

→ Decision Trees

- What is a decision tree

A decision tree is a supervised machine learning algorithm that models decisions using a tree-like structure, where:

- Internal nodes represent feature-based conditions
- Branches represent decision rules
- Leaf node represent final predictions.

It can be used for classification and regression.

- Purpose (When)

To make predictions by recursively splitting data into homogeneous subsets, mimicking human decision-making.

- When relationships are non-linear
- When interpretability is important
- When mixed data types exist (numerical + categorical)

- Why it is used

- Easy to understand and visualize
- Handles non-linear patterns naturally.
- No need for feature scaling
- Works with missing values

- How it works

- Start with entire dataset at the root node
- Select the best feature and split point
- Split data into child nodes
- ~~Stop when stopping~~ Repeat recursively for each node
- Stop when stopping criteria are met.
- Assign prediction at leaf nodes.

- Splitting Criteria

• For Classification:

- Gini Impurity : $Gini = 1 - \sum p_i^2$

- Entropy : $Entropy = - \sum p_i \log_2(p_i)$

- Information Gain

$$IG = Entropy(\text{parent}) - \sum \frac{n_k}{n} Entropy(\text{child}_k)$$

• For Regression:

- MSE

- MAE

- Variance Reduction

- Types

- Based on Task: Classification Tree

- Regression Tree

- Based on Algorithm: ID3 (Entropy)

- C4.5 (Entropy + Pruning)

- CART (Gini/MSE)

- Formula

- Prediction: Classification \rightarrow majority class in leaf

- Regression \rightarrow mean/median value in leaf

- Variance Reduction:

$$VR = Var(\text{parent}) - \sum \frac{n_k}{n} Var(\text{child}_k)$$

- Technical Details

- Assumptions: No strict statistical assumptions

- Works without linearity or normality

- Stopping Criteria: Max depth reached

- Minimum samples per node

- No further impurity reduction

- Tree Growth: Top-down, greedy algorithm

- Locally optimal splits (not global)

- Parameters

(Learned)

• Model Parameters: Tree structure

Split thresholds

Leaf values

• Hyperparameters (Tuning): max_depth

min_sample_split

min_sample_leaf

max_features

criterion (gini, entropy, mse)

- Pros

• Highly interpretable

• Handles non-linear data

• Works with mixed feature types

• No feature scaling required

• Fast inference

- Cons

• Prone to overfitting

• Unstable to small data changes

• Greedy (locally optimal splits)

• Poor generalization alone

• Biased toward dominant features

- Real-World Applications (Where)

• Business: Customer segmentation, churn prediction

• Finance: Credit risk analysis, Loan approval systems

• Healthcare: Disease diagnosis, Treatment decision support

• Engineering: Fault detection systems

• ML Systems: Base learner in Random Forest, XGBoost,

LightGBM.

→ Random Forest

- What is Random Forest

Random forest is a supervised ensemble machine learning algorithm that builds multiple decision trees and combines their outputs to produce a more accurate and robust prediction.

It works for both classification and regression.

- Purpose (When)

To improve prediction accuracy and generalization by reducing the overfitting and instability of individual tree.

- Why it is used

- Reduces overfitting of decision trees
- Handles non-linear relationships
- Works well with high-dimensional data
- Robust to noise and outliers
- Provides feature importance

- How it works

- Core ideas: Bagging (Bootstrap Aggregation)
Random Feature Selection

- Algorithm:
 - Draw multiple bootstrap samples from ^{dataset}
 - Train a decision tree on each sample
 - At every split, select a random subset of ^{features}
 - Grow trees independently and in parallel
 - Aggregate predictions

Classification → majority vote

Regression → mean of predictions

- Formula

Classification Prediction: $\hat{y} = \text{model}\{T_1(x), T_2(x), \dots, T_n(x)\}$

Regression Prediction: $\hat{y} = \frac{1}{n} \sum_{i=1}^n T_i(x)$

Where T_i is the prediction of the i -th tree

- Technical Details

- Ensemble Type: Bagging based ensemble
- Tree properties: Trees are deep and unpruned
High variance, low bias individually
Ensemble reduces variance
- Randomness Source: Data sampling (bootstrap)
Feature sampling at splits

- Parameters

- Model Parameters (Learned): Individual tree structures
Split thresholds
Leaf predictions
- Hyperparameters (Most important):
 - $n_estimators$: no. of trees
 - max_depth : max depth of each tree
 - $min_samples_split$: min samples to split
 - $min_samples_leaf$: min samples at leaf
 - $max_features$: features considered per split
 - $bootstrap$: Whether sampling is with replacement
 - $criterion$: Gini, Entropy, MSE

- Feature Importance

Random Forest estimates feature importance using:

- Mean Decrease in Impurity (MDI)
- Permutation Importance

- Assumptions

- No strict statistical assumptions
- Assumes trees are weakly correlated

- Pros

- High accuracy
- Handles non-linear data well
- Resistant to overfitting
- Works with missing values
- Scales well with large datasets

- Cons

- Less interpretable than single trees
- Computationally expensive
- Slower inference
- Large memory usage
- Feature importance can be biased

- Real World Application (Where)

- Finance: Credit scoring
Fraud detection
- Healthcare: Disease prediction
Risk stratification
- Marketing: Customer churn prediction
Recommendation systems
- Engineering: Fault detection
Predictive maintenance
- ML Pipelines: Baseline model for tabular data
Feature selection