## Ex. 1 FIND-S algorithm

## Aim:

Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples. Read the training data from a .CSV file.

```
import pandas as pd
import numpy as np
#to read the data in the csv file
data = pd.read_csv("C://users/siva/sport.csv")
print(data,"n")
#making an array of all the attributes
d = np.array(data)[:,:-1]
print("n The attributes are: ",d)
#segragating the target that has positive and negative examples
target = np.array(data)[:,-1]
print("n The target is: ",target)
#training function to implement find-s algorithm
def train(c,t):
  for i, val in enumerate(t):
     if val == "Yes":
       specific_hypothesis = c[i].copy()
       break
  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] = '?'
    else:
        pass

return specific_hypothesis

#obtaining the final hypothesis
print("n The final hypothesis is:",train(d,target))
```

```
Sky Temp Humidity Wind Water Forecast EnjoySport
0 1 Sunny Warm Normal Strong Warm
                                              Same
                                                         Yes
                     High Strong Warm
1 2 Sunny Warm
                                             Same
                                                       Yes
2 3 Rainy Cold
                   High Strong Warm Change
                                                       No
3 4 Sunny Warm
                     High Strong Cool Change
                                                       Yes n
n The attributes are: [[1 'Sunny ' 'Warm ' 'Normal ' 'Strong ' 'Warm ' 'Same ']
[2 'Sunny ' 'Warm ' 'High ' 'Strong ' 'Warm ' 'Same ']
[3 'Rainy ' 'Cold ' 'High ' 'Strong ' 'Warm ' 'Change ']
[4 'Sunny ' 'Warm ' 'High ' 'Strong ' 'Cool ' 'Change ']]
n The target is: ['Yes' 'Yes' 'No' 'Yes']
n The final hypothesis is: ['?' 'Sunny ' 'Warm ' '?' 'Strong ' '?' '?']
```

## Ex.2 Candidate-Elimination algorithm

**Aim:**For a given set of training data examples stored in a .CSV file, implement and demonstr ate the Candidate-Elimination algorithm to output a description of the set of all hypotheses consistent with the training examples.

## Algorithm:

## Code:

import numpy as np import pandas as pd

```
data = pd.read_csv("E:\Goms_Academic\AI & ML LAB\sport new.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")
```

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 Values are: ['yes' 'yes' 'no' 'yes']
Initialization of specific h and genearal h
Specific Boundary: ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
Generic Boundary: [['?', '?', '?', '?', '?'], ['?', '?', '?', '?'
, '?', '?'], ['\bar{2}', '?', '?', '?', '?'], ['?', '?', '?', '?', '?',
'?'], ['?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
Instance 1 is ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
Instance is Positive
Specific Bundary after 1 Instance is ['sunny' 'warm' 'normal' 'strong
' 'warm' 'same']
Generic Boundary after 1 Instance is [['?', '?', '?', '?', '?'],
?!, !?!, !?!, !?!, !?!], [!?!, !?!, !?!, !?!, !?!, !?!], [!?!, !?!, !?!
, '?', '?', '?']]
Instance 2 is ['sunny' 'warm' 'high' 'strong' 'warm' 'same']
Instance is Positive
Specific Bundary after 2 Instance is ['sunny' 'warm' '?' 'strong' 'wa
rm' 'same']
Generic Boundary after 2 Instance is [['?', '?', '?', '?', '?'],
[131, 131, 131, 131, 131, 131], [131, 131, 131, 131, 131, 131], [131, 1
?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?'], ['?', '?', '?'
, '?', '?', '?']]
Instance 3 is ['rainy' 'cold' 'high' 'strong' 'warm' 'change']
Instance is Negative
Specific Bundary after 3 Instance is ['sunny' 'warm' '?' 'strong' 'wa
rm' 'same']
Generic Boundary after 3 Instance is [['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?'], ['?', '?', '?', '?', '?'],
[131, 131, 131, 131, 131, 131], [131, 131, 131, 131, 131, 131], [131, 1
?', '?', '?', '?', 'same']]
Instance 4 is ['sunny' 'warm' 'high' 'strong' 'cool' 'change']
Instance is Positive
Specific Bundary 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', '?', '?', '?', '?']]
```

#### Ex.3Working of decision tree based ID3 algorithm

#### Aim:

Write a program to demonstrate the working of the decision tree based ID3 algorithm. Use an appro priate data set for building the decision tree and apply this knowledge to classify a new sample.

```
Code:
import pandas as pd
import math
import numpy as np
data = pd.read_csv("3-dataset.csv")
features = [feat for feat in data]
features.remove("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])
  #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["answer"])
       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)
Result:
outlook
        overcast -> ['yes']
        rain
                wind
                         strong -> ['no']
                         weak -> ['yes']
        sunny
                humidity
                         high -> ['no']
                         normal -> ['yes']
```

## Ex. 4 Back propagation algorithm

**Aim:**Build an Artificial Neural Network by implementing the Back propagation algorithm and test the same using appropriate data sets

```
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=5 #Setting training iterations
lr=0.1 #Setting learning rate
inputlayer neurons = 2 #number of features in data set
hiddenlayer_neurons = 3 #number of hidden layers neurons
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)
  hiddengrad = derivatives sigmoid(hlayer act)#how much hidden layer wts contributed to e
rror
  d_hiddenlayer = EH * hiddengrad
  wout += hlayer_act.T.dot(d_output) *lr # dotproduct of nextlayererror and currentlayerop
  wh += X.T.dot(d hiddenlayer) *lr
  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)
```

```
-----Epoch- 1 Starts-----
Input:
[[0.66666667 1. ]
[0.33333333 0.55555556]
           0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.81361748]
[0.80545255]
[0.80887549]]
-----Epoch- 1 Ends-----
-----Epoch- 2 Starts-----
Input:
[[0.66666667 1.
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.81464174]
[0.80640982]
[0.80987396]]
-----Epoch- 2 Ends-----
-----Epoch- 3 Starts-----
Input:
[[0.66666667 1.
[0.33333333 0.55555556]
          0.66666667]]
[1.
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.81564531]
[0.8073482]
[0.81085253]]
-----Epoch- 3 Ends-----
-----Epoch- 4 Starts-----
```

```
Input:
[[0.66666667 1.
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.81662881]
 [0.80826822]
[0.81181177]]
-----Epoch- 4 Ends-----
-----Epoch- 5 Starts-----
Input:
[[0.66666667 1.
[0.33333333 0.55555556]
           0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.81759282]
[0.80917043]
[0.81275225]]
-----Epoch- 5 Ends-----
Input:
[[0.66666667 1.
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
 [[0.81759282]
 [0.80917043]
 [0.81275225]]
```

## Naive Bayesian Classifier

**Aim:** Write a program to implement the Naive Bayesian Classifier for a sample training data set stor ed as a .CSV file. Compute the accuracy of the classifier, considering few test data sets.

```
Code:
```

```
# importing the libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
# importing the dataset
dataset = pd.read_csv("D://NaiveBayes.csv")
# split the data into inputs and outputs
X = dataset.iloc[:, [0,1]].values
y = dataset.iloc[:, 2].values
# training and testing data
from sklearn.model_selection import train_test_split
# assign test data size 25%
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
# scalling the input data
sc X = StandardScaler()
X_train = sc_X.fit_transform(X_train)
X_{test} = sc_X.fit_{transform}(X_{test})
# importing classifier
from sklearn.naive_bayes import BernoulliNB
# import Gaussian Naive Bayes classifier
from sklearn.naive bayes import GaussianNB
# create a Gaussian Classifier
classifer1 = GaussianNB()
# training the model
classifer1.fit(X_train, y_train)
# testing the model
y pred1 = classifer1.predict(X test)
# importing accuracy score
from sklearn.metrics import accuracy_score
# printing the accuracy of the model
print(accuracy score(y test,y pred1))
Output:
0.91
```

## Ex. 6 NaiveBayesian Classifier

#### Aim:

By assuming a set of documents that need to be classified, use the naive Bayesian classifier model to perform this task. Built in java classes / API can be used to write the program. Calculate the accuracy, precision and recall for your data set.

#### Code:

# importing the libraries

import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
# importing the dataset
dataset = pd.read\_csv("NaiveBayes.csv")

# split the data into inputs and outputs
X = dataset.iloc[:, [0,1]].values
y = dataset.iloc[:, 2].values
# training and testing data

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

# assign test data size 25%

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

# scalling the input data

sc\_X = StandardScaler()

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

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

# importing classifier

from sklearn.naive\_bayes import BernoulliNB

# import Gaussian Naive Bayes classifier from sklearn.naive\_bayes import GaussianNB

# create a Gaussian Classifier classifer1 = GaussianNB()

# training the model
classifer1.fit(X\_train, y\_train)

```
# testing the model
y_pred1 = classifer1.predict(X_test)
# importing accuracy score
from sklearn.metrics import accuracy_score

# printing the accuracy of the model
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 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))

Output:
0.91
Accuracy Metrics:
```

Accuracy: 0.91 Recall: 0.84375

Confusion Matrix:

[[64 4] [5 27]]

Precision: 0.8709677419354839

#### Ex. 7

#### **Bayesian network**

#### Aim:

Write a program to construct a Bayesian network considering medical data. Use this model to demo nstrate the diagnosis of heart patients using standard Heart Disease Data Set. You can use Java/Pyth on ML library classes/API.

## Code:

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('Exp 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', 'heartdisease'),('
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.query(variables=['heartdisease'],evidence={'cp':2})
print(q2)
Output:
Sample instances from the dataset are given below
   age gender cp trestbps chol fbs restecg thalach
                                                                             oldpe
                                                                     exang
ak
   \
0
    63
               1 1
                             145
                                    233
                                            1
                                                       2
                                                                150
                                                                          0
                                                                                   2
.3
1
    67
               1
                    4
                             160
                                    286
                                             0
                                                       2
                                                                108
                                                                          1
                                                                                   1
.5
2
    67
               1
                    4
                             120
                                    229
                                            0
                                                       2
                                                                129
                                                                          1
                                                                                   2
.6
    37
               1
                    3
                             130
                                                       0
                                                                          0
                                                                                   3
3
                                    250
                                            0
                                                                187
. 5
4
    41
               0
                    2
                             130
                                    204
                                            0
                                                       2
                                                                172
                                                                          0
. 4
   slope ca thal heartdisease
0
        3 0 6
        2 3
                                  2
1
                 3
                 7
2
        2 2
                                  1
        3 0
3
                 3
                                  0
Attributes and datatypes
gender
                     int64
                     int64
ср
trestbps
                     int64
chol
                     int64
                     int64
fbs
                    int64
restecg
                    int64
thalach
exang
                    int64
```

oldpeak float64
slope int64
ca object
thal object
heartdisease int64

dtype: object

Learning CPD using Maximum likelihood estimators

Inferencing with Bayesian Network:

1. Probability of HeartDisease given evidence= restecg

heartdisease	phi(heartdisease)    -===================================
heartdisease(0)	0.1012
heartdisease(1)	0.0000
heartdisease(2)	0.2392
heartdisease(3)	0.2015
heartdisease(4)	0.4581

2. Probability of HeartDisease given evidence= cp

+	++
heartdisease	phi(heartdisease)   
heartdisease(0)	0.3610
heartdisease(1)	0.2159
heartdisease(2)	0.1373
heartdisease(3)	0.1537
heartdisease(4)	0.1321

## **Result:**

## Ex. 8

## **EM Algorithm and K-Means Algorithm**

Aim: To apply EM algorithm to cluster a set of data stored in a .csv file. Use the same dataset for clustering using k-means algorithm.

## Code:

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-dataset.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'])
# REAL PLOT
plt.subplot(1,3,1)
plt.title('Real')
plt.scatter(X.Petal_Length,X.Petal_Width,c=colormap[y])
# K-PLOT
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 matrix of K-Mean:\n',metrics.confusion matrix(y, model.labels ))
# GMM PLOT
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))
Output:
The accuracy score of K-Mean: 0.09333333333333333
The Confusion matrix of K-Mean:
 [[ 0 50 0]
 [ 2 0 48]
 [36 0 14]]
The accuracy score of EM: 0.9666666666666667
The Confusion matrix of EM:
```

```
[[50 0 0]
[ 0 45 5]
[ 0 0 50]]
```

#### Ex. 9

#### k-Nearest Neighbour

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

```
Code:
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']
# Read dataset to pandas dataframe
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)
ypred = 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 ("-----")
Output:
  sepal length (cm) sepal width (cm) petal length (cm) petal width
(cm)
0
                5.1
                                 3.5
                                                    1.4
0.2
                4.9
                                 3.0
1
                                                    1.4
0.2
                4.7
                                3.2
2
                                                    1.3
0.2
3
                4.6
                                3.1
                                                    1.5
0.2
                5.0
                                 3.6
                                                    1.4
0.2
Original Label Predicted Label Correct/Wrong
0
                         0
                                                  Correct
0
                         0
                                                  Correct
0
                         0
                                                  Correct
1
                         1
                                                  Correct
2
                         2
                                                  Correct
1
                         1
                                                  Correct
0
                         0
                                                  Correct
2
                         2
                                                  Correct
0
                         0
                                                  Correct
                         2
2
                                                  Correct
1
                         1
                                                  Correct
                         2
2
                                                  Correct
0
                         0
                                                  Correct
0
                         0
                                                  Correct
                         2
Confusion Matrix:
[[7 0 0]
 [0 3 0]
[0 0 5]]
Classification Report:
              precision recall f1-score support

      1.00
      1.00
      1.00

      1.00
      1.00
      1.00

          0
                                                  3
          1
```

2	1.00	1.00	1.00	5	
accuracy macro avg weighted avg	1.00	1.00	1.00 1.00 1.00	15 15 15	
Accuracy of the	classifer	is 1.00			

#### Ex. 10

## **Study of PROLOG**

## **Prolog Study**

- o Prolog stands for programming in logic. In the logic programming paradigm, prolog language is most widely available. Prolog is a declarative language, which means that a program consists of data based on the facts and rules (Logical relationship) rather than computing how to find a solution. A logical relationship describes the relationships which hold for the given application.
- To obtain the solution, the user asks a question rather than running a program. When a
  user asks a question, then to determine the answer, the run time system searches through
  the database of facts and rules.
- Starting Prolog
- Prolog system is straightforward. From one person to other person, the precise details
  of Prolog will vary. Prolog will produce a number of lines of headings in the starting,
  which is followed by a line. It contains just
- · ?-
- The above symbol shows the system prompt. The prompt is used to show that the Prolog system is ready to specify one or more goals of sequence to the user. Using a full stop, we can terminate the sequence of goals.
- ?- write('Welcome to Javatpoint'),nl,write('Example of Prolog'),nl.
- o **nl** indicates 'start a new line'. When we press 'return' key, the above line will show the effect like this:
- **o** Welcome to Javatpoint
- Example of Prolog

- o yes
- **?- prompt** shows the sequence of goal which is entered by the user. The user will not type the prompt. Prolog system will automatically generate this prompt. It means that it is ready to receive a sequence of goals.
- o The above example shows a sequence of goals entered by the user like this:
- o write('Welcome to Javatpoint'), write('Example of Prolog'), nl(twice).

Consider the following sequence of goals:

## write('Welcome to AI'),nl,write('Example of Prolog'),nl.

The above sequence of goals has to succeed in order to be succeeded.

- o write('Welcome to AI')On the screen of the user, Welcome to AI has to be displayed
- $\circ$  nl

On the screen of the user, a new line has to be output

- o write('Example of Prolog')
- o On the screen of the user, Example of Prolog has to be displayed
- $\circ$  nl

On the screen of the user, a new line has to be output

All these goals will simply achieve by the Prolog system by outputting the line of text to the screen of the user. To show that the goals have succeeded, we will output **yes**.

The Prolog system predefined the meanings of **nl** and **write**. Write and nl are called as built-in predicates.

**Halt** and **statistics** are the two other built-in predicates. In almost all Prolog versions, these predicates are provided as standard.

#### o ?-halt.

The above command is used to terminate the Prolog system.

#### ?-statistics.

This command will cause the Prolog system statistics. This statistics feature is mainly used to experienced user. In statistics, the following things will generate:

# 8 queens problem

## Aim:

Write a program to solve 8 queens problem.

```
:- 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).
```

## **Depth First Search**

#### Aim:

Write a program to solve any problem using depth first search.

```
% 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).
```

#### Ex. 13

#### 8 Puzzle

Aim:

Write a program to solve any problem using 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('\sim n\sim w\sim n\sim n', [Move]),
 format('\sim w \sim w \sim w \sim n', [A,B,C]), \\format('\sim w \sim w \sim w \sim n', [D,E,F]), \\
 format('\simw \simw \simn',[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).
move( state(A,*,C,D,E,F,G,H,J), state(*,A,C,D,E,F,G,H,J), left ).
move(state(A,*,C,D,E,F,G,H,J), state(A,C,*,D,E,F,G,H,J), right).
move(state(A,*,C,D,E,F,G,H,J), state(A,E,C,D,*,F,G,H,J), down).
move( state(A,B,*,D,E,F,G,H,J), state(A,*,B,D,E,F,G,H,J), left ).
move(state(A,B,*,D,E,F,G,H,J), state(A,B,F,D,E,*,G,H,J), down).
move(\ state(A,B,C,^*,E,F,G,H,J),\ state(^*,B,C,A,E,F,G,H,J),\ up\ \ ).
move(state(A,B,C,*,E,F,G,H,J), state(A,B,C,E,*,F,G,H,J), right).
move(state(A,B,C,*,E,F,G,H,J), state(A,B,C,G,E,F,*,H,J), down).
move( state(A,B,C,D,*,F,G,H,J), state(A,*,C,D,B,F,G,H,J), up ).
move(state(A,B,C,D,*,F,G,H,J), state(A,B,C,D,F,*,G,H,J), right).
move(state(A,B,C,D,*,F,G,H,J), state(A,B,C,D,H,F,G,*,J), down).
```

```
move( state(A,B,C,D,*,F,G,H,J), state(A,B,C,*,D,F,G,H,J), left ). move( state(A,B,C,D,E,*,G,H,J), state(A,B,*,D,E,C,G,H,J), up ). move( state(A,B,C,D,E,*,G,H,J), state(A,B,C,D,*,E,G,H,J), left ). move( state(A,B,C,D,E,*,G,H,J), state(A,B,C,D,E,J,G,H,*), down ). move( state(A,B,C,D,E,F,*,H,J), state(A,B,C,D,E,F,H,*,J), left ). move( state(A,B,C,D,E,F,*,H,J), state(A,B,C,D,E,F,H,*,J), up ). move( state(A,B,C,D,E,F,G,*,J), state(A,B,C,D,E,F,*,G,J), left ). move( state(A,B,C,D,E,F,G,*,J), state(A,B,C,D,E,F,G,J), up ). move( state(A,B,C,D,E,F,G,*,J), state(A,B,C,D,E,F,G,J,*), right). move( state(A,B,C,D,E,F,G,H,*), state(A,B,C,D,E,F,G,*,H), left )
```

## traveling salesman

#### Aim:

Write a program to solve any problem using traveling 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,!.
```