1. A STAR

```
def aStarAlgo(start_node, stop_node):
  open_set = set(start_node)
  closed_set = set()
  g = {} #store distance from starting node
  parents = {}
  g[start\_node] = 0
  parents[start_node] = start_node
  while len(open_set) > 0:
     n = None
     for v in open_set:
       if n == None \text{ or } g[v] + heuristic(v) < g[n] + heuristic(n):
          n = v
     if n == stop_node or Graph_nodes[n] == None:
       pass
     else:
       for (m, weight) in get_neighbors(n):
          if m not in open_set and m not in closed_set:
             open_set.add(m)
            parents[m] = n
            g[m] = g[n] + weight
          else:
            if g[m] > g[n] + weight:
               g[m] = g[n] + weight
               parents[m] = n
               if m in closed_set:
                  closed_set.remove(m)
                  open_set.add(m)
     if n == stop node:
       path = []
       while parents[n] != n:
          path.append(n)
          n = parents[n]
       path.append(start node)
       path.reverse()
       print('Path found: {}'.format(path))
       return path
```

```
open_set.remove(n)
     closed_set.add(n)
  print('Path does not exist!')
  return None
def get_neighbors(v):
  if v in Graph_nodes:
     return Graph_nodes[v]
  else:
     return None
def heuristic(n):
  H_dist = {
  'A': 11,
  'B': 6,
  'C': 99,
  'D': 1,
  'E': 7,
  'G': 0,
 }
  return H_dist[n]
Graph_nodes = {
  'A': [('B', 2), ('E', 3)],
  'B': [('C', 1),('G', 9)],
  'C': None,
  'E': [('D', 6)],
  'D': [('G', 1)],
 }
aStarAlgo('A', 'G')
```

2. AO STAR SEARCH

```
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): # starts a recursive AO* algorithm
    self.aoStar(self.start, False)
  def getNeighbors(self, v): # gets the Neighbors of a given node
    return self.graph.get(v,")
  def getStatus(self,v): # return the status of a given node
    return self.status.get(v,0)
  def setStatus(self,v, val): # set the status of a given node
    self.status[v]=val
  def getHeuristicNodeValue(self, n):
    return self.H.get(n,0) # always return the heuristic value of a given node
  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]=[]
    flag=True
    for nodeInfoTupleList in self.getNeighbors(v):
       cost=0
       nodeList=[]
      for c, weight in nodeInfoTupleList:
         cost=cost+self.getHeuristicNodeValue(c)+weight
```

```
if flag==True:
       minimumCost=cost
       costToChildNodeListDict[minimumCost]=nodeList
       flag=False
    else: # checking the Minimum Cost nodes with the current Minimum Cost
       if minimumCost>cost:
         minimumCost=cost
         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) \geq= 0: # if status node v \geq= 0, compute Minimum Cost nodes of v
    minimumCost, childNodeList = self.computeMinimumCostChildNodes(v)
    print(minimumCost, childNodeList)
    self.setHeuristicNodeValue(v, minimumCost)
    self.setStatus(v,len(childNodeList))
    solved=True # check the Minimum Cost nodes of v are solved
    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: # check the current call is not for backtracking
       for childNode in childNodeList:
         self.setStatus(childNode,0)
         self.aoStar(childNode, False)
```

nodeList.append(c)

```
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.CANDIDATE ELIMINATION: (NUMPY AND PANDAS)
import numpy as np
import pandas as pd
# Loading Data from a CSV File
data = pd.DataFrame(pd.read_csv('finds.csv'))
# Separating concept features from Target
concepts = np.array(data.iloc[:,0:-1])
# Isolating target into a separate DataFrame
target = np.array(data.iloc[:,-1])
def learn(concepts, target):
  specific_h=[0,0,0,0,0,0]
  print ('s0',specific_h)
  specific h = concepts[0].copy()
  print('s1',specific_h)
  general_h = [["?" for i in range(len(concepts[0]))] for j in range(len(concepts[0]))]
  print('g0',general h)
  for i, h in enumerate(concepts):
     if target[i] == "Yes":
       for x in range(len(h)): # Change values in S & G only if values change
```

```
if h[x] != specific_h[x]:
             specific_h[x] = '?'
             general_h[x][x] = '?'
     if target[i] == "No":
        for x in range(len(h)):
          if h[x] != specific_h[x]:
             general_h[x][x] = specific_h[x]
          else:
             general h[x][x] = '?'
     print(f"s{i}",specific_h)
     print(f"g{i}",general_h)
     print()
  # find indices where we have empty rows, meaning those that are unchanged
  indices = [i for i,val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']
  for i in indices:
     # remove those rows from general h
     general_h.remove(['?', '?', '?', '?', '?', '?'])
  # Return final values
  return specific_h,general_h
s_final, g_final = learn(concepts, target)
print('Final specific Hypothesis : ',s_final)
print('Final general Hypothesis :',g_final)
```

4. ID3 (NUMPY, PANDAS, MATH)

```
import numpy as np
import pandas as pd
import math
class Node:
  def init (self,I):
     self.label=l
     self.branch={}
#Calculayes Entropy of a dataset
def entropy(data):
  total_ex=len(data)
  p_ex=len(data.loc[data['play']=='yes'])
  n_ex=len(data.loc[data['play']=='no'])
  en=0
  if(p ex>0):
     en = -(p_ex/float(total_ex))*(math.log(p_ex,2)- math.log(total_ex,2))
  if(n_ex>0):
     en += -(n_ex/float(total_ex))*(math.log(n_ex,2)- math.log(total_ex,2))
  return en
#Calculates Gain of an attribute
def gain(data s,attrib):
  values=set(data_s[attrib])
  gain=entropy(data_s)
  for val in values:
     gain= gain -
len(data_s.loc[data_s[attrib]==val])/float(len(data_s))*entropy(data_s[data_s[attrib]==val])
  return gain
#Gets attribute with highest gain
def get attr(data):
  attribute=" "
  max gain=0
  for attr in data.columns[:-1]:
     g=gain(data,attr)
     if g>max_gain:
       max_gain=g
```

```
attribute =attr
```

```
#Constructs Decision Tree
def decision tree(data):
  root=Node("NULL")
  #If Entropy is 0, All data is Yes/No.
  if(entropy(data)==0):
     if(len(data.loc[data['play']=='yes']) == len(data)):
       root.label='yes'
     else:
       root.label='no'
     return root
  #If only one attribute is left, Tree is complete
  if(len(data.columns)==1):
     return
  else:
     #Get the attribute with highest gain.
     attr=get_attr(data)
     root.label=attr
     values=set(data[attr])
     for value in values:
       root.branch[value]=decision tree(data.loc[data[attr]==value].drop(attr,axis=1))
     return root
def test(tree,test_str):
  if not tree.branch:
     return tree.label
  return test(tree.branch[str(test_str[tree.label])],test_str)
data=pd.read_csv('playtennis.csv')
print("Number of Records : ",len(data))
tree=decision_tree(data)
test str={
"outlook": "Sunny",
"temperature": "Hot",
"humidity": "high",
"wind": "Weak",
```

```
print(test(tree,test_str))
5. BACK PROPAGATION: (NUMPY, TIME, MATPLOTLIB. PYPLOT)
import numpy as np
import time
import matplotlib.pyplot as plt
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
Ir=0.1
inputlayer neurons=2
hiddenlayer neurons=3
output_neurons=1
def sigmoid(x):
  return 1/(1+np.exp(-x))
def derivatives_sigmoid(x):
  return x^*(1-x)
A=[]
B=[]
for i in range(5):
  epoch=700*i
  A.append(epoch)
  hidden_weight=np.random.uniform(size=(inputlayer_neurons,hiddenlayer_neurons))
  hidden_bias=np.random.uniform(size=(1,hiddenlayer_neurons))
  output weight=np.random.uniform(size=(hiddenlayer neurons,output neurons))
  output_bias=np.random.uniform(size=(1,output_neurons))
  start time=time.time()
  for i in range(epoch):
    hidden_input=np.dot(X,hidden_weight)+hidden_bias
```

```
hidden output=sigmoid(hidden input)
     output input=np.dot(hidden output,output weight)+output bias
     output output=sigmoid(output input)
     Output Error=y-output output
     Output Difference=Output Error*derivatives sigmoid(output output)
     Hidden Error=Output Difference.dot(output weight.T)
     Hidden Difference=Hidden Error*derivatives sigmoid(hidden output)
     output weight=output weight+hidden output.T.dot(Output Difference)*Ir
     output bias=output bias+np.sum(Output Difference,axis=0,keepdims=True)*Ir
     hidden weight=hidden weight+X.T.dot(Hidden Difference)*Ir
     hidden_bias=hidden_bias+np.sum(Hidden_Difference,axis=0,keepdims=True)*Ir
  B.append((time.time() - start time))
plt.plot(A, B)
plt.xlabel('x - axis')
plt.ylabel('y - axis')
plt.title('My first graph!')
plt.show()
print("input:\n" + str(X))
print("Actual output:\n" + str(y))
print("Predicted output \n"+str(output output))
```

6. NAIVE BAYESIAN CLASSIFIER: (MATH, NUMPY, PANDAS)

NOTE: LOADING OF CSV IS DONE USING PANDAS. SO,1st ROW OF CSV IS IGNORED. TAKE CARE IN DATASET(JUST COPY AND PASTE THE 1st ROW in the LAST ROW/INSERT A NEW ROW AT BEGINNING)

```
import math
import numpy as np
import pandas as pd
#Creates a dictionary where the keys are the target values
#and classifies the records based on the target value.
#Appending the whole record except the last column(The target value)
def separateByClass(dataset):
  separate={}
  for i in range(len(dataset)):
    if(dataset[i][-1] not in separate):
       separate[dataset[i][-1]]=[]
     separate[dataset[i][-1]].append(dataset[i][0:-1])
  return separate
def mean(numbers):
  return sum(numbers)/float(len(numbers))
def stdev(numbers):
  avg=mean(numbers)
  varience=sum([math.pow(x-avg,2) for x in numbers])/float(len(numbers)-1)
  return math.sqrt(varience)
#Takes in a set of class records, Here, Its "1.0" & "0.0".
#Summarizes the mean and stddev of each attribute for those set of class records.
def getclassdetails(class records):
  mean_std=[(mean(attribute),stdev(attribute)) for attribute in zip(*class_records)]
  return mean std
```

#Here,We are creating another dictionary after segregating the records

#The values for each key is the set of mean and stdev instead of set of records.

```
def summarizeByClass(dataset):
  separated=separateByClass(dataset)
  summaries={}
  #items returns key, value pair in the dictionary.
  for classval, class records in separated.items():
     summaries[classval]=getclassdetails(class records)
  return summaries
#Calculates the probabilty given an attribute x
def cal_attr_probability(mean,stdev,x):
  expo=math.exp(-(math.pow(x-mean,2)/(2*math.pow(stdev,2))))
  result=(1/(math.sqrt(2*math.pi)*stdev))*expo
  return result
#Calculates the probabilty of all classes(Target value) for
#one record based on summaries
def calprobability(summaries,record):
  probability={}
  #items returns key, value pair in the dictionary
  for classval, class_det in summaries.items():
     probability[classval]=1
     for i in range(len(class_det)):
       mean,stdev=class_det[i]
       attr=record[i]
       probability[classval]*=cal attr probability(mean,stdev,attr)
  return probability
#Takes in one record and classifies it
def predict_this(summaries,record):
  probability=calprobability(summaries,record)
  bestlabel, bestprob=None, -1
  #Finding the classval with the most probabilty
  for classval, prob in probability.items():
     if bestlabel is None or prob>bestprob:
```

```
bestprob=prob
bestlabel=classval
return bestlabel
```

```
#The real main function. Takes in testset, Returns Classfied class values array
def prediction(summaries,testset):
  predict=[]
  for i in range(len(testset)):
     result=predict_this(summaries,testset[i])
     predict.append(result)
  return predict
# Finds accuracy of prediction
def getaccuracy(testset,prediction):
  correct=0
  for i in range(len(testset)):
     if testset[i][-1]==prediction[i]:
       correct+=1
  return (float(correct)/float(len(testset)))*100.0
def main():
  trainingset=np.array(pd.DataFrame(pd.read_csv('NaiveBayesDiabetes.csv')),float)
  testset=np.array(pd.DataFrame(pd.read_csv('NaiveBayesDiabetes1.csv')),float)
  print("records in training dataset={0} and test dataset={1}
rows".format(len(trainingset),len(testset)))
  summaries=summarizeByClass(trainingset)
  predict=prediction(summaries,testset)
  print(predict)
  accuracy=getaccuracy(testset,predict)
  print(accuracy)
main()
```

7. EM AND KMEANS: (PANDAS, NUMPY, MATPLOTLIB. PYPLOT, 4 SKLEARN)

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn import preprocessing
from sklearn.cluster import KMeans
from sklearn.mixture import GaussianMixture
# import some data to play with
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']
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.show()
# K MEANS MODEL
model = KMeans(n_clusters=3)
model.fit(X)
```

Plot the Models Classifications

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')
plt.show()
# GMM MODEL
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
gmm = GaussianMixture(n_components=3)
gmm.fit(xs)
gmm_y = gmm.predict(xs)
# Plot the Models Classifications
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')
plt.show()
print('Observation: The GMM using EM algorithm based clustering matched the true labels more
closely than the Kmeans.')
8. K Nearest Neighbors : (4 SKLEARN)
from sklearn import datasets
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report, confusion_matrix
# Load dataset
iris=datasets.load iris()
print("Iris Data set loaded...")
```

```
# 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 testing 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
model = KNeighborsClassifier(n_neighbors=1)
# Perform Training
model.fit(x_train, y_train)
# Perform testing
y_pred=model.predict(x_test)
# Display the results
print("Results of Classification using K-nn with K=1")
for i in range(len(x_test)):
  print(" Sample:", str(x_test[i]), " Actual-label:", str(y_test[i]), " Predicted-label:", str(y_pred[i]))
print("Classification Accuracy :", model.score(x_test,y_test));
print('Confusion Matrix :\n',confusion_matrix(y_test,y_pred))
print('Accuracy Metrics : \n',classification_report(y_test,y_pred))
9 . Locally weighted Regression : (MATPLOTLIB.PYPLOT,PANDAS,NUMPY)
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
```

```
def kernel(point,xmat, k):
  m,n = np.shape(xmat)
  weights = np.mat(np.eye((m))) # eye - identity matrix
  for j in range(m):
     diff = point - X[j]
     weights[j,j] = np.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 = np.shape(xmat)
  ypred = np.zeros(m)
  for i in range(m):
     ypred[i] = xmat[i]*localWeight(xmat[i],xmat,ymat,k)
  return ypred
def graphPlot(X,ypred):
  #Argsort sorts the indices in ascending order of the values and returns the indices
  sortindex = X[:,1].argsort(0)
  xsort = X[sortindex][:,0]
  fig = plt.figure()
  ax = fig.add_subplot(1,1,1)
  #Plot clusters
  ax.scatter(bill,tip, color='green')
  #Plot the line
  ax.plot(xsort[:,1],ypred[sortindex], color = 'red', linewidth=5)
  plt.xlabel('Total bill')
  plt.ylabel('Tip')
  plt.show();
# load data points
data = pd.read_csv('program9_dataset.csv')
```

```
# We use only Bill amount and Tips data
bill = np.array(data.total_bill)
tip = np.array(data.tip)

# .mat will convert nd array to 2D array
mbill = np.mat(bill)
mtip = np.mat(tip)

#Getting number of records as m
m= len(bill)

#Creating a identity matrix of order m
one = np.mat(np.ones(m))

#Combining the Identity matrix with Bill matrix and craeating a matrix of mx2
X = np.hstack((one.T,mbill.T)) # 244 rows, 2 cols

# increase k to get smooth curves
ypred = localWeightRegression(X,mtip,3)
graphPlot(X,ypred)
```