IT496: Introduction to Data Mining



Lecture 19

Decision Tree Classifier

(Slides are created from the book Hands-on ML by Aurelien Geron)

Arpit Rana 5th October 2023

Decision Trees

A decision tree is a representation of a function that maps a vector of attribute values to a single output value—a "decision."

- Decision Trees can be used for *regression* and *classification*.
 - o for binary or multiclass classification (so you don't need one-versus-rest or one-versus-all)
- They are more complex than linear models and so can better fit complex datasets.
- Many people claim that they produce interpretable models.
- Random Forests are another popular model in Machine Learning, and they contain Decision Trees.

Decision Trees

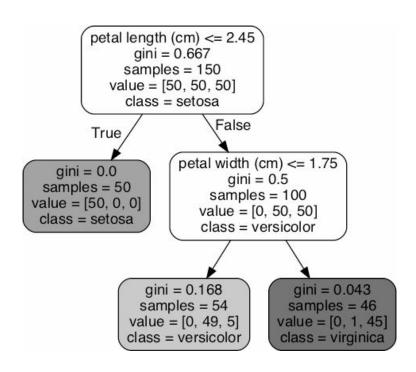
A decision tree reaches its decision by performing a sequence of tests, starting at the root and following the appropriate branch until a leaf is reached.

- Each *internal node* in the tree corresponds to a test of the value of one of the input attributes,
- the *branches* from the node are labeled with the possible values of the attribute, and
- the *leaf nodes* specify what value is to be returned by the function.

Decision Tree on the Iris Dataset

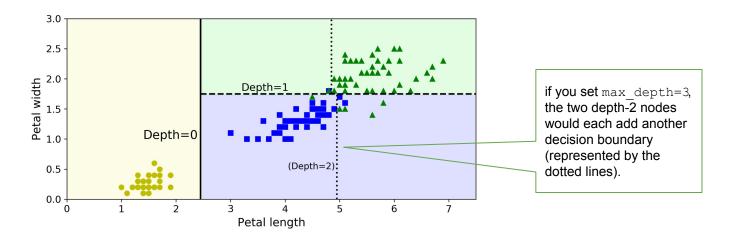
```
from sklearn.datasets import load_iris
from sklearn.tree import DecisionTreeClassifier
iris = load_iris()
X = iris.data[:, 2:] # petal length and width
y = iris.target

tree_clf = DecisionTreeClassifier(max_depth=2)
tree_clf.fit(X, y)
```



Making Predictions

- What about the flower with Petal length = 5 and Petal width = 1.5?
 - Start at the root: is the Petal length <= 2.45 cm?
 - No, so move right: is the Petal width <= 1.75 cm?</p>
 - Yes, so move left: it is an Iris-Versicolor.



Decision Tree decision boundaries

Reading the Nodes of the Tree

- samples: how many training examples the node applies to.
 - E.g. 100 training examples have a petal length > 2.45 cm.
- value: how many training examples of each class this node applies to.
 - o E.g. [0, 1, 45] means 0 Iris-Sentosa, 1 Iris-Versicolor, and 45 Iris-Virginica.
- gini: is the *Gini impurity* of a node (with values in [0.0, 1.0]):
- it measures how often an example would be incorrectly labeled if it was randomly labeled according to the distribution of labels for this node;
 - e.g. gini of 0 means no impurity: all examples that this node applies to belong to the same class;

Reading the Nodes of the Tree

The *Gini impurity/ Gini index* measures the impurity of X, a data partition or set of training tuples at a node, as

$$Gini(X) = 1 - \sum_{i=1}^n p_i^2$$

where p_i is the probability that a tuple in X belongs to the class C_i and is estimated by $|C_{i,X}|/|X|$. The sum is computed over n classes.

• For example, in the tree shown previously, the depth-2 left node has a gini score equal to $1 - (0/54)^2 - (49/54)^2 - (5/54)^2 \approx 0.168$.

Learning a Decision Tree

- scikit-learn uses the CART Algorithm (Classification and Regression Tree).
 - It only produces binary trees (hence yes/no questions in non-leaf nodes);
 - o it is *recursive* (repeats until it reaches some stopping criterion); and
 - it is *greedy* (It does not check whether or not the split will lead to the lowest possible impurity several levels down).
- There are other algorithms:
 - o ID3, which can produce non-binary trees, and
 - o C4.5, which is like CART but uses *entropy* in place of *Gini*, and
 - Others including ones that can directly handle nominal-valued features and missing values, which we will not study.

The CART Training Algorithm

- For each feature x_i and each value v, split the dataset into two:
 - \circ X_{left} are the examples in X for which $x_i \le v$
 - \circ X_{right} are the examples in X for which $x_i > v$

and calculate the **CART's loss function**:
$$\frac{|X_{left}|}{|X|}Gini(X_{left}) + \frac{|X_{right}|}{|X|}Gini(X_{right})$$

- From the above, choose the feature x_i and a value v with the lowest loss
- If a stopping criterion has been reached (e.g. maximum depth or if no split reduces impurity) then:
 - o return
- Else:
 - Recursively call CART on X_{left}
 - o Recursively call CART on X_{right}

Regularization Hyperparameters

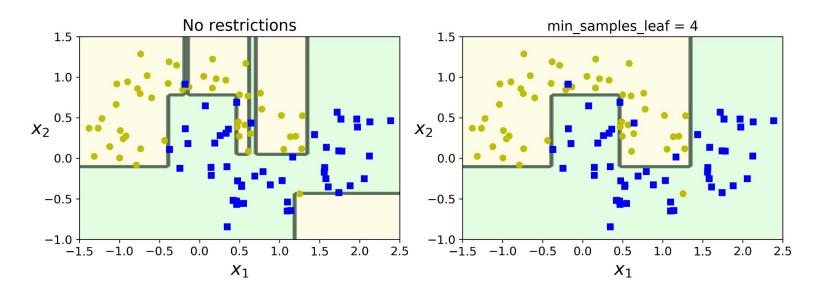
- max_depth is a hyperparameter.
 - o Increasing max_depth increases model's complexity and may result in overfitting.
 - If there is no constraint on tree depth, then branches will be grown until all leaves are pure.
- There are other hyperparameters:
 - min_samples_split, the minimum number of samples a node must have before it can be split,
 - min_samples_leaf, the minimum number of samples a leaf node must have,
 - $\verb|omax_leaf_nodes|, maximum number of leaf nodes|, and \\$
 - max_features, maximum number of features that are evaluated for splitting at each node.

Increasing min_* or reducing max_* hyperparameters will regularize the model.

Regularization Hyperparameters

Regularization using min_samples_leaf

It is quite obvious that the model on the left is overfitting, and the model on the right will probably generalize better.



Computational Complexity

- Learning is $O(mn \log_2 m)$ for the basic CART algorithm above.
 - o Of course, hyperparameters such as a maximum depth can speed-up learning.
- Prediction is roughly $O(\log_2 m)$ (which is the depth of tree, assuming the tree is balanced, which it often, approximately, is). This is fast!

Regression using a Decision Tree

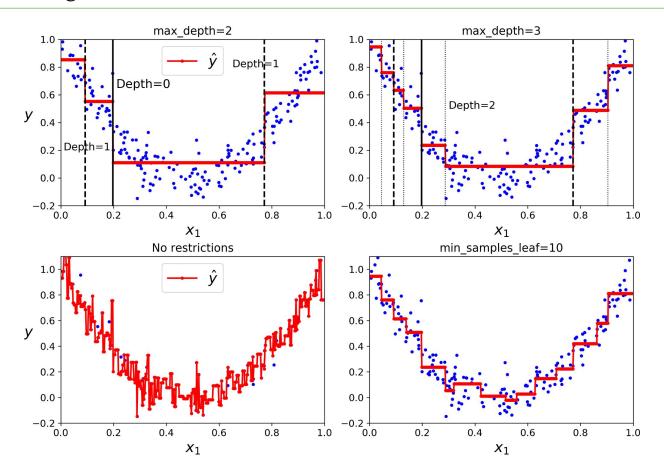
- What does the Decision Tree Regressor predict?
 - Start at the root and follow the decisions down to a leaf.
 - At any node, value is the prediction and is simply the mean target values of the training examples that the node applies to.
- For regression, the only difference is the loss function: in place of *Gini* it uses the *mean* squared error between the y-values of the training examples and their mean.

$$rac{|X_{left}|}{|X|}MSE(X_{left}) + rac{|X_{right}|}{|X|}MSE(X_{right})$$

Regression using a Decision Tree

```
from sklearn.tree import DecisionTreeRegressor
tree_reg = DecisionTreeRegressor(max_depth=2)
tree reg.fit(X, y)
                                                                       x1 \le 0.197
                                                                       mse = 0.098
                                                                      samples = 200
                                                                       value = 0.354
                                                                                 False
                                                                    True
                                                               x1 <= 0.092
                                                                                x1 \le 0.772
                                                               mse = 0.038
                                                                                mse = 0.074
                                                              samples = 44
                                                                               samples = 156
                                                              value = 0.689
                                                                               value = 0.259
                                               mse = 0.018
                                                                                mse = 0.015
                                                               mse = 0.013
                                                                                                 mse = 0.036
                                              samples = 20
                                                              samples = 24
                                                                               samples = 110
                                                                                                samples = 46
                                              value = 0.854
                                                              value = 0.552
                                                                               value = 0.111
                                                                                                value = 0.615
```

Regression using a Decision Tree



Decision Tree: Pros and Cons

- The model is interpretable:
 - we can display the tree and people can see what has been learned.
- An individual prediction is *explainable*:
 - we can display the path through the tree.
- Note we do not need to scale the data before using this algorithm.
- They are very sensitive to small variations in the training data (a.k.a. *instability*). Use of PCA or Random Forests can limit this issue.

Parametric vs. Non-parametric Learning

- Parametric learning: the number of parameters is known prior to training and is not affected by *m* the number of examples in the training set.
 - \circ E.g. Linear Regression the number of parameters is n + 1
 - E.g. Polynomial Regression the number of parameters is $\frac{(n+d)!}{n!d!}$
- Non-parametric learning: the number of parameters is not known in advance and may grow with the size of the training set.
 - E.g. For Decision Trees, in some sense, the nodes are the parameters: the structure of the model (tree) may grow to accommodate the complexity of the training data.
 - E.g. For kNN, in some sense, the neighbours are the parameters: the more training examples there are, the more different possible sets of neighbours there are.

Parametric vs. Non-parametric Learning

This does not mean that non-parametric models will have lower validation error.

- Unconstrained, they are prone to overfitting.
- So we want to impose some constraints on the CART algorithm to restrict the shape of the Decision Tree (such as maximum depth and others).
- Similarly, we avoid small values for *k* in *k*NN.

Dimensionality Reduction	
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6th October 2023

Next lecture