Solution 1:

a) The spam data is a binary classification task where the aim is to classify an email as spam or no-spam.

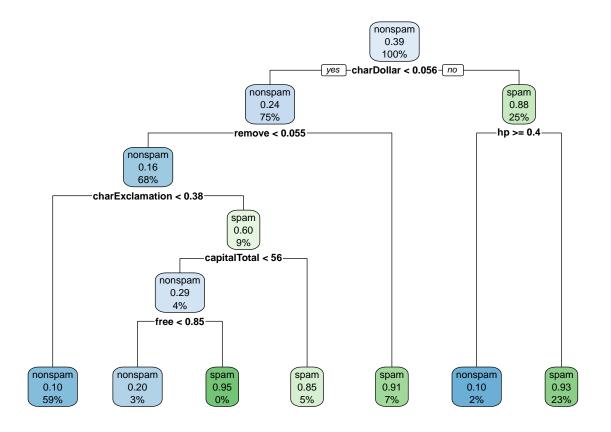
```
library(mlr)
spam.task
## Supervised task: spam-example
## Type: classif
## Target: type
## Observations: 4601
## Features:
## numerics factors ordered functionals ## 57 0 0 0
## Missings: FALSE
## Has weights: FALSE
## Has blocking: FALSE
## Has coordinates: FALSE
## Classes: 2
## nonspam spam
## 2788 1813
## Positive class: nonspam
```

```
b) library(rpart.plot)

## Loading required package: rpart

lrn = makeLearner("classif.rpart")
model = train(lrn, spam.task)
mod = getLearnerModel(model)
rpart.plot(mod)

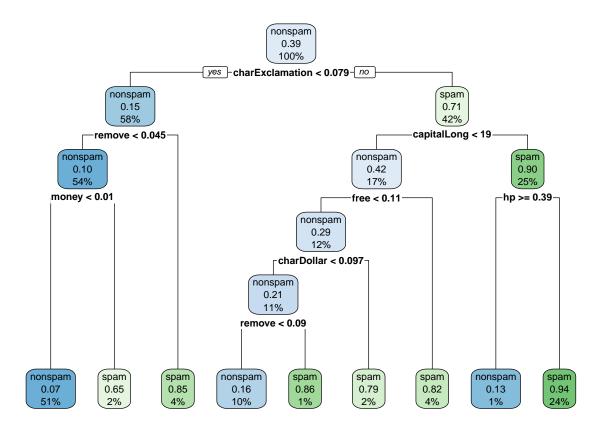
## Warning: Cannot retrieve the data used to build the model (so cannot determine roundint and is.binary for the variables).
## To silence this warning:
## Call rpart.plot with roundint=FALSE,
## or rebuild the rpart model with model=TRUE.
```



```
set.seed(42)
subset1 = sample.int(getTaskSize(spam.task), size = 0.8 * getTaskSize(spam.task))
subset2 = sample.int(getTaskSize(spam.task), size = 0.8 * getTaskSize(spam.task))

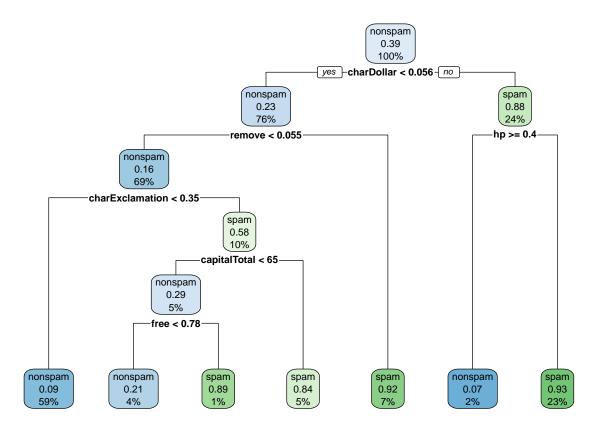
model = train(lrn, spam.task, subset = subset1)
mod = getLearnerModel(model)
rpart.plot(mod)

## Warning: Cannot retrieve the data used to build the model (so cannot determine roundint and is.binary for the variables).
## To silence this warning:
## Call rpart.plot with roundint=FALSE,
## or rebuild the rpart model with model=TRUE.
```



```
model = train(lrn, spam.task, subset = subset2)
mod = getLearnerModel(model)
rpart.plot(mod)

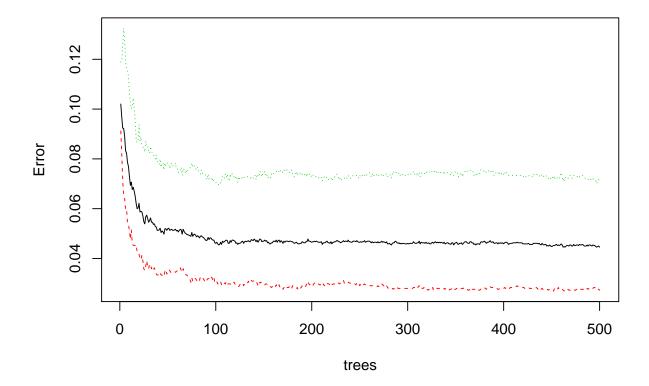
## Warning: Cannot retrieve the data used to build the model (so cannot determine
roundint and is.binary for the variables).
## To silence this warning:
## Call rpart.plot with roundint=FALSE,
## or rebuild the rpart model with model=TRUE.
```



Observation: Trees with different sample find different split points and variables, leading to different trees!

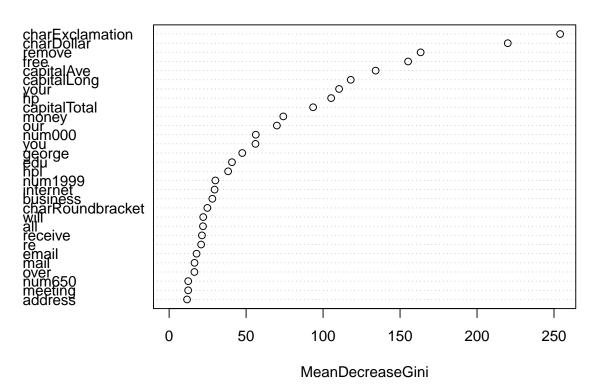
```
c) lrn = makeLearner("classif.randomForest")
  model = train(lrn, spam.task)
  mod = getLearnerModel(model)
  ##
  ## Call:
  ## randomForest(formula = f, data = data, classwt = classwt, cutoff = cutoff)
  ##
                     Type of random forest: classification
                           Number of trees: 500
  ##
  ## No. of variables tried at each split: 7
  ##
  ##
             OOB estimate of error rate: 4.46%
  ## Confusion matrix:
  ##
             nonspam spam class.error
  ## nonspam
                 2712
                        76
                               0.02726
  ## spam
                  129 1684
                               0.07115
  plot(mod)
```

mod



```
d) imp = getFeatureImportance(model)
  sort(imp$res, decreasing = TRUE)
  ##
       charExclamation charDollar remove free capitalAve capitalLong your
  ## 1
                 254
                             220 163.4 155.2
                                                  134.1
                                                                118 110.4
  ##
          hp capitalTotal money our num000 you george edu hpl num1999
  ## 1 105.2
               93.52 74.15 69.98 56.3 56.09 47.5 40.82 38.31
       internet business charRoundbracket will all receive re email mail
                 28.06
                                  24.82 22.13 21.98 21.31 20.77 17.82 16.49
          29.51
        over num650 meeting address charSemicolon order credit make labs
  ## 1 16.41 12.38
                    12.36 11.63
                                         10.85 9.045 8.255 8.01 7.917
       \verb|char| Hash | \verb|num85| | \verb|people| | technology | m | data | char Square bracket | font |
         7.689 7.663 7.415
                              6.834 6.589 5.838
                                                              5.443 5.09
       project report lab telnet original addresses conference direct
  ## 1 4.693 4.506 4.175 3.96
                                   3.63
                                               2.94
                                                         2.724 2.449 2.025
  ## num415 num3d num857 parts table
  ## 1 1.689 1.686 1.625 0.9221 0.5594
  # as alternative, the randomForest package provides a plotting function
  randomForest::varImpPlot(getLearnerModel(model))
```

getLearnerModel(model)



Solution 2: See R code randomForest_l_2.R

Solution 3:

- a) Proceed as follows:
 - (i) Split x in two groups using the following split points.
 - (1), (2, 7, 10, 20) (splitpoint 1.5)
 - (1,2), (7,10,20) (splitpoint 4.5)
 - (1,2,7), (10,20) (splitpoint 8.5)
 - (1, 2, 7, 10), (20) (splitpoint 15)
 - (ii) For each possible split point compute the sum of squares in both groups.
 - (iii) Use as split point the point that splits both groups best w.r.t. minimizing the sum of squares in both groups.

Here, we have only one split variable x. A split point t, leads to the following half-spaces:

$$\mathcal{N}_1(t) = \{(x, y) \in \mathcal{N} : x \le t\} \text{ and } \mathcal{N}_2(t) = \{(x, y) \in \mathcal{N} : x > t\}.$$

Remember the minimization Problem (here only for one split variable x):

$$\min_{t} \left(\min_{c_1} \sum_{(x,y) \in \mathcal{N}_1} (y - c_1)^2 + \min_{c_2} \sum_{(x,y) \in \mathcal{N}_2} (y - c_2)^2 \right).$$

The inner minimization is solved through: $\hat{c}_1 = \bar{y}_1$ and $\hat{c}_2 = \bar{y}_2$ Which results in:

$$\min_{t} \left(\sum_{(x,y)\in\mathcal{N}_{1}} (y - \bar{y}_{1})^{2} + \sum_{(x,y)\in\mathcal{N}_{2}} (y - \bar{y}_{2})^{2} \right).$$

The sum of squares error of the parent is:

$$Impurity_{parent} = MSE_{parent} = \frac{1}{5} \sum_{i=1}^{5} (y_i - 4.7)^2 = 22.56$$

Calculate the risk for each split point:

 $x \le 1.5$

$$\mathcal{R}(1, 1.5) = \frac{1}{5} \text{MSE}_{left} + \frac{4}{5} \text{MSE}_{right} =$$

$$= \frac{1}{5} \cdot \frac{1}{1} (1 - 1)^2 + \frac{4}{5} \cdot \frac{1}{4} ((1 - 5.625)^2 + (0.5 - 5.625)^2 + (10 - 5.625)^2 + (11 - 5.625)^2)$$

$$= 19.1375$$

```
x \le 4.5 \ \mathcal{R}(1, 4.5) = 13.43

x \le 8.5 \ \mathcal{R}(1, 8.5) = 0.13

x \le 15 \ \mathcal{R}(1, 15) = 12.64
```

Minimal empirical risk is obtained by choosing the split point 8.5.

Doing the same for the log-transformation gives:

```
x \le 0.3 \ \mathcal{R}(1, 0.3) = 19.14

x \le 1.3 \ \mathcal{R}(1, 1.3) = 13.43

x \le 2.1 \ \mathcal{R}(1, 2.1) = 0.13

x \le 2.6 \ \mathcal{R}(1, 2.6) = 12.64
```

Minimal empirical risk is obtained by choosing the split point 2.1.

```
b) x = c(1,2,7,10,20)
y = c(1,1,0.5,10,11)

calculateMSE = function (y) mean((y - mean(y))^2)
calculateTotalMSE = function (yleft, yright) {
    n_left = length(yleft)
    n_right = length(yright)

    w_mse_left = n_left / (n_left + n_right) * calculateMSE(yleft)
    w_mse_right = n_right / (n_left + n_right) * calculateMSE(yright)

    return(w_mse_left + w_mse_right)
}

split = function(x, y) {
```

```
# try out all points as potential split points and ...
  split_points = 0.5 * diff(sort(x)) + x[-length(x)]
  node_mses = lapply(split_points, function(i) {
   y_{\text{left}} = y[x \le i]
   y_right = y[x > i]
    # ... compute SS in both groups
    mse_split = calculateTotalMSE(y_left, y_right)
    print(sprintf("Split at %.1f: empirical Risk = %.2f", i, mse_split))
    return(mse_split)
  })
  # select the split point yielding the maximum impurity reduction
  best = which.min(node_mses)
  split_points[best]
X
## [1] 1 2 7 10 20
split(x, y) # the 3rd observation is the best split point
## [1] "Split at 1.5: empirical Risk = 19.14"
## [1] "Split at 4.5: empirical Risk = 13.43"
## [1] "Split at 8.5: empirical Risk = 0.13"
## [1] "Split at 15.0: empirical Risk = 12.64"
## [1] 8.5
log(x)
## [1] 0.0000 0.6931 1.9459 2.3026 2.9957
split(log(x), y) # also here, the 3rd observation is the best split point
## [1] "Split at 0.3: empirical Risk = 19.14"
## [1] "Split at 1.3: empirical Risk = 13.43"
## [1] "Split at 2.1: empirical Risk = 0.13"
## [1] "Split at 2.6: empirical Risk = 12.64"
## [1] 2.124
```

Solution 4:

The fractions of the classes k = 1, ..., g in node \mathcal{N} of a decision tree are $p(1|\mathcal{N}), ..., p(g|\mathcal{N})$. Assume we replace the classification rule in node t

$$\hat{k}|\mathcal{N} = \arg\max_{k} p(k|\mathcal{N})$$

with a randomizing rule, in which we draw the classes in one node from their estimated probabilities. Derive an estimator for the misclassification rate in node \mathcal{N} . What do you (hopefully) recognize?

Solution:

• For the target space $\mathcal{Y} = \{1, \dots, g\}$ we assume the distribution P_Y . The distribution $P_{\hat{Y}}$ of our classifier $\hat{Y} = h(x)$ is defined as the individual class frequencies in a node \mathcal{N} (estimated probability, that an object of class k is in node \mathcal{N}):

$$P(\hat{Y} = k \mid \mathcal{N}) = p(k|\mathcal{N})$$

• As estimate of the target distribution P_Y we use the distribution of the classifier h(x) which we assume to be independent of $P_{\hat{\mathbf{v}}}$:

$$P_Y \stackrel{\text{ind.}}{\sim} P_{\hat{V}}$$

• The individual error rate of wrongly predicting a true label k in node \mathcal{N} ($\mathsf{err}_{k|\mathcal{N}}$) can be written as probability that Y = k and $\hat{Y} \neq k$:

$$\begin{split} P(\mathsf{err}_{k|\mathcal{N}}) &= P(Y = k, \hat{Y} \neq k \mid \mathcal{N}) \\ &= P(Y = k \mid \mathcal{N}) P(\hat{Y} \neq k \mid \mathcal{N}) \\ &= p(k|\mathcal{N}) (1 - p(k|\mathcal{N})) \end{split}$$

• The error rate is the combination of all individual error rates:

$$\mathsf{err}_{\mathcal{N}} = igcup_{k=1}^g \mathsf{err}_{k|\mathcal{N}}$$

ullet Finally, we are interested in the probability of the error rate $err_{\mathcal{N}}$ as estimator for the missclassification rate:

$$\begin{split} P(\mathsf{err}_{\mathcal{N}}) &= P\left(\bigcup_{k=1}^g \mathsf{err}_{k|\mathcal{N}}\right) \\ &= \sum_{k=1}^g P\left(\mathsf{err}_{k|\mathcal{N}}\right) \\ &= \sum_{k=1}^g p(k|\mathcal{N})(1-p(k|\mathcal{N})) \\ &= \sum_{k=1}^g p(k|\mathcal{N}) - \sum_{k=1}^g p(k|\mathcal{N})^2 \\ &= 1 - \sum_{k=1}^g p(k|\mathcal{N})^2 \end{split}$$

This is exactly the Gini-Index that CART uses for splitting the tree.

Solution 5:

 $f(x) = \frac{1}{B} \sum_{b=1}^{B} f_b(x)$ is the bagging estimator based on B bootstrap samples. Then we can easily calculate:

$$\operatorname{Var}(f(x)) = \frac{1}{B^2} \left(\sum_{b=1}^B \operatorname{Var}(f_b(x)) + \sum_{i \neq j}^B \operatorname{Cov}(f_i(x), f_j(x)) \right)$$
$$= \frac{1}{B^2} \left(B\sigma^2 + (B^2 - B)\rho\sigma^2 \right)$$
$$= \frac{1}{B}\sigma^2 + \rho\sigma^2 - \frac{1}{B}\rho\sigma^2$$
$$= \rho\sigma^2 + \frac{\sigma^2}{B}(1 - \rho)$$

In the first line the rules for variance of a non-independent sum of random variables is used. All other steps are trivial.