

Department of Computer Science and Engineering

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Department of Computer Science and Engineering

BE - VII SEMESTER

ARTIFICIAL INTELLIGENCE AND MACHINE LEARNING LABORATORY

18CSL76

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1. Implement A* Search Algorithm.

```
class Node():
    """A node class for A* Pathfinding"""
    def init (self, parent=None, position=None):
         self.parent = parent
         self.position = position
         self.g = 0
         self.h = 0
         self.f = 0
    def __eq__(self, other):
         return self.position == other.position
# Simulate poping from a priority queue. The node with least f value is popped from queue
def pop_queue(que):
# Get the current node
        current node = que[0]
        current index = 0
        for index, item in enumerate(que):
            if item.f < current_node.f:</pre>
                 current node = item
                 current_index = index
        # Pop current off open list, add to closed list
        return que.pop(current_index)
def astar(maze, start, end):
    """Returns a list of tuples as a path from the given start to the given end in the given maze"""
   # Create start and end node
   start_node = Node(None, start)
   start_node.g = start_node.h = start_node.f = 0
   end_node = Node(None, end)
   end_node.g = end_node.h = end_node.f = 0
   # Initialize both open and closed list
   open_list = []
   closed_list = []
    # Add the start node
   open_list.append(start_node)
    # Loop until you find the end
   while len(open_list) > 0:
       current_node = pop_queue(open_list)
       closed_list.append(current_node)
       # Found the goal
       if current_node == end_node:
           path = []
           current = current node
           while current is not None:
               path.append(current.position)
               current = current.parent
           return path[::-1] # Return reversed path
```

```
# Generate children
children = []
for new_position in [(0, -1), (0, 1), (-1, 0), (1, 0), (-1, -1), (-1, 1), (1, -1), (1, 1)]: # Adjacent squares
    # Get node position
    node_position = (current_node.position[0] + new_position[0], current_node.position[1] + new_position[1])
    # Make sure within range
    if node_position[0] > (len(maze) - 1) or node_position[0] < 0 or node_position[1] >
    (len(maze[len(maze)-1]) -1) or node_position[1] < 0:
    # Make sure walkable terrain
    if maze[node_position[0]][node_position[1]] != 0:
        continue
    # Create new node
    new_node = Node(current_node, node_position)
    # Append
    children.append(new_node)
# Loop through children
for child in children:
    # if Child is on the closed list, skip it
    child_in_closed = False
    for closed child in closed list:
        if child == closed child:
                child_in_closed = True
                break
        if child in closed:
                continue
        # Create the f, g, and h values
        child.g = current_node.g + 1
        dx = abs(child.position[0] - end_node.position[0])
        dy = abs(child.position[1] - end_node.position[1])
        D = 1 # distance to next horizontal/vertical node
        D2 = 1 # distance to next diagonal node
        child.h = D * (dx + dy) + (D2 - 2 * D) * min(dx, dy)
        \#child.h = ((child.position[0] - end\_node.position[0]) ** 2) + ((child.position[1] - end\_node.position[1]) ** 2)
        child.f = child.g + child.h
        # Child is already in the open list
        discard child = False
        for open_node in open_list:
            if child == open node:
                if child.g < open node.g:
                    open node.g = child.g
                    open_node.f = open_node.g + open_node.h
                    open_node.parent = current_node
                discard child = True
                break
```

```
# Add the child to the open list
              if discard_child == False:
                  open_list.append(child)
def main():
    maze = [[0, 0, 0, 0, 1, 0, 0, 0, 0, 0],
              [0, 0, 0, 0, 1, 0, 0, 0, 0, 0],
              [0, 0, 0, 0, 1, 0, 0, 0, 0, 0],
              [0, 0, 0, 0, 1, 0, 1, 0, 0, 0], [0, 0, 0, 0, 1, 0, 0, 0, 0, 0],
              [0, 0, 0, 0, 0, 0, 1, 0, 0, 0],
              [0, 0, 0, 0, 1, 0, 1, 0, 0, 0],
              [0, 0, 0, 0, 1, 0, 1, 0, 0, 0],
              [0, 0, 0, 0, 1, 0, 1, 0, 0, 0],
[0, 0, 0, 0, 0, 0, 0, 0, 0.2, 0]]
    start = (0, 0)
    end = (0, 9)
    path = astar(maze, start, end)
    print(path)
if __name__ == '__main__':
    main()
```

Output:

[(0, 0), (1, 1), (2, 2), (3, 3), (4, 3), (5, 4), (4, 5), (4, 6), (3, 7), (2, 7), (1, 8), (0, 9)]

```
a = (1,2)
b = (1,2)
if a==b:
    print("Same")
```

Output:

Same

2. Implement AO* Search Algorithm.

```
class Node:
    def __init__(self, index,cost,visited=False,is solved=False,and map=False, or map=False):
        self.index=index
        self.cost=cost
        self.visited=visited
        self.is solved=is solved
        self.and map= and map
        self.or map = or map
        self.children=()
    def __str__(self):
        return f'{self.index}: {self.cost}'
    def set_children(self,ch):
        self.children=ch
         adj=[]
         n nodes = 21
         #heuristic costs
         cost=[None,0,40,2,4,1,2,3,50,60,70,80,4,5,8,9,6,7,90,90,90,90]
         and edges={}
         for i in range(n nodes+1):
             n=Node(i, cost[i])
             adj.append(n)
         adj[1].set_children((adj[2],adj[3],adj[4]))
         adj[2].set_children((adj[5],adj[6],adj[7]))
         adj[3].set_children((adj[8],adj[9]))
         adj[4].set children((adj[10],adj[11]))
         adj[5].set_children((adj[12],adj[13])); adj[6].set_children((adj[14],adj[15]))
         adj[7].set_children((adj[16],adj[17])); adj[8].set_children((adj[18],))
         adj[9].set_children((adj[19],)); adj[10].set_children((adj[20],)); adj[11].set_children((adj[21],))
         and_edges[adj[1]] = (adj[3],adj[4])
         adj[3].and map = adj[4].and map = True
         and_edges[adj[2]] = (adj[5],adj[6], adj[7])
         adj[5].and_map = adj[6].and_map = adj[7].and_map=True
         for a in adj:
            if len(a.children)==0: a.is_solved=True
             if a.and map==False: a.or map=True
             #print(f'{a.index} and {a.and map} or {a.or map}')
```

```
def get key(and edges, c):
    for idx, ae in and edges.items():
        if c in ae: return idx, ae
def explore head(head):
    print(f'Head: {head.index}, Cost: {head.cost}')
    head.visited=True; temp_cost = MAX; temp_map={}
    for c in head.children:
        if temp_map.get(c,False): continue;
        if c.and_map: # if the child is in the and edge
            temp solved=True
           #calculate the cost and check if there are more nodes in the and edge
            idx,ae = get_key(and_edges,c)
            cc=0
            for aek in ae:
                cc+=aek.cost+EDGE
                temp_map[aek]=True
                temp_solved=temp_solved and aek.is_solved
            temp_cost = min(temp_cost,cc)
            if temp_solved:
                head.is solved=True
        else: # else if child is in the or edge
            temp_cost = min(temp_cost,c.cost+EDGE)
            temp_map[c]=True
            if c.is_solved: head.is_solved=True
    #head is explored now update the best value of head i.e. temp cost
    if temp_cost < MAX:
       head.cost=temp cost
        print(f'Updated head cost {head.cost}')
def find best move(head):
     #find the best move
        isAnd=False
        bestCost=MAX;bestMove=None; bestAndIndex=-1
        temp map1={}
        for c in head.children:
            if temp_map1.get(c,False):continue
            if c.or map:
                if bestCost>c.cost+EDGE:
                    bestCost = c.cost+EDGE
                    bestMove=c; isAnd=False
                    temp map1[c]=True
                print(f'or edge {c.index}, {bestCost}')
            else:
                cc=0
                idx,ae = get_key(and_edges,c)
                for aek in ae:
                    cc+=aek.cost+EDGE
                    temp map1[aek]=True
                    print(f'and-pair {idx.index}-{c.index}')
                print(f'bestCost {bestCost} cc {cc}')
                if bestCost>cc and cc!=0:
                    bestCost = cc; bestAndIndex = idx; bestMove = c
                    isAnd=True
```

```
print(f'\nmoving forward, finding the best move,i>>{c.index}')
      if isAnd:
          print(f'and edge, cost: {bestCost}')
      else:
          print(f'or edge, cost: {bestCost}')
 if isAnd:
      for ae in and_edges[bestAndIndex]:
          print(f'isAnd: {isAnd} and, aoStarUtil {ae.index}')
          aostarUtil(ae)
 else:
      print(f'isAnd: {isAnd} or, aoStarUtil {bestMove.index}')
      aostarUtil(bestMove)
def check_update(head):
    temp_cost=MAX; temp_map={}
    for c in head.children:
        if temp_map.get(c,False):continue
        if c.or_map:
            if c.is_solved: head.is_solved=True
            temp_cost= min(temp_cost, c.cost+EDGE)
            temp_map[c]=True
        elif c.and_map:
            f=True;cc=0
            idx,ae = get_key(and_edges,c)
            for aek in ae:
                f = f and aek.is_solved
                cc+=aek.cost+EDGE
                temp_map[aek]=True
            temp_cost = min(temp_cost,cc)
            print(f'temp_cost=min({temp_cost},{cc})')
            if f:
                head.is_solved=True
                break
    if temp_cost<=MAX:</pre>
        head.cost = temp_cost
    print(f'Updated Cost of node {head.index} {head.cost}')
def aostarUtil(head):
    if head.visited ==False:
        explore_head(head)
    else:
        find best move(head)
        #check if any of the options were solved
        #if there are multiple solved nodes , select the best out of them
        #also update the current cost i.e. head cost while backtracking to the root
        check update(head)
def aostar(head):
    iter = 0
    while head.is solved==False and iter <MAX:
        print(f'\n **Iteration {iter}')
        aostarUtil(head)
        iter+=1
    for a in adj:
        print(a.index,': ',a.cost, end=" ")
```

```
MAX=1000
EDGE=5 #g cost of edge
aostar(adj[1])
```

Output:

```
**Iteration 0
Head: 1, Cost: 0
Updated head cost 16
  **Iteration 1
or edge 2, 45
moving forward, finding the best move, i>>2
or edge, cost: 45
and-pair 1-3
and-pair 1-3
bestCost 45 cc 16
moving forward, finding the best move,i>>3
and edge, cost: 16
isAnd: True and, aoStarUtil 3
Head: 3, Cost: 2
Updated head cost 55
isAnd: True and, aoStarUtil 4
Head: 4, Cost: 4
Updated head cost 75
temp_cost=min(45,140)
Updated Cost of node 1 45
  **Iteration 2
or edge 2, 45
moving forward, finding the best move,i>>2
or edge, cost: 45
and-pair 1-3
and-pair 1-3
bestCost 45 cc 140
moving forward, finding the best move,i>>3
or edge, cost: 45
isAnd: False or, aoStarUtil 2
Head: 2, Cost: 40
Updated head cost 21
temp_cost=min(26,140)
Updated Cost of node 1 26
  **Iteration 3
or edge 2, 26
moving forward, finding the best move, i>>2
or edge, cost: 26
and-pair 1-3
and-pair 1-3
bestCost 26 cc 140
moving forward, finding the best move, i>>3
or edge, cost: 26
isAnd: False or, aoStarUtil 2
and-pair 2-5
and-pair 2-5
and-pair 2-5
bestCost 1000 cc 21
```

```
moving forward, finding the best move,i>>5
and edge, cost: 21
isAnd: True and, aoStarUtil 5
Head: 5, Cost: 1
Updated head cost 9
isAnd: True and, aoStarUtil 6
Head: 6, Cost: 2
Updated head cost 13
isAnd: True and, aoStarUtil 7
Head: 7, Cost: 3
Updated head cost 11
temp_cost=min(48,48)
Updated Cost of node 2 48
temp_cost=min(53,140)
Updated Cost of node 1 53
0: None 1: 53 2: 48 3: 55 4: 75 5: 96: 13 7: 11 8: 50 9: 60 10: 70 11: 80 12: 4 13: 5 14: 8 15:
9 16 : 6 17 : 7 18 : 90 19 : 90 20 : 90 21 : 90
```

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.

```
import csv
a = []
print("\n The Given Training Data Set \n")
with open('data1.csv', 'r') as csvFile:
    reader = csv.reader(csvFile)
    for row in reader:
        a.append (row)
        print(row)
num attributes = len(a[0])-1 # we don't want Last col which is target concet ( ves/no)
print("\n The initial value of hypothesis: ")
S = ['0'] * num_attributes
G = ['?'] * num attributes
print ("\n The most specific hypothesis S0 : [0,0,0,0,0,0]\n")
print (" \n The most general hypothesis G0 : [?,?,?,?,?]\n")
for j in range(0,num_attributes):
       S[j] = a[0][j];
# Comparing with Remaining Training Examples of Given Data Set
print("\n Candidate Elimination algorithm Hypotheses Version Space Computation\n")
temp=[]
for i in range(0,len(a)):
    if a[i][num_attributes]=='1':
        for j in range(0,num_attributes):
            if a[i][j]!=S[j]:
                S[j]='?'
        for j in range(0,num_attributes):
            for k in range(0,len(temp)):
                if temp[k][j] != '?' and temp[k][j] != S[j]:
                    del temp[k] #remove it if it's not matching with the specific hypothesis
        print(" For Training Example No :{0} the hypothesis is S{0} ".format(i+1),S)
        if (len(temp)==0):
            print(" For Training Example No :{0} the hypothesis is G{0} ".format(i+1),G)
            print(" For Training Example No :{0} the hypothesis is G{0}".format(i+1),temp)
   if a[i][num_attributes]=='0':
        for j in range(0,num_attributes):
             if S[j] != a[i][j] and S[j]!= '?': #if not matching with the specific Hypothesis
                 G[j]=S[j]
                 temp.append(G) # this is the version space to store all Hypotheses
                 G = ['?'] * num_attributes
        print(" For Training Example No :{0} the hypothesis is S{0} ".format(i+1),S)
        print(" For Training Example No :{0} the hypothesis is G{0}".format(i+1),temp)
```

Dataset to be considered is as follows:-

sunny	warm	normal	strong	warm	same	1
sunny	warm	high	strong	warm	same	1
rainy	cold	high	strong	warm	change	0
sunny	warm	high	strong	cool	change	1

The output is as follows:-

```
The Given Training Data Set
['sunny', 'warm', 'normal', 'strong', 'warm', 'same', '1']
['sunny', 'warm', 'high', 'strong', 'warm', 'same', '1']
['rainy', 'cold', 'high', 'strong', 'warm', 'change', '0']
['sunny', 'warm', 'high', 'strong', 'cool', 'change', '1']
The initial value of hypothesis:
 The most specific hypothesis SO: [0,0,0,0,0,0]
 The most general hypothesis GO: [?,?,?,?,?]
 Candidate Elimination algorithm Hypotheses Version Space Computation
 For Training Example No :1 the hypothesis is S1 ['sunny', 'warm', 'normal', 'strong',
'warm', 'same']
 For Training Example No :1 the hypothesis is G1 ['?', '?', '?', '?', '?']
 For Training Example No :2 the hypothesis is S2 ['sunny', 'warm', '?', 'strong',
'warm', 'same']
 For Training Example No :2 the hypothesis is G2 ['?', '?', '?', '?', '?']
 For Training Example No :3 the hypothesis is S3 ['sunny', 'warm', '?', 'strong',
'warm', 'same']
For Training Example No :3 the hypothesis is G3 [['sunny', '?', '?', '?', '?'],
['?', 'warm', '?', '?', '?'], ['?', '?', '?', '?', '?', 'same']]
For Training Example No :4 the hypothesis is S4 ['sunny', 'warm', '?', 'strong', '?',
'?'1
 For Training Example No :4 the hypothesis is G4 [['sunny', '?', '?', '?', '?'],
['?', 'warm', '?', '?', '?', '?']]
```

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.

```
import pandas as pd
import math
import numpy as np
data = pd.read_csv("3_data.csv")
features = [feat for feat in data]
features.remove("Target")
class Node:
    def __init__(self):
         self.children = []
         self.value = ""
         self.isLeaf = False
self.pred = ""
def entropy(examples):
    pos = 0.0
    neg = 0.0
    for _, row in examples.iterrows():
    if row["Target"] == "yes":
             pos += 1
         else:
            neg += 1
     if pos == \overline{0}.0 or neg == 0.0:
         return 0.0
     else:
         p = pos / (pos + neg)
         return -(p * math.log(p, 2) + n * math.log(n, 2))
```

```
def info_gain(examples, attr):
    uniq = np.unique(examples[attr])
#print ("\n",uniq)
    gain = entropy(examples)
    #print ("\n",gain)
    for u in uniq:
         subdata = examples[examples[attr] == u]
#print ("\n", subdata)
         sub_e = entropy(subdata)
gain -= (float(len(subdata)) / float(len(examples))) * sub_e
         #print ("\n",gain)
    return gain
def ID3(examples, attrs):
    root = Node()
    max\_gain = 0
    max_feat = ""
    for feature in attrs:
         #print ("\n",examples)
         gain = info_gain(examples, feature)
         if gain > max_gain:
              max_gain = gain
max_feat = feature
    root.value = max_feat
    #print ("\nMax feature attr",max_feat)
    uniq = np.unique(examples[max_feat])
#print ("\n",uniq)
    for u in uniq:
#print ("\n",u)
         subdata = examples[examples[max_feat] == u]
#print ("\n",subdata)
         if entropy(subdata) == 0.0:
              newNode = Node()
```

```
newNode.isLeaf = True
    newNode.value = u
    newNode.pred = np.unique(subdata["Target"])
    root.children.append(newNode)
else:
    dummyNode = Node()
    dummyNode.value = u
    new_attrs = attrs.copy()
    new_attrs.remove(max_feat)
    child = ID3(subdata, new_attrs)
    dummyNode.children.append(child)
    root.children.append(dummyNode)
return root

def printTree(root: Node, depth=0):
    for i in range(depth):
        print("\t", end="")
    print(root.value, end="")
    if root.isLeaf:
        print(" -> ", root.pred)
    print()
    for child in root.children:
        printTree(child, depth + 1)

root = ID3(data, features)
printTree(root)
```

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The dataset to be considered is as follows:-

Outlook	Temperat	Humidity	Wind	Target
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

The output is:

Output:

The decision tree for the dataset using ID3 algorithm is

```
Outlook
rain
Wind
strong
no
weak
yes
overcast
yes
sunny
Humidity
normal
yes
high
no
```

The test instance: ['rain', 'cool', 'normal', 'strong']

The label for test instance: no

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

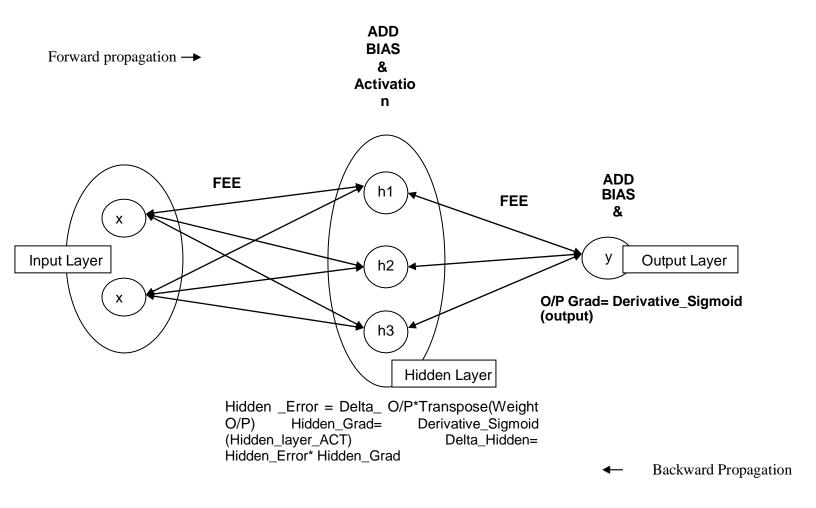
```
1 import numpy as np
 2 X = np.array(([2, 9], [1, 5], [3, 6]), dtype=float)
 3 y = np.array(([92], [86], [89]), dtype=float)
 4 X = X/np.amax(X,axis=0) # maximum of X array longitudinally
 5 y = y/100
 6 #Sigmoid Function
7 def sigmoid (x):
      return 1/(1 + np.exp(-x))
9 #Derivative of Sigmoid Function
10 def derivatives sigmoid(x):
11
       return x * (1 - x)
12 #Variable initialization
13 epoch=1
14 #Setting training iterations
15 lr=0.1 #Setting learning rate
16 inputlayer neurons = 2 #number of features in data set
17 hiddenlayer neurons = 3 #number of hidden layers neurons
18 output neurons = 1 #number of neurons at output layer
19 #weight and bias initialization
20 wh=np.random.uniform(size=(inputlayer neurons, hiddenlayer neurons))
21 bh=np.random.uniform(size=(1,hiddenlayer neurons))
22 wout=np.random.uniform(size=(hiddenlayer_neurons,output_neurons))
23 bout=np.random.uniform(size=(1,output_neurons))
 24 #draws a random range of numbers uniformly of dim x*y
 25 for i in range(epoch):
 26 #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)
 33 #Backpropagation
 34 EO = y-output
 35    outgrad = derivatives_sigmoid(output)
 d_output = EO* outgrad

EH = np.dot(d_output,wout.T)

#wout.T is used to calculate the transpose of wout

hiddengrad = derivatives_sigmoid(hlayer_act)
 40  #how much hidden layer wts contributed to error
41  d_hiddenlayer = EH * hiddengrad
 42 #Update weights and bias for next iteration
 43 wout += np.dot(hlayer_act.T,d_output) *lr
 44 # dotproduct of nextlayererror and currentlayerop
      bout += np.sum(d_output, axis=0) *lr
       wh += X.T.dot(d hiddenlayer) *lr
       bh += np.sum(d_hiddenlayer, axis=0) *lr
 48 print("Input: \n" + str(X))
 49 print("Actual Output: \n" + str(y))
 50 print("Predicted Output: \n" ,output)
 str(y-output))
51 print("Final Error In Predicted Output: \n" ,str(y-output))
```

For this we consider:



The output is as follows:

```
Input:
[[0.66666667 1.
 [0.33333333 0.55555556]
             0.66666667]]
Actual Output:
[[0.92]
 [0.86]
 [0.89]]
Predicted Output:
 [[0.88169538]
 [0.8710618]
 [0.88065579]]
Final Error In Predicted Output:
 [[ 0.03830462]
 [-0.0110618]
 [ 0.00934421]]
```

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.

The dataset to be considered is as follows:-

Outlook	Temperat	Humidity	Wind	Target
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

```
1 import numpy as np
 2 import csv
 3 def read data(filename):
      with open(filename, 'r') as csvfile:
 5
         datareader = csv.reader(csvfile)
 6
          metadata = next(datareader)
 7
          traindata=[]
        for row in datareader:
 8
              traindata.append(row)
9
         print(traindata,"\n")
10
          print("Metadata is:\n ",metadata)
11
     return (metadata, traindata)
12
13
14 def splitDataset(dataset, splitRatio):
15 trainSize = int(len(dataset) * splitRatio)
     trainSet = []
16
17 testset = list(dataset)
18
     i=0
     while len(trainSet) <trainSize:</pre>
19
          trainSet.append(testset.pop(i))
20
   return [trainSet, testset]
21
```

```
22 def classify(train,test):
23
      train rows = train.shape[0] #total training rows
      test rows=test.shape[0]
                               #total testina rows
24
25
      train_col = train.shape[1]
      test col = test.shape[1]
26
      print("training data size=",train rows)
27
      print("test data size=",test.shape[0])
28
29
      countYes,countNo,probYes,probNo=0,0,0,0
      print("target
                      count
                               probability")
30
31
      for x in range(train rows):
          if train[x,train col-1] == 'yes':
32
33
              countYes +=1
34
          if train[x,train col-1] == 'no':
              countNo +=1
35
36
      probYes=countYes/train rows
37
      probNo= countNo / train rows
38
      print('Yes',"\t",countYes,"\t",probYes)
      print('No',"\t",countNo,"\t",probNo)
39
      prob0 =np.zeros((test col-1))
40
41
      prob1 =np.zeros((test col-1))
42
      accuracy=0
43
      for t in range(test rows):
44
          for k in range (test.shape[1]-1): #test.shape[1] refers to columns
45
              count1,count0=0,0
              for j in range (train rows):
46
47
                 if test[t,k] == train[j,k] and train[j,train col-1]=='no':
48
                     count0+=1
                 if test[t,k]==train[j,k] and train[j,train col-1]=='yes':
49
50
                     count1+=1
              prob0[k]=count0/countNo
51
52
              prob1[k]=count1/countYes
53
       probno=probNo
        probyes=probYes
54
        for i in range(test_col-1):
55
56
            probno=probno*prob0[i]
57
            probyes=probyes*prob1[i]
58
            if probno>probyes:
59
                 predict='no'
60
            else:
                 predict='yes'
61
            if predict == test[t,test col-1]:
62
63
                 accuracy+=1
        final accuracy=(accuracy/test rows)*100
64
65
        print("accuracy",final_accuracy,"%")
        return
67 metadata, traindata = read data("data3.csv")
68 splitRatio=0.8
69 trainingset, testset=splitDataset(traindata, splitRatio)
70 training=np.array(trainingset)
71 testing=np.array(testset)
72 print("Training :\n",training,"\nTesting: \n",testing)
73 classify(training, testing)
```

The output is as follows:

```
[['sunny', 'hot', 'high', 'weak', 'no'], ['sunny', 'hot', 'high', 'strong', 'no'],
 ['overcast', ' hot', ' high', 'weak', 'yes'], ['rain', 'mild', ' high', 'weak',
 ['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']]
 Metadata is:
  ['Outlook', 'Temperature', 'Humidity', 'Wind', 'Target']
Training :
 [['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']]
Testing:
 [['overcast' 'mild' 'high' 'strong' 'yes']
 ['overcast' 'hot' 'normal' 'weak' 'yes']
 ['rain' 'mild' 'high' 'strong' 'no']]
training data size= 11
test data size= 3
target count probability
         7
              0.6363636363636364
Yes
No 4
                   0.36363636363636365
```

7. 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.

```
1 import numpy as np
 2 import matplotlib.pyplot as plt
 3 import pandas as pd
 4 from sklearn.cluster import KMeans
 5 from sklearn.mixture import GaussianMixture
 6 df1 = pd.read csv("data8.csv")
 7 print(df1)
 8 f1 = df1['Distance Feature'].values
 9 f2 = df1['Speeding Feature'].values
10 X = np.matrix(list(zip(f1,f2)))
11 plt.plot(1)
12 plt.subplot(511)
13 plt.xlim([0, 100])
14 plt.ylim([0, 50])
15 plt.title('Dataset')
16 plt.ylabel('speeding feature')
17 plt.xlabel('distance feature')
18 plt.scatter(f1,f2)
19 colors = ['b', 'g', 'r']
20 markers = ['o', 'v', 's']
21 # create new plot and data for K- means algorithm
22 plt.plot(2)
23 ax=plt.subplot(513)
24 kmeans model = KMeans(n clusters=3).fit(X)
25 for i, l in enumerate(kmeans model.labels ):
      plt.plot(f1[i], f2[i], color=colors[1],marker=markers[1])
27 plt.xlim([0, 100])
28 plt.ylim([0, 50])
29 plt.title('K- Means')
30 plt.ylabel('speeding feature')
31 plt.xlabel('distance feature')
```

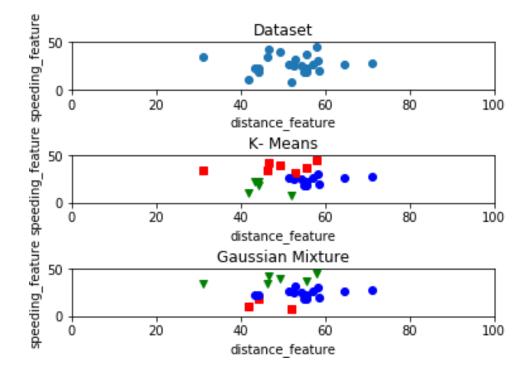
```
32 # create new plot and data for gaussian mixture i.e. EM Algorithm
33 plt.plot(3)
34 plt.subplot(515)
35 gmm=GaussianMixture(n_components=3).fit(X)
36 labels= gmm.predict(X)
37 for i, l in enumerate(labels):
38    plt.plot(f1[i], f2[i], color=colors[l], marker=markers[l])
39 plt.xlim([0, 100])
40 plt.ylim([0, 50])
41 plt.title('Gaussian Mixture')
42 plt.ylabel('speeding_feature')
43 plt.xlabel('distance_feature')
44 plt.show()
```

The dataset to be considered is:

1	Driver_ID	Distance_Feature	Speeding_Feature
2	3423311935	71.24	28
3	3423313212	52.53	25
4	3423313724	64.54	27
5	3423311373	55.69	22
6	3423310999	54.58	25
7	3423313857	41.91	10
8	3423312432	58.64	20
9	3423311434	52.02	8
10	3423311328	31.25	34
11	3423312488	44.31	19
12	3423311254	49.35	40
13	3423312943	58.07	45
14	3423312536	44.22	22
15	3423311542	55.73	19
16	3423312176	46.63	43
17	3423314176	52.97	32
18	3423314202	46.25	35
19	3423311346	51.55	27
20	3423310666	57.05	26
21	3423313527	58.45	30
22	3423312182	43.42	23
23	3423313590	55.68	37
24	3423312268	55.15	18

The output is as follows:

	_	Distance_Feature		Unnamed: 3
0	3423311935	71.24	28	NaN
1	3423313212	52.53	25	NaN
2	3423313724	64.54	27	NaN
3	3423311373	55.69	22	NaN
4	3423310999	54.58	25	NaN
5	3423313857	41.91	10	NaN
6	3423312432	58.64	20	NaN
7	3423311434	52.02	8	NaN
8	3423311328	31.25	34	NaN
9	3423312488	44.31	19	NaN
10	3423311254	49.35	40	NaN
11	3423312943	58.07	45	NaN
12	3423312536	44.22	22	NaN
13	3423311542	55.73	19	NaN
14	3423312176	46.63	43	NaN
15	3423314176	52.97	32	NaN
16	3423314202	46.25	35	NaN
17	3423311346	51.55	27	NaN
18	3423310666	57.05	26	NaN
19	3423313527	58.45	30	NaN
20	3423312182	43.42	23	NaN
21	3423313590	55.68	37	NaN
22	3423312268	55.15	18	NaN



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

Program:

```
1 from sklearn import datasets
 2 iris=datasets.load iris()
 3 iris data=iris.data
 4 iris labels=iris.target
 5 from sklearn.model selection import train test split
 6 x_train,x_test,y_train,y_test=train_test_split(iris_data,iris_labels,test_size=0.30)
 8 from sklearn.neighbors import KNeighborsClassifier
 9 classifier=KNeighborsClassifier(n neighbors=5)
10 classifier.fit(x train,y train)
11 y pred=classifier.predict(x test)
12
13 from sklearn.metrics import classification report, confusion matrix
14 print('Confusion matrix is as follows')
15 print(confusion matrix(y test,y pred))
16 print('Accuracy Matrics')
17 print(classification report(y test,y pred))
```

The data set to be considered is:

Iris Flower dataset from SKLearn is imported.

The output is as follows:

```
Confusion matrix is as follows
[[16 0 0]
[0193]
[0 1 6]]
Accuracy Matrics
            precision recall f1-score support
                1.00
         0
                         1.00
                                  1.00
                                            16
                0.95
                         0.86
                                  0.90
                                            22
         1
         2
                0.67
                         0.86
                                  0.75
                                            7
  micro avg
               0.91
                        0.91
                                 0.91
                                            45
  macro avg
               0.87
                        0.91
                                 0.88
                                            45
weighted avg
               0.92
                         0.91
                                  0.91
                                            45
```

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

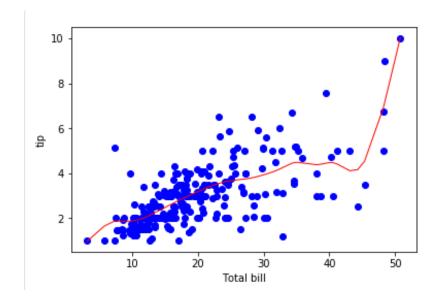
```
1 import matplotlib.pyplot as plt
 2 import pandas as pd
 3 import numpy as np1
 4 def kernel(point, xmat, k):
      m,n=np1.shape(xmat) #size of matrix m
      weights=np1.mat(np1.eye(m)) #np.eye returns mat with 1 in the diagonal
 6
 7
      for j in range(m):
 8
          diff=point-xmat[j]
          weights[j,j]=np1.exp(diff*diff.T/(-2.0*k**2))
 9
      return weights
10
11 def localWeight(point,xmat,ymat,k):
      wei=kernel(point,xmat,k)
12
13
      W=(xmat.T*(wei*xmat)).I*(xmat.T*(wei*ymat.T))
      return W
14
15 def localWeightRegression(xmat,ymat,k):
      row,col=np1.shape(xmat) #return 244 rows and 2 columns
16
      ypred=np1.zeros(row)
17
      for i in range(row):
18
          ypred[i]=xmat[i]*localWeight(xmat[i],xmat,ymat,k)
19
20
      return ypred
21 data=pd.read csv('data10.csv')
22 bill=np1.array(data.total bill)
23 tip=np1.array(data.tip)
24 mbill=np1.mat(bill)
25 mtip=np1.mat(tip)
26 mbillMatCol=np1.shape(mbill)[1] # 1 for vertical i.e columns
27 onesArray=np1.mat(np1.ones(mbillMatCol))
28 #hstack concate horizontal lists it takes one value from the fist and one from the second
29 xmat=np1.hstack((onesArray.T,mbill.T))
30 ypred=localWeightRegression(xmat,mtip,2)
31 SortIndex=xmat[:,1].argsort(0)
32 #argsort take the index of each and sort them according to the orginal value
33 xsort=xmat[SortIndex][:,0]
34 fig= plt.figure()
35 ax=fig.add subplot(1,1,1)
36 ax.scatter(bill,tip,color='blue')
37 ax.plot(xsort[:,1],ypred[SortIndex],color='red',linewidth=1)
38 plt.xlabel('Total bill')
39 plt.ylabel('tip')
40 plt.show();
```

The dataset considered is:

The overall dataset consists of over 200 hypothesis values. Out of this First 24 are given below:

-		
1	total_bill	tip
2	16.99	1.01
3	10.34	1.66
4	21.01	3.5
5	23.68	3.31
6	24.59	3.61
7	25.29	4.71
8	8.77	2
9	26.88	3.12
10	15.04	1.96
11	14.78	3.23
12	10.27	1.71
13	35.26	5
14	15.42	1.57
15	18.43	3
16	14.83	3.02
17	21.58	3.92
18	10.33	1.67
19	16.29	3.71
20	16.97	3.5
21	20.65	3.35
22	17.92	4.08
23	20.29	2.75
24	15.77	2.23
25	39.42	7.58

The output is:



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