

Regression Tree with HSAUR

Yamuna Dhungana

Exercises

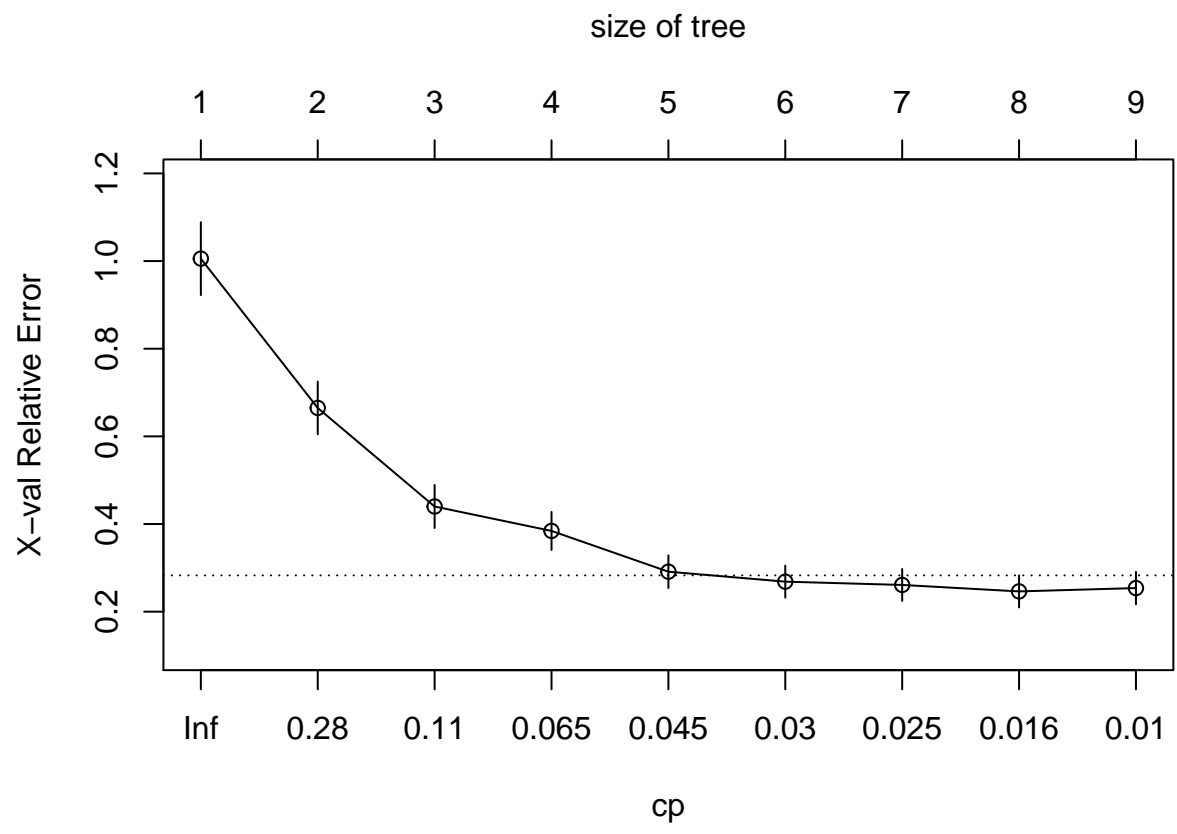
1. (Ex. 9.1 pg 186 in HSAUR, modified for clarity) The **BostonHousing** dataset reported by Harrison and Rubinfeld (1978) is available as a `data.frame` structure in the **mlbench** package (Leisch and Dimitriadou, 2009). The goal here is to predict the median value of owner-occupied homes (`medv` variable, in 1000s USD) based on other predictors in the dataset.

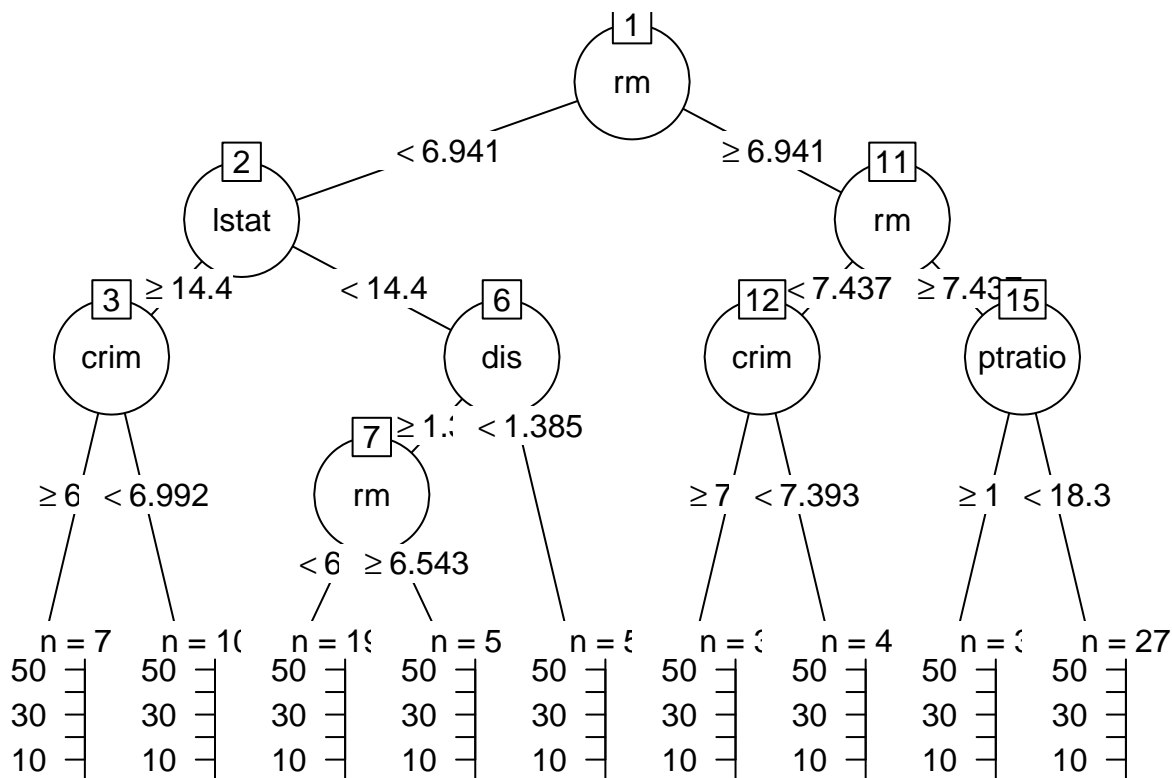
a) Construct a regression tree using `rpart()`. Discuss the results, including these key components:

- How many nodes does your tree have?
- Did you prune the tree? Did it decrease the number of nodes?
- What is the prediction error (MSE)?
- Plot the predicted vs. observed values.
- Plot the final tree.

First of all, installing all the required libraries for the exercise

```
##
## Regression tree:
## rpart(formula = medv ~ ., data = BostonHousing, control = rpart.control(minsplit = 10))
##
## Variables actually used in tree construction:
## [1] crim    dis    lstat  ptratio rm
##
## Root node error: 42716/506 = 84.42
##
## n= 506
##
##      CP nsplit rel error  xerror   xstd
## 1 0.452744     0  1.00000 1.00556 0.083187
## 2 0.171172     1  0.54726 0.66485 0.060191
## 3 0.071658     2  0.37608 0.43995 0.049160
## 4 0.059002     3  0.30443 0.38423 0.043400
## 5 0.033756     4  0.24542 0.29121 0.037190
## 6 0.026613     5  0.21167 0.26852 0.036445
## 7 0.023572     6  0.18506 0.26098 0.036438
## 8 0.010859     7  0.16148 0.24626 0.036535
## 9 0.010000     8  0.15062 0.25383 0.037104
```



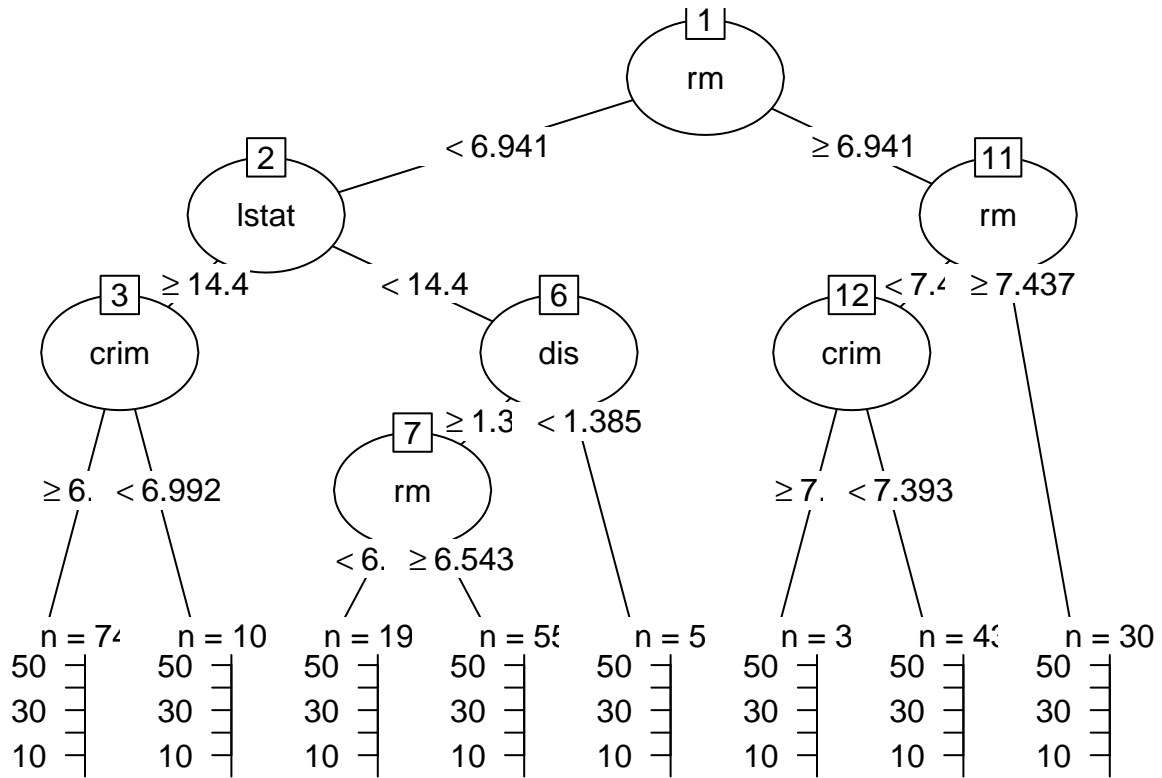


```

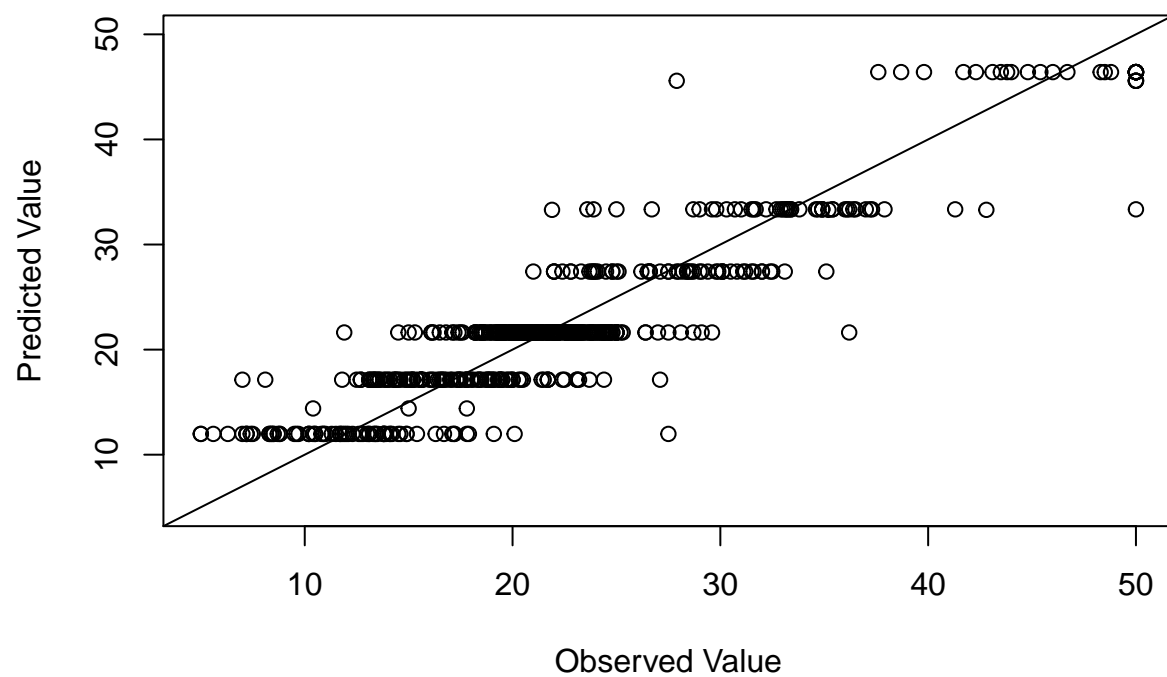
## n= 506
##
## node), split, n, deviance, yval
##      * denotes terminal node
##
## 1) root 506 42716.3000 22.53281
##    2) rm< 6.941 430 17317.3200 19.93372
##      4) lstat>=14.4 175 3373.2510 14.95600
##        8) crim>=6.99237 74 1085.9050 11.97838 *
##        9) crim< 6.99237 101 1150.5370 17.13762 *
##      5) lstat< 14.4 255 6632.2170 23.34980
##        10) dis>=1.38485 250 3721.1630 22.90520
##          20) rm< 6.543 195 1636.0670 21.62974 *
##          21) rm>=6.543 55 643.1691 27.42727 *
##        11) dis< 1.38485 5 390.7280 45.58000 *
##    3) rm>=6.941 76 6059.4190 37.23816
##      6) rm< 7.437 46 1899.6120 32.11304
##        12) crim>=7.393425 3 27.9200 14.40000 *
##        13) crim< 7.393425 43 864.7674 33.34884 *
##      7) rm>=7.437 30 1098.8500 45.09667
##        14) ptratio>=18.3 3 223.8200 33.30000 *
##        15) ptratio< 18.3 27 411.1585 46.40741 *
##
##      CP nsplit rel error      xerror      xstd
## 1 0.45274420      0 1.0000000 1.0055569 0.08318656
## 2 0.17117244      1 0.5472558 0.6648488 0.06019104
## 3 0.07165784      2 0.3760834 0.4399528 0.04916012

```

```
## 4 0.05900152      3 0.3044255 0.3842313 0.04339967
## 5 0.03375589      4 0.2454240 0.2912136 0.03718954
## 6 0.02661300      5 0.2116681 0.2685244 0.03644484
## 7 0.02357238      6 0.1850551 0.2609760 0.03643787
## 8 0.01085935      7 0.1614827 0.2462587 0.03653519
## 9 0.01000000      8 0.1506234 0.2538281 0.03710374
```

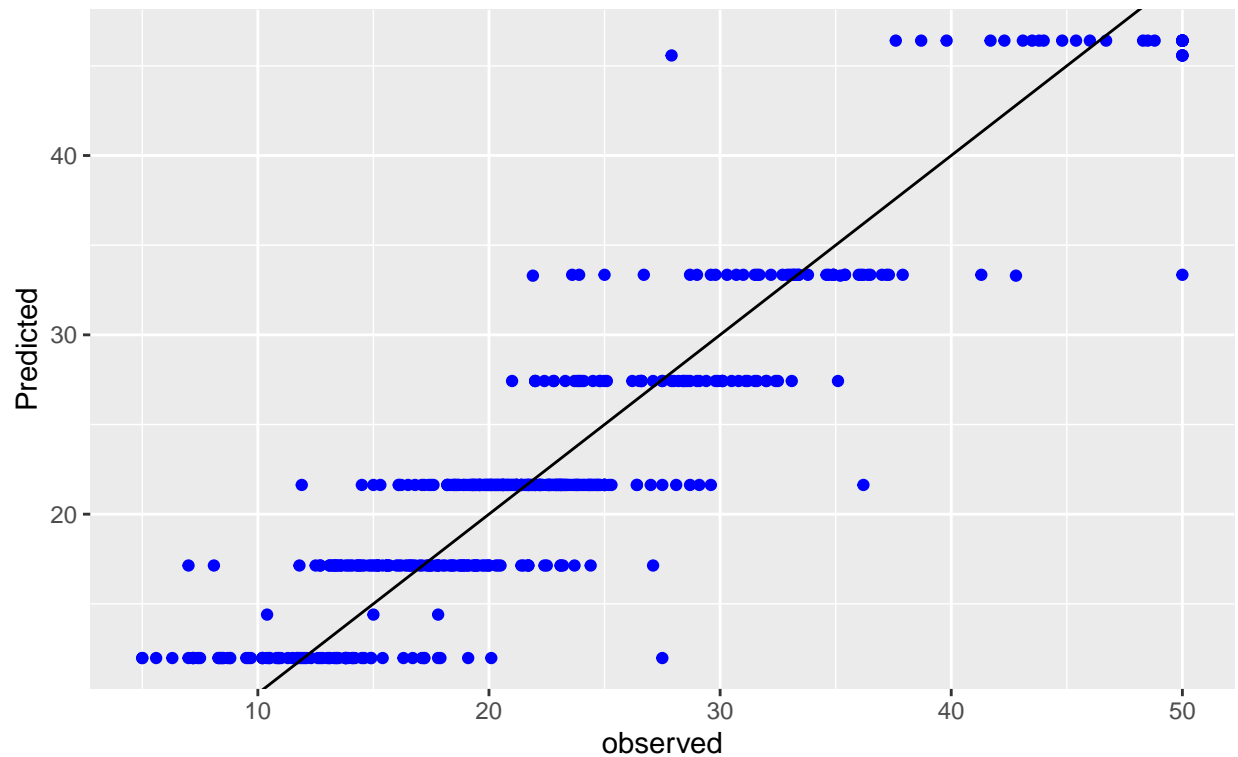


**Base R: Observed vs predicted values for
regression tree**



```
## integer(0)
```

ggplot: Observed vs predicted values for
Regression tree



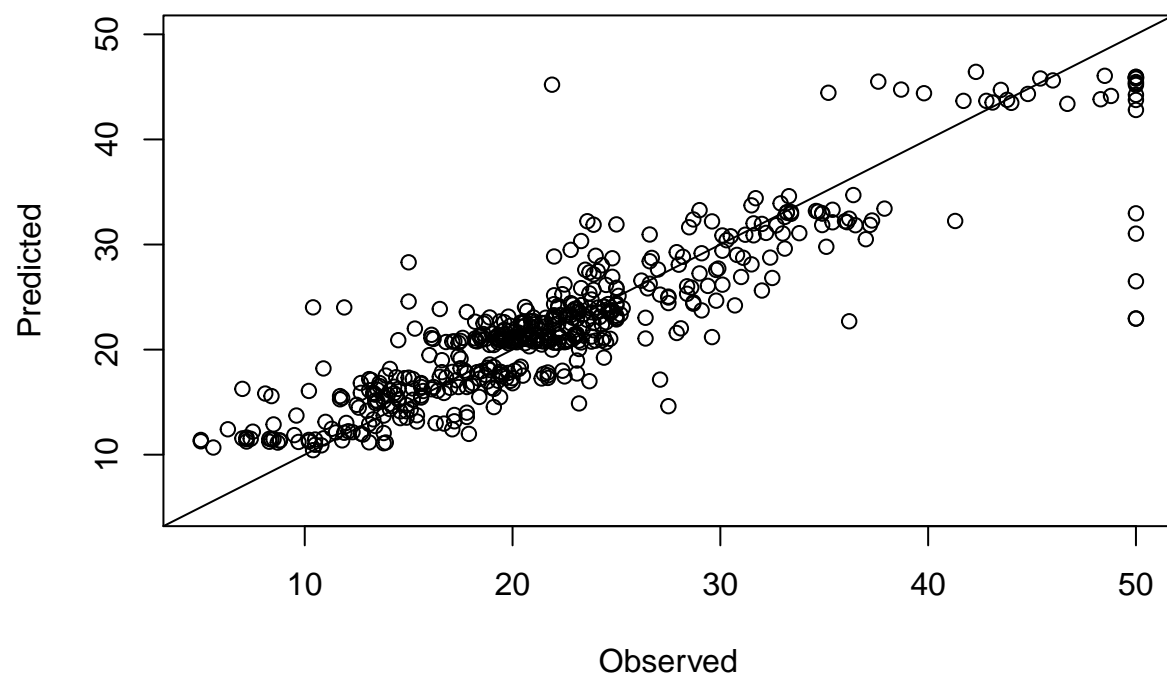
The MSE for Regression Tree is: 12.71556

My tree has nine(9) nodes. Yes I pruned the tree but I couldn't see any changes in my tree. It already has the minimum no of tree therefore, the prune was not so effective. The predictive error of the data is 12.71556

- b) Apply bagging with 50 trees. Report the prediction error (MSE) and plot the predicted vs observed values.

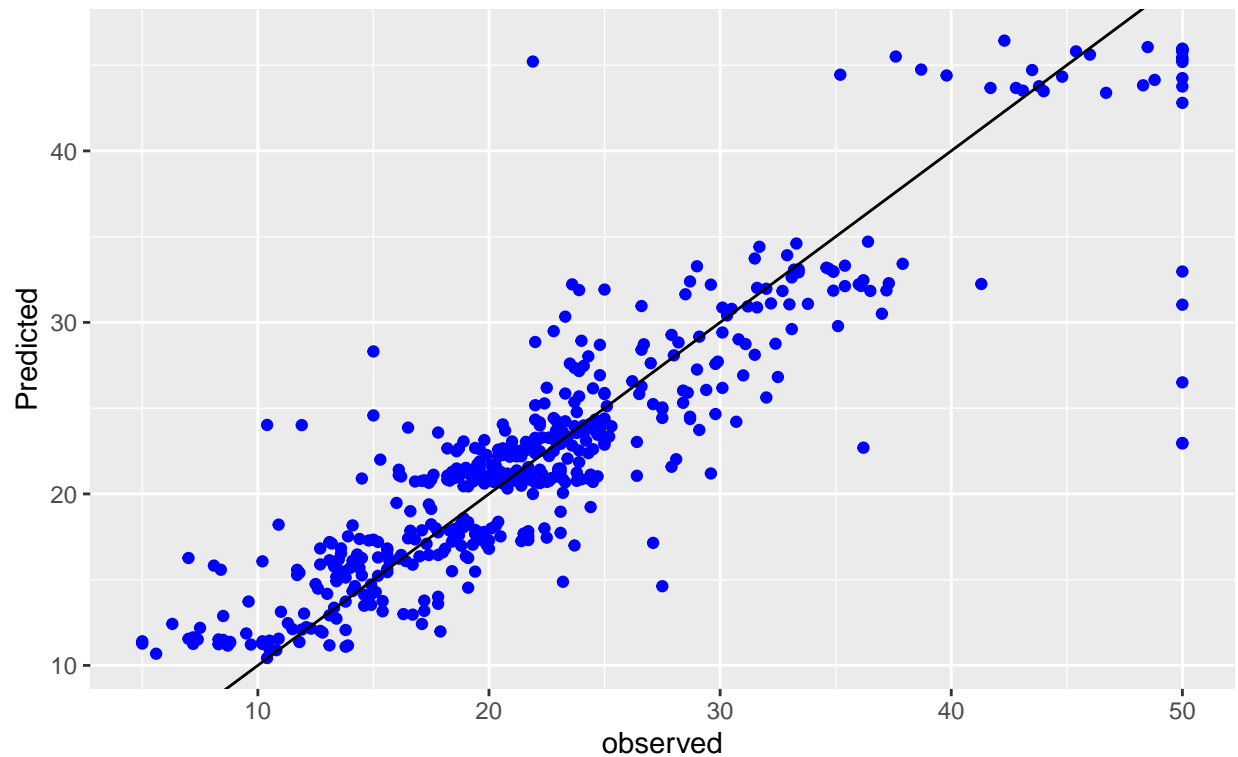
The MSE Bagging 50 is: 17.21584

**Base R: Observed vs predicted values for
bagging_50**



```
## integer(0)
```

ggplot: Observed vs predicted values for
bagging_50

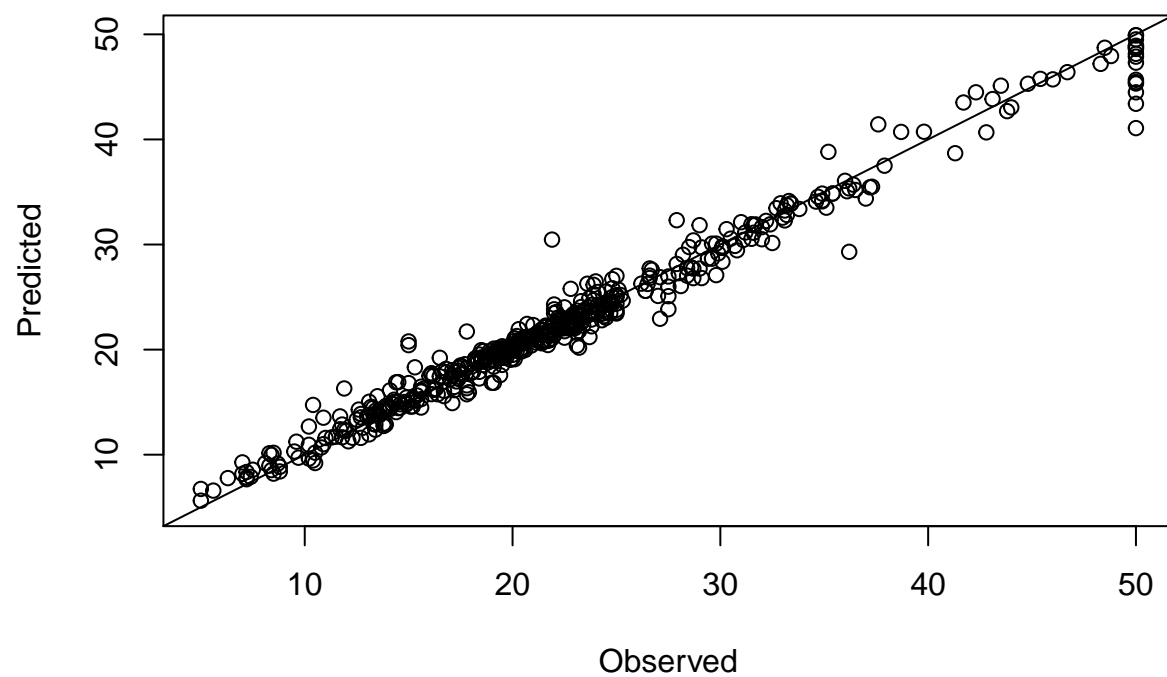


The MSE for the bagging of 50 data is 17.21584. Most of the predicted values are seen in between the 10 to 40 and very few near 50.

- c) Apply bagging using the `randomForest()` function. Report the prediction error (MSE). Was it the same as (b)? If they are different what do you think caused it? Plot the predicted vs. observed values.

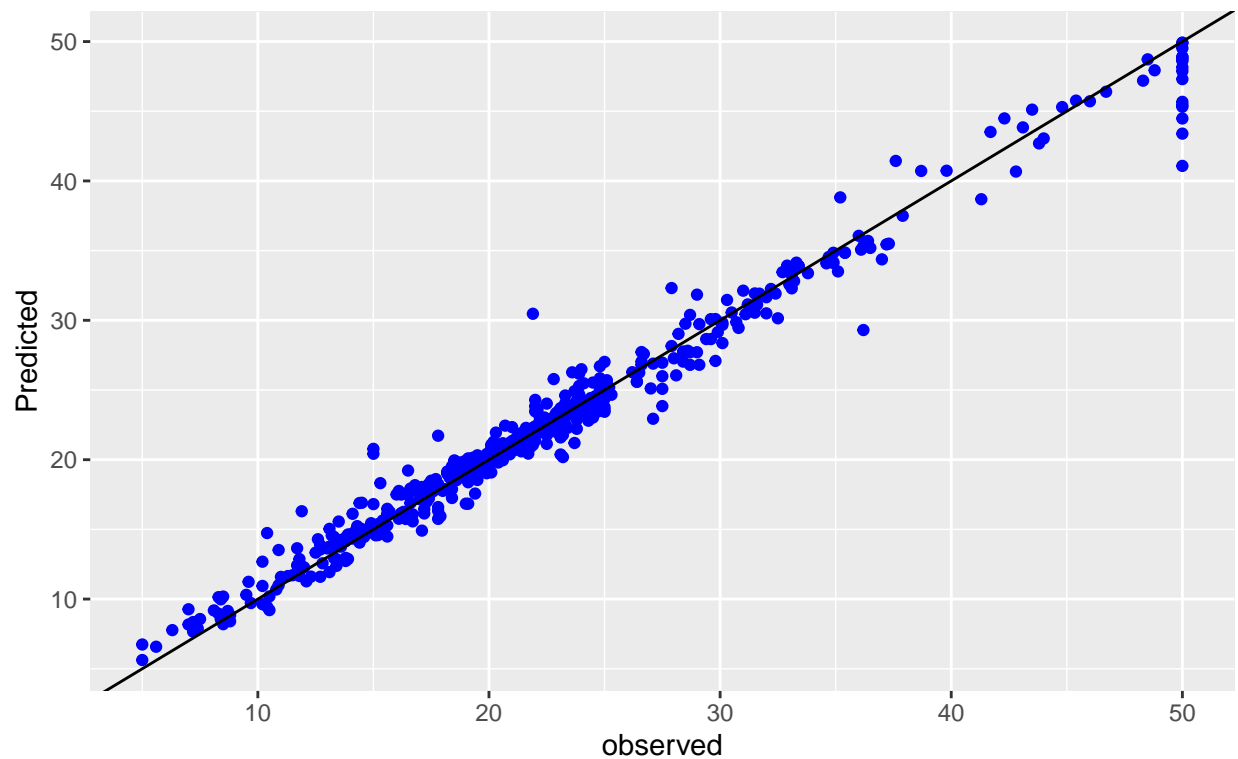
```
## MSE for Random Forest mtry13 : 2.044668
```


**Base R: Observed vs predicted values for
Randomforest_mtry13()**



```
## integer(0)
```

ggplot:Observed vs predicted values for
Randomforest_mtry13()

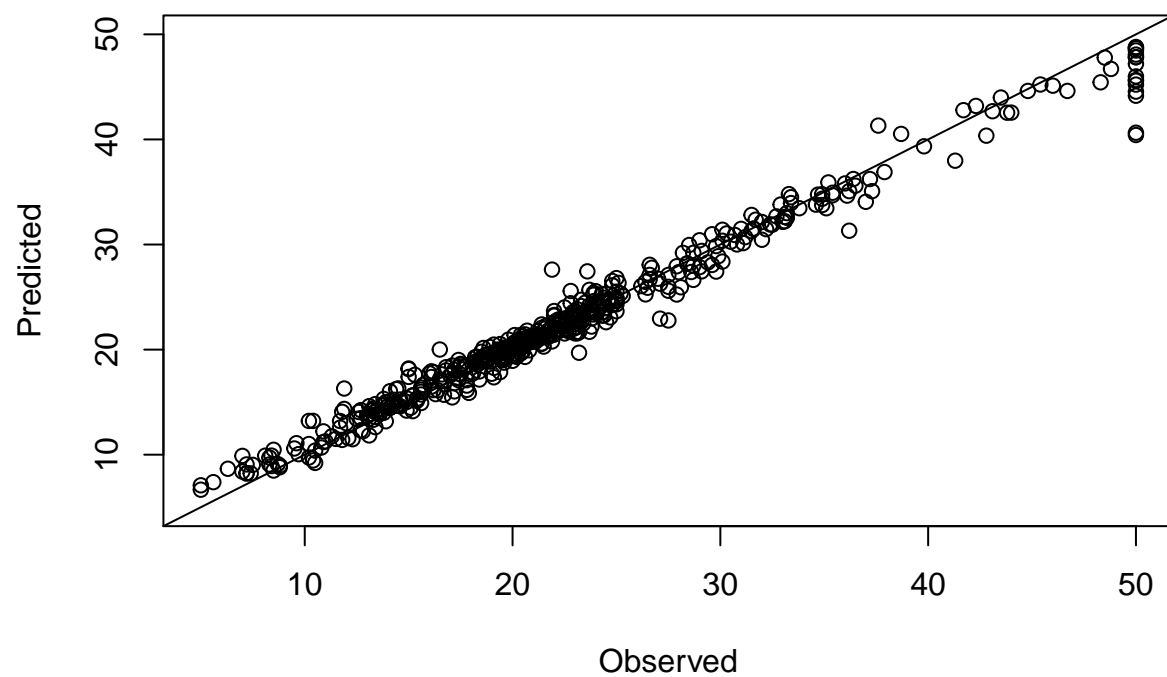


The MSE of the random_forest is 1.788203, I think the graph is some what different from the question-b . In question-b the graph is bit more scattered around the regression line. Whereas, in this question the sample are more closely packed within the lines. This might be because we used mtry =13 which means using all the variables with 50 trees.

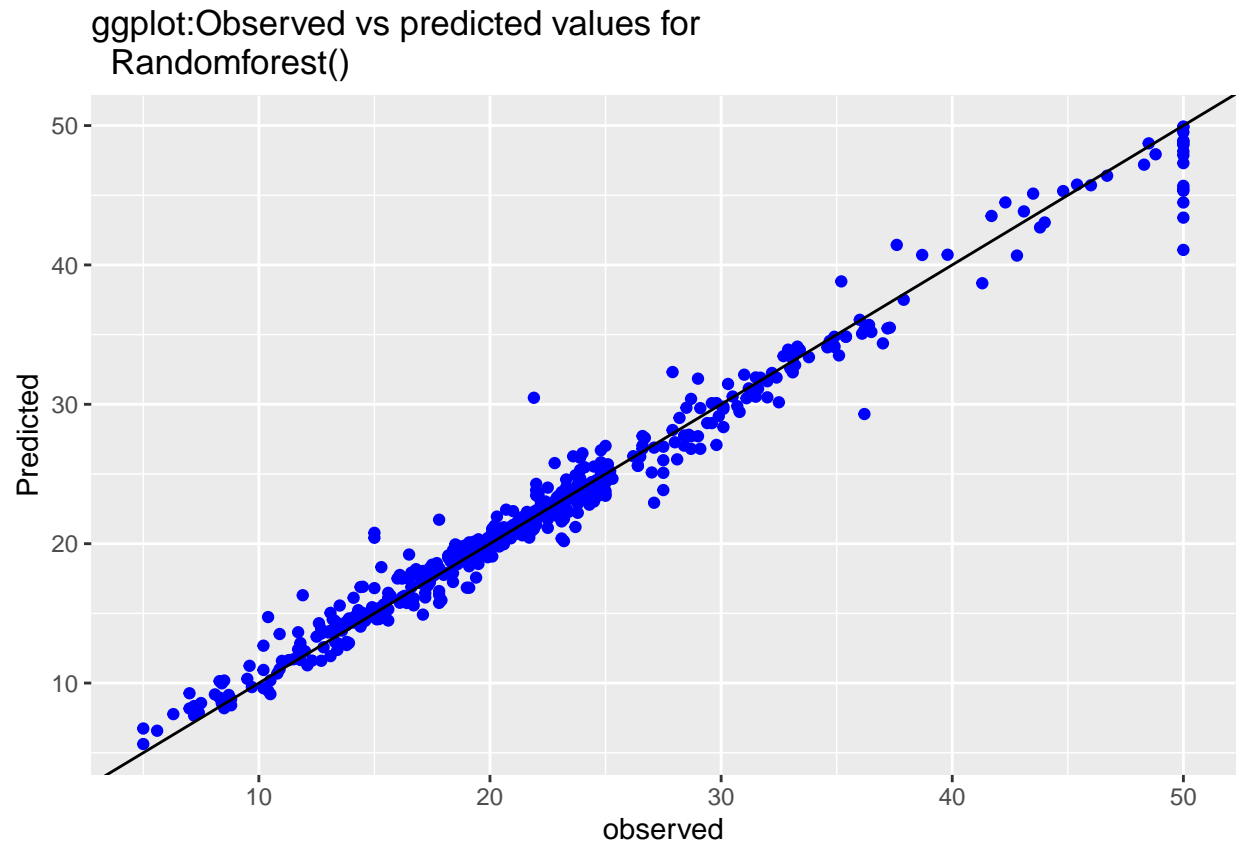
- d) Use the randomForest() function to perform random forest. Report the prediction error (MSE). Plot the predicted vs. observed values.

MSE for Random Forest: 1.994478

Base R: Observed vs predicted values for Randomforest()



```
## integer(0)
```



The MSE of the Random Forest whose default no of trees is 500 is 1.946478. The observed vs predicted data looks similar to the data with the random forest with 50 trees.

e) Include a table of each method and associated MSE. Which method is more accurate?

The error table for the data

	Regression_Tree	Bagging_50	Random_Forest_mtry13	Random_Forest
## 1	12.71556	17.21584	2.044668	1.994478

Since, more trees is always better because of less error, Here Random forest with the default no of trees of 500 has the lowest MSE. Therefore, Random forest is more accurate.