STAT406- Methods of Statistical Learning Lecture 11

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Overfitting...

Boston Example

Not surprisingly, when we overfit...

Pruning...

Cost pruning

$$\min_{T \subset T_0} \sum_{m=1}^{|T|} \sum_{\mathbf{x}_i \in R_m} (y_i - \hat{\mu}_m)^2 + \alpha |T|$$

- ullet We can compute the solution for all lpha
- Compare each subtree in this sequence using CV
- Pick the best subtree

Pruning...

- More specifically:
- Let $T_{\ell} \subset T_0$ be the solution to

$$\min_{T \subset T_0} \sum_{m=1}^{|T|} \sum_{\mathbf{x}_i \in R_m} (y_i - \hat{\mu}_m)^2 + \alpha |T|$$

when

$$\alpha \in [\alpha_{\ell}, \alpha_{\ell+1}) \subseteq [0, +\infty) \quad \ell = 1, 2, \dots, L$$

Pruning...

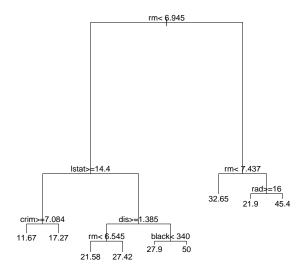
- Split the data into K folds
- For j = 1, ..., K
 - Build a tree without using the j-th fold
 - Prune it with penalties α_{ℓ} , $\ell=1,\ldots,L$
 - Use these *L* trees to predict the *j*-th fold
 - Record the prediction errors.
- Sum or average over the folds.
- We obtain K-fold CV-estimated prediction errors for the L trees corresponding to pruning with α_{ℓ} , $\ell = 1, \ldots, L$.

Boston Example

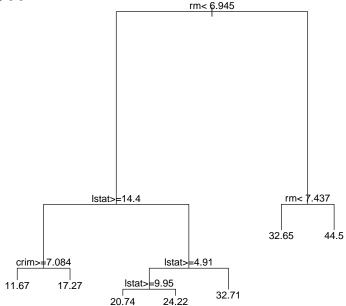
Pruning works...

```
> b <- ***cp with minimum xerror***
> bos.t3 <- prune(bos.to, cp=b)</pre>
> plot(bos.t3)
> pr.t3 <- predict(bos.t3,</pre>
        newdata=dat.te,
        type='vector')
> with(dat.te, mean((medv - pr.t3)^2))
[11 18.96988
```

Pruned tree

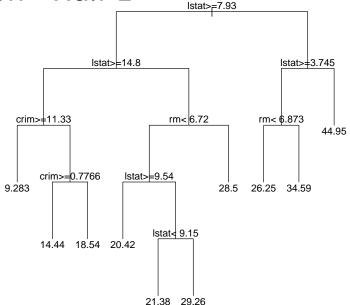


Boston

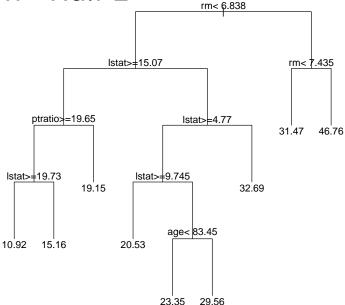


- Trees can be highly variable
- Trees computed on samples from the sample population can be quite different from each other
- For example, we split the Boston data in two...

Boston - Half 1



Boston - Half 2



- Linear regression, for example, is not so variable
- Estimated coefficients computed on the same two halfs

```
(Intercept) crim zn indus chas
[1,] 39.21 -0.13 0.04 0.04 2.72
[2,] 33.12 -0.10 0.05 -0.01 2.80

nox rm age dis rad tax
[1,] -20.07 3.45 0 -1.44 0.28 -0.01
[2,] -14.18 4.15 0 -1.46 0.34 -0.02

ptratio black lstat
[1,] -1.01 0.01 -0.56
[2,] -0.90 0.01 -0.50
```

- If we could average many trees trained on independent samples from the same population, we would obtain a predictor with lower variance
- If \hat{f}_1 , \hat{f}_2 , ..., \hat{f}_B are B regression trees, then their average is

$$\hat{f}_{av}(\mathbf{x}) = \frac{1}{B} \sum_{i=1}^{B} \hat{f}_{i}(\mathbf{x})$$

- However, we generally do not have B training sets...
- We can **bootstrap** the training set to obtain B pseudo-new-training sets
- Let (Y_1, \mathbf{X}_1) , (Y_2, \mathbf{X}_2) , ..., (Y_n, \mathbf{X}_n) be the training sample, where

$$(Y_j, \mathbf{X}_j) \sim F_0$$

- If we knew F₀, then we could generate / simulate new training sets, and average the resulting trees...
- We do not know F₀, but we have an estimate for it
- Let F_n be the empirical distribution of our only training set (Y_1, \mathbf{X}_1) , (Y_2, \mathbf{X}_2) , ..., (Y_n, \mathbf{X}_n)

We know that

$$F_n \xrightarrow[n\to\infty]{} F_0$$

(in what sense?)

- Bootstrap generates / simulates samples from F_n
- Taking a sample of size n from F_n is the same as sampling with replacement from the training set $(Y_1, \mathbf{X}_1), (Y_2, \mathbf{X}_2), \ldots, (Y_n, \mathbf{X}_n)$

- To apply bagging to a regression tree, take B independent samples (with replacement) from the training set
- Obtain the B trees: \hat{f}_1^* , \hat{f}_2^* , ..., \hat{f}_B^*
- and average their predictions

$$\hat{f}_{\mathsf{bag}}(\mathbf{x}) = \frac{1}{B} \sum_{j=1}^{B} \hat{f}_{j}^{*}(\mathbf{x})$$

- Generally, we apply bagging on "large" trees, without pruning them (try to retain their low-bias and reduce their variance by averaging)
- With the Boston data set, if we apply bagging to the regression tree computed on the training set, and then use it to predict on the test set, we obtain:

• B = 1> mean((dat.te\$medv - pr.ba)^2) [1] 16.44972 • B = 5> mean((dat.te\$medv - pr.ba)^2) [1] 15.12332 • B = 100> mean((dat.te\$medv - pr.ba)^2) [1] 12.30543 • B = 500> mean((dat.te\$medv - pr.ba)^2) [1] 12.32504

B = 2000
mean((dat.te\$medv - pr.ba)^2)
[1] 11.8116
B = 5000
mean((dat.te\$medv - pr.ba)^2)
[1] 11.85943

- This approach applies to any predictor (not only trees)
- It will be particularly useful for low-bias / high-variance predictors