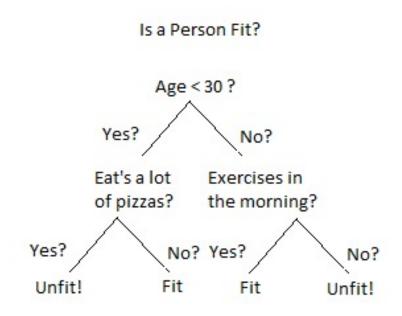
Lecture 9: decision trees

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Decision trees - introduction

- Decision tree is a simple and easy to interpret classifier.
- In decision trees, the data is continuously split according to a certain parameter.
- The tree can be explained by two entities, namely decision nodes and leaves. The leaves are the decisions or the final outcomes, and the decision nodes are where the data is split.



Decision tree learning

- How many trees? 2^m (exponential in the number of attributes m)
 - Trees containing only one attribute
 - Trees containing two attributes
 - ...
 - Trees containing m attributes
- Many algorithms: reasonably accurate, suboptimal, reasonable amount of time
 - Hunt's Algorithm (basis of many others)
 - CART (Classification and Regression Trees), a book by Breiman et al.
 - ID3, C4.5 by Quinlan

Hunt's algorithm

• Start from the tree root and split the data on the feature that results in the largest **Information Gain (IG)**.

• Input:

- D_t is the set of training records that reach a node t
 - Root node corresponds to the whole training dataset
- $y = \{y_1, y_2, \dots, y_c\}$ are class labels

General procedure

- If D_t contains records that belong to the same class y_t (i.e., the node is pure), then t is a leaf node labeled as y_t
- If D_t contains records that belong to more than one class
 - Use an attribute test to split the data into smaller subsets
 - Recursively apply the procedure to each subset

Hunt's algorithm – questions to ask

- Which attribute to choose to split the node?
- How to split the attribute?

Measures of node impurity

- Let p(i|t) represent the probability that an instance t in D_t belongs to class C_i , estimated by $\frac{n_i}{|D_t|}$, where n_i is the number of instances with class label C_i .
- Gini index

$$Gini(t) = 1 - \sum_{i=0}^{c-1} (p(i|t))^2$$

• Entropy (log uses base 2: information is encoded in bits)

$$Entropy(t) = -\sum_{i=0}^{c-1} p(i|t)log_2 p(i|t)$$

Classification error

Classification error(t) =
$$1 - max_i\{p(i|t)\}$$

Gini index

$$Gini(t) = 1 - \sum_{i=0}^{c-1} (p(i|t))^2$$

| C2 6 C2 5 C2 4 C2 | | | | | / | | , ., , | - 1 |
|---|------|-------|------|---|------|---|----------|-----|
| C1 0 C1 1 C1 2 C1 3 | | _ | | 1 | | _ | <u> </u> | |

Table 1: Data example 1

$$Gini(N_1) = 1 - \left(\frac{0}{6}\right)^2 - \left(\frac{6}{6}\right)^2 = 0$$

$$Gini(N_3) = 1 - \left(\frac{2}{6}\right)^2 - \left(\frac{4}{6}\right)^2 = \frac{16}{36} = 0.444$$

$$Gini(N_2) = 1 - \left(\frac{1}{6}\right)^2 - \left(\frac{5}{6}\right)^2 = \frac{10}{36} = 0.278$$

$$Gini(N_4) = 1 - 0.5^2 - 0.5^2 = 0.5$$

- Maximum: $1-\frac{1}{n_c}$ when records are equally distributed among all classes, implying least interesting information
- Minimum: (0.0) when all records belong to one class, implying most interesting information
- First used in CART, which allows only binary splitting

$$Entropy(t) = -\sum_{i=0}^{c-1} p(i|t)log_2p(i|t)$$

| | C1 | 0 | |
|----|--------|------|---|
| | C2 | 6 | |
| (8 | a) Noc | de N | 1 |

$$\begin{array}{c|ccccc} C1 & 2 & & C1 & 3 \\ \hline C2 & 4 & & C2 & 3 \\ \hline (c) Node N_3 (d) Node $N_4$$$

Table 1: Data example 1

$$Entropy(N_1) = -0log0 - 1log1 = 0$$

$$Entropy(N_3) = -\frac{2}{6}\log(\frac{2}{6}) - \frac{4}{6}\log(\frac{4}{6}) = 0.92$$

$$Entropy(N_2) = -\frac{1}{6}\log(\frac{1}{6}) - \frac{5}{6}\log(\frac{5}{6}) = 0.65$$

$$Entropy(N_4) = -0.5\log 0.5 - 0.5log 0.5 = -(-1) = 1$$

- prob. is 0 means it does not happen, let 0log0 = 0
- Measures homogeneity of a node.
 - Maximum: $\log(n_c)$ when records are equally distributed among all classes implying least information. Given n_c : the number of classes. Then, $-n_c * \left(\frac{1}{n_c}\right) * \log_2\left(\frac{1}{n_c}\right) = -\log_2\left(\frac{1}{n_c}\right) = \log_2(n_c)$
 - Minimum: 0.0 when all records belong to one class, implying most information

Determine the best split – Maximizing the information gain

• The attribute that we choose to split the data to smaller subsets need to gain the **maximum information gain**.

$$IG(D_p, f) = I(D_p) - \sum_{j=1}^{m} \frac{N_j}{N_p} I(D_j)$$

- *f*: the feature/attribute to perform the split.
- D_p : dataset corresponding to the parent node.
- D_i : dataset corresponding to the *j*th child node.
- N_p , N_j : number of samples in D_p and D_j respectively.
- $I(\cdot)$: impurity measure of a node.

Determine the best split

Most libraries implement binary decision tree.

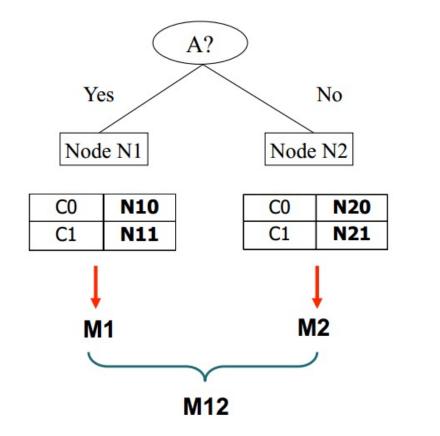
$$IG(D_p, f) = I(D_p) - \frac{N_{left}}{N_p} I(D_{left}) - \frac{N_{right}}{N_p} I(D_{right})$$

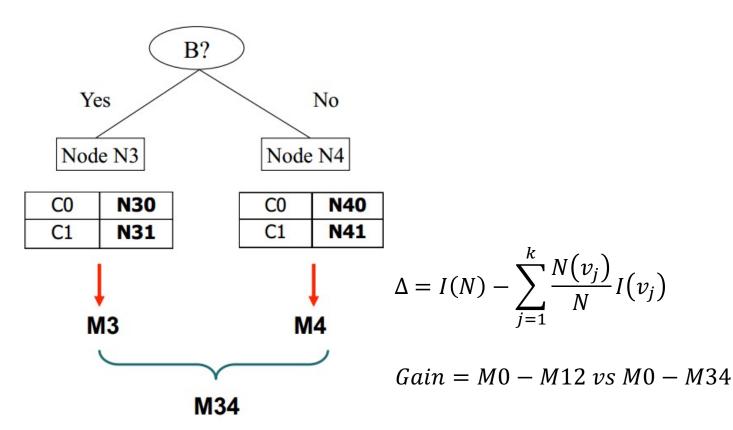
Binary decision tree - example

Before splitting:

| C0 | N00 | 1 |
|----|-----|-----|
| C1 | N01 | IV. |

M0 = I(N)





Binary decision tree - example

Splitting one attribute A

| | parent node N_p | left child node N_1 | right child node N ₂ |
|---------------------------------|-------------------|-----------------------|---------------------------------|
| Instances belonging to class C1 | 40 | 30 | 10 |
| Instances belonging to class C2 | 40 | 10 | 30 |

Splitting one attribute B

| | parent node N_p | left child node N ₃ | right child node N_4 |
|---------------------------------|-------------------|--------------------------------|------------------------|
| Instances belonging to class C1 | 40 | 20 | 20 |
| Instances belonging to class C2 | 40 | 40 | 0 |

Using **gini index**:

Splitting on A:
$$IG(N_p, A) = 0.5 - \frac{4}{8} * 0.375 - \frac{4}{8} * 0.375 = 0.125$$

Splitting on *B*:
$$IG(N_p, B) = 0.5 - \frac{6}{8} * 0.4 - 0 = 0.16$$

Thus, splitting on B is preferred.

Using **Entropy**

Splitting on A:
$$IG(N_p, A) = 1 - \frac{4}{8} * 0.81 - \frac{4}{8} * 0.81 = 0.19$$

Splitting on *B*:
$$IG(N_p, B) = 1 - \frac{6}{8} * 0.92 - 0 = 0.31$$

Thus, splitting on B is preferred.

Using classification error

Splitting on A:
$$IG(N_p, A) = 0.5 - \frac{4}{8} * 0.25 - \frac{4}{8} * 0.5 = 0.25$$

Splitting on *B*:
$$IG(N_p, B) = 0.5 - \frac{6}{8} * \frac{1}{3} - 0 = 0.25$$

Thus, no preference on splitting attribute.

Stopping criteria for tree induction

- Stop expanding a node when all the records belong to the same class
- Stop expanding a node when all the records have the same attribute values
- In practice, such stopping criteria may result in a deep tree with many nodes, which may lead to an issue called overfitting. Thus, the decision trees typically need to be pruned by setting a limit for the maximal depth of the tree.

Decision tree based classification - discussions

Advantages

- Inexpensive to construct
- Extremely fast at classifying unknown records
- Easy to interpret for small-sized trees
- Accuracy is comparable to other classification techniques for many simple data sets

Decision Boundary

- Decision boundary is a border line between two neighboring regions of different classes
- Decision boundary is parallel to axes because test condition involves a single attribute at-a-time.

Overfitting

Classification errors

- Training errors: the number of misclassification errors on training records
- **Generalization/test errors**: the expected error of the model on previously unseen records

Overfitting and Underfitting

- **Model overfitting**: A model fits the training data too well but has poorer generalization error than a model with a higher training error.
- **Model underfitting**: a model has not learned the true structure of the data. It has high training error and generalization error.
- Underfitting: when a model is too simple; both training and test errors are large

Variance & Bias

- When a model suffers from overfitting, we also say that the model has a high variance.
 - Which can be caused by having too many parameters, leading to a model that is too complex given the underlying data.
- When a model suffers from underfitting, we also say that the model has high bias.
 - The model is not complex enough to capture the pattern in the training data well and therefore also suffers from low performance on unseen data.

Overfitting reasons

- Possible reason 1: noise
- Possible reason 2: insufficient samples
- What is the primary reason for overfitting? a subject of debate
- Generally agreed: the complexity of a model has an impact on model overfitting

How to address overfitting: pre-pruning (early stropping rule)

- Stop the algorithm before it becomes a fully-grown tree
- Typical stopping conditions for a node:
 - Stop if all instances belong to the same class
 - Stop if all the attribute values are the same
- More restrictive conditions
 - Stop if the number of instances is less than some user-specified threshold
 - Stop if expanding the current node does not improve impurity measures or estimated generalization error.

How to address overfitting: post-pruning

- Grow decision tree to its entirety
- Trim the nodes of the decision tree in a bottom-up fashion
- Replace a subtree by
 - a leaf node: class label is determined from majority class of instances in the sub-tree (subtree replacement)
 - most frequently used branch of the subtree (subtree raising)
- Termination: no further improvement on generalization errors

Building a decision tree

• Feature scaling is not required for decision tree algorithms.

```
from sklearn import tree
from sklearn.tree import DecisionTreeClassifier
tree model = DecisionTreeClassifier(criterion='gini',
                   max_depth=4,
                   random state=1)
tree_model.fit(X_train, y_train)
tree.plot_tree(tree_model)
plt.show()
```

class sklearn.tree.DecisionTreeClassifier(criterion='gini', splitter='best', max_dept h=None, min_samples_split=2, min_samples_leaf=1, min_weight_fraction_leaf=0.0, max_features=None, random_state=None, max_leaf_nodes=None, min_impu rity_decrease=0.0, min_impurity_split=None, class_weight=None, presort='depre cated', ccp_alpha=0.0)

- criterion: The function to measure the quality of a split. Supported criteria are "gini" for the Gini impurity and "entropy" for the information gain.
- max_depth: the maximum depth of the tree. If None, then nodes are expanded until all leaves are pure or until all leaves contain less than min_samples_split samples.
- min_samples_split: the minimum number of samples required to split an internal node
- min_samples_leaf: the minimum number of samples required to be at a leaf node.

DecisionTreeClassifier parameter

min_impurity_decrease: A node will be split if this split induces a
decrease of the impurity greater than or equal to this value. The
weighted impurity decrease equation is the following

$$\frac{N_t}{N}*impurity - \frac{N_{t_R}}{N_t}*right_impurity - \frac{N_{t_L}}{N_t}*left_impurity$$

- Where N is the total number of samples
- N_t is the number of samples at the current node
- N_{t_L} is the number of samples in the left child
- N_{t_R} is the number of samples in the right child

Prediction

- predict(self, X, check_input=True): Predict class (or regression value) for X.
- **predict_proba**(*self*, *X*, *check_input=True*) Predict class probabilities of the input samples X.