```
import numpy as np
import pgmpy
import pandas as pd
import csv
from pgmpy.models import BayesianModel
print(pgmpy. version )
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
heartDisease = pd.read csv('heart.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'), ('sex', 'heartdisease'), ('exang', 'heartdisease'), ('cp', 'he
artdisease'),('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)
 Learning CPD using Maximum likelihood estimators
  Inferencing with Bayesian Network:
                                                 Probability of HeartDisease given evidence= cp
  1. Probability of HeartDisease given evidence= restecg
                                                 | heartdisease | phi(heartdisease)
 +----+
                                                 +=======+
  heartdisease | phi(heartdisease) |
                                                  heartdisease(0)
 +========+
  heartdisease(0)
                                                 heartdisease(1)
 +-----+
                                                 +----+
  heartdisease(1)
                                                 heartdisease(2)
 +-----+
  heartdisease(2)
                                                 heartdisease(3) 0.1541
  heartdisease(3)
                                                heartdisease(4)
 +----+
                                                 +-----+
  heartdisease(4)
 +-----
```

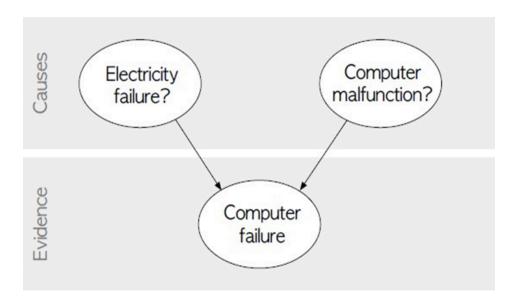
A Bayesian network is a directed acyclic graph in which each edge corresponds to a conditional dependency, and each node corresponds to a unique random variable.

Bayesian network consists of two major parts: a directed acyclic graph and a set of conditional probability distributions

- The directed acyclic graph is a set of random variables represented by nodes.
- The conditional probability distribution of a node (random variable) is defined for every possible outcome of the preceding causal node(s).

For illustration, consider the following example. Suppose we attempt to turn on our computer, but the computer does not start (observation/evidence). We would like to know which of the possible causes of computer failure is more likely. In this simplified illustration, we assume only two possible causes of this misfortune: electricity failure and computer malfunction.

The corresponding directed acyclic graph is depicted in below figure.



The goal is to calculate the posterior conditional probability distribution of each of the possible unobserved causes given the observed evidence, i.e. P [Cause | Evidence].

Sa	mple	insta	nces	from the	datase	t are	given be	low				
	age	sex		trestbps			_		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	
0 1 2 3 4	ca t 0 3 2 0	hal 6 3 7 3 3	heart	disease 0 2 1 0			Attributes and datatypes age int64 sex int64	stbps 1	restecg int64 thalach int64 exang int64 oldbeak float64	slope int64 ca object thal object heartdisease int64	dtype: object	

Data Set:

Title: Heart Disease Databases

The Cleveland database contains 76 attributes, but all published experiments refer to using a subset of 14 of them. In particular, the Cleveland database is the only one that has been used by ML researchers to this date. The "Heartdisease" field refers to the presence of heart disease in the patient. It is integer valued from 0 (no presence) to 4.

Database: 0 1 2 3 4 Total

Cleveland: 164 55 36 35 13 303

<u>Attribute Information:</u>

1. age: age in years

2. sex: sex (1 = male; 0 = female)

3. cp: chest pain type

Value 1: typical angina

· Value 2: atypical angina

· Value 3: non-anginal pain

· Value 4: asymptomatic

4. trestbps: resting blood pressure (in mm Hg on admission to the hospital)

5. chol: serum cholestoral in mg/dl

6. fbs: (fasting blood sugar > 120 mg/dl) (1 = true; 0 = false)

7. restecg: resting electrocardiographic results

· Value 0: normal

 Value 1: having ST-T wave abnormality (T wave inversions and/or ST elevation or depression of > 0.05 mV)

Value 2: showing probable or definite left ventricular hypertrophy by Estes' criteria

8. thalach: maximum heart rate achieved

9. exang: exercise induced angina (1 = yes; 0 = no)

10. oldpeak = ST depression induced by exercise relative to rest

11. slope: the slope of the peak exercise ST segment

Value 1: upsloping

· Value 2: flat

· Value 3: downsloping

12. thal: 3 = normal; 6 = fixed defect; 7 = reversable defect

13. Heartdisease: It is integer valued from 0 (no presence) to 4.

Some instance from the dataset:

age	sex	ср	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	Heartdisease
63	1	1	145	233	1	2	150	0	2.3	3	0	6	0
67	1	4	160	286	0	2	108	1	1.5	2	3	3	2
67	1	4	120	229	0	2	129	1	2.6	2	2	7	1
41	0	2	130	204	0	2	172	0	1.4	1	0	3	0
62	0	4	140	268	0	2	160	0	3.6	3	2	3	3
60	1	4	130	206	0	2	132	1	2.4	2	2	7	4