# **Decision trees**

## Fraida Fund

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## Math prerequisites for this lecture: None

## In this lecture

- · Decision trees
- · Training decision trees
- · Bias and variance of decision trees

#### Recap

## Flexible decisions with cheap prediction?

KNN was very flexible, but prediction is **slow**.

Next: flexible decisions, non-parametric approach, fast prediction

**Idea**: In KNN, we find the "neighborhood" of a test point and then give it the value of training points in that "neighborhood" - but it takes too long at inference time to define the "neighborhood".

What if we define "neighborhoods" and their values in advance, at training time? Then at inference time, we only need to determine which "neighborhood" a test point belongs in.

However, we run into another **computationally hard** problem! To *partition* the feature space into optimal neighborhoods is too expensive. Instead, we will rely on some heuristics and get a non-optimal, but good enough, partition.

## **Decision tree**

## **Tree terminology**

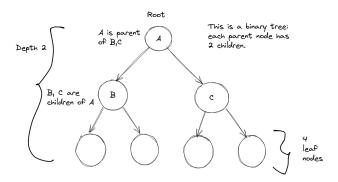


Figure 1: A binary tree.

- size of tree |T| (number of leaf nodes)
- depth (max length from root node to a leaf node)

#### **Note on notation**

Following notation of ISLR, Chapter 8:

- $X_i$  is feature j
- $x_i$  is sample i

## **Stratification of feature space (1)**

- Given set of possible predictors,  $X_1,\dots,X_p$
- ullet Training: Divide predictor space (set of possible values of X) into J non-overlapping regions:  $R_1,\dots,R_J$  , by splitting sequentially on one feature at a time.

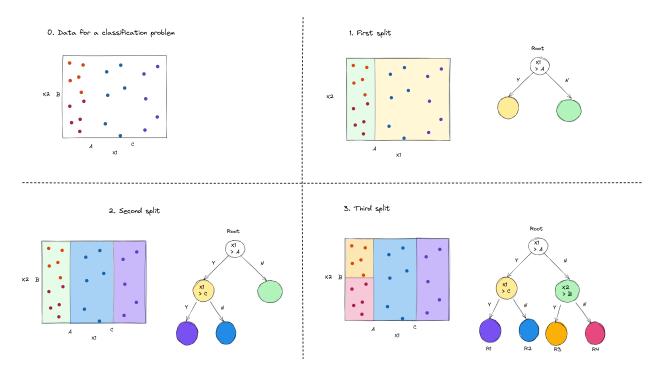


Figure 2: Dividing the feature space with a decision tree.

## **Stratification of feature space (2)**

- Prediction: For each observation that falls in region  $\boldsymbol{R}_{j}$  , predict

  - mean of labels of training points in  $R_j$  (regression) mode of labels of training points in  $R_j$  (classification)

## **Tree representation**

- At node that is not a leaf: test one feature  ${\cal X}_i$  Branch from node depending on value of  ${\cal X}_i$
- Each leaf node: predict  $\hat{y}_{R_m}$

## Stratification of feature space - illustration

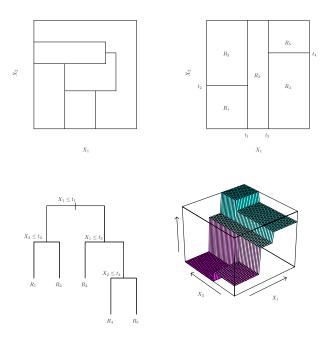


Figure 3: ISLR, Fig. 8.3.

The stratification on the top left cannot be produced by a decision tree using recursive binary splitting. The other three subfigures represent a single stratification. Note that the decision tree fits a piecewise step function!

## **Training a decision tree**

#### Basic idea (1)

- Goal: find the high-dimensional rectangles that minimize error
- Computationally expensive to consider every possible partition

### Basic idea (2)

- Instead: recursive binary splitting (top-down, greedy approach)
- Greedy: at each step, make the best decision at that step, without looking ahead and making a decision that might yield better results at future steps

## **Recursive binary splitting steps**

Start at root of the tree, considering all training samples.

- 1. At the current node,
- 2. Find feature  $\boldsymbol{X}_i$  and cutpoint  $\boldsymbol{s}$  that minimizes some loss function (?)
- 3. Split training samples at that node into two leaf nodes
- 4. Stop when no training error (?)
- 5. Otherwise, repeat at leaf nodes

At step 2, we apply a greedy heuristic - we are choosing the feature that minimizes a loss function in *this* iteration only.

## **Recursive binary splitting**

For any feature j and *cutpoint* s, define the regions

$$R_1(j,s) = \{X|X_j < s\}, \quad R_2(j,s) = \{X|X_j \geq s\}$$

where  $\{X|X_j < s\}$  is the region of predictor space in which  $X_j$  takes on a value less than s.

## Loss function for regression tree

For regression: look for feature j and cutpoint s that leads to the greatest possible reduction in squared error, where the "new" squared error is:

$$\sum_{i: x_i \in R_1(j,s)} (y_i - \hat{y}_{R_1})^2 \quad + \sum_{i: x_i \in R_2(j,s)} (y_i - \hat{y}_{R_2})^2$$

( $\hat{y}_{R_{i}}$  is the prediction for the samples in  $R_{j}$ .)

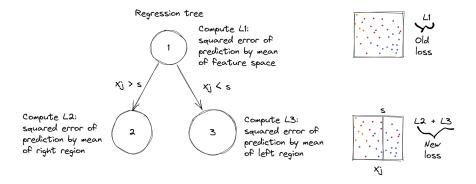


Figure 4: Training a regression tree.

#### Loss function for classification tree

For classification, find a split that minimizes some measure of node impurity:

- A node whose samples all belong to the same class most pure
- · A node whose samples are evenly distributed among all classes highly impure

#### **Classification error rate**

For classification: one possible way is to split on 0-1 loss or misclassification rate:

$$\sum_{x_i \in R_m} 1(y_i \neq \hat{y}_{R_m})$$

Not used often (this is a good explanation re: why), but used for pruning.

## **GINI** index

The GINI index is:

$$\sum_{k=1}^K \hat{p}_{mk} (1 - \hat{p}_{mk})$$

where  $\hat{p}_{mk}$  is the proportion of training samples in  $R_m$  belonging to class k.

You can see that this is small when all values of  $\hat{p}_{mk}$  are around 0 or 1.

## **Entropy**

Entropy as a measure of impurity on subset of samples:

$$-\sum_{k=1}^K \hat{p}_{mk} \log_2 \hat{p}_{mk}$$

where  $\hat{p}_{mk}$  is the proportion of training samples in  $R_m$  belonging to class k.

## **Comparison - measures of node impurity**

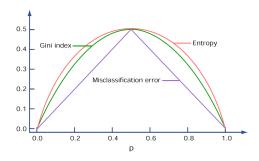


Figure 5: Measures of node "impurity".

#### **Conditional entropy**

- Splitting on feature X creates subsets  $S_1$  and  $S_2$  with different entropies
- · Conditional entropy:

$$\mathrm{Entropy}(S|X) = \sum_v \frac{|S_v|}{|S|} \mathrm{Entropy}(S_v)$$

## **Information gain**

• Choose feature to split so as to maximize information gain, the expected reduction in entropy due to splitting on X:

$$\mathrm{Gain}(S,X) := \mathrm{Entropy}(S) - \mathrm{Entropy}(S|X)$$

## Example: should I play tennis? (1)

| Day | Outlook  | Temperature | Humidity | Wind   | PlayTennis |
|-----|----------|-------------|----------|--------|------------|
| D1  | Sunny    | Hot         | High     | Weak   | No         |
| D2  | Sunny    | Hot         | High     | Strong | No         |
| D3  | Overcast | Hot         | High     | Weak   | Yes        |
| D4  | Rain     | Mild        | High     | Weak   | Yes        |
| D5  | Rain     | Cool        | Normal   | Weak   | Yes        |
| D6  | Rain     | Cool        | Normal   | Strong | No         |
| D7  | Overcast | Cool        | Normal   | Strong | Yes        |
| D8  | Sunny    | Mild        | High     | Weak   | No         |
| D9  | Sunny    | Cool        | Normal   | Weak   | Yes        |
| D10 | Rain     | Mild        | Normal   | Weak   | Yes        |
| D11 | Sunny    | Mild        | Normal   | Strong | Yes        |
| D12 | Overcast | Mild        | High     | Strong | Yes        |
| D13 | Overcast | Hot         | Normal   | Weak   | Yes        |
| D14 | Rain     | Mild        | High     | Strong | No         |

Figure 6: Via Tom Mitchell.

## Example: should I play tennis? (2)

For top node:  $S = \{9+, 5-\}, |S| = 14$ 

$$\mathrm{Entropy}(S) = -\frac{9}{14} \log_2 \frac{9}{14} - \frac{5}{14} \log_2 \frac{5}{14} = 0.94$$

## Example: should I play tennis? (3)

If we split on Wind:

Considering the Weak branch:

$$\begin{array}{l} \bullet \; S_{\rm weak} = \{6+,2-\}, |S_{\rm weak}| = 8 \\ \bullet \; {\rm Entropy}(S_{\rm weak}) = -\frac{6}{8} \log_2(\frac{6}{8}) - \frac{2}{8} \log_2(\frac{2}{8}) = 0.81 \end{array}$$

Considering the Strong branch:

- $$\begin{split} & \cdot \ S_{\rm strong} = \{3+,3-\}, |S_{\rm strong}| = 6 \\ & \cdot \ {\rm Entropy}(S_{\rm strong}) = 1 \end{split}$$

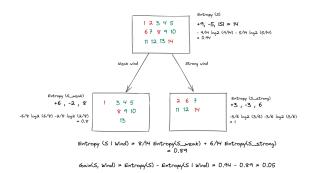


Figure 7: Considering the split on Wind.

## Example: should I play tennis? (4)

$$\mathrm{Entropy}(S) = -\tfrac{9}{14}\log_2\tfrac{9}{14} - \tfrac{5}{14}\log_2\tfrac{5}{14} = 0.94$$

$$\begin{split} & \operatorname{Entropy}(S|\operatorname{Wind}) = \tfrac{8}{14} \operatorname{Entropy}(S_{\operatorname{weak}}) + \tfrac{6}{14} \operatorname{Entropy}(S_{\operatorname{strong}}) = 0.89 \\ & \operatorname{Gain}(S,\operatorname{Wind}) = 0.94 - 0.89 = 0.05 \end{split}$$

## Example: should I play tennis? (5)

- Gain(S, Outlook) = 0.246
- Gain(S, Humidity) = 0.151
- Gain(S, Wind) = 0.048
- Gain(S, Temperature) = 0.029

#### $\rightarrow$ Split on Outlook!

In this example, the data had only categorical variables, and no missing values.

What if we had a continuous (not categorical) variable? We would need to also decide how to partition the continuous feature into a discrete set of intervals.

There are a few well-known algorithms for fitting decision trees - CART, ID3, C4.5 - that have different capabilities with respect to continuous features, features with missing values, and what measure of node impurity is used.

e.g. C4.5 introduces the idea that if a sample has a missing value for a feature,

- · when training, compute information gain using only samples where the feature is defined
- · when using, we decide which branch to follow based on which is more probable

## **Feature importance**

- ullet For each feature  $X_i$ , find all nodes where the feature was used as the split variable
- Add up information gain due to split (or for GINI index, difference in loss weighted by number of samples.)
- · This sum reflects feature importance

This feature importance can be used for feature selection or feature weighting!

It tends to do reasonable things both with (1) features that are only useful in combination and (2) features that are highly correlated.

#### **Bias and variance**

## **Managing tree depth**

- If tree is too deep likely to overfit (high variance)
- · If tree is not deep enough likely to have high bias

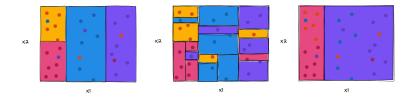


Figure 8: The depth/size of the tree (number of regions) controls the complexity of the regression line or decision boundaries, and the bias variance tradeoff.

## **Stopping criteria**

If we build tree until there is zero error on training set, we have "memorized" training data.

Other stopping criteria:

- · Max depth
- · Max size (number of leaf nodes)
- · Min number of samples to split
- Min number of samples in leaf node
- · Min decrease in loss function due to split

(Can select depth, etc. by CV)

## **Pruning**

- · Alternative to stopping criteria: build entire tree, then prune
- · With greedy algorithm a very good split may descend from a less-good split

## **Pruning classification trees**

We usually prune classification trees using classification error rate as loss function, even if tree was built using GINI or entropy.

## Weakest link pruning (1)

Prune a large tree from leaves to root:

- Start with full tree  $T_{
  m 0}$
- ullet Merge two adjacent leaf nodes into their parent to obtain  $T_1$  by minimizing:

$$\frac{Err(T_1) - Err(T_0)}{|T_0| - |T_1|}$$

#### Weakest link pruning (2)

- Iterate to produce a sequence of trees  $T_0, T_1, \dots, T_m$  where  $T_m$  is a tree of minimum size.
- · Select optimal tree by CV

#### **Cost complexity pruning**

Equivalent to: Minimize

$$\sum_{m=1}^{|T|}\sum_{x_i\in R_m}(y_i-\hat{y}_{R_m})^2+\alpha|T|$$

Choose  $\alpha$  by CV, 1-SE rule ( $\uparrow \alpha, \downarrow |T|$ ).

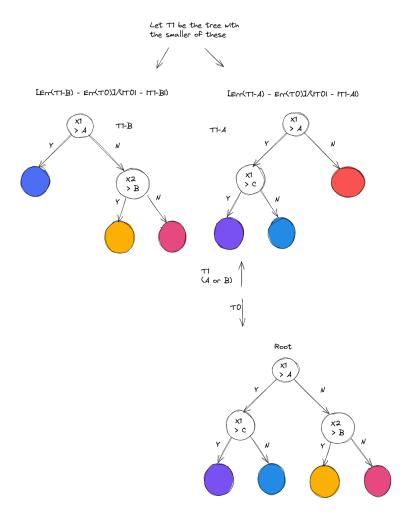


Figure 9: Weakest link pruning.

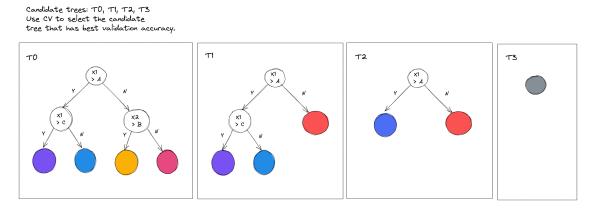


Figure 10: Selecting tree from the set of candidate trees.

## **Summary - so far**

## The good and the bad (1)

## Good:

- · Flexible with much faster inference time than KNN
- Easy to interpret, close to human decision-making
- Can derive feature importance
- Easily handles mixed types, different ranges

## The good and the bad (2)

## Bad:

- Need greedy heuristic to train
- Deep trees have large variance
- Non-robust: Small change in data can cause large change in estimated tree