# RAJALAKSHMI ENGINEERING COLLEGE

**RAJALAKSHMI NAGAR, THANDALAM – 602 105**

# AI23331 FUNDAMENTALS OF MACHINE LEARNING LAB

**Laboratory Record Notebook**

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**EXPT NO: 1** **A python program to implement univariate regression**

**DATE: 23.08.2024** **bivariate regression and multivariate regression.**

**AIM:**

To write a python program to implement univariate regression, bivariate regression and multivariate regression.

**PROCEDURE:**

Implementing univariate, bivariate, and multivariate regression using the Iris dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import numpy as np

import pandas as pd

import seaborn as sns

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: Load the Iris Dataset**

The Iris dataset can be loaded and display the first few rows of the dataset .

```
# Load the Iris dataset

iris = sns.load_dataset('iris')



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

```
print(iris.head())
```

**OUTPUT :**

```
   sepal_length  sepal_width  petal_length  petal_width species
0           5.1          3.5           1.4          0.2  setosa
1           4.9          3.0           1.4          0.2  setosa
2           4.7          3.2           1.3          0.2  setosa
3           4.6          3.1           1.5          0.2  setosa
4           5.0          3.6           1.4          0.2  setosa
```

### Step 3: Data Preprocessing

Ensure the data is clean and ready for modeling. Since the Iris dataset is clean, minimal preprocessing is needed.

```
# Check for missing values
print(iris.isnull().sum())

# Display the basic statistical details
print(iris.describe())
```

**OUTPUT :**

```
sepal_length    0
sepal_width     0
petal_length    0
petal_width     0
species         0
dtype: int64
       sepal_length  sepal_width  petal_length  petal_width
count    150.000000   150.000000    150.000000   150.000000
mean       5.843333     3.057333      3.758000     1.199333
std        0.828066     0.435866      1.765298     0.762238
min        4.300000     2.000000      1.000000     0.100000
25%        5.100000     2.800000      1.600000     0.300000
50%        5.800000     3.000000      4.350000     1.300000
75%        6.400000     3.300000      5.100000     1.800000
max        7.900000     4.400000      6.900000     2.500000
```

**Step 4: Univariate Regression**

Univariate regression involves predicting one variable based on a single predictor.

**4.1: Select the Features**

Choose one feature (e.g., sepal_length) and one target variable (e.g., sepal_width).

```
X_uni = iris[['sepal_length']]
y_uni = iris['sepal_width']
```

**4.2: Split the Data**

Split the data into training and testing sets.
Fit the linear regression model on the training data.

```
X_uni_train, X_uni_test, y_uni_train, y_uni_test = train_test_split(X_uni, y_uni,
test_size=0.2, random_state=42)
```

**4.3: Train the model**

```
uni_model = LinearRegression()
uni_model.fit(X_uni_train, y_uni_train)
```

```
LinearRegression
LinearRegression()
```

**4.4: Make Predictions**

Use the model to make predictions on the test data.

```
y_uni_pred = uni_model.predict(X_uni_test)
```

**4.5: Evaluate the Model**

Evaluate the model performance using metrics like Mean Squared Error (MSE) and R-squared.

```
print(f'Univariate MSE: {mean_squared_error(y_uni_test, y_uni_pred)}')
print(f'Univariate R-squared: {r2_score(y_uni_test, y_uni_pred)}')
```

**OUTPUT:**

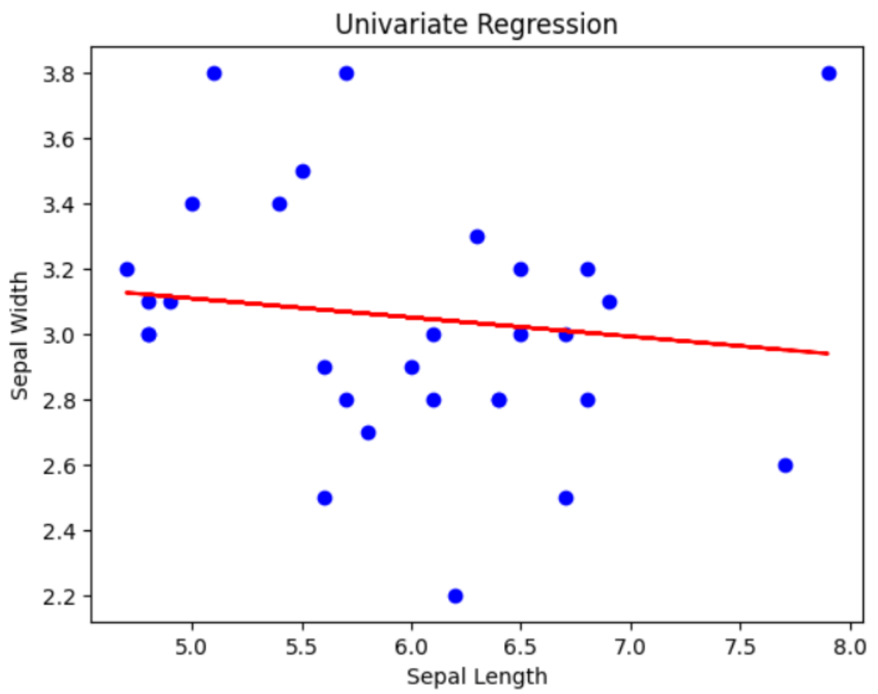
```
Univariate MSE: 0.13961895650579023
Univariate R-squared: 0.024098626473972984
```

### 4.6: Visualize the Results

Visualize the relationship between the predictor and the target variable.

```
plt.scatter(X_uni_test, y_uni_test, color='blue')
plt.plot(X_uni_test, y_uni_pred, color='red')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.title('Univariate Regression')
plt.show()
```

**OUTPUT :**




## Step 5 : Bivariate Regression

Bivariate regression involves predicting one variable based on two predictors.

### 5.1: Select the Features

Choose two features (e.g., sepal_length, petal_length) and one target variable (e.g., sepal_width).

```
X_bi = iris[['sepal_length', 'petal_length']]

y_bi = iris['sepal_width']
```

### 5.2: Split the Data

Split the data into training and testing sets.

```
X_bi_train, X_bi_test, y_bi_train, y_bi_test = train_test_split(X_bi, y_bi,

test_size=0.2, random_state=42)
```

### 5.3: Train the Model

Fit the linear regression model on the training data.

```
bi_model = LinearRegression()

bi_model.fit(X_bi_train, y_bi_train)
```

**OUTPUT :**

```
LinearRegression
LinearRegression()
```

### 5.4: Make Predictions

Use the model to make predictions on the test data.

```
y_bi_pred = bi_model.predict(X_bi_test)
```

### 5.5: Evaluate the Model

Evaluate the model performance using metrics like MSE and R-squared.

```
print(f'Bivariate MSE: {mean_squared_error(y_bi_test, y_bi_pred)}')
print(f'Bivariate R-squared: {r2_score(y_bi_test, y_bi_pred)}')
```

OUTPUT :

```
Bivariate MSE: 0.08308605032913309
Bivariate R-squared: 0.4192494152204116
```

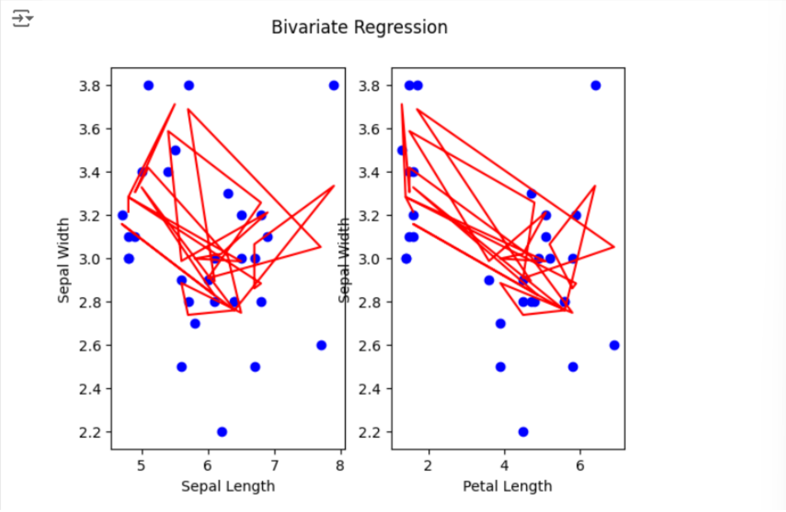
### 5.6: Visualize the Results

Since visualizing in 3D is challenging, we can plot the relationships between the target and each predictor separately.

```
# Sepal Length vs Sepal Width
```

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

plt.scatter(X_bi_test['sepal_length'], y_bi_test, color='blue')

plt.plot(X_bi_test['sepal_length'], y_bi_pred, color='red')

plt.xlabel('Sepal Length')

plt.ylabel('Sepal Width')

# Petal Length vs Sepal Width

plt.subplot(1, 2, 2)

plt.scatter(X_bi_test['petal_length'], y_bi_test, color='blue')

plt.plot(X_bi_test['petal_length'], y_bi_pred, color='red')

plt.xlabel('Petal Length')

plt.ylabel('Sepal Width')

plt.suptitle('Bivariate Regression')

plt.show()
```

**OUTPUT :**



**Step 6: Multivariate Regression**

Multivariate regression involves predicting one variable based on multiple predictors.

**6.1: Select the Features**

Choose multiple features (e.g., sepal_length, petal_length, petal_width) and one target variable (e.g., sepal_width).

```
X_multi = iris[['sepal_length', 'petal_length', 'petal_width']]

y_multi = iris['sepal_width']
```

**6.2: Split the Data**

Split the data into training and testing sets.

```
X_multi_train, X_multi_test, y_multi_train, y_multi_test =
train_test_split(X_multi,

y_multi, test_size=0.2, random_state=42)
```

**6.3: Train the Model**

Fit the linear regression model on the training data.

```
multi_model = LinearRegression()
multi_model.fit(X_multi_train, y_multi_train)
```

**OUTPUT :**

```
▾ LinearRegression
LinearRegression()
```

**6.4: Make Predictions**

Use the model to make predictions on the test data.

```
y_multi_pred = multi_model.predict(X_multi_test)
```

**6.5: Evaluate the Model**

Evaluate the model performance using metrics like MSE and R-squared.

```
    print(f'Multivariate MSE: {mean_squared_error(y_multi_test, y_multi_pred)}')
    print(f'Multivariate R-squared: {r2_score(y_multi_test, y_multi_pred)}')
```

**OUTPUT :**

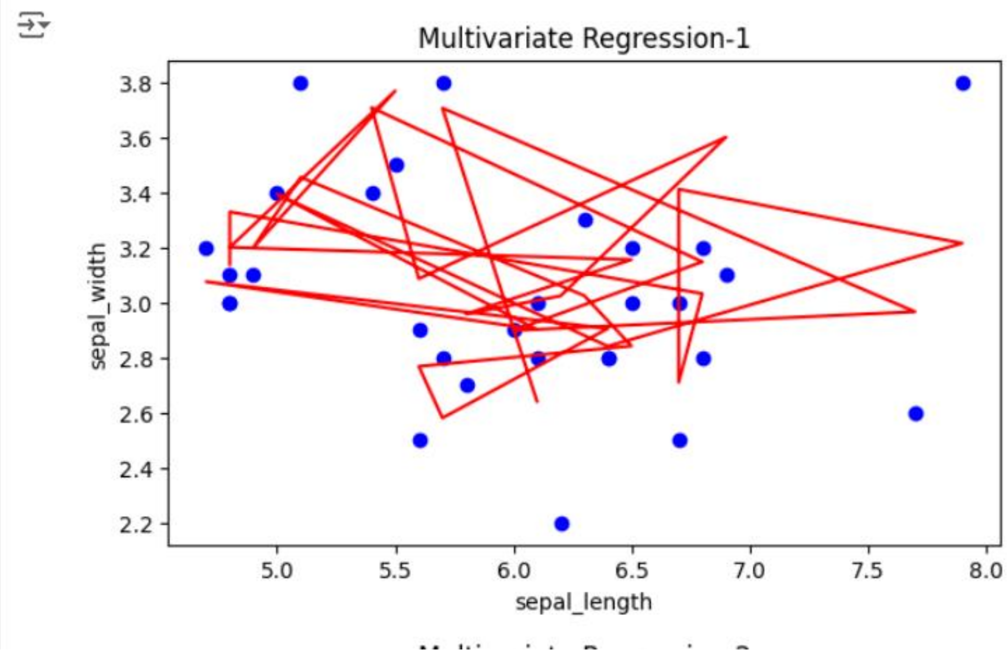
```
Multivariate MSE: 0.0868353771078583
Multivariate R-squared: 0.39304256448374897
```

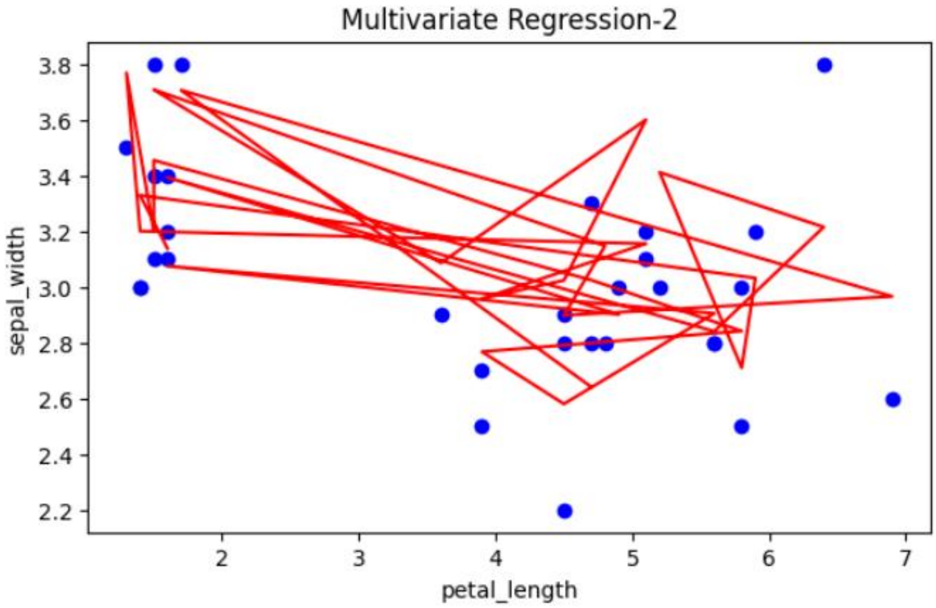
### Step 7: Visualize the multivariate regression

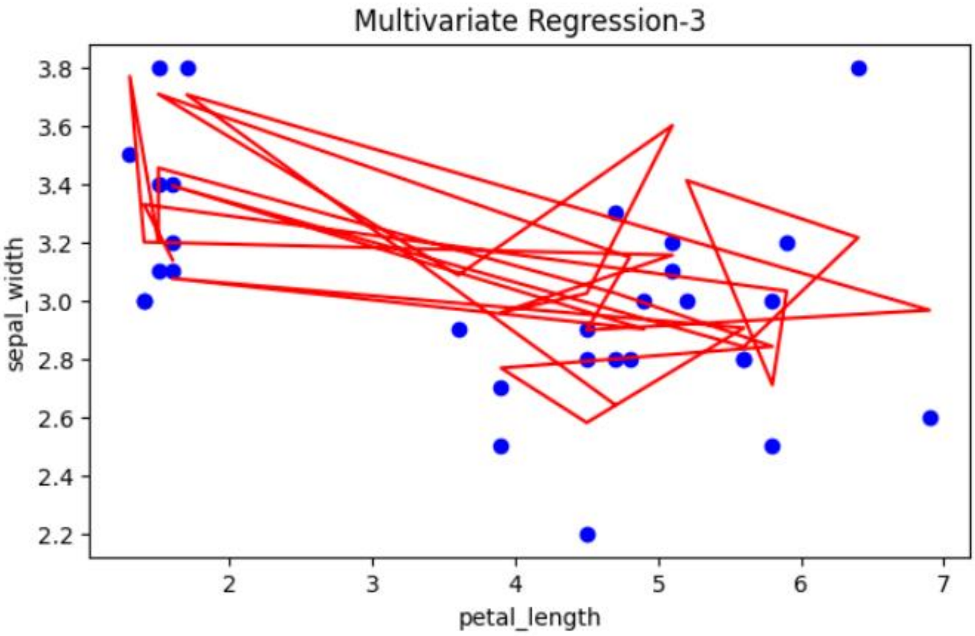
```
plt.figure(figsize=(15,4))

plt.subplot(1, 2, 1)

plt.scatter(X_multi_test['sepal_length'], y_multi_test, color='blue')

plt.plot(X_multi_test['sepal_length'], y_multi_pred, color='red')

plt.xlabel('sepal_length')

plt.ylabel('sepal_width')

plt.title('Multivariate Regression-1')

plt.show()

plt.figure(figsize=(15,4))

plt.subplot(1, 2, 1)

plt.scatter(X_multi_test['petal_length'], y_multi_test, color='blue')

plt.plot(X_multi_test['petal_length'], y_multi_pred, color='red')

plt.xlabel('petal_length')

plt.ylabel('sepal_width')

plt.title('Multivariate Regression-2')

plt.show()

plt.figure(figsize=(15,4))

plt.subplot(1, 2, 2 )

plt.scatter(X_multi_test['petal_length'], y_multi_test, color='blue')
```

```
plt.plot(X_multi_test['petal_length'], y_multi_pred, color='red')

plt.xlabel('petal_length')

plt.ylabel('sepal_width')

plt.title('Multivariate Regression-3')

plt.show()
```

**OUTPUT :**




Multivariate Regression-2
sepal_width
petal_length
3.8
3.6
3.4
3.2
3.0
2.8
2.6
2.4
2.2
2
3
4
5
6
7


Multivariate Regression-3
sepal_width
petal_length
3.8
3.6
3.4
3.2
3.0
2.8
2.6
2.4
2.2
2
3
4
5
6
7

**Step 8: Interpret the Results**

After implementing and evaluating the models, interpret the coefficients to understand the influence of each predictor on the target variable.

```
print('Univariate Coefficients:', uni_model.coef_)
print('Bivariate Coefficients:', bi_model.coef_)
print('Multivariate Coefficients:', multi_model.coef_)
```

**OUTPUT :**

```
Univariate Coefficients: [-0.05829418]
Bivariate Coefficients: [ 0.56420418 -0.33942806]
Multivariate Coefficients: [ 0.62934965 -0.63196673  0.6440201 ]
```

**RESULT:**

This step-by-step process will help us to implement univariate, bivariate, and multivariate regression models using the Iris dataset and analyse their performance.

**EXPT NO : 2** **A python program to implement Simple linear**

**DATE: 30.08.2024** **Regression using Least Square Method**

**AIM:**

To write a python program to implement Simple linear regression using Least Square Method.

**PROCEDURE:**

Implementing Simple linear regression using Least Square method using the headbrain dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import pandas as pd

import matplotlib.pyplot as plt

import numpy as np
```

**Step 2: Load the Iris Dataset**

The HeadBrain dataset can be loaded.

```
data = pd.read_csv('/content/headbrain.csv')
```

**Step 3: Data Preprocessing**

Ensure the data is clean and ready for modeling. Since the Iris dataset is clean, minimal preprocessing is needed.

```
x,y=np.array(list(data['Head Size(cm^3)'])),np.array(list(data['Brain Weight(grams)']))

print(x[:5],y[:5])
```

**OUTPUT :**

```
[4512 3738 4261 3777 4177] [1530 1297 1335 1282 1590]
```

**Step 4 :Compute the Least Squares Solution**

Apply the least squares formula to find the regression coefficients.

```
def get_line(x,y):

 x_m,y_m = np.mean(x), np.mean(y)

 print(x_m,y_m)

 x_d,y_d=x-x_m,y-y_m

 m = np.sum(x_d*y_d)/np.sum(x_d**2)

 c = y_m - (m*x_m)

 print(m, c)

 return lambda x : m*x+c

lin=get_line(x,y)
```

**OUTPUT :**

```
3633.9915611814345 1282.873417721519
0.2634293394893993 325.5734210494428
```

**Step 5 : Make Predictions**

Use the model to make predictions based on the independent variable.

```
def get_error(line_fuc, x, y):

 y_m = np.mean(y)

 y_pred = np.array([line_fuc(_) for _ in x])

 ss_t = np.sum((y-y_m)**2)

 ss_r = np.sum((y-y_pred)**2)

 return 1-(ss_r/ss_t)
```

```
get_error(lin, x, y)
```

```
from sklearn.linear_model import LinearRegression

x = x.reshape((len(x),1))

reg=LinearRegression()

reg=reg.fit(x, y)

print(reg.score(x, y))
```

**OUTPUT :**

```
1.0
```

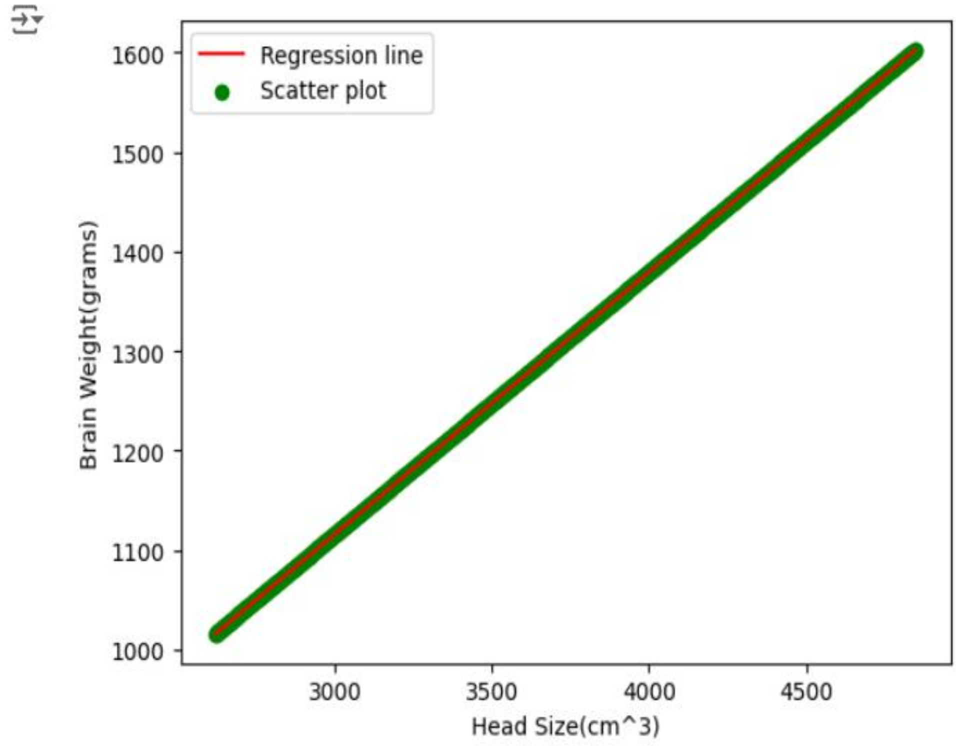
```
1.0
```

**Step 6 :Visualize the Results**

Plot the original data points and the fitted regression line.

```
x=np.linspace(np.min(x)-100,np.max(x)+100,1000)

y=np.array([lin(x)for x in x])

plt.plot(x, y, color='red', label='Regression line')

plt.scatter(x, y, color='green', label='Scatter plot')

plt.xlabel('Head Size(cm^3)')

plt.ylabel('Brain Weight(grams)')

plt.legend()

plt.show()
```

**OUTPUT :**




**RESULT:**

This step-by-step process will help us to implement least square regression models using the HeadBrain dataset and analyze their performance.

**EXPT NO : 3** **A python program to implement Logistic Model**

**DATE: 06.09.2024**

**AIM:**

To write a python program to implement a Logistic Model.

**PROCEDURE:**

Implementing Logistic method using the iris dataset involve the following steps:

### Step 1: Import Necessary Libraries

First, import the libraries that are essential for data manipulation, visualisation, and model building.

```
# Step 1: Import Necessary Libraries

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split

from sklearn.linear_model import LogisticRegression

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

### Step 2: Load the Iris Dataset

The iris dataset can be loaded.

```
# Step 2: Load the Dataset

# For this example, we'll use a built-in dataset from sklearn. You can
replace it with your dataset.

from sklearn.datasets import load_iris



# Load the iris dataset
```

```
data = load_iris()

X = data.data

y = (data.target == 0).astype(int)  # For binary classification
(classifying Iris-setosa)
```

### Step 3: Data Preprocessing

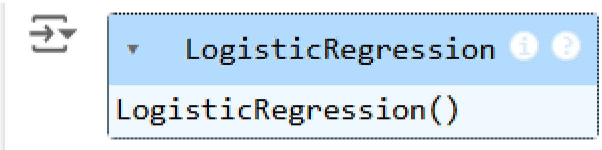
Ensure the data is clean and ready for modeling. Since the Iris dataset is clean, minimal preprocessing is needed.

```
# Step 3: Prepare the Data

# Split the dataset 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 4 : Train a Model

```
# Step 4: Create and Train the Model

model = LogisticRegression()

model.fit(X_train, y_train)
```

### OUTPUT :




### Step 5 : Make Predictions

Use the model to make predictions based on the independent variable.

```
# Step 5: Make Predictions

y_pred = model.predict(X_test)
```

### Step 6 : Evaluate the Model

Evaluate the model performance.

```
# Step 6: Evaluate the Model
```

```
accuracy = accuracy_score(y_test, y_pred)

conf_matrix = confusion_matrix(y_test, y_pred)

class_report = classification_report(y_test, y_pred)

# Print evaluation metrics

print(f"Accuracy: {accuracy}")

print("Confusion Matrix:")

print(conf_matrix)

print("Classification Report:")

print(class_report)
```

**OUTPUT :**

```
Accuracy: 1.0
Confusion Matrix:
[[20  0]
 [ 0 10]]
Classification Report:
              precision    recall  f1-score   support

           0       1.00      1.00      1.00        20
           1       1.00      1.00      1.00        10

    accuracy                           1.00        30
   macro avg       1.00      1.00      1.00        30
weighted avg       1.00      1.00      1.00        30
```

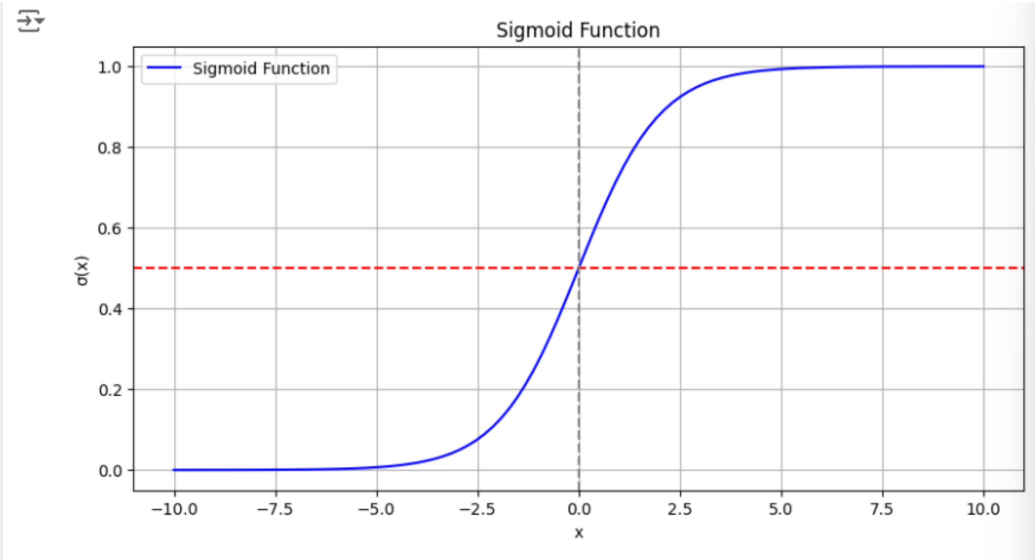
**Step 7 :Visualize the Results**

Plot the original data points and the fitted regression line.

```
# Step 7: Visualize Results (Optional)

x_values = np.linspace(-10, 10, 100)

sigmoid_values = 1 / (1 + np.exp(-x_values))


# Plot the sigmoid function

plt.figure(figsize=(10, 5))

plt.plot(x_values, sigmoid_values, label='Sigmoid Function', color='blue')
```

```
plt.title('Sigmoid Function')

plt.xlabel('x')

plt.ylabel('σ(x)')

plt.grid()

plt.axhline(0.5, color='red', linestyle='--')  # Line at y=0.5

plt.axvline(0, color='gray', linestyle='--')  # Line at x=0

plt.legend()

plt.show()
```

**OUTPUT :**



**RESULT:**

This step-by-step process will help us to implement Logistic models using the Iris dataset and analyze their performance.

**EXPT NO : 4** **A python program to implement Single Layer**

**DATE: 13.09.2024** **Perceptron**

**AIM:**

To write a python program to implement Single layer perceptron.

**PROCEDURE:**

Implementing Single layer perceptron method using the Keras dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import numpy as np

import pandas as pd

from tensorflow import keras

import matplotlib.pyplot as plt
```

**Step 2: Load the Keras Dataset**

The Keras dataset can be loaded.

```
(X_train,y_train),(X_test,y_test)=keras.datasets.mnist.load_data()
```

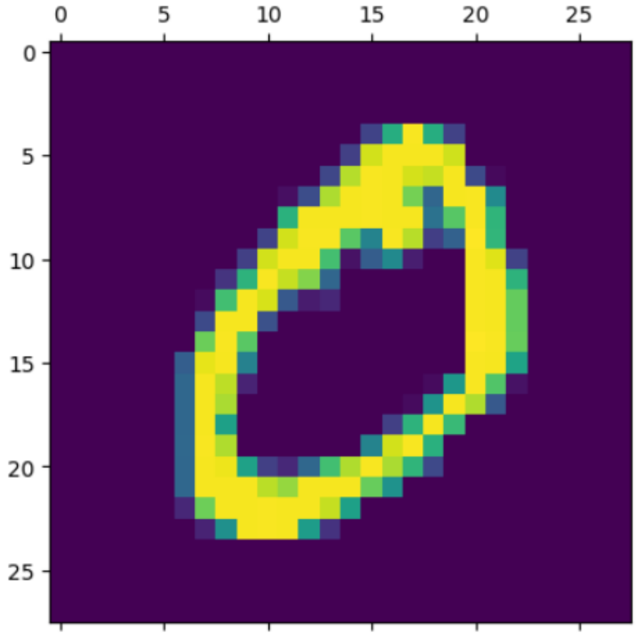
**Step 3: Data Preprocessing**

Ensure the data is clean and ready for modeling. Since the Iris dataset is clean, minimal preprocessing is needed.

```
print(f"Training set:{X_train.shape}")

print(f"Testing set:{X_test.shape}")



print(X_train[1].shape)

plt.matshow(X_train[1])
```

**OUTPUT :**

```
Training set:(60000, 28, 28)
Testing set:(10000, 28, 28)
(28, 28)
<matplotlib.image.AxesImage at 0x7b83d00435e0>
```




### Step 4 : Train a Model

```
#Normalizing the dataset

x_train=X_train/255

x_test=X_test/255


#Flatting the dataset in order to compute for model building

x_train_flatten=x_train.reshape(len(x_train),28*28)

x_test_flatten=x_test.reshape(len(x_test),28*28)

x_train_flatten.shape
```

### Step 5 : Make Predictions

Use the model to make predictions based on the independent variable.

```
model=keras.Sequential([
```

```
    keras.layers.Dense(10,input_shape=(784,),

                                activation='sigmoid')

])



model.compile(

     optimizer='adam',

     loss='sparse_categorical_crossentropy',

     metrics=['accuracy'])



model.fit(x_train_flatten,y_train,epochs=5



          )
```

**OUTPUT :**

```
Epoch 1/5
1875/1875 ━━━━━━━━━━━━━━━━━━━━ 3s 1ms/step - accuracy: 0.8180 - loss: 0.7118
Epoch 2/5
1875/1875 ━━━━━━━━━━━━━━━━━━━━ 3s 1ms/step - accuracy: 0.9148 - loss: 0.3101
Epoch 3/5
1875/1875 ━━━━━━━━━━━━━━━━━━━━ 4s 956us/step - accuracy: 0.9238 - loss: 0.2769
Epoch 4/5
1875/1875 ━━━━━━━━━━━━━━━━━━━━ 2s 940us/step - accuracy: 0.9250 - loss: 0.2744
Epoch 5/5
1875/1875 ━━━━━━━━━━━━━━━━━━━━ 3s 990us/step - accuracy: 0.9239 - loss: 0.2706
<keras.src.callbacks.history.History at 0x7b83d00c6a70>
```

### Step 6 : Evaluate the Model

Evaluate the model performance.

```
model.evaluate(x_test_flatten,y_test)
```

**OUTPUT :**

```
313/313 ━━━━━━━━━━━━━━━━━━━━ 0s 1ms/step - accuracy: 0.9138 - loss: 0.3021
[0.2668659687042236З, 0.9257000088691711]
```

**RESULT:**

This step-by-step process will help us to implement Single Layer Perceptron models using the Keras dataset and analyze their performance.

**EXPT NO : 5** **A python program to implement Multi Layer**

**DATE: 20.09.2024** **Perceptron With Backpropagation**

**AIM:**

To write a python program to implement Multilayer perceptron with backpropagation .

**PROCEDURE:**

Implementing Multilayer perceptron with backpropagation using the Keras dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
# importing modules

import tensorflow as tf

import numpy as np

from tensorflow.keras.models import Sequential

from tensorflow.keras.layers import Flatten

from tensorflow.keras.layers import Dense

from tensorflow.keras.layers import Activation

import matplotlib.pyplot as plt
```

**Step 2: Load the Keras Dataset**

The Keras dataset can be loaded.

```
 (x_train, y_train), (x_test, y_test) =
tf.keras.datasets.mnist.load_data()
```

**OUTPUT :**

```
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/mnist.npz
11490434/11490434 ━━━━━━━━━━━━━━━━━━━━ 0s 0us/step
```

### Step 3: Data Preprocessing

Ensure the data is clean and ready for modeling. Since the Iris dataset is clean, minimal preprocessing is needed.

```
# Cast the records into float values

x_train = x_train.astype('float32')

x_test = x_test.astype('float32')



# normalize image pixel values by dividing

# by 255

gray_scale = 255

x_train /= gray_scale

x_test /= gray_scale



print("Feature matrix:", x_train.shape)

print("Target matrix:", x_test.shape)

print("Feature matrix:", y_train.shape)

print("Target matrix:", y_test.shape)
```

### OUTPUT :

```
Feature matrix: (60000, 28, 28)
Target matrix: (10000, 28, 28)
Feature matrix: (60000,)
Target matrix: (10000,)
```

### Step 4 : Train a Model

```
model = Sequential([
```

```
  # reshape 28 row * 28 column data to 28*28 rows

  Flatten(input_shape=(28, 28)),



  # dense layer 1

  Dense(256, activation='sigmoid'),



  # dense layer 2

  Dense(128, activation='sigmoid'),



  # output layer

  Dense(10, activation='sigmoid'),

])
```

**OUTPUT:**

```
/usr/local/lib/python3.10/dist-packages/keras/src/layers/reshaping/flatten.py:37: UserWarning:
  super().__init__(**kwargs)
```

### Step 5 : Make Predictions

Use the model to make predictions based on the independent variable.

```
model.compile(optimizer='adam',

      loss='sparse_categorical_crossentropy',

      metrics=['accuracy'])

model.fit(x_train, y_train, epochs=10,

    batch_size=2000,

    validation_split=0.2)
```

**OUTPUT:**

```
Epoch 1/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 5s 115ms/step - accuracy: 0.3546 - loss: 2.1596 - val_accuracy: 0.68
Epoch 2/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 4s 53ms/step - accuracy: 0.7116 - loss: 1.3743 - val_accuracy: 0.82
Epoch 3/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 1s 53ms/step - accuracy: 0.8221 - loss: 0.8221 - val_accuracy: 0.87
Epoch 4/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 3s 65ms/step - accuracy: 0.8720 - loss: 0.5676 - val_accuracy: 0.89
Epoch 5/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 2s 99ms/step - accuracy: 0.8907 - loss: 0.4444 - val_accuracy: 0.90
Epoch 6/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 3s 102ms/step - accuracy: 0.8993 - loss: 0.3852 - val_accuracy: 0.91
Epoch 7/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 3s 104ms/step - accuracy: 0.9088 - loss: 0.3416 - val_accuracy: 0.91
Epoch 8/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 2s 92ms/step - accuracy: 0.9119 - loss: 0.3188 - val_accuracy: 0.92
Epoch 9/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 2s 92ms/step - accuracy: 0.9191 - loss: 0.2911 - val_accuracy: 0.92
Epoch 10/10
24/24 ━━━━━━━━━━━━━━━━━━━━ 3s 99ms/step - accuracy: 0.9245 - loss: 0.2704 - val_accuracy: 0.92
<keras.src.callbacks.history.History at 0x7d9ca1406a40>
```

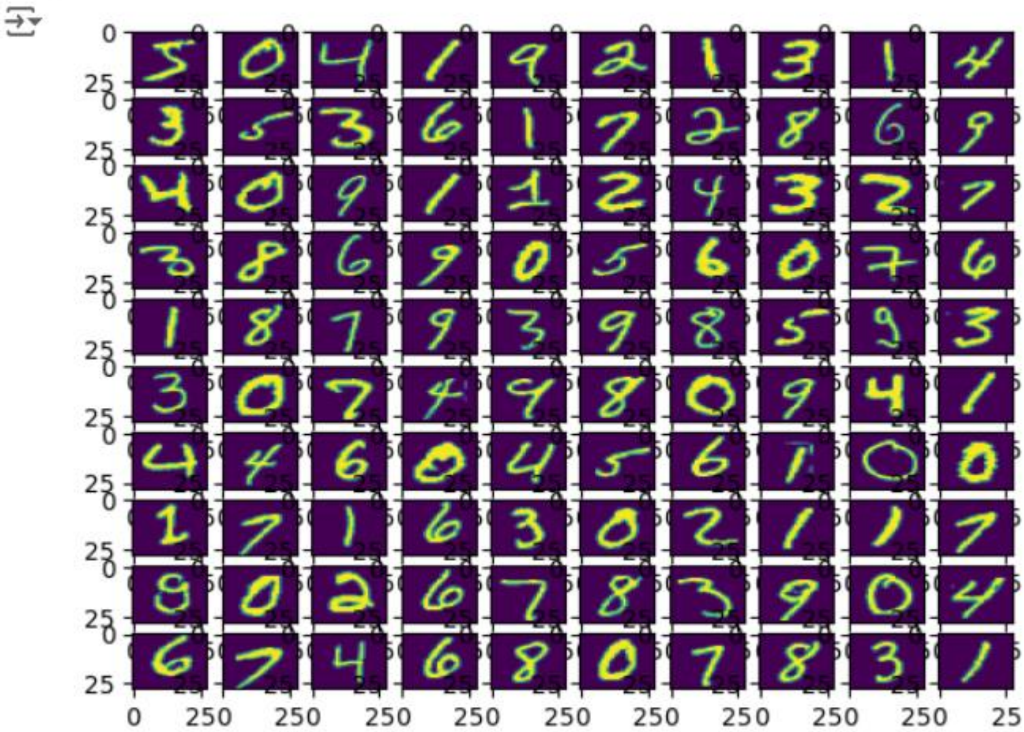
## Step 6 : Evaluate the Model

Evaluate the model performance.

```
results = model.evaluate(x_test, y_test, verbose = 0)

print('test loss, test acc:', results)

fig, ax = plt.subplots(10, 10)

k = 0

for i in range(10):

  for j in range(10):

    ax[i][j].imshow(x_train[k].reshape(28, 28),

            aspect='auto')

    k += 1

plt.show()
```

## OUTPUT :

```
test loss, test acc: [0.2589016258716583, 0.9277999997138977]
```



**RESULT:**

This step-by-step process will help us to implement MultiLayer Perceptron with Backpropagation models using the Keras dataset and analyze their performance.

**EXPT NO: 6** **A python program to do face recognition using**

**DATE: 27.09.2024** **SVM Classifier**

**AIM:**

To write a python program to implement face recognition using the SVM Classifier

**PROCEDURE:**

Implementing face recognition using the SVM Classifier using the cat and dog dataset involve the following steps:

### Step 1: Import Necessary Libraries

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import pandas as pd

import imageio

import os

from skimage.transform import resize

from skimage.io import imread

import numpy as np

import matplotlib.pyplot as plt

from sklearn import svm

from sklearn.model_selection import GridSearchCV

from sklearn.model_selection import train_test_split

from sklearn.metrics import accuracy_score

from sklearn.metrics import classification_report
```

### Step 2: Load theDog and cat  Dataset

The dog and cat dataset can be loaded.

```
Categories=['cats','dogs']

flat_data_arr=[] #input array
```

```
target_arr=[] #output array

datadir='/content/images'

#path which contains all the categories of images

for i in Categories:



  print(f'loading... category : {i}')

  path=os.path.join(datadir,i)

  for img in os.listdir(path):

    img_array=imread(os.path.join(path,img))

    img_resized=resize(img_array,(150,150,3))

    flat_data_arr.append(img_resized.flatten())

    target_arr.append(Categories.index(i))

  print(f'loaded category:{i} successfully')

flat_data=np.array(flat_data_arr)

target=np.array(target_arr)

#dataframe

df=pd.DataFrame(flat_data)

df['Target']=target

df.shape
```

**OUTPUT :**

```
(80, 67501)
```

**Step 3: Separate input features and targets.**

```
#input data
x=df.iloc[:,:-1]
#output data
y=df.iloc[:,-1]
```

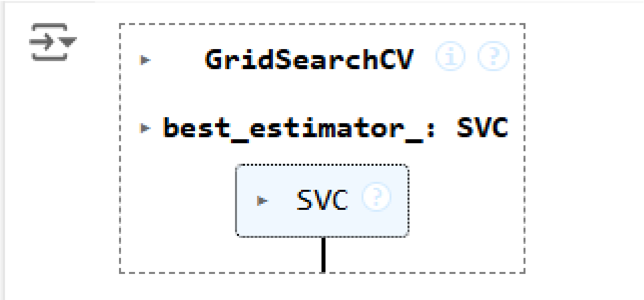
**Step 4 : Separate the input features and target**

```
# Splitting the data into training and testing sets
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.20,
random_state=77, stratify=y)
```

## Step 5 : Build and train the model

```
# Defining the parameters grid for GridSearchCV
param_grid={'C':[0.1,1,10,100],
      'gamma':[0.0001,0.001,0.1,1],
      'kernel':['rbf','poly']}

# Creating a support vector classifier
svc=svm.SVC(probability=True)

# Creating a model using GridSearchCV with the parameters grid
model=GridSearchCV(svc,param_grid)

# Training the model using the training data
model.fit(x_train,y_train)
```

**OUTPUT :**




## Step 6 : Model evaluation

```
# Testing the model using the testing data
y_pred = model.predict(x_test)

# Calculating the accuracy of the model
accuracy = accuracy_score(y_pred, y_test)

# Print the accuracy of the model
print(f"The model is {accuracy*100}% accurate")

print(classification_report(y_test, y_pred, target_names=['cat', 'dog']))
```

**OUTPUT :**

```
The model is 62.5% accurate
```

```
              precision    recall  f1-score   support

         cat       0.58      0.88      0.70         8
         dog       0.75      0.38      0.50         8

    accuracy                           0.62        16
   macro avg       0.67      0.62      0.60        16
weighted avg       0.67      0.62      0.60        16
```

## Step 7 : Prediction

```python
path='/content/cat.83.jpg'
img=imread(path)
plt.imshow(img)
plt.show()
img_resize=resize(img,(150,150,3))
l=[img_resize.flatten()]
probability=model.predict_proba(l)
for ind,val in enumerate(Categories):
  print(f'{val} = {probability[0][ind]*100}%')
print("The predicted image is : "+Categories[model.predict(l)[0]])
```

## OUTPUT :


```
cats = 52.70216647851706%
dogs = 47.29783352148294%
The predicted image is : cat
```

**RESULT :**

Thus the process helps us to implement the face recognition using SVM Classifier using python program.

## EXPT NO: 7 A python program to implement Decision tree

**DATE: 04.10.2024**

**AIM:**

To write a python program to implement a Decision tree.

**PROCEDURE:**

Implementing the decision tree using the Iris dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import numpy as np

import pandas as pd

from sklearn import datasets

from sklearn.model_selection import train_test_split

from sklearn.tree import DecisionTreeClassifier

from sklearn import metrics

import matplotlib.pyplot as plt

from sklearn.tree import plot_tree
```

**Step 2: Load the Iris Dataset**

The Iris dataset can be loaded and display the first few rows of the dataset .

```
# Load the Iris dataset

iris = datasets.load_iris()

X = iris.data  # Features

y = iris.target  # Target variable
```

## Step 3 : Split the data set into training and testing sets

```
# Split the dataset 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 4 : Create a decision tree classifier

```
# Create a Decision Tree classifier

clf = DecisionTreeClassifier(random_state=42)
```

## Step 5 : Train the model :

```
# Train the model

clf.fit(X_train, y_train)
```

## OUTPUT :

```
DecisionTreeClassifier
DecisionTreeClassifier(random_state=42)
```

## Step 6 : Make the predictions and evaluate the model

```
# Make predictions

y_pred = clf.predict(X_test)



# Evaluate the model

accuracy = metrics.accuracy_score(y_test, y_pred)

confusion = metrics.confusion_matrix(y_test, y_pred)

classification_report = metrics.classification_report(y_test, y_pred)



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

print("Confusion Matrix:")

print(confusion)
```

```
print("Classification Report:")

print(classification_report)
```

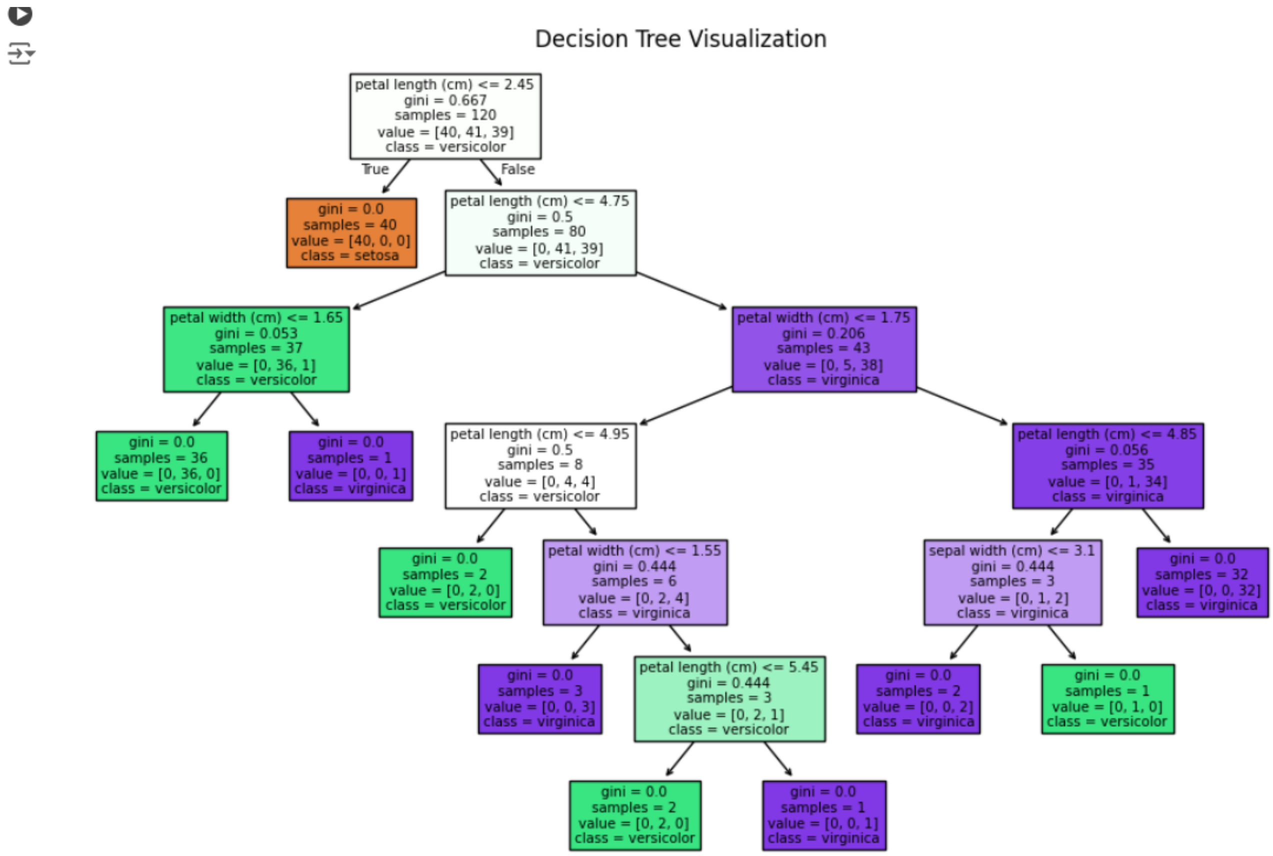
**OUTPUT :**

```
Accuracy: 1.00
Confusion Matrix:
[[10  0  0]
 [ 0  9  0]
 [ 0  0 11]]
Classification Report:
              precision    recall  f1-score   support

           0       1.00      1.00      1.00        10
           1       1.00      1.00      1.00         9
           2       1.00      1.00      1.00        11

    accuracy                           1.00        30
   macro avg       1.00      1.00      1.00        30
weighted avg       1.00      1.00      1.00        30
```

**Step 7 : Visualize the decision tree**

```
# Visualize the Decision Tree

plt.figure(figsize=(12,8))

plot_tree(clf, filled=True, feature_names=iris.feature_names,
class_names=iris.target_names)

plt.title("Decision Tree Visualization")

plt.show()
```

**OUTPUT :**




**RESULT :**

**This process helps us to implement the decision tree using a python program.**

**EX.NO: 8** **A PYTHON PROGRAM TO IMPLEMENT**

**DATE : 18.10.2024** **ADA BOOSTING**

**AIM:**

To write a python program to implement ADA Boosting.

**PROCEDURE:**

Implementing ADA Boosting using the dataset involve the following steps:

**Step 1: Import Necessary Libraries**

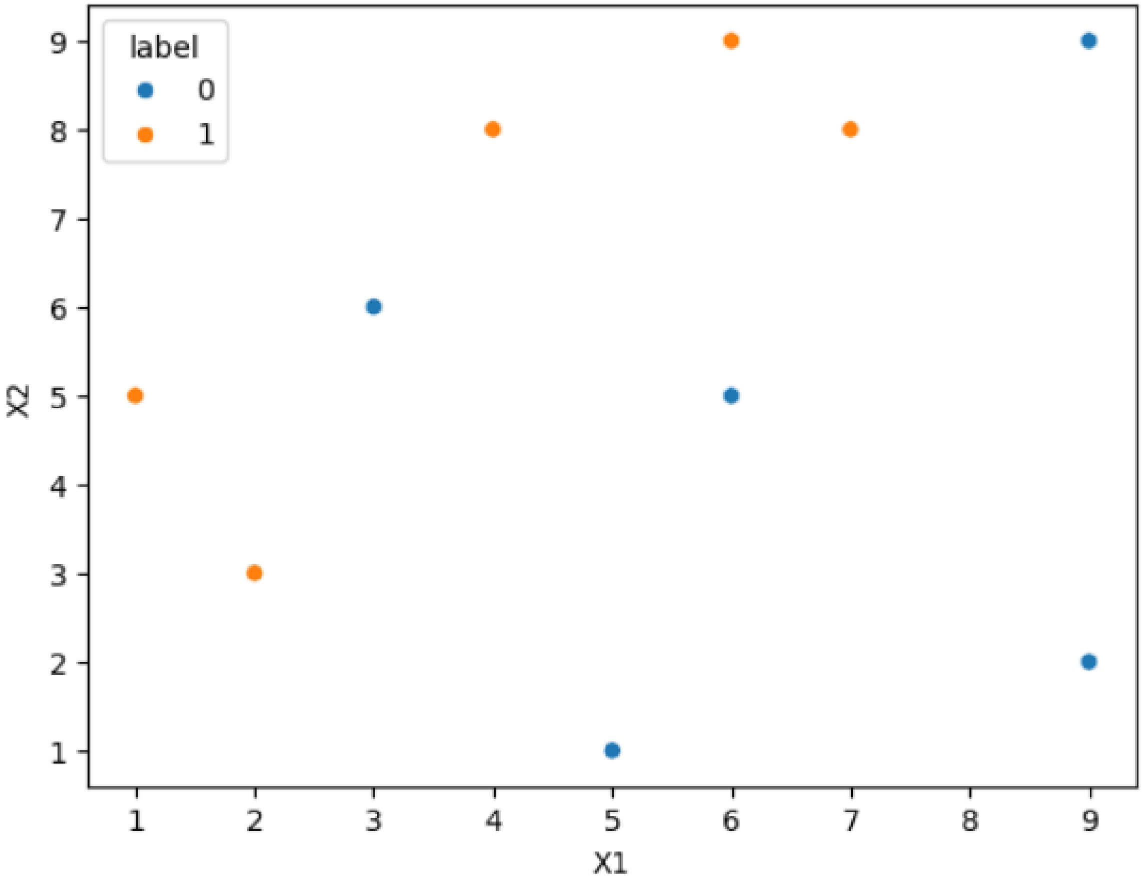
First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import numpy as np

import pandas as pd

from sklearn.tree import DecisionTreeClassifier

from mlxtend.plotting import plot_decision_regions

import seaborn as sns

from sklearn.metrics import accuracy_score
```

**Step 2 : Load and prepare data**

```
df = pd.DataFrame()

df['X1'] = [1, 2, 3, 4, 5, 6, 6, 7, 9, 9]

df['X2'] = [5, 3, 6, 8, 1, 9, 5, 8, 9, 2]

df['label'] = [1, 1, 0, 1, 0, 1, 0, 1, 0, 0]

sns.scatterplot(x=df['X1'], y=df['X2'], hue=df['label'])



df['weights'] = 1 / df.shape[0]

x = df.iloc[:, 0:2].values

y = df.iloc[:, 2].values
```

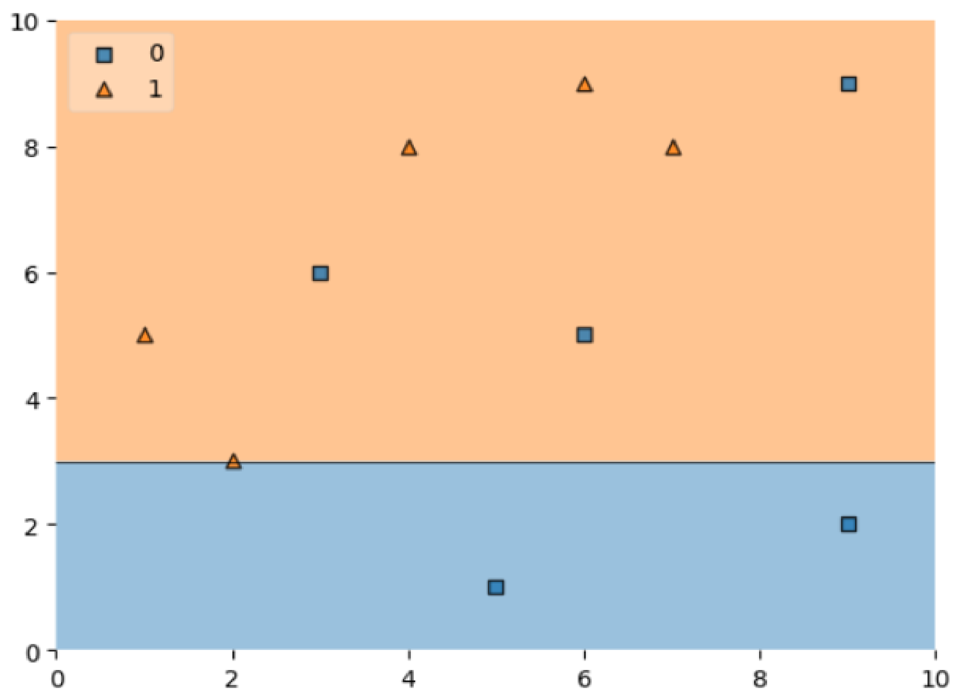
**OUTPUT :**




**Step 3 : Train the 1st model**

```
# Step 2: Train 1st Model

dt1 = DecisionTreeClassifier(max_depth=1)

dt1.fit(x, y)

plot_decision_regions(x, y, clf=dt1, legend=2)

df['y_pred'] = dt1.predict(x)
```

**OUTPUT :**




## Step 4 : Calculate model weight

```
# Step 4: Update Weights

def update_row_weights(row, alpha=0.423):

    if row['label'] == row['y_pred']:

        return row['weights'] * np.exp(-alpha)

    else:

        return row['weights'] * np.exp(alpha)



df['updated_weights'] = df.apply(update_row_weights, axis=1)

df['normalized_weights'] = df['updated_weights'] /
df['updated_weights'].sum()

df['cumsum_upper'] = np.cumsum(df['normalized_weights'])

df['cumsum_lower'] = df['cumsum_upper'] - df['normalized_weights']
```

## Step 5 : Create new dataset

```
# Step 5: Create New Dataset

def create_new_dataset(df):
```

```
    indices = []

    for i in range(df.shape[0]):

        a = np.random.random()

        for index, row in df.iterrows():

            if row['cumsum_upper'] > a and a > row['cumsum_lower']:

                indices.append(index)

    return indices



index_values = create_new_dataset(df)

second_df = df.iloc[index_values, [0, 1, 2, 3]]
```

## Step 6 : Train 2nd model

```
# Step 6: Train 2nd Model

dt2 = DecisionTreeClassifier(max_depth=1)

x = second_df.iloc[:, 0:2].values

y = second_df.iloc[:, 2].values

dt2.fit(x, y)
```

## OUTPUT :

```
DecisionTreeClassifier
DecisionTreeClassifier(max_depth=1)
```

## Step 7 : Plot decision tree and calculate model weights for 2nd model

```
# Plot the decision tree for the second model

plot_decision_regions(x, y, clf=dt2, legend=2)

second_df['y_pred'] = dt2.predict(x)
```

```
# Step 7: Calculate Model Weight for 2nd Model

alpha2 = calculate_model_weight(0.1)

print(f"Alpha2: {alpha2}")
```

## Step 8 : update weights for 2nd model

```
# Step 8: Update Weights for 2nd Model

def update_row_weights(row, alpha=1.09):

    if row['label'] == row['y_pred']:

        return row['weights'] * np.exp(-alpha)

    else:

        return row['weights'] * np.exp(alpha)



second_df['updated_weights'] = second_df.apply(update_row_weights, axis=1)

second_df['nomalized_weights'] = second_df['updated_weights'] /
second_df['updated_weights'].sum()

second_df['cumsum_upper'] = np.cumsum(second_df['nomalized_weights'])

second_df['cumsum_lower'] = second_df['cumsum_upper'] -
second_df['nomalized_weights']
```

## Step 9 : Calculate alpha for 3rd model

```
# Step 9: Calculate Alpha for 3rd Model

alpha3 = calculate_model_weight(0.7)

print(f"Alpha3: {alpha3}")



# Step 10: Accuracy Calculation

y_true = second_df['label'].values
```

```
y_pred = second_df['y_pred'].values


# Calculate accuracy for the AdaBoost model

accuracy = accuracy_score(y_true, y_pred)

print(f"Accuracy of the AdaBoost model: {accuracy:.4f}")
```

**OUTPUT :**

**ALPHA 3: -0.4236489301936017**

**Accuracy of the Ada Boosting model : 0.80000**

**RESULT :**

Thus the python program to implement Adaboosting has been executed successfully and the results have been verified.

**EXPT NO: 9A** **A python program to implement**

**DATE: 25.10.2024** **KNN MODEL .**

**AIM:**

To write a python program to implement KNN Model.

**PROCEDURE:**

Implementing KNN Model using the mall_customer dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import numpy as np

import matplotlib.pyplot as plt

import pandas as pd

from sklearn.model_selection import train_test_split

from sklearn.preprocessing import StandardScaler

from sklearn.neighbors import KNeighborsClassifier

from sklearn.metrics import classification_report, confusion_matrix

from sklearn.cluster import KMeans
```

**Step 2: Load the Dataset**

The mall_customer dataset can be loaded and display the first few rows of the dataset.

```
# Load the dataset

dataset = pd.read_csv('/content/Mall_Customers.csv')



# Display the first few rows of the dataset

print(dataset.head())



# Display the dimensions of the dataset

print(f"Dataset shape: {dataset.shape}")
```

```
# Display descriptive statistics of the dataset

print(dataset.describe())
```

## Step 3 : Separate the features (x) and target variable (y)

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

X = dataset.iloc[:, [3, 4]].values  # We use 'Annual Income' and 'Spending
Score'



# Standardize the features

scaler = StandardScaler()

X_scaled = scaler.fit_transform(X)
```

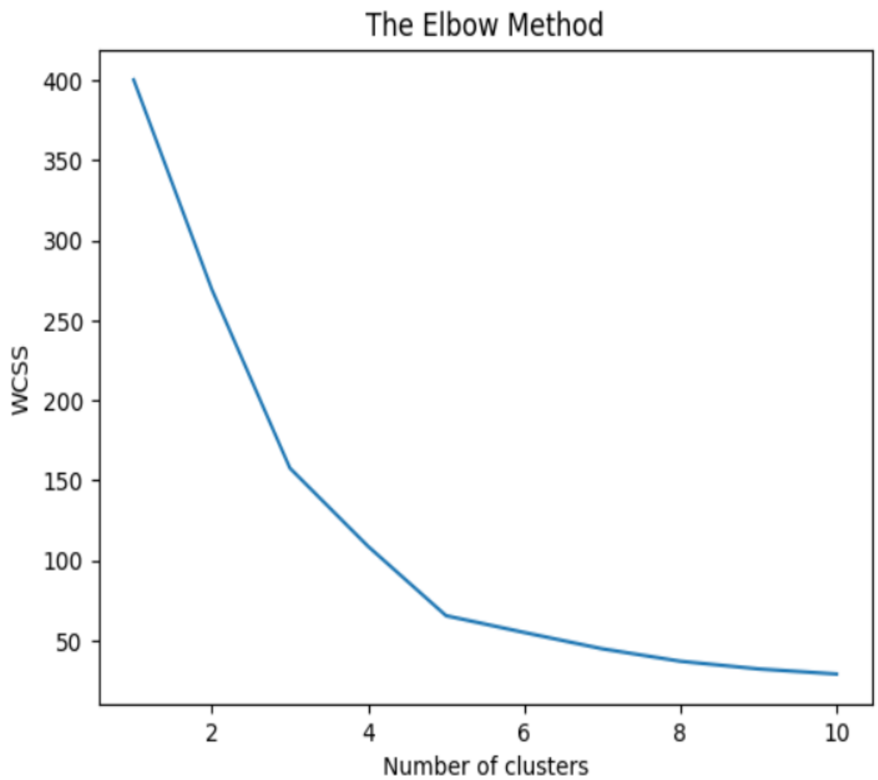
## Step 4 : Visualizing the cluster of customer

```
# Apply KMeans clustering using the Elbow Method to find the optimal
number of clusters

wcss = []  # Within-cluster sum of squares

for i in range(1, 11):

    kmeans = KMeans(n_clusters=i, init='k-means++', max_iter=300,
n_init=10, random_state=0)

    kmeans.fit(X_scaled)

    wcss.append(kmeans.inertia_)



# Plot the Elbow Method graph

plt.plot(range(1, 11), wcss)

plt.title('The Elbow Method')

plt.xlabel('Number of clusters')

plt.ylabel('WCSS')
```

```
plt.show()


# From the plot, we can observe that the optimal number of clusters is 5 (elbow point)
kmeans = KMeans(n_clusters=5, init='k-means++', max_iter=300, n_init=10, random_state=0)
y_kmeans = kmeans.fit_predict(X_scaled)


# Visualizing the clusters of customers
plt.scatter(X_scaled[y_kmeans == 0, 0], X_scaled[y_kmeans == 0, 1], s=100, c='red', label='Cluster 1')
plt.scatter(X_scaled[y_kmeans == 1, 0], X_scaled[y_kmeans == 1, 1], s=100, c='blue', label='Cluster 2')
plt.scatter(X_scaled[y_kmeans == 2, 0], X_scaled[y_kmeans == 2, 1], s=100, c='green', label='Cluster 3')
plt.scatter(X_scaled[y_kmeans == 3, 0], X_scaled[y_kmeans == 3, 1], s=100, c='cyan', label='Cluster 4')
plt.scatter(X_scaled[y_kmeans == 4, 0], X_scaled[y_kmeans == 4, 1], s=100, c='magenta', label='Cluster 5')


# Plot the centroids
plt.scatter(kmeans.cluster_centers_[:, 0], kmeans.cluster_centers_[:, 1], s=300, c='yellow', label='Centroids')


plt.title('Clusters of customers')
plt.xlabel('Annual Income (k$)')
plt.ylabel('Spending Score (1-100)')
plt.legend()
plt.show()
```

**OUTPUT :**


Clusters of customers
Spending Score (1-100)
Annual Income (k$)
Cluster 1
Cluster 2
Cluster 3
Cluster 4
Cluster 5
Centroids
0
20
40
60
80
100
20
40
60
80
100
120
140


The Elbow Method
WCSS
Number of clusters
400
350
300
250
200
150
100
50
2
4
6
8
10

**RESULT :**

Thus the python program to implement KNN model has been successfully implemented and the results have been verified.

**EXPT NO: 9B** **A python program to implement**

**DATE: 25.10.2024** **K-Means Model**

**AIM:**

To write a python program to implement the K-means Model.

**PROCEDURE:**

Implementing K - means Model using the mall_customer dataset involve the following steps:

**Step 1: Import Necessary Libraries**

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
import numpy as np

import pandas as pd

from math import sqrt
```

**Step 2 : load the Dataset**

```
data = pd.read_csv('/content/Mall_Customers.csv')

data.head(5)
```

**OUTPUT:**

| | CustomerID | Gender | Age | Annual Income (k$) | Spending Score (1-100) |
|---|---|---|---|---|---|
| **0** | 1 | Male | 19 | 15 | 39 |
| **1** | 2 | Male | 21 | 15 | 81 |
| **2** | 3 | Female | 20 | 16 | 6 |
| **3** | 4 | Female | 23 | 16 | 77 |
| **4** | 5 | Female | 31 | 17 | 40 |

**Step 3 : Preprocess the data**

```
req_data = data[['Age', 'Annual Income (k$)', 'Spending Score (1-100)']]



req_data.head(5)
```

## OUTPUT :

| | Age | Annual Income (k$) | Spending Score (1-100) |
|---|---|---|---|
| 0 | 19 | 15 | 39 |
| 1 | 21 | 15 | 81 |
| 2 | 20 | 16 | 6 |
| 3 | 23 | 16 | 77 |
| 4 | 31 | 17 | 40 |

## Step 4 : Assign the data points to clusters

```
shuffle_index = np.random.permutation(req_data.shape[0])  # Shuffle the dataset rows

req_data = req_data.iloc[shuffle_index]

req_data.head(5)
```

## OUTPUT :

| | Gender | Age | Annual Income (k$) | Spending Score (1-100) |
|---|---|---|---|---|
| 14 | Male | 37 | 20 | 13 |
| 102 | Male | 67 | 62 | 59 |
| 89 | Female | 50 | 58 | 46 |
| 181 | Female | 32 | 97 | 86 |
| 183 | Female | 29 | 98 | 88 |

## Step 5 : Update the clusters centers

```
train_size = int(req_data.shape[0]*0.7)  # Set 70% of the data for training

train_df = req_data.iloc[:train_size,:]

test_df = req_data.iloc[train_size:,:]

train = train_df.values  # Convert train data to numpy array
```

```
test = test_df.values  # Convert test data to numpy array
y_true = test[:,-1]  # The target values for the test set
print('Train_Shape: ', train_df.shape)
print('Test_Shape: ', test_df.shape)
from math import sqrt


def euclidean_distance(x_test, x_train):
    distance = 0
    for i in range(len(x_test)):  # Loop through all features
        distance += (x_test[i]-x_train[i])**2
    return sqrt(distance)


def get_neighbors(x_test, x_train, num_neighbors):
    distances = []
    data = []
    for i in x_train:
        distances.append(euclidean_distance(x_test, i))
        data.append(i)
    distances = np.array(distances)
    data = np.array(data)
    sort_indexes = distances.argsort()  # Sort distances in ascending order
    data = data[sort_indexes]  # Sort the data based on sorted distances
    return data[:num_neighbors]  # Return the closest 'num_neighbors' neighbors
def prediction(x_test, x_train, num_neighbors):
    classes = []
```

```python
    neighbors = get_neighbors(x_test, x_train, num_neighbors)
    for i in neighbors:
        classes.append(i[-1])  # The target value is the last column
    predicted = max(classes, key=classes.count)  # Return the most frequent class (the majority vote)
    return predicted


def predict_classifier(x_test):
    classes = []
    neighbors = get_neighbors(x_test, req_data.values, 5)  # Predict using the top 5 neighbors
    for i in neighbors:
        classes.append(i[-1])
    predicted = max(classes, key=classes.count)  # Return the majority vote
    print(predicted)
    return predicted
def accuracy(y_true, y_pred):
    num_correct = 0
    for i in range(len(y_true)):
        if y_true[i] == y_pred[i]:  # Compare true values to predicted values
            num_correct += 1
    accuracy = num_correct / len(y_true)  # Calculate accuracy as the ratio of correct predictions
    return accuracy
def accuracy(y_true, y_pred):
    num_correct = 0
```

```
    for i in range(len(y_true)):
        if y_true[i] == y_pred[i]:
            num_correct += 1
    return num_correct / len(y_true)
y_pred = []
for i in test:
    y_pred.append(prediction(i, train, 5))  # Make predictions for each test instance


# Calculate and print the accuracy
acc = accuracy(y_true, y_pred)
print(f"Accuracy: {acc * 1000:.2f}%")
```

**OUTPUT :**

```
Accuracy: 66.67%
```

**RESULT :**

Thus the python program implementing the k-means model is successful.

**EXPT NO: 10** **A python program to implement Dimensionality**

**DATE: 04.11.2024** **Reduction -PCA.**

**AIM:**

To write a python program to implement Dimensionality Reduction - PCA .

**PROCEDURE:**

ImplementingDimensionality reduction -pca  using the Iris dataset involve the following steps:

### Step 1: Import Necessary Libraries

First, import the libraries that are essential for data manipulation, visualization, and model building.

```
# Importing necessary libraries

from sklearn import datasets

import pandas as pd

from sklearn.preprocessing import StandardScaler

from sklearn.decomposition import PCA

import seaborn as sns

import matplotlib.pyplot as plt
```

### Step 2: Load the Iris Dataset

The Iris dataset can be loaded and display the first few rows of the dataset

```
# Load the Iris dataset

iris = datasets.load_iris()

df = pd.DataFrame(iris['data'], columns=iris['feature_names'])



# Display the first few rows of the dataset

df.head()
```

**OUTPUT :**

| | sepal length (cm) | sepal width (cm) | petal length (cm) | petal width (cm) |
|---|---|---|---|---|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 |

## Step 3 : Standardize the data

```
# Standardize the features using StandardScaler
scalar = StandardScaler()
scaled_data = pd.DataFrame(scalar.fit_transform(df))  # Scaling the data


# Display the scaled data (optional)
scaled_data.head()
```

## OUTPUT :

| | 0 | 1 | 2 | 3 |
|---|---|---|---|---|
| 0 | -0.900681 | 1.019004 | -1.340227 | -1.315444 |
| 1 | -1.143017 | -0.131979 | -1.340227 | -1.315444 |
| 2 | -1.385353 | 0.328414 | -1.397064 | -1.315444 |
| 3 | -1.506521 | 0.098217 | -1.283389 | -1.315444 |
| 4 | -1.021849 | 1.249201 | -1.340227 | -1.315444 |

## Step 4 : Apply PCA

```
# Apply PCA to reduce the data to 3 components
pca = PCA(n_components=3)
pca.fit(scaled_data)  # Fit PCA on scaled data
data_pca = pca.transform(scaled_data)  # Transform the data to principal
```

```
components


# Convert PCA data to a DataFrame for easier inspection
data_pca = pd.DataFrame(data_pca, columns=['PC1', 'PC2', 'PC3'])
data_pca.head()
```

**OUTPUT :**

| | PC1 | PC2 | PC3 |
|---|---|---|---|
| 0 | -2.264703 | 0.480027 | 0.127706 |
| 1 | -2.080961 | -0.674134 | 0.234609 |
| 2 | -2.364229 | -0.341908 | -0.044201 |
| 3 | -2.299384 | -0.597395 | -0.091290 |
| 4 | -2.389842 | 0.646835 | -0.015738 |

**Step 5 : Explained Variance Ratio**

```
# Calculate the explained variance ratio for each principal component
explained_variance = pca.explained_variance_ratio_
print(f"Explained Variance Ratio: {explained_variance}")


# This output shows how much variance each principal component explains.
```

**OUTPUT :**

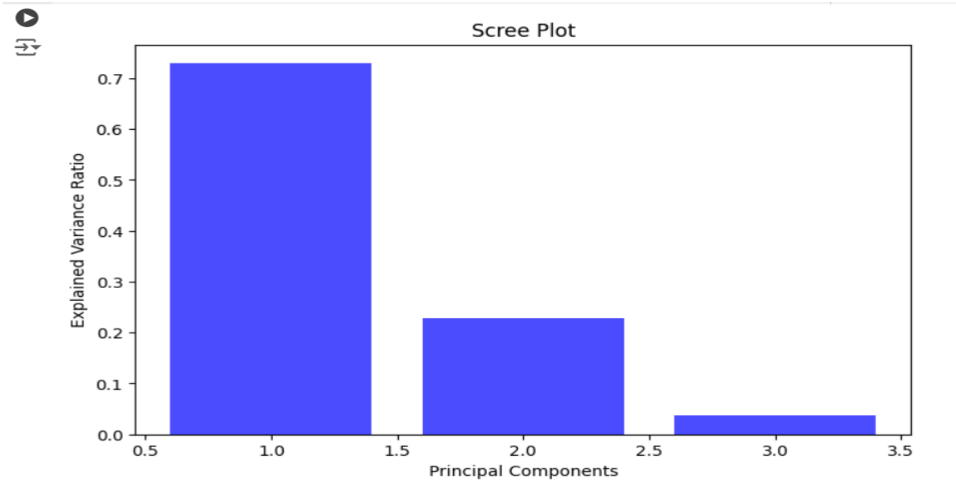
```
Explained Variance Ratio: [0.72962445 0.22850762 0.03668922]
```

**Step 6 :Visualize the reduced data.**

```
# Plotting the explained variance ratio as a scree plot
plt.figure(figsize=(8, 5))
```

```
plt.bar(range(1, len(explained_variance) + 1), explained_variance,
alpha=0.7, color='blue')

plt.ylabel('Explained Variance Ratio')

plt.xlabel('Principal Components')

plt.title('Scree Plot')

plt.show()
```

## OUTPUT :




## RESULT :

Thus the Dimensionality Reduction has been implemented using PCA in python program Successfully.