# Machine Learning

Decision Tree and Random Forest

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## **Decision Tree**

Decision tree = hypothesis/model  $h: \mathcal{X} \to \mathcal{Y}$  that predicts the target value  $h(\mathbf{x})$  for an instance  $\mathbf{x} \in \mathcal{X}$  using a *tree*.

- Prediction obtained starting from the root and traveling to a leaf of the tree.
- At each node, the successor (child) is chosen on the basis of a splitting of the input space
- Usually: splitting based on one of the features of x

# Example (for Classification)

 $\mathcal{Y} = \{0, 1\}$ ; leaf = predicted label

# Geometric Interpretation

## Expressiveness of Decision Trees

Assume  $\mathcal{X} = \{0,1\}^d \Rightarrow$  rules in each node are of the type " $x_i = 1$ " for some  $i \in \{1,\ldots,d\}$ 

### Proposition

Let  $\mathcal{X} = \{0,1\}^d$ , and let f be any function from  $\mathcal{X}$  to  $\{0,1\}$ . Then the set of hypothesis  $\mathcal{H} = \{\text{decision tree from } \{0,1\}^d \text{ to } \{0,1\} \}$  contains a tree h that for each  $\mathbf{x} \in \mathcal{X}$ :  $h(\mathbf{x}) = f(\mathbf{x})$ .

## Sample Complexity of Decision Trees

## Corollary

The VC-dimension of  $\mathcal{H} = \{\text{decision tree from } \{0,1\}^d \text{ to } \{0,1\} \}$  is  $2^d$ .

A large number of samples is required to learn decision trees!

**In practice**: need to limit the *complexity* of the tree.

### How? Various options

- fix a maximum depth for the tree;
- fix a minimum number of training samples "corresponding" to a leaf
- instead of looking for the tree in  $\mathcal{H}$  that minimizes the training error  $L_S(h)$  (S=training set), look for the set that minimizes  $L_S(h) + R(h)$ , where R(h) is some measure of complexity for the tree h (similar to regularization...)

## Algorithm to Learn A Decision Tree

How do we find the best decision tree?

Informally: computing the "best" decision tree is NP-hard

⇒ in practice: greedy (non-optimal) approaches are used

# Greedy Approach to Learn Decision Trees

Assume binary classification and binary features.

Training set 5

#### Overall scheme:

- start with a tree with a single leaf (the root); label of this leaf
  majority vote among all labels over the training set.
- perform a series of iterations, for each iteration
  - examine the effect of splitting a single leaf;
  - define some "gain" measure that quantifies the improvement due to the split
  - among all possible splits, either choose the one that maximizes the gain and perform it, or choose not to split the leaf at all.

**Note**: very easy to include constraints on the "complexity" of the tree (e.g.,build trees with depth < some value)

#### ID3(S,A)

INPUT: training set S, feature subset  $A \subseteq [d]$ 

if all examples in S are labeled by 1, return a leaf 1

if all examples in S are labeled by 0, return a leaf 0

if  $A = \emptyset$ , return a leaf whose value = majority of labels in S else :

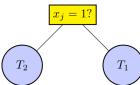
Let  $j = \operatorname{argmax}_{i \in A} \operatorname{Gain}(S, i)$ 

if all examples in S have the same value of  $\times_{\S}$ :

Return a leaf whose value = majority of labels in S

#### else

Let  $T_1$  be the tree returned by  $ID3(\{(\mathbf{x},y) \in S : x_j = 1\}, A \setminus \{j\})$ . Let  $T_2$  be the tree returned by  $ID3(\{(\mathbf{x},y) \in S : x_j = 0\}, A \setminus \{j\})$ . Return the tree:



## Gain Measures

There are several definitions, we consider the simplest one: **decrease in training error**.

Define 
$$C(a) = \min\{a, 1 - a\}$$
.

Let  $Pr_S[y=1]$  be the probability that a sample (uniformly at random) from S has label 1.

Then the training error before splitting the data S is  $C(Pr_S[y=1])$ .

Why? Because of the *majority of labels* rule.

The training error after splitting the data S using feature  $x_i$  is:

$$\Pr_{S}[x_{i} = 1]C(\Pr_{S}[y = 1|x_{i} = 1]) + \Pr_{S}[x_{i} = 0]C(\Pr_{S}[y = 1|x_{i} = 0])$$

Gain using feature  $x_i$  for splitting: difference in error before and after splitting

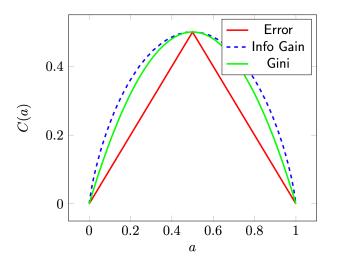
$$Gain(S, i) = C(\Pr_{S}[y = 1]) -$$

$$\left(\Pr_{S}[x_{i}=1]C(\Pr_{S}[y=1|x_{i}=1]) + \Pr_{S}[x_{i}=0]C(\Pr_{S}[y=1|x_{i}=0])\right)$$

## Other Gain Measures

Obtained by different definition of C(a):

- Information gain:  $C(a) = -a \log a (1-a) \log(1-a)$
- Gini index: C(a) = 2a(1-a)



## What About Real-valued Features?

Consider the *i* feature that takes real value  $(x_i \in \mathbb{R})$ .

Assume we have "sorted" the training set  $S = \{x_1, ..., x_m\}$  so that for the *i*-th feature we have  $x_{1,i} \le x_{2,i} \le \cdots \le x_{m,i}$ .

Define thresholds  $\theta_{0,i}, \theta_{1,i}, \dots, \theta_{m+1,i}$  with

- $\theta_{0,i} \in [-\infty, x_{1,i}]$
- $\theta_{j,i} \in [x_{j,i}, x_{j+1,i}]$  for all j = 1, ..., m
- $\theta_{m+1,i} \in [x_{m,i}, \infty]$

For every pair i, j, i = 1, ..., m, j = 0, ..., m+1 and every sample  $\mathbf{x} \in S$ , consider the sample  $\mathbf{x}'$  having features  $\mathbf{x}'_{i,j} = 1$  if  $\mathbf{x}_i < \theta_{j,i}$ , and  $\mathbf{x}'_{i,j} = 0$  otherwise.

Run the algorithm for binary features on the new samples  $\mathbf{x}'$ .

### Additional Notes

- in practice, *pruning* procedures are often applied after a tree is built to simplify the tree
- there are other algorithms to learn trees, with similar approaches (e.g., scikit-learn uses an implementation of CART)
- what about regression?
  - squared loss is often used
  - prediction of a leaf = average value of target for associated samples
  - error on a set of samples = mean squared error
- decision trees are interpretable models

## Random Forest

A different approach to prevent overfitting for decision trees.

It is an *ensemble method*: builds a *collection* of decision trees from the same data

#### Prediction:

- classification: majority vote over the predictions of the individual trees.
- regression: average of predictions of the individual trees.

How are the trees generated?

## Tree Generation in Random Forest

Let S be the training set of m samples, with d features.

To build one of the trees in the forest:

- sample a new training set S' of size m using the uniform distribution over S
- 2 construct a sequence  $\mathcal{I}_1$ ,  $\mathcal{I}_2$ , . . ., where each  $\mathcal{I}_t$  is a subset of size k of the d features, generated by sampling uniformly at random the features.

**Intution**: if **k** is small, it will be difficult to overfit