

Annand Module 07 Lab 01

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```
library(ISLR2)
library(MASS)
```

```
##
## Attaching package: 'MASS'

## The following object is masked from 'package:ISLR2':
##
## Boston
```

```
library(tree)
```

```
## Warning: package 'tree' was built under R version 4.3.2
```

Question 8

Part A

```
# Split data into training and test data
set.seed(1)
train <- sample(1:nrow(Carseats), nrow(Carseats) / 2)
sales.test <- Carseats[-train, "Sales"]
```

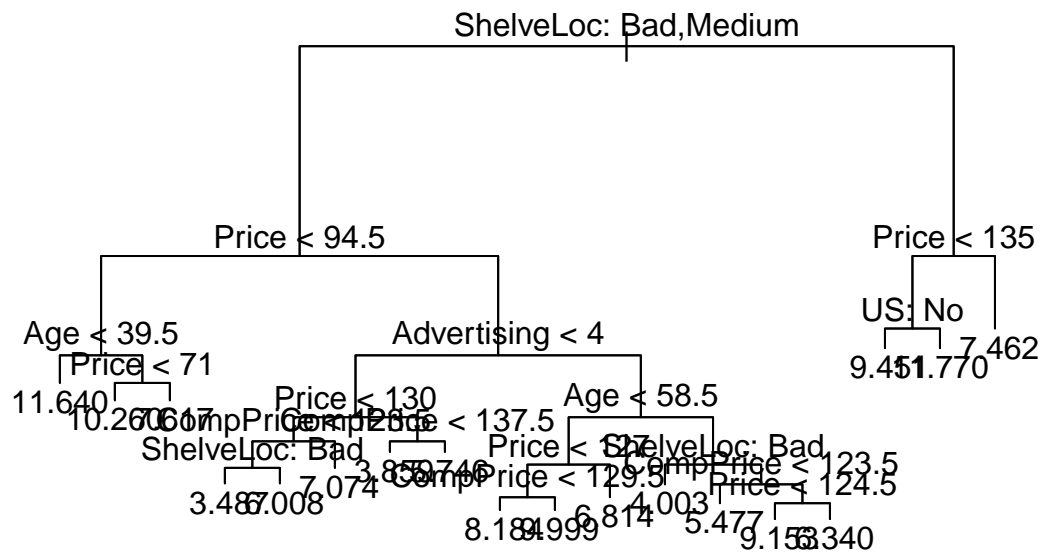
Part B

```
tree.sales <- tree(Sales ~ ., data = Carseats, subset = train)
summary(tree.sales)
```

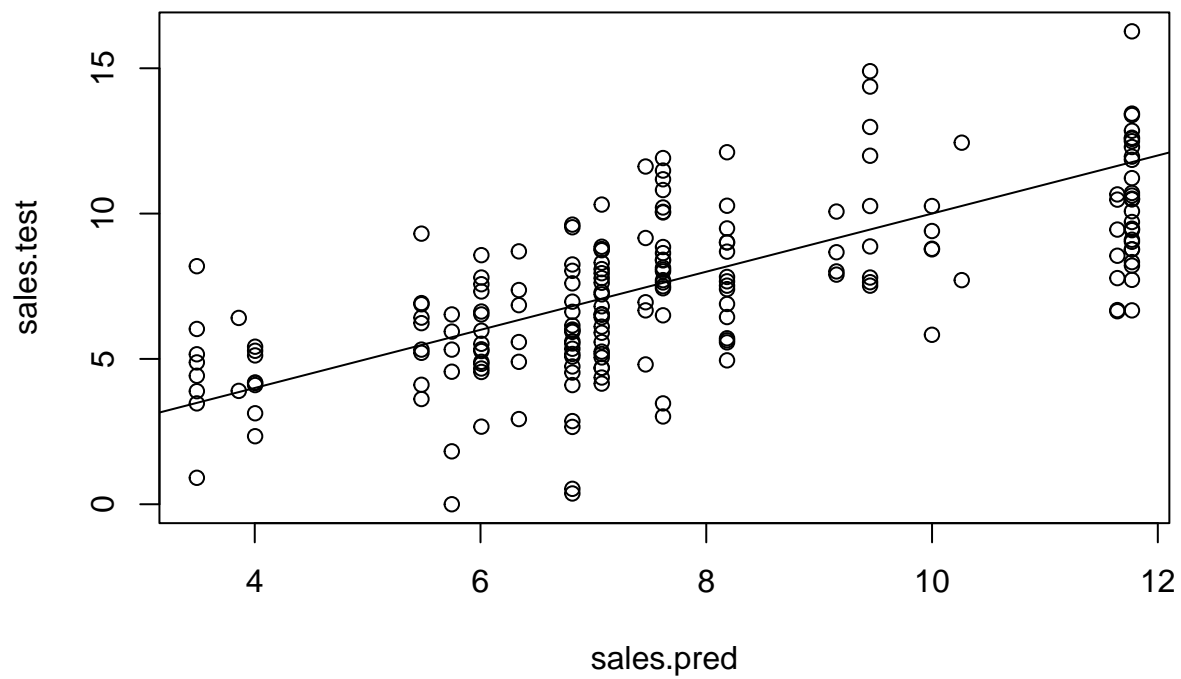
```
##
## Regression tree:
## tree(formula = Sales ~ ., data = Carseats, subset = train)
## Variables actually used in tree construction:
## [1] "ShelveLoc" "Price" "Age" "Advertising" "CompPrice"
## [6] "US"
## Number of terminal nodes: 18
## Residual mean deviance: 2.167 = 394.3 / 182
```

```
## Distribution of residuals:
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## -3.88200 -0.88200 -0.08712  0.00000  0.89590  4.09900
```

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



```
sales.pred <- predict(tree.sales, newdata = Carseats[-train, ])
plot(sales.pred, sales.test)
abline(0, 1)
```

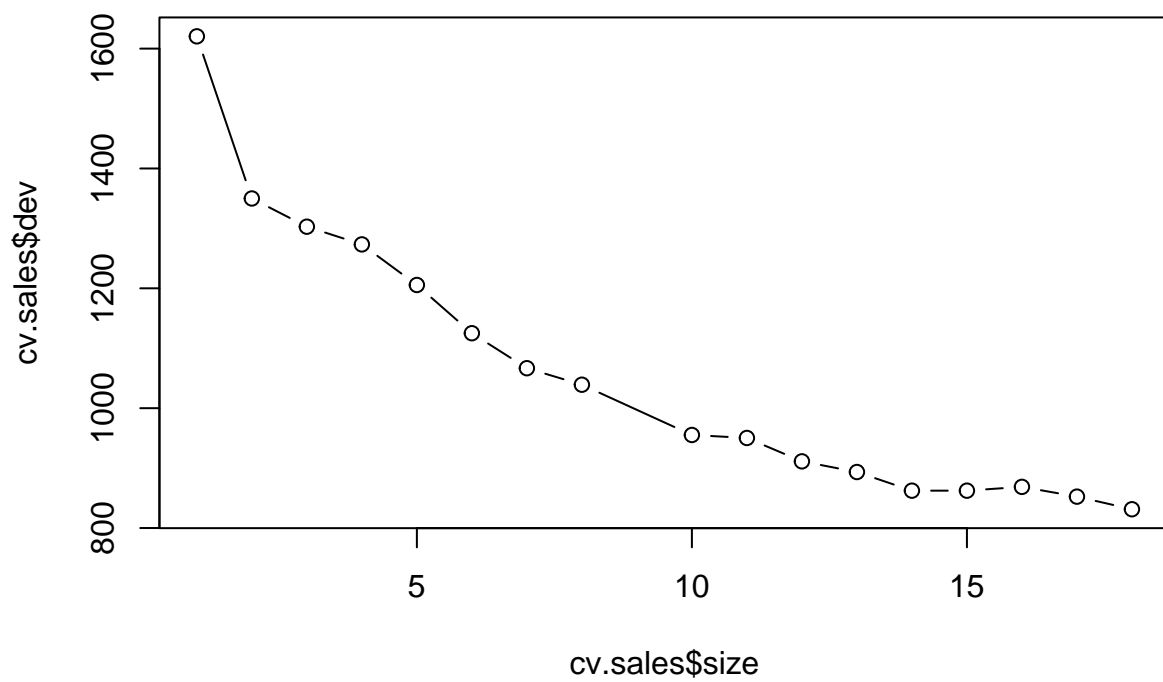


```
mean((sales.pred - sales.test)^2)
```

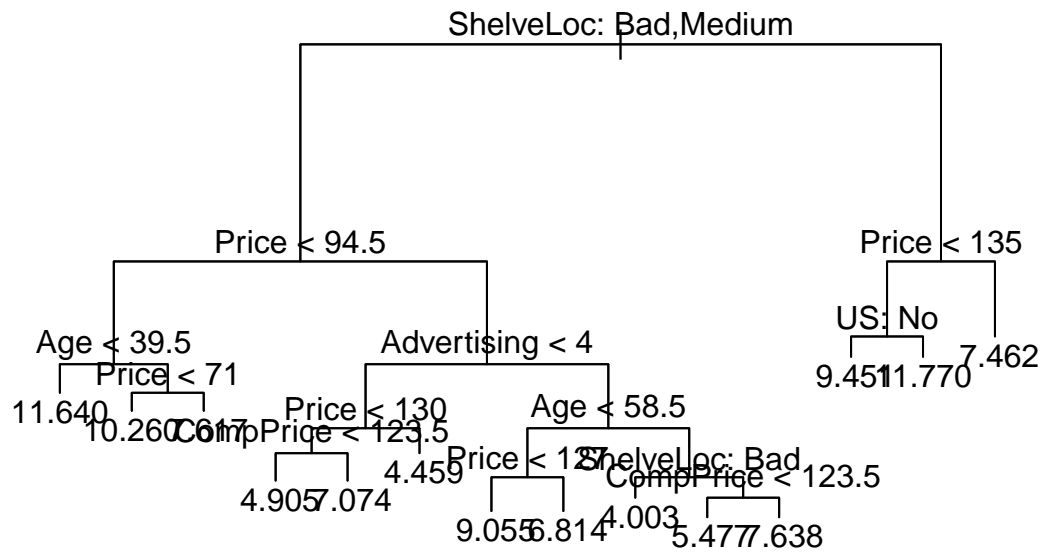
```
## [1] 4.922039
```

Part C

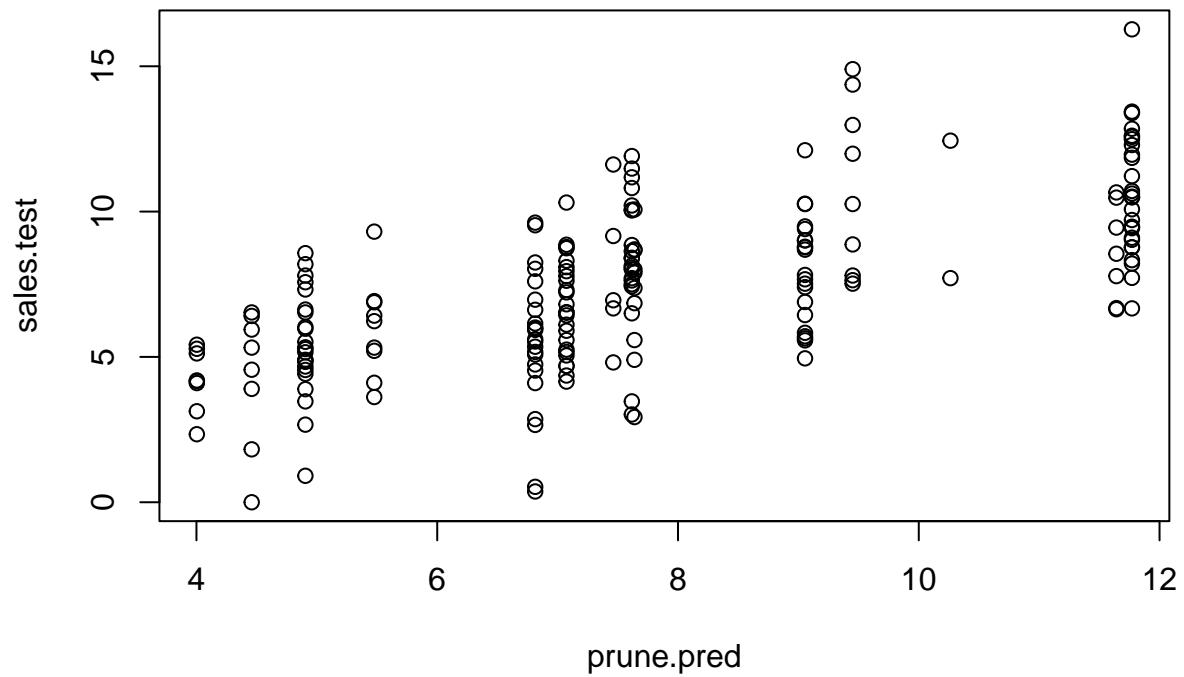
```
cv.sales <- cv.tree(tree.sales)
plot(cv.sales$size, cv.sales$dev, type = "b")
```



```
prune.sales <- prune.tree(tree.sales, best = 14)
plot(prune.sales)
text(prune.sales, pretty = 0)
```



```
prune.pred <- predict(prune.sales, newdata = Carseats[-train, ])
plot(prune.pred, sales.test)
```



```
mean((prune.pred - sales.test)^2)
```

```
## [1] 5.013738
```

Pruning the data set resulted in a higher MSE than when it was unpruned.

Part D

```
library(randomForest)
```

```
## Warning: package 'randomForest' was built under R version 4.3.2
```

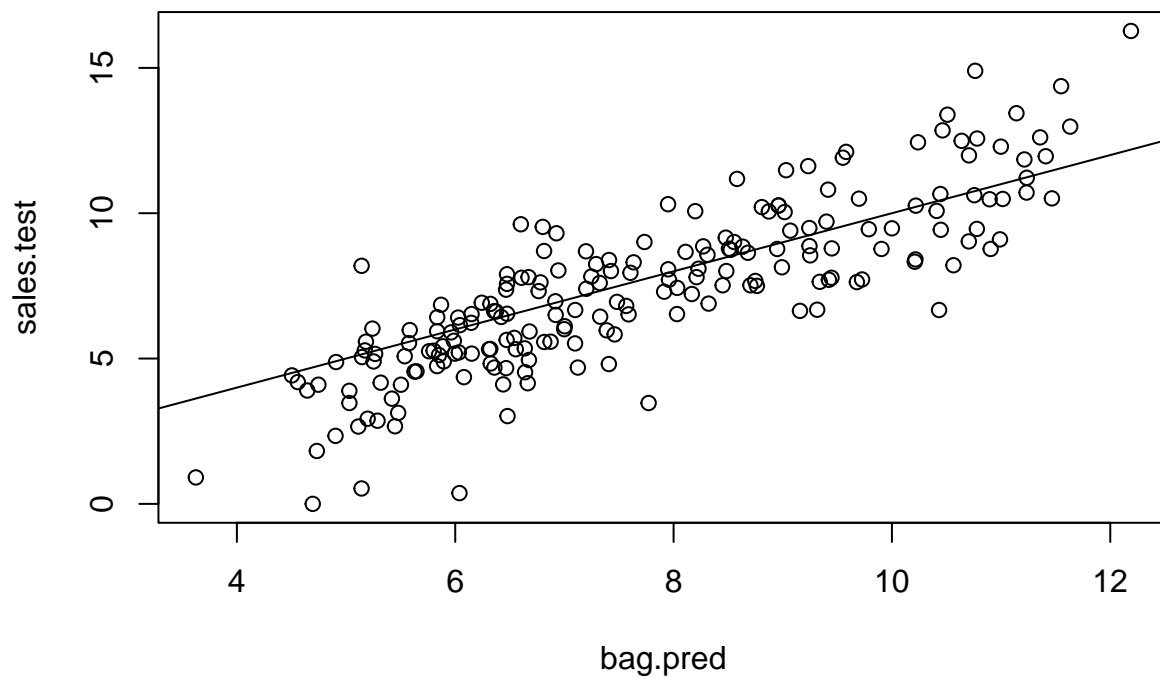
```
## randomForest 4.7-1.1
```

```
## Type rfNews() to see new features/changes/bug fixes.
```

```
set.seed(2)
bag.sales <- randomForest(Sales ~ ., data = Carseats, subset = train,
                          mtry = (ncol(Carseats) - 1), importance = T)
bag.sales
```

```
##
## Call:
## randomForest(formula = Sales ~ ., data = Carseats, mtry = (ncol(Carseats) - 1), importance = T
##           Type of random forest: regression
##           Number of trees: 500
## No. of variables tried at each split: 10
##
##           Mean of squared residuals: 2.908014
##           % Var explained: 63.02
```

```
bag.pred <- predict(bag.sales, newdata = Carseats[-train, ])
plot(bag.pred, sales.test)
abline(0, 1)
```



```
mean((bag.pred - sales.test)^2)
```

```
## [1] 2.586535
```

Part E

```
set.seed(2)
# Create random forest model using default m = p/3
rf.sales <- randomForest(Sales ~ ., data = Carseats, subset = train,
```

```

importance = T)
rf.pred <- predict(rf.sales, newdata = Carseats[-train, ])
mean((rf.pred - sales.test)^2)

```

```
## [1] 2.972319
```

```

set.seed(2)
# Create random forest model using default m = 6
rf.sales6 <- randomForest(Sales ~ ., data = Carseats, subset = train,
                          mtry = 6, importance = T)
rf.pred6 <- predict(rf.sales6, newdata = Carseats[-train, ])
mean((rf.pred6 - sales.test)^2)

```

```
## [1] 2.63335
```

```

set.seed(2)
# Create random forest model using default m = 9
rf.sales9 <- randomForest(Sales ~ ., data = Carseats, subset = train,
                          mtry = 9, importance = T)
rf.pred9 <- predict(rf.sales9, newdata = Carseats[-train, ])
mean((rf.pred9 - sales.test)^2)

```

```
## [1] 2.637777
```

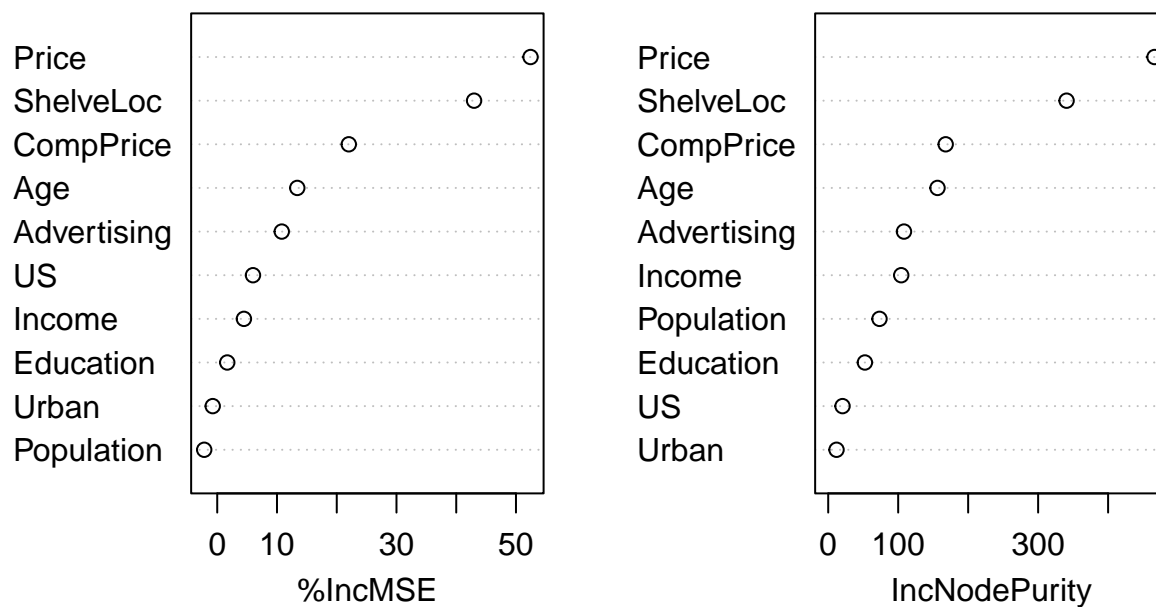
Setting m, the number of variables considered at each split,

```
importance(rf.sales6)
```

```
##           %IncMSE IncNodePurity
## CompPrice  22.0178739    167.89030
## Income     4.4666039    104.36139
## Advertising 10.8014745    108.29346
## Population -2.1801356     73.31029
## Price      52.4368458    466.35937
## ShelfLoc   42.9852979    340.65312
## Age        13.3967721    156.21255
## Education   1.6788660     52.43254
## Urban      -0.7559588     11.90340
## US         5.9833908     20.46648
```

```
varImpPlot(rf.sales6)
```


rf.sales6



Across all the trees considered, Price and ShelveLoc are the two most important variables.

Part F

```
# Analyze data using BART
library(BART)
```

```
## Warning: package 'BART' was built under R version 4.3.2
```

```
## Loading required package: nlme
```

```
## Loading required package: nnet
```

```
## Loading required package: survival
```

```
x <- Carseats[, 2:11]
y <- Carseats[, "Sales"]
xtrain <- x[train, ]
ytrain <- y[train]
xtest <- x[-train, ]
ytest <- y[-train]

# Run BART with default settings
set.seed(1)
bartfit <- gbart(xtrain, ytrain, x.test = xtest)
```

```
## *****Calling gbart: type=1
## *****Data:
## data:n,p,np: 200, 14, 200
## y1,yn: 2.781850, 1.091850
## x1,x[n*p]: 107.000000, 1.000000
## xp1,xp[np*p]: 111.000000, 1.000000
## *****Number of Trees: 200
## *****Number of Cut Points: 63 ... 1
## *****burn,nd,thin: 100,1000,1
## *****Prior:beta,alpha,tau,nu,lambda,offset: 2,0.95,0.273474,3,0.23074,7.57815
## *****sigma: 1.088371
## *****w (weights): 1.000000 ... 1.000000
## *****Dirichlet:sparse,theta,omega,a,b,rho,augment: 0,0,1,0.5,1,14,0
## *****printevery: 100
##
## MCMC
## done 0 (out of 1100)
## done 100 (out of 1100)
## done 200 (out of 1100)
## done 300 (out of 1100)
## done 400 (out of 1100)
## done 500 (out of 1100)
## done 600 (out of 1100)
## done 700 (out of 1100)
## done 800 (out of 1100)
## done 900 (out of 1100)
## done 1000 (out of 1100)
## time: 3s
## trcnt,tecnt: 1000,1000
```

```
bart.pred <- bartfit$yhat.test.mean
mean((ytest - bart.pred)^2)
```

```
## [1] 1.450842
```

BART yields a significantly lower test error than bagging and random forests.

Question 9

Part A

```
# Divide OJ data set into training and test data
set.seed(3)
train.oj <- sample(1:nrow(OJ), 800)
test.oj <- OJ[-train.oj, ]
```

Part B

```
# Fit a classification tree to OJ data with Purchase as the response
tree.oj <- tree(Purchase ~ ., data = OJ, subset = train.oj)
summary(tree.oj)
```

```
##
## Classification tree:
## tree(formula = Purchase ~ ., data = OJ, subset = train.oj)
## Variables actually used in tree construction:
## [1] "LoyalCH"      "PriceDiff"    "PriceMM"      "SalePriceMM"
## Number of terminal nodes: 9
## Residual mean deviance: 0.7247 = 573.2 / 791
## Misclassification error rate: 0.1812 = 145 / 800
```

The variables used in the classification tree construction are LoyalCH, PriceDiff, PriceMM, and SalePriceMM. There are 9 terminal nodes, and the training error rate is 18.12%.

Part C

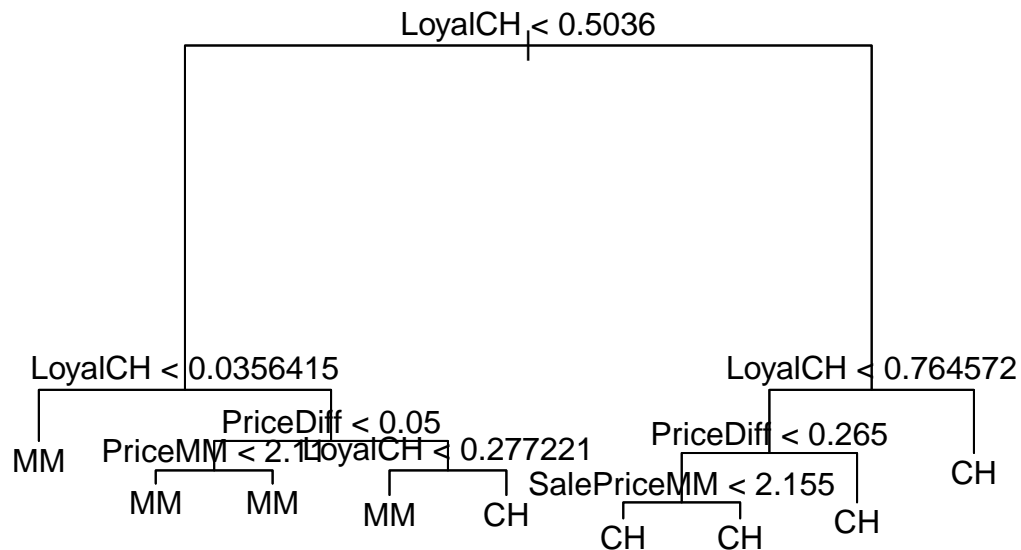
```
tree.oj
```

```
## node), split, n, deviance, yval, (yprob)
##      * denotes terminal node
##
## 1) root 800 1068.00 CH ( 0.61250 0.38750 )
##    2) LoyalCH < 0.5036 346 414.30 MM ( 0.28613 0.71387 )
##      4) LoyalCH < 0.0356415 57 0.00 MM ( 0.00000 1.00000 ) *
##      5) LoyalCH > 0.0356415 289 371.50 MM ( 0.34256 0.65744 )
##        10) PriceDiff < 0.05 114 105.90 MM ( 0.17544 0.82456 )
##          20) PriceMM < 2.11 89 94.84 MM ( 0.22472 0.77528 ) *
##          21) PriceMM > 2.11 25 0.00 MM ( 0.00000 1.00000 ) *
##        11) PriceDiff > 0.05 175 240.90 MM ( 0.45143 0.54857 )
##          22) LoyalCH < 0.277221 62 66.24 MM ( 0.22581 0.77419 ) *
##          23) LoyalCH > 0.277221 113 154.10 CH ( 0.57522 0.42478 ) *
##    3) LoyalCH > 0.5036 454 365.70 CH ( 0.86123 0.13877 )
##      6) LoyalCH < 0.764572 187 221.10 CH ( 0.72193 0.27807 )
##        12) PriceDiff < 0.265 113 154.70 CH ( 0.56637 0.43363 )
##          24) SalePriceMM < 2.155 102 141.20 CH ( 0.51961 0.48039 ) *
##          25) SalePriceMM > 2.155 11 0.00 CH ( 1.00000 0.00000 ) *
##        13) PriceDiff > 0.265 74 25.11 CH ( 0.95946 0.04054 ) *
##      7) LoyalCH > 0.764572 267 91.71 CH ( 0.95880 0.04120 ) *
```

Taking a look at node 24, the classification process can be described as followed: if LoyalCH is greater than 0.5036 and less than 0.764572, and if PriceDiff is less than 0.265, and if SalePriceMM is less than 2.155, then the response is classified as CH.

Part D

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



The decision tree shows that generally when customer brand loyalty is the sole predictor in the first two levels of the tree. Price difference is the next major predictor to be used in the tree. Price and sale price of Minute Maid are used in the last level of the tree.

Part E

```
# Use classification tree to predict responses of test data
oj.pred <- predict(tree.oj, test.oj, type = "class")
# Generate confusion matrix
table(oj.pred, test.oj$Purchase)
```

```
##
##  oj.pred  CH  MM
##      CH 148  31
##      MM  15  76
```

```
(15 + 31) / (148 + 31 + 15 + 76)
```

```
## [1] 0.1703704
```

The test error rate is 17%.

Part F

```
# Use cross-validation to determine if pruning the tree may result in better prediction error
set.seed(4)
cv.oj <- cv.tree(tree.oj, FUN = prune.misclass)
names(cv.oj)
```

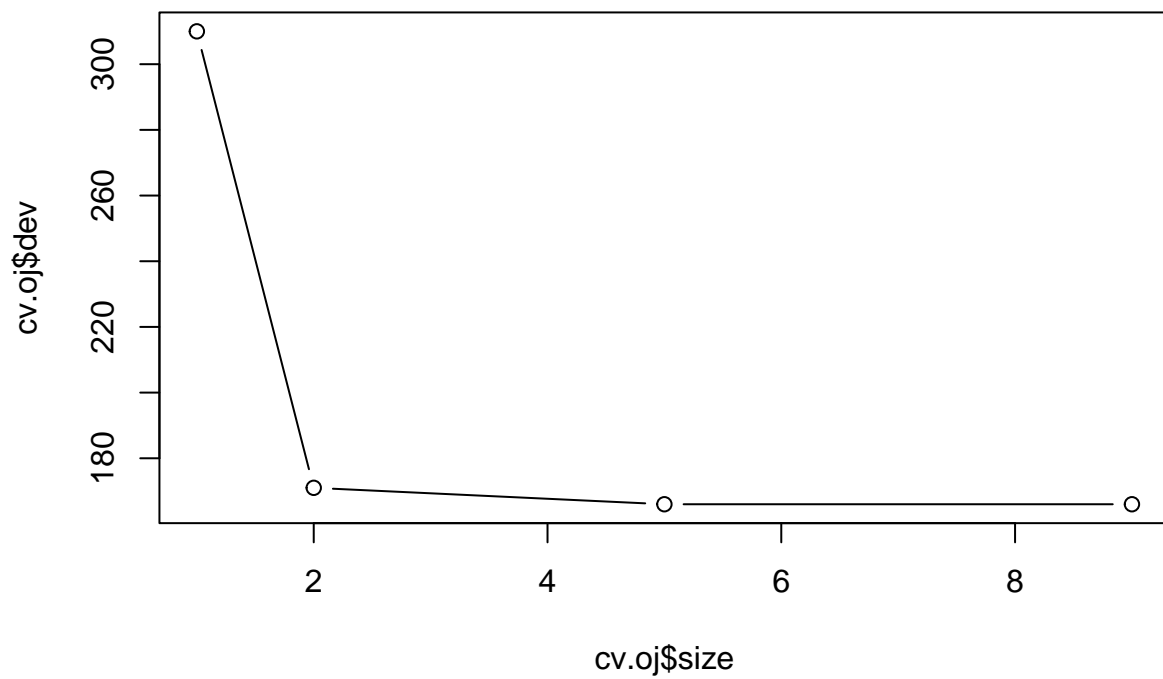
```
## [1] "size" "dev" "k" "method"
```

```
cv.oj
```

```
## $size
## [1] 9 5 2 1
##
## $dev
## [1] 166 166 171 310
##
## $k
## [1] -Inf 0.000000 5.666667 148.000000
##
## $method
## [1] "misclass"
##
## attr(,"class")
## [1] "prune" "tree.sequence"
```

Part G

```
# Create plot of cross-validated training error over tree size
plot(cv.oj$size, cv.oj$dev, type = "b")
```

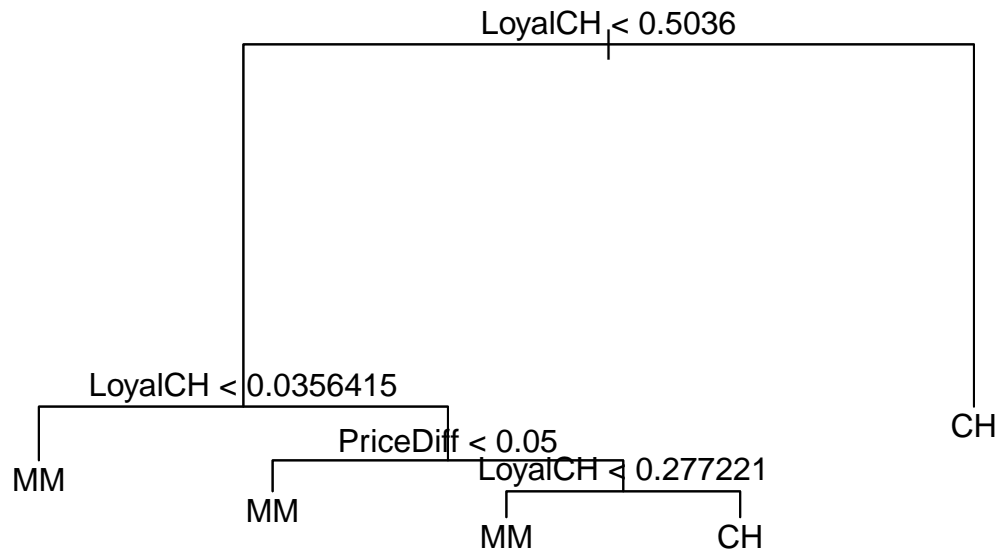


Part H

Tree size 5 corresponds to the lowest cross-validated classification error.

Part I

```
prune.oj <- prune.misclass(tree.oj, best = 5)
plot(prune.oj)
text(prune.oj, pretty = 0)
```



```
summary(prune.oj)
```

```
##
## Classification tree:
## snip.tree(tree = tree.oj, nodes = c(3L, 10L))
## Variables actually used in tree construction:
## [1] "LoyalCH" "PriceDiff"
## Number of terminal nodes: 5
## Residual mean deviance: 0.8703 = 691.9 / 795
## Misclassification error rate: 0.1812 = 145 / 800
```

Part J

The training errors are the same for the pruned tree and the original classification tree we created.

Part K

```
# Predict response with pruned tree
prune.oj.pred <- predict(prune.oj, test.oj, type = "class")
# Generate confusion matrix for pruned tree
table(prune.oj.pred, test.oj$Purchase)
```

```
##
## prune.oj.pred  CH  MM
##              CH 148 31
##              MM  15 76
```

```
(15 + 31) / (148 + 31 + 15 + 76)
```

```
## [1] 0.1703704
```

The pruned tree yields the same test error as the original tree.

Question 10

Part A

```
hitters.data <- na.omit(Hitters)
hitters.data$Salary <- log(hitters.data$Salary)
```

Part B

```
hitters.train <- c(1:200)
hitters.test  <- c(201:nrow(hitters.data))
```

Part C

```
library(gbm)
```

```
## Warning: package 'gbm' was built under R version 4.3.2
```

```
## Loaded gbm 2.1.8.1
```

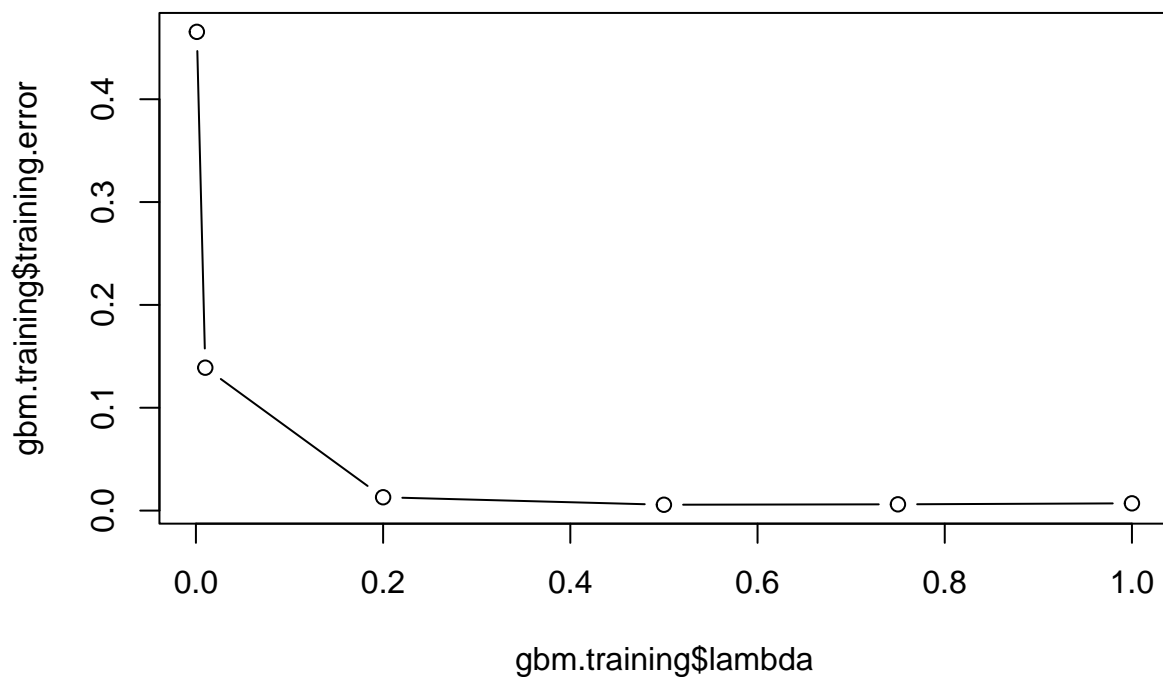
```
tunings <- c(0.001, 0.01, 0.2, 0.5, 0.75, 1.0)
gbm.training <- data.frame(lambda = tunings,
                           training.error = rep(NA, length(tunings)))
for (x in 1:length(tunings)) {
  set.seed(5)
  boost.hitters <- gbm(Salary ~ ., data = hitters.data[hitters.train, ],
                      distribution = "gaussian", n.trees = 1000,
                      interaction.depth = 4, shrinkage = tunings[x])
  gbm.training[x, "training.error"] <- mean(boost.hitters$train.error)
}

gbm.training
```



```
##   lambda training.error
## 1 0.001    0.465506603
## 2 0.010    0.138947034
## 3 0.200    0.012968059
## 4 0.500    0.005684627
## 5 0.750    0.006102646
## 6 1.000    0.007057622
```

```
plot(gbm.training$lambda, gbm.training$training.error, type = "b")
```



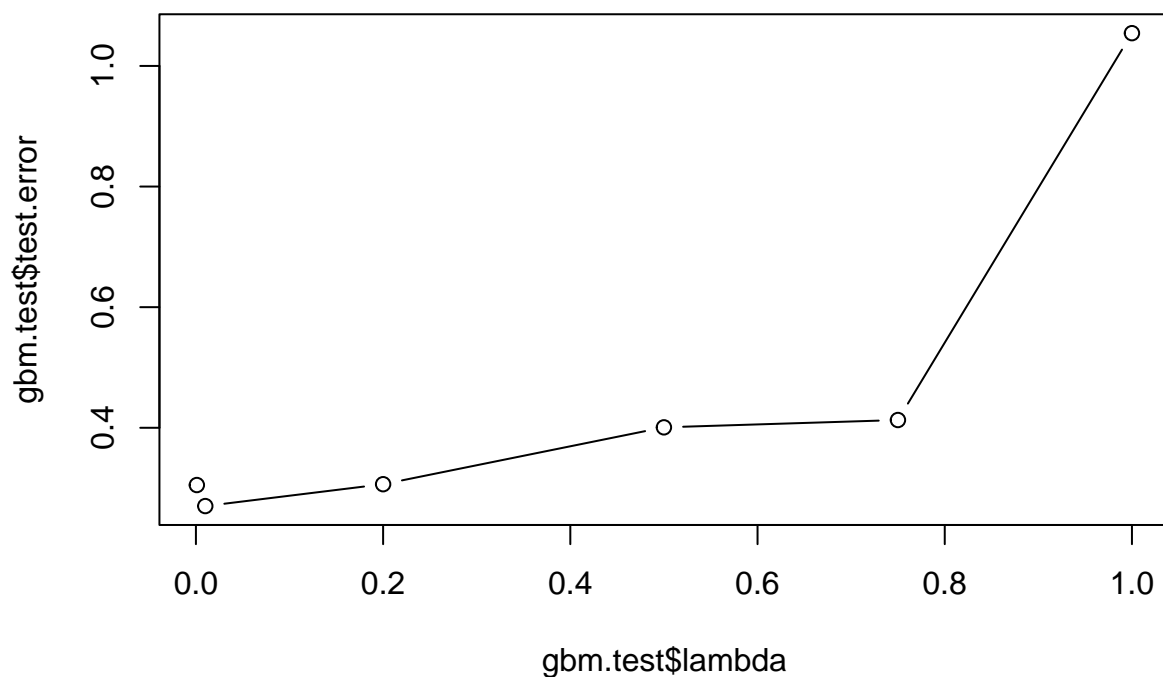
Part D

```
gbm.test <- data.frame(lambda = tunings,
                        test.error = rep(NA, length(tunings)))
for (x in 1:length(tunings)) {
  set.seed(5)
  boost.hitters <- gbm(Salary ~ ., data = hitters.data[hitters.train, ],
                       distribution = "gaussian", n.trees = 1000,
                       interaction.depth = 4, shrinkage = tunings[x])
  yhat.boost <- predict(boost.hitters, newdata = hitters.data[hitters.test, ],
                        n.trees = 1000)
  gbm.test[x, "test.error"] <- mean((yhat.boost - hitters.data$Salary[hitters.test])^2)
}
```

```
gbm.test
```

```
##   lambda test.error
## 1  0.001  0.3050198
## 2  0.010  0.2701642
## 3  0.200  0.3064144
## 4  0.500  0.4007037
## 5  0.750  0.4128302
## 6  1.000  1.0543288
```

```
plot(gbm.test$lambda, gbm.test$test.error, type = "b")
```



Part E

```
# Create LS regression with Salary as the predictor and determine test MSE
lm.hitters <- lm(Salary ~ ., data = hitters.data, subset = hitters.train)
lm.predict <- predict(lm.hitters, newdata = hitters.data[hitters.test, ])
lm.mse <- mean((lm.predict - hitters.data$Salary[hitters.test])^2)
lm.mse
```

```
## [1] 0.4917959
```

```
library(glmnet)
```

```
## Warning: package 'glmnet' was built under R version 4.3.2
```

```
## Loading required package: Matrix
```

```
## Loaded glmnet 4.1-8
```

```
x <- model.matrix(Salary ~ ., hitters.data)[, -1]
y <- hitters.data$Salary

lambda.grid <- 10^seq(10, -2, length = 100)
lasso.mod <- glmnet(x[hitters.train, ], y[hitters.train], alpha = 1,
                    lambda = lambda.grid)
set.seed(7)
cv.out <- cv.glmnet(x[hitters.train, ], y[hitters.train], alpha = 1)
bestlam <- cv.out$lambda.min

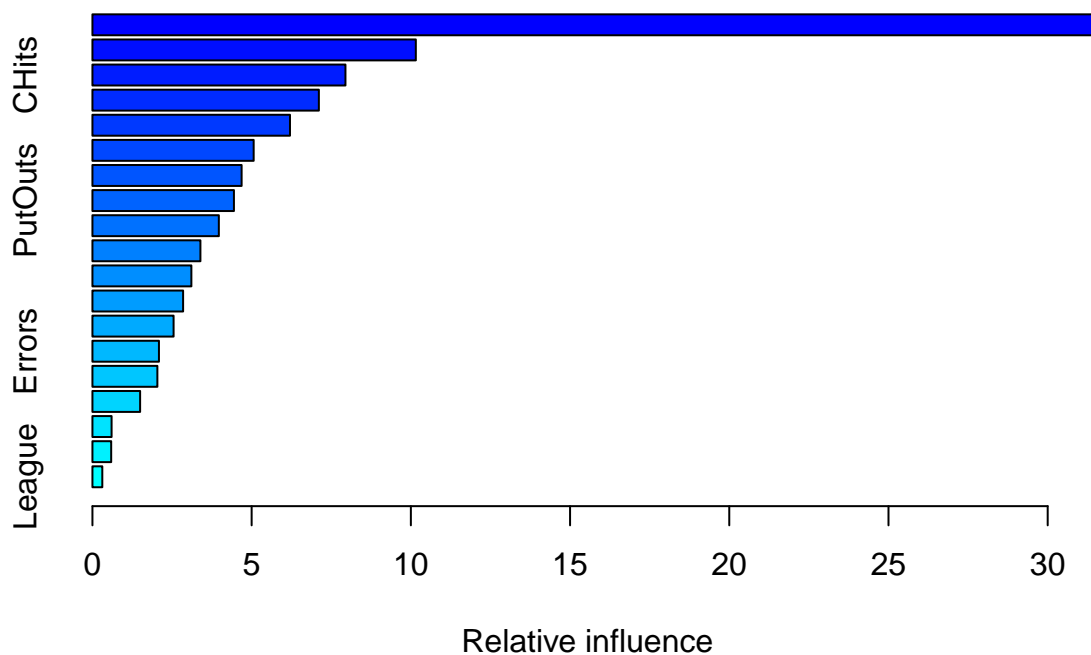
lasso.pred <- predict(lasso.mod, s = bestlam, newx = x[hitters.test, ])
lasso.mse <- mean((lasso.pred - y[hitters.test])^2)
lasso.mse
```

```
## [1] 0.470371
```

The boosting MSE when $\lambda = 0.01$ is much smaller than the MSE of a multiple linear regression and the Lasso.

Part F

```
# Create boosting model using shrinkage, or tuning parameter, equal to 0.01
set.seed(5)
boost.hitters <- gbm(Salary ~ ., data = hitters.data[hitters.train, ],
                     distribution = "gaussian", n.trees = 1000,
                     interaction.depth = 4, shrinkage = 0.01)
summary(boost.hitters)
```



##	var	rel.inf
##	CATBat	31.4028722
##	CWalks	10.1559784
##	CHits	7.9460562
##	CRBI	7.1113938
##	CRuns	6.2042978
##	Years	5.0629877
##	Walks	4.6850235
##	PutOuts	4.4432945
##	CHmRun	3.9699283
##	AtBat	3.3906305
##	Assists	3.1067186
##	RBI	2.8470835
##	Hits	2.5482569
##	Errors	2.0887654
##	HmRun	2.0395010
##	Runs	1.4972505
##	NewLeague	0.6009227
##	Division	0.5889558
##	League	0.3100826

CATBat is clearly the most important predictor in the boosting model. CWalks, CRuns, and CRBI are the next highest in importance.

Part G

```
set.seed(5)
bag.hitters <- randomForest(Salary ~ ., data = hitters.data,
                           subset = hitters.train, mtry = 19,
                           importance = T)
yhat.bag <- predict(bag.hitters, newdata = hitters.data[hitters.test, ])
mean((yhat.bag - hitters.data$Salary[hitters.test])^2)
```

```
## [1] 0.233566
```

Bagging yields a test MSE of 0.234. This is slightly lower than the test MSE of boosting with tuning parameter equal to 0.01, which was 0.270.