

MACHINE LEARNING

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Practical 1: Data Pre-processing and Exploration

1a. Load a CSV dataset. Handle missing values, inconsistent formatting, and outliers.

Code :

1. Import Libraries

Import necessary libraries

```
import pandas as pd import numpy as np  
import seaborn as sns  
import matplotlib.pyplot as plt
```

2. Load the Dataset

Load the Titanic dataset from a URL

```
url="https://raw.githubusercontent.com/datasets/master/titanic.csv" data =  
pd.read_csv(url)
```

Display the first few rows

```
print(data.head())
```

3. Handle Missing Values

Check for missing values

```
print("Missing values in each column:")
```

```
print(data.isnull().sum())
```

Fill missing values in 'Age' with the mean

```
data['Age'].fillna(data['Age'].mean(), inplace=True)
```

Fill missing values in 'Embarked' with the most common value

```
data['Embarked'].fillna(data['Embarked'].mode()[0], inplace=True)
```

Drop rows where 'Cabin' is missing (too many NaNs)

```
data.drop(columns=['Cabin'], inplace=True)
```

Verify missing values are handled

```
print("\nAfter handling missing values:")
print(data.isnull().sum())
```

4. Fix Inconsistent Formatting

```
# Fix inconsistent formatting in the 'Sex' column
```

```
data['Sex'] = data['Sex'].str.lower().str.strip()
```

```
# Verify unique values
```

```
print("\nUnique values in 'Sex' column after formatting:")
```

```
print(data['Sex'].unique())
```

```
5. Detect and Handle Outliers # Boxplot for the 'Fare' column sns.boxplot(data['Fare'],
color='skyblue') plt.title('Boxplot of Fare') plt.show()
```

```
# Detect outliers using the IQR method
```

```
Q1 = data['Fare'].quantile(0.25)
Q3 = data['Fare'].quantile(0.75)
IQR = Q3 - Q1
lower_bound =
Q1 - 1.5 * IQR
upper_bound =
Q3 + 1.5 * IQR
```

```
# Capping outliers
```

```
data['Fare'] = np.where(data['Fare'] > upper_bound, upper_bound, np.where(data['Fare'] <
lower_bound, lower_bound, data['Fare']))
```

```
# Verify with an updated boxplot
```

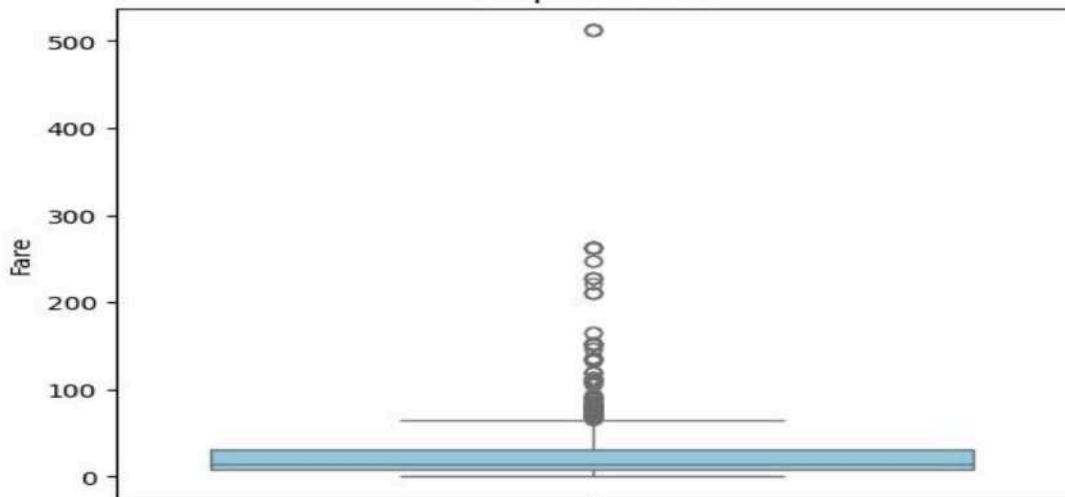
```
sns.boxplot(data['Fare'], color='lightgreen')
plt.title('Boxplot of Fare (After Handling Outliers)')
plt.show()
```

```
6. Save the Cleaned Dataset # Save the cleaned dataset
```

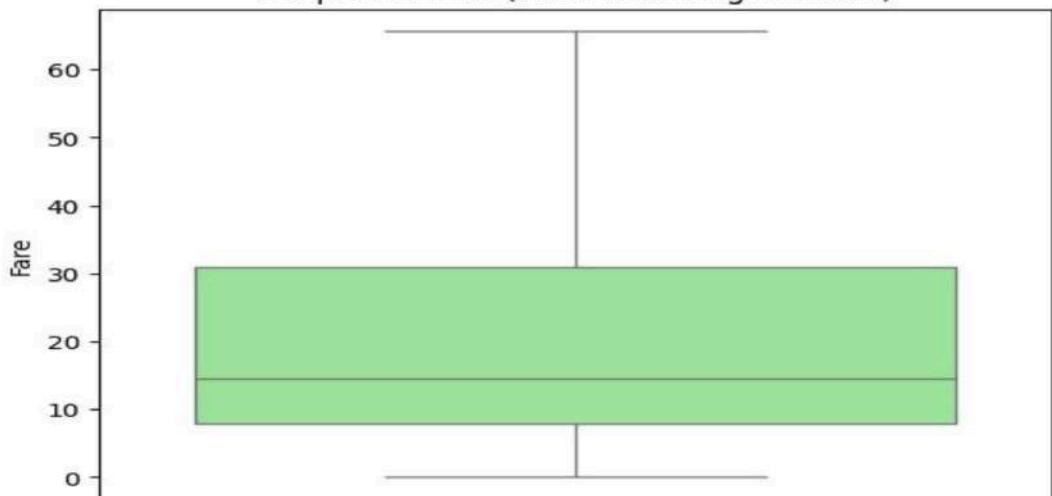
```
data.to_csv('cleaned_titanic.csv', index=False)
print("\nCleaned dataset saved as 'cleaned_titanic.csv'") .
```

Output :

Boxplot of Fare



Boxplot of Fare (After Handling Outliers)



1b. Load a dataset, calculate descriptive summary statistics, create visualizations using different graphs, and identify potential features and target variables Note:

Explore Univariate and Bivariate graphs (Matplotlib) and Seaborn for visualization

Code :

1. Import Necessary Libraries # Import required libraries

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import seaborn as sns
```

2. Load the Dataset

Load the dataset from the URL

```
url = "https://raw.githubusercontent.com/mwaskom/seaborn-data/master/iris.csv"  
data = pd.read_csv(url)  
  
# Display the first few rows  
  
print("First 5 rows of the dataset:")  
print(data.head())
```

3. Calculate Descriptive Summary Statistics # Dataset information

```
print("\nDataset Info:")  
print(data.info())  
  
# Summary statistics for numerical columns  
  
print("\nDescriptive Statistics for Numerical Columns:")  
print(data.describe())  
  
# Check unique values for categorical columns  
  
print("\nUnique values in 'species' column:")  
print(data['species'].value_counts())
```

4. Univariate Analysis

Histograms for numerical columns

```
data.hist(figsize=(10,8), color='skyblue', edgecolor='black')
plt.suptitle("Histograms of Numerical Features")
plt.show()

# Bar plot for 'species' column
sns.countplot(x='species', data=data, palette='pastel')
plt.title("Count of Each Species") plt.show()
```

5. Bivariate Analysis

Scatter plot for two features

```
plt.figure(figsize=(8, 6))

plt.scatter(data['sepal_length'], data['sepal_width'], alpha=0.7, c='blue')
plt.title("Sepal Length vs Sepal Width")
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.show()
```

Pairplot to visualize relationships between features

```
sns.pairplot(data, hue='species', palette='husl', diag_kind='kde')
plt.suptitle("Pairplot of Features by Species", y=1.02)
plt.show()
```

```
# Boxplot for petal_length across species sns.boxplot(x='species',
y='petal_length', data=data, palette='Set3')

plt.title("Boxplot of Petal Length by Species")
plt.show()
```

6. Identify Potential Features and Target Variables

Separate features and target

```
features = data.drop(columns=['species'])
```

Drop the target

```
column_target = data['species']
```

Target variable

```
print("\nFeatures:")
```

```
print(features.head())
```

```
print("\nTarget:")
```

```
print(target.head())
```

```
# Visualize target distribution
sns.countplot(x=target, palette='viridis')
plt.title("Target Variable Distribution")

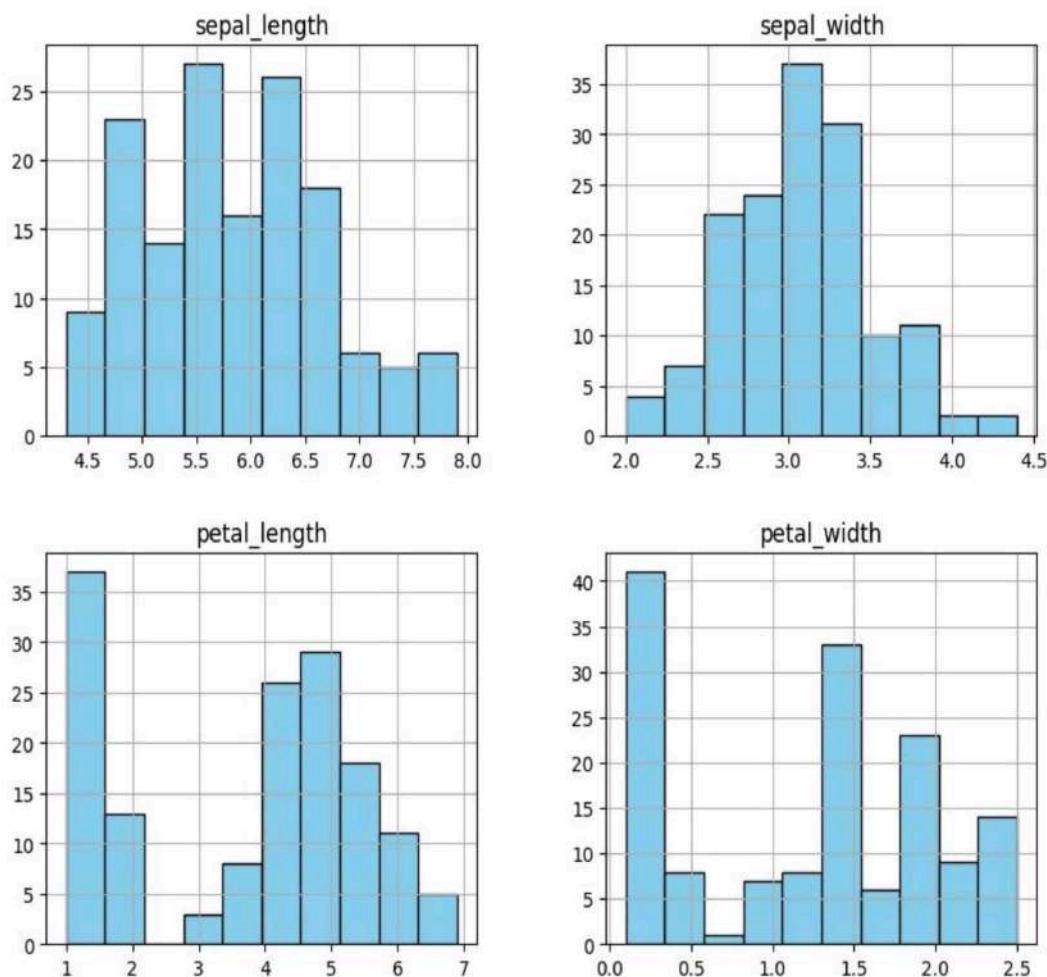
plt.show()
```

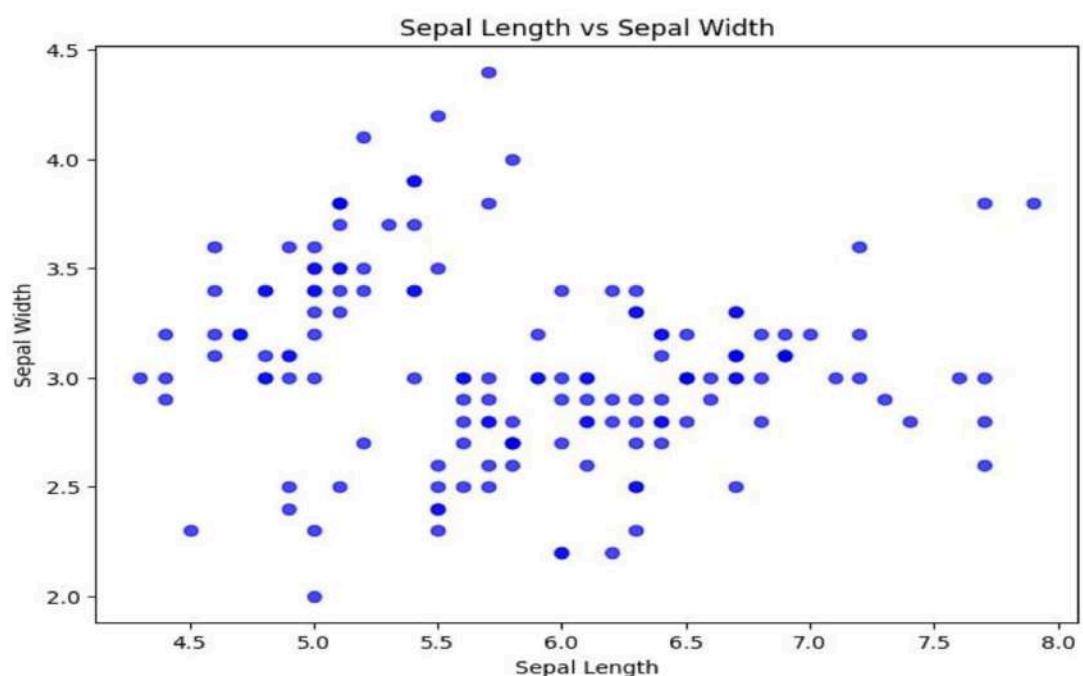
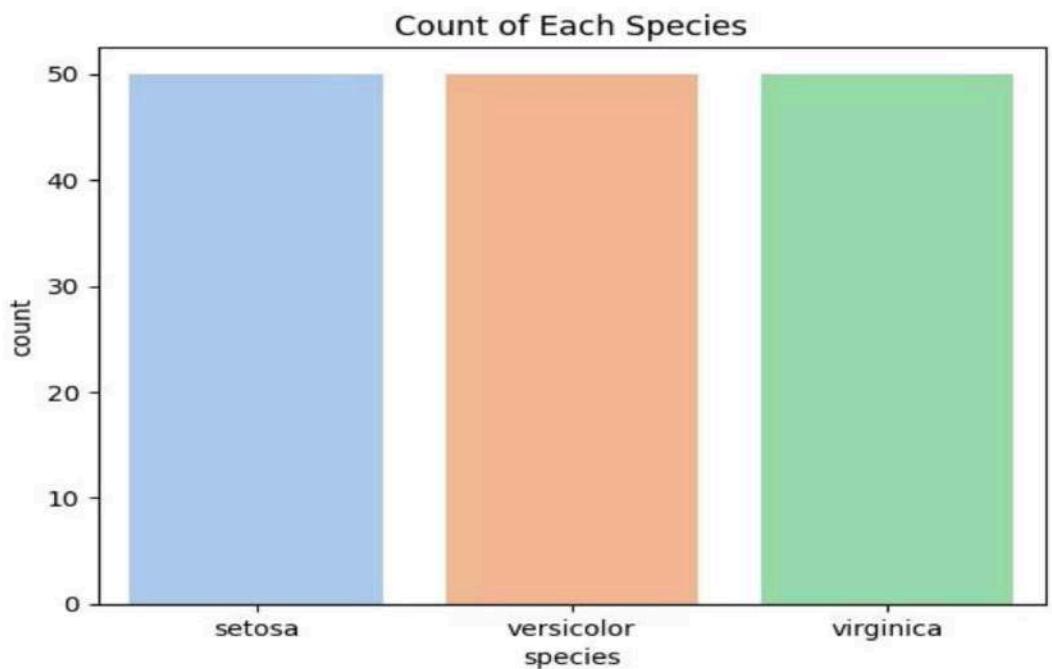
7. Save the Cleaned and Processed Dataset

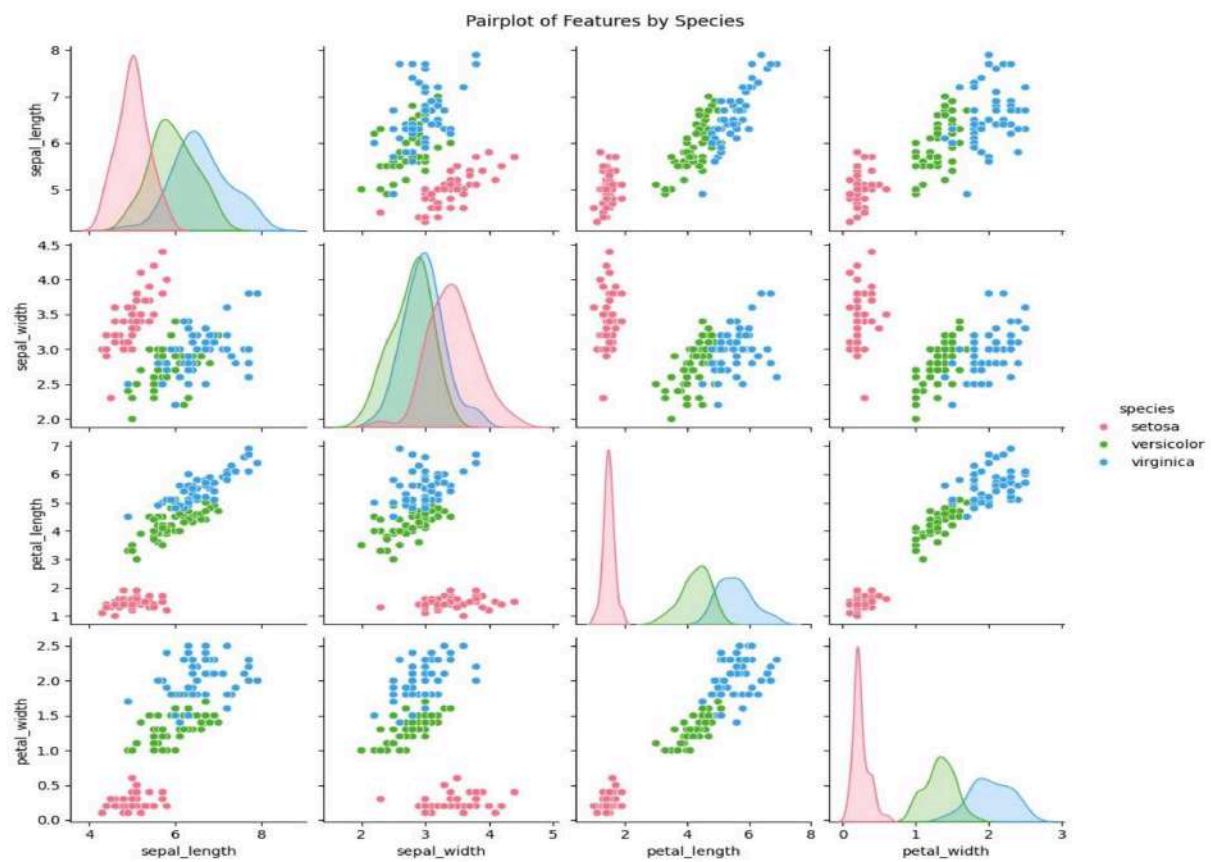
```
# Save the dataset
data.to_csv('processed_iris.csv', index=False)
print("\nProcessed dataset saved as 'processed_iris.csv'")
```

Output :

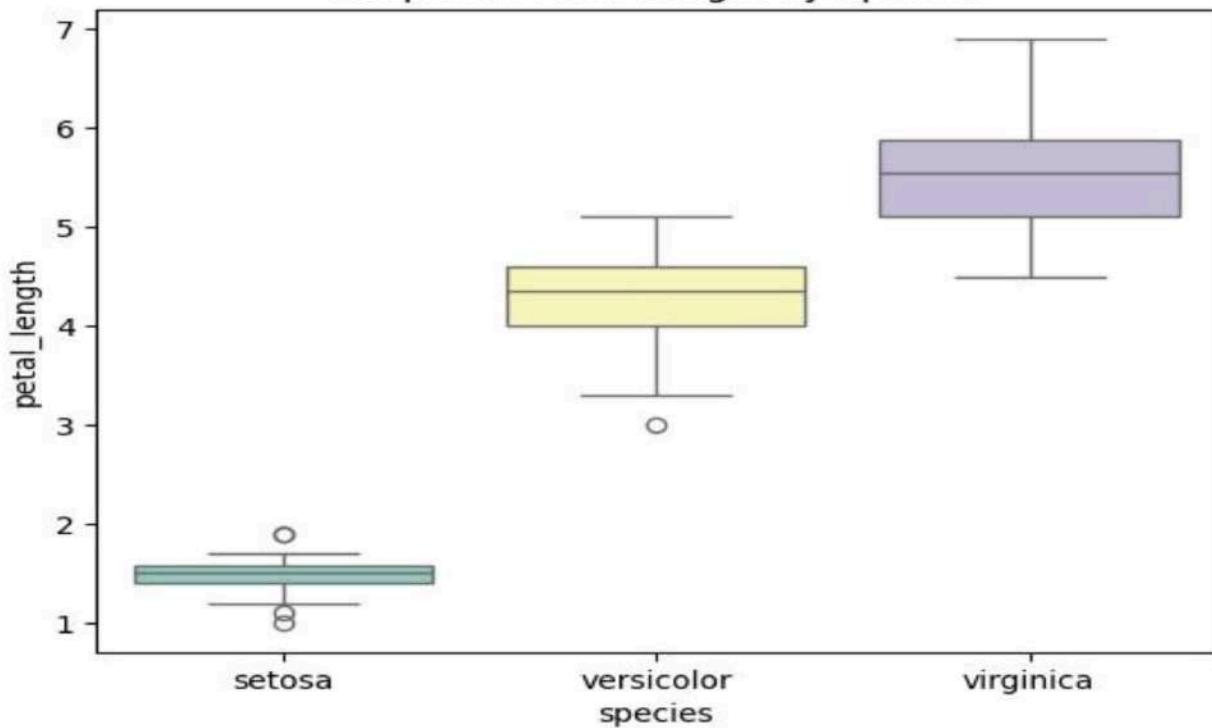
Histograms of Numerical Features







Boxplot of Petal Length by Species



1c. Create or Explore datasets to use all pre-processing routines like label encoding, scaling, and binarization.

Code :

1. Import Necessary Libraries # Import required libraries

```
import pandas as pd  
import numpy as np from sklearn.preprocessing  
import LabelEncoder, MinMaxScaler, StandardScaler, Binarizer
```

2. Create or Load a Dataset # Create a sample dataset

```
data = pd.DataFrame({  
    'Category': ['A', 'B', 'C', 'A', 'B', 'C'],  
    # Categorical variable  
    'Age': [23, 45, 31, 22, 35, 30],  
    # Numerical variable  
    'Income': [50000, 60000, 70000, 80000, 90000, 100000],  
    # Numerical variable 'Has_Car':  
    ['Yes', 'No', 'Yes', 'No', 'Yes', 'No']  
    # Binary categorical variable })
```

Display the dataset

```
print("Sample Dataset:")  
print(data)
```

3. Apply Pre-Processing Routines

```
# Label Encoding for 'Category' column  
label_encoder = LabelEncoder()  
data['Category_Encoded'] =  
label_encoder.fit_transform(data['Category'])  
# Label Encoding for binary column 'Has_Car'
```

```

data['Has_Car_Encoded'] =
label_encoder.fit_transform(data['Has_Car']) print("\nAfter Label
Encoding:")
print(data)

# Min-Max Scaling for 'Income'
min_max_scaler = MinMaxScaler()
data['Income_MinMax'] = min_max_scaler.fit_transform(data[['Income']])

# Standard Scaling for 'Age'
standard_scaler = StandardScaler()
data['Age_Standardized'] =
standard_scaler.fit_transform(data[['Age']]) print("\nAfter Scaling:")
print(data)

# Binarization for 'Income' with a threshold of 75,000
binarizer = Binarizer(threshold=75000)
data['Income_Binary'] =
binarizer.fit_transform(data[['Income']]) print("\nAfter
Binarization:")
print(data)

```

4. Save the Processed Dataset

```

# Save the processed dataset
data.to_csv('processed_data.csv', index=False)
print("\nProcessed dataset saved as
'processed_data.csv'")

```

Output :

Sample Dataset:					
	Category	Age	Income	Has_Car	
0	A	23	50000	Yes	
1	B	45	60000	No	
2	C	31	70000	Yes	
3	A	22	80000	No	
4	B	35	90000	Yes	
5	C	30	100000	No	



After Label Encoding:

	Category	Age	Income	Has_Car	Category_Encoded	Has_Car_Encoded
0	A	23	50000	Yes	0	1
1	B	45	60000	No	1	0
2	C	31	70000	Yes	2	1
3	A	22	80000	No	0	0
4	B	35	90000	Yes	1	1
5	C	30	100000	No	2	0



After Scaling:

	Category	Age	Income	Has_Car	Category_Encoded	Has_Car_Encoded	\
0	A	23	50000	Yes	0	1	
1	B	45	60000	No	1	0	
2	C	31	70000	Yes	2	1	
3	A	22	80000	No	0	0	
4	B	35	90000	Yes	1	1	
5	C	30	100000	No	2	0	

	Income_MinMax	Age_Standardized	
0	0.0	-1.035676	
1	0.2	1.812434	
2	0.4	0.000000	
3	0.6	-1.165136	
4	0.8	0.517838	
5	1.0	-0.129460	



After Binarization:

	Category	Age	Income	Has_Car	Category_Encoded	Has_Car_Encoded	\
0	A	23	50000	Yes	0	1	
1	B	45	60000	No	1	0	
2	C	31	70000	Yes	2	1	
3	A	22	80000	No	0	0	
4	B	35	90000	Yes	1	1	
5	C	30	100000	No	2	0	

	Income_MinMax	Age_Standardized	Income_Binary
0	0.0	-1.035676	0
1	0.2	1.812434	0
2	0.4	0.000000	0
3	0.6	-1.165136	1
4	0.8	0.517838	1
5	1.0	-0.129460	1

Practical 2 : Testing Hypothesis

Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples. Read the training data from a. CSV file and generate the final specific hypothesis. (Create your dataset)

CODE :

```
import pandas as pd

# Step 1: Create the Dataset and Load It

data = {'Outlook': ['Sunny', 'Sunny', 'Overcast', 'Rainy', 'Rainy', 'Rainy', 'Overcast', 'Sunny',
'Sunny', 'Rainy'],
'Temperature': ['Hot', 'Hot', 'Hot', 'Mild', 'Cool', 'Cool', 'Mild', 'Cool', 'Mild'],
'Humidity': ['High', 'High', 'High', 'High', 'Normal', 'Normal', 'Normal', 'High', 'Normal',
'Normal'],
'Wind': ['Weak', 'Strong', 'Weak', 'Weak', 'Weak', 'Strong', 'Strong', 'Weak', 'Weak', 'Weak'],
'PlayTennis': ['No', 'No', 'Yes', 'Yes', 'Yes', 'No', 'Yes', 'No', 'Yes', 'Yes']
}

}
```

Load dataset into a pandas DataFrame

```
df = pd.DataFrame(data)

# Step 2: Implementing the FIND-S Algorithm

def find_s_algorithm(data):

    # Get the positive examples (PlayTennis = 'Yes')
    positive_examples = data[data['PlayTennis'] == 'Yes']

    # Initialize hypothesis with the first positive example (most specific)
    hypothesis = positive_examples.iloc[0].drop('PlayTennis')

    # Loop through the rest of the positive examples and generalize the
    # hypothesis

    for index, row in positive_examples.iterrows():

        for feature in hypothesis.index:

            if hypothesis[feature] != row[feature]:
                hypothesis[feature] = '?'
```

```
    return hypothesis
```

Step 3: Apply FIND-S to the dataset

```
hypothesis = find_s_algorithm(df)

# Display the final specific hypothesis

print("The most specific hypothesis is:")

print(hypothesis)
```

Output :

```
→ Dataset:
   Sky Temperature Humidity      Wind Water Forecast Condition
0  Sunny          Warm  Normal  Strong  Warm    Same     Yes
1  Sunny          Cold   High   Strong  Warm    Same     No
2  Rainy          Warm   High   Weak   Cool   Change   No
3  Sunny          Warm  Normal  Strong  Warm    Same     Yes
4  Rainy          Cold  Normal   Weak   Cool   Change   No
```

```
→
Loaded Dataset:
   Sky Temperature Humidity      Wind Water Forecast Condition
0  Sunny          Warm  Normal  Strong  Warm    Same     Yes
1  Sunny          Cold   High   Strong  Warm    Same     No
2  Rainy          Warm   High   Weak   Cool   Change   No
3  Sunny          Warm  Normal  Strong  Warm    Same     Yes
4  Rainy          Cold  Normal   Weak   Cool   Change   No
```

```
→
Final Specific Hypothesis:
['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same']
```

Practical 3 : Linear Models

3a. Simple Linear Regression

Fit a linear regression model on a dataset. Interpret coefficients, make predictions, and evaluate performance using metrics like R-squared and MSE

Code :

Step 1: Import Libraries # Import required libraries

```
import pandas as pd  
  
import numpy as np  
  
import matplotlib.pyplot as plt  
  
from sklearn.model_selection import train_test_split  
  
from sklearn.linear_model import LinearRegression  
  
from sklearn.metrics import mean_squared_error, r2_score
```

Step 2: Create a Dataset and Save as CSV

Create a sample dataset

```
data = {  
  
    'House_Size': [750, 800, 850, 900, 1000, 1100, 1200, 1300, 1400, 1500],  
  
    'Price': [150000, 160000, 165000, 170000, 180000, 190000, 200000, 210000, 220000,  
    230000]  
}
```

Convert the dataset into a DataFrame

```
df = pd.DataFrame(data)
```

Save to CSV file

```
df.to_csv('house_prices.csv',  
index=False)
```

Display the dataset

```
print("Dataset:")  
print(df)
```

Step 3: Load the Dataset

```
# Load the dataset  
dataset = pd.read_csv('house_prices.csv')  
# Display the first few  
rows      print("\nLoaded  
Dataset:")  
print(dataset.head())
```

Step 4: Split the Dataset into Training and Test Sets

```
# Features and target variable  
X = dataset[['House_Size']] # Feature: House size  
y = dataset['Price']      # Target: Price  
  
# Split data into training and testing sets (80% train, 20% test)  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)  
print("\nTraining and Testing Data Sizes:")  
print("Training Data Size:", X_train.shape[0])  
print("Testing Data Size:", X_test.shape[0])
```

Step 5: Fit a Linear Regression Model

```
# Initialize and fit the linear regression model  
model = LinearRegression()  
model.fit(X_train, y_train)  
  
# Display the coefficients  
print("\nModel Coefficients:")  
print("Slope (m):", model.coef_[0])  
print("Intercept (b):", model.intercept_)
```

Step 6: Make Predictions

```
# Predict on the test set  
y_pred = model.predict(X_test)  
# Display predictions  
print("\nPredictions on Test Data:")  
print("Actual Prices:", y_test.values)  
print("Predicted Prices:", y_pred)
```

Step 7: Evaluate the Model

```
# Calculate evaluation metrics  
mse = mean_squared_error(y_test, y_pred) r2 = r2_score(y_test, y_pred)
```

Display metrics

```
print("\nModel Performance Metrics:")
print("Mean Squared Error (MSE):", mse)
print("R-squared (R2):", r2)
```

Step 8: Visualize the Results # Scatter plot of the training data

```
plt.scatter(X_train, y_train, color='blue', label='Training Data')
```

Plot the regression line

```
plt.plot(X_train, model.predict(X_train), color='red', label='Regression Line')
# Scatter plot of the test data
```

```
plt.scatter(X_test, y_test, color='green', label='Test Data')
```

```
plt.title("Simple Linear Regression")
```

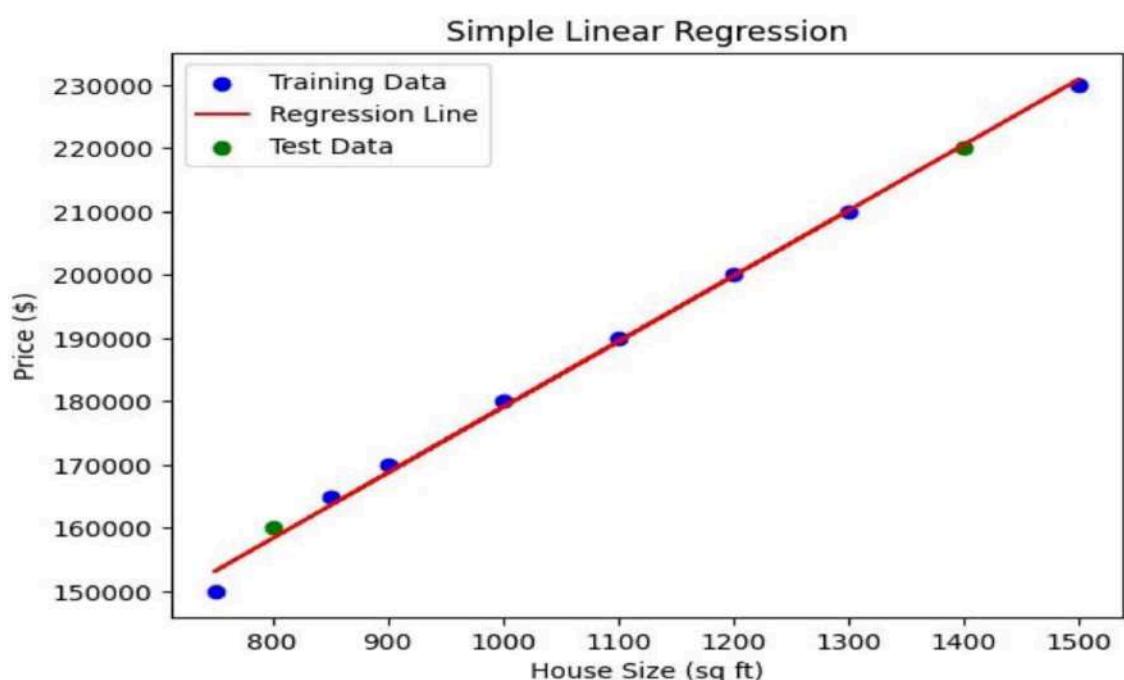
```
plt.xlabel("House Size (sq ft)")
```

```
plt.ylabel("Price ($)")
```

```
plt.legend()
```

```
plt.show()
```

Output :



3b. Multiple Linear Regression :

Extend linear regression to multiple feature. Handle feature selection and potential multicollinearity

Code :

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error, r2_score
from statsmodels.stats.outliers_influence import variance_inflation_factor
from sklearn.preprocessing import LabelEncoder

# Import LabelEncoder
from sklearn.impute import SimpleImputer

# Load dataset
from google.colab import files
uploaded = files.upload() # Upload your CSV file

# Read the CSV file
data = pd.read_csv(list(uploaded.keys())[0])

# Display the first few rows
print(data.head())

# Check for null values and basic statistics
print(data.info())
print(data.describe())

# Define a function to calculate VIF
def calculate_vif(df):

# Select only numeric features for VIF calculation
numeric_df = df.select_dtypes(include=np.number)

# Drop rows with infinite or missing values
```

```

numeric_df = numeric_df.replace([np.inf, -np.inf], np.nan).dropna()
vif_data = pd.DataFrame()
vif_data["feature"] = numeric_df.columns
vif_data["VIF"] = [variance_inflation_factor(numeric_df.values, i)
for i in range(numeric_df.shape[1])]

# Selecting features and target variable
X = data.drop("Survived", axis=1)
# Changed 'y' to 'Survived' y = data["Survived"]

# Handle categorical features (e.g., using Label Encoding)
for col in X.select_dtypes(include=['object']).columns:
    le = LabelEncoder()
    X[col] = le.fit_transform(X[col])

# Impute missing values using the mean (you can choose other strategies)
imputer = SimpleImputer(strategy='mean')
# Create an imputer instance
X = pd.DataFrame(imputer.fit_transform(X), columns=X.columns)
# Impute and update X

# Calculate VIF for initial features
print("VIF before handling multicollinearity:")
print(calculate_vif(X)) # Call the modified function

# Drop features based on VIF analysis (example: drop 'X1' if VIF is high)
# Check if the column exists before dropping
if 'X1' in X.columns:
    X = X.drop("X1", axis=1) # Replace 'X1' with the actual high VIF feature name
else:
    print("Column 'X1' not found in the DataFrame.")

# Recalculate VIF
print("VIF after handling multicollinearity:")
print(calculate_vif(X))
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Initialize and fit the model
model = LinearRegression()
model.fit(X_train, y_train)

# Get coefficients and intercept
print("Coefficients:", model.coef_)
print("Intercept:", model.intercept_)

# Predictions
y_pred = model.predict(X_test)

# Evaluation metrics

```

```

rmse = np.sqrt(mean_squared_error(y_test, y_pred)) r2 = r2_score(y_test, y_pred)
print(f"RMSE: {rmse}")
print(f"R^2: {r2}")
from sklearn.feature_selection import RFE

# Recursive Feature Elimination
rfe = RFE(estimator=LinearRegression(), n_features_to_select=5)

# Adjust features
rfe.fit(X_train, y_train)

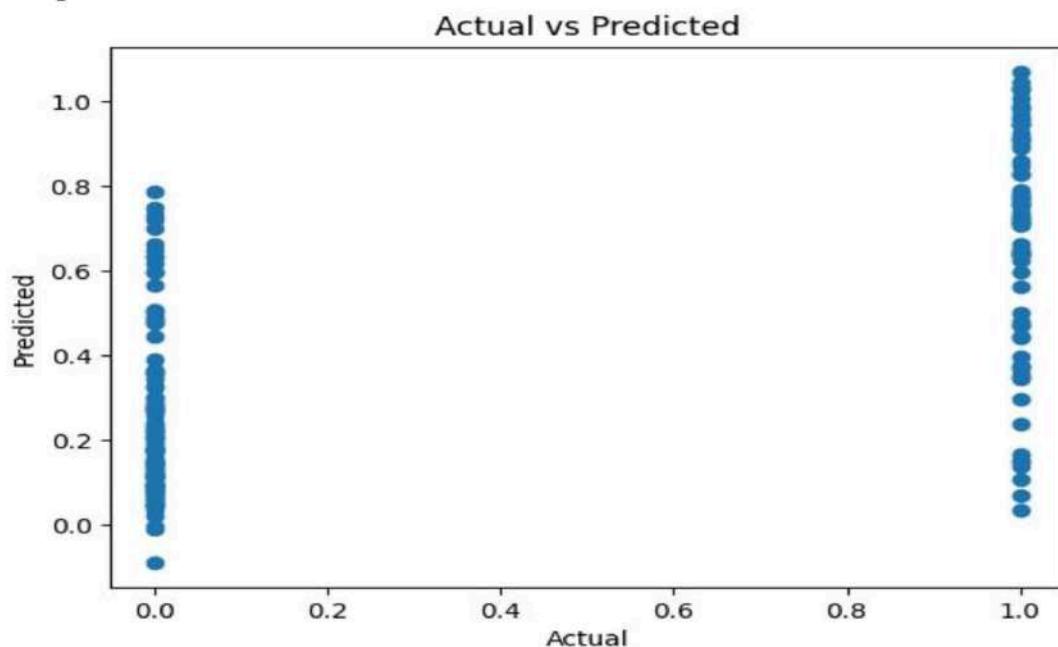
# Selected features
print("Selected Features:", X.columns[rfe.support_])

# Scatter plot of actual vs predicted values
plt.scatter(y_test, y_pred) plt.xlabel("Actual")
plt.ylabel("Predicted")
plt.title("Actual vs Predicted")
plt.show()

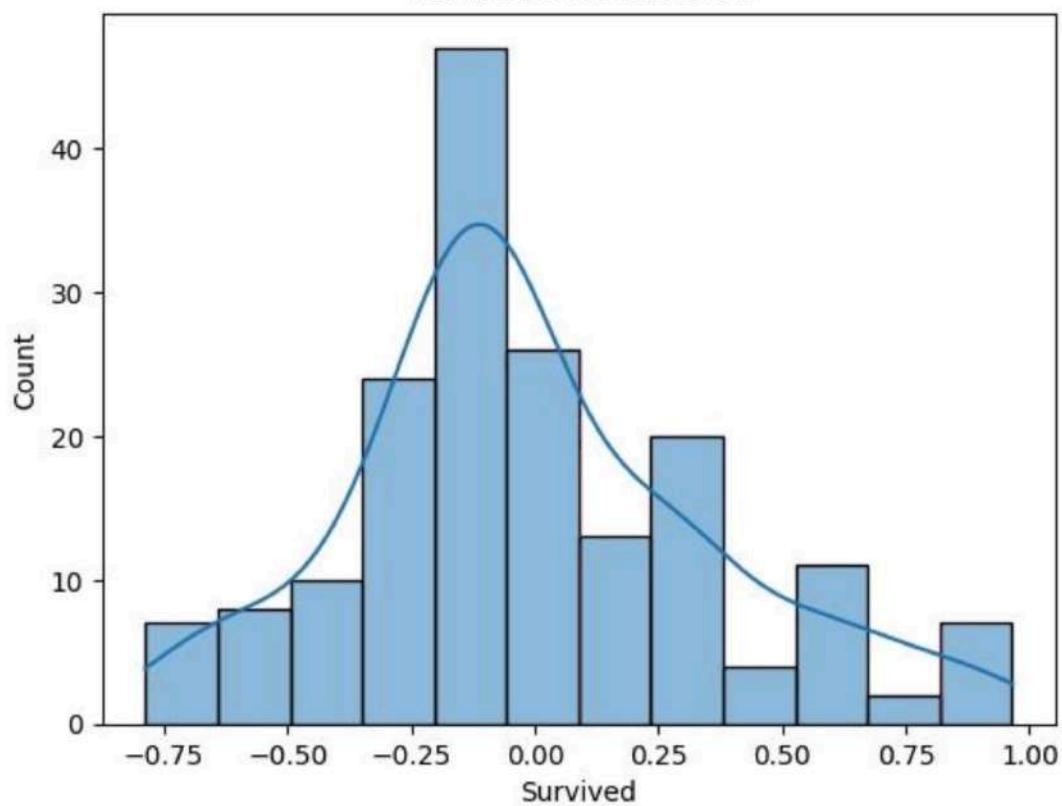
# Residuals
residuals = y_test - y_pred sns.histplot(residuals, kde=True)
plt.title("Residuals Distribution")
plt.show()

```

Output :



Residuals Distribution



3c. Regularized Linear Models :

Implement Regression variants like LASSO and Ridge on any generated dataset

Code :

1. Set Up the Environment

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.linear_model import Ridge, Lasso, ElasticNet
from sklearn.metrics import mean_squared_error, r2_score
from sklearn.datasets import make_regression
# Set random seed for reproducibility
np.random.seed(42)
```

2. Generate a Synthetic Dataset

Generate synthetic data

```
X, y = make_regression(n_samples=1000,
```

Number of samples

```
n_features=10,
```

Number of features

```
noise=15,
```

Add some noise

```
random_state=42
```

```
)
```

Convert to DataFrame for exploration

```
data = pd.DataFrame(X, columns=[f"X{i}"
```

```
for i in range(1, 11)]) data["y"] = y
```

Display the first few rows

```
print(data.head())
```

3. Split the Dataset

```
# Split data into training and testing sets
```

```
X_train, X_test, y_train, y_test = train_test_split(data.drop("y", axis=1),
```

```
# Features
```

```
data["y"],
```

```
# Target variable
```

```
test_size=0.2,
```

```
# 20% for testing
```

```
random_state=42
```

```
)
```

4. Train and Evaluate Ridge Regression

```
# Initialize Ridge Regression with a regularization parameter (alpha)
```

```
ridge = Ridge(alpha=1.0)
```

```
# Train the model
```

```
ridge.fit(X_train, y_train)
```

```
# Predictions
```

```
ridge_pred = ridge.predict(X_test)
```

```
# Evaluate Ridge Regression
```

```
ridge_rmse = np.sqrt(mean_squared_error(y_test, ridge_pred))
```

```
ridge_r2 = r2_score(y_test, ridge_pred)
```

```
print(f'Ridge RMSE: {ridge_rmse}')
```

```
print(f'Ridge R^2: {ridge_r2}')
```

5. Train and Evaluate Lasso Regression

```
# Initialize Lasso Regression
```

```
lasso = Lasso(alpha=0.1)
```

```
# Train the model
```

```
lasso.fit(X_train, y_train)
```

```
# Predictions  
lasso_pred = lasso.predict(X_test)
```

```
# Evaluate Lasso Regression  
lasso_rmse = np.sqrt(mean_squared_error(y_test, lasso_pred))  
lasso_r2 = r2_score(y_test, lasso_pred)  
print(f'Lasso RMSE: {lasso_rmse}')  
print(f'Lasso R^2: {lasso_r2}')  
# Features shrunk to  
zero print("Lasso Coefficients:", lasso.coef_)
```

6. Train and Evaluate ElasticNet Regression

```
# Initialize ElasticNet  
elastic_net = ElasticNet(alpha=0.1, l1_ratio=0.5) # l1_ratio balances L1 and L2 penalties
```

```
# Train the model  
elastic_net.fit(X_train, y_train)
```

```
# Predictions  
elastic_net_pred = elastic_net.predict(X_test)  
# Evaluate  
ElasticNet Regression elastic_net_rmse = np.sqrt(mean_squared_error(y_test,  
elastic_net_pred)) elastic_net_r2 = r2_score(y_test, elastic_net_pred)  
print(f'ElasticNet RMSE: {elastic_net_rmse}')  
print(f'ElasticNet R^2: {elastic_net_r2}')
```

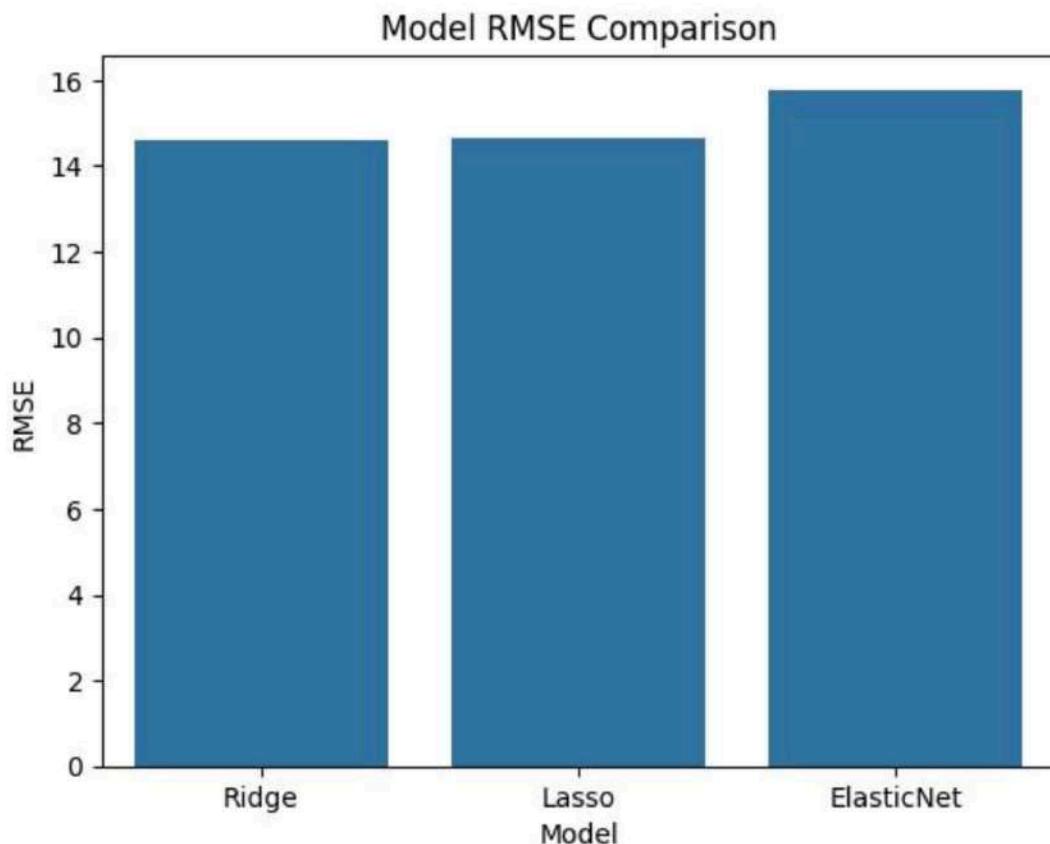
7. Compare Results

```
# Collect metrics  
metrics = pd.DataFrame({  
    "Model": ["Ridge", "Lasso", "ElasticNet"],  
    "RMSE": [ridge_rmse, lasso_rmse, elastic_net_rmse],  
    "R^2": [ridge_r2, lasso_r2, elastic_net_r2]})
```

```
})
print(metrics)

# Plot RMSE comparison
sns.barplot(data=metrics, x="Model", y="RMSE")
plt.title("Model RMSE Comparison")
plt.show()
```

Output :



Practical 4 : Discriminative Models

4a. Logistic Regression :

Perform binary classification using logistic regression. Calculate accuracy, precision, recall, and understand the ROC curve."

Code :

Step 1: Import Required Libraries

Import necessary libraries

```
import numpy as np
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, precision_score, recall_score, roc_curve, auc
import matplotlib.pyplot as plt
```

Step 2: Prepare the Dataset

```
from sklearn.datasets import make_classification
```

Create a synthetic dataset

```
X, y = make_classification(n_samples=1000, n_features=10, n_classes=2, random_state=42)
```

Split data into training and testing sets

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
```

Step 3: Train the Logistic Regression Model

Initialize the logistic regression model

```
logreg = LogisticRegression()
```

Train the model on the training data

```
logreg.fit(X_train, y_train)
```

Step 4: Make Predictions

```
# Predict labels for the test set
```

```
y_pred = logreg.predict(X_test)
```

```
# Predict probabilities for the ROC curve
```

```
y_prob = logreg.predict_proba(X_test)[:, 1]
```

Step 5: Evaluate the Model

```
# Calculate metrics
```

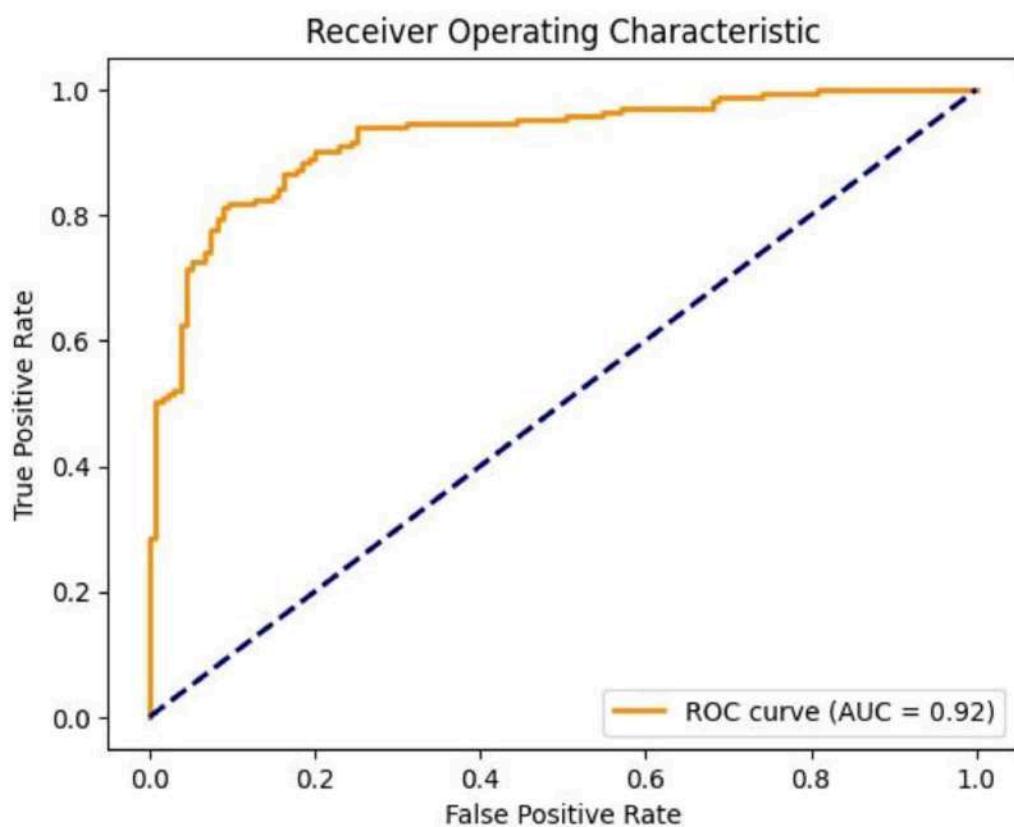
```
accuracy = accuracy_score(y_test, y_pred)
```

```
precision = precision_score(y_test, y_pred)
```

```
recall = recall_score(y_test, y_pred)
```

```
print(f'Accuracy: {accuracy:.2f}')
```

Output :



4b .Implement and demonstrate k-nearest Neighbor algorithm. Read the training data from a .CSV file and build the model to classify a test sample. Print both correct and wrong predictions.

Code :

```
Step 1: Import Required Libraries # Import necessary libraries
import pandas as pd

import numpy as np

from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score

from google.colab import files
```

Step 2: Create or Upload the CSV File

```
# Check if the user wants to create a dataset or upload one
print("Do you have a CSV file to upload? (yes/no)")
response = input().lower()

if response == "yes":
    uploaded = files.upload()
    filename = list(uploaded.keys())[0]
else:

    # Create a synthetic dataset
    from sklearn.datasets import make_classification

    # Generate synthetic data
    X,y=make_classification(n_samples=200,n_features=5, n_classes=2, random_state=42)

    # Combine features and target into a single DataFrame
    data = pd.DataFrame(X, columns=[f"Feature_{i}" for i in range(X.shape[1])])
    data["Target"] = y

    # Save the dataset to a CSV file
    filename = "synthetic_data.csv"
    data.to_csv(filename, index=False)

    print(f"Synthetic dataset saved as {filename}.")
```

Step 3: Load the CSV File into a DataFrame

```
# Load the dataset into a DataFrame  
data = pd.read_csv(filename)  
# Display the first few rows of the dataset  
print("Loaded Dataset:")  
print(data.head())
```

Step 4: Preprocess the Data**# Separate features (X) and labels (y)**

```
X = data.iloc[:, :-1].values # All columns except the last one  
y = data.iloc[:, -1].values # Last column as the target
```

Split the dataset into training and testing sets (80% train, 20% test)

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

Step 5: Train the k-NN Model #**Initialize the k-NN model with k=3 knn**

```
= KNeighborsClassifier(n_neighbors=3) #
```

Train the model on the training data

```
knn.fit(X_train, y_train)
```

Step 6: Predict Test Samples #**Predict the labels for the test set**

```
y_pred = knn.predict(X_test)
```

Step 7: Evaluate and Print Predictions #**Calculate and display the accuracy**

```
accuracy = accuracy_score(y_test, y_pred)  
print(f"\nModel Accuracy:  
{accuracy:.2f}\n")
```

Display correct and incorrect predictions

```
print("Correct Predictions:")  
for i in range(len(y_test)):  
    if y_pred[i] == y_test[i]:  
        print(f"Sample {i}: Predicted={y_pred[i]}, Actual={y_test[i]}")  
print("\nIncorrect Predictions:")  
  
for i in range(len(y_test)):
```

```
if y_pred[i] != y_test[i]:  
    print(f"Sample {i}: Predicted={y_pred[i]}, Actual={y_test[i]}")
```

Output :

```
[ ] Model Accuracy: 0.88  
→ Correct Predictions:  
Sample 0: Predicted=0, Actual=0  
Sample 1: Predicted=1, Actual=1  
Sample 2: Predicted=1, Actual=1  
Sample 3: Predicted=0, Actual=0  
Sample 4: Predicted=1, Actual=1  
Sample 5: Predicted=1, Actual=1  
Sample 6: Predicted=0, Actual=0  
Sample 7: Predicted=0, Actual=0  
Sample 9: Predicted=1, Actual=1  
Sample 10: Predicted=1, Actual=1  
Sample 11: Predicted=1, Actual=1  
Sample 12: Predicted=0, Actual=0  
Sample 13: Predicted=0, Actual=0  
Sample 14: Predicted=0, Actual=0  
Sample 15: Predicted=0, Actual=0  
Sample 16: Predicted=0, Actual=0  
Sample 17: Predicted=1, Actual=1  
Sample 18: Predicted=1, Actual=1  
Sample 19: Predicted=0, Actual=0  
Sample 20: Predicted=0, Actual=0  
Sample 22: Predicted=1, Actual=1  
Sample 23: Predicted=1, Actual=1  
Sample 24: Predicted=1, Actual=1  
Sample 25: Predicted=1, Actual=1  
Sample 26: Predicted=1, Actual=1  
Sample 27: Predicted=0, Actual=0  
Sample 28: Predicted=0, Actual=0  
Sample 30: Predicted=1, Actual=1  
Sample 31: Predicted=1, Actual=1  
Sample 32: Predicted=1, Actual=1  
Sample 34: Predicted=0, Actual=0  
Sample 35: Predicted=1, Actual=1  
Sample 36: Predicted=1, Actual=1  
Sample 38: Predicted=1, Actual=1  
Sample 39: Predicted=1, Actual=1
```

```
Incorrect Predictions:  
Sample 8: Predicted=1, Actual=0  
Sample 21: Predicted=1, Actual=0  
Sample 29: Predicted=0, Actual=1  
Sample 33: Predicted=1, Actual=0  
Sample 37: Predicted=1, Actual=0
```

4c. Build a decision tree classifier or regressor. Control hyperparameters like tree depth to avoid overfitting. Visualize the tree.

Code :

Step 1: Import Required Libraries

Import necessary libraries

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier, DecisionTreeRegressor,
plot_tree
from sklearn.metrics import accuracy_score, mean_squared_error
import matplotlib.pyplot as plt
from google.colab import files
```

Step 2: Create or Upload the CSV File

Check if the user wants to upload a file or generate one

```
print("Do you have a CSV file to upload? (yes/no)")
```

```
response = input().lower()
```

```
if response == "yes":
```

Upload the CSV file

```
uploaded = files.upload()
```

```
filename = list(uploaded.keys())[0]
```

```
else:
```

Generate synthetic data (classification or regression)

```
from sklearn.datasets import make_classification, make_regression
print("Choose a task: (1) Classification (2) Regression")
```

```
task = int(input())
```

```
if task == 1:
```

Generate synthetic classification data

```
X, y = make_classification(n_samples=200, n_features=5, random_state=42)
```

```
task_type = "classification"
```

```

else:
    # Generate synthetic regression data
    X, y = make_regression(n_samples=200, n_features=5, random_state=42)
    task_type = "regression"

    # Combine features and target into a single DataFrame
    data = pd.DataFrame(X, columns=[f"Feature_{i}" for i in range(X.shape[1])])
    data['Target'] = y

    # Save the dataset to a CSV file
    filename = "synthetic_data.csv"
    data.to_csv(filename, index=False)
    print(f"Synthetic {task_type} dataset saved as {filename}.")

```

Step 3: Load the Dataset

```

# Load the dataset
data = pd.read_csv(filename)

# Display the first few rows of the dataset
print("Dataset Preview:")
print(data.head())

```

Step 4: Preprocess the Data

```

# Separate features and target
X = data.iloc[:, :-1].values # All columns except the last one
y = data.iloc[:, -1].values # Last column as the target

# Split data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

```

Step 5: Build the Decision Tree

```

# Define the tree depth to avoid overfitting
max_depth = 3

```

```

# Initialize the model
if task_type == "classification":
    model = DecisionTreeClassifier(max_depth=max_depth, random_state=42)
else:
    model = DecisionTreeRegressor(max_depth=max_depth, random_state=42)

# Train the model
model.fit(X_train, y_train)

```

Step 6: Make Predictions

```

# Predict on the test set
y_pred = model.predict(X_test)
# Evaluate the model
if task_type == "classification":
    accuracy = accuracy_score(y_test, y_pred)
    print(f"Accuracy: {accuracy:.2f}")
else:
    mse = mean_squared_error(y_test, y_pred)
    print(f"Mean Squared Error: {mse:.2f}")

```

Step 7: Visualize the Tree

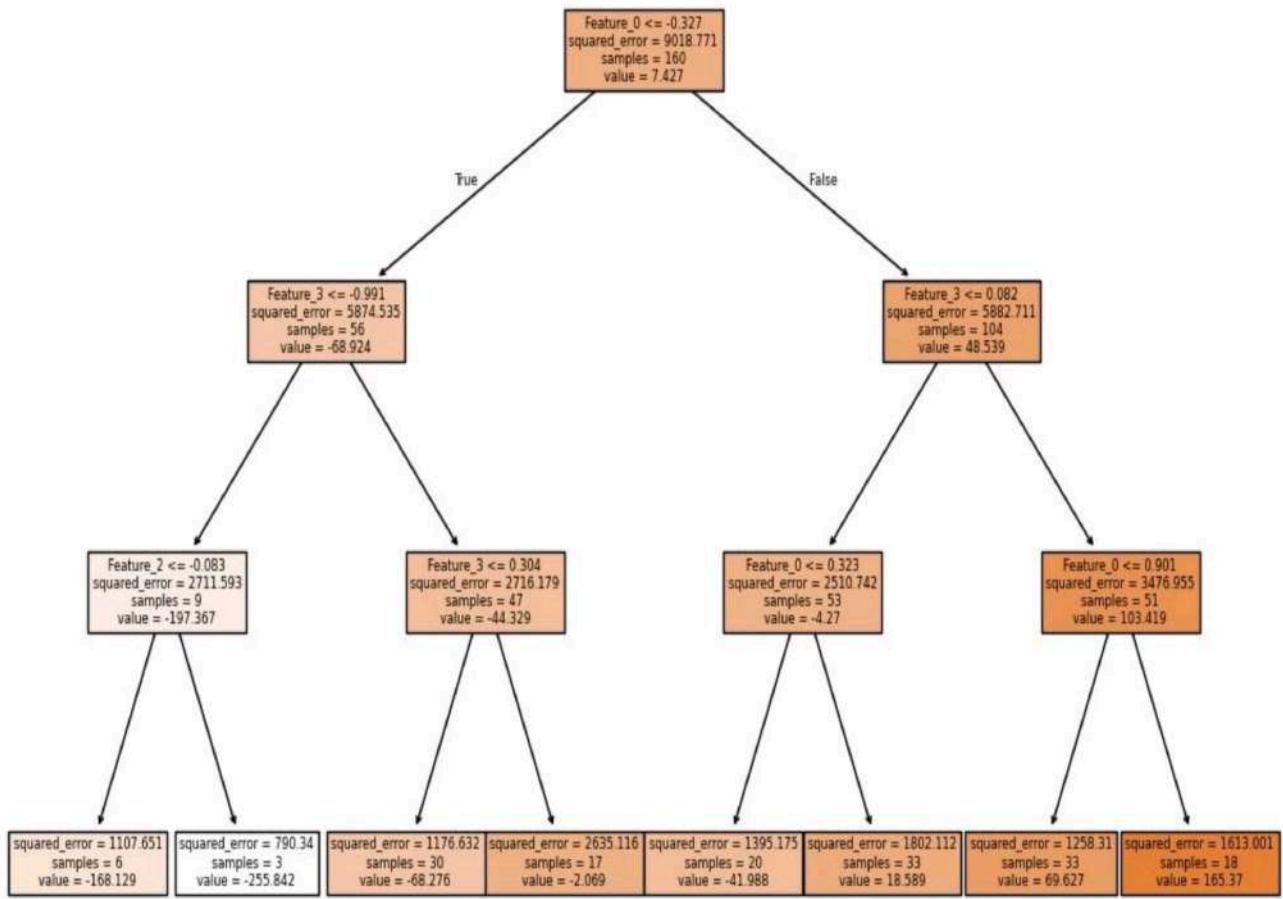
```

# Visualize the decision tree
plt.figure(figsize=(12, 8))
plot_tree(model, feature_names=data.columns[:-1], class_names=str(np.unique(y)))
if task_type == "classification" else None, filled=True)
plt.title("Decision Tree Visualization")
plt.show()

```

Output :

Decision Tree Visualization



4d. Implement a Support Vector Machine for any relevant dataset.

Code:

Step 1: Import Required Libraries

```
# Import necessary libraries
```

```
import pandas as pd
```

```
import numpy as np
```

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.svm import SVC
```

```
from sklearn.metrics import accuracy_score, classification_report
```

```
from google.colab import files
```

Step 2: Create or Upload a Dataset

Check if the user wants to upload a file or generate one

```
print("Do you have a CSV file to upload? (yes/no)")
```

```
response = input().lower()
```

```
if response == "yes":
```

Upload the CSV file

```
uploaded = files.upload()
```

```
filename = list(uploaded.keys())[0]
```

```
else:
```

Generate synthetic classification data

```
from sklearn.datasets import make_classification
```

```
X, y = make_classification(n_samples=200, n_features=5, n_classes=2, random_state=42)
```

Combine features and target into a DataFrame

```
data = pd.DataFrame(X, columns=[f"Feature_{i}"
```

```
for i in range(X.shape[1])])
```

```
data['Target'] = y
```

#Save the synthetic dataset to a CSV file

```
filename="synthetic_data.csv"
data.to_csv(filename,index=False)
print(f"Synthetic dataset saved as {filename}.")
```

Step 3: Load the Dataset

```
# Load the dataset into a DataFrame
```

```
data = pd.read_csv(filename)
```

```
# Display the first few rows of the dataset
```

```
print("Dataset Preview:")
```

```
print(data.head())
```

Step 4: Preprocess the Data

```
# Separate features (X) and target (y)
```

```
X = data.iloc[:, :-1].values # All columns except the last one
```

```
y = data.iloc[:, -1].values # Last column as the target
```

```
# Split the dataset into training (80%) and testing (20%) sets
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

Step 5: Train the Support Vector Machine

```
# Initialize the SVM model (use RBF kernel as default)
```

```
svm_model = SVC(kernel='rbf', C=1.0, gamma='scale', random_state=42)
```

```
# Train the SVM model on the training data
```

```
svm_model.fit(X_train, y_train)
```

Step 6: Make Predictions

```
# Predict the labels for the test set
```

```
y_pred = svm_model.predict(X_test)
```

Step 7: Evaluate the Model

```
# Calculate and print the accuracy
```

```
accuracy = accuracy_score(y_test, y_pred)
```

```
print(f"Model Accuracy: {accuracy:.2f}")
```

```
# Print a detailed classification report
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
```

Step 8: Visualize the Decision Boundary (Optional for 2D Data)

```
import matplotlib.pyplot as plt

# Generate 2D synthetic data
from sklearn.datasets import make_blobs
X, y = make_blobs(n_samples=100, centers=2, random_state=42, cluster_std=1.5)

# Fit the SVM on this data
svm_model.fit(X, y)

# Plot the decision boundary
plt.figure(figsize=(8, 6))

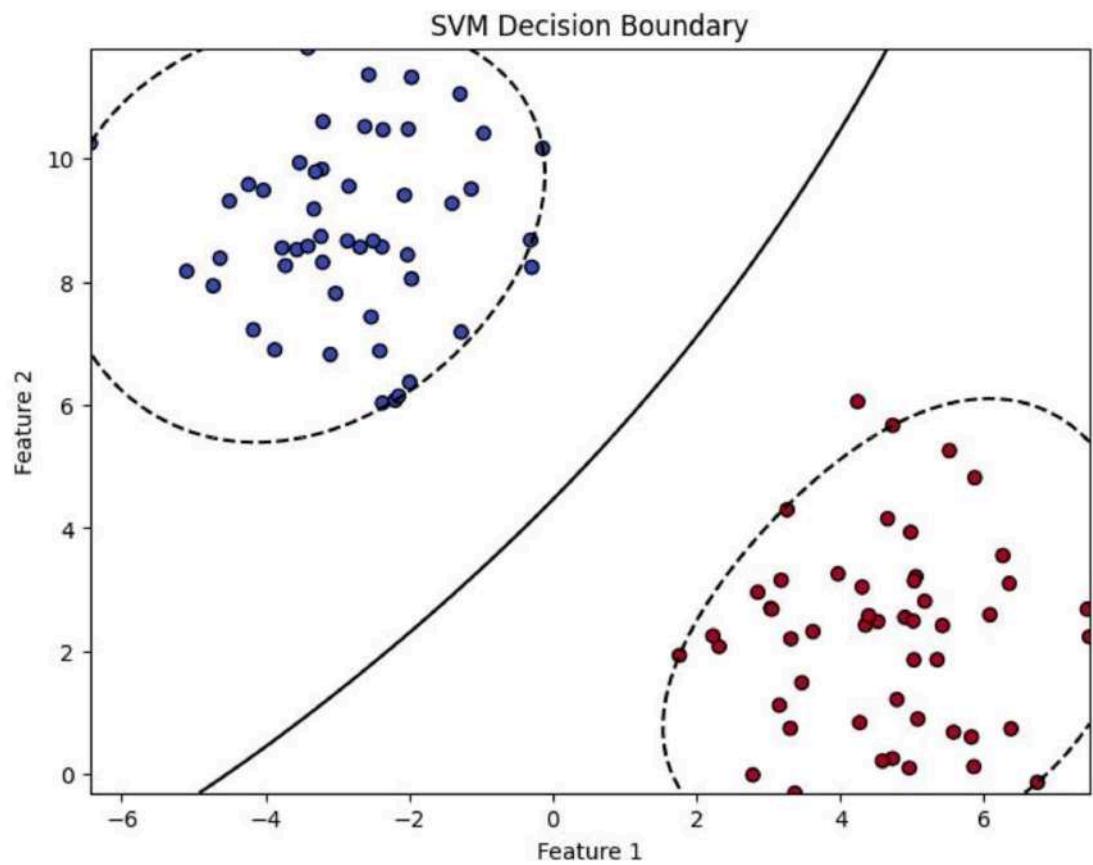
plt.scatter(X[:, 0], X[:, 1], c=y, cmap='coolwarm', edgecolor='k')
# Create a grid to evaluate the model

xx, yy = np.meshgrid(np.linspace(X[:, 0].min(), X[:, 0].max(), 100), np.linspace(X[:, 1].min(), X[:, 1].max(), 100))
Z = svm_model.decision_function(np.c_[xx.ravel(), yy.ravel()])

Z = Z.reshape(xx.shape)

# Plot the decision boundary and margins
plt.contour(xx, yy, Z, levels=[-1, 0, 1], linestyles=['--', ':', '-'], colors='k')
plt.title("SVM Decision Boundary")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.show()
```

Output :



4e. Train a random forest ensemble. Experiment with the number of trees and feature sampling. Compare performance to a single decision tree.

Code :

Step 1: Import Required Libraries # Import necessary libraries

```
import pandas as pd

import numpy as np

from sklearn.model_selection import train_test_split

from sklearn.tree import DecisionTreeClassifier

from sklearn.ensemble import RandomForestClassifier

from sklearn.metrics import accuracy_score, classification_report

from google.colab import files
```

Step 2: Create or Upload a Dataset

```
# Check if the user wants to upload a file or generate one
print("Do you have a CSV file to upload? (yes/no)")
response = input().lower()
if response == "yes":
# Upload the CSV file
uploaded = files.upload()
filename = list(uploaded.keys())[0]
else:
# Generate synthetic classification data
from sklearn.datasets import make_classification

X, y = make_classification(n_samples=300, n_features=10, n_classes=2, random_state=42)

# Combine features and target into a DataFrame
data = pd.DataFrame(X, columns=[f'Feature_{i}' for i in range(X.shape[1])])
data['Target'] = y

# Save the synthetic dataset to a CSV file
filename = "synthetic_data.csv"
data.to_csv(filename, index=False)
print(f'Synthetic dataset saved as {filename}.')
```

Step 3: Load the Dataset

```
# Load the dataset
```

```
data = pd.read_csv(filename)

# Display the first few rows of the dataset
print("Dataset Preview:")
print(data.head())
```

Step 4: Preprocess the Data

```
# Separate features (X) and target (y)
```

```
X = data.iloc[:, :-1].values # All columns except the last one
y = data.iloc[:, -1].values # Last column as the target
```

```
# Split the dataset into training (80%) and testing (20%) sets
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

Step 5: Train a Single Decision Tree Classifier

```
# Initialize and train the Decision Tree model
```

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

```
# Predict and evaluate
```

```
y_pred_tree = decision_tree.predict(X_test)
accuracy_tree = accuracy_score(y_test, y_pred_tree)
print(f"Decision Tree Accuracy: {accuracy_tree:.2f}")
```

Step 6: Train a Random Forest Classifier

```
# Initialize the Random Forest model with hyperparameter tuning
```

```
random_forest = RandomForestClassifier(n_estimators=100, max_features='sqrt',
random_state=42)
```

```
# Train the model
```

```
random_forest.fit(X_train, y_train)
```

```
# Predict and evaluate
```

```
y_pred_rf = random_forest.predict(X_test)
accuracy_rf = accuracy_score(y_test, y_pred_rf)
print(f"Random Forest Accuracy (100 trees, sqrt features): {accuracy_rf:.2f}")
```

Step 7: Experiment with Random Forest Hyperparameters

```
# Experiment with fewer trees and different feature sampling
```

```
rf_experiment = RandomForestClassifier(n_estimators=50, max_features=3,
random_state=42)
```

```
rf_experiment.fit(X_train, y_train)
```

Predict and evaluate

```
y_pred_rf_exp = rf_experiment.predict(X_test)  
accuracy_rf_exp = accuracy_score(y_test, y_pred_rf_exp)  
print(f"Random Forest Accuracy (50 trees, max_features=3): {accuracy_rf_exp:.2f}")
```

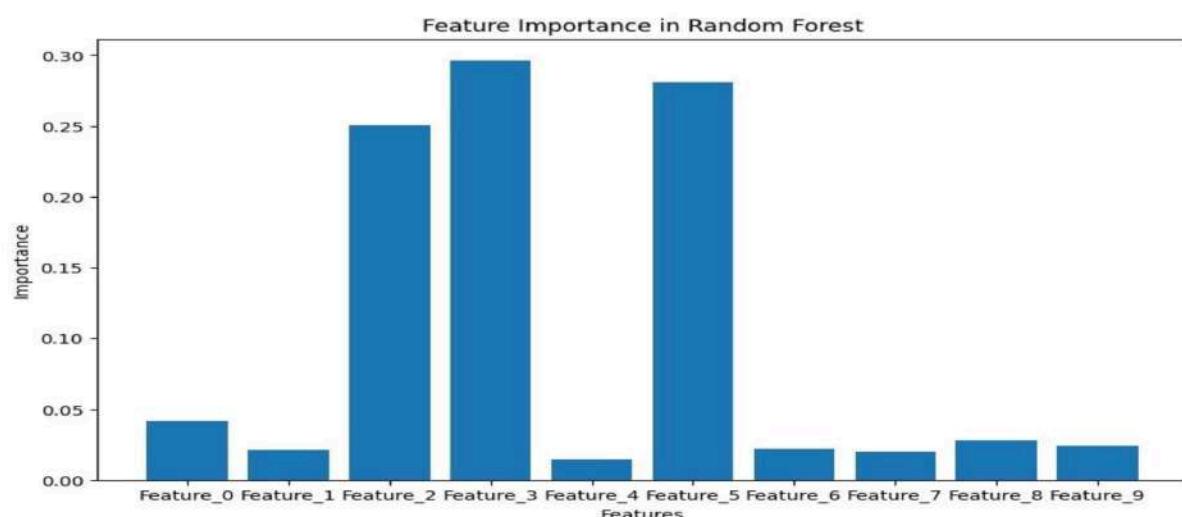
Step 8: Compare the Models

```
print("\nModel Comparison:")  
print(f"Decision Tree Accuracy: {accuracy_tree:.2f}")  
print(f"Random Forest Accuracy (100 trees): {accuracy_rf:.2f}")  
print(f"Random Forest Accuracy (50 trees, max_features=3): {accuracy_rf_exp:.2f}")
```

Step 9: Visualize Feature Importance (Optional)

```
import matplotlib.pyplot as plt  
  
# Extract feature importance from the Random Forest model  
feature_importances = random_forest.feature_importances_  
  
# Plot the feature importance  
plt.figure(figsize=(10, 6))  
plt.bar(range(len(feature_importances)), feature_importances, tick_label=data.columns[:-1])  
plt.title("Feature Importance in Random Forest")  
plt.xlabel("Features")  
plt.ylabel("Importance")  
plt.show()
```

Output :



4f. Implement a gradient boosting machine (e.g., XGBoost). Tune hyperparameters and explore feature importance.

Code :

Step 1: Import Required Libraries

Import necessary libraries

```
import pandas as pd
```

```
import numpy as np
```

```
from sklearn.model_selection import train_test_split, GridSearchCV  
from sklearn.metrics import accuracy_score, classification_report  
from xgboost import XGBClassifier, plot_importance
```

```
import matplotlib.pyplot as plt
```

```
from google.colab import files
```

Step 2: Create or Upload a Dataset

Check if the user wants to upload a file or generate

```
one print("Do you have a CSV file to upload? (yes/no)")
```

```
response = input().lower()
```

```
if response == "yes":
```

```
# Upload the CSV file
```

```
uploaded=files.upload()
```

```
filename = list(uploaded.keys())[0]
```

```
else:
```

Generate synthetic classification data

```
from sklearn.datasets import make_classification
```

```
X, y = make_classification(n_samples=300, n_features=10, n_classes=2, random_state=42)
```

Combine features and target into a DataFrame

```
data = pd.DataFrame(X, columns=[f"Feature_{i}"
```

```
for i in range(X.shape[1])])data['Target'] = y
```

Save the synthetic dataset to a CSV file

```
filename="synthetic_data.csv"
data.to_csv(filename,index=False)
print(f"Synthetic dataset saved as {filename}.")
```

Step 3: Load the Dataset

Load the dataset

```
data = pd.read_csv(filename)
# Display the first few rows of the
dataset print("Dataset Preview:")
print(data.head())
```

Step 4: Preprocess the Data

Separate features (X) and target (y)

```
X = data.iloc[:, :-1].values # All columns except the last one
y = data.iloc[:, -1].values # Last column as the target
```

Split the dataset into training (80%) and testing (20%) sets

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

Step 5: Train a Basic XGBoost Model

Initialize and train the XGBoost model with default parameters

```
xgb = XGBClassifier(random_state=42)
xgb.fit(X_train, y_train)
```

Predict and evaluate the model

```
y_pred = xgb.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)

print(f"XGBoost Accuracy (Default Parameters): {accuracy:.2f}")
```

Step 6: Tune Hyperparameters with GridSearchCV

Define a grid of hyperparameters

```
param_grid = { 'n_estimators': [50, 100, 150], 'learning_rate': [0.01, 0.1, 0.2], 'max_depth': [3, 5, 7] }
```

Initialize GridSearchCV

```
grid_search =
GridSearchCV(estimator=XGBClassifier(random_state=42), param_grid=param_grid,
scoring='accuracy', cv=3, verbose=1)
# Fit the model with grid search
grid_search.fit(X_train, y_train)
```

```

# Best parameters from GridSearch
print(f'Best Parameters: {grid_search.best_params_}')
# Train the final model with the best parameters
best_xgb = grid_search.best_estimator_
# Predict and evaluate

y_pred_best = best_xgb.predict(X_test)

accuracy_best = accuracy_score(y_test, y_pred_best)
print(f'XGBoost Accuracy (Tuned Parameters): {accuracy_best:.2f}')

```

Step 7: Explore Feature Importance

Plot feature importance for the tuned model

```

plt.figure(figsize=(10, 6))

plot_importance(best_xgb, importance_type='weight', xlabel="Importance",
                ylabel="Features")

plt.title("XGBoost Feature Importance")
plt.show()

```

Step 8: Evaluate the Model

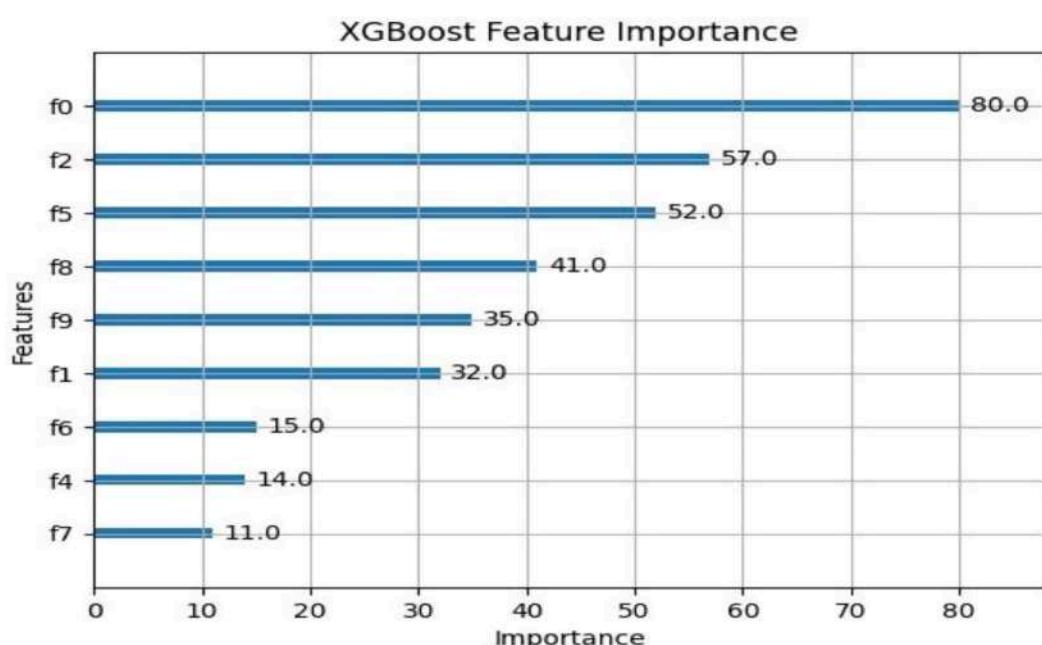
Print a detailed classification report

```

print("Classification Report:")
print(classification_report(y_test, y_pred_best))

```

Output :



Practical 5

5a. Implement and demonstrate the working of a Naive Bayesian classifier using a sample data set. Build the model to classify a test sample.

Step 1: Import Required Libraries

Import necessary libraries

```
import pandas as pd  
import numpy as np  
from sklearn.model_selection import train_test_split  
from sklearn.metrics import accuracy_score,classification_report  
from sklearn.naive_bayes import GaussianNB  
from google.colab import files
```

Step 2: Create or Upload a Dataset

Ask if the user wants to upload a file or generate one

```
print("Do you have a CSV file to upload? (yes/no)")
```

```
response = input().lower()
```

```
if response == "yes":
```

Upload the CSV file

```
uploaded = files.upload()
```

```
filename = list(uploaded.keys())[0]
```

```
else:
```

Generate synthetic classification data

```
from sklearn.datasets import make_classification
```

```
X, y = make_classification(n_samples=300, n_features=8, n_classes=2, random_state=42)
```

Combine features and target into a DataFrame

```
data = pd.DataFrame(X, columns=[f'Feature_{i}' for i in range(X.shape[1])])  
data['Target'] = y
```

Save the synthetic dataset to a CSV file

```
filename = "synthetic_naive_bayes_data.csv"
```

```
data.to_csv(filename, index=False)
```

```
print(f'Synthetic dataset saved as {filename}.')
```

Step 3: Load the Dataset

```
# Load the dataset  
data = pd.read_csv(filename)  
  
# Display the first few rows of the dataset  
print("Dataset Preview:")  
print(data.head())
```

Step 4: Preprocess the Data

```
# Separate features (X) and target (y)  
X = data.iloc[:, :-1].values # All columns except the last one  
y = data.iloc[:, -1].values # Last column as the target  
  
# Split the dataset into training (80%) and testing (20%) sets  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

Step 5: Train a Naive Bayes Classifier

```
# Initialize the Gaussian Naive Bayes classifier  
naive_bayes = GaussianNB()  
  
# Train the model  
naive_bayes.fit(X_train, y_train)
```

Step 6: Make Predictions and Evaluate

```
# Predict on the test set  
y_pred = naive_bayes.predict(X_test)  
  
# Evaluate the model  
accuracy = accuracy_score(y_test, y_pred)  
print(f'Naive Bayes Accuracy:  
{accuracy:.2f}')  
  
# Detailed classification report  
print("Classification Report:")  
print(classification_report(y_test, y_pred))
```

Step 7: Test the Model with a Custom Sample

```

# Define a sample test input (replace with meaningful values based on your dataset)
test_sample = [X_test[0]]

# Taking the first test sample for demonstration

# Predict the class for the test sample

predicted_class = naive_bayes.predict(test_sample)

print(f"Test Sample: {test_sample}")

print(f"Predicted Class: {predicted_class[0]}")

```

Output :

Dataset Preview:						
	Feature_0	Feature_1	Feature_2	Feature_3	Feature_4	Feature_5
0	-1.274158	1.317988	-2.423879	0.906946	-1.583903	-0.331811
1	1.607963	-1.649959	0.299293	-0.891720	1.301741	1.508502
2	-0.154167	0.161033	2.210523	0.139400	-0.557492	0.087713
3	-0.920991	0.949136	-1.613561	0.588410	1.471170	-0.529287
4	1.013304	-1.038578	-0.305225	-0.539334	-0.609512	1.048078
	Feature_6	Feature_7	Target			
0	-0.452306	0.760415	1			
1	0.742095	1.561511	0			
2	0.963879	-1.369803	0			
3	-1.371901	-0.209324	0			
4	-1.065114	-0.186971	0			

```

→ Test Sample: [array([-0.90320608,  0.9220511 , -1.32308979,  0.41081065,  1.64201516,
   -1.23559176, -0.63896175,  1.00981709])]

Predicted Class: 1

```

5b. Implement Hidden Markov Models using hmmlearn

Code :

Step 1: Install Required Libraries

```
# Install hmmlearn
```

```
!pip install hmmlearn
```

Step 2: Import Required Libraries

```
# Import necessary libraries
```

```
import numpy as np
```

```
import pandas as pd
```

```
from hmmlearn import hmm
```

```
import matplotlib.pyplot as plt
```

Step 3: Create or Load a Dataset

```
# Generate synthetic observable data
```

```
np.random.seed(42)
```

```
# Create a sequence of observations and hidden states
```

```
observations = np.random.choice(['A', 'B', 'C'], size=100, p=[0.5, 0.3, 0.2])
```

```
hidden_states = np.random.choice(['X', 'Y'], size=100, p=[0.6, 0.4])
```

```
# Save the data in a DataFrame for analysis
```

```
data = pd.DataFrame({'Observations': observations, 'Hidden States': hidden_states})
```

```
print("Generated Data:")
```

```
print(data.head())
```

Step 4: Encode Observations

```
# Encode the observations into integers
```

```
observation_mapping = {obs: idx for idx, obs in enumerate(np.unique(observations))}
```

```
encoded_observations = np.array([observation_mapping[obs] for obs in observations])
```

```
# Print the mapping
```

```
print("Observation Encoding:")
```

```
print(observation_mapping)
```

Step 5: Initialize and Configure the HMM

Initialize the HMM model

```
n_states = 2 # Number of hidden states
```

```
n_observations = len(observation_mapping)
```

Number of unique observations

```
model = hmm.MultinomialHMM(n_components=n_states, random_state=42, n_iter=100, tol=0.01)
```

Define start probabilities (initial distribution of states)

```
start_probs = np.array([0.6, 0.4]) # Assumed probabilities
```

```
model.startprob_ = start_probs
```

Define transition probabilities between states

```
trans_probs = np.array([
```

```
    [0.7, 0.3], # From state X
```

```
    [0.4, 0.6], # From state Y])
```

```
model.transmat_ = trans_probs
```

Define emission probabilities (probability of observations given states)

```
emission_probs = np.array([
```

```
    [0.5, 0.4, 0.1], # State X emits A, B, C
```

```
    [0.2, 0.3, 0.5], # State Y emits A, B, C
```

```
])
```

```
model.emissionprob_ = emission_probs
```

Print the configured model parameters

```
print("Start Probabilities:", model.startprob_)
```

```
print("Transition Matrix:", model.transmat_)
```

```
print("Emission Probabilities:",
```

```
model.emissionprob_)
```

Step 6: Train the Model

Reshape the data for HMM (requires 2D array)

```
encoded_observations = encoded_observations.reshape(-1, 1)
```

Fit the model

```
model.fit(encoded_observations)
```

Predict hidden states for the observations

```
predicted_states = model.predict(encoded_observations)
```

```
# Print the predicted states
print("Predicted States:")
print(predicted_states)
```

Step 7: Visualize the Results

```
# Map predicted states back to their original labels
```

```
state_mapping = {0: 'X', 1: 'Y'}
```

```
predicted_state_labels = [state_mapping[state] for state in predicted_states]
```

```
# Add predicted states to the DataFrame
```

```
data['Predicted States'] = predicted_state_labels
```

```
# Display the first few rows with predicted states
```

```
print("Data with Predicted States:")
```

```
print(data.head())
```

```
# Plot the observations and predicted states
```

```
plt.figure(figsize=(12, 6))
```

```
plt.plot(data['Observations'], label='Observations', marker='o', linestyle='-', alpha=0.7)
```

```
plt.plot(data['Predicted States'], label='Predicted States', marker='x', linestyle='--', alpha=0.7)
```

```
plt.legend()
```

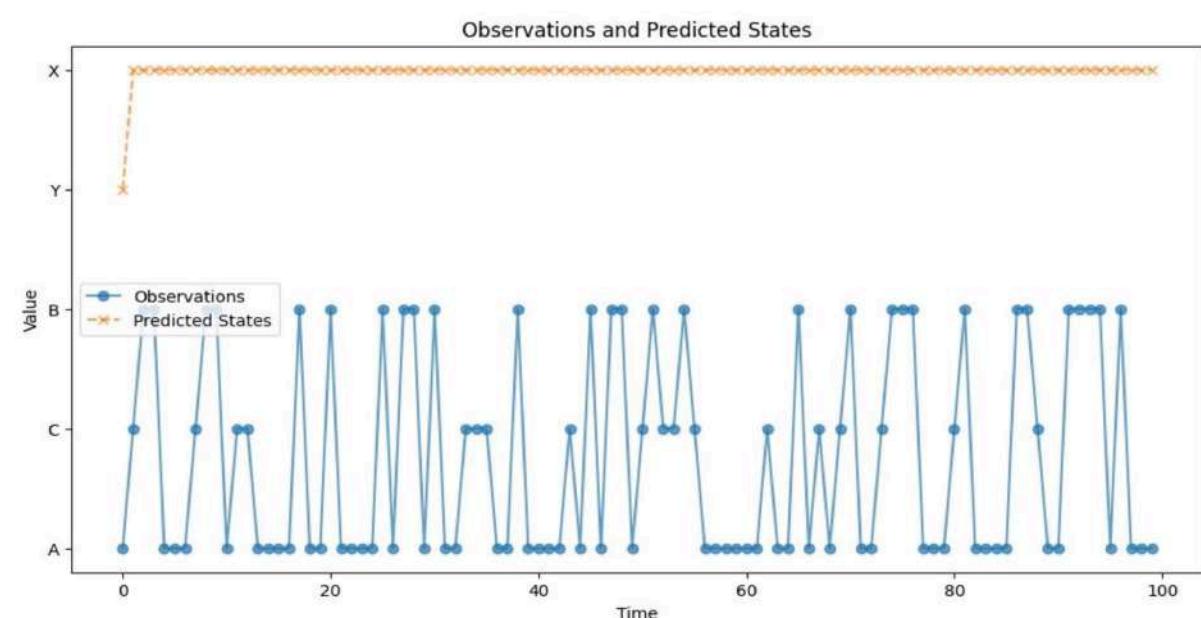
```
plt.title("Observations and Predicted States")
```

```
plt.xlabel("Time")
```

```
plt.ylabel("Value")
```

```
plt.show()
```

Output :



Practical 6 : Probabilistic Model

6a. Implement Bayesian Linear Regression to explore prior and posterior distribution.

Bayesian Linear Regression is a probabilistic approach to linear regression that incorporates uncertainty in the model parameters. Instead of estimating point values for parameters (as in traditional linear regression), we estimate distributions over the parameters.

Code :

Step 1: Install Required Libraries

Install necessary libraries

```
!pip install matplotlib seaborn scikit-learn
```

Step 2: Import Required Libraries

Import necessary libraries

```
import numpy as np  
import pandas as pd  
import matplotlib.pyplot as plt  
import seaborn as sns  
from sklearn.linear_model import BayesianRidge  
from sklearn.model_selection import train_test_split  
from sklearn.metrics import mean_squared_error  
from google.colab import files
```

Step 3: Create or Upload a Dataset

Upload a CSV file if you have one

```
print("Do you have a CSV file to upload? (yes/no)")  
response = input().lower()  
  
if response == "yes":  
  
    # Upload the CSV file  
  
    uploaded = files.upload()
```

```

filename = list(uploaded.keys())[0]
else:

# Generate synthetic data for demonstration
np.random.seed(42)

X = np.random.rand(100, 1) * 10

# Random data between 0 and 10

y = 2 * X + 1 + np.random.randn(100, 1) * 2

# y = 2x + 1 with some noise

# Convert to a DataFrame

data = pd.DataFrame(np.hstack((X, y)), columns=["X", "y"])

# Save to CSV for convenience

filename="synthetic_data.csv"

data.to_csv(filename,index=False)

print(f"Synthetic dataset saved as {filename}.")

```

Step 4: Load and Explore the Data

```

# Load the dataset (for CSV file)
data = pd.read_csv(filename)

# Display first few rows
print("Dataset Preview:")
print(data.head())

```

Step 5: Preprocess the Data

```

# Separate features (X) and target (y)

X = data["X"].values.reshape(-1, 1) # Feature matrix
y = data["y"].values # Target vector

# Split the dataset into training (80%) and testing (20%) sets

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

```

Step 6: Implement Bayesian Linear Regression Model

```

# Initialize the BayesianRidge model (which implements Bayesian Linear Regression)

bayesian_regressor = BayesianRidge()

# Fit the model on the training data

bayesian_regressor.fit(X_train, y_train)

```

```
# Predict on the test data  
y_pred = bayesian_regressor.predict(X_test)
```

Step 7: Visualize the Prior and Posterior Distributions

Plot the prior and posterior distributions of the parameters

```
fig, ax = plt.subplots(1, 2, figsize=(12, 6))  
  
# Plot prior distribution (assuming the model starts with a standard prior)  
ax[0].set_title("Prior Distribution (Assumed)") ax[0].hist(np.random.normal(0, 1, 1000),  
bins=50, alpha=0.7, color='blue', label="Prior") ax[0].legend()  
  
# Plot posterior distribution (after model fitting)  
  
ax[1].set_title("Posterior Distribution (After Fitting)")  
ax[1].hist(bayesian_regressor.coef_, bins=50, alpha=0.7, color='green',  
label="Posterior") ax[1].legend()  
  
plt.show()
```

Step 8: Evaluate the Model Performance

```
Calculate the Mean Squared Error (MSE)  
mse = mean_squared_error(y_test, y_pred)  
print(f'Mean Squared Error (MSE):  
{mse:.2f}')
```

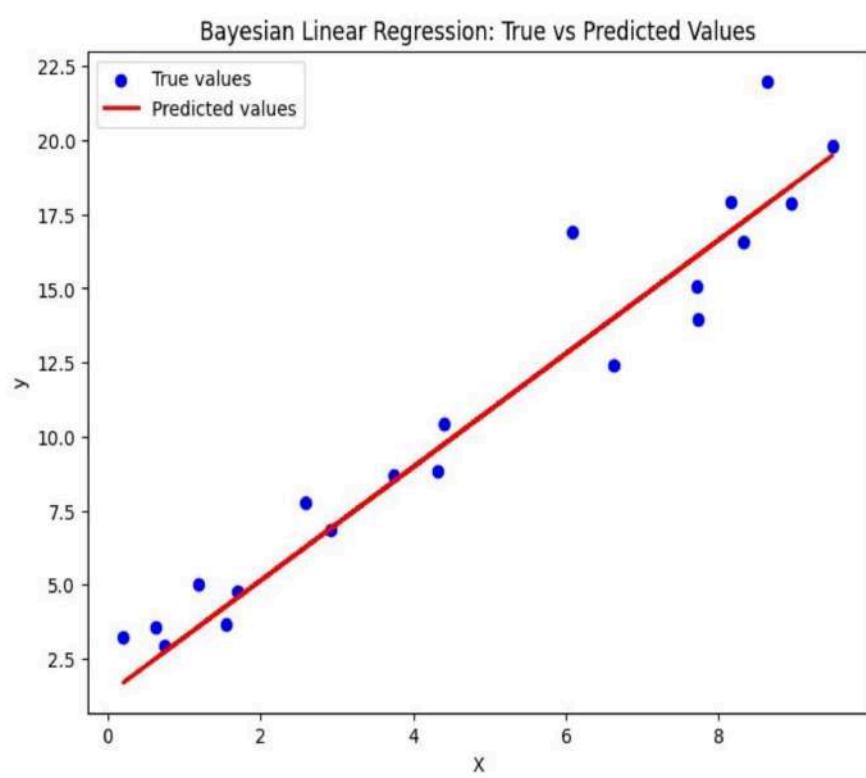
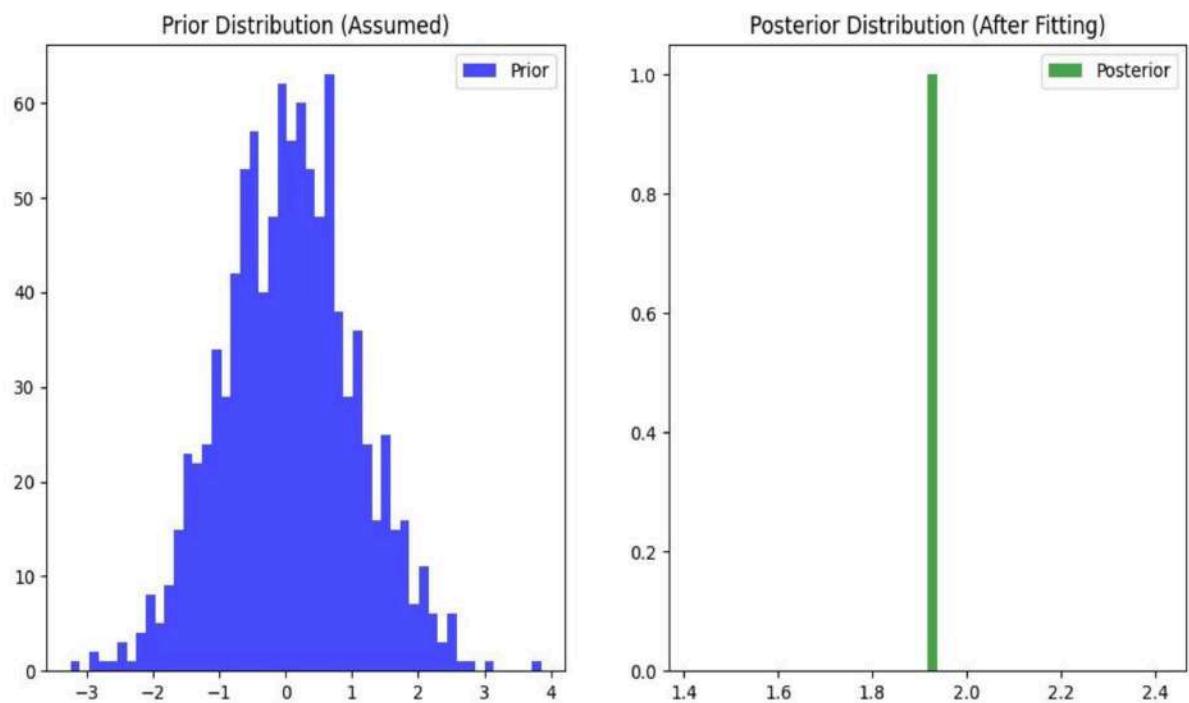
Step 9: Visualize the Fit of the Model

Plot the true values and the predicted values

```
plt.figure(figsize=(8, 6))  
  
plt.scatter(X_test, y_test, color="blue", label="True values")  
plt.plot(X_test, y_pred, color="red", label="Predicted values",  
linewidth=2)  
  
plt.title("Bayesian Linear Regression: True vs Predicted Values")  
plt.xlabel("X")  
plt.ylabel("y")  
plt.legend()  
plt.show()
```

Mean Squared Error (MSE): 3.9

Output :



6b. Implement Gaussian Mixture Models for density estimation and unsupervised clustering.

Code :

Step 1: Install Required Libraries

Install required libraries

```
!pip install matplotlib seaborn scikit-learn
```

Step 2: Import Required Libraries

Import necessary libraries

```
import numpy as np
```

```
import pandas as pd
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
from sklearn.mixture import GaussianMixture
```

```
from sklearn.model_selection import train_test_split
```

```
from google.colab import files
```

Step 3: Create or Upload a Dataset

#Ask if the user has a CSV file to upload

```
print("Do you have a CSV file to upload? (yes/no)")
```

```
response = input().lower()
```

```
if response == "yes":
```

Upload the CSV file

```
uploaded = files.upload()
```

```
filename = list(uploaded.keys())[0]
```

```
else:
```

Generate synthetic 2D data with two clusters for demonstration

```
np.random.seed(42)
```

Generate data for two Gaussian distributions

```
X1 = np.random.normal(loc=0, scale=1, size=(300, 2)) # Cluster 1: mean = 0, std = 1
```

```
X2 = np.random.normal(loc=5, scale=1, size=(300, 2)) # Cluster 2: mean = 5, std = 1 #
```

Stack the data to create a dataset

```
X = np.vstack([X1, X2])
```

```
# Create DataFrame to simulate the CSV file for consistency
data = pd.DataFrame(X, columns=["Feature_1", "Feature_2"])
filename = "synthetic_data.csv"
data.to_csv(filename, index=False)
print(f"Synthetic dataset saved as {filename}.")
```

Step 4: Load and Explore the Dataset

```
# Load the dataset (if CSV file is uploaded)
data = pd.read_csv(filename)
# Display the first few rows
print("Dataset Preview:")
print(data.head())
# Plot the data to visualize its structure
sns.scatterplot(data=data, x="Feature_1", y="Feature_2")
plt.title("Synthetic Data")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.show()
```

Step 5: Fit a Gaussian Mixture Model (GMM)

```
# Define the GMM model
n_components = 2 # Number of Gaussian distributions (clusters)
gmm = GaussianMixture(n_components=n_components, covariance_type='full',
random_state=42)

# Fit the GMM model to the data
gmm.fit(data)

# Predict the cluster labels for each data point
labels = gmm.predict(data)

# Add the cluster labels to the dataset for visualization
data['Cluster'] = labels

# Plot the clustered data
sns.scatterplot(data=data, x="Feature_1", y="Feature_2", hue="Cluster", palette="viridis",
marker="o")

plt.title("Gaussian Mixture Model Clustering")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
```

```
plt.legend()
```

```
plt.show()
```

Step 6: Visualize the Gaussian Mixture Model (GMM) Components

```
# Extract the means and covariances of the Gaussian components
```

```
means = gmm.means_
```

```
covariances = gmm.covariances_
```

```
# Plot the GMM components on top of the data
```

```
plt.figure(figsize=(8, 6))
```

```
# Plot data points
```

```
sns.scatterplot(data=data, x="Feature_1", y="Feature_2", hue="Cluster",  
palette="viridis", marker="o", s=60, alpha=0.7)
```

```
# Plot the GMM ellipses for mean, covar in zip(means, covariances):
```

```
# Plot the Gaussian components as ellipses
```

```
v, w = np.linalg.eigh(covar)
```

```
v = 2.0 * np.sqrt(2.0) * np.sqrt(v)
```

```
# Scaling factor for the ellipse
```

```
u = w[0] / np.linalg.norm(w[0])
```

```
# Normalize the eigenvector
```

```
angle = np.arctan(u[1] / u[0])
```

```
# Create the ellipse
```

```
angle = angle * 180.0 / np.pi # Convert to degrees
```

```
ellipse = plt.matplotlib.patches.Ellipse(means, v[0], v[1], angle=angle, color='red', alpha=0.3)  
plt.gca().add_patch(ellipse)
```

```
plt.title("GMM Clustering with Gaussian Components")
```

```
plt.xlabel("Feature 1")
```

```
plt.ylabel("Feature 2")
```

```
plt.legend()
```

```
plt.show()
```

Step 7: Model Evaluation (Optional)

```
# Compute the log-likelihood of the data under the fitted GMM model
```

```
log_likelihood = gmm.score(data)
```

```
print(f"Log-Likelihood of the data: {log_likelihood:.2f}")
```

Step 8: Predict New Data Points

```
# Example of predicting the cluster for new data points
new_data = np.array([[1.5, 2.5], [4.5, 5.5], [7.0, 8.0]])

new_labels = gmm.predict(new_data)

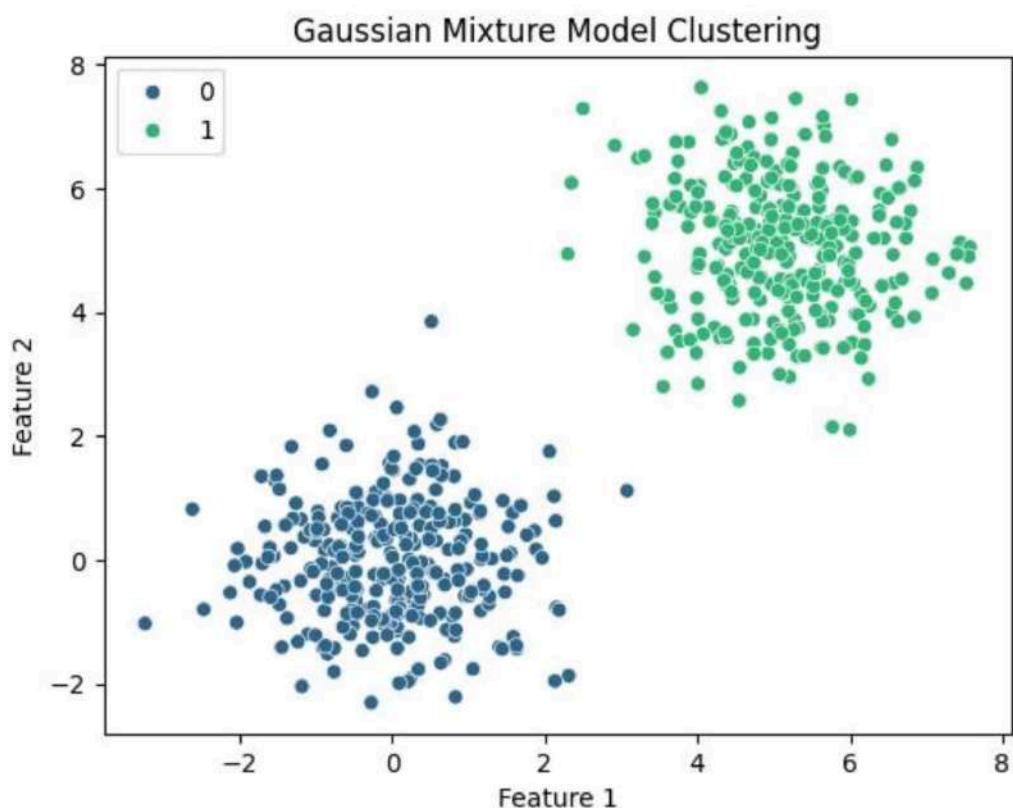
# Print the predicted clusters for the new data
# points

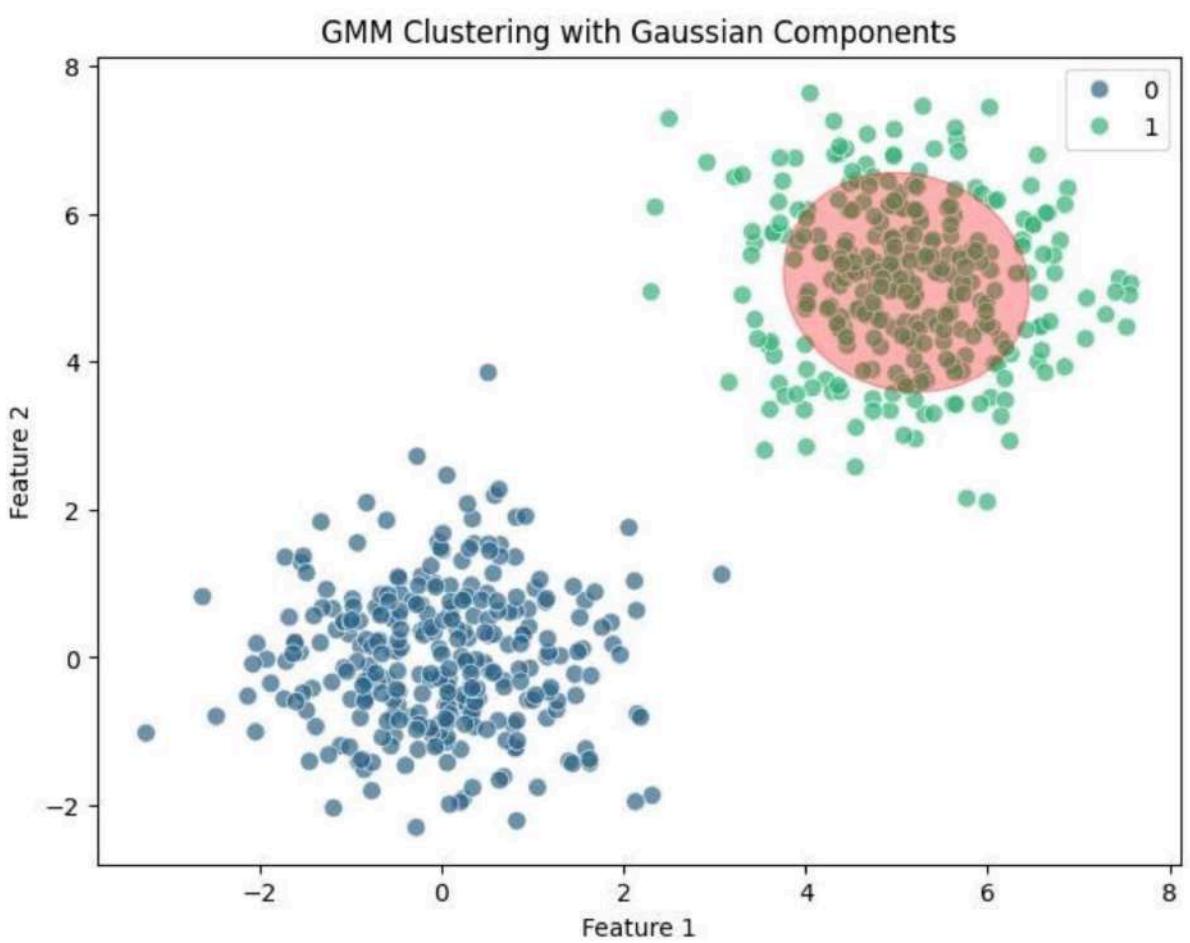
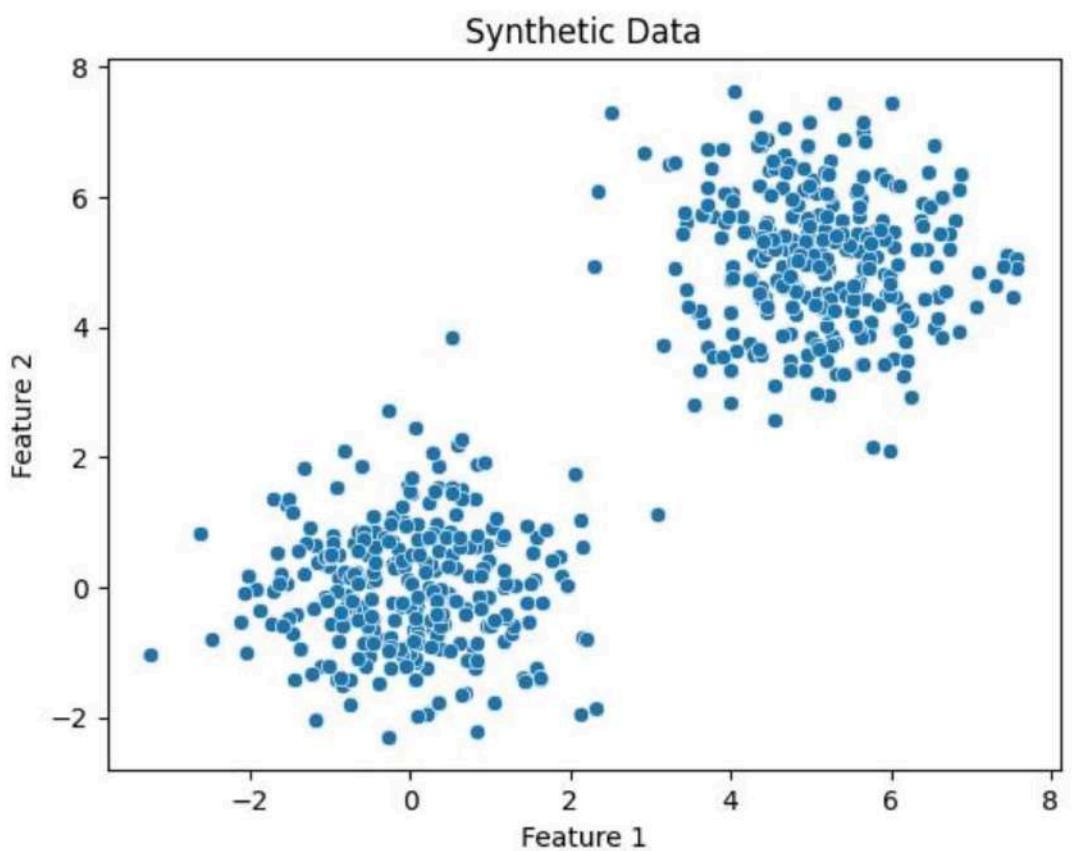
print("Predicted Clusters for New Data Points:")

for i, label in enumerate(new_labels):

    print(f'Data point {new_data[i]} is in Cluster {label}'")
```

Output :





Practical 7 : Model Evaluation and Hyperparameter Tuning

7a. Implement cross-validation techniques (k-fold, stratified, etc.) for robust model evaluation

Code :

1. Import Necessary Libraries

```
import numpy as np
import pandas as pd
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split, KFold, StratifiedKFold, GridSearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
import matplotlib.pyplot as plt
import seaborn as sns
```

2. Generate a Synthetic Dataset

Create a synthetic dataset with 2 classes

```
X, y = make_classification(
    n_samples=1000, n_features=10, n_informative=8, n_redundant=2,
    n_clusters_per_class=1, random_state=42
)
```

Convert to a DataFrame for visualization

```
df = pd.DataFrame(X, columns=[f'Feature_{i}' for i in range(1, 11)])
df['Target'] = y
```

Display the first few rows

```
print(df.head())
```

3. Split Data into Train and Test Sets

Split data into 80% training and 20% testing

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42, stratify=y)
```

4. Define k-Fold Cross-Validation

```
kf = KFold(n_splits=5, shuffle=True, random_state=42)
print("k-Fold Cross-Validation:")
for train_index, val_index in kf.split(X_train):
    print("TRAIN:", train_index, "VALIDATION:", val_index)
```

5. Define Stratified k-Fold Cross-Validation

```
skf = StratifiedKFold(n_splits=5, shuffle=True, random_state=42)
print("\nStratified k-Fold Cross-Validation:")
for train_index, val_index in skf.split(X_train, y_train):
    print("TRAIN:", train_index, "VALIDATION:", val_index)
```

6. Train and Evaluate Using k-Fold Cross-Validation

Initialize model

```
model = RandomForestClassifier(random_state=42)
```

Perform k-Fold Cross-Validation

```
accuracies = []
```

```
for train_index, val_index in kf.split(X_train):
```

```
    X_kf_train, X_kf_val = X_train[train_index], X_train[val_index]
```

```
    y_kf_train, y_kf_val = y_train[train_index], y_train[val_index]
```

Train model

```
    model.fit(X_kf_train, y_kf_train)
```

Validate model

```
    y_pred = model.predict(X_kf_val)
```

```
    accuracy = accuracy_score(y_kf_val, y_pred)
```

```
    accuracies.append(accuracy)
```

```
print(f"Average Accuracy from k-Fold: {np.mean(accuracies):.2f}")
```

7. Hyperparameter Tuning Using GridSearchCV

```
# Define parameter grid
param_grid = {
    'n_estimators': [50, 100, 200],
    'max_depth': [None, 10, 20, 30],
    'min_samples_split': [2, 5, 10],
}

# Perform GridSearchCV with Stratified k-Fold
grid_search = GridSearchCV(
    estimator=RandomForestClassifier(random_state=42),
    param_grid=param_grid,
    cv=StratifiedKFold(n_splits=5, shuffle=True, random_state=42),
    scoring='accuracy',
    n_jobs=-1,
    verbose=1
)

# Fit to training data
grid_search.fit(X_train, y_train)
print("Best Parameters:", grid_search.best_params_)
print("Best Cross-Validation Accuracy:", grid_search.best_score_)
```

8. Evaluate the Final Model

```
# Use the best model for evaluation
best_model = grid_search.best_estimator_
# Predict on test data
y_test_pred = best_model.predict(X_test)
# Evaluate performance
print("\nTest Accuracy:", accuracy_score(y_test, y_test_pred))
print("\nClassification Report:\n", classification_report(y_test, y_test_pred))
```

Confusion matrix

```
conf_matrix = confusion_matrix(y_test, y_test_pred)

sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=['Class 0', 'Class 1'], yticklabels=['Class 0', 'Class 1'])

plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.title('Confusion Matrix')
plt.show()
```

Output :

```
[[ 0 -3.358483  3.159918  0.827163  0.069658 -6.715639 -2.708559
  1  2.071819 -4.055419 -2.615948 -2.599432  3.053752  0.366795
  2 -0.633469  0.712482  2.024390 -0.432639 -1.307929  0.419320
  3 -0.464478  0.892442  2.521018  2.766580 -1.933734 -1.418018
  4  1.842426 -1.192605 -2.071386 -0.131231  0.545377  0.379060

   Feature_7  Feature_8  Feature_9  Feature_10 Target
0  0.183206  1.113502  1.730759  1.228394      1
1 -0.392171 -1.191720 -1.220516  1.899925      0
2 -1.469518 -0.719051  1.155005  2.018026      0
3  1.391760 -2.430279  1.308295 -0.270896      1
4 -0.062978 -1.325591  2.037936  0.115414      0

k-Fold Cross-Validation:
TRAIN: [ 0  1  3  4  5  6  8  9 11 12 13 14 15 16 17 18 19 20
21 22 24 25 26 27 28 32 34 35 36 37 38 40 41 42 43 44
45 46 47 48 50 51 52 53 55 56 57 58 59 60 61 62 64 68
69 70 71 73 74 75 79 80 82 83 85 87 88 89 90 91 92 93
94 95 98 99 100 102 103 104 105 106 107 108 111 112 113 114 115 116
117 119 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136
138 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 156 157
158 159 160 161 162 163 164 165 166 167 169 170 171 172 173 175 176 177
178 179 180 181 182 183 184 185 186 187 188 189 190 191 193 194 195 196
197 198 201 202 203 205 206 207 212 213 214 215 217 218 220 221 222 223
224 225 226 227 228 229 230 232 233 234 236 237 238 239 240 241 242 243
245 246 247 248 249 251 252 253 255 256 257 258 259 261 262 263 264 267
268 269 270 271 272 273 274 276 277 278 279 280 282 283 284 285 287 288
289 290 291 292 293 295 297 298 299 300 301 303 304 305 307 308 309 310
311 312 313 315 317 318 319 320 321 322 324 325 328 329 330 331 332 334

785 708 789 796 791 792 793 784 797 799] VALIDATION: [- 2  0  7  8 10  18 23 29 30 31 33 39 49 54 63 65 66 67 72 76 77
78 81 86 87 91 92 100 101 102 120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136
137 138 139 140 141 142 143 144 145 146 147 148 150 151 152 153 154 155 156 157 159 160 161
162 166 167 168 170 171 172 173 174 175 176 180 183 184 185 186 187 188
189 190 191 192 194 195 197 199 200 201 202 203 204 205 206 207 208
209 210 211 214 215 216 217 218 219 221 222 224 225 226 228 229 230 231
232 233 235 236 237 238 240 241 242 243 244 245 246 249 250 251 252 253
254 255 256 257 258 260 261 262 263 265 266 267 268 269 270 271 272 273
274 275 276 277 278 279 280 281 282 283 284 286 287 288 289 293 294 295
296 297 298 301 302 303 304 305 306 307 308 310 312 313 314 315 316 317
318 320 321 322 323 324 325 326 327 330 333 335 336 337 338 339 340 341

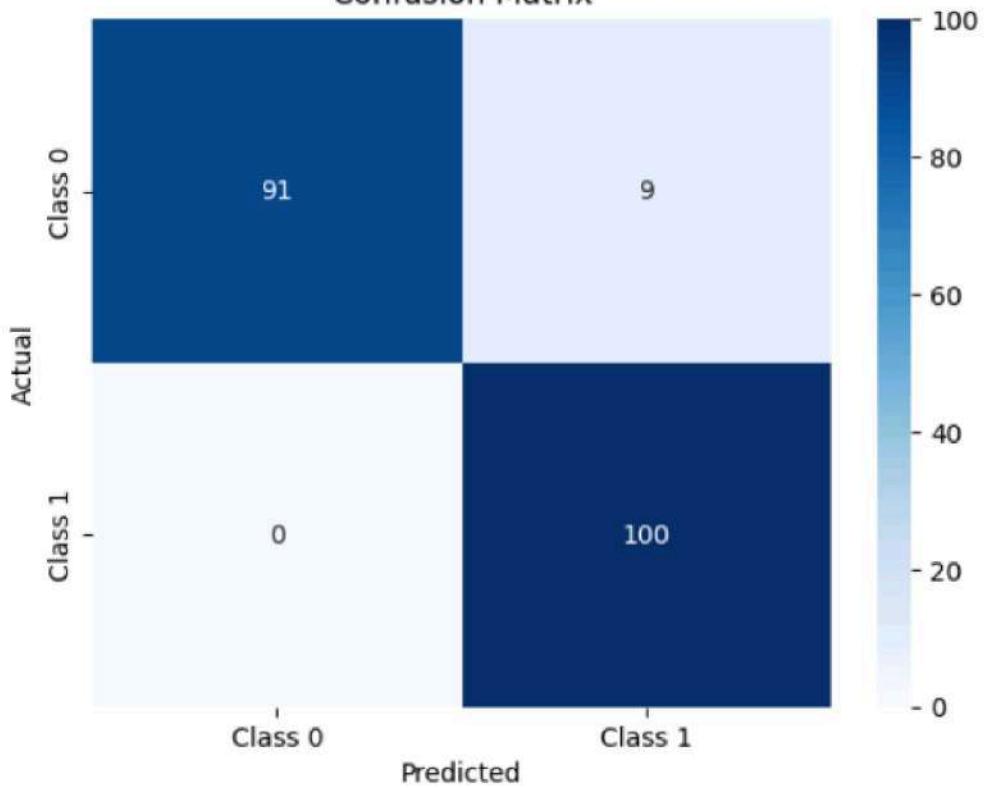
450 455 459 468 463 470 472 475 486 481 483 494 496 502 503 505 514 519
521 523 529 533 553 555 560 563 565 579 585 590 594 600 603 620 630 631
634 635 637 638 644 646 648 649 651 673 675 686 691 693 708 709 715 720
729 730 738 747 753 759 767 771 774 776 777 780 782 783 784 796
Average Accuracy from k-Fold: 0.95
Fitting 5 folds for each of 36 candidates, totalling 180 fits
Best Parameters: {'max_depth': None, 'min_samples_split': 5, 'n_estimators': 50}
Best Cross-Validation Accuracy: 0.96

Test Accuracy: 0.95

Classification Report:
precision    recall    f1-score   support
0          1.00     0.91     0.95     100
1          0.92     1.00     0.96     100

accuracy          0.95     200
macro avg       0.96     0.96     0.95     200
weighted avg    0.96     0.95     0.95     200
```

Confusion Matrix



7b. Systematically explore combinations of hyperparameters to optimize model performance.(use grid and randomized search)

Code :

1. Import Necessary Libraries

```
import numpy as np  
import pandas as pd  
from sklearn.datasets import make_classification  
from sklearn.model_selection import train_test_split, GridSearchCV, RandomizedSearchCV, StratifiedKFold  
from sklearn.ensemble import RandomForestClassifier  
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix  
import matplotlib.pyplot as plt  
import seaborn as sns
```

2. Generate a Synthetic Dataset

Generate a binary classification dataset

```
X, y = make_classification(  
    n_samples=1000, n_features=12, n_informative=8, n_redundant=2,  
    n_clusters_per_class=1, flip_y=0.03, random_state=42  
)
```

Convert to a DataFrame for visualization

```
df = pd.DataFrame(X, columns=[f'Feature_{i}' for i in range(1, 13)])  
df['Target'] = y
```

Display the first few rows

```
print(df.head())
```

3. Split Data into Train and Test Sets

Split data into training and testing sets

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42, stratify=y)
```

4. Define the Model

```
# Initialize a Random Forest classifier
```

```
model = RandomForestClassifier(random_state=42)
```

5. Hyperparameter Tuning Using Grid Search

```
# Define a parameter grid for Grid Search
```

```
param_grid = {
```

```
    'n_estimators': [50, 100, 200],
```

```
    'max_depth': [None, 10, 20],
```

```
    'min_samples_split': [2, 5, 10],
```

```
    'min_samples_leaf': [1, 2, 4]
```

```
}
```

```
# GridSearchCV with 5-fold cross-validation
```

```
grid_search = GridSearchCV(
```

```
    estimator=model,
```

```
    param_grid=param_grid,
```

```
    scoring='accuracy',
```

```
    cv=StratifiedKFold(n_splits=5, shuffle=True, random_state=42),
```

```
    verbose=1,
```

```
    n_jobs=-1
```

```
)
```

```
# Fit the model
```

```
grid_search.fit(X_train, y_train)
```

```
# Best parameters and score from Grid Search
```

```
print("Best Parameters from Grid Search:", grid_search.best_params_)
```

```
print("Best Cross-Validation Accuracy from Grid Search:", grid_search.best_score_)
```

6. Hyperparameter Tuning Using Randomized Search

```
from scipy.stats import randint
```

```

# Define a parameter distribution for Randomized Search
param_dist = {
    'n_estimators': randint(50, 300),
    'max_depth': [None, 10, 20, 30],
    'min_samples_split': randint(2, 15),
    'min_samples_leaf': randint(1, 10)
}

# RandomizedSearchCV with 5-fold cross-validation
random_search = RandomizedSearchCV(
    estimator=model,
    param_distributions=param_dist,
    n_iter=50, # Number of random combinations to try
    scoring='accuracy',
    cv=StratifiedKFold(n_splits=5, shuffle=True, random_state=42),
    verbose=1,
    n_jobs=-1,
    random_state=42
)

# Fit the model
random_search.fit(X_train, y_train)

# Best parameters and score from Randomized Search
print("Best Parameters from Randomized Search:", random_search.best_params_)
print("Best Cross-Validation Accuracy from Randomized Search:",
      random_search.best_score_)

```

7. Evaluate the Best Model

```

# Select the best model from Grid Search and Randomized Search
best_model = random_search.best_estimator_ # Or use grid_search.best_estimator_
# Predict on test data
y_test_pred = best_model.predict(X_test)
# Evaluate the performance

```

```

print("\nTest Accuracy:", accuracy_score(y_test, y_test_pred))
print("\nClassification Report:\n", classification_report(y_test, y_test_pred))

# Confusion Matrix

conf_matrix = confusion_matrix(y_test, y_test_pred)

sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=['Class 0', 'Class 1'], yticklabels=['Class 0', 'Class 1'])

plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.title('Confusion Matrix')
plt.show()

```

Output :

```

Feature_1  Feature_2  Feature_3  Feature_4  Feature_5  Feature_6  \
0  0.013650  0.607473 -2.096916  2.867232  2.504360  0.784101
1  0.107199  0.105735 -3.843343  1.524052 -1.619824  0.778334
2  -1.779086 -5.219831 -0.738488  2.108084 -0.803833 -3.431122
3  -4.310656 -2.268569  1.864943 -1.246116  1.268794 -2.007664
4  -3.195179 -0.671327  3.720485  0.356661  0.819486  2.670238

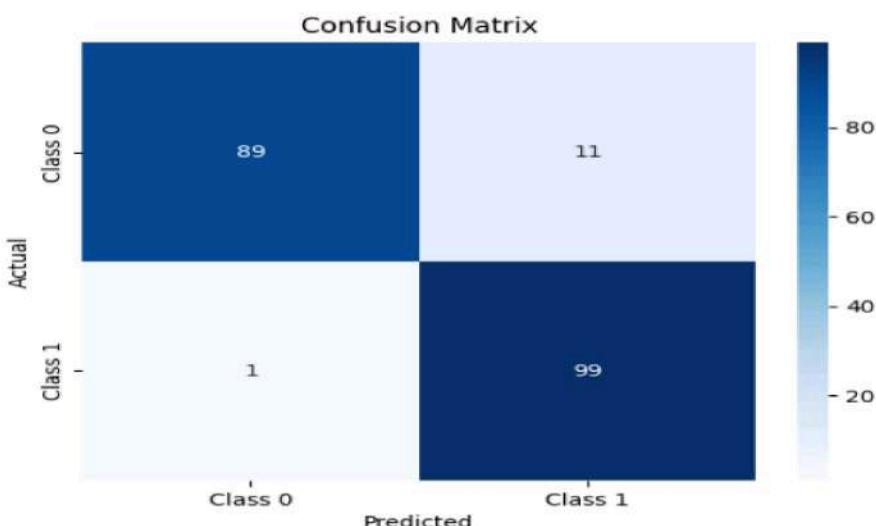
Feature_7  Feature_8  Feature_9  Feature_10  Feature_11  Feature_12  Target
0  -0.497744 -0.482072  1.112773  1.641637 -2.689832 -0.488311  1
1  0.551177 -1.843583 -0.110132 -0.494739 -0.985276 -0.978400  0
2  1.346120 -0.858351 -0.792415 -2.260815  0.238780  3.029952  0
3  -0.824133 -2.277449  0.936206  1.255903  1.386278 -0.321200  0
4  1.857477 -3.410944 -1.773719  0.656476  3.534189 -1.704889  0

Fitting 5 folds for each of 81 candidates, totalling 405 fits
Best Parameters from Grid Search: {'max_depth': None, 'min_samples_leaf': 1, 'min_samples_split': 2, 'n_estimators': 100}
Best Cross-Validation Accuracy from Grid Search: 0.9487499999999999
Fitting 5 folds for each of 50 candidates, totalling 250 fits
Best Parameters from Randomized Search: {'max_depth': None, 'min_samples_leaf': 2, 'min_samples_split': 9, 'n_estimators': 285}
Best Cross-Validation Accuracy from Randomized Search: 0.9487499999999999

Test Accuracy: 0.94

```

Classification Report:					
	precision	recall	f1-score	support	
0	0.99	0.89	0.94	100	
1	0.90	0.99	0.94	100	
accuracy			0.94	200	
macro avg	0.94	0.94	0.94	200	
weighted avg	0.94	0.94	0.94	200	



Practical 8 : Bayesian Learning

Implement Bayesian Learning using inferences

Code :

1. Import Necessary Libraries

```
import numpy as np
import pandas as pd
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt
```

2. Generate a Synthetic Dataset

We create a dataset suitable for classification problems.

Generate a dataset with 2 classes

```
X, y = make_classification(
    n_samples=1000, n_features=8, n_informative=6, n_redundant=2,
    n_classes=2, random_state=42)
```

Convert to DataFrame for visualization

```
df = pd.DataFrame(X, columns=[f'Feature_{i}' for i in range(1, 9)])
df['Target'] = y
```

Display the first few rows

```
print(df.head())
```

3. Split the Dataset

Divide the data into training and testing sets.

Split data into 80% training and 20% testing

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42, stratify=y)
```

4. Bayesian Learning with Naive Bayes

Here, we implement Bayesian Learning using the Gaussian Naive Bayes classifier.

Initialize the Gaussian Naive Bayes model

```
model = GaussianNB()
```

Fit the model to the training data

```
model.fit(X_train, y_train)
```

Predict on the test data

```
y_pred = model.predict(X_test)
```

5. Evaluate the Model

We evaluate the model's performance using accuracy, classification report, and confusion matrix.

Calculate accuracy

```
accuracy = accuracy_score(y_test, y_pred)
```

```
print(f'Test Accuracy: {accuracy:.2f}')
```

Print classification report

```
print("\nClassification Report:\n", classification_report(y_test, y_pred))
```

Generate and plot confusion matrix

```
conf_matrix = confusion_matrix(y_test, y_pred)
```

```
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=['Class 0', 'Class 1'], yticklabels=['Class 0', 'Class 1'])
```

```
plt.xlabel('Predicted')
```

```
plt.ylabel('Actual')
```

```
plt.title('Confusion Matrix')
```

```
plt.show()
```

6. Understanding Bayesian Inference

In Bayesian Learning, the model predicts based on the probabilities:

- **Prior Probability ($P(C)P(C)P(C)$):** The likelihood of each class based on historical data.
- **Likelihood ($P(X|C)P(X|C)P(X|C)$):** The probability of the data given a class.
- **Posterior Probability ($P(C|X)P(C|X)P(C|X)$):** Calculated using Bayes' theorem:

$$P(C|X) = P(X|C) \cdot P(C) / P(X) = \frac{P(X|C)}{\sum P(X|C)}$$

Example: Compute posterior probabilities for the first test sample

```
sample = X_test[0].reshape(1, -1)
posterior_probs = model.predict_proba(sample)
print(f"Sample Features: {sample}")
print(f"Posterior Probabilities: {posterior_probs}")
print(f"Predicted Class: {model.predict(sample)})")
```

Output :

```
Feature_1  Feature_2  Feature_3  Feature_4  Feature_5  Feature_6  \
0 -1.732538  5.260112 -2.952194 -4.603768  2.235848  1.928893
1  2.072914  2.240572 -1.385104 -2.514962 -0.984756  1.436260
2 -0.263106  1.527781 -1.872414 -0.028009  1.612809  3.264194
3 -0.164349 -0.550131 -0.019503 -0.765000  2.273523  2.084217
4 -1.419423  1.015324 -0.864441 -0.009297  0.385404  0.449093

Feature_7  Feature_8  Target
0 -0.101845  3.193487    0
1 -1.255271  2.089872    0
2 -1.296421  1.537870    0
3 -0.321931  0.426253    0
4 -0.029007 -1.982917    1
Test Accuracy: 0.84

Classification Report:
precision      recall   f1-score   support
          0       0.88      0.80      0.84      99
          1       0.82      0.89      0.85     101

accuracy           0.84      200
macro avg       0.85      0.84      0.84      200
weighted avg     0.85      0.84      0.84      200
```

