

HST.953x Workshop 2.09: Ensemble Methods Exercises

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21 Jul 2021

Contents

Description	1
Dataset	1
Test-Train Split the Data	2
Initial Tree Model of Median Value versus all other Predictors	2
Initial Linear Model of Median Value versus all other Predictors	3
Bagging	4
Random Forest	5
Boosting	6
Results	8
Linear Model of Median Value versus two most significant predictors from boosting (lstat and rm)	9

(Note: Updated and modified from “R for Statistical Learning”).

Description

This section contains examples of ensemble - bagging and boosting - methods.

Dataset

We will be using the **Boston** dataset from the **MASS** package for these examples. The dataset contains housing value data for the suburbs of Boston.

- **crim** - per capita crime rate by town.
- **zn** - proportion of residential land zoned for lots over 25,000 sq.ft.
- **indus** - proportion of non-retail business acres per town.
- **chas** - Charles River dummy variable (= 1 if tract bounds river; 0 otherwise).
- **nox** - nitrogen oxides concentration (parts per 10 million).
- **rm** - average number of rooms per dwelling.
- **age** - proportion of owner-occupied units built prior to 1940.
- **dis** - weighted mean of distances to five Boston employment centres.
- **rad** - index of accessibility to radial highways.
- **tax** - full-value property-tax rate per \$10,000.
- **prratio** - pupil-teacher ratio by town.

- **black** - $1000(\text{Bk} - 0.63)^2$ where Bk is the proportion of blacks by town.
- **lstat** - lower status of the population (percent).
- **medv** - median value of owner-occupied homes in \$1000s.

```
base_dir <- here::here("")

calc_rmse = function(actual, predicted) {
  sqrt(mean((actual - predicted) ^ 2))
}

str(Boston)
#> 'data.frame':  506 obs. of  14 variables:
#> $ crim   : num  0.00632 0.02731 0.02729 0.03237 0.06905 ...
#> $ zn     : num  18 0 0 0 0 12.5 12.5 12.5 12.5 ...
#> $ indus  : num  2.31 7.07 7.07 2.18 2.18 2.18 7.87 7.87 7.87 7.87 ...
#> $ chas   : int   0 0 0 0 0 0 0 0 0 0 ...
#> $ nox    : num  0.538 0.469 0.469 0.458 0.458 0.458 0.524 0.524 0.524 0.524 ...
#> $ rm     : num  6.58 6.42 7.18 7 7.15 ...
#> $ age    : num  65.2 78.9 61.1 45.8 54.2 58.7 66.6 96.1 100 85.9 ...
#> $ dis    : num  4.09 4.97 4.97 6.06 6.06 ...
#> $ rad    : int   1 2 2 3 3 3 5 5 5 5 ...
#> $ tax    : num  296 242 242 222 222 222 311 311 311 311 ...
#> $ ptratio: num  15.3 17.8 17.8 18.7 18.7 18.7 15.2 15.2 15.2 15.2 ...
#> $ black  : num  397 397 393 395 397 ...
#> $ lstat  : num  4.98 9.14 4.03 2.94 5.33 ...
#> $ medv   : num  24 21.6 34.7 33.4 36.2 28.7 22.9 27.1 16.5 18.9 ...
```

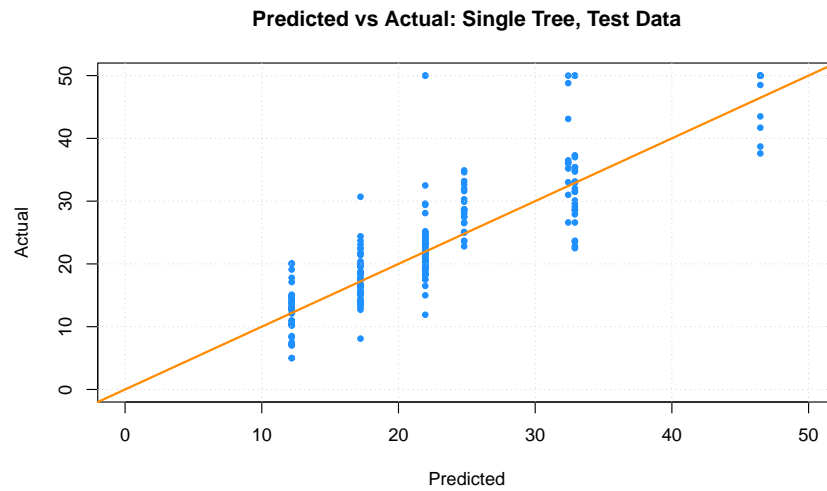
Test-Train Split the Data

```
set.seed(18)
boston_idx <- sample(1:nrow(Boston), nrow(Boston) / 2)
boston_trn <- Boston[boston_idx,]
boston_tst <- Boston[-boston_idx,]
```

Initial Tree Model of Median Value versus all other Predictors

```
boston_tree <- rpart(medv ~ ., data = boston_trn)

boston_tree_tst_pred <- predict(boston_tree, newdata = boston_tst)
plot(boston_tree_tst_pred, boston_tst$medv,
     xlim = c(0,50), ylim = c(0,50),
     xlab = "Predicted", ylab = "Actual",
     main = "Predicted vs Actual: Single Tree, Test Data",
     col = "dodgerblue", pch = 20)
grid()
abline(0, 1, col = "darkorange", lwd = 2)
```

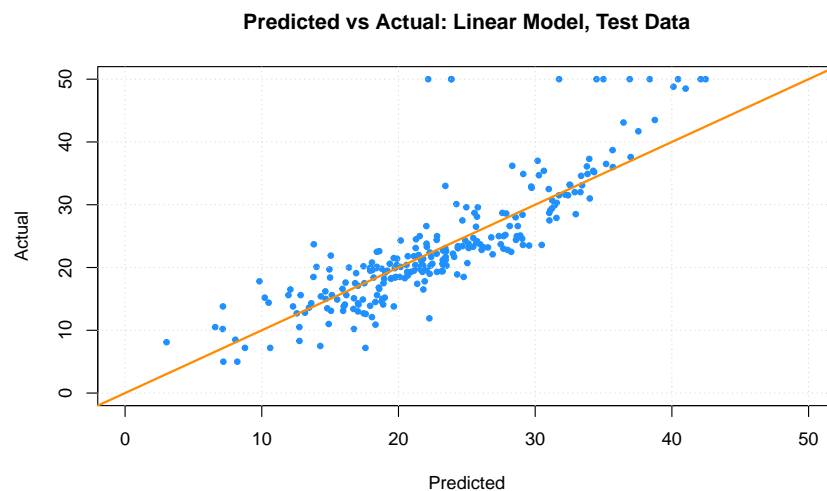


```
# RMSE
(tree_tst_rmse <- calc_rmse(boston_tree_tst_pred, boston_tst$medv))
#> [1] 5.051138
```

Initial Linear Model of Median Value versus all other Predictors

```
boston_lm <- lm(medv ~ ., data = boston_trn)

boston_lm_tst_pred <- predict(boston_lm, newdata = boston_tst)
plot(boston_lm_tst_pred, boston_tst$medv,
     xlim = c(0,50), ylim = c(0,50),
     xlab = "Predicted", ylab = "Actual",
     main = "Predicted vs Actual: Linear Model, Test Data",
     col = "dodgerblue", pch = 20)
grid()
abline(0, 1, col = "darkorange", lwd = 2)
```



```
# RMSE
(lm_tst_rmse <- calc_rmse(boston_lm_tst_pred, boston_tst$medv))
#> [1] 5.016083
```

Bagging

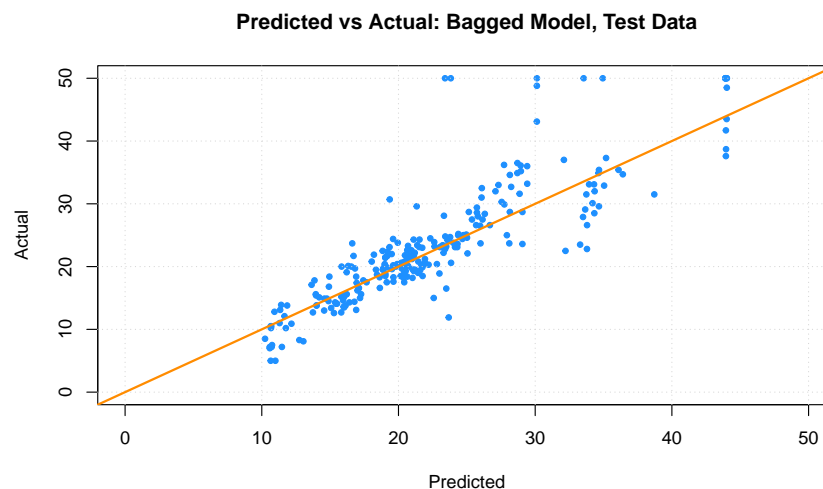
(The reference book used `randomForest` but this is not yet available for M1 machines. I am trying the `cforest` function from the `party` package as a replacement.)

Bagging is actually a special case of a random forest where `mtry` is equal to the number of predictors - 13 in this case.

```
boston_bag <- cforest(medv ~ ., data = boston_trn, controls = cforest_unbiased(mtry = 13, ntree = 500))

boston_bag
#>
#> Random Forest using Conditional Inference Trees
#>
#> Number of trees: 500
#>
#> Response: medv
#> Inputs: crim, zn, indus, chas, nox, rm, age, dis, rad, tax, ptratio, black, lstat
#> Number of observations: 253

boston_bag_tst_pred <- predict(boston_bag, newdata = boston_tst)
plot(boston_bag_tst_pred, boston_tst$medv,
     xlim = c(0, 50), ylim = c(0, 50),
     xlab = "Predicted", ylab = "Actual",
     main = "Predicted vs Actual: Bagged Model, Test Data",
     col = "dodgerblue", pch = 20)
grid()
abline(0, 1, col = "darkorange", lwd = 2)
```



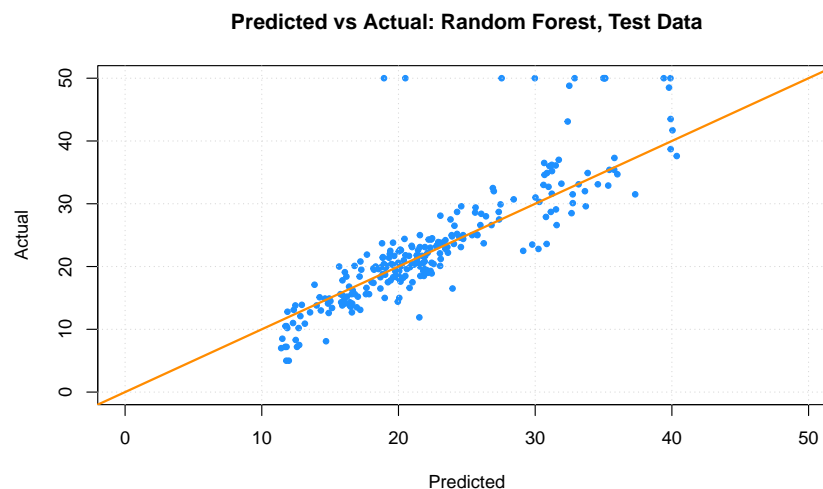
```
(bag_tst_rmse <- calc_rmse(boston_bag_tst_pred, boston_tst$medv))  
#> [1] 4.710894
```

Here we see two interesting results. First, the predicted versus actual plot no longer has a small number of predicted values. Second, our test error has dropped dramatically - note, not as dramatically as the original example did using `randomForest`.

Random Forest

We now try a random forest. For regression, the suggestion for `mtry` is number of predictors divided by 3 - 4 in this case.

```
boston_forest <- cforest(medv ~ ., data = boston_trn, controls = cforest_unbiased(mtry = 4, ntree = 500))  
  
boston_forest  
#>  
#> Random Forest using Conditional Inference Trees  
#>  
#> Number of trees: 500  
#>  
#> Response: medv  
#> Inputs: crim, zn, indus, chas, nox, rm, age, dis, rad, tax, ptratio, black, lstat  
#> Number of observations: 253  
  
boston_forest_tst_pred <- predict(boston_forest, newdata = boston_tst)  
plot(boston_forest_tst_pred, boston_tst$medv,  
     xlim = c(0,50), ylim = c(0,50),  
     xlab = "Predicted", ylab = "Actual",  
     main = "Predicted vs Actual: Random Forest, Test Data",  
     col = "dodgerblue", pch = 20)  
grid()  
abline(0, 1, col = "darkorange", lwd = 2)
```



```
(forest_tst_rmse <- calc_rmse(boston_forest_tst_pred, boston_tst$medv))
#> [1] 5.042991
```

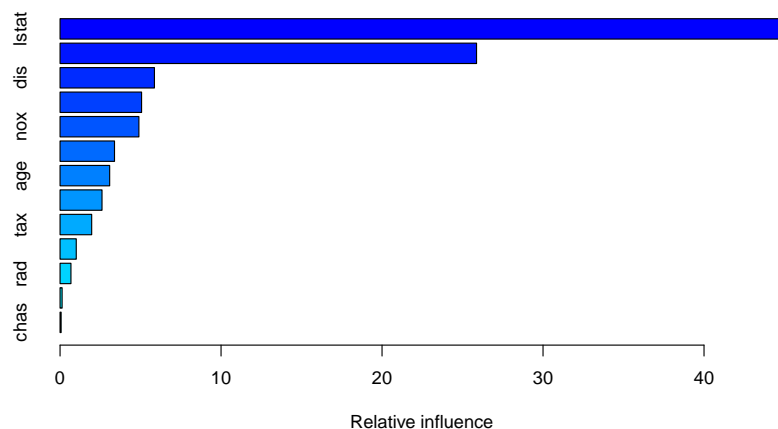
Boosting

```
booston_boost <- gbm(medv ~ ., data = boston_trn, distribution = "gaussian",
  n.trees = 5000, interaction.depth = 4, shrinkage = 0.01)
```

```
booston_boost
```

```
#> gbm(formula = medv ~ ., distribution = "gaussian", data = boston_trn,
#>      n.trees = 5000, interaction.depth = 4, shrinkage = 0.01)
#> A gradient boosted model with gaussian loss function.
#> 5000 iterations were performed.
#> There were 13 predictors of which 13 had non-zero influence.
```

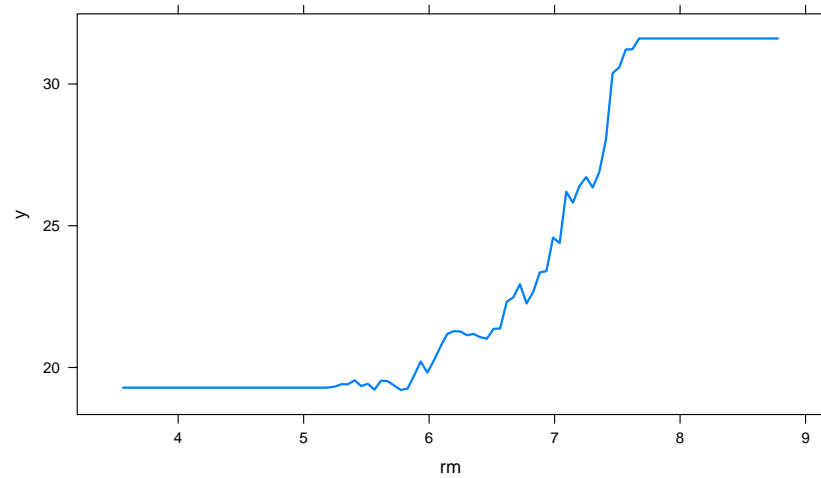
```
tibble::as_tibble(summary(booston_boost))
```



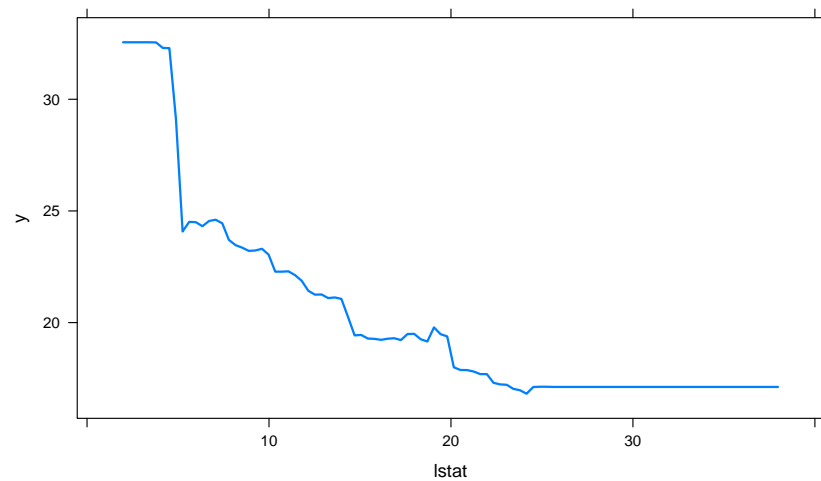
```
#> # A tibble: 13 x 2
#>   var      rel.inf
#>   <chr>      <dbl>
#> 1 lstat    45.4
#> 2 rm       25.9
#> 3 dis       5.86
#> 4 crim      5.06
#> 5 nox       4.90
#> 6 black     3.38
#> 7 age       3.08
#> 8 ptratio   2.61
#> 9 tax       1.96
#> 10 indus    1.01
#> 11 rad      0.676
#> 12 zn       0.127
```

```
#> 13 chas      0.0659
```

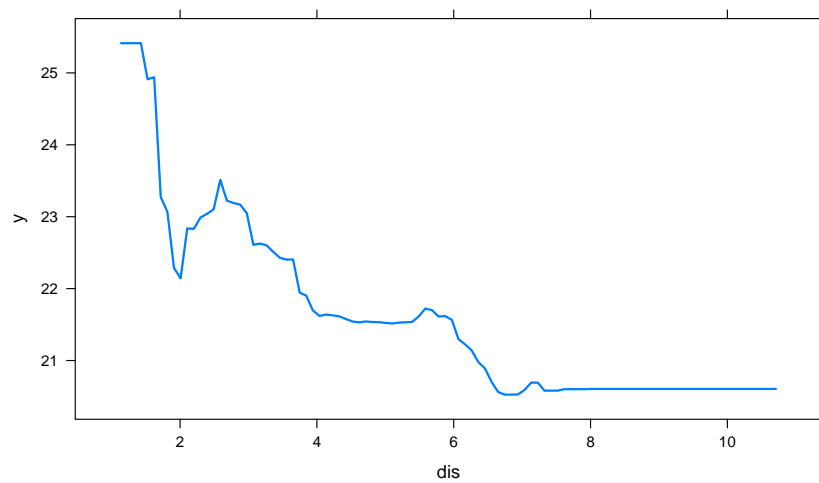
```
par(mfrow = c(1, 3))  
plot(booston_boost, i = "rm", col = "dodgerblue", lwd = 2)
```



```
plot(booston_boost, i = "lstat", col = "dodgerblue", lwd = 2)
```

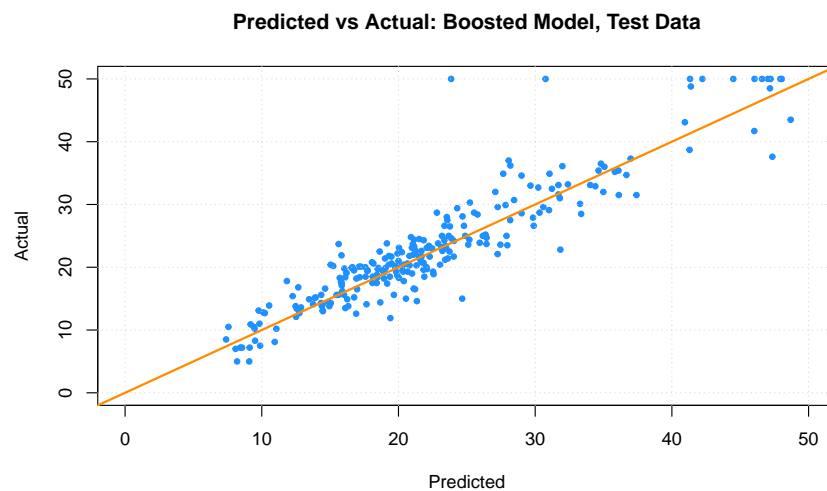


```
plot(booston_boost, i = "dis", col = "dodgerblue", lwd = 2)
```



```
boston_boost_tst_pred = predict(booston_boost, newdata = boston_tst, n.trees = 5000)
(boost_tst_rmse <- calc_rmse(boston_boost_tst_pred, boston_tst$medv))
#> [1] 3.643578
```

```
plot(boston_boost_tst_pred, boston_tst$medv,
     xlim = c(0,50), ylim = c(0,50),
     xlab = "Predicted", ylab = "Actual",
     main = "Predicted vs Actual: Boosted Model, Test Data",
     col = "dodgerblue", pch = 20)
grid()
abline(0, 1, col = "darkorange", lwd = 2)
```



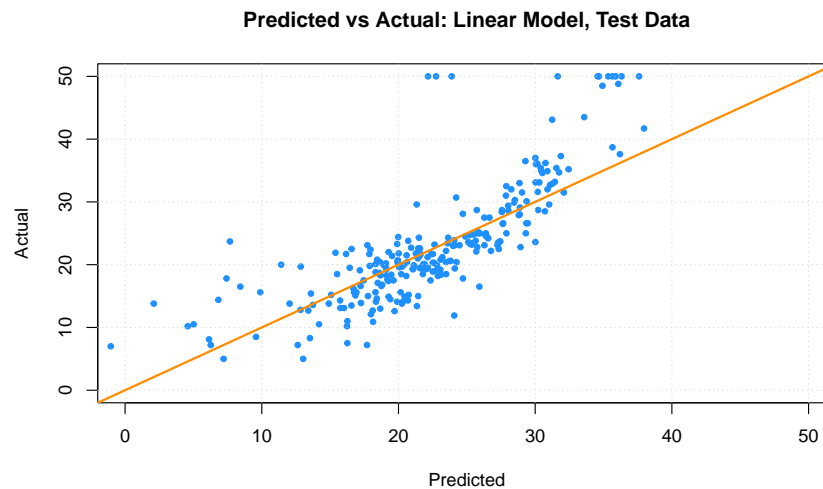
Results

Model	TestError
Single Tree	5.051138
Linear Model	5.016083
Bagging	4.710894
Random Forest	5.042992
Boosting	3.643578

Linear Model of Median Value versus two most significant predictors from boosting (lstat and rm)

```
boston_lm2 <- lm(medv ~ lstat + rm, data = boston_trn)

boston_lm2_tst_pred <- predict(boston_lm2, newdata = boston_tst)
plot(boston_lm2_tst_pred, boston_tst$medv,
     xlim = c(0,50), ylim = c(0,50),
     xlab = "Predicted", ylab = "Actual",
     main = "Predicted vs Actual: Linear Model, Test Data",
     col = "dodgerblue", pch = 20)
grid()
abline(0, 1, col = "darkorange", lwd = 2)
```



```
# RMSE
(lm2_tst_rmse <- calc_rmse(boston_lm_tst_pred, boston_tst$medv))
#> [1] 5.016083
```