

**Source Code:**

```
import pandas as pd
data=pd.read_csv("heart_disease_data1.csv")
heart_disease=pd.DataFrame(data)
print(heart_disease)

from pgmpy.models import BayesianModel
model=BayesianModel([
('age','Lifestyle'),
('Gender','Lifestyle'),
('Family','heartdisease'),
('diet','cholesterol'),
('Lifestyle','diet'),
('cholesterol','heartdisease'),
('diet','cholesterol')
])
from pgmpy.estimators import MaximumLikelihoodEstimator
model.fit(heart_disease, estimator=MaximumLikelihoodEstimator)

from pgmpy.inference import VariableElimination
HeartDisease_infer = VariableElimination(model)

print('For age enter SuperSeniorCitizen:0, SeniorCitizen:1, MiddleAged:2, Youth:3,
Teen:4')

print('For Gender Enter Male:0, Female:1')
print('For Family History Enter yes:1, No:0')
print('For diet Enter High:0, Medium:1')
print('for lifeStyle Enter Athlete:0, Active:1, Moderate:2, Sedentary:3')
print('for cholesterol Enter High:0, BorderLine:1, Normal:2')

q = HeartDisease_infer.query(variables=["heartdisease"], evidence={
    'age':int(input('enter age')),
    'Gender':int(input('enter Gender')),
    'Family':int(input('enter Family history')),
    'diet':int(input('enter diet')),
    'Lifestyle':int(input('enter Lifestyle')),
    'cholesterol':int(input('enter cholesterol'))
})

print(q['heartdisease'])
```

**Sample Dataset:**

1	age	Gender	Family	diet	Lifestyle	cholesterol	heartdisease
2	0	0	1	1	3	0	1
3	0	1	1	1	3	0	1
4	1	0	0	0	2	1	1
5	4	0	1	1	3	2	0
6	3	1	1	0	0	2	0
7	2	0	1	1	1	0	1
8	4	0	1	0	2	0	1
9	0	0	1	1	3	0	1
10	3	1	1	0	0	2	0
11	1	1	0	0	0	2	1
12	4	1	0	1	2	0	1
13	4	0	1	1	3	2	0
14	2	1	0	0	0	0	0
15	2	0	1	1	1	0	1
16	3	1	1	0	0	1	0
17	0	0	1	0	0	2	1
18	1	1	0	1	2	1	1
19	3	1	1	1	0	1	0
20	4	0	1	1	3	2	0

**Output:**

For age enter SuperSeniorCitizen:0, SeniorCitizen:1, MiddleAged:2, Youth:3, Teen:4

For Gender Enter Male:0, Female:1

For Family History Enter yes:1, No:0

For diet Enter High:0, Medium:1

for lifeStyle Enter Athlete:0, Active:1, Moderate:2, Sedentary:3

for cholesterol Enter High:0, BorderLine:1, Normal:2

enter age2

enter Gender1

enter Family history1

enter diet1

enter Lifestyle0

enter cholesterol2

```

+-----+-----+
| heartdisease | phi(heartdisease) |
+=====+=====+
| heartdisease_0 | 0.8333 |
+-----+-----+
| heartdisease_1 | 0.1667 |
+-----+-----+

```

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**Source Code:**

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.mixture import GaussianMixture
import pdb
df1 = pd.read_csv("clusterdata.csv")
print(df1)
f1 = df1['Distance_Feature'].values
f2 = df1['Speeding_Feature'].values

X = np.matrix(list(zip(f1,f2)))
plt.plot(1)
plt.subplot(511)
plt.xlim([0, 100])
plt.ylim([0, 50])
plt.title('Dataset')
plt.ylabel('speeding_feature')
plt.xlabel('distance_feature')
plt.scatter(f1,f2)

colors = ['b', 'g', 'r']
markers = ['o', 'v', 's']
# create new plot and data for K- means algorithm
plt.plot(2)
ax=plt.subplot(513)
kmeans_model = KMeans(n_clusters=3).fit(X)

for i, l in enumerate(kmeans_model.labels_):
    plt.plot(f1[i], f2[i], color=colors[l],marker=markers[l])

plt.xlim([0, 100])
plt.ylim([0, 50])
plt.title('K- Means')
plt.ylabel('speeding_feature')
plt.xlabel('distance_feature')

# create new plot and data for gaussian mixture
plt.plot(3)
plt.subplot(515)
gmm=GaussianMixture(n_components=3).fit(X)
labels= gmm.predict(X)
```

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for i, l in enumerate(labels):
    plt.plot(f1[i], f2[i], color=colors[l], marker=markers[l])

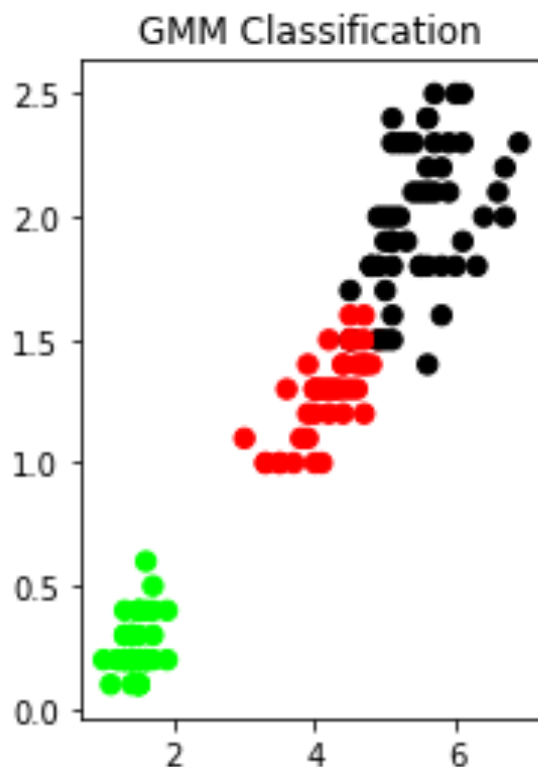
plt.xlim([0, 100])
plt.ylim([0, 50])
plt.title("Gaussian Mixture")
plt.ylabel('speeding_feature')
plt.xlabel('distance_feature')
plt.show()
pdb.set_trace()
```

### Sample Dataset:

1	age	Gender	Family	diet	Lifestyle	cholesterol	heartdisease
2	0	0	1	1	3	0	1
3	0	1	1	1	3	0	1
4	1	0	0	0	2	1	1
5	4	0	1	1	3	2	0
6	3	1	1	0	0	2	0
7	2	0	1	1	1	0	1
8	4	0	1	0	2	0	1
9	0	0	1	1	3	0	1
10	3	1	1	0	0	2	0
11	1	1	0	0	0	2	1
12	4	1	0	1	2	0	1
13	4	0	1	1	3	2	0
14	2	1	0	0	0	0	0
15	2	0	1	1	1	0	1
16	3	1	1	0	0	1	0
17	0	0	1	0	0	2	1
18	1	1	0	1	2	1	1
19	3	1	1	1	0	1	0
20	4	0	1	1	3	2	0



```
Text(0.5,1,'GMM Classification')
```



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