```
PROGRAM 1:
#!/usr/bin/env python
# coding: utf-8
# In[1]:
def aStarAlgo(start node, stop node):
        open set = set(start node)
        closed set = set()
        g = {} #store distance from starting node
        parents = {}# parents contains an adjacency map of all nodes
        #ditance of starting node from itself is zero
        g[start node] = 0
        #start node is root node i.e it has no parent nodes
        #so start node is set to its own parent node
        parents[start node] = start node
        while len(open set) > 0:
            n = None
            #node with lowest f() is found
            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):
                   #nodes 'm' not in first and last set are added to first
                    #n is set its parent
                    if m not in open_set and m not in closed set:
                        open set.add(m)
                        parents[m] = n
                        g[m] = g[n] + weight
                    #for each node m, compare its distance from start i.e g(m) to
the
                    #from start through n node
                    else:
                        if g[m] > g[n] + weight:
                            #update g(m)
                            g[m] = g[n] + weight
                             #change parent of m to n
                            parents[m] = n
                             #if m in closed set, remove and add to open
                            if m in closed set:
                                closed set.remove(m)
                                open set.add(m)
            if n == None:
                print('Path does not exist!')
                return None
            # if the current node is the stop node
            # then we begin reconstructin the path from it to the start node
            if n == stop node:
                path = []
```

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while parents[n] != n:
                     path.append(n)
                     n = parents[n]
                 path.append(start node)
                path.reverse()
                print('Path found: {}'.format(path))
                 return path
            # remove n from the open list, and add it to closed list
            # because all of his neighbors were inspected
            open set.remove(n)
            closed set.add(n)
        print('Path does not exist!')
        return None
#define fuction to return neighbor and its distance
#from the passed node
def get neighbors(v):
    if v in Graph nodes:
        return Graph nodes[v]
    else:
        return None
#for simplicity we ll consider heuristic distances given
#and this function returns heuristic distance for all nodes
def heuristic(n):
        H dist = {
            'A': 10,
             'B': 8,
             'C': 5,
             'D': 7,
             'E': 3,
             'F': 6,
             'G': 5,
            'H': 3,
            'I': 1,
             'J': 0
        }
        return H_dist[n]
#Describe your graph here
Graph nodes = {
    'A': [('B', 6), ('F', 3)],
    'B': [('C', 3), ('D', 2)],
    'C': [('D', 1), ('E', 5)],
    'D': [('C', 1), ('E', 8)],
    'E': [('I', 5), ('J', 5)], 
'F': [('G', 1),('H', 7)],
    'G': [('I', 3)],
    'H': [('I', 2)],
    'I': [('E', 5), ('J', 3)],
aStarAlgo('A', 'J')
PROGRAM 2:
# Recursive implementation of AO* aglorithm by Dr. K PARAMESHA, Professor, VVCE,
Mysuru, INDIA
class Graph:
```

```
def init (self, graph, heuristicNodeList, startNode): #instantiate graph
object with graph topology, heuristic values, start node
       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,'')
   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
                               # set the revised heuristic value of a given
node
   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): # Computes the Minimum Cost of
child nodes of a given node v
       minimumCost=0
       costToChildNodeListDict={}
       costToChildNodeListDict[minimumCost] = []
       flag=True
       for nodeInfoTupleList in self.getNeighbors(v): # iterate over all the set
of child node/s
          cost=0
          nodeList=[]
          for c, weight in nodeInfoTupleList:
              cost=cost+self.getHeuristicNodeValue(c)+weight
              nodeList.append(c)
          if flag==True:
                                           # initialize Minimum Cost with the
cost of first set of child node/s
              minimumCost=cost
              costToChildNodeListDict[minimumCost]=nodeList
                                                           # set the
Minimum Cost child node/s
              flag=False
                                           # checking the Minimum Cost nodes
          else:
with the current Minimum Cost
              if minimumCost>cost:
                 minimumCost=cost
                 costToChildNodeListDict[minimumCost]=nodeList # set the
Minimum Cost child node/s
```

```
return minimumCost, costToChildNodeListDict[minimumCost] # return Minimum
Cost and Minimum Cost child node/s
```

```
def aoStar(self, v, backTracking): # AO* algorithm for a start node and
backTracking status flag
       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)
           self.setHeuristicNodeValue(v, minimumCost)
           self.setStatus(v,len(childNodeList))
                                         # check the Minimum Cost nodes of v are
           solved=True
solved
           for childNode in childNodeList:
               self.parent[childNode]=v
               if self.getStatus(childNode)!=-1:
                   solved=solved & False
           if solved==True:
                                       # if the Minimum Cost nodes of v are
solved, set the current node status as solved(-1)
               self.setStatus(v,-1)
               self.solutionGraph[v]=childNodeList # update the solution graph
with the solved nodes which may be a part of solution
           if v!=self.start:
                                       # check the current node is the start node
for backtracking the current node value
               self.aoStar(self.parent[v], True) # backtracking the current node
value with backtracking status set to true
           if backTracking==False: # check the current call is not for
backtracking
               for childNode in childNodeList: # for each Minimum Cost child
node
                   self.setStatus(childNode,0) # set the status of child node to
0 (needs exploration)
                   self.aoStar(childNode, False) # Minimum Cost child node is
further explored with backtracking status as false
h1 = {'A': 1, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J':
1, 'T': 3}
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()
h2 = {'A': 1, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7} #
Heuristic values of Nodes
                                                 # Graph of Nodes and Edges
graph2 = {
    'A': [[('B', 1), ('C', 1)], [('D', 1)]], # Neighbors of Node 'A', B, C & D
with repective weights
```

```
'B': [[('G', 1)], [('H', 1)]],  # Neighbors are included in a
list of lists
    'D': [[('E', 1), ('F', 1)]]  # Each sublist indicate a "OR"
node or "AND" nodes
}

G2 = Graph(graph2, h2, 'A')  # Instantiate Graph object with
graph, heuristic values and start Node
G2.applyAOStar()  # Run the AO* algorithm
G2.printSolution()  # Print the solution graph as
output of the AO* algorithm search
```

PROGRAM 3:

```
import random
import csv
def g_0(n):
    return ("?",)*n
def s 0(n):
    return ('\Phi',)*n
def more general(h1, h2):
    more general parts = []
    for x, y in zip(h1, h2):
        mg = x == "?" \text{ or } (x != "\Phi" \text{ and } (x == y \text{ or } y == "\Phi"))
        more general parts.append(mg)
    return all (more general parts)
def fulfills (example, hypothesis):
    ### the implementation is the same as for hypotheses:
    return more_general(hypothesis, example)
def min generalizations (h, x):
    h new = list(h)
    for i in range(len(h)):
        if not fulfills(x[i:i+1], h[i:i+1]):
             h \text{ new}[i] = '?' \text{ if } h[i] != '\Phi' \text{ else } x[i]
    return [tuple(h new)]
def min specializations(h, domains, x):
    results = []
    for i in range(len(h)):
        if h[i] == "?":
             for val in domains[i]:
                 if x[i] != val:
                     h new = h[:i] + (val,) + h[i+1:]
                     results.append(h new)
        elif h[i] != "\Phi":
             h \text{ new} = h[:i] + ('\Phi',) + h[i+1:]
             results.append(h new)
    return results
with open('trainingexamples.csv') as csvFile:
        examples = [tuple(line) for line in csv.reader(csvFile)]
def get_domains(examples):
    d = [set() for i in examples[0]]
    for x in examples:
        for i, xi in enumerate(x):
             d[i].add(xi)
    return [list(sorted(x)) for x in d]
get domains(examples)
def candidate elimination (examples):
    domains = get domains(examples)[:-1]
    G = set([g 0(len(domains))])
    S = set([s 0(len(domains))])
```

```
i = 0
    print("\n G[{0}]:".format(i), G)
    print("\n S[{0}]:".format(i), S)
    for xcx in examples:
        i = i + 1
        x, cx = xcx[:-1], xcx[-1] # Splitting data into attributes and decisions
        if cx == 'Y': # x is positive example
            G = \{g \text{ for } g \text{ in } G \text{ if fulfills}(x, g)\}
            S = generalize S(x, G, S)
        else: # x is negative example
            S = \{s \text{ for } s \text{ in } S \text{ if not fulfills}(x, s)\}
            G = specialize_G(x, domains, G, S)
        print("\n G[{0}]:".format(i), G)
        print("\n S[{0}]:".format(i), S)
def generalize S(x, G, S):
    S prev = list(S)
    for s in S_prev:
        if s not in S:
            continue
        if not fulfills (x, s):
            S.remove(s)
            Splus = min generalizations(s, x)
            \#\# keep only generalizations that have a counterpart in G
            S.update([h for h in Splus if any([more general(g,h)
                                                 for g in G])])
            ## remove hypotheses less specific than any other in S
            S.difference update([h for h in S if
                                   any([more general(h, h1)
                                        for h1 in S if h != h1])])
    return S
def specialize G(x, domains, G, S):
    G prev = list(G)
    for g in G_prev:
        if g not in G:
            continue
        if fulfills (x, g):
            G.remove(g)
            Gminus = min specializations(g, domains, x)
            ## keep only specializations that have a conuterpart in S
            G.update([h for h in Gminus if any([more general(h, s)
                                                  for s in S])])
            ## remove hypotheses less general than any other in G
            G.difference update([h for h in G if
                                   any([more_general(g1, h)
                                        for g1 in G if h != g1])])
    return G
candidate elimination(examples)
PROGRAM 4:
# -*- coding: utf-8 -*-
"""ID3 2.ipynb
Automatically generated by Colaboratory.
Original file is located at
    https://colab.research.google.com/drive/1amD6hMNBxoXXebhuF70NJfNnQu18n9QZ
import pandas as pd
from pandas import DataFrame
```

df_tennis = pd.read_csv('tennis.csv')

#df tennis.columns[0]

print("\n Given Play Tennis Data Set:\n\n",df tennis)

```
df tennis.keys()[0]
#Function to calculate the entropy of probaility of observations
# -p*log2*p
def entropy(probs):
    import math
    return sum( [-prob*math.log(prob, 2) for prob in probs] )
#Function to calulate the entropy of the given Data Sets/List with respect to
target attributes
def entropy_of_list(a_list):
    from collections import Counter
    cnt = Counter(x for x in a list)
                                       # Counter calculates the propotion of class
    num instances = len(a list)*1.0 \# = 14
    print("\n Number of Instances of the Current Sub Class is
{0}:".format(num instances ))
    probs = [x / num instances for x in cnt.values()] # x means no of YES/NO
    print("\n Classes:", min(cnt), max(cnt))
    print(" \n Probabilities of Class {0} is {1}:".format(min(cnt), min(probs)))
    print(" \ n \ Probabilities of Class {0} is {1}:".format(max(cnt), max(probs)))
    return entropy(probs) # Call Entropy :
    # The initial entropy of the YES/NO attribute for our dataset.
print("\n INPUT DATA SET FOR ENTROPY CALCULATION:\n", df tennis['PlayTennis'])
total_entropy = entropy_of_list(df_tennis['PlayTennis'])
print("\n Total Entropy of PlayTennis Data Set:",total entropy)
def information gain(df, split attribute name, target attribute name, trace=0):
    print("Information Gain Calculation of ", split attribute name)
    Takes a DataFrame of attributes, and quantifies the entropy of a target
    attribute after performing a split along the values of another attribute.
    1 1 1
    # Split Data by Possible Vals of Attribute:
    df split = df.groupby(split attribute name)
    for name, group in df_split:
            print("Name:\n", name)
           print("Group:\n", group)
    # Calculate Entropy for Target Attribute, as well as
    # Proportion of Obs in Each Data-Split
    nobs = len(df.index) * 1.0
    print("NOBS", nobs)
    df_agg_ent = df_split.agg({target_attribute_name : [entropy_of_list, lambda x:
len(x)/nobs] })[target attribute name]
    print([target attribute name])
    print(" Entropy List ", entropy of list)
    print("DFAGGENT", df agg ent)
    df agg ent.columns = ['Entropy', 'PropObservations']
    if trace: # helps understand what fxn is doing:
        print(df_agg_ent)
    # Calculate Information Gain:
    new entropy = sum( df agg ent['Entropy'] * df agg ent['PropObservations'] )
    old entropy = entropy of list(df[target attribute name])
    return old_entropy - new_entropy
print('Info-gain for Outlook is : '+str( information gain(df tennis, 'Outlook',
'PlayTennis')),"\n")
print('\n Info-gain for Humidity is: ' + str( information gain(df tennis,
'Humidity', 'PlayTennis')),"\n")
print('\n Info-gain for Wind is:' + str( information gain(df tennis, 'Wind',
'PlayTennis')),"\n")
print('\n Info-gain for Temperature is:' + str( information gain(df tennis,
'Temperature','PlayTennis')),"\n")
```

```
def id3(df, target attribute name, attribute names, default class=None):
    ## Tally target attribute:
    from collections import Counter
    cnt = Counter(x for x in df[target attribute name])# class of YES /NO
    ## First check: Is this split of the dataset homogeneous?
    if len(cnt) == 1:
        return next(iter(cnt)) # next input data set, or raises StopIteration when
EOF is hit.
    ## Second check: Is this split of the dataset empty?
    # if yes, return a default value
    elif df.empty or (not attribute names):
        return default class # Return None for Empty Data Set
    ## Otherwise: This dataset is ready to be devied up!
        # Get Default Value for next recursive call of this function:
        default class = max(cnt.keys()) #No of YES and NO Class
        # Compute the Information Gain of the attributes:
        gainz = [information gain(df, attr, target attribute name) for attr in
attribute names] #
        index of max = gainz.index(max(gainz)) # Index of Best Attribute
        # Choose Best Attribute to split on:
        best attr = attribute names[index of max]
        \# Create an empty tree, to be populated in a moment
        tree = {best attr:{}} # Iniiate the tree with best attribute as a node
        remaining attribute names = [i for i in attribute names if i != best attr]
        # Split dataset
        # On each split, recursively call this algorithm.
        # populate the empty tree with subtrees, which
        # are the result of the recursive call
        for attr val, data subset in df.groupby(best attr):
            \overline{\text{subtree}} = id3(\overline{\text{data subset}},
                        target attribute name,
                        remaining attribute names,
                        default class)
            tree[best_attr][attr_val] = subtree
        return tree
# Get Predictor Names (all but 'class')
attribute_names = list(df_tennis.columns)
print("List of Attributes:", attribute names)
attribute names.remove('PlayTennis') #Remove the class attribute
print("Predicting Attributes:", attribute names)
# Run Algorithm:
from pprint import pprint
tree = id3(df tennis, 'PlayTennis', attribute names)
print("\n\nThe Resultant Decision Tree is :\n")
#print(tree)
pprint(tree)
attribute = next(iter(tree))
print("Best Attribute :\n",attribute)
print("Tree Keys:\n", tree[attribute].keys())
def classify(instance, tree, default=None): # Instance of Play Tennis with
Predicted
    print("Instance:",instance)
    attribute = next(iter(tree)) # Outlook/Humidity/Wind
    print("Attribute:",attribute) # [Key /Attribute Both are same ]
    # print("Insance of Attribute :",instance[attribute],attribute)
```

```
if instance[attribute] in tree[attribute].keys(): # Value of the attributs in
set of Tree keys
        result = tree[attribute][instance[attribute]]
        print("Instance Attribute:",instance[attribute],"TreeKeys
:", tree[attribute].keys())
        if isinstance (result, dict): # this is a tree, delve deeper
            return classify(instance, result)
        else:
            return result # this is a label
    else:
        return default
df_tennis['predicted'] = df_tennis.apply(classify, axis=1, args=(tree,'No') )
    # classify func allows for a default arg: when tree doesn't have answer for a
particular
    # combitation of attribute-values, we can use 'no' as the default guess
df tennis[['PlayTennis', 'predicted']]
training data = df tennis.iloc[1:-4] # all but last four instances
test data = df tennis.iloc[-4:] # just the last four
train tree = id3(training_data, 'PlayTennis', attribute_names)
test data['predicted2'] = test data.apply(
                                                                            # <----
test data source
                                           classify,
                                           axis=1,
                                           args=(train tree, 'Yes') ) # <----</pre>
train data tree
print ('\n\n Accuracy is : ' + str(
sum(test data['PlayTennis'] == test data['predicted2'] ) / (1.0*len(test data.index))
) )
```

PROGRAM 5:

```
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) \# 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=5000
             #Setting training iterations
lr=0.1
              #Setting learning rate
                                    #number of features in data set
inputlayer neurons = 2
                           #number of hidden layers neurons
hiddenlayer neurons = 3
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)
#how much hidden layer wts contributed to error
    hiddengrad = derivatives sigmoid(hlayer act)
    d hiddenlayer = EH * hiddengrad
# dotproduct of nextlayererror and currentlayerop
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("Predicted Output: \n" ,output)
```

PROGRAM 6:

```
#Write a program to implement the naïve Bayesian classifier for a sample training
#stored as a .CSV file. Compute the accuracy of the classifier, considering few
test data sets.
print("\nNaive Bayes Classifier for concept learning problem")
import csv
import random
import math
import operator
def safe div(x,y):
if y == 0:
 return 0
return x/y
# 1.Data Handling
# 1.1 Loading the Data from csv file of ConceptLearning dataset.
def loadCsv(filename):
  lines = csv.reader(open(filename))
  dataset = list(lines)
  for i in range(len(dataset)):
    dataset[i] = [float(x) for x in dataset[i]]
  return dataset
#1.2 Splitting the Data set into Training Set
def splitDataset(dataset, splitRatio):
 trainSize = int(len(dataset) * splitRatio)
 trainSet = []
 copy = list(dataset)
  i=0
 while len(trainSet) < trainSize:</pre>
  #index = random.randrange(len(copy))
    trainSet.append(copy.pop(i))
  return [trainSet, copy]
```

```
#2.Summarize Data
#The naive bayes model is comprised of a
#summary of the data in the training dataset.
#This summary is then used when making predictions.
#involves the mean and the standard deviation for each attribute, by class value
#2.1: Separate Data By Class
#Function to categorize the dataset in terms of classes
#The function assumes that the last attribute (-1) is the class value.
#The function returns a map of class values to lists of data instances.
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
#The mean is the central middle or central tendency of the data,
# and we will use it as the middle of our gaussian distribution
# when calculating probabilities
#2.2 : Calculate Mean
def mean (numbers):
  return safe div(sum(numbers), float(len(numbers)))
#The standard deviation describes the variation of spread of the data,
#and we will use it to characterize the expected spread of each attribute
#in our Gaussian distribution when calculating probabilities.
#2.3 : Calculate Standard Deviation
def stdev(numbers):
  avg = mean(numbers)
  variance = safe div(sum([pow(x-avg,2) for x in numbers]),float(len(numbers)-1))
 return math.sqrt(variance)
#2.4 : Summarize Dataset
#Summarize Data Set for a list of instances (for a class value)
#The zip function groups the values for each attribute across our data instances
#into their own lists so that we can compute the mean and standard deviation values
#for the attribute.
def summarize(dataset):
 summaries = [(mean(attribute), stdev(attribute)) for attribute in zip(*dataset)]
  del summaries[-1]
  return summaries
#2.5 : Summarize Attributes By Class
#We can pull it all together by first separating our training dataset into
#instances grouped by class. Then calculate the summaries for each attribute.
def summarizeByClass(dataset):
  separated = separateByClass(dataset)
  summaries = {}
  for classValue, instances in separated.items():
    summaries[classValue] = summarize(instances)
  print("Summarize Attributes By Class")
  print(summaries)
  print(" ")
  return summaries
#3.Make Prediction
#3.1 Calculate Probaility Density Function
def calculateProbability(x, mean, stdev):
```

```
exponent = math.exp(-safe div(math.pow(x-mean,2),(2*math.pow(stdev,2))))
  final = safe div(1 , (math.sqrt(2*math.pi) * stdev)) * exponent
 return final
#3.2 Calculate Class Probabilities
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
#3.3 Prediction : look for the largest probability and return the associated class
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
#4.Make Predictions
# Function which return predictions for list of predictions
# For each instance
def getPredictions(summaries, testSet):
 predictions = []
  for i in range(len(testSet)):
   result = predict(summaries, testSet[i])
   predictions.append(result)
 return predictions
#5. Computing Accuracy
def getAccuracy(testSet, predictions):
 correct = 0
  for i in range(len(testSet)):
    if testSet[i][-1] == predictions[i]:
      correct += 1
 accuracy = safe div(correct,float(len(testSet))) * 100.0
 return accuracy
def main():
 filename = 'diabetes2.csv'
 splitRatio = 0.9
 dataset = loadCsv(filename)
 trainingSet, testSet = splitDataset(dataset, splitRatio)
 print('Split {0} rows into'.format(len(dataset)))
 print('Number of Training data: ' + (repr(len(trainingSet))))
 print('Number of Test Data: ' + (repr(len(testSet))))
 print("\nThe values assumed for the concept learning attributes are\n")
 print("OUTLOOK=> Sunny=1 Overcast=2 Rain=3\nTEMPERATURE=> Hot=1 Mild=2
Cool=3\nHUMIDITY=> High=1 Normal=2\nWIND=> Weak=1 Strong=2")
 print("TARGET CONCEPT:PLAY TENNIS=> Yes=10 No=5")
 print("\nThe Training set are:")
 for x in trainingSet:
   print(x)
   print("\nThe Test data set are:")
  for x in testSet:
   print(x)
 print("\n")
# prepare model
  summaries = summarizeByClass(trainingSet)
```

```
# test model
  predictions = getPredictions(summaries, testSet)
  actual = []
  for i in range(len(testSet)):
    vector = testSet[i]
    actual.append(vector[-1])

# Since there are five attribute values, each attribute constitutes to 20%
  accuracy. So if all attributes
#match with predictions then 100% accuracy
    print('Actual values: {0}%'.format(actual))
    print('Predictions: {0}%'.format(predictions))
    accuracy = getAccuracy(testSet, predictions)
    print('Accuracy: {0}%'.format(accuracy))
main()
```

```
PROGRAM 7:
#Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same
#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
from sklearn.cluster import KMeans
import sklearn.metrics as sm
import pandas as pd
import numpy as np
11 = [0, 1, 2]
def rename(s):
       12 = []
       for i in s:
              if i not in 12:
                      12.append(i)
       for i in range(len(s)):
              pos = 12.index(s[i])
               s[i] = 11[pos]
       return s
# import some data to play with
iris = datasets.load iris()
print("\n IRIS DATA :", iris.data);
print("\n IRIS FEATURES :\n",iris.feature names)
print("\n IRIS TARGET :\n",iris.target)
print("\n IRIS TARGET NAMES:\n",iris.target names)
# Store the inputs as a Pandas Dataframe and set the column names
X = pd.DataFrame(iris.data)
X.columns = ['Sepal Length','Sepal Width','Petal Length','Petal Width']
#print(X.columns) #print("X:",x)
#print("Y:",y)
```

```
y = pd.DataFrame(iris.target)
y.columns = ['Targets']
# Set the size of the plot
plt.figure(figsize=(14,7))
# Create a colormap
colormap = np.array(['red', 'lime', 'black'])
# Plot Sepal
plt.subplot(1,2,1)
plt.scatter(X.Sepal_Length, X.Sepal Width, c=colormap[y.Targets], s=40)
plt.title('Sepal')
plt.subplot(1,2,2)
plt.scatter(X.Petal Length, X.Petal Width, c=colormap[y.Targets], s=40)
plt.title('Petal')
plt.show()
print("Actual Target is:\n", iris.target)
# K Means Cluster
model = KMeans(n clusters=3)
model.fit(X)
# Set the size of the plot
plt.figure(figsize=(14,7))
# Create a colormap
colormap = np.array(['red', 'lime', 'black'])
# Plot the Original Classifications
plt.subplot(1,2,1)
plt.scatter(X.Petal Length, X.Petal Width, c=colormap[y.Targets], s=40)
plt.title('Real Classification')
# Plot the Models Classifications
plt.subplot(1,2,2)
plt.scatter(X.Petal Length, X.Petal Width, c=colormap[model.labels], s=40)
plt.title('K Mean Classification')
plt.show()
km = rename(model.labels)
print("\nWhat KMeans thought: \n", km)
print("Accuracy of KMeans is ",sm.accuracy_score(y, km))
print("Confusion Matrix for KMeans is \n", sm.confusion matrix(y, km))
#The GaussianMixture scikit-learn class can be used to model this problem
#and estimate the parameters of the distributions using the expectation-
maximization algorithm.
from sklearn import preprocessing
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
print("\n", xs.sample(5))
from sklearn.mixture import GaussianMixture
gmm = GaussianMixture(n components=3)
gmm.fit(xs)
y cluster gmm = gmm.predict(xs)
plt.subplot(1, 2, 1)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y_cluster_gmm], s=40)
plt.title('GMM Classification')
```

```
plt.show()

em = rename(y_cluster_gmm)
print("\nWhat EM thought: \n", em)
print("Accuracy of EM is ",sm.accuracy_score(y, em))
print("Confusion Matrix for EM is \n", sm.confusion matrix(y, em))
```

PROGRAM 8:

```
# Python program to demonstrate # KNN classification algorithm # on IRISdataset
#Write a program to implement k-Nearest Neighbour algorithm to classify the iris
#Print both correct and wrong predictions. Java/Python ML library classes can be
used for
#this problem.
#import the dataset and library files
from sklearn.datasets import load iris
from sklearn.neighbors import KNeighborsClassifier
import numpy as np
from sklearn.model selection import train test split
iris dataset=load iris()
#display the iris dataset
print("\n IRIS FEATURES \ TARGET NAMES: \n ", iris dataset.target names)
for i in range(len(iris dataset.target names)):
    print("\n[{0}]:[{1}]".format(i,iris_dataset.target_names[i]))
print("\n IRIS DATA :\n",iris_dataset["data"])
#split the data into training and testing data
X train, X test, y train, y test = train test split(iris dataset["data"],
iris dataset["target"], random state=0)
print("\n Target :\n",iris dataset["target"])
print("\n X TRAIN \n", X_train)
print("\n X TEST \n", X_test)
print("\n Y TRAIN \n", y_train)
print("\n Y TEST \n", y test)
#train and fit the model
kn = KNeighborsClassifier(n neighbors=5)
kn.fit(X train, y train)
#predicting from model
x new = np.array([[5, 2.9, 1, 0.2]])
print("\n XNEW \n", x new)
prediction = kn.predict(x new)
print("\n Predicted target value: {}\n".format(prediction))
print("\n Predicted feature name:
{}\n".format(iris_dataset["target_names"][prediction]))
i=1
x= X test[i]
x new = np.array([x])
print("\n XNEW \n", x new)
for i in range(len(X_test)):
  x = X \text{ test[i]}
  x new = np.array([x])
  prediction = kn.predict(x new)
  print("\n Actual : {0} {1}, Predicted
:{2}{3}".format(y test[i],iris dataset["target names"][y test[i]],prediction,iris d
ataset["target names"][ prediction]))
```

PROGRAM 9:

```
import numpy as np
import matplotlib.pyplot as plt
def local regression(x0, X, Y, tau):
    x0 = [1, x0]
    X = [[1, i] \text{ for } i \text{ in } X]
   X = np.asarray(X)
    xw = (X.T) * np.exp(np.sum((X - x0) ** 2, axis=1) / (-2 * tau))
    beta = np.linalg.pinv(xw @ X) @ xw @ Y @ x0
   return beta
def draw(tau):
    prediction = [local_regression(x0, X, Y, tau) for x0 in domain]
    plt.plot(X, Y, 'o', color='black')
    plt.plot(domain, prediction, color='red')
    plt.show()
X = np.linspace(-3, 3, num=1000)
domain = X
Y = np.log(np.abs(X ** 2 - 1) + .5)
draw(10)
draw(0.1)
draw(0.01)
draw(0.001)
```