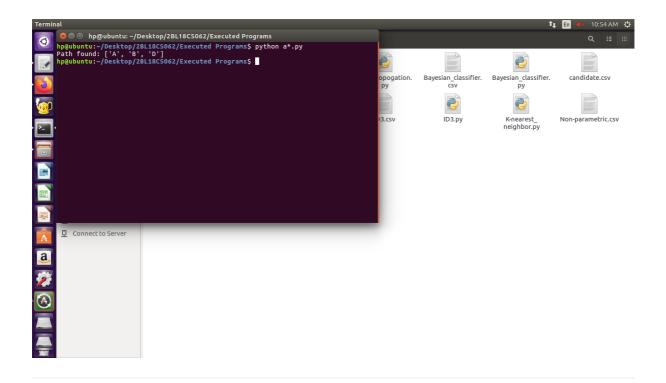
AI-ML Lab Programs and Outputs

1. Implement A* Search algorithm.

```
Algorithm:
01: Create a node containing the goal state node goal
02: Create a node containing the start state node start
03: Put node start on the open list
04: while the OPEN list is not empty
05: {
06: Get the node off the open list with the lowest f and call it node_current
07: if node current is the same state as node goal we have found the
solution; break from the while loop
08: Generate each state node successor that can come after node current
09: for each node_successor of node_current
10: {
11:
         Set the cost of node successor to be the cost of node current plus
the cost to get to node successor from node current
12:
         find node successor on the OPEN list
13:
         if node successor is on the OPEN list but the existing one is as
good or better than discard this successor and continue
         if node successor is on the CLOSE list but the existing one is as
good or better than discard this successor and continue
15:
         Remove occurrences of node successor from OPEN and CLOSED
16:
         Set the parent of node successor to node current
17:
         Set h to be the estimated distance to node_goal(Using the heuristic
function)
18:
         Add node successor to the OPEN list
19: }
20: Add node_current to the CLOSED list
21: }
```

```
from collections import deque
class Graph:
    def __init__(self, adjac_lis):
        self.adjac_lis = adjac_lis
    def get_neighbors(self, v):
```

```
return self.adjac_lis[v]
    def h(self, n):
       H = {
            'A': 1,
           'B': 1,
           'C': 1,
           'D': 1
        }
        return H[n]
    def a_star_algorithm(self, start, stop):
       open_lst = set([start])
        closed_lst = set([])
       poo = {}
        poo[start] = 0
        par = \{\}
        par[start] = start
        while len(open_lst) > 0:
           n = None
            for v in open_lst:
               if n == None or poo[v] + self.h(v) < poo[n] + self.h(n):
                  n = v;
            if n == None:
               print('Path does not exist!')
                return None
            if n == stop:
               reconst_path = []
                while par[n] != n:
                   reconst_path.append(n)
                    n = par[n]
                reconst_path.append(start)
                reconst_path.reverse()
                print('Path found: {}'.format(reconst_path))
                return reconst_path
            for (m, weight) in self.get_neighbors(n):
                if m not in open_lst and m not in closed_lst:
                   open_lst.add(m)
                    par[m] = n
                    poo[m] = poo[n] + weight
                else:
                    if poo[m] > poo[n] + weight:
                       poo[m] = poo[n] + weight
                        par[m] = n
                        if m in closed_lst:
                            closed_lst.remove(m)
                            open_lst.add(m)
            open_lst.remove(n)
            closed_lst.add(n)
        print('Path does not exist!')
       return None
adjac_lis = {
    'A': [('B', 1), ('C', 3), ('D', 7)],
    'B': [('D', 5)],
'C': [('D', 12)]
graph1 = Graph(adjac_lis)
graph1.a_star_algorithm('A', 'D')
```



2. Implement AO* Search algorithm.

Algorithm:

OPEN:

It contains the nodes that has been traversed but yet not been marked solvable or unsolvable.

CLOSE:

It contains the nodes that have already been processed.

- Step 1: Place the starting node into OPEN.
- Step 2: Compute the most promising solution tree say T0.
- Step 3: Select a node n that is both on OPEN and a member of T0. Remove it from OPEN and place it in CLOSE
- Step 4: If n is the terminal goal node then leveled n as solved and leveled all the ancestors of n as solved. If the starting node is marked as solved then success and exit.
- Step 5: If n is not a solvable node, then mark n as

unsolvable. If starting node is marked as unsolvable, then return failure and exit.

Step 6: Expand n. Find all its successors and find their h (n) value, push them into OPEN.

Step 7: Return to Step 2.

Step 8: Exit.

```
class Graph:
   def __init__(self, graph, heuristicNodeList, startNode):
       self.graph = graph
       self.H=heuristicNodeList
       self.start=startNode
       self.parent={}
       self.status={}
       self.solutionGraph={}
   def applyAOStar(self):
       self.aoStar(self.start, False)
   def getNeighbors(self, v):
       return self.graph.get(v,'')
   def getStatus(self,v):
       return self.status.get(v,0)
   def setStatus(self,v, val):
       self.status[v]=val
   def getHeuristicNodeValue(self, n):
       return self.H.get(n,0)
   def setHeuristicNodeValue(self, n, value):
       self.H[n]=value
   def printSolution(self):
       print("FOR GRAPH SOLUTION, TRAVERSE THE GRAPH FROM THE START NODE:", self.start)
       print("----")
       print(self.solutionGraph)
       print("-----
   def computeMinimumCostChildNodes(self, v):
       minimumCost=0
       costToChildNodeListDict={}
       costToChildNodeListDict[minimumCost]=[]
       for nodeInfoTupleList in self.getNeighbors(v):
           nodeList=[]
           for c, weight in nodeInfoTupleList:
              cost=cost+self.getHeuristicNodeValue(c)+weight
               nodeList.append(c)
           if flag==True:
              minimumCost=cost
               costToChildNodeListDict[minimumCost]=nodeList
               flag=False
           else:
               if minimumCost>cost:
```

```
minimumCost=cost
                    \verb|costToChildNodeListDict[minimumCost] = nodeList|
        return minimumCost, costToChildNodeListDict[minimumCost]
    def aoStar(self, v, backTracking):
        print("HEURISTIC VALUES :", self.H)
        print("SOLUTION GRAPH :", self.solutionGraph)
        print("PROCESSING NODE :", v)
        print("-----
        if self.getStatus(v) >= 0:
           minimumCost, childNodeList = self.computeMinimumCostChildNodes(v)
            print(minimumCost, childNodeList)
            self.setHeuristicNodeValue(v, minimumCost)
            self.setStatus(v,len(childNodeList))
            solved=True
            for childNode in childNodeList:
                self.parent[childNode]=v
                if self.getStatus(childNode)!=-1:
                   solved=solved & False
            if solved==True:
               self.setStatus(v,-1)
                self.solutionGraph[v]=childNodeList
            if v!=self.start:
                self.aoStar(self.parent[v], True)
            if backTracking==False:
                for childNode in childNodeList:
                   self.setStatus(childNode,0)
                   self.aoStar(childNode, False)
print ("Graph - 1")
h1 = {'A': 1, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1}
graph1 = {
    'A': [[('B', 1), ('C', 1)], [('D', 1)]],
    'B': [[('G', 1)], [('H', 1)]],
    'C': [[('J', 1)]],
    'D': [[('E', 1), ('F', 1)]],
    'G': [[('I', 1)]]
}
G1= Graph(graph1, h1, 'A')
G1.applyAOStar()
G1.printSolution()
```

3. 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.

Algorithm:

For each training example d, do:

If d is positive example

Remove from G any hypothesis h inconsistent with d

For each hypothesis s in S not consistent with d:

Remove s from S

Add to S all minimal generalizations of s consistent with d and having a generalization in G

Remove from S any hypothesis with a more specific h in S

If d is negative example

Remove from S any hypothesis h inconsistent with d

For each hypothesis g in G not consistent with d:

Remove g from G

Add to G all minimal specializations of g consistent with d and

having a specialization in S

Remove from G any hypothesis having a more general hypothesis

in G

Source code:

```
import csv
import numpy as np
with open('candidate.csv','r') as f:
   reads=csv.reader(f)
   tmp_lst=np.array(list(reads))
concept=np.array(tmp_lst[:,:-1])
target=np.array(tmp_lst[:,-1])
for i in range(len(target)):
   if(target[i]=='yes'):
       specific_h=concept[i]
h=[]
generic_h=[['?' for i in range (len(specific_h))]for i in range (len(specific_h))]
print(type(generic_h))
for i in range(len(target)):
   if(target[i]=='yes'):
       for j in range (len(specific_h)):
           if(specific_h[j]!=concept[i][j]):
               specific_h[j]='?'
              generic_h[j][j]='?'
       for j in range(len(specific_h)):
           if(specific_h[j]!=concept[i][j]):
               generic_h[j][j]=specific_h[j]
               generic_h[j][j]='?'
   print("Step ",i+1)
   print("The most generic is : ",generic_h)
    print("The most specific is : ",specific_h)
```

4. 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.

Algorithm:

ID3(Examples, Target attribute, Attributes)

Examples are the training examples.

Target_attribute is the attribute whose value is to be predicted by the tree. Attributes is a list of other attributes that may be tested by the learned decision tree.

Returns a decision tree that correctly classifies the given Examples.

Create a Root node for the tree

If all Examples are positive, Return the single-node tree Root, with label = + If all Examples are negative, Return the single-node tree Root, with label = - If Attributes is empty, Return the single-node tree Root, with label = most common value of Target attribute in Examples

Otherwise Begin

A ← the attribute from Attributes that best* classifies Examples

```
The decision attribute for Root ← A

For each possible value, vi, of A,

Add a new tree branch below Root, corresponding to the test A = vi

Let Examples vi, be the subset of Examples that have value vi for A

If Examples vi , is empty

Then below this new branch add a leaf node with

label = most common value of Target_attribute in Examples

Else

below this new branch add the subtree

ID3(Examples vi, Target_attribute, Attributes – {A}))

End

Return Root
```

```
import numpy as np
import math
import csv
class Node:
 def __init__ (self,attribute):
   self.attribute=attribute
   self.children=[]
   self.answer=" "
def read_data(filename):
 with open(filename, 'r') as csvfile:
    datareader=csv.reader(csvfile,delimiter=',')
   headers=next(datareader)
   metadata=[]
    traindata=[]
    for name in headers:
     metadata.append(name)
    for row in datareader:
      traindata.append(row)
 return(metadata, traindata)
def subtables(data,col,delete):
 items=np.unique(data[:,col])
 count=np.zeros((items.shape[0],1),dtype=np.int32)
  for x in range(items.shape[0]):
   for y in range(data.shape[0]):
     if data[y,col]==items[x]:
       count[x]+=1
  for x in range(items.shape[0]):
    dict[items[x]]=np.empty((int (count[x]),data.shape[1]),dtype="|S32")
    for y in range(data.shape[0]):
     if data[y,col]==items[x]:
       dict[items[x]][pos]=data[y]
        pos+=1
    if delete:
     dict[items[x]]=np.delete(dict[items[x]],col,1)
  return items, dict
def entropy(S):
  items=np.unique(S)
  if items.size==1:
    return 0
```

```
counts = np.zeros((items.shape[0],1))
  sums = 0
  for x in range(items.shape[0]):
   counts[x] = sum(S ==items[x])/(S.size*1.0)
  for count in counts:
   sums +=-1*count*math.log(count,2)
def gain_ratio(data,col):
 items, dict=subtables(data, col, delete=False)
  total_size=data.shape[0]
  entropies=np.zeros((items.shape[0],1))
  intrinsic=np.zeros((items.shape[0],1))
  for x in range(items.shape[0]):
    ratio=dict[items[x]].shape[0]/(total_size*1.0)
    entropies[x]=ratio*entropy(dict[items[x]][:,-1])
   intrinsic[x]=ratio*math.log(ratio,2)
    total_entropy=entropy(data[:,-1])
    iv=-1*sum(intrinsic)
    for x in range(entropies.shape[0]):
     total_entropy-=entropies[x]
    return total_entropy/iv
def create_node(data,metadata):
 if(np.unique(data[:,-1])).shape[0]==1:
    node = Node(" ")
    node.answer = np.unique(data[:,-1])[0]
    return node
  gains = np.zeros((data.shape[1]-1,1))
  for col in range(data.shape[1]-1):
    gains[col]=gain_ratio(data,col)
  split=np.argmax(gains)
  node=Node(metadata[split])
  metadata=np.delete(metadata,split,0)
  items, dict=subtables(data, split, delete=True)
  for x in range(items.shape[0]):
    child = create_node(dict[items[x]], metadata)
   node.children.append((items[x],child))
  return node
def empty(size):
 S = " "
  for x in range(size):
    S+=" "
  return S
def print_tree(node, level):
  if node.answer!=" ":
    print(empty(level), node.answer)
  print(empty(level), node.attribute)
  for value, n in node.children:
    print(empty(level+1), value)
    print_tree(n, level+2)
metadata, traindata=read_data("ID3.csv")
data=np.array(traindata)
node=create_node(data, metadata)
print_tree(node,0)
```

5. Build an Artificial Neural Network by implementing the Backpropagation algorithm and test the same using appropriate data sets.

Algorithm:

- 1: Run the network forward with your input data to get the network output
- 2: For each output node compute

$$\delta_k = \mathcal{O}_k \left(1 - \mathcal{O}_k
ight) \left(\mathcal{O}_k - t_k
ight)$$

3: For each hidden node calculate

$$\delta_j = \mathcal{O}_j \left(1 - \mathcal{O}_j
ight) \sum_{k \in K} \delta_k W_{jk}$$

4: Update the weights and biases as follows Given

$$egin{array}{ll} \Delta W = & -\eta \delta_\ell \mathcal{O}_{\ell-1} \ \Delta heta = \eta \delta_\ell \end{array}$$

Apply

$$W + \Delta W > W$$
 $heta + \Delta heta > heta$

```
import numpy as np
X=np.array(([2,9],[1,5],[3,6]),dtype=float)
y=np.array(([92],[86],[89]),dtype=float)
X=X/np.amax(X,axis=0)
y=y/100
def sigmoid(x):
 return 1/(1+np.exp(-x))
def derivatives_sigmoid(x):
 return x*(1-x)
epoch=7000
lr=0.25
inputlayer_neurons=2
hiddenlayer_neurons=3
output_neurons=1
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))
for i in range(epoch):
 hinp1=np.dot(X,wh)
 hinp=hinp1+bh
 hlayer_act=sigmoid(hinp)
 outinp1=np.dot(hlayer_act,wout)
 outinp=outinp1+bout
 output=sigmoid(outinp)
 E0=y-output
 outgrad=derivatives_sigmoid(output)
 d_output=E0*outgrad
 EH=d_output.dot(wout.T)
 hiddengrad=derivatives_sigmoid(hlayer_act)
 d_hiddenlayer=EH*hiddengrad
 wout+=hlayer_act.T.dot(d_output)*lr
 wh+=X.T.dot(d_hiddenlayer)*lr
print("Input=\n"+str(X))
print("Actual output:\n"+str(y))
print("predicated output:",output)
```

```
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```

6. 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.

Naïve Bayesian Classifier:

Bayes' Theorem is stated as:

$$P(h\mid D) = \frac{P(D\mid h)P(h)}{P(D)}$$

Steps to implement Naïve Bayesian Classifier:

Step 1: Separate By Class.

Step 2: Summarize Dataset.

Step 3: Summarize Data By Class.

Step 4: Gaussian Probability Density Function.

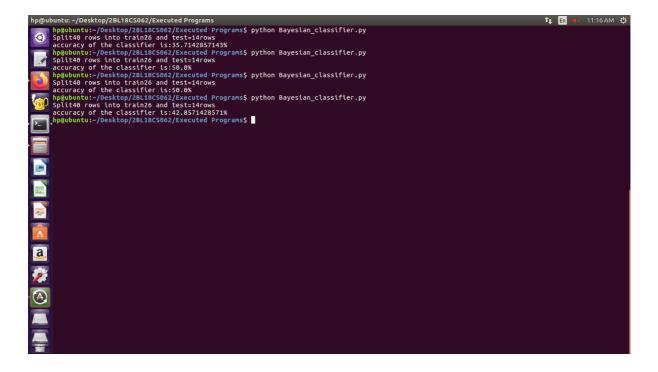
Step 5: Class Probabilities.

```
import csv
import random
```

```
import math
def loadCsv(filename):
  lines = csv.reader(open(filename, "r"));
  dataset = list(lines)
  for i in range(len(dataset)):
    dataset[i] = [float(x) for x in dataset[i]]
def splitDataset(dataset, splitRatio):
  trainSize = int(len(dataset) * splitRatio);
  trainSet = []
  copy = list(dataset);
  while len(trainSet) < trainSize:</pre>
    index = random.randrange(len(copy));
    trainSet.append(copy.pop(index))
  return [trainSet, copy]
def separateByClass(dataset):
  separated = {}
  for i in range(len(dataset)):
    vector = dataset[i]
    if (vector[-1] not in separated):
      separated[vector[-1]] = []
    separated[vector[-1]].append(vector)
 return separated
def mean(numbers):
  return sum(numbers)/float(len(numbers))
def stdev(numbers):
 avg = mean(numbers)
  variance = sum([pow(x-avg, 2) for x in numbers])/float(len(numbers)-1)
  return math.sqrt(variance)
def summarize(dataset):
 summaries = [(mean(attribute), stdev(attribute)) for attribute in zip(*dataset)];
  del summaries[-1]
  return summaries
def summarizeByClass(dataset):
  separated=separateByClass(dataset)
  summaries={}
  for classValue, instances in separated.items():
    summaries[classValue]=summarize(instances)
  return summaries
{\tt def\ calculateProbability(x,mean,stdev):}
  exponent = math.exp(-(math.pow(x-mean,2)/(2*math.pow(stdev,2))))
  return (1 / (math.sqrt(2*math.pi) * stdev)) * exponent
def calculateClassProbabilities(summaries, inputVector):
  probabilities = {}
  for classValue, classSummaries in summaries.items():
    probabilities[classValue] = 1
    for i in range(len(classSummaries)):
      mean, stdev = classSummaries[i]
      x = inputVector[i]
      probabilities[classValue] *= calculateProbability(x, mean, stdev);
    return probabilities
def predict(summaries, inputVector):
  probabilities = calculateClassProbabilities(summaries, inputVector)
  bestLabel, bestProb = None, -1
  for classValue, probability in probabilities.items():
    if bestLabel is None or probability > bestProb:
      bestProb = probability
      bestLabel=classValue
  return bestLabel
def getPredictions(summaries,testSet):
  predictions = []
  for i in range(len(testSet)):
    result = predict(summaries, testSet[i])
    predictions.append(result)
 return predictions
def getAccuracy(testSet, predictions):
  correct = 0
  for i in range(len(testSet)):
    if testSet[i][-1] == predictions[i]:
```

```
correct += 1
return (correct/float(len(testSet))) * 100.0

def main():
    filename="Bayesian_classifier.csv"
    splitRatio=0.67
    dataset=loadCsv(filename)
    trainingSet,testSet=splitDataset(dataset,splitRatio)
    print('Split{0} rows into train{1} and test={2}rows'.format(len(dataset),len(trainingSet),len(testSet)))
    summaries = summarizeByClass(trainingSet);
    predictions=getPredictions(summaries,testSet)
    accuracy=getAccuracy(testSet,predictions)
    print('accuracy of the classifier is:{0}%'.format(accuracy))
main()
```



7. Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same data set for clustering using the 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.

Algorithm:

K-Means:

K-Means clustering algorithm produces a Minimum Variance Estimate (MVE) of the state of the identified clusters in the data.

$$J = \sum_{k \in K} \sum_i z_k^i \left| x_i - \mu_k
ight|^2$$

$$\left(\mu_k,z_k^i
ight)= \operatorname{argmin}_{\left(z_k^i,\mu_k
ight)} J$$

EM Algorithm:

Step 01: Initial guess is made for the model's parameters and a probability distribution is created. This is sometimes called **<u>E-Step</u>** for the expected distribution.

Step 02: Newly observed data is fed into a model.

Step 03: The probability distributed the <u>E-step</u> is drawn to include the new data which is sometimes called **M-step**.

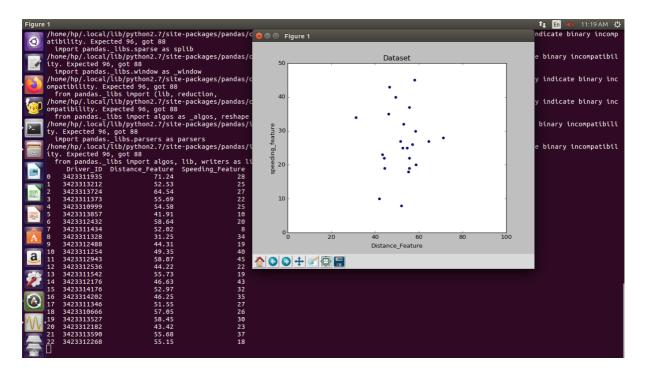
Step 04: <u>Step-02</u> through <u>Step-04</u> are repeated until S with normal distribution.

That is:

$$egin{aligned} E\left[z_{ji}
ight] &= rac{p\left(x = x_i \mid \mu = \mu j
ight)}{\sum^2 p\left(x = x_i \mid \mu = \mu(n)
ight)} \ &orall & \left(\mu_j \leftarrow rac{\sum_{i=1}^M E\left[z_{ij}
ight]x_i.}{\sum_{i=1}^M E\left[z_n
ight].} \end{aligned}$$

```
from sklearn.cluster import KMeans
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
data=pd.read_csv("EM_Algorithm.csv")
df1=pd.DataFrame(data)
print(df1)
f1 = df1['Distance_Feature'].values
f2 = df1['Speeding_Feature'].values
X=np.matrix(list(zip(f1,f2)))
plt.plot()
plt.xlim([0, 100])
plt.ylim([0, 50])
plt.title('Dataset')
plt.ylabel('speeding_feature')
plt.xlabel('Distance_Feature')
plt.scatter(f1,f2)
plt.show()
plt.plot()
colors = ['b', 'g', 'r']
markers = ['o', 'v', 's']
kmeans_model = KMeans(n_clusters=3).fit(X)
plt.plot()
```

```
for i, l in enumerate(kmeans_model.labels_):
  plt.plot(f1[i], f2[i], color=colors[l], marker=markers[l],ls='None')
  plt.xlim([0, 100])
  plt.ylim([0, 50])
plt.show()
```



8. Write a program to implement the 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 this problem.

Algorithm:

- Step-1: Select the number K of the neighbors
- Step-2: Calculate the Euclidean distance of K number of neighbors
- Step-3: Take the K nearest neighbors as per the calculated Euclidean distance.
- Step-4: Among these k neighbors, count the number of the data points in each category.
- Step-5: Assign the new data points to that category for which the number of the neighbor is maximum.
- Step-6: The model is ready.

```
from \ sklearn.model\_selection \ import \ train\_test\_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report, confusion_matrix
from sklearn import datasets
iris=datasets.load_iris()
iris_data=iris.data
iris_labels=iris.target
print(iris_data)
print(iris_labels)
x\_train, \ x\_test, \ y\_train, \ y\_test=train\_test\_split(iris\_data, iris\_labels, test\_size=0.30)
classifier=KNeighborsClassifier(n_neighbors=5)
classifier.fit(x_train,y_train)
y_pred=classifier.predict(x_test)
print('confusion matrix is as follows')
print(confusion_matrix(y_test,y_pred))
print('Accuracy metrics')
print(classification_report(y_test,y_pred))
```

```
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from ...swnlight_format inport _load_swnlight_file

[5. 13.5 1.4 0.2]

[6. 3.5 1.4 0.2]

[6. 3.5 1.4 0.2]

[6. 3.4 1.4 0.3]

[6. 3.4 1.4 0.3]

[6. 3.4 1.5 0.2]

[6. 3.4 1.5 0.2]

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[6. 3.4 1.5 0.2]

[6. 3.5 1.4 0.3]

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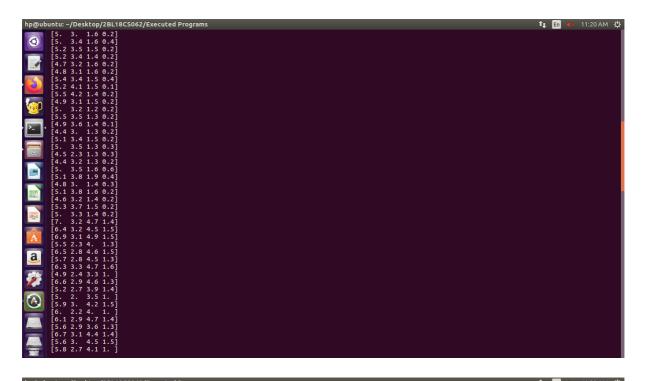
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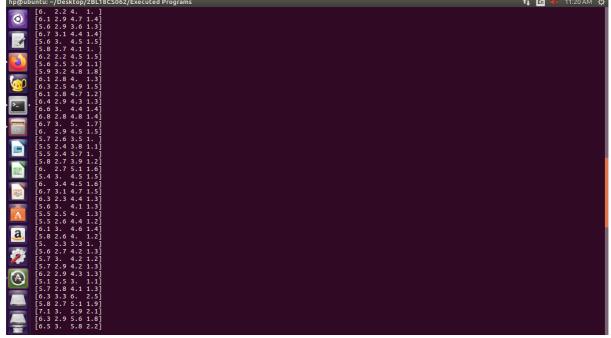
[6. 3.6 1. 0.2]

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```





9. Implement the non-parametric Locally Weighted Regression algorithm in order to fit data points. Select the appropriate data set for your experiment and draw graphs.

Algorithm:

- 01: Read the Given data Sample to X and the curve (linear or non linear) to Y
- 02: Set the value for Smoothening parameter or Free parameter say τ
- 03: Set the bias /Point of interest set x0 which is a subset of X
- 04: Determine the weight matrix using :

$$w\left(x,x_{o}
ight)=e^{-rac{\left(x-x_{o}
ight)^{2}}{2 au^{2}}}$$

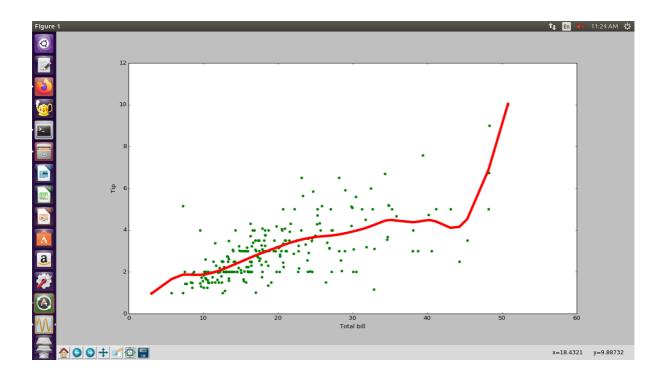
05: Determine the value of model term parameter β using:

$$\hat{eta}\left(x_{o}
ight)=\left(X^{T}WX
ight)^{-1}X^{T}Wy$$

06: Prediction = $x0*\beta$

```
from numpy import *
import operator
from os import listdir
```

```
import matplotlib
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np1
import numpy.linalg as np
from scipy.stats.stats import pearsonr
def kernel(point,xmat, k):
 m, n = np1.shape(xmat)
 weights = np1.mat(np1.eye((m)))
 for j in range(m):
   diff = point - X[j]
   weights[j,j] = np1.exp(diff*diff.T/(-2.0*k**2))
 return weights
def localWeight(point,xmat,ymat,k):
 wei = kernel(point,xmat,k)
 W=(X.T*(wei*X)).I*(X.T*(wei*ymat.T))
  return W
def localWeightRegression(xmat,ymat,k):
 m, n = np1.shape(xmat)
 ypred = np1.zeros(m)
  for i in range(m):
   ypred[i] = xmat[i]*localWeight(xmat[i],xmat,ymat,k)
 return ypred
# load data points
data = pd.read_csv('Non-Parametric.csv')
bill = np1.array(data.total_bill)
tip = np1.array(data.tip)
#preparing and add 1 in bill
mbill = np1.mat(bill)
mtip = np1.mat(tip)
m= np1.shape(mbill)[1]
one = np1.mat(np1.ones(m))
X= np1.hstack((one.T,mbill.T))
#set k here
ypred = localWeightRegression(X,mtip,2)
SortIndex = X[:,1].argsort(0)
xsort = X[SortIndex][:,0]
fig = plt.figure()
ax = fig.add_subplot(1,1,1)
ax.scatter(bill,tip, color='green')
ax.plot(xsort[:,1],ypred[SortIndex], color = 'red', linewidth=5)
plt.xlabel('Total bill')
plt.ylabel('Tip')
plt.show();
```



The below file contains all the python programs and the respective .csv datasets and also the output printout pdf for 7th and 9th program.

 $\underline{https://drive.google.com/drive/folders/1dtFdFmEc7G23VRtscFxDZMZt5tlazQkX?usp=sharing}$