# Artificial Intelligence And Machine Learning Laboratory Manual (18CSL76)

#### Program-1

Implement A\* Search Algorithm

#### Algorithm:

- 1. Place the starting node list in "OPEN" list.
- 2. Check if the OPEN list is empty or not

If list is empty then,

return failure and stop

3. Select the node from "OPEN" set which has the smallest value of evaluation f.

If node n is goal node then,

Return success and stop

- 4. Expand node n and generate all of its successors and put n to the CLOSED list
- For each successor 'n' check whether its already there in OPEN and CLOSED list.
- ➤ If not compute f value and place into OPEN

Else if node n is already there in OPEN and CLOSE then it should be attached to back pointer which should reflect lowest g(n) value.

5. Return Step 2.

#### **Program:**

```
#!/usr/bin/env python
#coding utf-8
#ln[1]:

def aStarAlgo(start_node,stop_node):
   open_set=set(start_node)
   closed_set=set()
   g={}
```

```
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 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==None:
    print("path does not exist")
    return None
  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 doesnot exist")
  return None
def get_neighbors(v):
  if v in Graph_nodes:
     return Graph_nodes[v]
  else:
     return None
def heuristic(n):
  H_dist={
     'S':7,
     'A':6,
     'B':2,
     'C':1,
     'D':0
  }
  return H_dist[n]
Graph nodes={
  'S':[('A',1),('B',4)],
  'A':[('B',2),('C',5),('D',12)],
  'B':[('C',2)],
  'C':[('D',3)],
```

```
}
aStarAlgo('S','D')

Output:
path found:['S', 'A', 'B', 'C', 'D']
['S', 'A', 'B', 'C', 'D']
```

#### Implement AO\* Algorithm

## Algorithm:

- 1. Let G be a graph with only starting node INIT.
- 2. Repeat the following until INIT is labeled as SOLVED or h(INIT)>FUTILITY.
- Select an unexpanded node from the most promising path from INIT (call it NODE).
- Generate SUCCESSORS of NODE.

If ther are none,

Set h(NODE)=FUTILITY(i.e NODE is unsolvable)

Otherwise,

For each SUCCESSORS that is not an ancestor of NODE

Do the following:

- 1. Add SUCCESSOR toG.
- 2. If SUCCESSOR is a terminal node,

label it SOLVED and set h(SUCCESSOR)=0

3. If SUCCESSOR is not a terminal node,

Compute its h value

- Propagate the newly discovered information up the graph by doing the following:
  - 1. Let S be set of SOLVED node or nodes whose h values have been changed and need to have values propagated back to their parents.
  - 2. Initialize S to node, until S is empty.
  - 3. Repeat the following:

Remove a node from S and call it CURRENT.

#### Program:

```
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
                         # 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
      else:
                             # checking the Minimum Cost nodes 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))
      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:
                             # 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()
Output:
HEURISTIC VALUES: {'A': 1, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1, 'T':
3}
SOLUTION GRAPH : {}
PROCESSING NODE: A
______
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1, '
T': 3
SOLUTION GRAPH : {}
PROCESSING NODE : B
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1, '
T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1, '
T': 3
SOLUTION GRAPH : {}
PROCESSING NODE : G
```

```
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 7, 'J': 1, '
T': 3
SOLUTION GRAPH : {}
PROCESSING NODE: B
HEURISTIC VALUES: {'A': 10, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 7, 'J': 1, '
T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 12, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 7, 'J': 1, '
T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE: I
HEURISTIC VALUES: {'A': 12, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 0, 'J': 1, '
T': 3
SOLUTION GRAPH : {'I': []}
PROCESSING NODE: G
HEURISTIC VALUES: {'A': 12, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1, '
T': 3}
SOLUTION GRAPH : {'I': [], 'G': ['I']}
PROCESSING NODE: B
HEURISTIC VALUES: {'A': 12, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1, '
T': 3}
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE : A
______
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1, 'T':
3}
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE : C
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1, 'T':
3}
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1, 'T':
3}
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE: J
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 0, 'T':
```

3}

 $SOLUTION \ GRAPH \quad : \{'I': [\ ], 'G': ['I'], 'B': ['G'], 'J': [\ ]\}$ 

PROCESSING NODE : C

-----

HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 1, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 0, 'T':

3}

SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G'], 'J': [], 'C': ['J']}

PROCESSING NODE: A

-----

FOR GRAPH SOLUTION, TRAVERSE THE GRAPH FROM THE START NODE: A

\_\_\_\_\_

 $\{'I': [], 'G': ['I'], 'B': ['G'], 'J': [], 'C': ['J'], 'A': ['B', 'C']\}$ 

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:

- 1. Initialize G to the set of normally general hypothesis in H.
- 2. Initialize S to the set of maximally specific hypothesis in H.
- 3. For each training example d, do:
  - a. 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 & having a generalization in G
      - Remove from S any hypothesis with a more specific h i
  - b. 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 &having a specialization in S
      - Remove from G any hypothesis having a more general hypothesis in G

#### File:Enjoysport.csv

sky	airtemp	humidity	wind	water	forcast	enjoysport
sunny	warm	normal	strong	warm	same	yes
sunny	warm	high	strong	warm	same	yes
rainy	cold	high	strong	warm	change	no
sunny	warm	high	strong	cool	change	yes

#### Program:

import numpy as np import pandas as pd

```
data=pd.read csv("enjoysport.csv")
concepts=np.array(data.iloc[:,0:-1])
print("\nInstances are;\n",concepts)
target=np.array(data.iloc[:,-1])
print("\nTarget value are:",target)
def learn(concepts,target):
  specific h=concepts[0].copy()
  print("\n Initialization of specific h and general h")
  print("\n Specific Boundary:",specific h)
  general h=[["?" for i in range(len(specific h))] for i in range(len(specific h))]
  print("\n Generic Boundary:",general h)
  for i,h in enumerate(concepts):
     print("\n instance ",i+1,"is",h)
     if target[i]=="yes":
       print("\n Instance is Positive")
       for x in range(len(specific h)):
          if h[x]!=specific h[x]:
            specific h[x]='?'
            general h[x][x]="?"
     if target[i]=="no":
       print("Instance is negative")
       for x in range(len(specific h)):
          if h[x]!=specific h[x]:
            general h[x][x]=specific h[x]
          else:
            general h[x][x]='?'
     print("Specific Boundary after",i+1,"Instance is",specific h)
     print("Generic Boundary after ",i+1,"Instance is",general h)
     print("\n")
  indices=[i for i,val in enumerate(general h) if val==['?','?','?','?','?','?']]
  for i in indices:
     general h.remove(['?','?','?','?','?'])
  return specific h,general h
s final,g final=learn(concepts,target)
print("Final Specific h:",s final,sep="\n")
print("Final General h:",g final,sep="\n")
Output:
Instances are;
[['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'high' 'strong' 'warm' 'same']
```

['rainy' 'cold' 'high' 'strong' 'warm' 'change'] ['sunny' 'warm' 'high' 'strong' 'cool' 'change']] Target value are: ['yes' 'yes' 'no' 'yes'] Initialization of specific h and general h Specific Boundary: ['sunny' 'warm' 'normal' 'strong' 'warm' 'same'] ', '?', '?', '?'], ['?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']] instance 1 is ['sunny' 'warm' 'normal' 'strong' 'warm' 'same'] Instance is Positive Specific Boundary after 1 Instance is ['sunny' 'warm' 'normal' 'strong' 'warm' 'same'] 1?', 1?'], [1?', 1?', 1?', 1?', 1?', 1?'], [1?', 1?', 1?', 1?', 1?'], [1?', 1?', 1?', 1?', 1?'] instance 2 is ['sunny' 'warm' 'high' 'strong' 'warm' 'same'] Instance is Positive Specific Boundary after 2 Instance is ['sunny' 'warm' '?' 'strong' 'warm' 'same'] Generic Boundary after 2 Instance is [['?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?'], !?!, !?!], [!?!, !?!, !?!, !?!, !?!], [!?!, !?!, !?!, !?!, !?!], [!?!, !?!, !?!] instance 3 is ['rainy' 'cold' 'high' 'strong' 'warm' 'change'] Instance is negative Specific Boundary after 3 Instance is ['sunny' 'warm' '?' 'strong' 'warm' 'same'] Generic Boundary after 3 Instance is [['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?'], ['?', instance 4 is ['sunny' 'warm' 'high' 'strong' 'cool' 'change'] Instance is Positive Specific Boundary after 4 Instance is ['sunny' 'warm' '?' 'strong' '?' '?']

Generic Boundary after 4 Instance is [['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?'], ['?',

Final Specific h:

['sunny' 'warm' '?' 'strong' '?' '?']
Final General\_h:
[['sunny', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?']]

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.

- 1.Create a Root node for the tree
- 2.If all Examples are positive,

Return the single-node tree Root,

with label = +

3.If all Examples are negative,

Return the single-node tree Root,

with label = -

4.If Attributes is empty,

Return the single-node tree Root,

with label = most common value of Target\_attribute in Examples

#### Otherwise Begin,

- $\bullet$  A  $\leftarrow$  the attribute from Attributes that best classifies Examples
- The decision attribute for Root  $\leftarrow$  A
- For each possible value, vi, of A,
  - $\triangleright$  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, Targe tattribute, Attribute – {A}))

5.End

6.Return Root

# Files:

id3\_test\_1.csv

outlook	temparture	humidity	wind
rain	cool	normal	strong
sunny	mild	normal	strong

# id.csv

Outlook	Temperature	Humidity	Wind	Answer
sunny	hot	high	weak	no
sunny	hot	high	strong	no
overcast	hot	high	weak	yes
rain	mild	high	weak	yes
rain	cool	normal	weak	yes
rain	cool	normal	strong	no
overcast	cool	normal	strong	yes
sunny	mild	high	weak	no
sunny	cool	normal	weak	yes
rain	mild	normal	weak	yes
sunny	mild	normal	strong	yes
overcast	mild	high	strong	yes
overcast	hot	normal	weak	yes
rain	mild	high	strong	no

# Program:

```
import math
import csv
def load_csv(filename):
    lines=csv.reader(open(filename,'r'));
    dataset=list(lines)
```

```
headers=dataset.pop(0)
  return dataset, headers
class Node:
  def init (self,attribute):
     self.attribute=attribute
     self.children=[]
     self.answer=""
def subtables(data,col,delete):
  dic=\{\}
  coldata=[row[col] for row in data]
  attr=list(set(coldata))
  for k in attr:
     dic[k]=[]
  for y in range(len(data)):
     key=data[y][col]
     if delete:
       del data[y][col]
     dic[key].append(data[y])
  return attr,dic
def entropy(S):
  attr=list(set(S))
  if len(attr)==1:
    return 0
  counts=[0,0]
  for i in range(2):
     counts[i]=sum([1 for x in S if attr[i]==x])/(len(S)*1.0)
  sums=0
  for cnt in counts:
     sums+=-1*cnt*math.log(cnt,2)
  return sums
def compute_gain(data,col):
  attValues,dic=subtables(data,col,delete=False)
  total entropy=entropy([row[-1] for row in data])
  for x in range(len(attValues)):
     ratio=len(dic[attValues[x]])/(len(data)*1.0)
     entro=entropy([row[-1] for row in dic[attValues[x]]])
     total entropy-=ratio*entro
  return total entropy
def build tree(data,features):
  lastcol=[row[-1] for row in data]
  if(len(set(lastcol)))==1:
     node=Node("")
     node.answer=lastcol[0]
     return node
  n=len(data[0])-1
  gains=[compute gain(data,col) for col in range(n)]
  split=gains.index(max(gains))
```

```
node=Node(features[split])
  fea=features[:split]+features[split+1:]
  attr,dic=subtables(data,split,delete=True)
  for x in range(len(attr)):
     child=build tree(dic[attr[x]],fea)
     node.children.append((attr[x],child))
  return node
def print tree(node,level):
  if node.answer!="":
     print(" "*level,node.answer)
     return
  print(" "*level,node.answer)
  for value,n in node.children:
     print(""*(level+1),value)
     print tree(n,level+2)
def classify(node,x test,features):
  if node.answer!="":
     print(node.answer)
     return
  pos=features.index(node.attribute)
  for value,n in node.children:
     if x test[pos]==value:
       classify(n,x test,features)
dataset, features=load csv("id3.csv")
node = build tree(dataset, features) # Build decision tree
print ("The decision tree for the dataset using ID3 algorithm is")
print tree(node, 0)
testdata, features =load csv("id3 test 1.csv")
for xtest in testdata:
  print("The test instance xtest")
  print("The predicted labe1 ", end="" )
  classify(node, xtest, features)
Output:
The decision tree for the dataset using ID3 algorithm is
overcast
 yes
sunny
normal
   yes
high
```

no

rain

weak yes strong

no

The test instance xtest
The predicted label no
The test instance xtest
The predicted label yes

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

#### Algorithm:

# BACKPROPAGATION (training\_example, η, n<sub>in</sub>, n<sub>out</sub>, n<sub>hidden</sub>)

Each training example is a pair of the form  $(\vec{x}, \vec{t})$ , where  $(\vec{x})$  is the vector of network input values,  $(\vec{t})$  and is the vector of target network output values.

 $\eta$  is the learning rate (e.g., .05).  $n_b$  is the number of network inputs,  $n_{hidden}$  the number of units in the hidden layer, and  $n_{out}$  the number of output units.

The input from unit i into unit j is denoted  $x_{jb}$  and the weight from unit i to unit j is denoted  $w_{ji}$ 

- Create a feed-forward network with n<sub>i</sub> inputs, n<sub>hidden</sub> hidden units, and n<sub>out</sub> output units.
- · Initialize all network weights to small random numbers
- · Until the termination condition is met, Do
  - For each  $(\vec{x}, \vec{t})$ , in training examples, Do

Propagate the input forward through the network:

Input the instance into the network and compute the output ou of every unit u in the network.

Propagate the errors backward through the network:

2. For each network output unit k, calculate its error term  $\,\delta_k$ 

$$\delta_k \leftarrow o_k(1-o_k)(t_k-o_k)$$

3. For each hidden unit h, calculate its error term  $\delta_h$ 

$$\delta_h \leftarrow o_h(1 - o_h) \sum_{k \in outputs} w_{h,k} \delta_k$$

4. Update each network weight wji

$$w_{ji} \leftarrow w_{ji} + \Delta w_{ji}$$

Where

$$\Delta w_{ji} = \eta \delta_j x_{i,j}$$

```
Program:
```

```
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 sigmoid grad(x):
  return x*(1-x)
epoch=1000
eta=0.2
input_neurons=2
hidden neurons=3
output_neurons=1
wh=np.random.uniform(size=(input neurons,hidden neurons))
bh=np.random.uniform(size=(1,hidden_neurons))
wout=np.random.uniform(size=(hidden_neurons,output_neurons))
bout=np.random.uniform(size=(1,output_neurons))
for i in range(epoch):
  h ip=np.dot(X,wh)+bh
  h_act=sigmoid(h_ip)
  o ip=np.dot(h act,wout)+bout
  output=sigmoid(o ip)
  Eo=y-output
```

```
outgrad=sigmoid_grad(output)
  d_output=Eo*outgrad
  Eh=d_output.dot(wout.T)
  hiddengrad=sigmoid grad(h act)
  d_hidden=Eh*hiddengrad
  wout+=h act.T.dot(d output)*eta
  why=X.T.dot(d_hidden)*eta
print("Normalized input:\n"+str(X))
print("Actual output:\n"+str(y))
print("Predicted Output:\n",output)
Output:
Normalized input:
[[2. 9.]
[1. 5.]
[3. 6.]]
Actual output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.89224518]
[0.88542877]
[0.89224316]]
```

Write a program to implement the navier Bayesian classifier for a sample training data set stored as a .csv file. Compute the accuracy of the classifier, considering few test data sets.

#### **Explanation:**

Bayes' Theorem is stated as:

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

Where,

P(h|D) is the probability of hypothesis h given the data D. This is called the **posterior** probability.

P(D|h) is the probability of data d given that the hypothesis h was true.

**P(h)** is the probability of hypothesis h being true. This is called the **prior probability of h. P(D)** is the probability of the data. This is called the **prior probability of D** 

After calculating the posterior probability for a number of different hypotheses h, and is interested in finding the most probable hypothesis  $h \in H$  given the observed data D. Any such maximally probable hypothesis is called a *maximum a posteriori (MAP) hypothesis*.

Bayes theorem to calculate the posterior probability of each candidate hypothesis is *hMAP* is a MAP hypothesis provided.

$$h_{MAP} = \arg \max_{h \in H} P(h|D)$$

$$= \arg \max_{h \in H} \frac{P(D|h)P(h)}{P(D)}$$

$$= \arg \max_{h \in H} P(D|h)P(h)$$

(Ignoring P(D) since it is a constant)

#### **Gaussian Naive Bayes**

A Gaussian Naive Bayes algorithm is a special type of Naïve Bayes algorithm. It's specifically used when the features have continuous values. It's also assumed that all the features are following a Gaussian distribution i.e., normal distribution

#### Representation for Gaussian Naive Bayes

We calculate the probabilities for input values for each class using a frequency. With real-valued inputs, we can calculate the mean and standard deviation of input values (x) for each class to summarize the distribution.

This means that in addition to the probabilities for each class, we must also store the mean and standard deviations for each input variable for each class.

#### Gaussian Naive Bayes Model from Data

The probability density function for the normal distribution is defined by two parameters (mean and standard deviation) and calculating the mean and standard deviation values of each input variable (x) for each class value.

$$\mu = \frac{1}{n} \sum_{i=1}^{n} x_{i}$$
 Mean 
$$\sigma = \left[ \frac{1}{n-1} \sum_{i=1}^{n} (x_{i} - \mu)^{2} \right]^{0.5}$$
 Standard deviation 
$$f(x) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^{2}}{2\sigma^{2}}}$$
 Normal distribution

#### **Files:**

naviedata.csv:

6	148	72	35	0	33.6	0.627	50	1
1	85	66	29	0	26.6	0.351	31	0
8	183	64	0	0	23.3	0.672	32	1
1	89	66	23	94	28.1	0.167	21	0
0	137	40	35	168	43.1	2.288	33	1

#### **Program:**

import csv

import random

```
import math
```

```
def loadcsv(filename):
       lines = csv.reader(open(filename, "r"));
       dataset = list(lines)
       for i in range(len(dataset)):
    #converting strings into numbers for processing
               dataset[i] = [float(x) for x in dataset[i]]
       return dataset
def splitdataset(dataset, splitratio):
  #67% training size
       trainsize = int(len(dataset) * splitratio);
       trainset = []
       copy = list(dataset);
       while len(trainset) < trainsize:
#generate indices for the dataset list randomly to pick ele for training data
               index = random.randrange(len(copy));
               trainset.append(copy.pop(index))
       return [trainset, copy]
def separatebyclass(dataset):
       separated = \{\} #dictionary of classes 1 and 0
#creates a dictionary of classes 1 and 0 where the values are
#the instances belonging to each class
       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): #creates a dictionary of classes
       summaries = [(mean(attribute), stdev(attribute)) for attribute in zip(*dataset)];
       del summaries[-1] #excluding labels +ve or -ve
       return summaries
def summarizebyclass(dataset):
       separated = separatebyclass(dataset);
  #print(separated)
       summaries = \{\}
       for classvalue, instances in separated.items():
#for key, value in dic.items()
#summaries is a dic of tuples(mean,std) for each class value
              summaries[classvalue] = summarize(instances) #summarize is used to cal to
mean and std
       return summaries
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 = {} # probabilities contains the all prob of all class of test data
       for classvalue, classsummaries in summaries.items():#class and attribute information
as mean and sd
               probabilities[classvalue] = 1
               for i in range(len(classsummaries)):
                       mean, stdev = classsummaries[i] #take mean and sd of every attribute
for class 0 and 1 seperaely
                       x = inputvector[i] #testvector's first attribute
                       probabilities[classvalue] *= calculateprobability(x, mean, stdev);#use
normal dist
       return probabilities
def predict(summaries, inputvector): #training and test data is passed
       probabilities = calculateclassprobabilities(summaries, inputvector)
       bestLabel, bestProb = None, -1
       for classvalue, probability in probabilities.items():#assigns that class which has he
highest prob
               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 = 'naivedata.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)))
       # prepare model
       summaries = summarizebyclass(trainingset);
       #print(summaries)
  # test model
       predictions = getpredictions(summaries, testset) #find the predictions of test data with
the training data
       accuracy = getaccuracy(testset, predictions)
       print('Accuracy of the classifier is : {0}%'.format(accuracy))
main()
Output:
Split 768 rows into train=514 and test=254 rows
```

Accuracy of the classifier is: 73.62204724409449%

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

## **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
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']
model=KMeans(n clusters=3)
model.fit(X)
plt.figure(figsize=(14,7))
colormap=np.array(['red','lime','black'])
plt.subplot(1,3,1)
plt.scatter(X.Petal Length,X.Petal Width,c=colormap[y.Targets],s=40)
plt.title('Real Clasification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
```

```
plt.subplot(1,3,2)
plt.scatter(X.Petal_Length,X.Petal_Width,c=colormap[model.labels_],s=40)
plt.title('K Mean Classification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
print('The accuracy score of K-Mean:',sm.accuracy score(y,model.labels ))
print('The Confusion matrix of K-Mean:',sm.confusion matrix(y,model.labels ))
from sklearn import preprocessing
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)
y gmm=gmm.predict(xs)
plt.subplot(1,3,3)
plt.scatter(X.Petal_Length,X.Petal_Width,c=colormap[y_gmm],s=40)
plt.title('GMM Classification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
print('The accuracy score of EM:',sm.accuracy score(y,y gmm)*100)
print('The confusion matrix of EM:',sm.confusion matrix(y,y gmm))
Output:
```

The accuracy score of K-Mean: 0.24

The Confusion matrix of K-Mean: [[ 0 50 0]

[2 0 48]

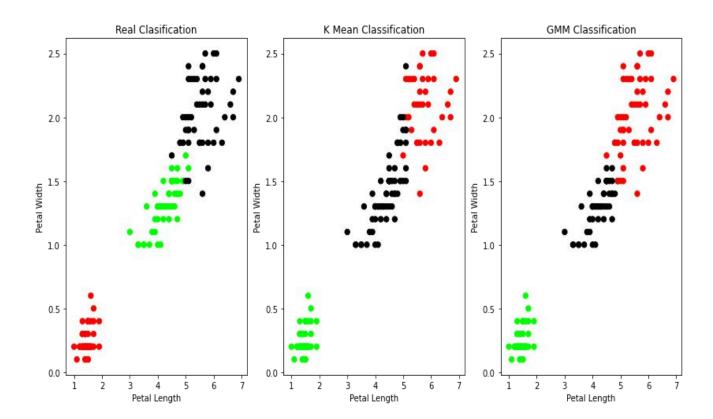
[36 0 14]]

The accuracy score of EM:30.0

The confusion matrix of EM: [[50 0 0]

[0 5 45]

[0500]]



Write a program to implement K-Nearest Neighbour 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:

#### K-Nearest Neighbor Algorithm

Training algorithm:

- For each training example (x, f (x)), add the example to the list training examples Classification algorithm:
  - Given a query instance xq to be classified,
    - Let  $x_1 cdots x_k$  denote the k instances from training examples that are nearest to  $x_q$
    - Return

$$\hat{f}(x_q) \leftarrow \frac{\sum_{i=1}^k f(x_i)}{k}$$

 Where, f(xi) function to calculate the mean value of the k nearest training examples.

## **Program:**

from sklearn.datasets import load iris

iris=load iris()

print("Features Names:",iris.feature\_names,"Iris Data:",iris.data,"Target Names:",iris.target names,"Target:",iris.target)

from sklearn.model\_selection import train\_test\_split

X\_train,X\_test,y\_train,y\_test=train\_test\_split(iris.data,iris.target,test\_size=.25)

from sklearn.neighbors import KNeighborsClassifier

clf=KNeighborsClassifier()

```
clf.fit(X_train,y_train)
print("Accuracy=",clf.score(X_test,y_test))
print("Predicted Data")
print(clf.predict(X_test))
prediction=clf.predict(X_test)
print("Test Data:")
print(y_test)
diff=prediction-y_test
print("Result is")
print(diff)
print("Total no of samples misclassified=",sum(abs(diff)))
Output:
Features Names: ['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)', 'petal width (cm)']
Iris Data: [[5.1 3.5 1.4 0.2]
[4.9 3. 1.4 0.2]
[4.7 3.2 1.3 0.2]
[4.6 3.1 1.5 0.2]
[5. 3.6 1.4 0.2]
[5.4 3.9 1.7 0.4]
[4.6 3.4 1.4 0.3]
[5. 3.4 1.5 0.2]
[4.4 2.9 1.4 0.2]
[4.9 3.1 1.5 0.1]
[5.4 3.7 1.5 0.2]
[4.8 3.4 1.6 0.2]
[4.8 3. 1.4 0.1]
[4.3 3. 1.1 0.1]
[5.8 4. 1.2 0.2]
[5.7 4.4 1.5 0.4]
[5.4 3.9 1.3 0.4]
[5.1 3.5 1.4 0.3]
```

- [5.7 3.8 1.7 0.3]
- [5.1 3.8 1.5 0.3]
- [5.4 3.4 1.7 0.2]
- [5.1 3.7 1.5 0.4]
- [4.6 3.6 1. 0.2]
- [5.1 3.3 1.7 0.5]
- [4.8 3.4 1.9 0.2]
- [5. 3. 1.6 0.2]
- [5. 3.4 1.6 0.4]
- [5.2 3.5 1.5 0.2]
- [5.2 3.4 1.4 0.2]
- [4.7 3.2 1.6 0.2]
- [4.8 3.1 1.6 0.2]
- [5.4 3.4 1.5 0.4]
- [5.2 4.1 1.5 0.1]
- [5.5 4.2 1.4 0.2]
- [4.9 3.1 1.5 0.2]
- [5. 3.2 1.2 0.2]
- [5.5 3.5 1.3 0.2]
- [4.9 3.6 1.4 0.1]
- [4.4 3. 1.3 0.2]
- [5.1 3.4 1.5 0.2]
- [5. 3.5 1.3 0.3]
- [4.5 2.3 1.3 0.3]
- [4.4 3.2 1.3 0.2]
- [5. 3.5 1.6 0.6]
- [5.1 3.8 1.9 0.4]
- [4.8 3. 1.4 0.3]
- [5.1 3.8 1.6 0.2]
- [4.6 3.2 1.4 0.2]

- [5.3 3.7 1.5 0.2]
- [5. 3.3 1.4 0.2]
- [7. 3.2 4.7 1.4]
- [6.4 3.2 4.5 1.5]
- [6.9 3.1 4.9 1.5]
- [5.5 2.3 4. 1.3]
- [6.5 2.8 4.6 1.5]
- [5.7 2.8 4.5 1.3]
- [6.3 3.3 4.7 1.6]
- [4.9 2.4 3.3 1.]
- [6.6 2.9 4.6 1.3]
- [5.2 2.7 3.9 1.4]
- [5. 2. 3.5 1.]
- [5.9 3. 4.2 1.5]
- [6. 2.2 4. 1.]
- [6.1 2.9 4.7 1.4]
- [5.6 2.9 3.6 1.3]
- [6.7 3.1 4.4 1.4]
- [5.6 3. 4.5 1.5]
- [5.8 2.7 4.1 1.]
- [6.2 2.2 4.5 1.5]
- [5.6 2.5 3.9 1.1]
- [5.9 3.2 4.8 1.8]
- [6.1 2.8 4. 1.3]
- [6.3 2.5 4.9 1.5]
- [6.1 2.8 4.7 1.2]
- [6.4 2.9 4.3 1.3]
- [6.6 3. 4.4 1.4]
- [6.8 2.8 4.8 1.4]
- [6.7 3. 5. 1.7]

- [6. 2.9 4.5 1.5]
- [5.7 2.6 3.5 1.]
- [5.5 2.4 3.8 1.1]
- [5.5 2.4 3.7 1.]
- [5.8 2.7 3.9 1.2]
- [6. 2.7 5.1 1.6]
- [5.4 3. 4.5 1.5]
- [6. 3.4 4.5 1.6]
- [6.7 3.1 4.7 1.5]
- [6.3 2.3 4.4 1.3]
- [5.6 3. 4.1 1.3]
- [5.5 2.5 4. 1.3]
- [5.5 2.6 4.4 1.2]
- [6.1 3. 4.6 1.4]
- [5.8 2.6 4. 1.2]
- [5. 2.3 3.3 1.]
- [5.6 2.7 4.2 1.3]
- [5.7 3. 4.2 1.2]
- [5.7 2.9 4.2 1.3]
- [6.2 2.9 4.3 1.3]
- [5.1 2.5 3. 1.1]
- [5.7 2.8 4.1 1.3]
- [6.3 3.3 6. 2.5]
- [5.8 2.7 5.1 1.9]
- [7.1 3. 5.9 2.1]
- [6.3 2.9 5.6 1.8]
- [6.5 3. 5.8 2.2]
- [7.6 3. 6.6 2.1]
- [4.9 2.5 4.5 1.7]
- [7.3 2.9 6.3 1.8]

- [6.7 2.5 5.8 1.8]
- [7.2 3.6 6.1 2.5]
- [6.5 3.2 5.1 2.]
- [6.4 2.7 5.3 1.9]
- [6.8 3. 5.5 2.1]
- [5.7 2.5 5. 2.]
- [5.8 2.8 5.1 2.4]
- [6.4 3.2 5.3 2.3]
- [6.5 3. 5.5 1.8]
- [7.7 3.8 6.7 2.2]
- [7.7 2.6 6.9 2.3]
- [6. 2.2 5. 1.5]
- [6.9 3.2 5.7 2.3]
- [5.6 2.8 4.9 2.]
- [7.7 2.8 6.7 2.]
- [6.3 2.7 4.9 1.8]
- [6.7 3.3 5.7 2.1]
- [7.2 3.2 6. 1.8]
- [6.2 2.8 4.8 1.8]
- [6.1 3. 4.9 1.8]
- [6.4 2.8 5.6 2.1]
- [7.2 3. 5.8 1.6]
- [7.4 2.8 6.1 1.9]
- [7.9 3.8 6.4 2.]
- [6.4 2.8 5.6 2.2]
- [6.3 2.8 5.1 1.5]
- [6.1 2.6 5.6 1.4]
- [7.7 3. 6.1 2.3]
- [6.3 3.4 5.6 2.4]
- [6.4 3.1 5.5 1.8]

```
[6. 3. 4.8 1.8]
[6.9 3.1 5.4 2.1]
[6.7 3.1 5.6 2.4]
[6.9 3.1 5.1 2.3]
[5.8 2.7 5.1 1.9]
[6.8 3.2 5.9 2.3]
[6.7 3.3 5.7 2.5]
[6.7 3. 5.2 2.3]
[6.3 2.5 5. 1.9]
[6.5 3. 5.2 2.]
[6.2 3.4 5.4 2.3]
[5.9 3. 5.1 1.8]] Target Names: ['setosa' 'versicolor' 'virginica'] Target: [0 0 0 0 0 0 0 0 0 0 0 0
2 2]
Accuracy= 1.0
Predicted Data
[2\ 2\ 1\ 2\ 2\ 1\ 2\ 0\ 0\ 2\ 0\ 0\ 2\ 2\ 0\ 2\ 1\ 2\ 1\ 1\ 0\ 0\ 2\ 2\ 1\ 2\ 2\ 0\ 1\ 0\ 2\ 1\ 2\ 1\ 1\ 2\ 0
1]
Test Data:
[2\ 2\ 1\ 2\ 2\ 1\ 2\ 0\ 0\ 2\ 0\ 0\ 2\ 2\ 0\ 2\ 1\ 2\ 1\ 1\ 0\ 0\ 2\ 2\ 1\ 2\ 2\ 0\ 1\ 0\ 2\ 1\ 2\ 1\ 1\ 2\ 0
1]
Result is
0]
Total no of samples misclassified= 0
```

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

#### Algorithm:

Locally Weighted Regression Algorithm:

- 1. Read the Given data Sample to X and the curve (linear or non linear) to Y
- 2. Set the value for Smoothening parameter or Free parameter say  $\tau$
- 3. Set the bias /Point of interest set x0 which is a subset of X
- 4. Determine the weight matrix using:

$$w(x, x_o) = e^{-\frac{(x - x_o)^2}{2\tau^2}}$$

4. Determine the value of model term parameter  $\beta$  using:

$$\hat{\beta}(x_o) = (X^T W X)^{-1} X^T W y$$

5. Prediction =  $x0*\beta$ 

#### **CSV Files:**

tips.csv

total_bill	tip	sex	smoker	day	time	size
16.99	1.01	Female	No	Sun	Dinner	2
10.34	1.66	Male	No	Sun	Dinner	3
21.01	3.5	Male	No	Sun	Dinner	3
23.68	3.31	Male	No	Sun	Dinner	2

#### Program:

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)))
  for j in range(m):
    diff = point - X[i]
    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
# load data points
data = pd.read_csv('tips.csv')
bill = np.array(data.total bill)
tip = np.array(data.tip)
#preparing and add 1 in bill
mbill = np.mat(bill)
mtip = np.mat(tip)
m= np.shape(mbill)[1]
```

```
one = np.mat(np.ones(m))
X = np.hstack((one.T,mbill.T))

#set k here

ypred = localWeightRegression(X,mtip,0.5)

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();
```

# **Output:**

