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

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

when

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

Pruning...

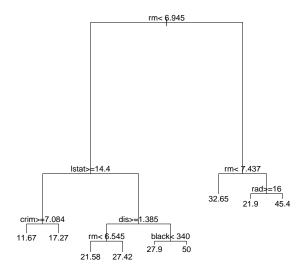
- Split the data into K folds
- For j = 1, ..., K
 - Build a tree without using the j-th fold
 - Prune it with penalties α_{ℓ} , $\ell=1,\ldots,L$
 - Use these *L* trees to predict the *j*-th fold
 - Record the prediction errors.
- Sum or average over the folds.
- We obtain K-fold CV-estimated prediction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

Boston Example

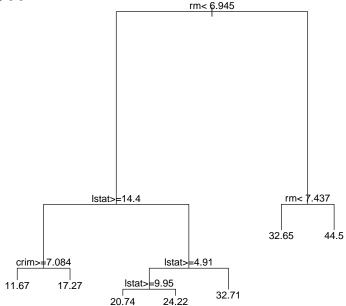
Pruning works...

```
> 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



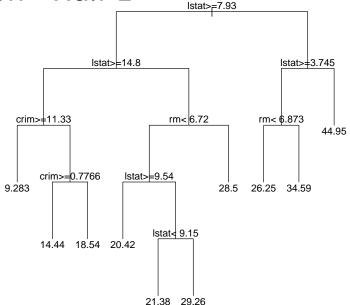
Boston



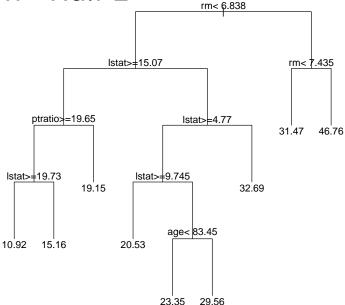
Bagging

- 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



Bagging

- 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
```

Bagging

- 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

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