * k)
for (i in 1:tree_count) {
    newdata <- data[sample(nrow(data), replace=TRUE),]</pre>
    datasets <- suppressWarnings(split(newdata, 1:k))</pre>
    train1 <- rbind(datasets[[1]], datasets[[2]])</pre>
    test1 <- datasets[[3]]</pre>
    train2 <- rbind(datasets[[1]], datasets[[3]])</pre>
    test2 <- datasets[[2]]</pre>
    train3 <- rbind(datasets[[2]], datasets[[3]])</pre>
    test3 <- datasets[[1]]</pre>
    fit1 <- tree(Bodyfat ~ ., data=train1, split="deviance")</pre>
    error1 <- mean((predict(fit1, test1) - test1$Bodyfat)^2)</pre>
```

```
fit2 <- tree(Bodyfat ~ ., data=train2, split="deviance")</pre>
    error2 <- mean((predict(fit2, test2) - test2$Bodyfat)^2)</pre>
    fit3 <- tree(Bodyfat ~ ., data=train3, split="deviance")</pre>
    error3 <- mean((predict(fit3, test3) - test3$Bodyfat)^2)</pre>
    errors[(i-1)*k+1] \leftarrow error1
    errors[(i-1)*k+2] \leftarrow error2
    errors[(i-1)*k+3] \leftarrow error3
mean(errors)
bagging.regtrees <- function(formula, data, newdata, b) {</pre>
    predictions <- matrix(0, nrow=nrow(newdata), ncol=b)</pre>
    for (i in 1:k) {
         bootstrap_sample <- data[sample(nrow(data), replace=TRUE),]</pre>
         fit <- tree(formula, data=bootstrap_sample, split="deviance")</pre>
        predictions[, i] <- predict(fit, newdata)</pre>
    }
    rowMeans(predictions)
}
cv.regtrees <- function(formula, data, newdata, b, k) {</pre>
    predictions <- matrix(0, nrow(nrow(newdata)), ncol=b*k)</pre>
    for (i in 1:tree_count) {
        bootstrap_sample <- data[sample(nrow(data), replace=TRUE),]</pre>
         datasets <- suppressWarnings(split(bootstrap_sample, 1:k))</pre>
        train1 <- rbind(datasets[[1]], datasets[[2]])</pre>
        test1 <- datasets[[3]]</pre>
        train2 <- rbind(datasets[[1]], datasets[[3]])</pre>
        test2 <- datasets[[2]]</pre>
        train3 <- rbind(datasets[[2]], datasets[[3]])</pre>
        test3 <- datasets[[1]]</pre>
        fit1 <- tree(Bodyfat ~ ., data=train1, split="deviance")</pre>
        prediction1 <- predict(fit1, newdata)</pre>
        fit2 <- tree(Bodyfat ~ ., data=train2, split="deviance")</pre>
        prediction2 <- predict(fit2, newdata)</pre>
        fit3 <- tree(Bodyfat ~ ., data=train3, split="deviance")</pre>
        prediction2 <- predict(fit2, newdata)</pre>
        predictions[, (i - 1) * k + 1] <- predcition1</pre>
        predictions[, (i - 1) * k + 2] <- prediction2</pre>
        predictions[, (i - 1) * k + 3] <- prediction3
    }
```

```
rowMeans(predictions)
}
```

#### Code for Assignment 2b

#### Code for Assignment 3a

```
library(mboost)
data <- read.csv2("../data/bodyfatregression.csv")</pre>
fit <- blackboost(Bodyfat_percent ~ Waist_cm + Weight_kg, data=data)
cvf <- cv(model.weights(fit), type="kfold")</pre>
cvm <- cvrisk(fit, folds=cvf, grid=1:100)</pre>
plot(cvm)
set.seed(1234567890)
train_idx <- sample(nrow(data), floor(nrow(data) * (2 / 3)))</pre>
train <- data[train_idx,]</pre>
test <- data[-train_idx,]</pre>
fit <- blackboost(Bodyfat_percent ~ Waist_cm + Weight_kg, data=train,</pre>
                   control=boost_control(mstop=mstop(cvm)))
test_error <- sum((predict(fit, test) - test$Bodyfat_percent)^2)</pre>
train_error <- sum((predict(fit, train) - train$Bodyfat_percent)^2)</pre>
test_error
train_error
```

#### Code for Assignment 4a

```
library(mboost)
library(randomForest)
library(ggplot2)
library(reshape2)
data <- read.csv2("../data/spambase.csv")</pre>
data$Spam <- as.factor(data$Spam)</pre>
set.seed(1234567890)
train_idx <- sample(nrow(data), floor(nrow(data) * (2 / 3)))</pre>
train <- data[train_idx,]</pre>
test <- data[-train_idx,]</pre>
tree_counts <- seq(10, 100, by=10)
test_errors <- rep(0, length(tree_counts))</pre>
train_errors <- rep(0, length(tree_counts))</pre>
for (i in 1:length(tree_counts)) {
    fit <- blackboost(Spam ~ ., data=train, family=AdaExp(),</pre>
                        control=boost_control(mstop=tree_counts[i]))
```

```
test_error <- 1 - (sum(predict(fit, test, type="class") == test$Spam) / nrow(test))</pre>
    train_error <- 1 - (sum(predict(fit, train, type="class") == train$Spam) / nrow(train))</pre>
    test_errors[i] <- test_error</pre>
    train_errors[i] <- train_error</pre>
plot_data <- data.frame(Trees=tree_counts, test=test_errors, train=train_errors)</pre>
plot_data <- melt(plot_data, id="Trees", value.name="Error", variable.name="Data")</pre>
ggplot(plot_data) +
    xlab("Number of Regression Trees") +
    ylab("Misclassification Rate") +
    geom_line(aes(x=Trees, y=Error, color=Data)) +
    scale_x_discrete(limits=tree_counts)
test_errors <- rep(0, length(tree_counts))</pre>
train_errors <- rep(0, length(tree_counts))</pre>
for (i in 1:length(tree_counts)) {
    fit <- randomForest(Spam ~ ., data=train, ntree=tree_counts[i])</pre>
    test_error <- 1 - (sum(predict(fit, test, type="class") == test$Spam) / nrow(test))</pre>
    train_error <- 1 - (sum(predict(fit, train, type="class") == train$Spam) / nrow(train))</pre>
    test_errors[i] <- test_error</pre>
    train_errors[i] <- train_error</pre>
}
plot_data <- data.frame(Trees=tree_counts, test=test_errors, train=train_errors)</pre>
plot_data <- melt(plot_data, id="Trees", value.name="Error", variable.name="Data")</pre>
ggplot(plot_data) +
    xlab("Number of Regression Trees") +
    ylab("Misclassification Rate") +
    geom_line(aes(x=Trees, y=Error, color=Data)) +
    scale_x_discrete(limits=tree_counts)
```