

ML LAB REPORT

1BM18CS061

NIKHIL A S

Week 1:

i) Find s algorithm

*# This Python 3 environment comes with many helpful analytics libraries installed
It is defined by the kaggle/python Docker image: <https://github.com/kaggle/docker-python>
For example, here's several helpful packages to load*

```
import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
```

*# Input data files are available in the read-only "../input/" directory
For example, running this (by clicking run or pressing Shift+Enter) will list all files under the input directory*

```
import os
for dirname, _, filenames in os.walk('/kaggle/input'):
    for filename in filenames: print(os.path.join(dirname,
                                                    filename))
```

You can write up to 20GB to the current directory (/kaggle/working/) that gets preserved as output when you create a version using "Save & Run All" # You can also write temporary files to /kaggle/temp/, but they won't be saved outside of the current session

/kaggle/input/datasetcsv/data.csv

```
data = pd.read_csv("/kaggle/input/datasetcsv/data.csv")
print("The entered data is \n")
print(data, "\n")
d = np.array(data)[:,:-1]
print("\n The attributes are: \n", d)
target = np.array(data)[:,-1]
print("\n The target is: ", target)
def training(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
print("\n The final hypothesis is:", training(d, target))

```

Output:

The entered data is

	Weather	Temperature	Humidity	Goes
0	Sunny	Warm	Mild	Yes
1	Rainy	Cold	Mild	No
2	Sunny	Moderate	Nomal	Yes
3	Sunny	Cold	High	Yes

The attributes are:

['Sunny ' 'Warm ' 'Mild']

['Rainy' 'Cold' 'Mild']

['Sunny ' 'Moderate' 'Nomal']

['Sunny ' 'Cold' 'High ']

The target is: ['Yes' 'No' 'Yes' 'Yes']

The final hypothesis is: ['Sunny ' '?' '?']

Week 2:

ii) Candidate elimination algorithm:

This Python 3 environment comes with many helpful analytics libraries installed

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import numpy as np # linear algebra

import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)

Input data files are available in the read-only "../input/" directory

For example, running this (by clicking run or pressing Shift+Enter) will list all files under the input directory

import os

```
for dirname, _, filenames in os.walk('/kaggle/input'):
    for filename in filenames: print(os.path.join(dirname,
        filename))
```

You can write up to 20GB to the current directory (/kaggle/working/) that gets preserved as output when you create a version using "Save & Run All" # You can also write temporary files to /kaggle/temp/, but they won't be saved outside of the current session

/kaggle/input/candidatecsv/candidate.csv

```
data = pd.read_csv("/kaggle/input/candidatecsv/candidate.csv") print("Entered data
is")
print(data)
concepts = np.array(data)[:,-1] print("\n The
attributes are: \n",d) target = np.array(data)[:,-1]
print("\n The target is: ",target)
```

Entered data is

	sky	airtemp	humidity	wind	water	forecast	enjoysport
0	sunny	warm	normal	strong	warm	same	yes
1	sunny	warm	high	strong	warm	same	yes
2	rainy	cold	high	strong	warm	change	no
3	sunny	warm	high	strong	cool	change	yes

The attributes 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']]
```

The target is: ['yes' 'yes' 'no' 'yes']

```
#training function to implement candidate_elimination algorithm
def learn(concepts, target):
    specific_h = concepts[0].copy()
    print("\n Initialization of specific_h and general_h")
    print(specific_h)
    general_h = [["?" for i in range(len(specific_h))] for i in
range(len(specific_h))]
    print(general_h)
    for i, h in enumerate(concepts):
        if target[i] == "yes":
            for x in range(len(specific_h)):
                if h[x] != specific_h[x]:
                    specific_h[x] = '?'
                    general_h[x][x] = '?'
            print(specific_h)
    print(specific_h)
    if target[i] == "no":
        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("\n Steps of Candidate Elimination Algorithm", i+1)
    print(specific_h)
    print(general_h)
    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)

#obtaining the final hypothesis
print("\nFinal Specific_h:", s_final, sep="\n")
print("\nFinal General_h:", g_final, sep="\n")

```

Output:

Initialization of specific_h and general_h ['sunny' 'warm' 'normal'

```

'strong' 'warm' 'same']
[['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?',
 '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?',
 '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']

```

Steps of Candidate Elimination Algorithm 1 ['sunny' 'warm'

```

'normal' 'strong' 'warm' 'same']
[['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?',
 '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?',
 '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
['sunny' 'warm' '?' 'strong' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']

```

Steps of Candidate Elimination Algorithm 2 ['sunny' 'warm'

```

'?' '?' 'warm' 'same']
[['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?',
 '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?',
 '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
['sunny' 'warm' '?' '?' 'warm' 'same']

```

Steps of Candidate Elimination Algorithm 3 ['sunny' 'warm'

```

'?' '?' 'warm' 'same']

```

```

[['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?'],
['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '
?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', 'same']]
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' 'warm' 'same']
['sunny' 'warm' '?' '?' '?' 'same']
['sunny' 'warm' '?' '?' '?' '?']
['sunny' 'warm' '?' '?' '?' '?']

```

Steps of Candidate Elimination Algorithm 4 ['sunny' 'warm'
'?' '?' '?' '?']

```

[['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?'],
['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '
?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

```

Final Specific_h:

```
['sunny' 'warm' '?' '?' '?' '?']
```

Final General_h:

```
[['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?']]
```

Week 3:

ID3 algorithm:

```

import
t math

import csv
def load_csv(filename):
    lines=csv.reader(open(filename,"r"))
    dataset = list(lines)
    headers = dataset.pop(0)
    return dataset,headers

class Node:
    def __init__(self,attribute):
        self.attribute=attribute
        self.children=[]
        self.answer=""

def subtables(data,col,delete):
    dic={}
    coldata=[row[col] for row in data]

```

```

attr=list(set(coldata))

counts=[0]*len(attr)
r=len(data)
c=len(data[0])
for x in range(len(attr)):
    for y in range(r):
        if data[y][col]==attr[x]:
            counts[x]+=1

for x in range(len(attr)):
    dic[attr[x]]=[[0 for i in range(c)] for j in
range(counts[x])]
    pos=0
    for y in range(r):
        if data[y][col]==attr[x]:
            if delete:
                del data[y][col]
            dic[attr[x]][pos]=data[y]
            pos+=1
return attr,dic

def entropy(S):
    attr=list(set(S))
    if len(attr)==1:
        return 0

    counts=[0,0]
    for i in range(2):
        counts[i]=sum([1 for x in S if attr[i]==x])/(len(S)*1.0)

    sums=0
    for cnt in counts:
        sums+=-1*cnt*math.log(cnt,2)
    return sums

def compute_gain(data,col):
    attr,dic = subtables(data,col,delete=False)

    total_size=len(data)
    entropies=[0]*len(attr)
    ratio=[0]*len(attr)

    total_entropy=entropy([row[-1] for row in data])
    for x in range(len(attr)):
        ratio[x]=len(dic[attr[x]])/(total_size*1
.0)

```

```

        entropies[x]=entropy([row[-1] for row in dic[attr[x]]])
        total_entropy-=ratio[x]*entropies[x]
    return total_entropy

def build_tree(data,features):
    lastcol=[row[-1] for row in data]
    if(len(set(lastcol))==1):
        node=Node("")
        node.answer=lastcol[0]
        return node

    n=len(data[0])-1
    gains=[0]*n
    for col in range(n):
        gains[col]=compute_gain(data,col)
    split=gains.index(max(gains))
    node=Node(features[split])
    fea = features[:split]+features[split+1:]

    attr,dic=subtables(data,split,delete=True)

    for x in range(len(attr)):
        child=build_tree(dic[attr[x]],fea)
        node.children.append((attr[x],child))
    return node

def print_tree(node,level):
    if node.answer!="":
        print("  "*level,node.answer)
        return

    print("  "*level,node.attribute)
    for value,n in node.children:
        print("  "*(level+1),value)
        print_tree(n,level+2)

def classify(node,x_test,features):
    if node.answer!="":
        print(node.answer)
        return

```

```

        pos=features.index(node.attribute)
        for value, n in node.children:
            if x_test[pos]==value:
                classify(n,x_test,features)

'''Main program'''
dataset,features=load_csv("id3.csv")
node1=build_tree(dataset,features)

print("The decision tree for the dataset using ID3 algorithm is")
print_tree(node1,0)
testdata,features=load_csv("id3_test.csv")

for xtest in testdata:
    print("The test instance:",xtest)
    print("The label for test instance:")
    classify(node1,xtest,features)

```

Output:

```

bmsce@bmsce-Precision-T1700:~/Documents/LAB - 3 - DECISION TREE$ python ml3.py
The decision tree for the dataset using ID3 algorithm is
(' ', 'Outlook')
(' ', 'overcast')
(' ', 'yes')
(' ', 'sunny')
(' ', 'Humidity')
(' ', 'high')
(' ', 'no')
(' ', 'normal')
(' ', 'yes')
(' ', 'rain')
(' ', 'Wind')
(' ', 'strong')
(' ', 'no')
(' ', 'weak')
(' ', 'yes')
('The test instance:', ['rain', 'cool', 'normal', 'strong'])
The label for test instance:
no
('The test instance:', ['sunny', 'mild', 'normal', 'strong'])
The label for test instance:
yes

```


Week 4:

iv) Naïve bayes classifier:

```
import pandas as pd
```

```
data = pd.read_csv('PlayTennis.csv')  
data.head()
```

```
y = list(data['PlayTennis'].values) X =
```

```
data.iloc[:,1:].values print(f'Target Values:
```

```
{y}') print(f'Features: \n{X}')
```

Target Values: ['No', 'No', 'Yes', 'Yes', 'Yes', 'No', 'Yes', 'No', 'Yes', 'Yes', 'Yes', 'Yes', 'Yes', 'No']

Features:

[['Sunny' 'Hot' 'High' 'Weak']

['Sunny' 'Hot' 'High' 'Strong']

['Overcast' 'Hot' 'High' 'Weak']

['Rain' 'Mild' 'High' 'Weak']

['Rain' 'Cool' 'Normal' 'Weak']

['Rain' 'Cool' 'Normal' 'Strong'] ['Overcast' 'Cool'

'Normal' 'Strong'] ['Sunny' 'Mild' 'High' 'Weak']

['Sunny' 'Cool' 'Normal' 'Weak']

['Rain' 'Mild' 'Normal' 'Weak']

['Sunny' 'Mild' 'Normal' 'Strong']

['Overcast' 'Mild' 'High' 'Strong']

['Overcast' 'Hot' 'Normal' 'Weak']

['Rain' 'Mild' 'High' 'Strong']]

```
y_train = y[:8]
```

```
y_val = y[8:]
```

```
X_train = X[:8]
```

```
X_val = X[8:]
```

```
print(f"Number of instances in training set: {len(X_train)}")
```

```
print(f"Number of instances in testing set: {len(X_val)}")
```

Number of instances in training set: 8

Number of instances in testing set: 6

```
class NaiveBayesClassifier:
```

```
    def __init__(self, X, y):
```

```
        self.X, self.y = X, y
```

```
        self.N = len(self.X)
```

```

self.dim = len(self.X[0])

self.attrs = [[] for _ in range(self.dim)]

self.output_dom = {}

self.data = []

for i in range(len(self.X)):
    for j in range(self.dim):
        if not self.X[i][j] in self.attrs[j]:
            self.attrs[j].append(self.X[i][j])

    if not self.y[i] in self.output_dom.keys():
        self.output_dom[self.y[i]] = 1

    else:
        self.output_dom[self.y[i]] += 1

    self.data.append([self.X[i], self.y[i]])
def classify(self, entry):

    solve = None
    max_arg = -1

    for y in self.output_dom.keys():

        prob = self.output_dom[y]/self.N

        for i in range(self.dim):
            cases = [x for x in self.data if x[0][i] == entry[i] and
d x[1] == y]

            n = len(cases)
            prob *= n/self.N

        if prob > max_arg:
            max_arg = prob
            solve = y

    return solve

nbc = NaiveBayesClassifier(X_train, y_train)
total_cases = len(y_val)

good = 0
bad = 0
predictions = []

for i in range(total_cases):
    predict = nbc.classify(X_val[i])
    predictions.append(predict)

    if y_val[i] == predict:
        good += 1
    else:
        bad += 1

```

```

print('Predicted values:', predictions)
print('Actual values:', y_val)
print()
print('Total number of testing instances in the dataset:', total_cases)
print('Number of correct predictions:', good)
print('Number of wrong predictions:', bad)
print()
print('Accuracy of Bayes Classifier:', good/total_cases)

```

Output:

Predicted values: ['No', 'Yes', 'No', 'Yes', 'Yes', 'No']

Actual values: ['Yes', 'Yes', 'Yes', 'Yes', 'Yes', 'No']

Total number of testing instances in the dataset: 6 Number of correct predictions: 4

Number of wrong predictions: 2

Accuracy of Bayes Classifier: 0.6666666666666666

Week 5:

v) Bayesian network:

This Python 3 environment comes with many helpful analytics libraries installed # It is defined by the kaggle/python Docker image: <https://github.com/kaggle/docker-python> # For example, here's several helpful packages to load **import numpy as np**

linear algebra

import pandas as pd

import pgmpy as pgmpy from pgmpy.estimators

import MaximumLikelihoodEstimator **from pgmpy.models import**

BayesianModel **from pgmpy.inference**

import VariableElimination

import os for dirname, _ filenames **in** os.walk('/kaggle/input'):

for filename **in** filenames: **print**(os.path.join(dirname, filename)) *# You can write up to 20GB to the current directory (/kaggle/working/) that gets preserved as output when you create a version using "Save & Run All" # You can also write temporary files to /kaggle/temp/, but they won't be saved outside of the current session*

#read Cleveland Heart Disease data

heartDisease = pd.read_csv("/kaggle/input/bayesiannetwork/heart.csv")

heartDisease = heartDisease.replace('?', np.nan)

#display the data

print('Sample instances from the dataset are given below')

print(heartDisease.head())

```

#display the Attributes names and datatypes
print('\n Attributes and datatypes')
print(heartDisease.dtypes)
#Creat Model- Bayesian Network
model = BayesianModel([('age', 'heartdisease'), ('sex', 'heartdisease'), ('exang', 'heartdisease'), ('cp', 'heartdisease'), ('heartdisease', 'restecg'), ('heartdisease', 'chol')])
#Learning CPDs using Maximum Likelihood Estimators
print('\n Learning CPD using Maximum likelihood estimators')
model.fit(heartDisease, estimator=MaximumLikelihoodEstimator)
# Inferencing with Bayesian Network
print('\n Inferencing with Bayesian Network:')
HeartDiseasetest_infer = VariableElimination(model)
#computing the Probability of HeartDisease given restecg
print('\n 1.Probability of HeartDisease given evidence= restecg :1')
q1=HeartDiseasetest_infer.query(variables=['heartdisease'], evidence={'restecg':1})
print(q1)
#computing the Probability of HeartDisease given cp
print('\n 2.Probability of HeartDisease given evidence= cp:2 ')
q2=HeartDiseasetest_infer.query(variables=['heartdisease'], evidence={'cp':2})
print(q2)

```

Output:

Sample instances from the dataset are given below

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak
slope										
0	63	1	1	145	233	1	2	150	0	2.3
3										
1	67	1	4	160	286	0	2	108	1	1.5
2										
2	67	1	4	120	229	0	2	129	1	2.6
2										
3	37	1	3	130	250	0	0	187	0	3.5
3										
4	41	0	2	130	204	0	2	172	0	1.4
1										

	ca	thal	heartdisease
0	0	6	0
1	3	3	2
2	2	7	1
3	0	3	0
4	0	3	0

Attributes and datatypes

age	int64
sex	int64
cp	int64
trestbps	int64
chol	int64
fbs	int64
restecg	int64
thalach	int64

```

exang          int64
oldpeak        float64
slope          int64
ca             object
thal           object
heartdisease    int64
dtype: object

```

Learning CPD using Maximum likelihood estimators

```

Finding Elimination Order: :          0%|          | 0/5 [00:00<?, ?it/s] 0%|  |
      0/5 [00:00<?, ?it/s]

```

```

Eliminating: age:          0%|          | 0/5 [00:00<?, ?it/s] Eliminating: chol:
                        0%|          | 0/5 [00:00<?, ?it/s] Eliminating: cp:
                        0%|          | 0/5 [00:00<?, ?it/s] Eliminating: sex:
                        0%|          | 0/5 [00:00<?, ?it/s] Eliminating: exang:

```

```

100%|■■■■■■■■■■■■■■■■| 5/5 [00:00<00:00, 189.65it/s]

```

```

Finding Elimination Order: : 100%|■■■■■■■■■■■■■■■■| 5/5 [00:00<00:00, 132.81it/s] Finding

```

```

Elimination Order: :          0%|          | 0/5 [00:00<?, ?it/s]
      0%|          | 0/5 [00:00<?, ?it/s]

```

```

Eliminating: age:          0%|          | 0/5 [00:00<?, ?it/s] Eliminating: chol:
                        0%|          | 0/5 [00:00<?, ?it/s] Eliminating: restecg:
                        0%|          | 0/5 [00:00<?, ?it/s] Eliminating: sex:
                        0%|          | 0/5 [00:00<?, ?it/s] Eliminating: exang:

```

```

100%|■■■■■■■■■■■■■■■■| 5/5 [00:00<00:00, 230.00it/s]

```

Inferencing with Bayesian Network:

1 . Probability of HeartDisease given evidence= restecg :1

```

+-----+-----+-----+-----+
| 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:2

```

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

```

WEEK 6:

vi) Inferring from Bayesian model:

```
from pgmpy.models import BayesianModel
```

```
from pgmpy.factors.discrete import TabularCPD from
```

```
pgmpy.inference import VariableElimination
```

```
cancer_model = BayesianModel([ ('Pollution', 'Cancer'),
                                ('Smoker', 'Cancer'),
                                ('Cancer', 'Xray'),
                                ('Cancer', 'Dyspnoea') ])

print('Bayesian network nodes are:')
print("\t",cancer_model.nodes())
print('Bayesian network edges are:')
print("\t",cancer_model.edges())

cpd_poll = TabularCPD(variable='Pollution',variable_card=2,values=[[0.9
], [0.1]])
cpd_smoke = TabularCPD(variable='Smoker',variable_card=2,values=[[0.3],
[0.7]])
cpd_cancer = TabularCPD(variable='Cancer',variable_card=2,values=[[0.03
,0.05,0.001,0.02],
                                                                    [0.97,
0.95,0.999,0.98]],
                        evidence=['Smoker','Pollution'],
                        evidence_card=[2,2])
cpd_xray = TabularCPD(variable='Xray',variable_card=2,values=[[0.9,0.2]
,[0.1,0.8]],
                        evidence=['Cancer'],evidence_card=[2])
cpd_dysp = TabularCPD(variable='Dyspnoea',variable_card=2,values=[[0.65
,0.3], [0.35,0.7]],
                        evidence=['Cancer'],evidence_card=[2])
```

Bayesian network nodes are:

['Pollution', 'Cancer', 'Smoker', 'Xray', 'Dyspnoea'] Bayesian network edges

are:

[('Pollution', 'Cancer'), ('Cancer', 'Xray'), ('Cancer', 'Dyspnoea'), ('Smoker', 'Cancer')]

```
cancer_model.add_cpds(cpd_poll,cpd_smoke,cpd_cancer,cpd_xray,cpd_dysp)
print('Model generated by adding cpts(cpds)')
print('Checking correctness of model:',end='')
print(cancer_model.check_model())
```

Model generated by adding cpts(cpds) Checking
correctness of model:True

```
print('All local dependencies are as follows')
```

All local dependencies are as follows

Out[10]:

(Pollution \perp Smoker)
 (Pollution \perp Dyspnoea, Xray | Cancer) (Pollution \perp Xray |
 Dyspnoea, Cancer) (Pollution \perp Dyspnoea | Cancer, Xray)
 (Pollution \perp Dyspnoea, Xray | Cancer, Smoker) (Pollution \perp Xray
 | Dyspnoea, Cancer, Smoker) (Pollution \perp Dyspnoea | Cancer,
 Xray, Smoker) (Smoker \perp Pollution)
 (Smoker \perp Dyspnoea, Xray | Cancer) (Smoker \perp
 Xray | Dyspnoea, Cancer)
 (Smoker \perp Dyspnoea, Xray | Pollution, Cancer) (Smoker \perp
 Dyspnoea | Cancer, Xray)
 (Smoker \perp Xray | Dyspnoea, Pollution, Cancer) (Smoker \perp
 Dyspnoea | Pollution, Cancer, Xray) (Xray \perp Dyspnoea,
 Pollution, Smoker | Cancer) (Xray \perp Pollution, Smoker |
 Dyspnoea, Cancer) (Xray \perp Dyspnoea, Smoker | Pollution,
 Cancer) (Xray \perp Dyspnoea, Pollution | Cancer, Smoker) (Xray \perp
 Smoker | Dyspnoea, Pollution, Cancer) (Xray \perp Pollution |
 Dyspnoea, Cancer, Smoker) (Xray \perp Dyspnoea | Pollution,
 Cancer, Smoker) (Dyspnoea \perp Pollution, Xray, Smoker | Cancer)
 (Dyspnoea \perp Xray, Smoker | Pollution, Cancer) (Dyspnoea \perp
 Pollution, Smoker | Cancer, Xray) (Dyspnoea \perp Pollution, Xray |
 Cancer, Smoker) (Dyspnoea \perp Smoker | Pollution, Cancer, Xray)
 (Dyspnoea \perp Xray | Pollution, Cancer, Smoker) (Dyspnoea \perp
 Pollution | Cancer, Xray, Smoker)

```
print('Displaying CPDs')
print(cancer_model.get_cpds('Pollution'))
print(cancer_model.get_cpds('Smoker'))
print(cancer_model.get_cpds('Cancer'))
print(cancer_model.get_cpds('Xray'))
print(cancer_model.get_cpds('Dyspnoea'))
```

Smoker	Smoker(0)	Smoker(0)	Smoker(1)	Smoker(1)
Pollution	Pollution(0)	Pollution(1)	Pollution(0)	Pollution(1)
Cancer(0)	0.03	0.05	0.001	0.02
Cancer(1)	0.97	0.95	0.999	0.98
Xray(0)	0.9	0.2		
Xray(1)	0.1	0.8		
Dyspnoea(0)	0.65	0.3		
Dyspnoea(1)	0.35	0.7		

```

cancer_infer=VariableElimination(cancer_model)
print('\n Inferencing with bayesian network')
print("\n Probability of Cancer given smoker")
q=cancer_infer.query(variables=['Cancer'],evidence={'Smoker':1})
print(q)

print("\n Probability of Cancer given smoker,pollution")
q=cancer_infer.query(variables=['Cancer'],evidence={'Smoker':1,'Pollution':1})
print(q)

```

Output:

```

Finding Elimination Order: : 0%| 0/3 [00:00<?, ?it/s] 0%|
0/3 [00:00<?, ?it/s]
Eliminating: Dyspnoea: 0%| 0/3 [00:00<?, ?it/s] Eliminating:
Pollution: 0%| 0/3 [00:00<?, ?it/s] Eliminating:
Xray: 100%|■■■■■■■■■■| 3/3 [00:00<00:00, 359.52it/s]

0%| 0/2 [00:00<?, ?it/s]
Finding Elimination Order: : 0%| 0/2 [00:00<?, ?it/s] 0%|
0/2 [00:00<?, ?it/s]

```


Eliminating: Dyspnoea: 0% | 0/2 [00:00<?, ?it/s]

Eliminating: Xray: 100%|■■■■■■■■■■| 2/2 [00:00<00:00, 333.49it/s]A Inferencing
with bayesian network

Probability of Cancer given smoker

+-----+	
Cancer	phi(Cancer)
+=====+	
Cancer(0)	0.0029
+-----+	
Cancer(1)	0.9971

Probability of Cancer given smoker,pollution

+-----+	
Cancer	phi(Cancer)
+=====+	
Cancer(0)	0.0200
+-----+	
Cancer(1)	0.9800
+-----+	