1.Implement and demonstrate the FIND-S algorithm to finding the most specific hypothesis based ona given set of data samples. Read the training data from a .CSV file.

```
import pandas as pd
d=pd.read_csv("/content/ENJOYSPORT.csv")
d=pd.DataFrame(d)
print(d)
h=[]
t=0
for i in range(len(d)):
l=list(d.loc[i])
if I[len(I)-1]==1:
 I.pop()
  h.extend(I)
  k=i
  break
print("H",t+1,":",h)
t=t+2
I=[]
for i in range(k+1,len(d)):
 l=list(d.loc[i])
  if I[len(I)-1]==1:
  for j in range(len(h)):
    if h[j]!=l[j]:
    h[j]='?'
  print("H",t,":",h)
  t=t+1
OUTPUT:-
H 1 : ['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same']
H 2 : ['Sunny', 'Warm', '?', 'Strong', 'Warm', 'Same']
H 3 : ['Sunny', 'Warm', '?', 'Strong', 'Warm', 'Same']
H 4 : ['Sunny', 'Warm', '?', 'Strong', '?', '?']
```

2. For a given set of training data examples stored in a .CSV file, implement and demonstrate the Candidate-Elimination algorithm to output a description of the set of all hypotheses consistent with the training examples.

```
from re import S
import pandas as pd
d=pd.read_csv("/content/ENJOYSPORT.csv")
# print(len(d.columns))
d=pd.DataFrame(d)
print("DATA SET: ")
print(d)
# print(len(d.columns))
s=[]
t=['?' for i in range(len(d.columns)-1)]
g=[]
l=list(d.loc[0])
I.pop()
s.extend(I)
for i in range(len(d)):
  l=list(d.loc[i])
  if I[len(I)-1]==1:
   for j in range(len(s)):
    if s[j]!=l[j]:
     s[j]='?'
   if g!=[]:
    for p in range(len(g)):
     for q in range(len(s)):
       if g[p][q] == s[q] or g[p][q] == '?':
          continue
```

```
else:
          g.pop(p)
  else:
   for j in range(len(s)):
    if I[j]!=s[j] and s[j]!='?':
     t[j]=s[j]
     g.append(t)
     t=['?' for i in range(len(d.columns)-1)]
# print(g)
# print('\n')
# print(s)
def versionspace(g,s):
 vs=[]
 for i in g:
  for j in range(len(i)):
   if i[j]=='?' and s[j]!='?':
    m=i[:]
    m[j]=s[j]
    if m not in vs:
     vs.append(m)
 return vs
r=[]
r.append(s)
r1=versionspace(g,s)
r.extend(r1)
r.extend(g)
print("\nGeneral Hypothesis : ",g)
```

```
print("\nSpecific Hypothesis : ",s)
print("\nVersion Space :")
for i in r:
 print(i)
OUTPUT:-
DATA SET:
  Sky AirTemp Humidity Wind Water Forecast EnjoySport
O Sunny Warm Normal Strong Warm Same
1 Sunny Warm High Strong Warm Same
                                                      1
2 Rainy Cold High Strong Warm Change
                                                     0
3 Sunny Warm High Strong Cool Change
General Hypothesis: [['Sunny', '?', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?', '?']]
Specific Hypothesis: ['Sunny', 'Warm', '?', 'Strong', '?', '?']
Version Space:
['Sunny', 'Warm', '?', 'Strong', '?', '?']
['Sunny', 'Warm', '?', '?', '?', '?']
['Sunny', '?', '?', 'Strong', '?', '?']
['?', 'Warm', '?', 'Strong', '?', '?']
['Sunny', '?', '?', '?', '?', '?']
['?', 'Warm', '?', '?', '?', '?']
3. Write a program to implement the naïve Bayesian classifier for a sample training data set stored as
a .CSV file. Compute the accuracy of the classifier, considering few test data sets.
import pandas as pd
df=pd.read_csv('/content/play_tennis.csv')
df=df.iloc[:,1:]
```

```
print("DATA FRAME:")
print(df)
train=df[:10]
yn=train.iloc[:,-1].value_counts()
y=yn['Yes']
n=yn['No']
t=len(train)
py=y/t
pn=n/t
c=list(train.iloc[:,:].columns)
dict={}
for i in c:
 dict[i]={}
 for j in train[i]:
   if j not in dict[i]:
    dict[i][j]=[]
for k in dict:
   for I in dict[k]:
    ky=len(train[(train[c[-1]]=='Yes') & (train[k]==I)][k])
    kn=len(train[(train[c[-1]]=='No') & (train[k]==I)][k])
    dict[k][l].extend([(ky/y),(kn/n)])
# print(dict)
test=df[10:]
test=test.iloc[:,:]
test1=test
print(test1)
```

```
test=test.iloc[:,:-1]
o=[]
for i in range(len(test)):
 yes=py
 no=pn
 l=list(test.iloc[i,:])
 print("Test row : ",I)
 for j in range(len(l)):
  yes=yes*dict[c[j]][l[j]][0]
  no=no*dict[c[j]][l[j]][1]
 print("Yes:",yes,"No:",no)
 if yes>no:
  print("Yes")
  o.append('Yes')
 elif no>yes:
  print("No")
  o.append('No')
test2=pd.DataFrame(test)
test2[c[-1]]=o
print(test2)
import sklearn
from sklearn.metrics import accuracy_score
actual=list(test1[c[-1]])
predicted=o
accuracy=sklearn.metrics.accuracy_score(actual,predicted)
print(accuracy)
OUTPUT:-
```

```
DATA FRAME :
                          wind play
    outlook temp humidity
\cap
      Sunny Hot High
                            Weak No
     Sunny Hot High Strong No ercast Hot High Weak Yes Rain Mild High Weak Yes
1
2
   Overcast
3
      Rain Cool Normal
4
                           Weak Yes
      Rain Cool Normal Strong
5
                                  Nο
  Overcast Cool Normal Strong Yes
6
    Sunny Mild High Weak No
Sunny Cool Normal Weak Yes
7
8
      Rain Mild Normal
                           Weak Yes
9
     Sunny Mild Normal Strong Yes
10
11 Overcast Mild High Strong Yes
                           Weak Yes
12 Overcast Hot Normal
    Rain Mild High Strong
1.3
   outlook temp humidity wind play
10
    Sunny Mild Normal Strong Yes
11 Overcast Mild High Strong Yes
12 Overcast Hot Normal Weak Yes
1.3
      Rain Mild
                    High Strong
                                  No
Test row: ['Sunny', 'Mild', 'Normal', 'Strong']
Yes: 0.0037037037037037025 No: 0.00937500000000001
Test row : ['Overcast', 'Mild', 'High', 'Strong']
Yes: 0.0037037037037025 No: 0.0
Test row : ['Overcast', 'Hot', 'Normal', 'Weak']
Yes: 0.018518518518518514 No:
Test row : ['Rain', 'Mild', 'High', 'Strong']
Yes: 0.005555555555555554 No: 0.00937500000000001
    outlook temp humidity
                           wind play
10
     Sunny Mild Normal Strong No
11 Overcast Mild High Strong Yes
12 Overcast Hot Normal Weak Yes
     Rain Mild High Strong No
1.3
```

4. Assuming a set of documents that need to be classified, use the naïve Bayesian classifier model to perform this task. Built-in Java classes /API can be used to write the program. Calculate the accuracy precision and recall for your data set.

```
import pandas as pd
d=pd.read_csv("/content/text_classification.csv")
# d=d.iloc[:,:-1]
print("DATA FRAME : ")
print(d)
```

0.75

```
train=d[:10]
yn=train.iloc[:,-1].value_counts()
y=yn['pos']
n=yn['neg']
t=len(train)
py=y/t
pn=n/t
# print(train)
dict={}
n1,n2=0,0
for i in range(len(train)):
 dict[i]={}
 l=list(train.iloc[i])
 if(I[-1]=='pos'):
 sl=I[0].split()
 for j in sl:
   if j not in dict:
    dict[i][j]=1
    n1=n1+1
   else:
    dict[i][j]=dict[i][j]+1
 else:
  sl=I[0].split()
  for j in sl:
   if j not in dict:
    dict[i][j]=1
    n2=n2+1
   else:
    dict[i][j]=dict[i][j]+1
# print(dict)
```

```
dict2={}
dict3={}
for i in range(len(train)):
  l=list(train.iloc[i])
  if(I[-1]=='pos'):
   sl=l[0].split()
   for j in sl:
    if j not in dict2:
      dict2[j]=1
    else:
      dict2[j]=dict2[j]+1
  else:
   sl=l[0].split()
   for j in sl:
    if j not in dict3:
      dict3[j]=1
    else:
      dict3[j]=dict3[j]+1
# print(dict2) # pos words
# print(dict3) # neg words
#total vocabulary
v=0
for i in dict3:
 if i not in dict2:
  v=v+1
v=v+len(dict2)
# print(v)
pp={}
```

```
np={}
for i in dict2:
pp[i]=(dict2[i]+1)/(v+n1)
# print(pp)
for i in dict3:
np[i]=(dict3[i]+1)/(v+n2)
# print(np)
test=d[10:]
actual=list(test.iloc[:,-1])
predicted=[]
for i in range(len(test)):
 l=list(test.iloc[i])
 sl=l[0].split()
 p_prob=py
 n_prob=pn
 for j in sl:
  if j in pp.keys():
  p_prob*=dict2[j]
  else:
  p_prob*=(1)/(v+n1)
  if j in np.keys():
  n_prob*=dict3[j]
  else:
  n_{prob}*=(1)/(v+n2)
 # print(I[0],p_prob,n_prob)
 if p_prob>n_prob:
   predicted.append('pos')
 else:
 predicted.append('neg')
# print(actual)
```

print(predicted)

OUTPUT:-

```
[[3 1]
[0 4]]
accuracy 0.875
```

precision 0.8 recall 1.0

5. Write a program to construct a Bayesian network considering medical data. Use this model to demonstrate the diagnosis the of heart patients using standard heart disease data set. You can use Java or Python ML Library classes /API.

```
import pandas as pd
import numpy as np
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
heart=pd.read csv("/content/heart.csv")
heart
heart = heart.replace('?', np.nan)
print('Sample instances from the dataset are given below')
print (heart.head())
print("\n Attributes and datatypes")
print (heart.dtypes)
model =BayesianModel([('age', 'target'), ('sex', 'target'), ('exang', 'target'),
('cp','target'),('target','restecg'), ('target', 'chol')])
model.fit(heart, estimator=MaximumLikelihoodEstimator)
print("\n Inferencing with Bayesian Network:")
HeartDiseasetest_infer= VariableElimination(model)
print("\n 1. Probability of HeartDisease given evidence-restecg: 1 and age=40")
q1=HeartDiseasetest_infer.query(variables=['target'], evidence={'restecg':1,'age':50})
print(q1)
print("\n 2.Probability of HeartDisease given evidence- cp:2 and sex-1")
q2=HeartDiseasetest infer.query(variables=['target'], evidence={'cp':2,'sex':1})
print(q2)
```

OUTPUT:-

```
Inferencing with Bayesian Network:
```

1. Probability of HeartDisease given evidence-restecg :1 and age=40

2. Probability of HeartDisease given evidence- cp:2 and sex-1

```
| target | phi(target) |
| target | phi(target) |
| target(0) | 0.4076 |
| target(1) | 0.5924 |
```

6. Write a program to demonstrate the working of the decision tree based ID3 algorithm. Use an appropriate data set for building the decision tree and apply this knowledge to classify a new sample.

import pandas as pd

import numpy as np

from collections import Counter

```
def entropy(labels):
```

```
counter = Counter(labels)
probabilities = [count / len(labels) for count in counter.values()]
return -np.sum(probabilities * np.log2(probabilities))
```

def information_gain(data, attribute, labels):

```
total_entropy = entropy(labels)
attribute_values = set(data[attribute])
weighted_entropy = 0
for value in attribute_values:
    subset_labels = labels[data[attribute] == value]
    weighted_entropy += len(subset_labels) / len(labels) * entropy(subset_labels)
return total_entropy - weighted_entropy
```

```
def id3(data, attributes, labels):
  # Base cases
  if len(set(labels)) == 1: # All instances have the same label
    return labels[0]
  if len(attributes) == 0: # No more attributes to split on
    most_common_label = Counter(labels).most_common(1)[0][0]
    return most_common_label
  # Attribute selection
  information_gains = [information_gain(data, attribute, labels) for attribute in attributes]
  best_attribute_index = np.argmax(information_gains)
  best_attribute = attributes[best_attribute_index]
  # Recursion
  attribute_values = set(data[best_attribute])
  tree = {best_attribute: {}}
  for value in attribute_values:
    subset_indices = data[best_attribute] == value
    subset_data = data[subset_indices]
    subset_labels = labels[subset_indices]
    if len(subset data) == 0:
      most_common_label = Counter(labels).most_common(1)[0][0]
      tree[best_attribute][value] = most_common_label
    else:
      remaining attributes = attributes[:best attribute index] + attributes[best attribute index+
1:]
      tree[best_attribute][value] = id3(subset_data, remaining_attributes, subset_labels)
  return tree
def predict(instance, tree):
  if isinstance(tree, dict)==False:
```

return tree

```
attribute_value = instance[list(tree.keys())[0]]
  if attribute_value not in list(tree.values())[0]:
    return None
  child = tree[list(tree.keys())[0]][attribute_value]
  return predict(instance, child)
# Example usage
data = pd.read_csv('/content/play_tennis.csv')
attributes = data.columns[1:-1].tolist()
labels = data.iloc[:, -1].values
decision_tree = id3(data, attributes, labels)
# Example prediction
instance = {'outlook': 'Sunny', 'temp': 'Cool', 'humidity': 'High', 'wind': 'Weak'}
prediction = predict(instance, decision_tree)
print("Prediction:", prediction)
print(decision_tree)
OUTPUT:-
Prediction: No
{'outlook': {'Rain': {'wind': {'Weak': 'Yes', 'Strong': 'No'}}, 'Sunny': {'humidity': {'Normal': 'Yes', 'High':
'No'}}, 'Overcast': 'Yes'}}
7. Build an Artificial Neural Network by implementing the Backpropagation algorithm and
test the same using appropriate data sets.
import numpy as np
```

```
X = \text{np.array}(([2, 9], [1, 5], [3, 6]), \text{dtype=float})
y = np.array(([92], [86], [89]), dtype=float)
X = X/np.amax(X,axis=0) #maximum of X array longitudinally
y = y/100
#Sigmoid Function
def sigmoid (x):
  return 1/(1 + np.exp(-x))
#Derivative of Sigmoid Function
def derivatives_sigmoid(x):
  return x * (1 - x)
#Variable initialization
epoch=5 #Setting training iterations
lr=0.1 #Setting learning rate
inputlayer_neurons = 2 #number of features in data set
hiddenlayer_neurons = 3 #number of hidden layers neurons
output_neurons = 1 #number of neurons at output layer
#weight and bias initialization
wh=np.random.uniform(size=(inputlayer_neurons,hiddenlayer_neurons))
bh=np.random.uniform(size=(1,hiddenlayer_neurons))
wout=np.random.uniform(size=(hiddenlayer_neurons,output_neurons))
```

```
bout=np.random.uniform(size=(1,output_neurons))
#draws a random range of numbers uniformly of dim x*y
for i in range(epoch):
  #Forward Propogation
  hinp1=np.dot(X,wh)
  hinp=hinp1 + bh
  hlayer_act = sigmoid(hinp)
  outinp1=np.dot(hlayer_act,wout)
  outinp= outinp1+bout
  output = sigmoid(outinp)
  #Backpropagation
  EO = y-output
  outgrad = derivatives_sigmoid(output)
  d_output = EO * outgrad
  EH = d\_output.dot(wout.T)
  hiddengrad = derivatives_sigmoid(hlayer_act)#how much hidden layer wts contributed to
error
  d_hiddenlayer = EH * hiddengrad
  wout += hlayer_act.T.dot(d_output) *lr # dotproduct of nextlayererror and currentlayerop
  wh += X.T.dot(d_hiddenlayer) *lr
  print ("------")
  print("Input: \n'' + str(X))
```

```
print("Actual Output: \n" + str(y))
  print("Predicted Output: \n",output)
  print ("------Epoch-", i+1, "Ends -----\n")
print("Input: \n" + str(X))
print("Actual Output: \n" + str(y))
print("Predicted Output: \n" ,output)
OUTPUT:-
Input:
[[0.66666667 1.
 [0.33333333 0.55555556]
 [1. 0.66666667]]
Actual Output:
[[0.92]
 [0.86]
 [0.89]]
Predicted Output:
 [[0.91755203]
 [0.90180959]
 [0.91814824]]
8. Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same data set for
clustering using K-Means algorithm. Compare the results of these two algorithms and
comment on the quality of clustering. You can add Java / Python ML library classes/API in
the program.
import matplotlib.pyplot as plt
from sklearn import datasets
import pandas as pd
from sklearn.cluster import KMeans
import numpy as np
```

iris = datasets.load_iris()

X = pd.DataFrame(iris.data)

```
X.columns = ['Sepal_Length', 'Sepal_width', 'Petal_Length', 'Petal_width']
y = pd.DataFrame(iris.target)
y.columns = ['Targets']
# Build the K Means Model
model = KMeans (n_clusters=3)
model.fit(X)
plt.figure(figsize=(14,14))
colormap = np.array(['red', 'lime', 'black'])
# Plot the original Classifications using Petal features
plt.subplot(2, 2, 1)
plt.scatter(X.Petal_Length, X.Petal_width, c=colormap[y.Targets], s=40)
plt.title('Real Clusters')
plt.xlabel('Petal Length')
plt.ylabel('Petal width')
plt.subplot(2, 2, 2)
plt.scatter(X.Petal_Length, X.Petal_width, c=colormap[model.labels_], s=40)
plt.title('K-Means Clustering')
plt.xlabel('Petal Length')
plt.ylabel('Petal width')
from sklearn import preprocessing
# transform your data such that its distribution will have a # mean value 0 and standard
deviation of 1.
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
```

from sklearn.mixture import GaussianMixture

gmm=GaussianMixture(n_components=3)

gmm.fit(xs)

gmm_y=gmm.predict(xs)

plt.subplot(2, 2, 3)

plt.scatter (X.Petal_Length, X.Petal_width, c=colormap[gmm_y], s=40)

plt.title('GMM Clustering')

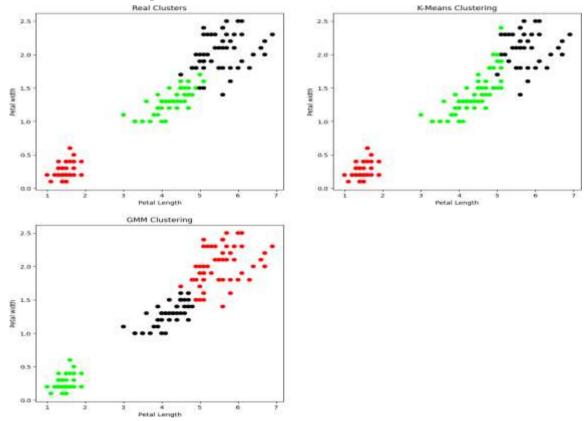
plt.xlabel('Petal Length')

plt.ylabel('Petal width')

print('Observation: The GMM using EM algorithm based clustering matched the true labels more closely than the Kmeans.')

OUTPUT:-

Observation: The GMM using EM algorithm based clustering matched the true labels more closely than the Kmeans.



9. Write a program to implement k-Nearest Neighbor algorithm to classify the iris data set. Print both correct and wrong predictions. Java/Python ML library classes can be used for thisproblem.

```
from sklearn.model selection import train test split
from sklearn.neighbors import KNeighborsClassifier
from sklearn import datasets
#Load dataset
iris=datasets.load_iris()
print("Iris Data set loaded...")
# print(iris)
# Split the data into train and test samples
x train, x test, y train, y test = train test split(iris.data,iris.target,test size=0.1)
print("Dataset is split into training and testing...")
print("size of training data and its label",x train.shape,y train.shape)
print("Size of training data and its label",x test.shape, y test.shape)
# Prints Label no. and their names
for i in range(len(iris.target_names)):
   print("Label", i, "-",str(iris.target_names[i]))
   # Create object of KNN classifier
   classifier = KNeighborsClassifier(n neighbors=1)
#Perform Training
classifier.fit(x_train, y_train)
# Perform testing
y_pred=classifier.predict(x_test)
# Display the results
print("Results of Classification using K-nn with K=1")
for r in range(0,len(x_test)):
   print(" Sample:", str(x_test[r]), " Actual-label:", str(y_test[r]), "Predicted-label:", str(y_pred[r]))
   print("Classification Accuracy:", classifier.score(x_test,y_test));
```

```
from sklearn.metrics import classification report, confusion matrix
print('Confusion Matrix')
print(confusion matrix(y test,y pred))
print('Accuracy Metrics')
print(classification report(y test,y pred))
OUTPUT:-
Iris Data set loaded...
Dataset is split into training and testing...
size of training data and its label (135, 4) (135,)
Size of training data and its label (15, 4) (15,)
Label 0 - setosa
KNeighborsClassifier(n neighbors=1)
Label 1 - versicolor
KNeighborsClassifier(n neighbors=1)
Label 2 - virginica
KNeighborsClassifier(n neighbors=1)
Results of Classification using K-nn with K=1
Sample: [6.1 2.9 4.7 1.4] Actual-label: 1 Predicted-label: 1
Classification Accuracy: 0.9333333333333333
Sample: [6. 2.2 5. 1.5] Actual-label: 2 Predicted-label: 1
Classification Accuracy: 0.9333333333333333
Sample: [4.5 2.3 1.3 0.3] Actual-label: 0 Predicted-label: 0
Classification Accuracy: 0.9333333333333333
Sample: [7.7 2.8 6.7 2.] Actual-label: 2 Predicted-label: 2
Classification Accuracy: 0.9333333333333333
Sample: [6.7 3.1 4.4 1.4] Actual-label: 1 Predicted-label: 1
Classification Accuracy: 0.93333333333333333
Sample: [6.3 2.3 4.4 1.3] Actual-label: 1 Predicted-label: 1
Classification Accuracy: 0.93333333333333333
Sample: [5.1 3.7 1.5 0.4] Actual-label: 0 Predicted-label: 0
Classification Accuracy: 0.93333333333333333
Sample: [5.5 2.6 4.4 1.2] Actual-label: 1 Predicted-label: 1
Sample: [5.4 3.9 1.3 0.4] Actual-label: 0 Predicted-label: 0
Sample: [4.8 3. 1.4 0.3] Actual-label: 0 Predicted-label: 0
Sample: [7.2 3. 5.8 1.6] Actual-label: 2 Predicted-label: 2
Sample: [6.5 3.
                5.2 2. | Actual-label: 2 Predicted-label: 2
Sample: [5.6 2.8 4.9 2.] Actual-label: 2 Predicted-label: 2
Sample: [4.9 3.1 1.5 0.2] Actual-label: 0 Predicted-label: 0
Classification Accuracy: 0.9333333333333333
Sample: [6.4 3.1 5.5 1.8] Actual-label: 2 Predicted-label: 2
Confusion Matrix
[[5 0 0]
 [0 4 0]
 [0 1 5]]
Accuracy Metrics
            precision recall f1-score
                                         support
                1.00
                         1.00
                                  1.00
                                              5
```

```
1
                              1.00
                   0.80
                                        0.89
                                                     4
                   1.00
                              0.83
                                        0.91
                                                     6
                                        0.93
                                                    15
    accuracy
                              0.94
                                        0.93
                                                    15
   macro avg
                   0.93
                                        0.93
                                                    15
weighted avg
                   0.95
                              0.93
10. Implement the non-parametric Locally Weighted Regression algorithm in
order to fit data points. Select appropriate data set your experiment and
draw graphs.
import numpy as np
import pandas as pd
# kernel smoothing function
def kernel(point, xmat, k):
   m, n= np.shape(xmat)
   weights =np.mat(np.eye((m)))
    for j in range(m):
        diff =point-xmat[j]
        weights[j, j] = np.exp(diff* diff.T/(-2.0 * k**2)) # smooth = exp
-(x-x0)^2/2k^2
    return weights
# function to return local weight of eah traiining example
def localweight (point, xmat, ymat, k):
    wt =kernel(point, xmat, k)
    W = ((xmat.T * (wt*xmat)).I)* (xmat.T* wt* ymat.T) # beta = (xtrans *
smooth * x)inverse *(xtrans * smooth * y)
    return W
def localweightRegression(xmat, ymat, k):
    m, n= np. shape(xmat)
    ypred = np.zeros(m)
    for i in range(m):
        ypred[i] = xmat[i] *localweight(xmat[i], xmat, ymat, k)
                                                                   # y = x0
* beta
    return ypred
import matplotlib.pyplot as plt
#import data
data = pd.read csv('10-dataset.csv')
# place them in suitable data types
colA = np.array(data.total bill)
colB= np.array(data.tip)
# print(colA)
# print(colB)
mcolA = np.mat (colA)
mcolB= np.mat (colB)
# print(mcolA)
# print(mcolB)
m = np.shape(mcolB) [1]
# print(m)
one = np.ones((1, m), dtype = int)
# print(one)
# print(one.T, mcolA.T)
# horizontal stacking
X = np.hstack((one. T, mcolA.T))
```

```
# print(X)
print(X. shape)
# predicting values using LWLR
ypred = localweightRegression(X, mcolB, 0.8)
# plotting the predicted graph
xsort = X.copy()
xsort.sort(axis=0)
plt.scatter (colA, colB, color='blue')
plt.plot(xsort[:, 1], ypred[X[:, 1].argsort(0)], color='yellow',
linewidth=5)
plt.xlabel('Total Bill')
plt.ylabel('Tip')
plt.show()
OUTPUT: -
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