#### **Decision Trees**

#### Tal Linzen Slides based on Lecture 10 from David Rosenberg's course materials (https://github.com/davidrosenberg/mlcourse)

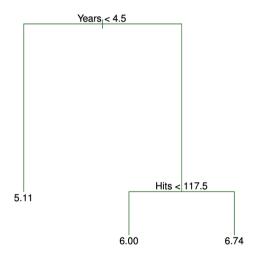
CDS, NYU

April 5, 2022

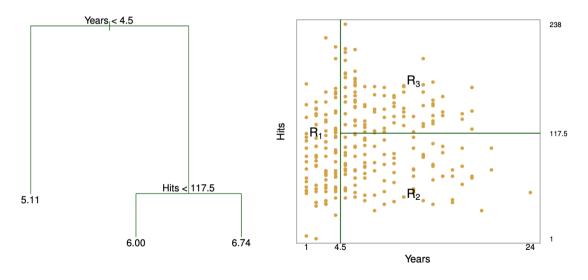
- Our first inherently non-linear classifier: decision trees.
- Ensemble methods: bagging and boosting.

### **Decision Trees**

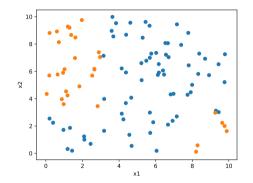
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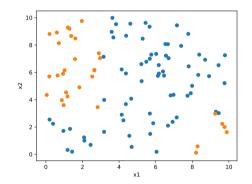


### Classification trees

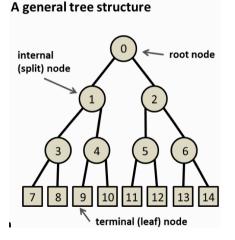


• Can we classify these points using a linear classifier?

# Classification trees

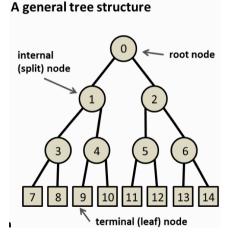


- Can we classify these points using a linear classifier?
- Partition the data into axis-aligned regions recursively (on the board)



• We focus on *binary* trees (as opposed to multiway trees where nodes can have more than two children)

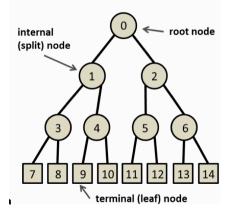
From Criminisi et al. MSR-TR-2011-114, 28 October 2011.



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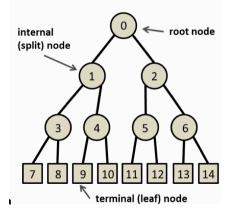




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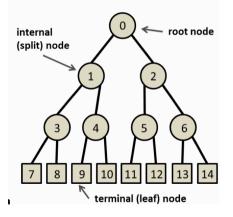




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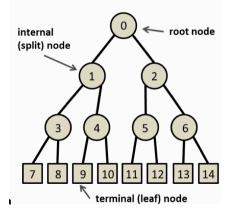




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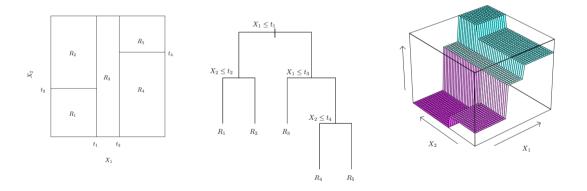
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- A greedy algorithm is the one that make the best **local** decisions, without lookahead to evaluate their downstream consequences
- This procedure is not very likely to result in the globally optimal tree

### Prediction in a Regression Tree



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- It is common to split half way between two adjacent values:

$$s_j \in \left\{ \frac{1}{2} \left( x_{j(r)} + x_{j(r+1)} \right) \mid r = 1, \dots, n-1 \right\}.$$
  $n-1$  splits (1)

# Decision Trees and Overfitting

• What will happen if we keep splitting the data into more and more regions?

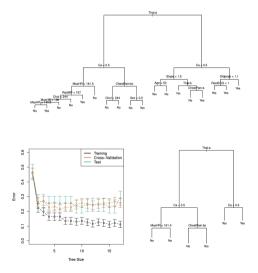
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- What will happen if we keep splitting the data into more and more regions?
  - Every data point will be in its own region-overfitting.
- When should we stop splitting? (Controlling the complexity of the hypothesis space)
  - Limit total number of nodes.
  - Limit number of terminal nodes.
  - Limit tree depth.
  - Require minimum number of data points in a terminal node.
  - Backward pruning (the approach used in CART; Breiman et al 1984):
    - **(**) Build a really big tree (e.g. until all regions have  $\leq 5$  points).
    - Prune the tree back greedily, potentially all the way to the root, until validation performance starts decreasing.

# Pruning: Example



# What Makes a Good Split for Classification?

Our plan is to predict the majority label in each region.

Which of the following splits is better?

Split 1 $R_1: 8 + /2 R_2: 2 + /8 -$ Split 2 $R_1: 6 + /4 R_2: 4 + /6 -$ 

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Intuition: we want to produce *pure* nodes, i.e. nodes where most instances have the same class.

#### Misclassification error in a node

• Let's consider the multiclass classification case:  $\mathcal{Y} = \{1, 2, \dots, K\}$ .

- Let node m represent region  $R_m$ , with  $N_m$  observations
- We denote the proportion of observations in  $R_m$  with class k by

$$\hat{p}_{mk} = \frac{1}{N_m} \sum_{\{i: x_i \in R_m\}} 1(y_i = k).$$

• We predict the majority class in node *m*:

$$k(m) = \underset{k}{\arg\max} \hat{p}_{mk}$$

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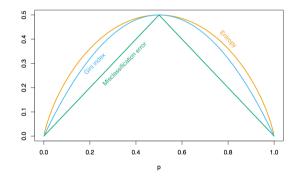
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• The Gini index and entropy are numerically similar to each other, and both work better in practice than the misclassification error.

Tal Linzen Slides based on Lecture 10 from David F

#### Impurity Measures for Binary Classification

(p is the relative frequency of class 1)



#### HTF Figure 9.3

Tal Linzen Slides based on Lecture 10 from David F

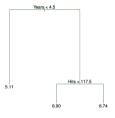
# Quantifying the Impurity of a Split

Scoring a potential split that produces the nodes  $R_L$  and  $R_R$ :

- Suppose we have  $N_L$  points in  $R_L$  and  $N_R$  points in  $R_R$ .
- Let  $Q(R_L)$  and  $Q(R_R)$  be the node impurity measures for each node.
- We aim to find a split that minimizes the weighted average of node impurities:

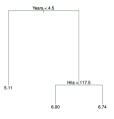
 $\frac{N_L Q(R_L) + N_R Q(R_R)}{N_L + N_R}$ 

#### Discussion: Interpretability of Decision Trees



• Trees are easier to visualize and explain than other classifiers (even linear regression)

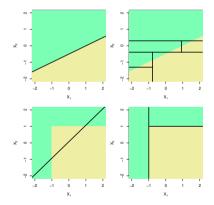
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- Trees are easier to visualize and explain than other classifiers (even linear regression)
- Small trees are interpretable large trees, maybe not so much

#### Discussion: Trees vs. Linear Models

Trees may have to work hard to capture linear decision boundaries, but can easily capture certain nonlinear ones:



#### Discussion: Review

Decision trees are:

- Non-linear: the decision boundary that results from splitting may end up being quite complicated
- Non-metric: they do not rely on the geometry of the space (inner products or distances)
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Additional pros:

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Cons:

- Struggle to capture linear decision boundaries
- They have high variance and tend to overfit: they are sensitive to small changes in the training data (The ensemble techniques we discuss next can mitigate these issues)

# Bagging and Random Forests

• We observe data  $\mathcal{D} = (x_1, x_2, \dots, x_n)$  sampled i.i.d. from a parametric distribution  $p(\cdot | \theta)$ 

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- A statistic  $\hat{\theta} = \hat{\theta}(\mathcal{D})$  is a point estimator of  $\theta$  if  $\hat{\theta} \approx \theta$

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- Statistics are random, so they have probability distributions.
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$$\begin{split} \text{Bias } \mathsf{Bias}(\hat{\theta}) \stackrel{\text{def}}{=} \mathbb{E}\left[\hat{\theta}\right] - \theta.\\ \text{Variance } \mathsf{Var}(\hat{\theta}) \stackrel{\text{def}}{=} \mathbb{E}\left[\hat{\theta}^2\right] - \mathbb{E}^2\left[\hat{\theta}\right]. \end{split}$$

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Variance  $\operatorname{Var}(\hat{\theta}) \stackrel{\text{def}}{=} \mathbb{E}\left[\hat{\theta}^{2}\right] - \mathbb{E}^{2}\left[\hat{\theta}\right].$ 

• Why does variance matter if an estimator is unbiased?

- Let  $\hat{\theta}(\mathcal{D})$  be an unbiased estimator with variance  $\sigma^2$ :  $\mathbb{E}\left[\hat{\theta}\right] = \theta$ ,  $Var(\hat{\theta}) = \sigma^2$ .
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- The average has the same expected value but smaller standard error (recall that  $Var(cX) = c^2 Var(X)$ , and that the  $\hat{\theta}_i$ -s are uncorrelated):

$$\mathbb{E}\left[\frac{1}{n}\sum_{i=1}^{n}\hat{\theta}_{i}\right] = \theta \qquad \text{Var}\left[\frac{1}{n}\sum_{i=1}^{n}\hat{\theta}_{i}\right] = \frac{\sigma^{2}}{n}$$
(2)

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- Our learning algorithm gives us *B* prediction functions:  $\hat{f}_1(x), \hat{f}_2(x), \dots, \hat{f}_B(x)$
- We will define the average prediction function as:

$$\hat{f}_{\mathsf{avg}} \stackrel{\text{def}}{=} rac{1}{B} \sum_{b=1}^{B} \hat{f}_{b}$$

(3)

#### Averaging Reduces Variance of Predictions

• The average prediction for  $x_0$  is

$$\hat{f}_{avg}(x_0) = \frac{1}{B} \sum_{b=1}^{B} \hat{f}_b(x_0).$$

- $\hat{f}_{avg}(x_0)$  and  $\hat{f}_b(x_0)$  have the same expected value, but
- $\hat{f}_{avg}(x_0)$  has smaller variance:

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• Problem: in practice we don't have B independent training sets!

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• For large n,

$$\left(1-\frac{1}{n}\right)^n \approx \frac{1}{e} \approx .368. \tag{4}$$

• So we expect ~63.2% of elements of  $\mathcal{D}_n$  will show up at least once.

### The Bootstrap Method

#### Definition

A **bootstrap method** simulates *B* independent samples from *P* by taking *B* bootstrap samples from the sample  $\mathcal{D}_n$ .

- Given original data  $\mathcal{D}_n$ , compute *B* bootstrap samples  $D_n^1, \ldots, D_n^B$ .
- For each bootstrap sample, compute some function

 $\phi(D_n^1),\ldots,\phi(D_n^B)$ 

- Use these values as though  $D_n^1, \ldots, D_n^B$  were i.i.d. samples from P.
- This often ends up being very close to what we'd get with independent samples from P!

## Independent Samples vs. Bootstrap Samples

- Point estimator  $\hat{\alpha} = \hat{\alpha}(\mathcal{D}_{100})$  for samples of size 100, for a synthetic case where the data generating distribution is known
- Histograms of  $\hat{\alpha}$  based on
  - 1000 independent samples of size 100 (left), vs.
  - 1000 bootstrap samples of size 100 (right)

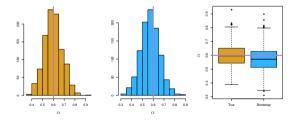


Figure 5.10 from ISLR (Springer, 2013) with permission from the authors: G. James, D. Witten, T. Hastie and R. Tibshirani.

#### Ensemble Methods

Key ideas:

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- Sequential ensemble (e.g., boosting): models are built sequentially
  - We try to find new learners that do well where previous learners fall short

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- The **bagged prediction function** is a *combination* of these:

$$\hat{f}_{avg}(x) = \text{Combine}\left(\hat{f}_1(x), \hat{f}_2(x), \dots, \hat{f}_B(x)\right)$$

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- For classification, averaging doesn't make sense; we can take a majority vote instead
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- Is there a downside, compared to having a single decision tree?
- Yes: if we have many trees, the bagged predictor is much less interpretable

### Aside: Out-of-Bag Error Estimation

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- The remaining 37% are called **out-of-bag (OOB)** observations.
- For *i*th training point, let

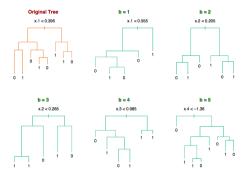
 $S_i = \{ b \mid D^b \text{ does not contain } i \text{th point} \}$ 

• The **OOB prediction** on x<sub>i</sub> is

$$\hat{f}_{\text{OOB}}(x_i) = \frac{1}{|S_i|} \sum_{b \in S_i} \hat{f}_b(x_i)$$

- The OOB error is a good estimate of the test error
- Similar to cross validation error: both are computed on the training set

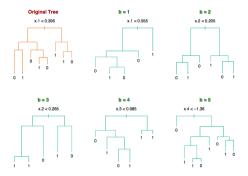
• Input space  $\mathfrak{X} = \mathbb{R}^5$  and output space  $\mathfrak{Y} = \{-1, 1\}$ . Sample size n = 30.



From HTF Figure 8.9

Tal Linzen Slides based on Lecture 10 from David F

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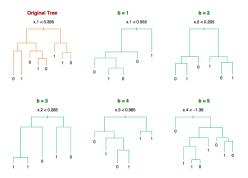


• Each bootstrap tree is quite different: different splitting variable at the root!

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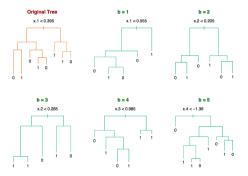


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- Each bootstrap tree is quite different: different splitting variable at the root!
- **High variance**: small perturbations of the training data lead to a high degree of model variability
- Bagging helps most when the base learners are relatively unbiased but have high variance (exactly the case for decision trees)

From HTF Figure 8.9

Tal Linzen Slides based on Lecture 10 from David I

• For 
$$\hat{\theta}_1, \dots, \hat{\theta}_n$$
 *i.i.d.* with  $\mathbb{E}\left[\hat{\theta}\right] = \theta$  and  $\operatorname{Var}\left[\hat{\theta}\right] = \sigma^2$ ,  
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Recall the motivating principle of bagging:

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- Can we reduce the dependence between  $\hat{f}_i$ 's?

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Use bagged decision trees, but modify the tree-growing procedure to reduce the dependence between trees.

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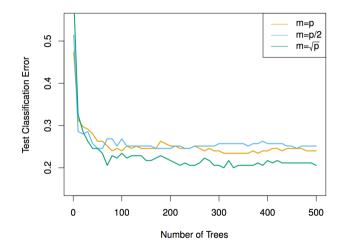
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- If m = p, this is just bagging

#### Random Forests: Effect of m



From An Introduction to Statistical Learning, with applications in R (Springer, 2013) with permission from the authors: G. James, D. Witten, T. Hastie and R. Tibshirani.

Tal Linzen Slides based on Lecture 10 from David F



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- But bootstrap samples (and the induced models) are correlated
- Ensembling works better when we combine a diverse set of prediction functions
  - ullet  $\Longrightarrow$  Random forests: select a random subset of features for each decision tree

# Boosting

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- Boosting Reduce the error rate of a high bias estimator by ensembling many estimators trained in sequence (without bootstrapping).
  - Like bagging, boosting is a general method that is particularly popular with decision trees.
  - Main intuition: instead of fitting the data very closely using a large decision tree, train gradually, using a sequence of simpler trees

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- We'll focus on a specific implementation, AdaBoost (Freund & Schapire, 1997)

### AdaBoost: Setting

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- Typical base hypothesis spaces:
  - Decision stumps (tree with a single split)
  - Trees with few terminal nodes
  - Linear decision functions

# Weighted Training Set

Each base learner is trained on weighted data.

- Training set  $\mathcal{D} = ((x_1, y_1), \dots, (x_n, y_n)).$
- Weights  $(w_1, \ldots, w_n)$  associated with each example.

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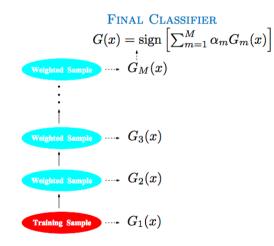
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- Weighted empirical risk:

$$\hat{R}_n^w(f) \stackrel{\text{def}}{=} rac{1}{W} \sum_{i=1}^n w_i \ell(f(x_i), y_i) \quad ext{where } W = \sum_{i=1}^n w_i$$

• Examples with larger weights affect the loss more.

#### AdaBoost: Schematic



From ESL Figure 10.1

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### AdaBoost: Classifier Weights

• Our final prediction is 
$$G(x) = \operatorname{sign}\left[\sum_{m=1}^{M} \alpha_m G_m(x)\right]$$
.

- We would like  $\alpha_m$  to be:
  - Nonnegative
  - Larger when  $G_m$  fits its weighted training data well
- The weighted 0-1 error of  $G_m(x)$  is

$$\operatorname{err}_{m} = \frac{1}{W} \sum_{i=1}^{n} w_{i} \mathbb{1}(y_{i} \neq G_{m}(x_{i})) \quad \text{where } W = \sum_{i=1}^{n} w_{i}.$$

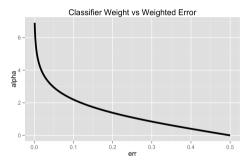
● err<sub>m</sub> ∈ [0, 1]

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• Higher weighted error  $\implies$  lower weight

• We train  $G_m$  to minimize weighted error; the resulting error rate is  $err_m$ 

• Then 
$$\alpha_m = \ln\left(\frac{1 - err_m}{err_m}\right)$$
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=  $w_i \left( \frac{1 - \operatorname{err}_m}{\operatorname{err}_m} \right)$ 

• If  $G_m$  is a strong classifier overall, then its  $\alpha_m$  will be large; this means that if  $x_i$  is misclassified,  $w_i$  will increase to a greater extent

## AdaBoost with Decision Stumps

• After 1 round:

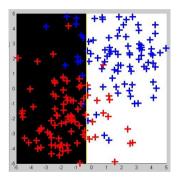


Figure: Size of plus sign represents weight of example. Blackness represents preference for red class; whiteness represents preference for blue class.

KPM Figure 16.10

## AdaBoost with Decision Stumps

• After 3 rounds:

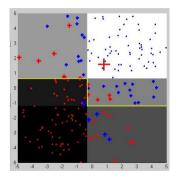


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KPM Figure 16.10

## AdaBoost with Decision Stumps

• After 120 rounds:

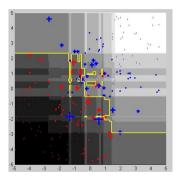
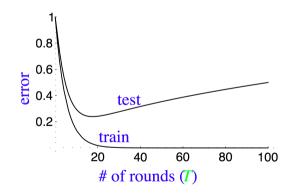


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KPM Figure 16.10

### Does AdaBoost overfit?

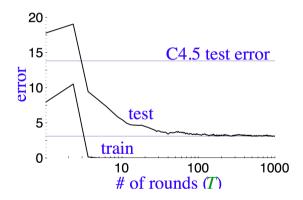
- Does a large number of rounds of boosting lead to overfitting?
- If we were overfitting, the learning curves would look like:



From Rob Schapire's NIPS 2007 Boosting tutorial.

#### Learning Curves for AdaBoost

- AdaBoost is usually quite resistant to overfitting
- The test error continues to decrease even after the training error drops to zero!



From Rob Schapire's NIPS 2007 Boosting tutorial.

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