## 1. FIND S

```
import pandas as pd
import numpy as np
data = pd.read_csv("2_1.csv")
print("Data:\n", data, "\n")
data array = np.array(data)
d = data_array[:, :-1]
print("The attributes are:\n", d, "\n")
target = data array[:, -1]
print("The target is:\n", target, "\n")
def train(c, t):
  specific hypothesis = None
  for i, val in enumerate(t):
    if val == "Yes":
       specific_hypothesis = c[i].copy()
       break
  if specific_hypothesis is None:
    return "No positive examples found."
  for i, val in enumerate(c):
    if t[i] == "Yes":
       for x in range(len(specific_hypothesis)):
         if val[x] != specific_hypothesis[x]:
           specific hypothesis[x] = '?'
  return specific_hypothesis
final hypothesis = train(d, target)
print("The final hypothesis is:\n", final hypothesis)
```

## 2. CANDIDATE ELIMINATION

```
import numpy as np
import pandas as pd
data = pd.read csv("2 1.csv")
concepts = np.array(data.iloc[:,0:-1])
print("\nInstances are:\n",concepts)
target = np.array(data.iloc[:,-1])
print("\nTarget Values are: ",target)
def learn(concepts, target):
  specific_h = concepts[0].copy()
  print("\nInitialization of specific_h and genearal h")
  print("\nSpecific Boundary: ", specific h)
  general h = [["?" for i in range(len(specific h))] for i in range(len(specific h))]
  print("\nGeneric Boundary: ",general_h)
  for i, h in enumerate(concepts):
    print("\nInstance", i+1, "is ", h)
    if target[i] == "Yes":
       print("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 Bundary 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")
```

```
3. ID3 DECISION TREE
data = pd.read csv("ws1.csv")
print("Columns in the dataset:", data.columns)
if 'answer' not in data.columns:
  raise KeyError("The 'answer' column is not found in the dataset. Please check
the column names.")
features = [feat for feat in data.columns if feat != "answer"]
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["answer"] == "yes":
      pos += 1
    else:
      neg += 1
  if pos == 0.0 or neg == 0.0:
    return 0.0
  else:
    p = pos / (pos + neg)
    n = neg / (pos + neg)
    return -(p * math.log(p, 2) + n * math.log(n, 2))
```

```
def info gain(examples, attr):
  uniq = np.unique(examples[attr])
  gain = entropy(examples)
  for u in uniq:
    subdata = examples[examples[attr] == u]
    sub_e = entropy(subdata)
    gain -= (float(len(subdata)) / float(len(examples))) * sub_e
  return gain
def ID3(examples, attrs):
  root = Node()
  max_gain = -1
  max feat = None
  for feature in attrs:
    gain = info_gain(examples, feature)
    if gain > max_gain:
      max gain = gain
      max_feat = feature
  if max_feat is None:
    return None
  root.value = max_feat
  uniq = np.unique(examples[max_feat])
  for u in uniq:
    subdata = examples[examples[max feat] == u]
    if entropy(subdata) == 0.0:
      newNode = Node()
      newNode.isLeaf = True
```

```
newNode.value = u
      newNode.pred = np.unique(subdata["answer"])[0]
      root.children.append(newNode)
    else:
      dummyNode = Node()
      dummyNode.value = u
      new_attrs = [attr for attr in attrs if attr != max_feat]
      child = ID3(subdata, new_attrs)
      if child: # Only add the child if it's not None
         dummyNode.children.append(child)
      root.children.append(dummyNode)
  return root
def printTree(root: Node, depth=0):
  if root is None:
    return
  for i in range(depth):
    print("\t", end="")
  print(root.value, end="")
  if root.isLeaf:
    print(" -> ", root.pred)
  else:
    print()
  for child in root.children:
    printTree(child, depth + 1)
root = ID3(data, features)
printTree(root)
```

```
4. BACK PROPOGATION
import numpy as np
X = np.array(([2, 9], [1, 5], [3, 6]), dtype=float)
y = np.array(([92], [86], [89]), dtype=float)
X = X/np.amax(X,axis=0)
y = y/100
def sigmoid (x):
  return 1/(1 + np.exp(-x))
def derivatives sigmoid(x):
  return x * (1 - x)
epoch=5
Ir=0.1
inputlayer neurons = 2
hiddenlayer neurons = 3
output_neurons = 1
wh=np.random.uniform(size=(inputlayer neurons,hiddenlayer neurons))
bh=np.random.uniform(size=(1,hiddenlayer_neurons))
wout=np.random.uniform(size=(hiddenlayer_neurons,output_neurons))
bout=np.random.uniform(size=(1,output_neurons))
for i in range(epoch):
  hinp1=np.dot(X,wh)
  hinp=hinp1 + bh
  hlayer act = sigmoid(hinp)
  outinp1=np.dot(hlayer act,wout)
  outinp= outinp1+bout
  output = sigmoid(outinp)
```

```
EO = y-output
  outgrad = derivatives sigmoid(output)
  d_output = EO * outgrad
  EH = d_output.dot(wout.T)
  hiddengrad = derivatives sigmoid(hlayer act)
  d hiddenlayer = EH * hiddengrad
  wout += hlayer_act.T.dot(d_output) *Ir
  wh += X.T.dot(d hiddenlayer) *Ir
  print ("------Epoch-", i+1, "Starts-----")
  print("Input: \n" + str(X))
  print("Actual Output: \n" + str(y))
  print("Predicted Output: \n",output)
  print ("--------Epoch-", i+1, "Ends-----\n")
print("Input: \n" + str(X))
print("Actual Output: \n" + str(y))
print("Predicted Output: \n", output)
```

```
5. NAIVE BAYESIAN
```

import numpy as np

import matplotlib.pyplot as plt

import pandas as pd

import seaborn as sns

dataset = pd.read csv("naive.csv")

X = dataset.iloc[:, [0,1]].values

y = dataset.iloc[:, 2].values

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

X\_train, X\_test, y\_train, y\_test = train\_test\_split(X,y,test\_size= 0.25, random state=0)

# importing standard scaler

from sklearn.preprocessing import StandardScaler

sc X = StandardScaler()

X\_train = sc\_X.fit\_transform(X\_train)

X\_test = sc\_X.fit\_transform(X\_test)

from sklearn.naive\_bayes import BernoulliNB

from sklearn.naive\_bayes import GaussianNB

classifer1 = GaussianNB()

classifer1.fit(X\_train, y\_train)

y\_pred1 = classifer1.predict(X\_test)

from sklearn.metrics import accuracy\_score

print(accuracy\_score(y\_test,y\_pred1))

```
6. NAIVE BAYESAIN - CLASSIFICATION
import matplotlib.pyplot as plt
import seaborn as sns
dataset = pd.read csv("naive.csv")
X = dataset.iloc[:, [0,1]].values
y = dataset.iloc[:, 2].values
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X,y,test_size= 0.25,
random state=0)
from sklearn.preprocessing import StandardScaler
sc X = StandardScaler()
X train = sc X.fit transform(X train)
X test = sc X.fit transform(X test)
from sklearn.naive bayes import BernoulliNB
from sklearn.naive bayes import GaussianNB
classifer1 = GaussianNB(
classifer1.fit(X_train, y_train)
y pred1 = classifer1.predict(X test)
from sklearn.metrics import accuracy score
print(accuracy_score(y_test,y_pred1))
from sklearn.metrics import accuracy score, confusion matrix,
precision score, recall score
print('Accuracy Metrics: \n')
print('Accuracy: ', accuracy score(y test, y pred1))
print('Recall: ', recall score(y test, y pred1))
print('Precision: ', precision_score(y_test, y_pred1))
print('Confusion Matrix: \n', confusion matrix(y test, y pred1))
```

```
7. BAYESIAN NETWORK
import numpy as np
import pandas as pd
import csv
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
heartDisease = pd.read csv('7.csv')
heartDisease = heartDisease.replace('?',np.nan)
print('Sample instances from the dataset are given below')
print(heartDisease.head())
print('\n Attributes and datatypes')
print(heartDisease.dtypes)
model=
BayesianModel([('age','heartdisease'),('gender','heartdisease'),('exang','heartdi
sease'),('cp','heartdisease'),('heartdisease','restecg'),('heartdisease','chol')])
print('\nLearning CPD using Maximum likelihood estimators')
model.fit(heartDisease,estimator=MaximumLikelihoodEstimator)
print('\n Inferencing with Bayesian Network:')
HeartDiseasetest infer = VariableElimination(model)
print('\n 1. Probability of HeartDisease given evidence= restecg')
q1=HeartDiseasetest infer.query(variables=['heartdisease'],evidence={'restecg'
:1})
print(q1)
print('\n 2. Probability of HeartDisease given evidence= cp ')
q2=HeartDiseasetest infer.guery(variables=['heartdisease'],evidence={'cp':2})
print(q2)
```

## 8. K MEANS AND EM ALGORITHM

```
from sklearn.cluster import KMeans
from sklearn.mixture import GaussianMixture
import sklearn.metrics as metrics
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
names = ['Sepal Length', 'Sepal Width', 'Petal Length', 'Petal Width', 'Class']
dataset = pd.read_csv("8.csv", names=names)
X = dataset.iloc[:, :-1]
label = {'Iris-setosa': 0,'Iris-versicolor': 1, 'Iris-virginica': 2}
y = [label[c] for c in dataset.iloc[:, -1]]
plt.figure(figsize=(14,7))
colormap=np.array(['red','lime','black']
plt.subplot(1,3,1)
plt.title('Real')
plt.scatter(X.Petal Length, X.Petal Width, c=colormap[y])
model=KMeans(n clusters=3, random state=0).fit(X)
plt.subplot(1,3,2)
plt.title('KMeans')
plt.scatter(X.Petal Length, X.Petal Width, c=colormap[model.labels ])
print('The accuracy score of K-Mean: ',metrics.accuracy_score(y,
model.labels ))
print('The Confusion matrixof K-Mean:\n',metrics.confusion matrix(y,
model.labels ))
gmm=GaussianMixture(n components=3, random state=0).fit(X)
y cluster gmm=gmm.predict(X)
plt.subplot(1,3,3)
plt.title('GMM Classification')
plt.scatter(X.Petal_Length,X.Petal_Width,c=colormap[y_cluster_gmm])
print('The accuracy score of EM: ',metrics.accuracy_score(y, y_cluster_gmm))
print('The Confusion matrix of EM:\n',metrics.confusion matrix(y,
y cluster_gmm))
```

```
9. KNN
```

```
import numpy as np
import pandas as pd
from sklearn.neighbors import KNeighborsClassifier
from sklearn.model selection import train test split
from sklearn import metrics
from sklearn.datasets import load iris
iris = load iris()
names = ['sepal-length', 'sepal-width', 'petal-length', 'petal-width', 'Class']
df = pd.DataFrame(iris.data,columns=iris.feature names)
df['target'] = iris.target
X = df.iloc[:, :-1]
y = df.iloc[:, -1]
print(X.head())
Xtrain, Xtest, ytrain, ytest = train test split(X, y, test size=0.10)
classifier = KNeighborsClassifier(n neighbors=5).fit(Xtrain, ytrain)
vpred = classifier.predict(Xtest)
i = 0
print ("\n-----")
print ('%-25s %-25s %-25s' % ('Original Label', 'Predicted Label',
'Correct/Wrong'))
print ("-----")
for label in ytest:
 print ('%-25s %-25s' % (label, ypred[i]), end="")
 if (label == ypred[i]):
   print (' %-25s' % ('Correct'))
 else:
   print (' %-25s' % ('Wrong'))
 i = i + 1
print ("-----")
print("\nConfusion Matrix:\n",metrics.confusion matrix(ytest, ypred))
print ("-----")
print("\nClassification Report:\n",metrics.classification report(ytest, ypred))
print ("-----")
print('Accuracy of the classifer is %0.2f' % metrics.accuracy score(ytest,ypred))
print ("-----")
```

```
11. 8 QUEENS
use_module(library(clpfd)).
n_queens(N, Qs):-
      length(Qs, N),
      Qs ins 1..N,
      safe_queens(Qs).
safe_queens([]).
safe queens([Q|Qs]):-
      safe_queens(Qs, Q, 1),
      safe_queens(Qs).
safe_queens([], _, _).
safe_queens([Q|Qs], Q0, D0) :-
      Q0 \# Q
      abs(Q0 - Q) \#= D0,
      D1 #= D0 + 1,
      safe_queens(Qs, Q0, D1).
Query:
queens(8, Qs), labeling([ff], Qs).
12.DFS
% solve( Node, Solution):
% Solution is an acyclic path (in reverse order) between Node and a goal
solve(Node, Solution):-
 depthfirst([], Node, Solution).
% depthfirst( Path, Node, Solution):
% extending the path [Node | Path] to a goal gives Solution
depthfirst( Path, Node, [Node | Path] ) :-
 goal( Node).
```

```
depthfirst( Path, Node, Sol) :-
 s(Node, Node1),
 \+ member( Node1, Path),
                                     % Prevent a cycle
 depthfirst( [Node | Path], Node1, Sol).
depthfirst2( Node, [Node], _):-
 goal( Node).
depthfirst2( Node, [Node | Sol], Maxdepth) :-
 Maxdepth > 0,
 s(Node, Node1),
 Max1 is Maxdepth - 1,
 depthfirst2( Node1, Sol, Max1).
goal(f).
goal(j).
s(a,b).
s(a,c).
s(b,d).
s(b,e).
s(c,f).
s(c,g).
s(d,h).
s(e,i).
s(e,j).
```

```
13. BFS
import heapq
class Node:
def init (self, state, parent, cost, heuristic):
      self.state = state
      self.parent = parent
      self.cost = cost
      self.heuristic = heuristic
def It (self, other):
      return self.heuristic < other.heuristic
def best_first_search(start, goal, heuristic_fn, get_neighbors_fn):
      open list = []
      closed list = set()
      start_node = Node(start, None, 0, heuristic_fn(start, goal))
      heapq.heappush(open_list, start_node)
      while open list:
            current_node = heapq.heappop(open_list)
            if current node.state == goal:
                   return reconstruct path(current node)
            closed list.add(current node.state)
      for neighbor, cost in get_neighbors_fn(current_node.state):
            if neighbor in closed list:
                   continue
            neighbor_node = Node(neighbor, current_node,
            current node.cost + cost, heuristic fn(neighbor, goal))
            for open node in open list:
                   if open node.state == neighbor and open node.cost <=
neighbor node.cost:
```

```
break
                   else:
                         heapq.heappush(open_list, neighbor_node)
      return None
def reconstruct path(node):
      path = []
      while node:
            path.append(node.state)
            node = node.parent
      return path[::-1]
def manhattan_distance(state, goal):
      return abs(state[0] - goal[0]) + abs(state[1] - goal[1])
def get_neighbors(state):
      neighbors = []
      x, y = state
      moves = [(-1, 0), (1, 0), (0, -1), (0, 1)]
      for move in moves:
            neighbor = (x + move[0], y + move[1])
            if 0 <= neighbor[0] < 5 and 0 <= neighbor[1] < 5:
                   neighbors.append((neighbor, 1))
            return neighbors
```

path = best\_first\_search(start, goal, manhattan\_distance, get\_neighbors)

start = (0, 0)

goal = (4, 4)

print("Path found:", path)

```
14. 8 PUZZLE
ids:-
 start(State),
 length(Moves, N),
 dfs([State], Moves, Path), !,
 show([start|Moves], Path),
 format('\simnmoves = \simw\simn', [N]).
dfs([State|States], [], Path):-
 goal(State), !,
 reverse([State|States], Path).
dfs([State|States], [Move|Moves], Path):-
 move(State, Next, Move),
 not(memberchk(Next, [State|States])),
 dfs([Next,State|States], Moves, Path).
show([], _).
show([Move|Moves], [State|States]):-
 State = state(A,B,C,D,E,F,G,H,I),
 format('~n~w~n~n', [Move]),
 format('~w ~w ~w~n',[A,B,C]),
 format('~w ~w ~w~n',[D,E,F]),
 format('~w ~w ~w~n',[G,H,I]),
 show(Moves, States).
% Empty position is marked with '*'
start( state(6,1,3,4,*,5,7,2,0) ).
goal( state(*,0,1,2,3,4,5,6,7) ).
move( state(*,B,C,D,E,F,G,H,J), state(B,*,C,D,E,F,G,H,J), right).
move(state(*,B,C,D,E,F,G,H,J), state(D,B,C,*,E,F,G,H,J), down).
```

## 15. TRAVELLING SALESMAN

```
Production Rules:-
route(Town1,Town2,Distance) road(Town1,Town2,Distance).
route(Town1,Town2,Distance)
road(Town1,X,Dist1),route(X,Town2,Dist2),Distance=Dist1+Dist2,
domains
town = symbol
distance = integer
predicates
nondeterm road(town,town,distance)
nondeterm route(town,town,distance)
clauses
road("tampa", "houston", 200).
road("gordon","tampa",300).
road("houston", "gordon", 100).
road("houston", "kansas city", 120).
road("gordon","kansas_city",130).
route(Town1,Town2,Distance):-
road(Town1,Town2,Distance).
route(Town1,Town2,Distance):-
road(Town1,X,Dist1),
route(X,Town2,Dist2),
Distance=Dist1+Dist2,!.
```