

# Decision Tree

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# Introduction

- Decision tree is one of the most widely used supervised learning method in machine learning
- Its big appeal is that the decision process is very much akin to how we humans make decisions
- Therefore it is easy to understand and accept the results coming from the tree-style decision process
- It used to be an baseline model when constructing some other algorithms (e.g. Random Forest, Boosting)

# Decision Tree in a nutshell

## Decision Tree Construction Process

- 1 Recursively partition the feature space  $\mathcal{X}$  into the subregions  $t_1, \dots, t_m$  based on the specific **splitting rule**
- 2 Stop splitting the feature space (or stop growing tree) when the **stopping rule** holds
- 3 Prune the grown tree based on the **pruning rule**
- 4 Make a (local) prediction on each subregion in pruned tree based on the specific **estimation method**

# Four Ingredients in Decision Tree Construction

- **Splitting rule**

- How does the split works?
- What is the criterion for splitting the feature space?

- **Stopping rule**

- What kind of threshold is used for stopping?

- **Pruning rule**

- Why prune the tree?
- How does it works?

- **Estimation method**

- What kind of estimation method is used?
- How does it works?

## Notation

- $\mathfrak{D} = \{(x^{(i)}, y^{(i)})\}_{i=1}^N$ : the dataset,
- Where  $x^{(i)} = (x_1^{(i)}, \dots, x_d^{(i)}) \in \mathfrak{X}$  and  $y^{(i)} \in \mathfrak{Y}$ , where  $\mathfrak{Y} = \{1, \dots, K\}$
- Let  $t$  be a node, which is also identified with a subregion of  $\mathfrak{X}$
- $\mathfrak{D}(t) = \{(x^{(i)}, y^{(i)}) \in \mathfrak{D} : (x^{(i)}, y^{(i)}) \in t\}$ : the set of data points in the node  $t$
- Let  $T$  be a tree
- $N = |\mathfrak{D}|$ : the total number of data points
- $N(t) = |\mathfrak{D}(t)|$ : the number of data points in the node  $t$
- $N_j(t) = |\{(x^{(i)}, y^{(i)}) \in \mathfrak{D}(t) : y^{(i)} = j\}|$ : the number of data points in the node  $t$  with class label  $j \in \mathfrak{Y}$
- $p(j, t) = \frac{N_j}{N} \cdot \frac{N_j(t)}{N_j} = \frac{N_j(t)}{N}$
- $p(t) = \sum_j p(j, t) = \frac{N(t)}{N}$
- $p(j|t) = \frac{p(j, t)}{p(t)} = \frac{N_j(t)}{N(t)}$

## Splitting Rule

- For each node, determine a splitting variable  $x_j$  and splitting criterion  $c$
- For continuous splitting variable, splitting criterion  $c$  is a number.
  - For example, if an observation  $x^{(i)}$  is the case that its splitting variable  $x_j^{(i)} < c$
  - Then tree assigns it to the left child node. Otherwise tree assigns it to the right child
- For categorical variable, the splitting criterion divides the range of the splitting variable in two parts
  - For example, let splitting variable  $x_j \in \{1, 2, 3, 4\}$
  - And let the splitting criterion is  $\{1, 2, 4\}$
  - If  $x_j^{(i)} \in \{1, 2, 4\}$ , tree assigns it to the left child. Otherwise, tree assigns it to the right child

## Splitting Rule

- A split is determined based on the impurity
- Impurity (or purity) is the measure of homogeneity for a given node
- For each node, we select a splitting variable and a splitting criterion which minimizes the sum of impurities of the two child nodes
- Impurity is calculated by an *impurity function*  $\phi$  which satisfies the following conditions



## Splitting Rule

- **Definition 1.** An **impurity function**  $\phi$  is a function  $\phi(p_1, \dots, p_K)$  defined for  $p_1, \dots, p_K$  with  $p_j \geq 0$  for all  $j$  and  $p_1 + \dots + p_K = 1$  such that

(i)  $\phi(p_1, \dots, p_K) \geq 0$

(ii)  $\phi(1/K, \dots, 1/K)$  is the maximum value of  $\phi$

(iii)  $\phi(p_1, \dots, p_K)$  is symmetric with regard to  $p_1, \dots, p_K$

(iv)  $\phi(1, 0, \dots, 0) = \phi(0, 1, \dots, 0) = \phi(0, \dots, 0, 1) = 0$

- **Definition 2.** For node  $t$ , its impurity (measure)  $i(t)$  is defined as

$$i(t) = \phi(p(1|t), \dots, p(K|t))$$

# Splitting Rule

## Examples of impurity functions

- *Entropy impurity*

$$\phi(p_1, \dots, p_K) = - \sum_j p_j \log p_j,$$

(Where we use the convention  $0 \log 0 = 0$ )

- *Gini impurity*

$$\phi(p_1, \dots, p_K) = \frac{1}{2} \sum_j p_j (1 - p_j)$$

## Splitting Rule

- **Definition 3.** The decrease in impurity
  - Let  $t$  be a node and let  $s$  be split of  $t$  into two child nodes  $t_L$  and  $t_R$

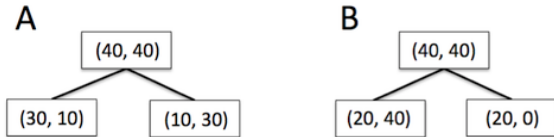
$$\Delta i(s, t) = i(t) - p_L i(t_L) - p_R i(t_R)$$

- Where  $p_L = \frac{p(t_L)}{p(t)}$  and  $p_R = \frac{p(t_R)}{p(t)}$ ,  $p_L + p_R = 1$
- Then  $\Delta i(t) \geq 0$  (see Proposition 4.4. in Breiman et al. (1984))
- Hence the splitting rule at  $t$  is  $s^*$  such that we take the split  $s^*$  among all possible candidate splits that decreases the cost most

$$s^* = \operatorname{argmax}_s \Delta i(s, t)$$

## Splitting Rule

Why does not use misclassification error as an impurity function?



- Recall that  $\Delta i(s, t) = i(t) - p_L i(t_L) - p_R i(t_R)$  and  $s^* = \operatorname{argmax}_s \Delta i(s, t)$
- Misclassification error:  $i_M(t) = 1 - \max_j p_j$ , (where  $p_j = p(j|t)$ )
  - A:  $\Delta i_M(s_A, t) = (1 - \frac{1}{2}) - (\frac{40}{80}) * (1 - \frac{3}{4}) - (\frac{40}{80}) * (1 - \frac{3}{4}) = \frac{1}{4}$
  - B:  $\Delta i_M(s_B, t) = (1 - \frac{1}{2}) - (\frac{60}{80}) * (1 - \frac{4}{6}) - (\frac{20}{80}) * (1 - 1) = \frac{1}{4}$
- Entropy impurity:  $i_E(t) = -\sum_j p_j \log(p_j)$ 
  - A:  $\Delta i_E(s_A, t) = 0.130812$
  - B:  $\Delta i_E(s_B, t) = 0.2157616$

## Stopping Rule

- Stopping rules terminate further splitting
- For example
  - All observations in a node are contained in one group
  - The number of observations in a node is small
  - The decrease of impurity is small
  - The depth of a node is larger than a given number

## Pruning Rule

- A tree with too many nodes will have large prediction error rate for new observations
- It is appropriate to prune away some branch of tree for good prediction error rate
- To determine the size of tree, we estimate prediction error using validation set or cross validation

# Pruning Rule

## Pruning Process

- For a given tree  $T$  and positive number  $\alpha$ , cost-complexity pruning is defined by

$$\text{cost-complexity}(\alpha) = \text{error rate of } T + \alpha|T|$$

(Where  $|T|$  is the number of nodes)

- In general, the larger tree(the larger  $|T|$ ), the smaller error rate (see Proposition 1. in Hyeongin Choi (2017)). But cost-complexity does not decrease as  $|T|$  increases.
- For the grown tree  $T_{max}$ ,  $T(\alpha)$  is a subtree which minimizes cost-complexity( $\alpha$ )
- In general, the larger  $\alpha$ , the smaller  $|T(\alpha)|$

## Pruning Rule

### Pruning Process

- One important property of  $T(\alpha)$  is that if  $\alpha_1 \leq \alpha_2$ , then  $T(\alpha_1) \geq T(\alpha_2)$  which means  $T(\alpha_2)$  is a subtree of  $T(\alpha_1)$
- Let  $T_1 = T(\alpha_1)$ ,  $T_2 = T(\alpha_2), \dots$  then we can get the following sequence of pruned subtrees

$$T_1 \geq T_2 \geq \dots \geq \{t_1\},$$

$$\text{where } 0 = \alpha_1 < \alpha_2 < \dots$$

- For a given  $\alpha$ , we estimate the generalization error of  $T(\alpha)$  by validation set or cross-validation
- Choose  $\alpha^*$  (and corresponding  $T(\alpha^*)$ ) which minimizes the (estimated) generalization error. (see Hyeonjin Choi (2017) for details)



## Estimation Method

- Once a tree is fixed, a prediction at each subregion (terminal node) can be determined from that tree
- Since the estimation is operated at each subregion, it is called a local estimation
- For this local estimation, any estimation method can be obtained. What kind of method to use is dependent on the model assumption.
- For example, for classification, *majority vote* or *logistic regression* is obtained by modeler's probability assumption

## Some Algorithms for Decision Tree

### **CHAID** (CHi-squared Automatic Interaction Detector)

- by J. A. Hartigan 1975
- Employs  $\chi^2$  statistic as impurity
- No pruning process, it stops growing at a certain size

### **CART** (Classification And Regression Tree)

- by Breiman and et al. 1984
- Only binary split is operated
- Cost-complexity pruning is an important unique feature

### **C5.0** (successor of ID3 and C4.5)

- by J. Ross Quinlan 1993
- Multisplit is available
- For categorical input variable, a node splits into the number of categories.

# Advantages and Disadvantages

## **Advantages**

- Easy to interpret and explain
- Trees can be displayed graphically
- Trees can easily handle both continuous and categorical variables without the need to create dummy variables

## **Disadvantages**

- Poor prediction accuracy compared to other models
- When depth is large, not only accuracy but interpretation are bad

## References

- 1 Breiman, L. Friedman, J.H, Olshen, R.A. and Stone, C.J., *Classification And Regression Trees*, Chapman & Hall/CRC (1984)
- 2 Hyeongin Choi (2017), Lecture 9: Classification and Regression Tree (CART), <http://www.math.snu.ac.kr/~hichoi/machinelearning/lecturenotes/CART.pdf>
- 3 Yongdai Kim (2017), Chapter 6. Decision Tree, <https://stat.snu.ac.kr/ydkim/courses/2017-1/addm/Chap6-DecisionTree.pdf>