Descision Trees

YOUR NAME

This is from James, Witten, Hastie, and Ibshirani (2021) section 8.3 beginning on page 353.

The tree library is used to construct classification and regression trees.

```
library(tree)
```

```
## Warning: package 'tree' was built under R version 4.4.1
```

We first use classification trees to analyze the Carseats data set. In these data, Sales is a continuous variable, and so we begin by recoding it as a binary variable. We use the ifelse() function to create a variable, called High, which takes on a value of Yes if the Sales variable exceeds 8, and takes on a value of No otherwise.

```
library(ISLR2)
attach(Carseats)
High <- factor(ifelse(Sales <= 8, "No", "Yes"))</pre>
```

Finally, we use the data.frame() function to merge High with the rest of the Carseats data.

```
Carseats <- data.frame(Carseats, High)</pre>
```

We now use the tree() function to fit a classification tree in order to predict High using all variables but Sales. The syntax of the tree() function is quite similar to that of the lm() function.

```
tree.carseats <- tree(High ~ . - Sales, Carseats)</pre>
```

The summary() function lists the variables that are used as internal nodes in the tree, the number of terminal nodes, and the (training) error rate.

```
summary(tree.carseats)
```

```
##
## Classification tree:
## tree(formula = High ~ . - Sales, data = Carseats)
## Variables actually used in tree construction:
## [1] "ShelveLoc" "Price" "Income" "CompPrice" "Population"
## [6] "Advertising" "Age" "US"
## Number of terminal nodes: 27
## Residual mean deviance: 0.4575 = 170.7 / 373
## Misclassification error rate: 0.09 = 36 / 400
```

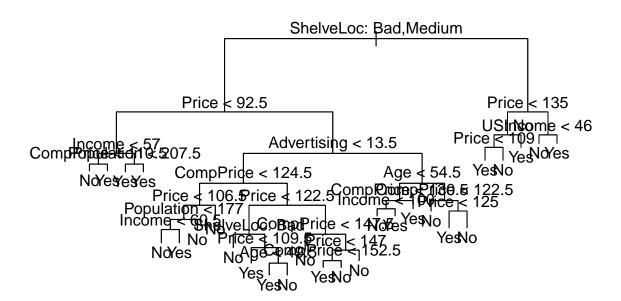
We see that the training error rate is 9%. For classification trees, the deviance reported in the output of summary() is given by

$$-2\sum_{m}\sum_{k}n_{mk}\log\hat{p}_{mk},$$

where n_{mk} is the number of observations in the *m*th terminal node that belong to the *k*th class. This is closely related to the entropy, defined in (8.7). A small deviance indicates a tree that provides a good fit to the (training) data. The *residual mean deviance* reported is simply the deviance divided by $n - |T_0|$, which in this case is 400 - 27 = 373.

One of the most attractive properties of trees is that they can be graphically displayed. We use the plot() function to display the tree structure, and the text() function to display the node labels. The argument pretty = 0 instructs R to include the category names for any qualitative predictors, rather than simply displaying a letter for each category.

```
plot(tree.carseats)
text(tree.carseats, pretty = 0)
```



The most important indicator of Sales appears to be shelving location, since the first branch differentiates Good locations from Bad and Medium locations.

If we just type the name of the tree object, R prints output corresponding to each branch of the tree. R displays the split criterion (e.g. Price < 92.5), the number of observations in that branch, the deviance, the overall prediction for the branch (Yes or No), and the fraction of observations in that branch that take on values of Yes and No. Branches that lead to terminal nodes are indicated using asterisks.

```
## node), split, n, deviance, yval, (yprob)
         * denotes terminal node
##
     1) root 400 541.500 No ( 0.59000 0.41000 )
##
       2) ShelveLoc: Bad, Medium 315 390.600 No (0.68889 0.31111)
##
         4) Price < 92.5 46 56.530 Yes ( 0.30435 0.69565 )
##
##
          8) Income < 57 10 12.220 No ( 0.70000 0.30000 )
##
            16) CompPrice < 110.5 5
                                    0.000 No ( 1.00000 0.00000 ) *
                                     6.730 Yes ( 0.40000 0.60000 ) *
##
            17) CompPrice > 110.5 5
##
           9) Income > 57 36 35.470 Yes (0.19444 0.80556)
##
            18) Population < 207.5 16 21.170 Yes (0.37500 0.62500) *
##
            19) Population > 207.5 20
                                       7.941 Yes ( 0.05000 0.95000 ) *
##
         5) Price > 92.5 269 299.800 No ( 0.75465 0.24535 )
##
          10) Advertising < 13.5 224 213.200 No ( 0.81696 0.18304 )
##
            20) CompPrice < 124.5 96 44.890 No ( 0.93750 0.06250 )
##
              40) Price < 106.5 38 33.150 No ( 0.84211 0.15789 )
##
                80) Population < 177 12 16.300 No (0.58333 0.41667)
##
                 160) Income < 60.5 6 0.000 No (1.00000 0.00000) *
                 161) Income > 60.5 6 5.407 Yes ( 0.16667 0.83333 ) *
##
##
                81) Population > 177 26 8.477 No ( 0.96154 0.03846 ) *
##
              41) Price > 106.5 58
                                   0.000 No ( 1.00000 0.00000 ) *
            21) CompPrice > 124.5 128 150.200 No ( 0.72656 0.27344 )
##
##
              42) Price < 122.5 51 70.680 Yes ( 0.49020 0.50980 )
##
                                       6.702 No ( 0.90909 0.09091 ) *
                84) ShelveLoc: Bad 11
##
                85) ShelveLoc: Medium 40 52.930 Yes (0.37500 0.62500)
                                       7.481 Yes ( 0.06250 0.93750 ) *
##
                 170) Price < 109.5 16
##
                 171) Price > 109.5 24 32.600 No ( 0.58333 0.41667 )
##
                   342) Age < 49.5 13 16.050 Yes ( 0.30769 0.69231 ) *
##
                   343) Age > 49.5 11
                                       6.702 No ( 0.90909 0.09091 ) *
##
              43) Price > 122.5 77 55.540 No ( 0.88312 0.11688 )
##
                86) CompPrice < 147.5 58 17.400 No ( 0.96552 0.03448 ) *
##
                87) CompPrice > 147.5 19 25.010 No ( 0.63158 0.36842 )
##
                 174) Price < 147 12 16.300 Yes ( 0.41667 0.58333 )
##
                   348) CompPrice < 152.5 7
                                             5.742 Yes ( 0.14286 0.85714 ) *
##
                                              5.004 No ( 0.80000 0.20000 ) *
                   349) CompPrice > 152.5 5
##
                 175) Price > 147 7
                                    0.000 No (1.00000 0.00000) *
##
          11) Advertising > 13.5 45 61.830 Yes ( 0.44444 0.55556 )
##
            22) Age < 54.5 25 25.020 Yes ( 0.20000 0.80000 )
##
              44) CompPrice < 130.5 14 18.250 Yes ( 0.35714 0.64286 )
##
                88) Income < 100 9 12.370 No ( 0.55556 0.44444 ) *
##
                                   0.000 Yes ( 0.00000 1.00000 ) *
                89) Income > 100 5
##
              45) CompPrice > 130.5 11
                                       0.000 Yes ( 0.00000 1.00000 ) *
##
            23) Age > 54.5 20 22.490 No ( 0.75000 0.25000 )
##
                                        0.000 No ( 1.00000 0.00000 ) *
              46) CompPrice < 122.5 10
##
              47) CompPrice > 122.5 10 13.860 No ( 0.50000 0.50000 )
##
                                    0.000 Yes ( 0.00000 1.00000 ) *
                94) Price < 125 5
##
                95) Price > 125 5
                                    0.000 No ( 1.00000 0.00000 ) *
##
       3) ShelveLoc: Good 85 90.330 Yes ( 0.22353 0.77647 )
##
         6) Price < 135 68 49.260 Yes (0.11765 0.88235)
##
          12) US: No 17 22.070 Yes ( 0.35294 0.64706 )
##
            24) Price < 109 8  0.000 Yes ( 0.00000 1.00000 ) *
            25) Price > 109 9 11.460 No ( 0.66667 0.33333 ) *
##
```

```
## 13) US: Yes 51 16.880 Yes ( 0.03922 0.96078 ) *
## 7) Price > 135 17 22.070 No ( 0.64706 0.35294 )
## 14) Income < 46 6 0.000 No ( 1.00000 0.00000 ) *
## 15) Income > 46 11 15.160 Yes ( 0.45455 0.54545 ) *
```

In order to properly evaluate the performance of a classification tree on these data, we must estimate the test error rather than simply computing the training error. We split the observations into a training set and a test set, build the tree using the training set, and evaluate its performance on the test data. The predict() function can be used for this purpose. In the case of a classification tree, the argument type = "class" instructs R to return the actual class prediction. This approach leads to correct predictions for around 77% of the locations in the test data set.

```
set.seed(2)
train <- sample(1:nrow(Carseats), 200)</pre>
Carseats.test <- Carseats[-train, ]</pre>
High.test <- High[-train]</pre>
tree.carseats <- tree(High ~ . - Sales, Carseats,</pre>
    subset = train)
tree.pred <- predict(tree.carseats, Carseats.test,</pre>
    type = "class")
table(tree.pred, High.test)
##
             High.test
## tree.pred
              No Yes
##
          No
              104
                    33
##
          Yes
              13
                    50
(104 + 50) / 200
```

```
## [1] 0.77
```

(If you re-run the predict() function then you might get slightly different results, due to "ties": for instance, this can happen when the training observations corresponding to a terminal node are evenly split between Yes and No response values.)

Next, we consider whether pruning the tree might lead to improved results. The function cv.tree() performs cross-validation in order to determine the optimal level of tree complexity; cost complexity pruning is used in order to select a sequence of trees for consideration. We use the argument FUN = prune.misclass in order to indicate that we want the classification error rate to guide the cross-validation and pruning process, rather than the default for the cv.tree() function, which is deviance. The cv.tree() function reports the number of terminal nodes of each tree considered (size) as well as the corresponding error rate and the value of the cost-complexity parameter used (k, which corresponds to α in (8.4)).

```
set.seed(7)
cv.carseats <- cv.tree(tree.carseats, FUN = prune.misclass)
names(cv.carseats)

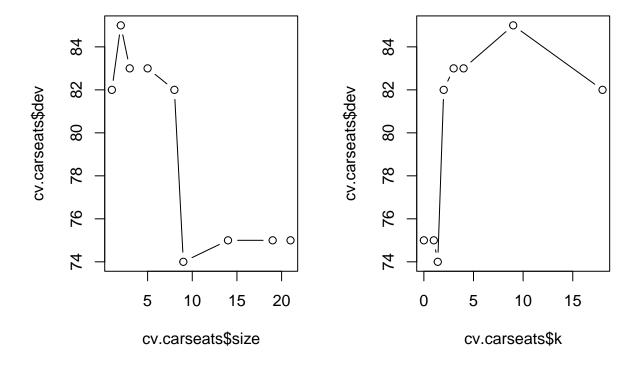
## [1] "size" "dev" "k" "method"

cv.carseats</pre>
```

```
## $size
##
  [1] 21 19 14 9
                  8 5
                         3
                            2 1
##
## $dev
   [1] 75 75 75 74 82 83 83 85 82
##
##
## $k
            0.0 1.0 1.4 2.0 3.0 4.0 9.0 18.0
## [1] -Inf
##
## $method
  [1] "misclass"
##
## attr(,"class")
  [1] "prune"
                       "tree.sequence"
```

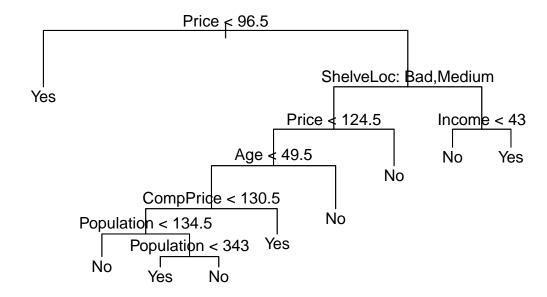
Despite its name, dev corresponds to the number of cross-validation errors. The tree with 9 terminal nodes results in only 74 cross-validation errors. We plot the error rate as a function of both size and k.

```
par(mfrow = c(1, 2))
plot(cv.carseats$size, cv.carseats$dev, type = "b")
plot(cv.carseats$k, cv.carseats$dev, type = "b")
```



We now apply the prune.misclass() function in order to prune the tree to obtain the nine-node tree.

```
prune.carseats <- prune.misclass(tree.carseats, best = 9)
plot(prune.carseats)
text(prune.carseats, pretty = 0)</pre>
```



How well does this pruned tree perform on the test data set? Once again, we apply the predict() function.

```
tree.pred <- predict(prune.carseats, Carseats.test,
    type = "class")
table(tree.pred, High.test)

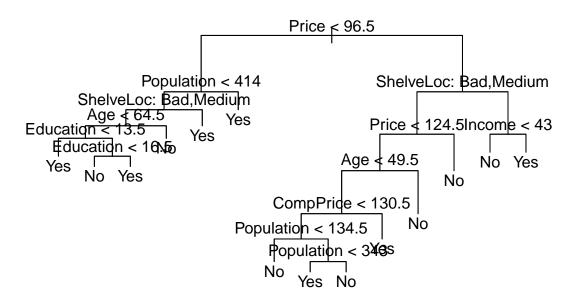
## High.test
## tree.pred No Yes
## No 97 25
## Yes 20 58</pre>
(97 + 58) / 200
```

[1] 0.775

Now 77.5% of the test observations are correctly classified, so not only has the pruning process produced a more interpretable tree, but it has also slightly improved the classification accuracy.

If we increase the value of best, we obtain a larger pruned tree with lower classification accuracy:

```
prune.carseats <- prune.misclass(tree.carseats, best = 14)
plot(prune.carseats)
text(prune.carseats, pretty = 0)</pre>
```



```
tree.pred <- predict(prune.carseats, Carseats.test,
    type = "class")
table(tree.pred, High.test)

## High.test
## tree.pred No Yes
## No 102 31
## Yes 15 52</pre>
(102 + 52) / 200
```

[1] 0.77

Fitting Regression Trees

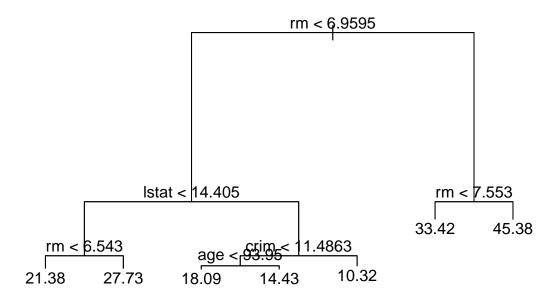
Here we fit a regression tree to the Boston data set. First, we create a training set, and fit the tree to the training data.

```
set.seed(1)
train <- sample(1:nrow(Boston), nrow(Boston) / 2)
tree.boston <- tree(medv ~ ., Boston, subset = train)
summary(tree.boston)</pre>
```

```
##
## Regression tree:
## tree(formula = medv ~ ., data = Boston, subset = train)
## Variables actually used in tree construction:
## [1] "rm"
               "lstat" "crim"
## Number of terminal nodes: 7
## Residual mean deviance: 10.38 = 2555 / 246
## Distribution of residuals:
##
             1st Qu.
       Min.
                       Median
                                        3rd Qu.
                                                     Max.
                                  Mean
## -10.1800 -1.7770 -0.1775
                                0.0000
                                         1.9230
                                                  16.5800
```

Notice that the output of summary() indicates that only four of the variables have been used in constructing the tree. In the context of a regression tree, the deviance is simply the sum of squared errors for the tree. We now plot the tree.

```
plot(tree.boston)
text(tree.boston, pretty = 0)
```



The variable lstat measures the percentage of individuals with {lower socioeconomic status}, while the variable rm corresponds to the average number of rooms. The tree indicates that larger values of rm, or lower

values of lstat, correspond to more expensive houses. For example, the tree predicts a median house price of \$45,400 for homes in census tracts in which rm >= 7.553.

It is worth noting that we could have fit a much bigger tree, by passing control = tree.control(nobs = length(train), mindev = 0) into the tree() function.

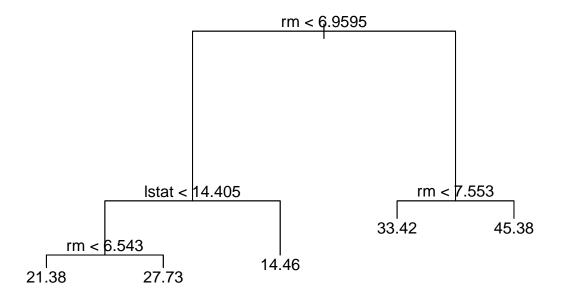
Now we use the cv.tree() function to see whether pruning the tree will improve performance.

```
cv.boston <- cv.tree(tree.boston)
plot(cv.boston$size, cv.boston$dev, type = "b")</pre>
```



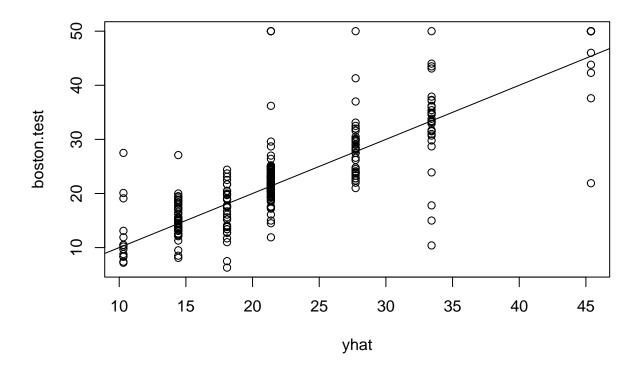
In this case, the most complex tree under consideration is selected by cross-validation. However, if we wish to prune the tree, we could do so as follows, using the prune.tree() function:

```
prune.boston <- prune.tree(tree.boston, best = 5)
plot(prune.boston)
text(prune.boston, pretty = 0)</pre>
```



In keeping with the cross-validation results, we use the unpruned tree to make predictions on the test set.

```
yhat <- predict(tree.boston, newdata = Boston[-train, ])
boston.test <- Boston[-train, "medv"]
plot(yhat, boston.test)
abline(0, 1)</pre>
```



mean((yhat - boston.test)^2)

[1] 35.28688

In other words, the test set MSE associated with the regression tree is 35.29. The square root of the MSE is therefore around 5.941, indicating that this model leads to test predictions that are (on average) within approximately \$5,941 of the true median home value for the census tract.

On Your Own

1. This problem involves the OJ data set which is part of the ISLR2 package.

```
data(OJ, package = 'ISLR2')
```

- (a) Create a training set containing a random sample of 800 observations, and a test set containing the remaining observations.
- (b) Fit a tree to the training data, with Purchase as the response and the other variables as predictors. Use the summary() function to produce summary statistics about the tree, and describe the results obtained. What is the training error rate? How many terminal nodes does the tree have?
- (c) Type in the name of the tree object in order to get a detailed text output. Pick one of the terminal nodes, and interpret the information displayed.
- (d) Create a plot of the tree, and interpret the results.
- (e) Predict the response on the test data, and produce a confusion matrix comparing the test labels to the predicted test labels. What is the test error rate?
- (f) Apply the cv.tree() function to the training set in order to determine the optimal tree size.
- (g) Produce a plot with tree size on the x-axis and cross-validated classification error rate on the y-axis.
- (h) Which tree size corresponds to the lowest cross-validated classification error rate?
- (i) Produce a pruned tree corresponding to the optimal tree size obtained using cross-validation. If cross-validation does not lead to selection of a pruned tree, then create a pruned tree with five terminal nodes.
- (j) Compare the training error rates between the pruned and unpruned trees. Which is higher?
- (k) Compare the test error rates between the pruned and unpruned trees. Which is higher?