Ch6. Decision Tree



01 Decision Tree

Algorithms that enable classification, regression, and multiple output operations

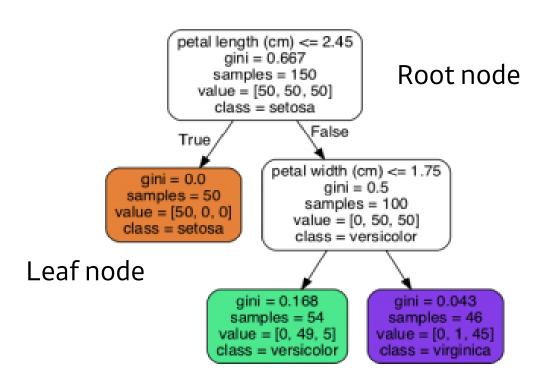
Training

```
from sklearn.datasets import load_iris
from sklearn.tree import DecisionTreeClassifier

iris = load_iris()
X = iris.data[:, 2:]
y = iris.target

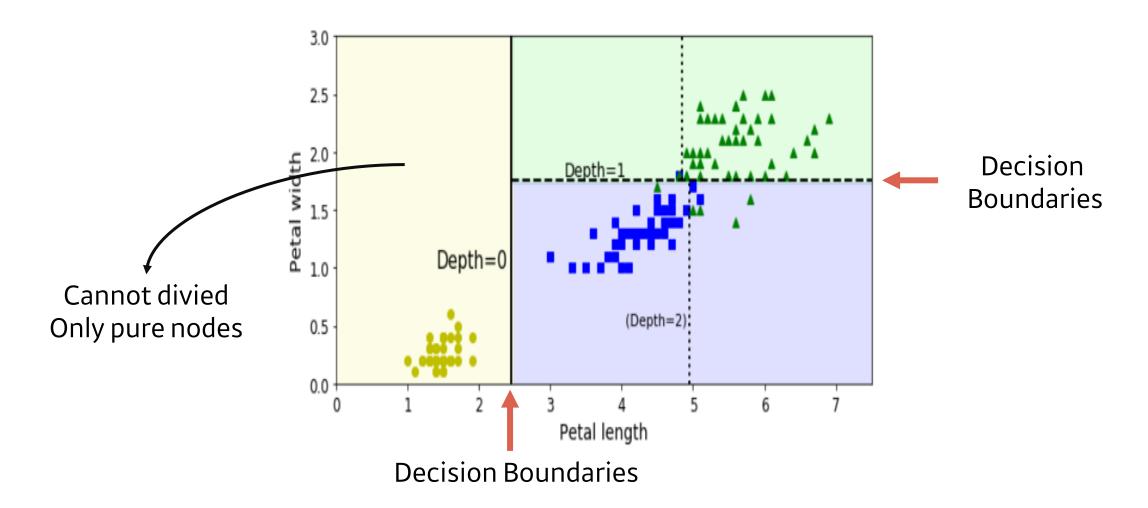
tree_clf = DecisionTreeClassifier(max_depth=2, random_state=42)
tree_clf.fit(X,y)
```

O2 Predict



- Sample: Number of data in training sample applied to the tree
- Value: How many training samples are present on the node by class
- Gini: To determine how many data belong to same class on that node by impurity

$$G_i = 1 - \sum_{k=1}^n p_{i,k}^2$$



Decision tree is called white box model => intuitive and easy to understand how to make decisions



03 Estimate class probability

Calculate the probability that a sample belongs to a particular class

- Explore the tree to find leaf node for sample
- 2) Return the percentage of the training samples of class k of that node



04 CART algorithm

Divide into 2 subsets with binary algorithm used to train the decision tree

Cost function to be minimized

$$J(k,t_k) = rac{m_{ ext{left}}}{m}G_{ ext{left}} + rac{m_{ ext{right}}}{m}G_{ ext{right}}$$
 - tk: d - tk: d - Stop 여기에서 $\begin{cases} G_{ ext{left/right}}$ 는 왼쪽/오른쪽 서브셋의 불순도 $m_{ ext{left/right}}$ 는 왼쪽/오른쪽 서브셋의 샘플 수

- k:training set
- tk: divide subsets into 2 by using this threshold
- Stop dividing
 - 1) At maximum depth
 - 2) When no segmentation to reduce impurity is found

=> Finding the optimal tree is NP complete problem



05 Computational Complexity

Decision tree is balanced + Predicted by checking only one characteristic value

=> Total complexity is independent of the number of attributes

: O (log₂(m))

Training Algorithm: Comparing all characteristics of all training samples

=> Small training set can be data- aligned to speed up training Large training set slow down a lot



To measure the disorder of a molecule => Stable & orderly, entropy = 0

$$H_{i} = -\sum_{\substack{k=1 \ p_{i,k} \neq 0}}^{n} p_{i,k} \log_{2}(p_{i,k})$$

- Gini impurity: Faster
 but, tendency to isolate the most
 frequent class to one side
- Entropy: Create a more balanced tree
- => No significant difference and create similar tree



07 Regulatory Hyperparameter

If not restricted to parameters, Overfitting is likely to occur

- Hyper parameter type
 - max_depth
 - min_samples_split
 - min_samples_leaf
 - min_weight_fraction_leaf
 - max_leaf_nodes
 - max_features

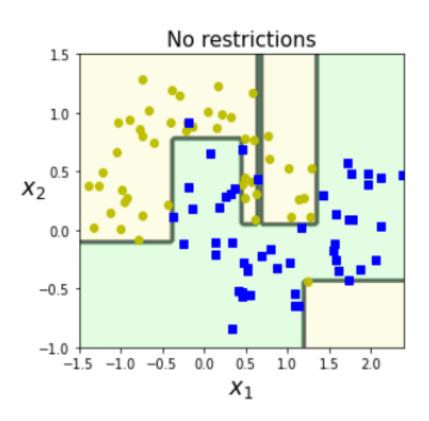
⇒ Min : increase paratemers

⇒ Max: decrease parameters

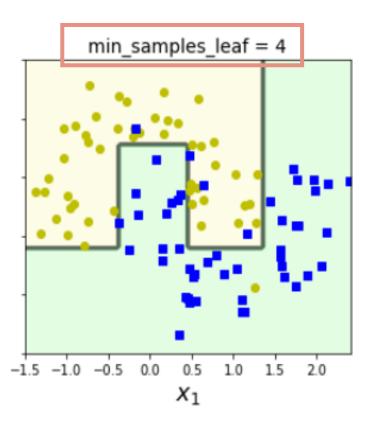
To increase regulation of model



07 Regulatory Hyperparameter



Overfitting No regulatory variables



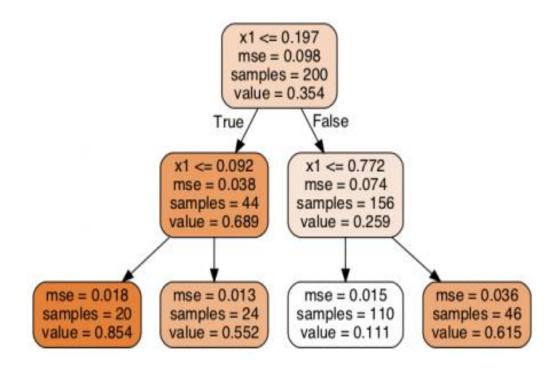
Good generalization Applying regulatory variables

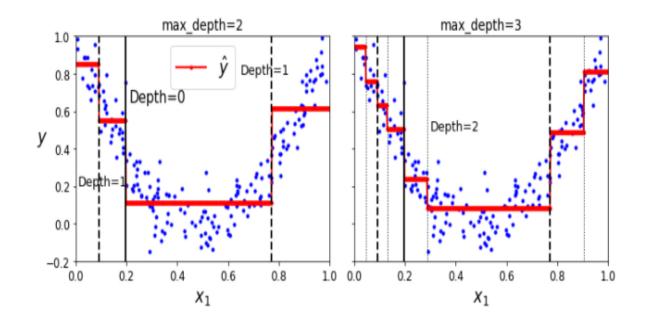


```
from sklearn.tree import DecisionTreeRegressor
tree_reg = DecisionTreeRegressor(max_depth=2, random_state=42)
tree_reg.fit(X, y)
```

Difference from classification tree

: Instead of predicting classes on each node, Predicting a value





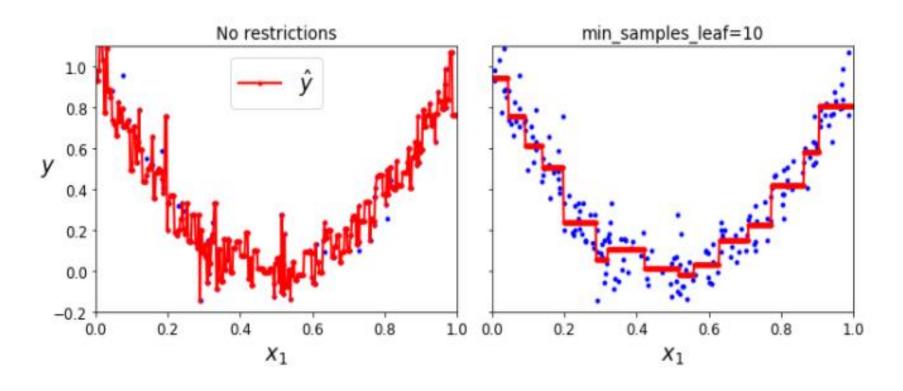
- Predicted values : Average of target values
 - => Algorithm divides the region so that as many samples as possible are close together
- CART: Split MSE to minimize

$$J(k, t_k) = \frac{m_{\text{left}}}{m} \text{MSE}_{\text{left}} + \frac{m_{\text{right}}}{m} \text{MSE}_{\text{right}}$$

where
$$\begin{cases} MSE_{node} = \sum_{i \in node} (\hat{y}_{node} - y^{(i)})^2 \\ \hat{y}_{node} = \frac{1}{m_{node}} \sum_{i \in node} y^{(i)} \end{cases}$$



Without regulation in regression, decision trees are likely to be over-fitting

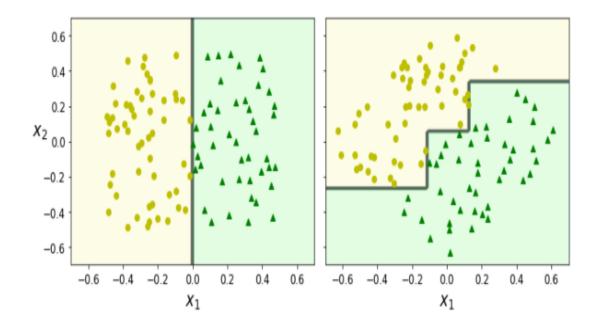


Over-fitting unregulated

Avoid over-fitting & Derive plausible model Regulation



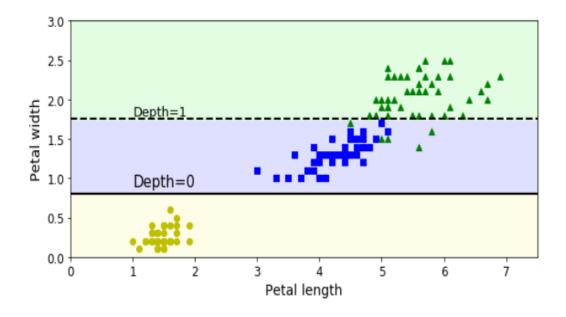
1) Sensitive to rotation of training sets



Right side will not generalize well

⇒ Training data in better direction Using PCA to rotate

2) Sensitive to small changes in training data



Remove data from training set

⇒ Can confirm that it is different from before

THANK YOU