A PYTHON PROGRAM TO IMPLEMENT K-MEANS MODEL

Ex.No.: 9B

Date of Experiment: 25/10/2024

AIM:-

To implement a python program using a K-Means Algorithm in a model.

ALGORITHM:-

Step1: Import all the other necessary libraries(numpy as np, matplotlib.pyplot as plt and

sklearn.tree,pandas as pd and seaborn as sns).

Step2: Select the number K to decide the number of clusters.

Step3: Select random K points or centroids. (It can be different from the input dataset). Step4:

Assign each data point to their closest centroid, which will form the predefined K clusters. Step5:

Calculate the variance and place a new centroid of each cluster.

Step6: Repeat the fourth steps, which means assign each datapoint to the new closest centroid of

each cluster.

Step7: If any reassignment occurs, then go to step-5 else go to FINISH.

Step8: Train the model and plot the graph using scatterplot() function.

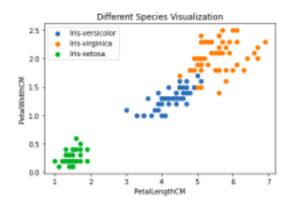
Page No.:

IMPLEMENTATION:-

231501106 [AIML -B]

data = pd.read_csv('../input/k-means-clustering/KNN (3).csv')
data.head(5)

Text(0.5, 1.0, 'Different Species Visualization')



req_data = data.iloc[:,1:]
req_data.head(5)

| | SepalLengthCm | SepalWidthCm | PetalLengthCm | PetalWidthCm | Species |
|---|---------------|--------------|---------------|--------------|-------------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 | Iris-setosa |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 | Iris-setosa |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 | Iris-setosa |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 | Iris-setosa |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 | Iris-setosa |

shuffle_index = np.random.permutation(req_data.shape[0]) #shuffling the row index of our dataset

req_data = req_data.iloc[shuffle_index]
req_data.head(5)

| | SepalLengthCm | SepalWidthCm | PetalLengthCm | PetalWidthCm | Species |
|-----|---------------|--------------|---------------|--------------|-----------------|
| 45 | 4.8 | 3.0 | 1.4 | 0.3 | Iris-setosa |
| 50 | 7.0 | 3.2 | 4.7 | 1.4 | Iris-versicolor |
| 135 | 7.7 | 3.0 | 6.1 | 2.3 | Iris-virginica |
| 49 | 5.0 | 3.3 | 1.4 | 0.2 | Iris-setosa |
| 89 | 5.5 | 2.5 | 4.0 | 1.3 | Iris-versicolor |

```
train size = int(req data.shape[0]*0.7)
train_df = req_data.iloc[:train_size,:]
test_df = req_data.iloc[train_size:,:]
train = train_df.values
test = test df.values
y_{true} = test[:,-1]
print('Train_Shape: ',train_df.shape)
print('Test_Shape: ',test_df.shape)
  Train_Shape: (105, 5)
  Test_Shape: (45, 5)
from math import sqrt
def euclidean_distance(x_test, x_train):
  distance = 0
  for i in range(len(x_test)-1):
     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() #argsort() function returns indices by sorting distances
data in ascending order
  data = data[sort indexes] #modifying our data based on sorted indices, so that we can get the
nearest neighbors
  return data[:num_neighbors]
def prediction(x test, x train, num neighbors):
  classes = []
  neighbors = get neighbors(x test, x train, num neighbors)
  for i in neighbors:
     classes.append(i[-1])
  predicted = max(classes, key=classes.count) #taking the most repeated class return
  predicted
def predict classifier(x test):
  classes = []
  neighbors = get neighbors(x test, req data.values, 5)
  for i in neighbors:
     classes.append(i[-1])
  predicted = max(classes, key=classes.count)
  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]:
        num_correct+=1
    accuracy = num_correct/len(y_true)
    return accuracy

y_pred = []
for i in test:
    y_pred.append(prediction(i, train, 5))
y_pred
```

```
['Iris-virginica',
'Iris-versicolor',
'Iris-versicolor',
'Iris-setosa',
'Iris-virginica',
'Iris-setosa',
'Iris-setosa',
'Iris-setosa',
'Iris-virginica',
'Iris-versicolor',
'Iris-setosa',
 'Iris-versicolor',
'Iris-versicolor',
'Iris-virginica',
'Iris-setosa',
'Iris-setosa',
'Iris-versicolor',
'Iris-virginica',
'Iris-virginica',
'Iris-setosa',
'Iris-virginica',
'Iris-versicolor',
'Iris-setosa',
'Iris-setosa',
'Iris-versicolor',
'Iris-setosa',
'Iris-setosa',
'Iris-versicolor',
'Iris-virginica',
'Iris-versicolor',
'Iris-virginica',
'Iris-versicolor',
'Iris-versicolor',
'Iris-virginica',
'Iris-virginica',
'Iris-versicolor',
'Iris-virginica',
'Iris-setosa',
'Iris-setosa',
'Iris-virginica',
'Iris-virginica',
'Iris-setosa',
'Iris-versicolor',
'Iris-virginica',
 'Iris-versicolor']
```

accuracy = accuracy(y_true, y_pred)

accuracy

0.95555555555556

test_df.sample(5)

| | SepalLengthCm | SepalWidthCm | PetalLengthCm | PetalWidthCm | Species |
|-----|---------------|--------------|---------------|--------------|-----------------|
| 113 | 5.7 | 2.5 | 5.0 | 2.0 | Iris-virginica |
| 125 | 7.2 | 3.2 | 6.0 | 1.8 | Iris-virginica |
| 149 | 5.9 | 3.0 | 5.1 | 1.8 | Iris-virginica |
| 94 | 5.6 | 2.7 | 4.2 | 1.3 | Iris-versicolor |
| 99 | 5.7 | 2.8 | 4.1 | 1.3 | Iris-versicolor |

RESULT:-

Thus the python program to implement the K-Means model has been successfully implemented and the results have been verified and analyzed.