

Decision Trees, Random Forest, XGBoost

1. How is best split chosen in decision trees/ Random Forests:

a. What metrics are used?

i. Gini Gain or Gini Impurity - Represents quality of feature to create a split

ii. Information Gain - Represents quality of feature to create a split

b. How is Gini Gain or Gini Impurity calculated?

Example: we have 20 balls – 10 blue and 10 green – all balls have same weight but blue balls are larger in diameter than green balls. We want to find which feature is right to classify a ball as B or G.

Impurity in the dataset:

Dataset has equal number of B and G balls. Hence Probab of picking B = 0.5 and Probab of picking G = 0.5

=>Probability of wrongly classifying a ball = Probab(picking a Blue ball)xProbab(classifying as Green Ball) + Probab(picking a Green ball)xProbab(classifying as Blue Ball)

=>Probability of wrongly classifying a ball = $P(\text{Blue}) \times P(\text{Green}) + P(\text{Green}) \times P(\text{Blue})$

=> Probability of wrongly classifying a ball = $0.5 \times 0.5 + 0.5 \times 0.5 = 0.5$

Hence Gini impurity of **Dataset** is 0.5.

0.5 is the worst Gini Impurity you can have.

Imperfect Split:

Now if we use Weight as our feature and split the 20 balls into 2 branches; because all 20 balls have same weight – each branch would end up with 10 B & 10 G balls (most likely).

Now take branch 1:

=>Probability of wrongly classifying a ball = Probab(picking a Blue ball)xProbab(classifying as Green Ball) + Probab(picking a Green ball)xProbab(classifying as Blue Ball)

=>Probability of wrongly classifying a ball = $P(\text{Blue}) \times P(\text{Green}) + P(\text{Green}) \times P(\text{Blue})$

=> Probability of wrongly classifying a ball = $0.5 \times 0.5 + 0.5 \times 0.5 = 0.5$

Hence Gini impurity of Branch 1 is 0.5.

Similarly, we can show that Gini impurity of Branch 2 is also 0.5.

0.5 is the worst Gini Impurity you can have.

Weighted Gini impurity = Fraction of total elements in branch1 * Gini-Impurity of branch1 + Fraction of total elements in branch2 * Gini-Impurity of branch2

Weighted Gini impurity = $(10/20) \times 0.5 + (10/20) \times 0.5 = 0.5$

Gini gain = Impurity before splitting – Weighted Impurity after splitting
= 0.5 (of raw dataset) – 0.5 (weighted) = 0

Hence, we didn't witness any Gini gain => Weight is probably not the right criteria.

Perfect Split:

Suppose if we take weight as feature, then we would correctly able to classify all Blue balls in branch1 and all Green balls in branch 2.

Now take branch 1:

=>Probability of wrongly classifying a ball = Probab(picking a Blue ball)xProbab(classifying as Green Ball) + Probab(picking a Green ball)xProbab(classifying as Blue Ball)

=>Probability of wrongly classifying a ball = $P(\text{Blue}) \times P(\text{Green}) + P(\text{Green}) \times P(\text{Blue})$

=> Probability of wrongly classifying a ball = $1 \times 0 + 0 \times 1 = 0$

Hence Gini impurity of Branch 1 is 0 => there is no impurity

Similarly, we can show that Gini impurity of Branch 2 is also 0.

0 is the best/ideal Gini Impurity you can have.

Weighted Gini impurity = Fraction of total elements in branch1 * Gini-Impurity of branch1 + Fraction of total elements in branch2 * Gini-Impurity of branch2

Weighted Gini impurity = $(10/20) \times 0 + (10/20) \times 0 = 0$

Gini gain = Impurity before splitting – Weighted Impurity after splitting
= 0.5 (of raw dataset) – 0 (weighted) = 0.5

Hence, we witnessed a significant Gini gain => Diameter is probably a very good right criteria for splitting.

Semi-Perfect split:

Suppose 2 G balls have same diameter has B balls. Now If we use diameter as feature to split the dataset, we would have 12 balls (10B + 2G) in branch1 and branch2 has all 8G balls.

Now take branch 1:

=>Probability of wrongly classifying a ball = $P(\text{Blue}) \times P(\text{Green}) + P(\text{Green}) \times P(\text{Blue})$

=> Probability of wrongly classifying a ball = $(10/12) \times (2/12) + (2/12) \times (10/12) = 40/144 = 0.278$

Hence Gini impurity of Branch 1 is 0.278

Now take branch 2:

=>Probability of wrongly classifying a ball = $P(\text{Blue}) \times P(\text{Green}) + P(\text{Green}) \times P(\text{Blue})$

=> Probability of wrongly classifying a ball = $(0/8) \times (8/8) + (8/8) \times (0/8) = 0$

Hence Gini impurity of Branch 2 is 0

Weighted Gini impurity = Fraction of total elements in branch1 * Gini-Impurity of branch1 + Fraction of total elements in branch2 * Gini-Impurity of branch2

Weighted Gini impurity = $(12/20)*0.278 + (8/20)*0 = 0.167$

Gini gain = Impurity before splitting – Weighted Impurity after splitting
 = 0.5 (of raw dataset) – 0.167 (weighted) = 0.333

Hence, we witnessed some Gini gain => Diameter may be a good criteria for splitting.

c. What is Information Gain?

It is similar to Gini Impurity. It uses entropy concept to quantify randomness in the split.

$$E = - \sum_i^C p_i \log_2 p_i$$

So, continuing the same example from above for Semi-perfect split.

Entropy in the Dataset:

As dataset has 10B and 10G = $P(B) = 0.5$, $P(G) = 0.5$

Entropy = $-(0.5*\log_2(0.5) + 0.5*\log_2(0.5)) = 1$

Entropy of 1 is the worst and Entropy of 0 is the best

Now take branch 1:

Branch 1 has 12 balls – 10B + 2G; $P(B) = 10/12=0.83$ & $P(G) = 2/12=0.167$

Entropy of Branch1 = $-(0.83*\log_2(0.83) + 0.167*\log_2(0.167)) = 0.65$

Hence Entropy of Branch 1 is 0.65

Now take branch 2:

Branch 2 has 8 balls – 0B + 8G; $P(B) = 0$ & $P(G) = 1$

Entropy of Branch2 = $-(0*\log_2(0) + 1*\log_2(1)) = 0$

Hence Entropy of Branch 2 is 0

Weighted Entropy = Fraction of total elements in branch1 * Entropy of branch1 + Fraction of total elements in branch2 * Entropy of branch2

Weighted Entropy = $(12/20)*0.65 + (8/20)*0 = 0.39$

Information gain = Entropy before splitting – Entropy after splitting
 = 1 (of raw dataset) – 0.39 (weighted) = 0.61

Hence, we witnessed some Information gain => Diameter may be a good criteria for splitting.

2. How is feature importance calculated?

- a. It is calculated by measuring the total decrease in node impurity averaged across all the trees. (Refer Q1 for more details on these metrics are calculated)
- 3. Boosting vs Bagging
 - Bagging – **Bootstrap Aggregating**
 - a. What is bootstrapping
 - i. It is a sampling procedure to create subset of data. Particularly, in RF, we create m subsets with all n examples with replacement.
 - b. Differences
 - i. Boosting trains different machine learning models one after another (**sequentially**) to get the final result, while bagging trains them in **parallel**.
 - ii. We know, Error in a model = Bias² + Variance + Noise. **Bagging reduces Variance while Boosting reduces Bias**. (We know Decision Trees have high variance. So, we perform bagging to reduce variance)
 - iii. Each tree in Bagging has same weight, while Boosting involves a set of weights which it assigns to each tree. So, at prediction stage, in Bagging – final prediction is average of all trees (or majority vote) while for Boosting it is weighted average of all trees predictions
 - iv. In **Bagging** different training data subsets are randomly drawn with replacement from the entire training dataset. In **Boosting** every new subsets contains the elements that were misclassified by previous models.
 - v. If the classifier is unstable (high variance), then we should apply **Bagging**. If the classifier is stable and simple (high bias) then we should apply **Boosting**.
 - vi. **Bagging** is extended to Random forest model while **Boosting** is extended to **Gradient boosting**.
 - c. When to choose Boosting and when to choose Bagging?

Normally their selection depends on problem at hand.

 - i. Bagging and Boosting decrease the variance of your single estimate as they combine several estimates from different models. So, the result may be a model with higher stability.
 - ii. If the problem is that the single model gets a very low performance (aka underfitting, high bias), Bagging will rarely get a better bias. However, Boosting could generate a combined model with lower errors as it optimizes the advantages and reduces pitfalls of the single model.
 - iii. By contrast, if the difficulty of the single model is over-fitting, then Bagging is the best option. Boosting for its part doesn't help to avoid over-fitting.
 - iv. In fact, this technique is faced with this problem itself. For this reason, Bagging is effective more often than Boosting.
 - d. What are similarities between Bagging and Boosting
 - i. Both are ensemble learning method
 - ii. Both makes Decision Trees stable
 - iii. Both are good at reducing variance
 - e. What is gradient boosting?
- 4. Can Boosting reduce bias?

- a. Yes. This is why they are popular.
- 5. Difference between GBDT and XGBoost. How do you parameter tune them?
- 6. Adaboost
- 7. More resources:

<https://victorzhou.com/blog/gini-impurity/>

<https://victorzhou.com/blog/information-gain/>

<https://www.kaggle.com/code/prashant111/bagging-vs-boosting>