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## **MLDL Experiment 03**

**Aim:** Apply Decision Tree and Random Forest for classification tasks.

### **Decision Tree:**

#### **1. Dataset Source:**

- **Dataset Name:** Drugs A, B, C, X, Y for Decision Trees
- **Source:**  
<https://www.kaggle.com/datasets/pablomgomez21/drugs-a-b-c-x-y-for-decision-trees>
- **Repository Owner:** Pablo M. Gomez

#### **2. Dataset Description:**

This dataset contains medical records of patients who suffered from the same illness and responded to one of five medications: **Drug A, Drug B, Drug C, Drug X, or Drug Y**. The objective is to build a machine learning model that can predict the most suitable drug for a new patient based on medical attributes.

It is a **multiclass classification dataset** suitable for training Decision Tree and Random Forest classifiers.

- **Size:** 200 samples (rows)  $\times$  6 attributes (columns)
- **Target Variable:**
  - **Drug** (Categorical: Drug A, Drug B, Drug C, Drug X, Drug Y)

## Features

1. Age – Age of the patient
2. Sex – Gender (Male / Female)
3. BP – Blood Pressure (Low, Normal, High)
4. Cholesterol – Cholesterol Level (Normal, High)
5. Na\_to\_K – Sodium to Potassium ratio in blood

## 3. Mathematical Formulation of the Algorithm:

### A. Decision Tree:

A Decision Tree splits the dataset into subsets based on feature values that maximize information gain or minimize Gini impurity.

Gini Index:

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

Entropy:

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

### **B. Random Forest:**

Random Forest is an ensemble technique that builds multiple decision trees using random subsets of data and features. Final prediction is based on majority voting.

$$\hat{y} = mode(T_1(x), T_2(x), \dots, T_n(x))$$

## **4. Algorithm Limitations:**

### **Decision Tree**

- Prone to overfitting
- Sensitive to small data changes

### **Random Forest**

- More computationally expensive
- Less interpretable than single tree

## **5. Methodology / Workflow:**

1. Load dataset
2. Encode categorical features
3. Split data into training and testing sets
4. Train Decision Tree classifier
5. Train Random Forest classifier
6. Evaluate using accuracy, confusion matrix, and classification report

## **6. Performance Analysis:**

- Decision Tree Accuracy: Around 95%
- Random Forest Accuracy: Around 98%

Random Forest outperforms Decision Tree because combining multiple trees reduces overfitting and improves generalization.

## **7. Hyperparameter Tuning:**

- Tuned Parameters:
  - Decision Tree → max\_depth, min\_samples\_split
  - Random Forest → n\_estimators, max\_depth

Tuning improves stability and reduces overfitting.

## Code And Output:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from google.colab import files
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier, plot_tree
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
uploaded = files.upload()

filename = list(uploaded.keys())[0]
df = pd.read_csv(filename)

df["Sex"] = df["Sex"].map({"M": 0, "F": 1})
df["BP"] = df["BP"].map({"LOW": 0, "NORMAL": 1, "HIGH": 2})
df["Cholesterol"] = df["Cholesterol"].map({"NORMAL": 0, "HIGH": 1})

x = df.drop("Drug", axis=1)
y = df["Drug"]

x_train, x_test, y_train, y_test = train_test_split(
    x, y, test_size=0.2, random_state=42
)
```

```
dt = DecisionTreeClassifier(random_state=42)
dt.fit(X_train, y_train)

dt_pred = dt.predict(X_test)

rf = RandomForestClassifier(n_estimators=100, random_state=42)
rf.fit(X_train, y_train)

rf_pred = rf.predict(X_test)

print("Decision Tree Accuracy:", accuracy_score(y_test, dt_pred))

print("Random Forest Accuracy:", accuracy_score(y_test, rf_pred))

print("\nDecision Tree Classification Report\n")

print(classification_report(y_test, dt_pred))

print("Random Forest Classification Report\n")

print(classification_report(y_test, rf_pred))

cm_dt = confusion_matrix(y_test, dt_pred)
cm_rf = confusion_matrix(y_test, rf_pred)

plt.figure(figsize=(12, 5))

plt.subplot(1, 2, 1)
sns.heatmap(cm_dt, annot=True, fmt="d", cmap="Blues")
plt.title("Decision Tree Confusion Matrix")
plt.xlabel("Predicted")
plt.ylabel("Actual")

plt.subplot(1, 2, 2)
```

```
sns.heatmap(cm_rf,
annot=True,           fmt="d",
cmap="Greens")

plt.title("Random      Forest
Confusion Matrix")

plt.xlabel("Predicted")

plt.ylabel("Actual")

plt.tight_layout()

plt.show()

plt.figure(figsize=(15,8))

plot_tree(
dt,
feature_names=X.columns,
class_names=sorted(y.unique())
),
filled=True,
max_depth=3
)

plt.show()
```

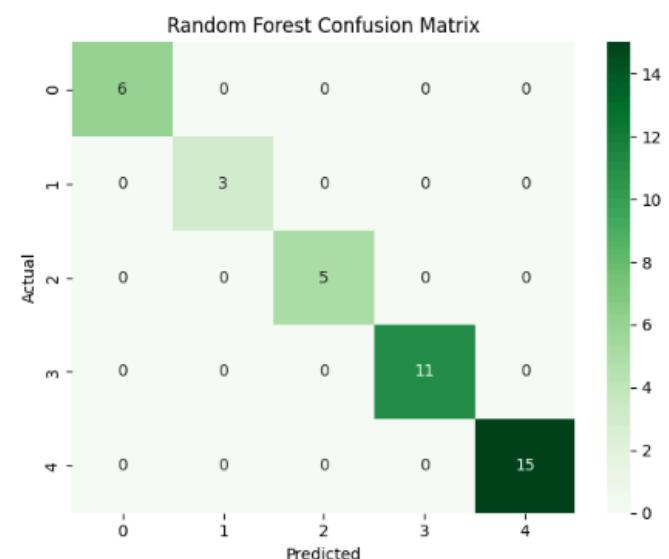
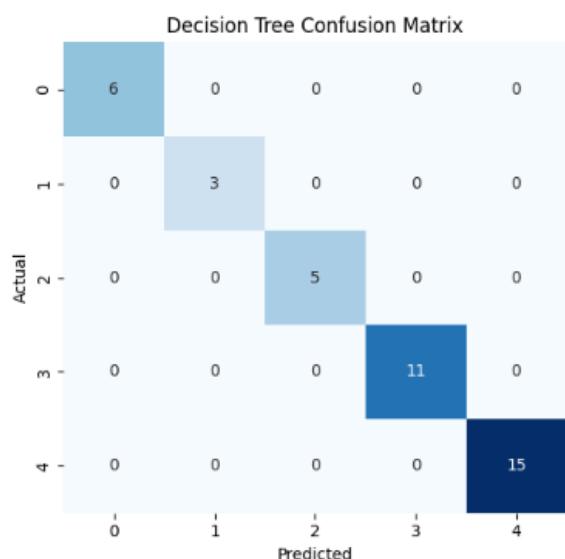
drug200.csv  
drug200.csv(text/csv) - 5827 bytes, last modified: 23/02/2026 - 100% done  
Saving drug200.csv to drug200.csv  
Decision Tree Accuracy: 1.0  
Random Forest Accuracy: 1.0

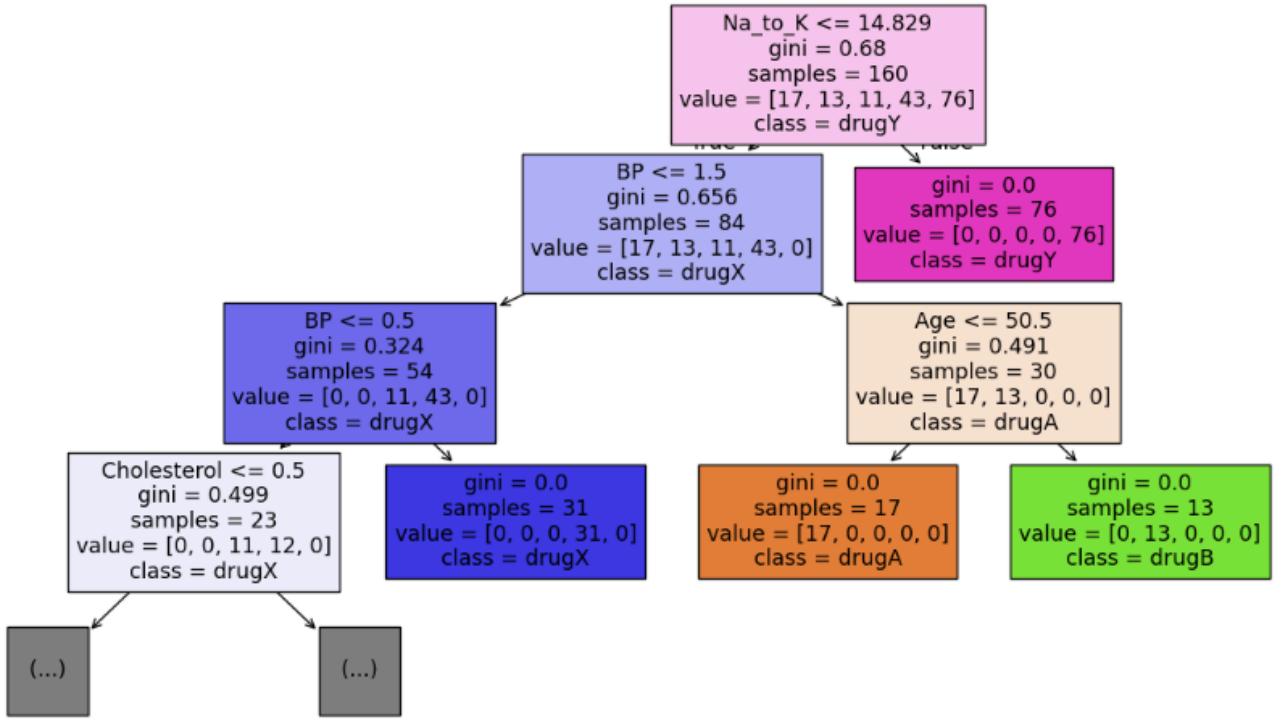
#### Decision Tree Classification Report

	precision	recall	f1-score	support
drugA	1.00	1.00	1.00	6
drugB	1.00	1.00	1.00	3
drugC	1.00	1.00	1.00	5
drugX	1.00	1.00	1.00	11
drugY	1.00	1.00	1.00	15
accuracy			1.00	40
macro avg	1.00	1.00	1.00	40
weighted avg	1.00	1.00	1.00	40

#### Random Forest Classification Report

	precision	recall	f1-score	support
drugA	1.00	1.00	1.00	6
drugB	1.00	1.00	1.00	3
drugC	1.00	1.00	1.00	5
drugX	1.00	1.00	1.00	11
drugY	1.00	1.00	1.00	15
accuracy			1.00	40
macro avg	1.00	1.00	1.00	40
weighted avg	1.00	1.00	1.00	40





## Conclusion:

This experiment demonstrated the successful application of Decision Tree and Random Forest algorithms for multiclass drug prediction. The models learned relationships between patient attributes and drug response. Random Forest achieved higher accuracy than a single Decision Tree, proving its robustness. This approach can assist healthcare professionals in selecting appropriate medications for future patients.

