

Compulsory exercise 2: Group 5

TMA4268 Statistical Learning V2022

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Problem 1

```
set.seed(1)
boston <- scale(Boston, center = T, scale = T)

train.ind = sample(1:nrow(boston), 0.8 * nrow(boston))
boston.train = data.frame(boston[train.ind, ])
boston.test = data.frame(boston[-train.ind, ])
```

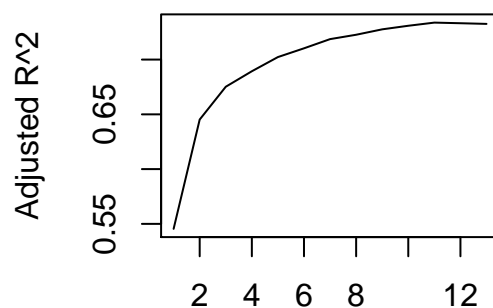
a)

```
set.seed(1)
forward_stepwise = regsubsets(medv ~ ., data = boston.train, nvmax = 13, method = 'forward')
backward_stepwise = regsubsets(medv ~ ., data = boston.train, nvmax = 13, method = 'backward')

forward_stepwise_summary = summary(forward_stepwise)
backward_stepwise_summary = summary(backward_stepwise)
#forward_stepwise_summary
#backward_stepwise_summary

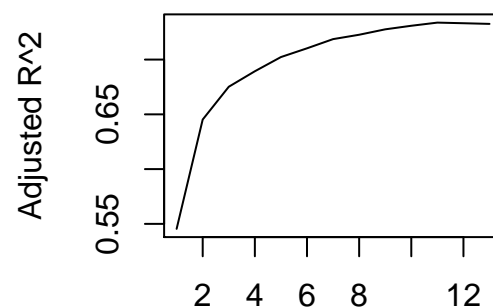
par(mfrow=c(1,2))
plot(forward_stepwise_summary$adjr2, xlab = '# variables', ylab = 'Adjusted R^2', type='l', main='Forwards')
plot(backward_stepwise_summary$adjr2, xlab = '# variables', ylab = 'Adjusted R^2', type='l', main='Backwards')
```

Forwards



variables

Backwards



variables

b)

```
forward_stepwise_summary$outmat
```

```
##          crim zn  indus chas nox rm  age dis rad tax ptratio black lstat
## 1  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 2  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 3  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 4  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 5  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 6  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 7  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 8  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 9  ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 10 ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 11 ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 12 ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
## 13 ( 1 )  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "  " "
```

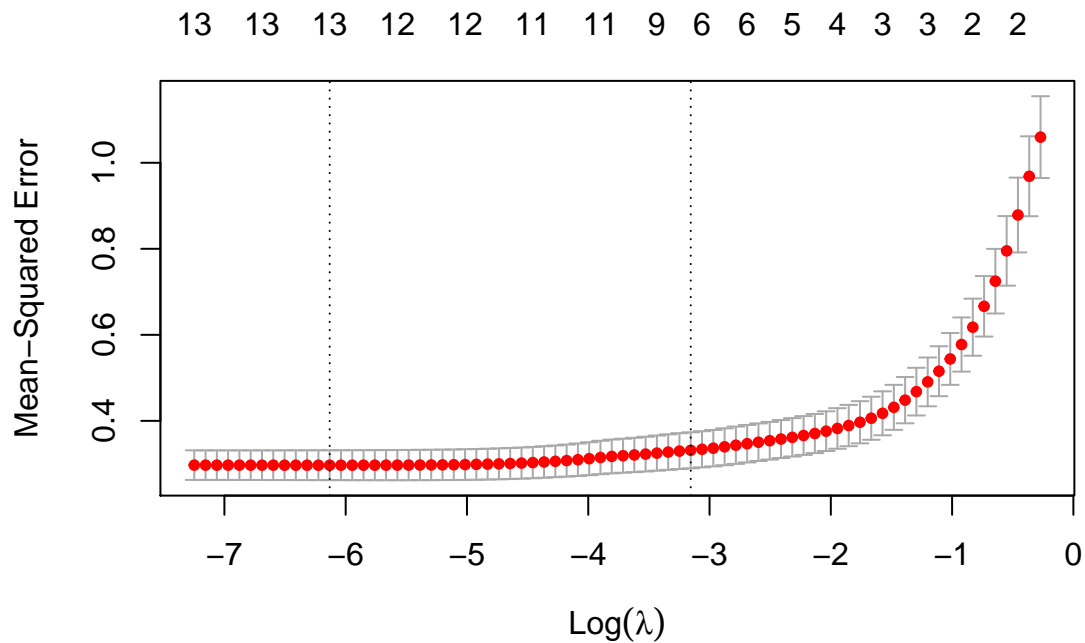
We choose the predictors 'rm', 'dis', 'ptratio' and 'lstat'.

c)

i)

```
set.seed(1)
y = boston.train$medv
x = data.matrix(boston.train[, -14])

cv_lasso = cv.glmnet(x, y, alpha=1, nfolds=5)
plot(cv_lasso)
```



ii)

```
lasso_best_lambda = cv_lasso$lambda.min  
lasso_best_lambda
```

```
## [1] 0.002172032
```

iii)

```
coef(glmnet(x, y, alpha=1, lambda=lasso_best_lambda))
```

```
## 14 x 1 sparse Matrix of class "dgCMatrix"  
##                s0  
## (Intercept)  0.023622904  
## crim        -0.081992849  
## zn           0.094717791  
## indus        0.002619428  
## chas         0.087341100  
## nox          -0.175365927  
## rm           0.312648954  
## age         -0.011212120  
## dis         -0.317143728  
## rad          0.270168177  
## tax         -0.207314714  
## ptratio     -0.204052488  
## black        0.102877803  
## lstat       -0.428298373
```

d)

i) TRUE

ii) FALSE

iii) FALSE

iv) TRUE

Problem 2

a)

b)

Problem 3

a)

i) TRUE

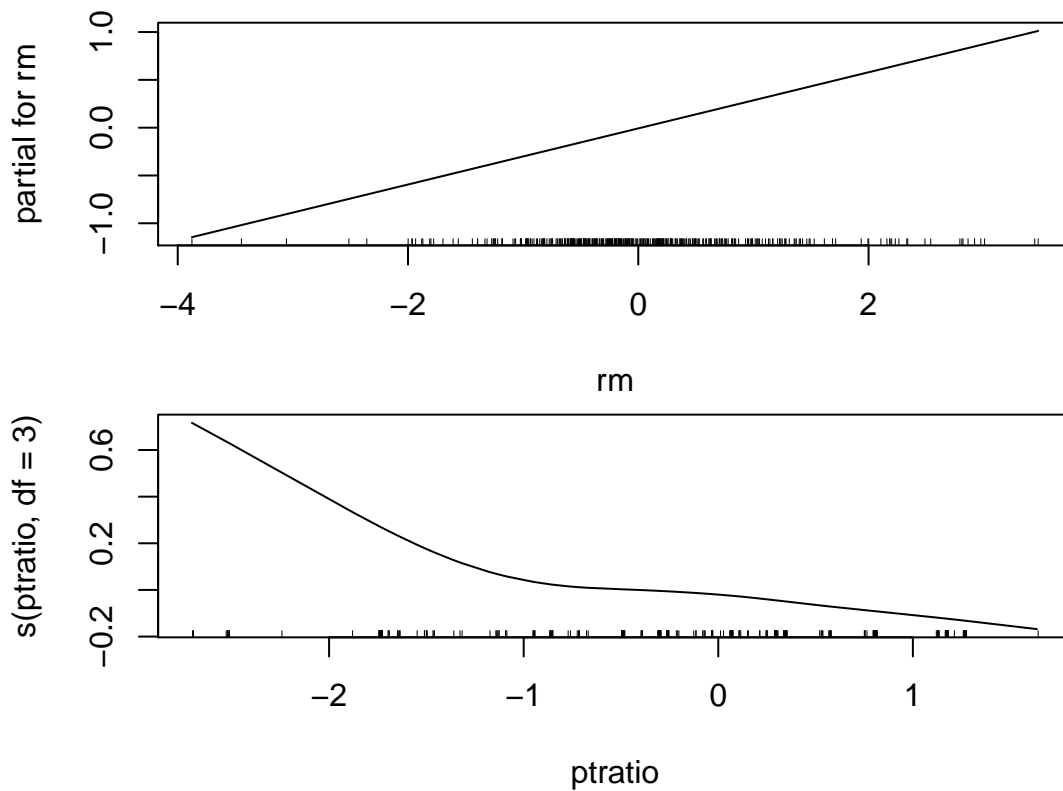
ii) FALSE

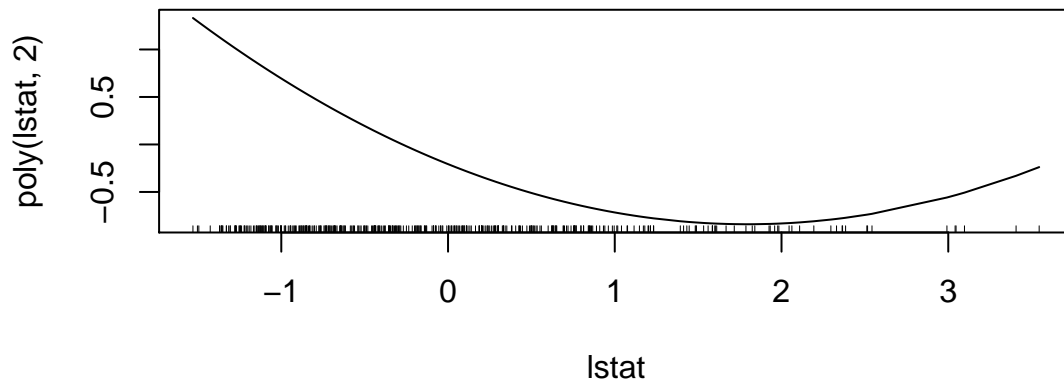
iii) FALSE

iv) TRUE

b)

```
additive_model = gam(medv ~ rm + s(ptratio, df=3) + poly(lstat, 2), data=boston.train)
plot(additive_model)
```





Problem 4

a)

i) FALSE

ii) TRUE

iii) TRUE

iv) TRUE

b)

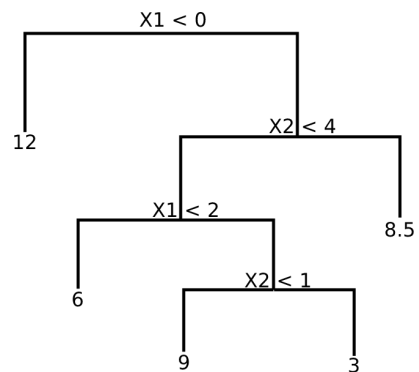


Figure 1: Tree

c)

```

library(tidyverse)
library(palmerpenguins) # Contains the data set "penguins".
data(penguins)

names(penguins) <- c("species", "island", "billL", "billD", "flipperL", "mass", "sex", "year")

Penguins_reduced <- penguins %>% dplyr::mutate(mass = as.numeric(mass), flipperL = as.numeric(flipperL))

# We do not want "year" in the data (this will not help for future predictions)
Penguins_reduced <- Penguins_reduced[,-c(8)]
  
```

```

set.seed(4268)
# 70% of the sample size for training set
training_set_size <- floor(0.7 * nrow(Penguins_reduced))
train_ind <- sample(seq_len(nrow(Penguins_reduced)), size = training_set_size)
train <- Penguins_reduced[train_ind, ]
test <- Penguins_reduced[-train_ind, ]

```

i)

```

penguin.tree = tree(formula=species ~ ., data=train, split='gini' )
summary(penguin.tree)

```

```

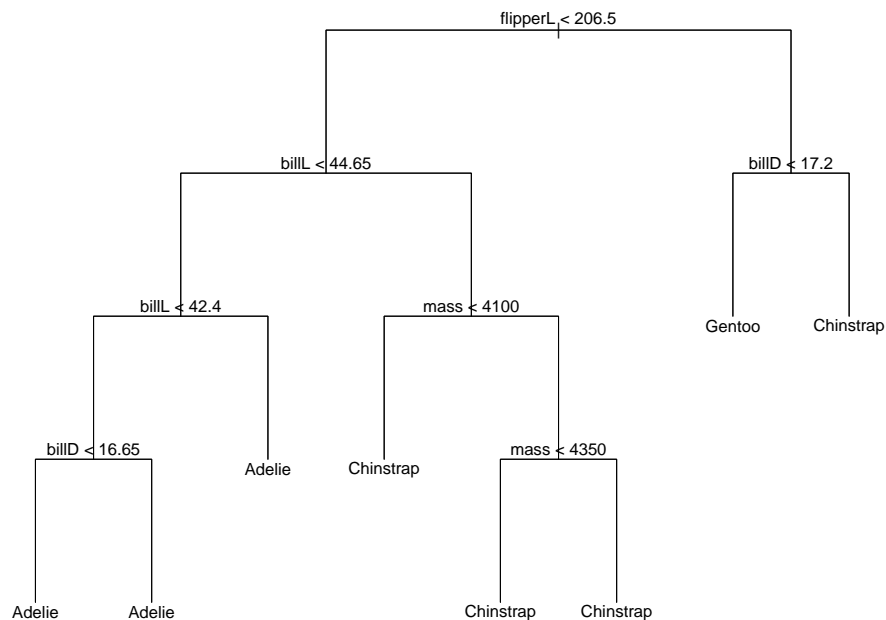
##
## Classification tree:
## tree(formula = species ~ ., data = train, split = "gini")
## Variables actually used in tree construction:
## [1] "flipperL" "billL"    "billD"    "mass"
## Number of terminal nodes: 8
## Residual mean deviance: 0.1869 = 42.06 / 225
## Misclassification error rate: 0.04292 = 10 / 233

```

```

plot(penguin.tree, type='uniform')
text(penguin.tree, pretty=0)

```



ii)

```

set.seed(123)
cv.penguins = cv.tree(penguin.tree, K=10)
cv.penguins$dev

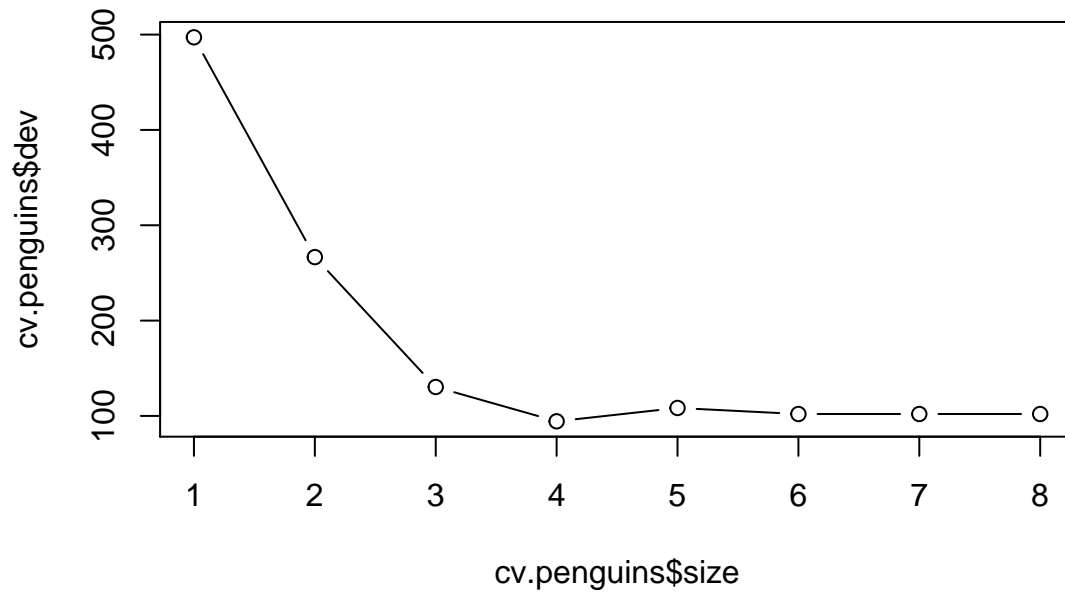
```

```

## [1] 102.00797 102.00797 102.00797 108.34150 94.33622 130.34165 266.60537
## [8] 497.15841

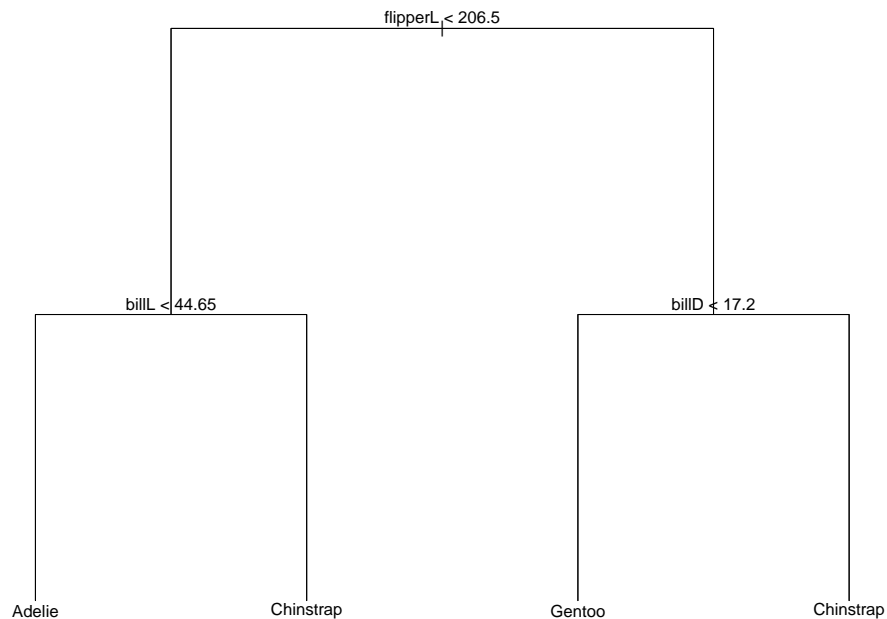
```

```
plot(cv.penguins$dev ~ cv.penguins$size, type='b')
```



iii)

```
prune.penguins = prune.tree(penguin.tree, best=4)
plot(prune.penguins, type='uniform')
text(prune.penguins, pretty=0)
```



```
tree.predict = predict(prune.penguins, test, type='class')
misclass = table(tree.predict, test$species)
misclass
```

```
##
## tree.predict Adelie Chinstrap Gentoo
##    Adelie      42         5         1
```

```
##   Chinstrap      0      15      0
##   Gentoo        0       0     37
```

```
1-sum(diag(misclass))/sum(misclass)
```

```
## [1] 0.06
```