

<b>Ex. No: 1</b>	<b>A python program to implement Univariate Regression, Bivariate Regression and Multivariate Regression</b>
<b>Date:24/01/25</b>	

### AIM:

To implement a python program using univariate, bivariate and multivariate regression features for a given iris dataset.

### ALGORITHM:

**STEP 1:** Start the program

**STEP 2:** Import the necessary libraries: pandas for data manipulation, numpy for numerical operations, matplotlib.pyplot for plotting, and LinearRegression from sklearn.linear\_model.

**STEP 3:** Read the dataset using pandas.read\_csv() and store it in a variable such as data.

**STEP 4:** Prepare the data by extracting the independent variable(s) X and the dependent variable y, and reshape them into 2D arrays if required.

**STEP 5:** Perform univariate, bivariate, and multivariate linear regression using either numpy.polyfit or sklearn's LinearRegression, then make predictions and calculate the R-squared value to assess model accuracy.

**STEP 6:** Visualize the results using 2D/3D scatter plots and regression lines or planes, and display the coefficients, intercepts, and R-squared values for each model.

**STEP 7:** Stop

### PROGRAM:

```
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np
df=pd.read_csv(r'C:\Users\221801049\Iris.csv')
df.head(150)
df.shape
```

### OUTPUT:

```
(150, 6)
```

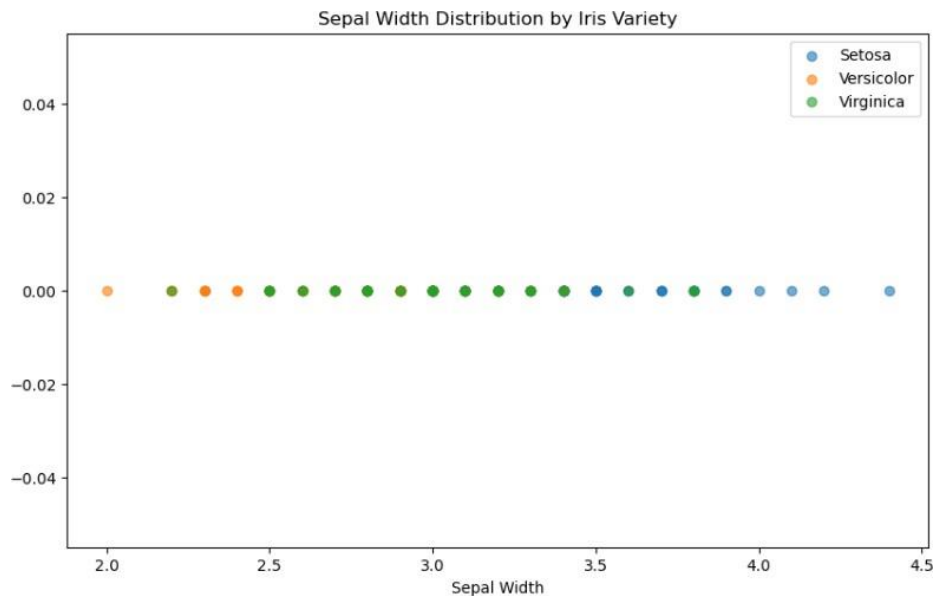
### UNIVARIATE FOR SEPAL WIDTH

```
df_Setosa = df[df['Species'] == 'Iris-setosa']
df_Virginica = df[df['Species'] == 'Iris-virginica']
df_Versicolor = df[df['Species'] == 'Iris-versicolor']
plt.figure(figsize=(10, 6))
```

```
plt.scatter(df_Setosa['SepalWidthCm'], np.zeros_like(df_Setosa['SepalWidthCm']), label='Setosa',
                                                    alpha=0.6)
plt.scatter(df_Versicolor['SepalWidthCm'], np.zeros_like(df_Versicolor['SepalWidthCm']),
                                                    label='Versicolor', alpha=0.6)
plt.scatter(df_Virginica['SepalWidthCm'], np.zeros_like(df_Virginica['SepalWidthCm']),
                                                    label='Virginica', alpha=0.6)

plt.xlabel('Sepal Width')
plt.title('Sepal Width Distribution by Iris Variety')
plt.legend()
plt.show()
```

## OUTPUT:

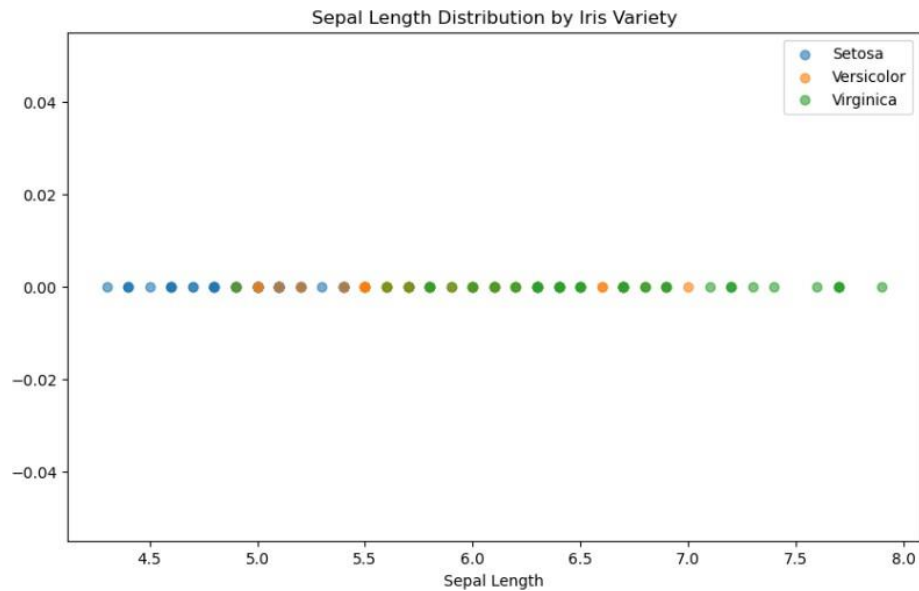


## UNIVARIATE FOR SEPAL LENGTH

```
plt.figure(figsize=(10, 6))
plt.scatter(df_Setosa['SepalLengthCm'], np.zeros_like(df_Setosa['SepalLengthCm']),
                                                    label='Setosa',alpha=0.6)
plt.scatter(df_Versicolor['SepalLengthCm'], np.zeros_like(df_Versicolor['SepalLengthCm']),
                                                    label='Versicolor', alpha=0.6)
plt.scatter(df_Virginica['SepalLengthCm'], np.zeros_like(df_Virginica['SepalLengthCm']),
                                                    label='Virginica', alpha=0.6)

plt.xlabel('Sepal Length')
plt.title('Sepal Length Distribution by Iris Variety')
plt.legend()
plt.show()
```

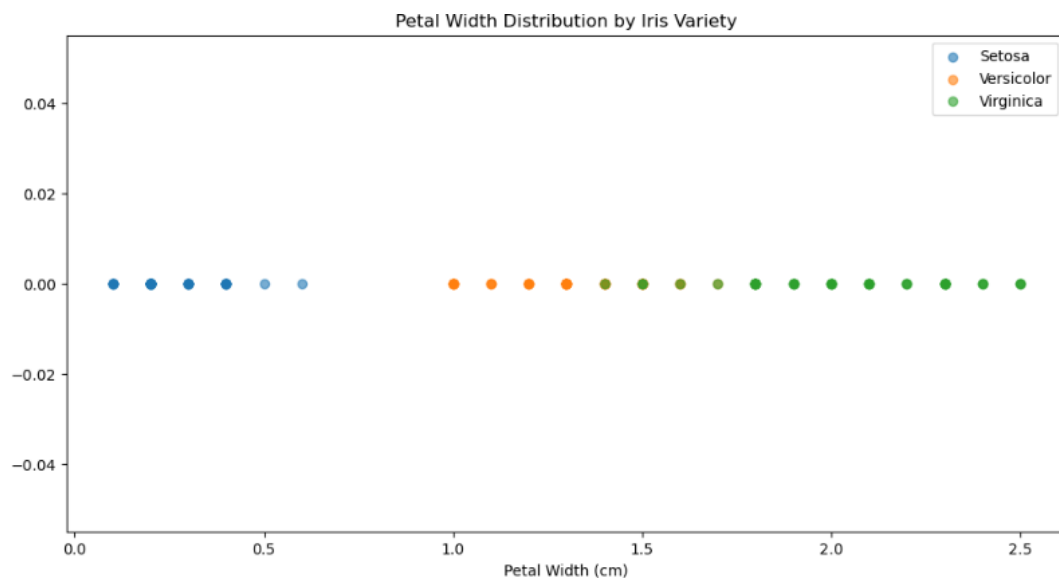
## OUTPUT:



## UNIVARIATE FOR PETAL WIDTH

```
plt.figure(figsize=(12, 6))
plt.scatter(df_Setosa['PetalWidthCm'], np.zeros_like(df_Setosa['PetalWidthCm']), label='Setosa', alpha=0.6)
plt.scatter(df_Versicolor['PetalWidthCm'], np.zeros_like(df_Versicolor['PetalWidthCm']), label='Versicolor',
alpha=0.6)
plt.scatter(df_Virginica['PetalWidthCm'], np.zeros_like(df_Virginica['PetalWidthCm']), label='Virginica',
alpha=0.6)
plt.xlabel('Petal Width (cm)')
plt.title('Petal Width Distribution by Iris Variety')
plt.legend()
plt.show()
```

## OUTPUT:

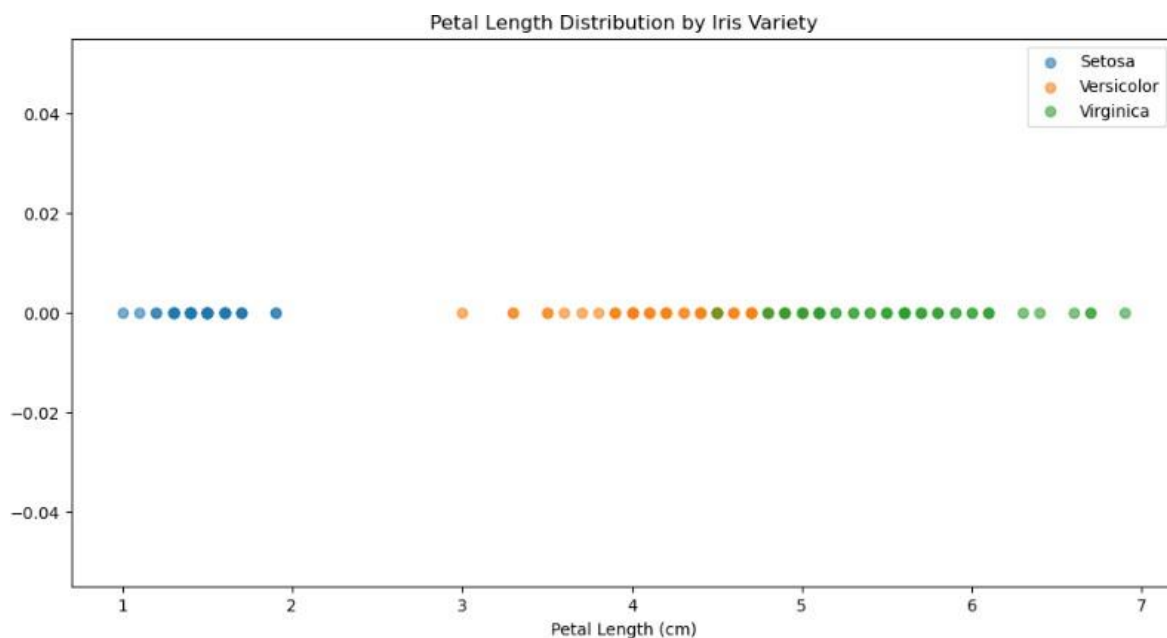


## UNIVARIATE FOR PETAL LENGTH

```
plt.figure(figsize=(12, 6))
plt.scatter(df_Setosa['PetalLengthCm'], np.zeros_like(df_Setosa['PetalLengthCm']), label='Setosa', alpha=0.6)
plt.scatter(df_Versicolor['PetalLengthCm'], np.zeros_like(df_Versicolor['PetalLengthCm']), label='Versicolor',
            alpha=0.6)
plt.scatter(df_Virginica['PetalLengthCm'], np.zeros_like(df_Virginica['PetalLengthCm']), label='Virginica',
            alpha=0.6)

plt.xlabel('Petal Length (cm)')
plt.title('Petal Length Distribution by Iris Variety')
plt.legend()
plt.show()
```

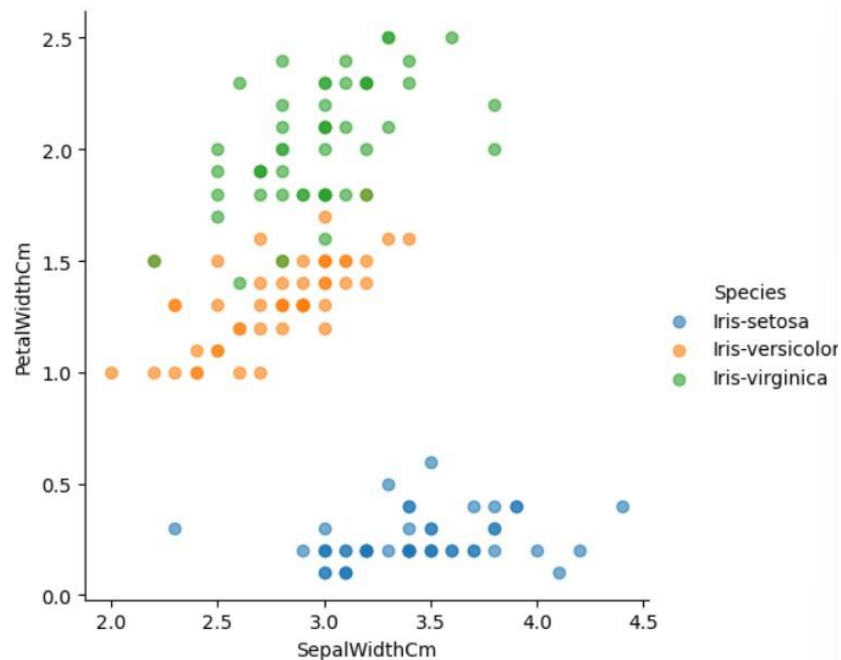
## OUTPUT:



## BIVARIATE SEPAL.WIDTH VS PETAL.WIDTH

```
g = sns.FacetGrid(df, hue='Species', height=5)
g.map(plt.scatter, 'SepalWidthCm', 'PetalWidthCm', alpha=0.6)
g.add_legend()
plt.show()
```

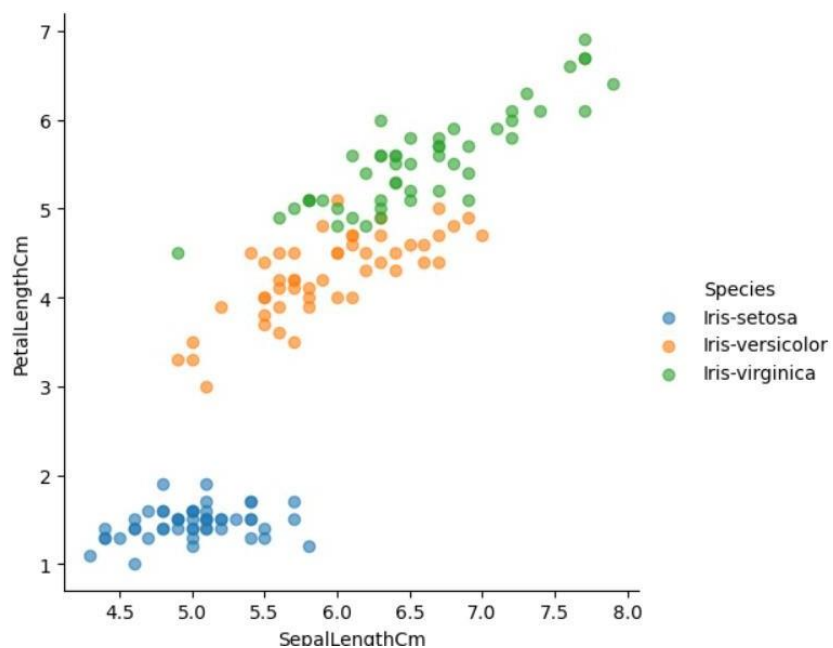
## OUTPUT:



## BIVARIATE SEPAL.LENGTH VS PETAL.LENGTH

```
g = sns.FacetGrid(df, hue='Species', height=5)
g.map(plt.scatter, 'SepalLengthCm', 'PetalLengthCm', alpha=0.6)
g.add_legend()
plt.show()
```

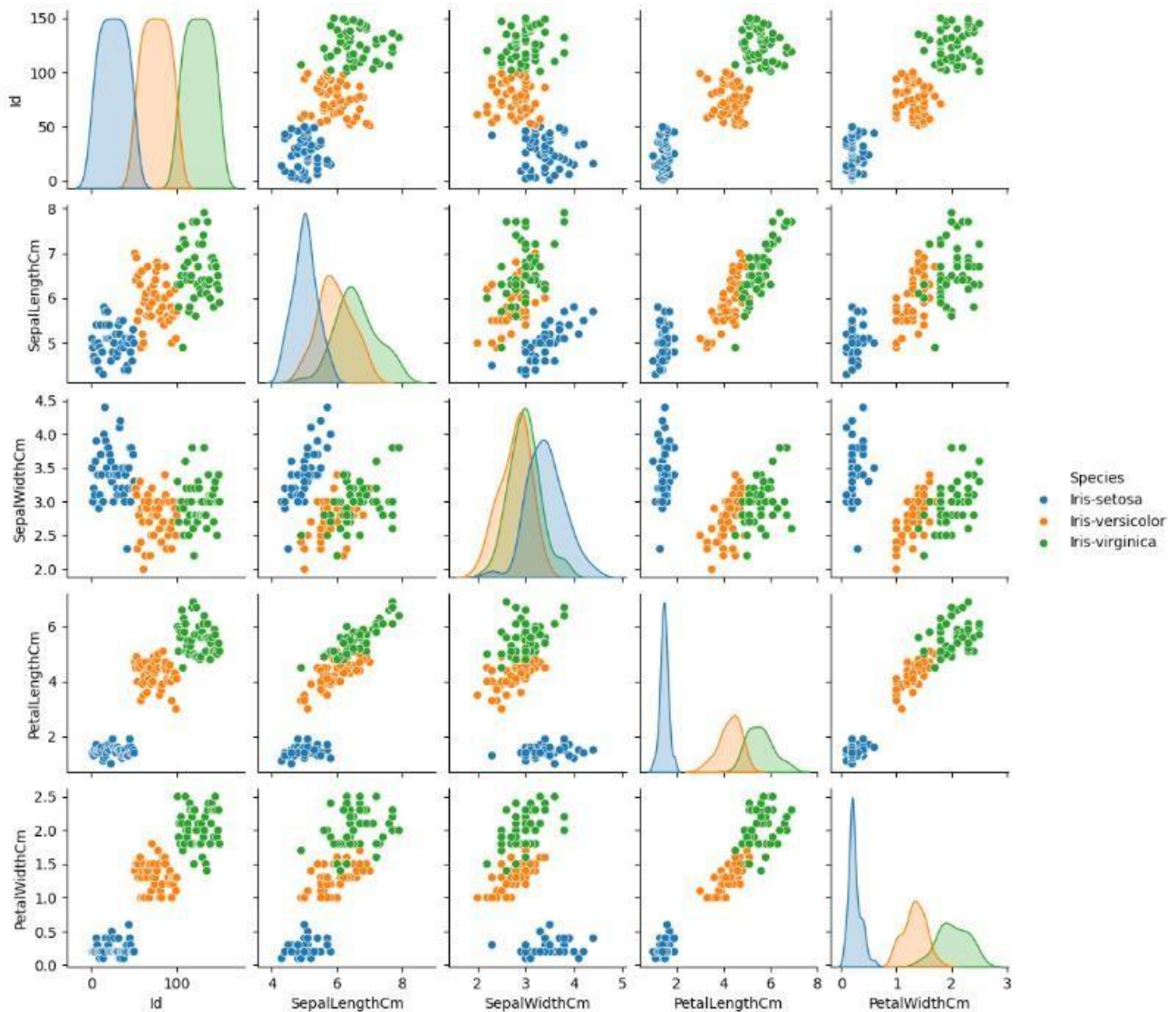
## OUTPUT:



## MULTIVARIATE ALL THE FEATURES

```
g=sns.pairplot(df,hue='Species',height=2)  
plt.show()
```

### OUTPUT:



### RESULT:

Thus, the python program to implement univariate, bivariate and multivariate regression features for the given iris dataset is analyzed and the features are plotted using scatter plot.

<b>Ex. No: 2</b>	<b>A python program to implement Simple linear regression using Least Square Method</b>
<b>Date:31/01/25</b>	

**AIM:**

To implement a python program for constructing a simple linear regression using least square method.

**ALGORITHM:**

**STEP 1:** Start the program

**STEP 2:** Import pandas for data handling and matplotlib.pyplot for plotting. Read the dataset using pandas.read\_csv() and store it in a variable like data.

**STEP 3:** Extract the independent variable X and dependent variable y, and reshape them into 2D arrays if necessary.

**STEP 4:** Calculate the mean of X and y, then compute the slope (m) and intercept (b) using the formulas for simple linear regression.

**STEP 5:** Use the slope and intercept to make predictions for each X value. Calculate the Total Sum of Squares (TSS), Residual Sum of Squares (RSS), and the R-squared value to assess model accuracy.

**STEP 6:** Plot the original data points as a scatter plot and the regression line using the predicted values.

**STEP 7:** Print the calculated slope, intercept, and R-squared value to complete the program.

**STEP 8:** Stop

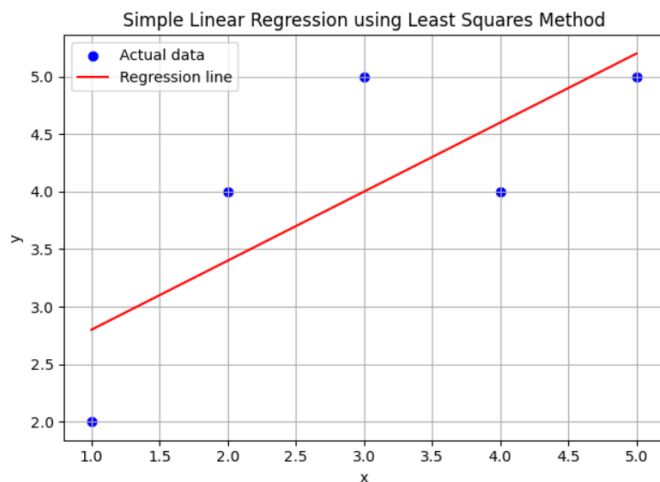
**PROGRAM:**

```
import numpy as np
import matplotlib.pyplot as plt
# Sample data
x = np.array([1, 2, 3, 4, 5], dtype=float)
y = np.array([2, 4, 5, 4, 5], dtype=float)
# Calculate means
x_mean = np.mean(x)
y_mean = np.mean(y)
# Calculate coefficients
numerator = np.sum((x - x_mean) * (y - y_mean))
denominator = np.sum((x - x_mean)**2)
slope = numerator / denominator
intercept = y_mean - slope * x_mean
# Prediction function
def predict(x_val):
    return slope * x_val + intercept
# Predict y values
```

```
y_pred = predict(x)
# Print results
print(f'Slope (m): {slope}')
print(f'Intercept (b): {intercept}')
print("Regression Line Equation: y = {:.2f}x + {:.2f}".format(slope, intercept))
# Plotting
plt.scatter(x, y, color='blue', label='Actual data')
plt.plot(x, y_pred, color='red', label='Regression line')
plt.xlabel('x')
plt.ylabel('y')
plt.title('Simple Linear Regression using Least Squares Method')
plt.legend()
plt.grid(True)
plt.show()
```

### OUTPUT:

```
Slope (m): 0.6
Intercept (b): 2.2
Regression Line Equation: y = 0.60x + 2.20
```



### RESULT:

Thus, the python program for constructing a simple linear regression using least square method has been implemented and executed successfully.



<b>Ex. No: 3</b>	<b>A python program to implement Logistic Model</b>
<b>Date:07/02/25</b>	

**AIM:**

To implement python program for the logistic model using DigitalAd dataset.

**ALGORITHM:**

**STEP 1:** Start the program

**STEP 2:** Load the dataset

**STEP 3:** Display the shape and first 5 rows of the dataset to understand its structure

**STEP 4:** Separate features (X) and target (Y) and split the data into training and testing sets

**STEP 5:** Train a Logistic Regression model and Make predictions on the test set

**STEP 6:** Evaluate the model using confusion matrix and Print the accuracy of the model

**STEP 7:** Predict for a new customer and Output the prediction result

**STEP 8:** Stop the program

**PROGRAM:**

```
import pandas as pd
import numpy as np
data = pd.read_csv("DigitalAd_dataset.csv")
print(data.shape)
print(data.head(5))
X = data.iloc[:, :-1].values # All columns except the last one (features)
Y = data.iloc[:, -1].values # Last column as target

from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, Y, test_size=0.25, random_state=0)

from sklearn.preprocessing import StandardScaler # Feature scaling (standardizing the features)
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

from sklearn.linear_model import LogisticRegression # Train a Logistic Regression model
model = LogisticRegression(random_state=0)
model.fit(X_train, y_train)

y_pred = model.predict(X_test) # Make predictions on the test set
from sklearn.metrics import confusion_matrix, accuracy_score
cm = confusion_matrix(y_test, y_pred) # Evaluate the model using confusion matrix
print("Confusion Matrix: ")
```

```

print(cm)
print("Accuracy of the Model: {0}%".format(accuracy_score(y_test, y_pred) * 100))
age = int(input("Enter New Customer Age: "))      # Predict for a new customer
sal = int(input("Enter New Customer Salary: "))
newCust = [[age, sal]]
result = model.predict(sc.transform(newCust))      # Output the prediction result
print(result)
if result == 1:
    print("Customer will Buy")
else:
    print("Customer won't Buy")

```

### OUTPUT:

```

(400, 3)
  Age  Salary  Status
0   18   82000       0
1   29   80000       0
2   47   25000       1
3   45   26000       1
4   46   28000       1
Confusion Matrix:
[[61  0]
 [20 19]]
Accuracy of the Model: 80.0%
Enter New Customer Age: 45
Enter New Customer Salary: 45000
[0]
Customer won't Buy

```

### RESULT:

Thus, the python program to implement logistic regression for the given dataset is analyzed and the performance of the developed model is measured successfully.

<b>Ex. No: 4</b>	<b>A python program to implement Single Layer Perceptron</b>
<b>Date:14/02/25</b>	

**AIM:**

To implement python program for the single layer perceptron.

**STEP 1:** Start the program.

**STEP 2:** Import the necessary packages: numpy and matplotlib.

**STEP 3:** Prepare or read a simple dataset (input features X and labels Y).

**STEP 4:** Initialize the perceptron parameters: weights and bias.

**STEP 5:** Define the activation function (step function).

**STEP 6:** Train the perceptron. For each epoch (iteration), Calculate the output.

**STEP 7:** After training, display the final weights and bias.

**STEP 8:** Visualize the decision boundary using matplotlib.

**STEP 9:** Stop the program.

**PROGRAM:**

```
import numpy as np
import matplotlib.pyplot as plt
X = np.array([
    [0, 0],
    [0, 1],
    [1, 0],
    [1, 1]])
Y = np.array([0, 1, 1, 1]) # OR gate outputs
weights = np.zeros(X.shape[1])
bias = 0
learning_rate = 0.1
epochs = 10
def activation_fn(x):
    return 1 if x >= 0 else 0
for epoch in range(epochs):
    print(f"\nEpoch {epoch+1}")
    for i in range(len(X)):
        linear_output = np.dot(X[i], weights) + bias
        y_predicted = activation_fn(linear_output)
        error = Y[i] - y_predicted
        weights += learning_rate * error * X[i]
        bias += learning_rate * error
```

```

    print(f"Input: {X[i]}, Predicted: {y_predicted}, Error: {error}, Updated Weights: {weights}, Updated
Bias: {bias}")
print("\nFinal Weights:", weights)
print("Final Bias:", bias)
for i in range(len(X)):
    if Y[i] == 0:
        plt.scatter(X[i][0], X[i][1], color='red', marker='o')
    else:
        plt.scatter(X[i][0], X[i][1], color='blue', marker='x')
x_values = [np.min(X[:, 0] - 1), np.max(X[:, 0] + 1)]
y_values = -(weights[0] * np.array(x_values) + bias) / weights[1]
plt.plot(x_values, y_values, label='Decision Boundary')
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('Single Layer Perceptron Decision Boundary')
plt.legend()
plt.grid(True)
plt.show()

```

## OUTPUT

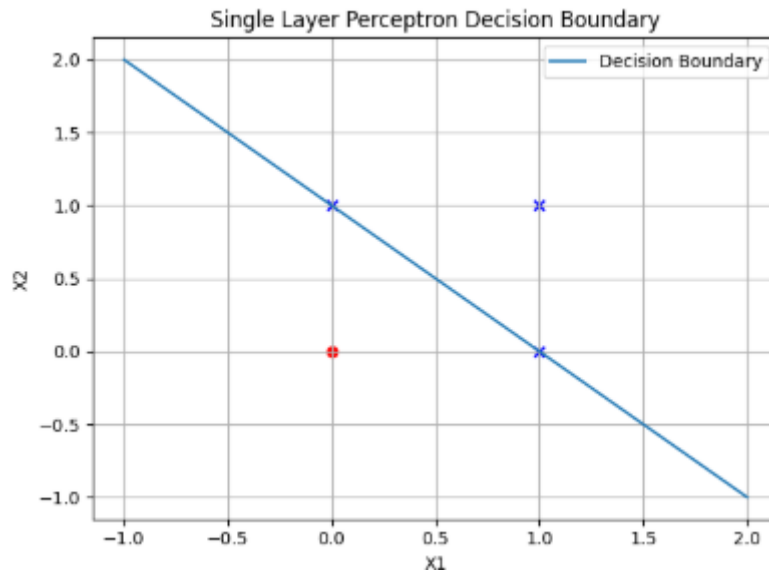
```

Epoch 1
Input: [0 0], Predicted: 1, Error: -1, Updated Weights: [0. 0.], Updated Bias: -0.1
Input: [0 1], Predicted: 0, Error: 1, Updated Weights: [0. 0.1], Updated Bias: 0.0
Input: [1 0], Predicted: 1, Error: 0, Updated Weights: [0. 0.1], Updated Bias: 0.0
Input: [1 1], Predicted: 1, Error: 0, Updated Weights: [0. 0.1], Updated Bias: 0.0

Epoch 2
Input: [0 0], Predicted: 1, Error: -1, Updated Weights: [0. 0.1], Updated Bias: -0.1
Input: [0 1], Predicted: 1, Error: 0, Updated Weights: [0. 0.1], Updated Bias: -0.1
Input: [1 0], Predicted: 0, Error: 1, Updated Weights: [0.1 0.1], Updated Bias: 0.0
Input: [1 1], Predicted: 1, Error: 0, Updated Weights: [0.1 0.1], Updated Bias: 0.0

Epoch 3
Input: [0 0], Predicted: 1, Error: -1, Updated Weights: [0.1 0.1], Updated Bias: -0.1
Input: [0 1], Predicted: 1, Error: 0, Updated Weights: [0.1 0.1], Updated Bias: -0.1
Input: [1 0], Predicted: 1, Error: 0, Updated Weights: [0.1 0.1], Updated Bias: -0.1
Input: [1 1], Predicted: 1, Error: 0, Updated Weights: [0.1 0.1], Updated Bias: -0.1

```

**RESULT:**

Thus, the python program Single Layer Perceptron was successfully implemented using Python.

<b>Ex. No: 5</b>	<b>A python program to implement Multilayer Perceptron with Back Propagation</b>
<b>Date:21/02/25</b>	

**AIM:**

To implement multilayer perceptron with back propagation using python.

**PROCEDURE:**

**STEP 1:** Start the program

**STEP 2:** Create a class MLP with an `__init__` method that initializes weights and biases for the input-to-hidden and hidden-to-output layers.

**STEP 3:** Include activation functions like sigmoid and its derivative and Implement the `train()` method which performs forward propagation to calculate predictions, backward propagation to compute gradients, and updates weights and biases using gradient descent over a number of epochs.

**STEP 4:** Set up an XOR dataset with 2 inputs and 1 output. Instantiate the MLP with 2 input nodes, 4 hidden nodes, and 1 output node. Train the model using the XOR data for 10,000 epochs.

**STEP 5:** After training, pass the input data through the network again using the learned weights to generate predictions.

**STEP 6:** print the outputs to observe how well the model learned the XOR logic.

**STEP 7:** Stop the program

**PROGRAM:**

```
import numpy as np
class MLP:
    def __init__(self, input_size, hidden_size, output_size):
        self.input_size = input_size
        self.hidden_size = hidden_size
        self.output_size = output_size

        # initialize weights matrix and biases
        self.W_input_hidden = np.random.rand(self.input_size, self.hidden_size)
        self.b_input_hidden = np.zeros((1, self.hidden_size))
        self.W_hidden_output = np.random.rand(self.hidden_size, self.output_size)
        self.b_hidden_output = np.zeros((1, self.output_size))
    def sigmoid(self, x):
        return 1 / (1 + np.exp(-x))
    def d_sigmoid(self, x):
        return x * (1 - x)
    def train(self, input_data, target, epochs=1000, lr=0.2):
        for epoch in range(epochs):
            # Forward propagation
```

```

hidden_layer_input = np.dot(input_data, self.W_input_hidden) + self.b_input_hidden
hidden_layer_output = self.sigmoid(hidden_layer_input)
output_layer_input = np.dot(hidden_layer_output, self.W_hidden_output) +
                                                                self.b_hidden_output
output = self.sigmoid(output_layer_input)

# Backward propagation
output_error = target - output
output_grad = output_error * self.d_sigmoid(output)
hidden_error = np.dot(output_grad, self.W_hidden_output.T)
hidden_grad = hidden_error * self.d_sigmoid(hidden_layer_output)

# Update weights and biases using gradient descent
self.W_hidden_output += np.dot(hidden_layer_output.T, output_grad) * lr
self.b_hidden_output += np.sum(output_grad, axis=0, keepdims=True) * lr
self.W_input_hidden += np.dot(input_data.T, hidden_grad) * lr
self.b_input_hidden += np.sum(hidden_grad, axis=0, keepdims=True) * lr

# Optionally, print error every 1000 epochs
if epoch % 1000 == 0:
    error = np.mean(np.square(target - output)) # Mean Squared Error
    print(f'Epoch {epoch}, Error: {error}')

# Example usage:
if __name__ == "__main__":
    # XOR problem: 4 samples, 2 input features, 1 output
    X = np.array([[0, 0], [0, 1], [1, 0], [1, 1]])
    y = np.array([[0], [1], [1], [0]])

    # Create MLP with 2 input nodes, 4 hidden nodes, and 1 output node
    mlp = MLP(input_size=2, hidden_size=4, output_size=1)

    mlp.train(X, y, epochs=10000) # Train the model

    # Test the model after training
    print("Predictions after training:")
    hidden_layer_input = np.dot(X, mlp.W_input_hidden) + mlp.b_input_hidden
    hidden_layer_output = mlp.sigmoid(hidden_layer_input)
    output_layer_input = np.dot(hidden_layer_output, mlp.W_hidden_output) + mlp.b_hidden_output
    predictions = mlp.sigmoid(output_layer_input)
    print(predictions)

```

## OUTPUT:

```
Epoch 0, Error: 0.3525487770976961
Epoch 1000, Error: 0.16546836123157288
Epoch 2000, Error: 0.022134675143509714
Epoch 3000, Error: 0.006844220579949708
Epoch 4000, Error: 0.0036914169964121224
Epoch 5000, Error: 0.0024534156381819903
Epoch 6000, Error: 0.001812478058253225
Epoch 7000, Error: 0.0014263384514248863
Epoch 8000, Error: 0.0011704001423583428
Epoch 9000, Error: 0.000989268984991053
Predictions after training:
[[0.03327479]
 [0.97289057]
 [0.97188183]
 [0.02804434]]
```

## RESULT:

Thus, the Python program to implement a simple multi-layer perceptron (MLP) for the XOR problem is analyzed, and the model successfully predicts the output after training by learning the non-linear relationships in the data.



<b>Ex. No: 6</b>	<b>A python program to do face recognition using SVM classifier</b>
<b>Date: 28/02/25</b>	

**AIM:**

To implement a SVM classifier model using python and determine its accuracy.

**PROCEDURE:**

**STEP 1:** Start the program.

**STEP 2:** Import the necessary libraries and Load the dataset.

**STEP 3:** Flatten the images and apply PCA

**STEP 4:** Split the dataset into train & test sets and train SVM model

**STEP 5:** Make prediction and evaluate the accuracy.

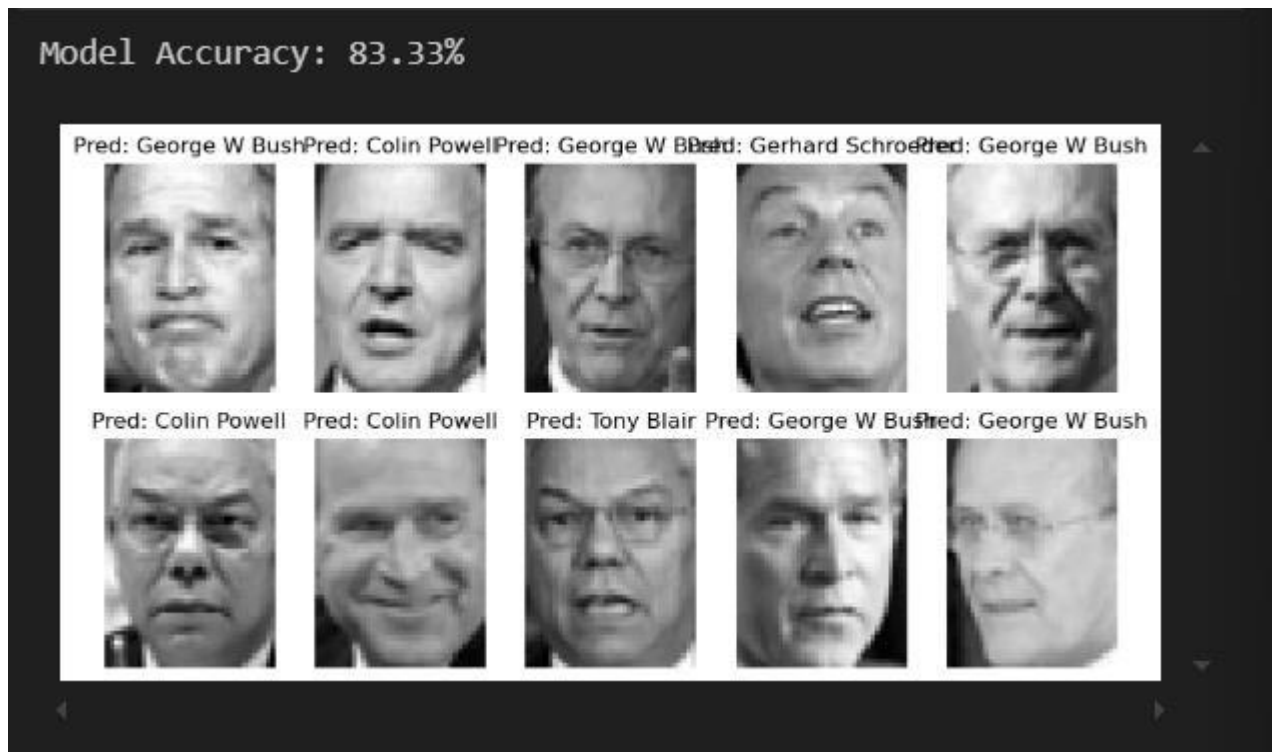
**STEP 6:** Print the model accuracy and visualize some test images with predictions.

**STEP 7:** Stop the program.

**PROGRAM:**

```
import cv2
import numpy as np
import matplotlib.pyplot as plt
from sklearn.svm import SVC
from sklearn.datasets import fetch_lfw_people
from sklearn.model_selection import train_test_split
from sklearn.decomposition import PCA
from sklearn.metrics import accuracy_score
lfw = fetch_lfw_people(min_faces_per_person=100, resize=0.4, download_if_missing=True)
X, y = lfw.images, lfw.target # Images and labels
X_flat = X.reshape(X.shape[0], -1) # Convert images to 1D array
pca = PCA(n_components=100).fit(X_flat) # Reduce dimensions to 100 principal components
X_pca = pca.transform(X_flat)
X_train, X_test, y_train, y_test = train_test_split(X_pca, y, test_size=0.2, random_state=42)
svm = SVC(kernel='linear')
svm.fit(X_train, y_train)
y_pred = svm.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'Model Accuracy: {accuracy * 100:.2f}%')
fig, axes = plt.subplots(2, 5, figsize=(10, 5)) # Create a grid of 2 rows, 5 columns
for i, ax in enumerate(axes.ravel()):
    ax.imshow(X[i], cmap='gray') # Show actual face image
    ax.set_title(f'Pred: {lfw.target_names[y_pred[i]]}') # Predicted name
    ax.axis('off')
plt.show()
```

## OUTPUT:



## RESULT:

Thus the python program to implement SVM classifier model has been executed successfully and the classified output has been analyzed for the given dataset.\\

<b>Ex. No: 7</b>	<b>A python program to implement Decision Tree</b>
<b>Date:07/03/2025</b>	

### AIM:

To build, train, and visualize a Decision Tree Classifier using the Iris dataset in Python, and to evaluate the model's performance based on its accuracy on the test data.

### ALGORITHM:

**STEP 1:** Start the program.

**STEP 2:** Import necessary modules: datasets, train\_test\_split, DecisionTreeClassifier, accuracy\_score, and matplotlib.pyplot.

**STEP 3:** Load the Iris dataset and separate it into features (X) and target labels (y).

**STEP 4:** Split the dataset into training and testing sets using train\_test\_split().

**STEP 5:** Create an instance of DecisionTreeClassifier, train (fit) it using the training data (X\_train, y\_train).

**STEP 6:** Predict the target labels for the test set using the trained classifier.

**STEP 7:** Calculate and print the model's accuracy using accuracy\_score().

**STEP 8:** Visualize the trained Decision Tree using plot\_tree() and display it with plt.show().

**STEP 9:** Stop the program.

### PROGRAM:

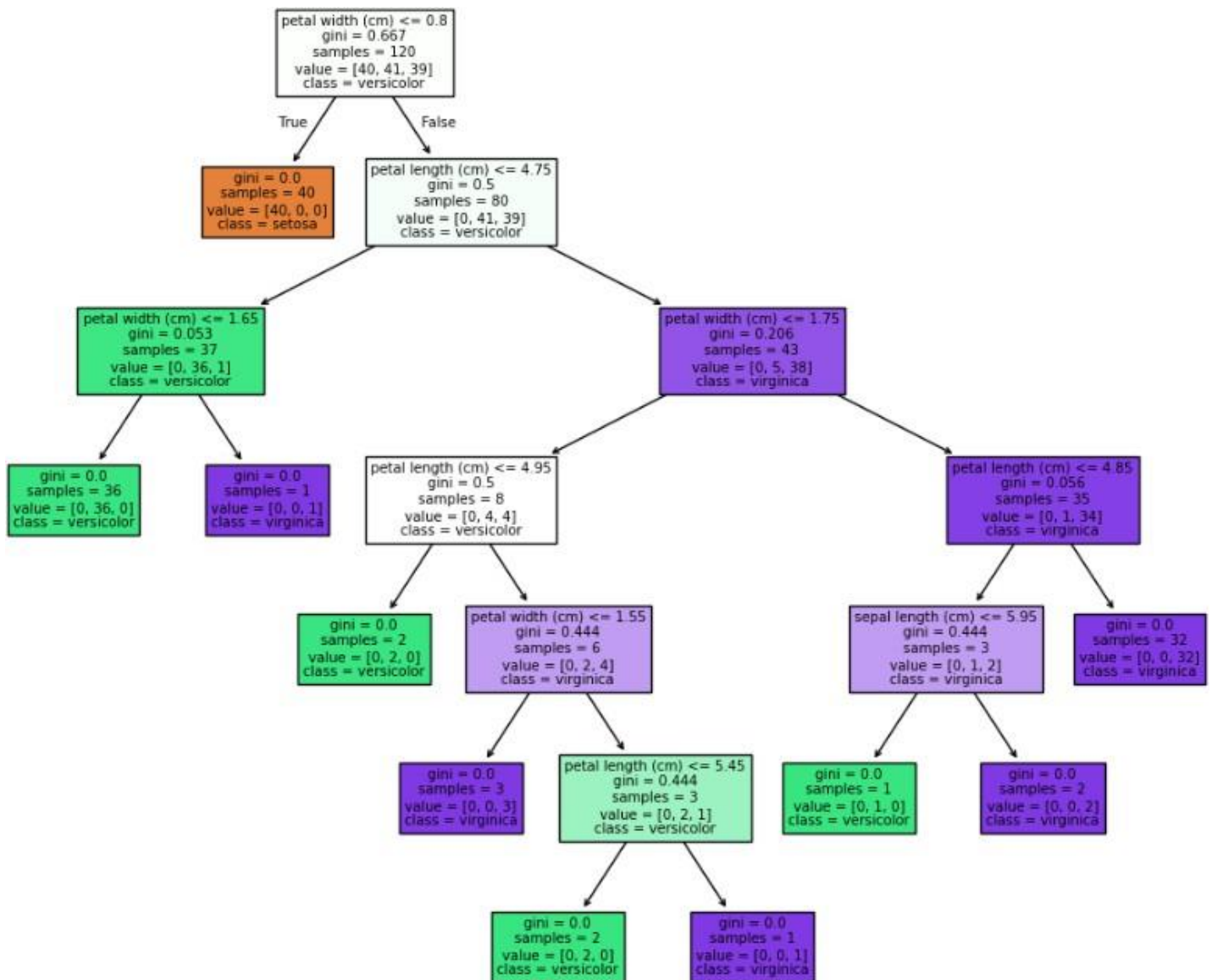
```

from sklearn import datasets
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier, plot_tree
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
iris = datasets.load_iris()
X, y = iris.data, iris.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
clf = DecisionTreeClassifier()
clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print(f'Accuracy: {accuracy_score(y_test, y_pred) * 100:.2f}%')
plt.figure(figsize=(12, 10))
plot_tree(clf, filled=True, feature_names=iris.feature_names, class_names=iris.target_names)
plt.show()

```

## OUTPUT:

Accuracy: 100.00%



## RESULT:

Thus, the Python program to build, train, and visualize a Decision Tree Classifier on the Iris dataset was successfully implemented.

Ex. No: 8	A python program to implement Boosting
Date:28/03/25	

**AIM:**

To build and evaluate a python program to implement boosting.

**ALGORITHM:**

**STEP 1:** Start the program.

**STEP 2:** Import required libraries: sklearn, numpy, and matplotlib.

**STEP 3:** Generate a synthetic binary classification dataset using make\_classification().

**STEP 4:** Split the dataset into training and testing sets using train\_test\_split().

**STEP 5:** Initialize the AdaBoostClassifier with a base estimator (DecisionTreeClassifier) and train it on the training data.

**STEP 6:** Use the trained model to make predictions on the test set.

**STEP 7:** Calculate and print the accuracy of the model using accuracy\_score().

**STEP 8:** Visualize the decision boundaries by predicting over a meshgrid and plotting the results.

**STEP 9:** Stop the program.

**PROGRAM:**

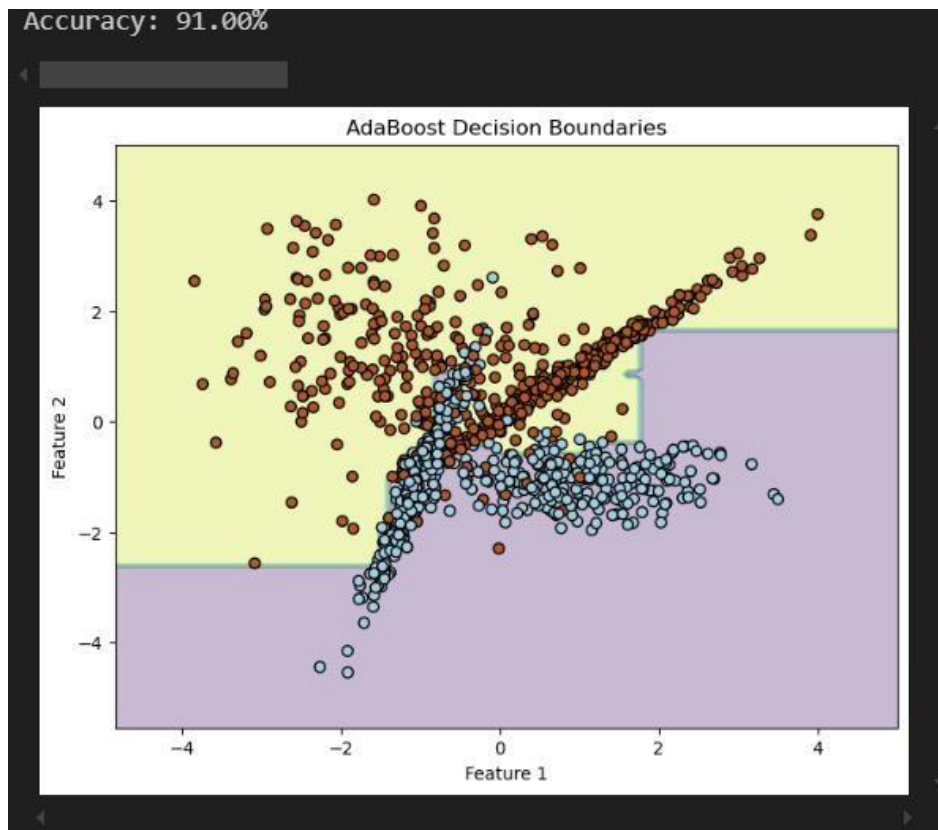
```

from sklearn.ensemble import AdaBoostClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
import numpy as np
X, y = make_classification(n_samples=1000, n_features=2, n_informative=2, n_redundant=0,
random_state=42)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
boost = AdaBoostClassifier(DecisionTreeClassifier(max_depth=1), n_estimators=50,
random_state=42)
boost.fit(X_train, y_train)
y_pred = boost.predict(X_test)          # Predict and evaluate model performance
accuracy = accuracy_score(y_test, y_pred)
print(f"Accuracy: {accuracy * 100:.2f}%")
x_min, x_max = X[:, 0].min() - 1, X[:, 0].max() + 1          # Visualization - Decision Boundaries
y_min, y_max = X[:, 1].min() - 1, X[:, 1].max() + 1
xx, yy = np.meshgrid(np.linspace(x_min, x_max, 100), np.linspace(y_min, y_max, 100))
Z = boost.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)

```

```
plt.figure(figsize=(8, 6))
plt.contourf(xx, yy, Z, alpha=0.3)
plt.scatter(X[:, 0], X[:, 1], c=y, edgecolors="k", cmap=plt.cm.Paired)
plt.title("AdaBoost Decision Boundaries")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.show()
```

### OUTPUT:



### RESULT:

Thus, the Python program for Boosting using AdaBoostClassifier was successfully executed.

<b>Ex. No: 9</b>	<b>A python program to implement KNN and K-Means</b>
<b>Date:04/04/25</b>	

**AIM:**

To implement the K-Nearest Neighbors (KNN) and K Means algorithm on the Iris dataset for classification and evaluate its accuracy.

**ALGORITHM:**

**STEP 1:** Start the program.

**STEP 2:** Import necessary libraries such as sklearn.datasets, train\_test\_split, KNeighborsClassifier, and accuracy\_score.

**STEP 3:** Load the Iris dataset and extract the features (X) and target labels (y).

**STEP 4:** Split the dataset into training and testing sets.

**STEP 5:** Create a KNeighborsClassifier model with a defined value of k (e.g., k = 3).

**STEP 6:** Train the model using the training data.

**STEP 7:** Predict the labels for the test data.

**STEP 8:** Evaluate the model by calculating the accuracy score.

**STEP 9:** Display the results.

**STEP 10:** Stop the program.

**STEP 11:** Import necessary libraries such as sklearn.datasets, KMeans, matplotlib.pyplot, and PCA from sklearn.decomposition.

**STEP 12:** Load the Iris dataset and extract the feature matrix (X).

**STEP 13:** Choose the number of clusters (k = 3 for Iris)

**STEP 14:** Create a KMeans object.

**STEP 15:** Fit the KMeans model on the dataset to form clusters.

**STEP 16:** Retrieve and store the cluster labels.

**STEP 17:** Reduce dimensionality using PCA for visualization (2 components).

**STEP 18:** Plot the clustered data on a 2D scatter plot using the PCA-reduced values.

**STEP 19:** Display the visualization.

**STEP 20:** Stop the program.

**PROGRAM:**

```

from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score
# Load dataset
iris = load_iris()
X, y = iris.data, iris.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
knn = KNeighborsClassifier(n_neighbors=3)

```

```

knn.fit(X_train, y_train)
y_pred = knn.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'KNN Accuracy: {accuracy * 100:.2f}%")
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score
iris = load_iris()
X, y = iris.data, iris.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
knn = KNeighborsClassifier(n_neighbors=3)
knn.fit(X_train, y_train)
y_pred = knn.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'KNN Accuracy: {accuracy * 100:.2f}%")

```

## OUTPUT:

**KNN Accuracy: 100.00%**



## RESULT:

Thus the python program for building classification model using KNN algorithm is executed and verified successfully.



<b>Ex. No: 10</b>	<b>A python program to implement Dimensionality Reduction – PCA</b>
<b>Date:11/04/25</b>	

**AIM:**

To develop a Python program to implement Principal Component Analysis (PCA) for dimensionality reduction using the Digits dataset.

**ALGORITHM:**

**STEP 1:** Start the program.

**STEP 2:** Import the required libraries: numpy, matplotlib.pyplot, sklearn.decomposition.PCA, and sklearn.datasets.load\_digits.

**STEP 3:** Load the Digits dataset using load\_digits() and store the data in variable X.

**STEP 4:** Apply PCA with n\_components=2 using the PCA() class and fit-transform the dataset.

**STEP 5:** Store the result of PCA in X\_pca and visualize the 2D projection of the data using a scatter plot.

**STEP 6:** Color the points in the plot based on their digit labels using a color map for better interpretation.

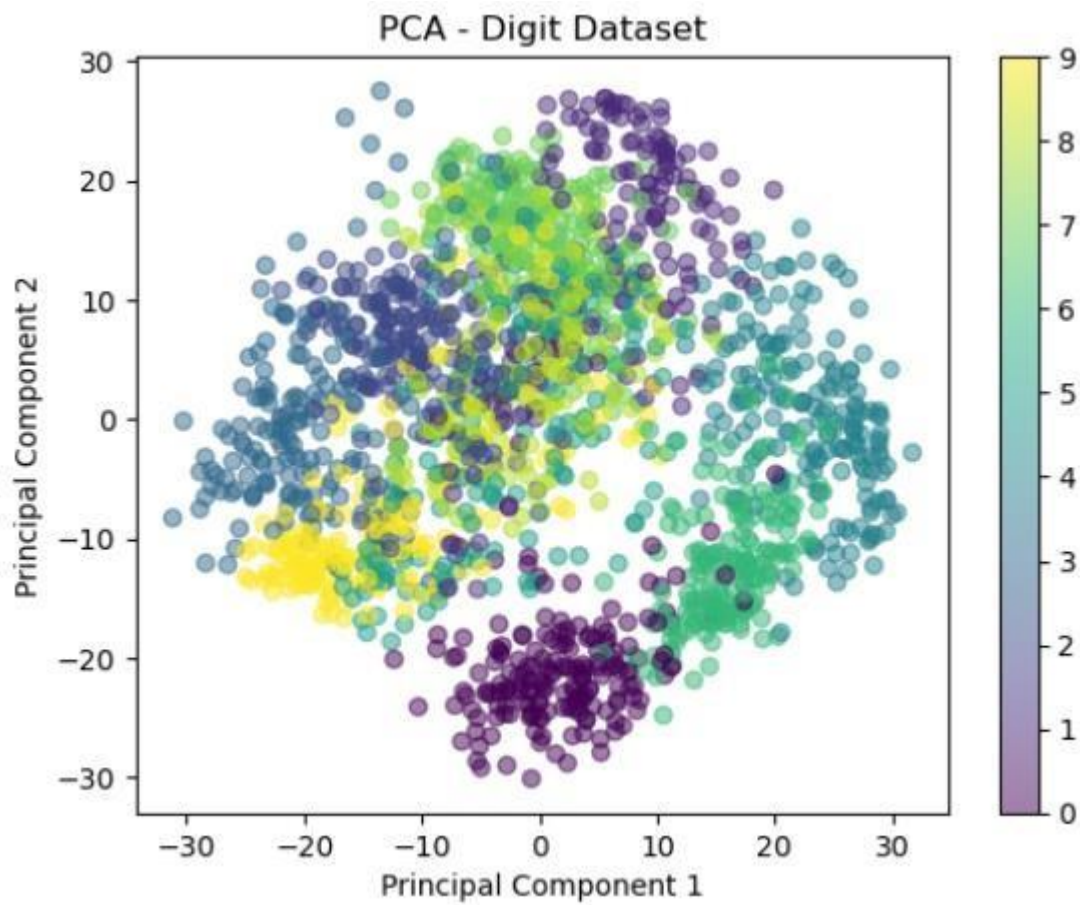
**STEP 7:** Add labels, a colorbar, and a title to the plot for clarity.

**STEP 8:** Stop the program.

**PROGRAM:**

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.decomposition import PCA
from sklearn.datasets import load_digits
# Load dataset
digits = load_digits()
X = digits.data
y = digits.target
# Apply PCA
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X)
# Plot PCA results
plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='viridis', alpha=0.5)
plt.colorbar()
plt.title("PCA - Digit Dataset")
plt.xlabel("Principal Component 1")
plt.ylabel("Principal Component 2")
plt.show()
```

## OUTPUT:



## RESULT:

Thus, the Python program to perform dimensionality reduction using PCA was successfully executed and visualized using the Digits dataset.

Ex. No: 11	Mini project – Develop a simple application using TensorFlow / Keras
Date:11/04/25	

**AIM:**

To develop a Removing Noise from Images Using the TensorFlow.

**ALGORITHM:**

**STEP 1:** Start the program.

**STEP 2:** Add random Gaussian noise to the images and clip values to stay between 0 and 1.

**STEP 3:** Load the MNIST dataset and split it into training and testing sets..

**STEP 4:** Build a Convolutional autoencoder model:

- Use Conv2D and Maxpooling layers for encoding.
- Use Con2D Transpose and Conv2D Layers for Decoding.

**STEP 5:** Compile the model using the adam optimizer and binary crossentropy loss.

**STEP 6:** Train the model with noisy images as input and original images as the output

**STEP 7:** Predict and Display the denoised images alongside their noisy versions

**STEP 8:** Stop the program.

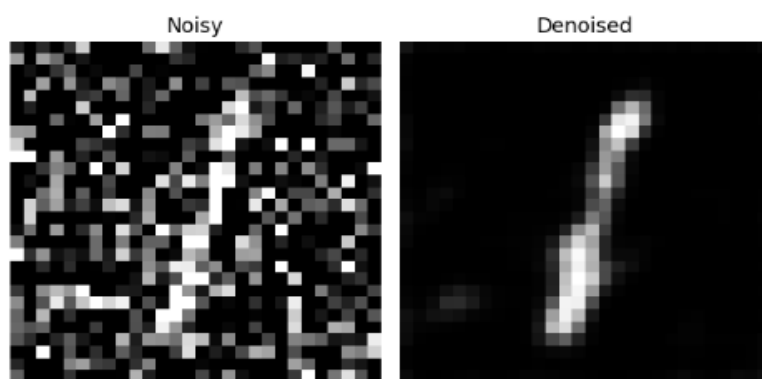
**PROGRAM:**

```
import tensorflow as tf, matplotlib.pyplot as plt
(x,*), (xt,*) = tf.keras.datasets.mnist.load_data()
x = xt[..., None] / 255.
xn = tf.clip_by_value(x + .5*tf.random.normal(tf.shape(x)), 0, 1)
m = tf.keras.Sequential([
    tf.keras.layers.Input((28,28,1)),
    tf.keras.layers.Conv2D(16, 3, padding='same', activation='relu'),
    tf.keras.layers.MaxPool2D(),
    tf.keras.layers.Conv2DTranspose(16, 3, strides=2, padding='same', activation='relu'),
    tf.keras.layers.Conv2D(1, 3, padding='same', activation='sigmoid')
])
m.compile('adam', 'binary_crossentropy')
m.fit(xn, x, epochs=3, batch_size=128, verbose=0)

for i in range(5):
    noisy_img = xn[i].numpy().squeeze()
    denoised_img = m.predict(xn[i:i+1], verbose=0).squeeze()
    plt.subplot(1,2,1)
```

```
plt.imshow(noisy_img, cmap='gray')
plt.axis('off')
plt.title('Noisy')
plt.subplot(1,2,2)
plt.imshow(denoised_img, cmap='gray')
plt.axis('off')
plt.title('Denoised')
plt.show()
```

### OUTPUT:



### RESULT:

Thus, the TensorFlow based application for Removing Noise from Images Using the TensorFlow is developed, executed successfully, and the model's performance is visualized and verified.