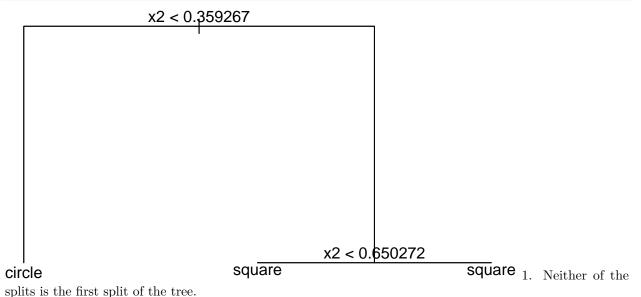
lab7 YILIN LI 2019/11/8

Data generation

```
set.seed(75)
n <- 16
x1 <- runif(n)
x2 <- runif(n)
group <- as.factor(sample(1:3, n, replace = TRUE))
levels(group) <- c("circle", "triangle", "square")
df <- data.frame(x1, x2, group)
df[1, 2] <- .765 # tweaks to make a more interesting configuration
df[9, 1] <- .741
df <- df[-7, ]</pre>
```

$\mathbf{Q}\mathbf{1}$

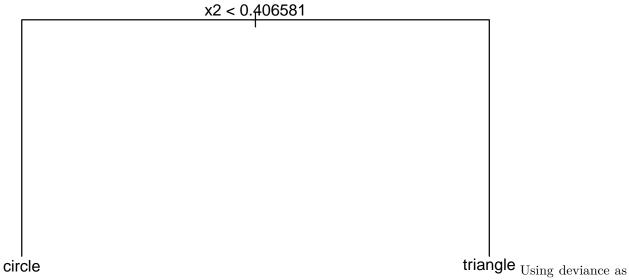
```
library(tree)
t1 = tree(group~., data = df, split = "gini")
plot(t1)
text(t1,pretty = 0)
```



- nos is the first spire of the tree.
- 2. Even though the second split predicts all data as square, it's useful because this split could futher increase the purity of all the terminals.
- 3. It would predict "square"

 $\mathbf{Q2}$

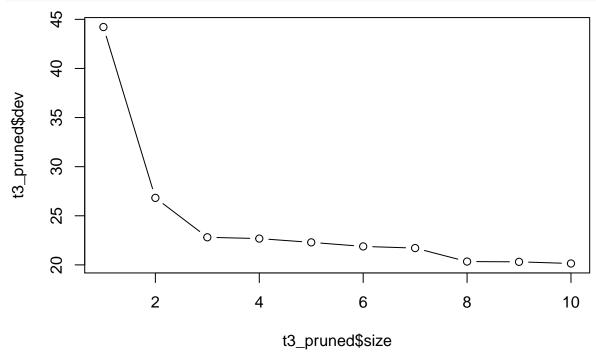
```
t2 = tree(group~., data = df, split = "deviance")
plot(t2)
text(t2,pretty = 0)
```



the split method is different from using gini index because with deviance, the tree will try to lower the misclassification rate, but with gini, the tree will try to increase purity among terminals.

Q3

```
d = read.csv("http://andrewpbray.github.io/data/crime-train.csv")
d["NumIllegsr"] = sqrt(d$NumIlleg)
t3 = tree(ViolentCrimesPerPop ~ racePctWhite + PctKids2Par +NumIllegsr,data = d)
t3_pruned = cv.tree(t3)
plot(t3_pruned$size, t3_pruned$dev, type = 'b')
```



```
t3_best = prune.tree(t3,method = "deviance",best = 10)
plot(t3_best)
text(t3_best,pretty = 0)
```

```
PctKids2Par < 0.295
Numlllegsr < 0.295
0.5233@.7811@.6689@.41970 Numlllegsr < 0.186603 PctKids2Par < 0.615
0.2955@.18730.4002@.2057@.11320.07251
```

$\mathbf{Q4}$

```
test_data <- read.csv("https://bit.ly/2PYS8Ap")
test_data["NumIllegsr"] = sqrt(test_data$NumIlleg)
pd_tree = predict(t3_best, test_data)
MSE_tree = mean((pd_tree-test_data$ViolentCrimesPerPop)^2)
m1 = lm(ViolentCrimesPerPop ~ racePctWhite + PctKids2Par +NumIllegsr, data = d)
pd_lm = predict(m1, test_data)
MSE_lm = mean((pd_lm - test_data$ViolentCrimesPerPop)^2)
MSE_tree
## [1] 0.01935877
MSE_lm
## [1] 0.0205459</pre>
```

So, clearly, the pruned tree has a lower test MSE than our linear fitting.

$\mathbf{Q5}$

```
library(randomForest)

## randomForest 4.6-14

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

rf = randomForest(ViolentCrimesPerPop ~ racePctWhite + PctKids2Par +NumIllegsr, data = d, mtry = 3, imp
pd_rf = predict(rf,test_data)

MSE_rf1 = mean((pd_rf-test_data$ViolentCrimesPerPop)^2)

rf2 = randomForest(ViolentCrimesPerPop ~ racePctWhite + PctKids2Par +NumIllegsr, data = d, mtry = 1, im
pd_rf2 = predict(rf2,test_data)
```

```
MSE_rf2 = mean((pd_rf2-test_data$ViolentCrimesPerPop)^2)
MSE_rf1
## [1] 0.00600712
MSE_rf2
```

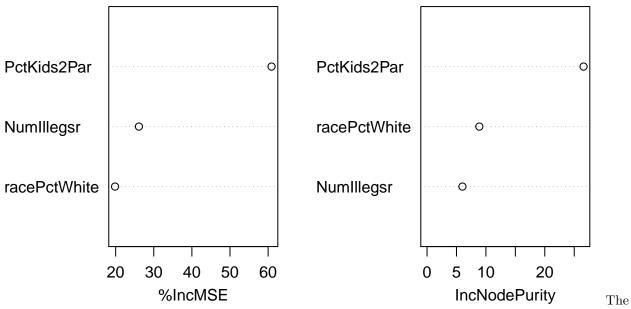
[1] 0.009030661

It turns out both the bagging trees and the random forest have lower test MSE than the pruned tree and linear regression model.

$\mathbf{Q6}$

```
importance = varImpPlot(rf)
```

rf



result turns out to be quite similar to our analysis in our linear model. "PctKids2Par" has the highest correlation with the "ViolentCrimesPerPop" and here also influences the randomForest model the most.