# Ex. 1 FIND-S algorithm

for i, val in enumerate(c):

## 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.

```
Code:
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
```



```
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))
```

#### Result:

```
Sky Temp Humidity Wind Water Forecast EnjoySport
0 1 Sunny Warm Normal Strong Warm
                                             Same
                                                        Yes
1 2 Sunny Warm High Strong Warm
                                            Same
                                                      Yes
2 3 Rainy Cold High Strong Warm Change
                                                     Nο
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 de monstrate the Candidate-Elimination algorithm to output a description of the set of a II 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:
            qeneral_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, q_final = learn(concepts, target)
print("Final Specific_h: ", s_final, sep="\n")
```



print("Final General\_h: ", g\_final, sep="\n")

#### Result:

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

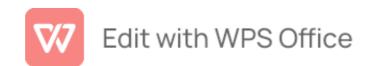
Specific Boundary: ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']

Instance 1 is ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
Instance is Positive

'?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

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

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



## Aim:

Write a program to demonstrate the working of the decision tree based ID3 algorithm. Use a n appropriate data set for building the decision tree and apply this knowledge to classify a ne w 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 unig:
    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",unig)
  for u in unig:
    #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)
```



# 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

```
Code:
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 contrib
uted to error
  d_hiddenlayer = EH * hiddengrad
  wout += hlayer_act.T.dot(d_output) *Ir # dotproduct of nextlayererror and currentl
ayerop
  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)
Result:
-----Epoch- 1 Starts-----
Input:
[[0.66666667 1. ]
```



```
[0.33333333 0.55555556]
[1.
       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]
[1.
       0.66666667]]
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]
[1.
       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]]
```

#### Ex.5

# Naive Bayesian Classifier

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

## 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
v = 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
Result:
```

NaiveBayesian Classifier

Edit with WPS Office

Ex. 6

#### 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. Cal culate 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: ', 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

Precision: 0.8709677419354839

Confusion Matrix:

[[64 4] [ 5 27]]

Result:

#### Ex. 7

# Bayesian network

#### Aim:

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

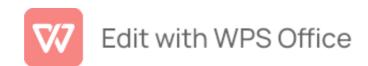
#### Code:

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','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(a1)
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 exang oldpeak \
               145 233 1
                                    150
0 63
        1 1
                               2
                                         0
                                               2.3
        1 4
                160 286 0
                                    108
                                               1.5
1 67
                                2
                                           1
        1 4

    1
    4
    120
    229
    0
    2
    129
    1
    2.6

    1
    3
    130
    250
    0
    0
    187
    0
    3.5

2 67
3 37
4 41
        0 2 130 204 0 2 172 0 1.4
 slope ca thal heartdisease
   3 0 6
                 0
1
    2 3 3
                 2
  227
2
                 1
3 3 0 3
                 0
  1 0 3
                 0
Attributes and datatypes
         int64
age
gender
           int64
ср
          int64
trestbps
          int64
chol
           int64
fbs
          int64
          int64
restecq
thalach
            int64
            int64
exang
oldpeak
           float64
slope
           int64
```

Learning CPD using Maximum likelihood estimators

object

object

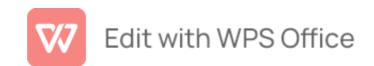
int64

ca

thal

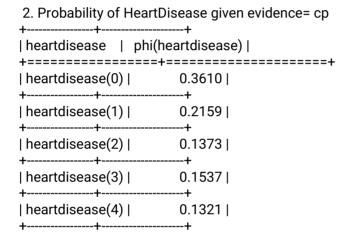
heartdisease

dtype: object



Inferencing with Bayesian Network:

1. Probability of HeartDisease given evidence= restect	
heartdisease   phi(heartdisease)	
+======+=	+
heartdisease(0)	0.1012
+	+
heartdisease(1)   	0.0000
heartdisease(2)	0.2392
	t
heartdisease(3)	0.2015
+	+
heartdisease(4)	0.4581
++	+



## 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 datase t 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(v, 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: 0.093333333333333333
The Confusion matrix of K-Mean:
[[0 50 0]
[2 0 48]
[36 0 14]]
The accuracy score of EM: 0.966666666666667
The Confusion matrix of EM:
[[50 0 0]
[0.45.5]
[0\ 0\ 50]
```



Result:

# Ex. 9 k-Nearest Neighbour

**Aim:** Write a program to implement k-Nearest Neighbour algorithm to classify the iris data s et. Print 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 ('%-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'))
    print (' %-25s' % ('Wrong'))
  i = i + 1
print("\nConfusion Matrix:\n",metrics.confusion_matrix(ytest, ypred))
```



```
print("\nClassification Report:\n",metrics.classification_report(ytest, ypred))
print('Accuracy of the classifer is %0.2f' % metrics.accuracy_score(ytest,ypred))
Output:
 sepal length (cm) sepal width (cm) petal length (cm) petal width (cm)
          5.1
                     3.5
                                 1.4
                                             0.2
1
          4.9
                     3.0
                                  1.4
                                             0.2
2
          4.7
                     3.2
                                  1.3
                                             0.2
3
          4.6
                                             0.2
                     3.1
                                  1.5
4
          5.0
                                  1.4
                                             0.2
                     3.6
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
               2
                              Correct
Confusion Matrix:
[[7 \ 0 \ 0]]
[0 \ 3 \ 0]
[0\ 0\ 5]]
Classification Report:
        precision recall f1-score support
      0
           1.00
                   1.00
                          1.00
                                    7
      1
           1.00
                          1.00
                                    3
                   1.00
      2
                                    5
           1.00
                   1.00
                          1.00
  accuracy
                          1.00
                                   15
 macro avg
                       1.00
                               1.00
                                        15
                1.00
weighted avg
                 1.00
                        1.00
                               1.00
                                         15
Accuracy of the classifer is 1.00
```



Result:

# **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.
- o 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.
- o Starting Prolog
- o 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
- o ?-
- o 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.
- o ?- 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
- o 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:
- 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.

 write('Welcome to Al')On the screen of the user, Welcome to Al has to be displayed

o **n** 

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

- write('Example of Prolog')
- o On the screen of the user, Example of Prolog has to be displayed
- o n

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.

o ?-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:

Ex. 11 Aim: 8 queens problem



Write a program to solve 8 queens problem.

# Code:

```
:- 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).
```

# Ex. 12

# **Depth First Search**

## Aim:

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

## Code:



```
% 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).
Result:
Ex. 13
                                       8 Puzzle
```



Aim:

## Write a program to solve any problem using 8 puzzle.

#### Code:

```
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('\simw \simw \simw\simn',[A,B,C]),
  format('\simw \simw \simw\simn',[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,*,E,F,D,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,*,F,G,E,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,*,G,H,F), up).
```



 $move(\ state(A,B,C,D,E,F,G,H,*),\ state(A,B,C,D,E,F,G,*,H),\ left\ )$ 

Result:

Ex. 15

traveling salesman



## Aim:

Write a program to solve any problem using traveling salesman.

#### Code:

```
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,!.
```