CSE 527 Problem Set 1

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1.

a)

The chances that you actually have the disease is denoted by, (D=Disease, P=Possitive)

$$\begin{split} P(D|P) &= \frac{P(PD)}{P(P)} = \frac{P(P|D) \times P(D)}{P(\bar{D}) \cdot P(P|\bar{D}) + P(D) \cdot P(P|D)} = \frac{P(P|D) \times P(D)}{(1 - P(D)) \cdot (1 - P(\bar{P}|\bar{D})) + P(D) \cdot P(P|D)} \\ &= \frac{l \cdot P(D)}{k + m \cdot P(D)} \end{split}$$

Where, l = P(P|D), $k = 1 - P(\bar{P}|\bar{D})$, $m = P(P|D) + P(\bar{P}|\bar{D}) - 1$. Thus, k, l, m are all positive numbers.

Take derivative,

$$rac{dP(D|P)}{dP(D)} = rac{l\cdot (k+m\cdot P(D)) - m\cdot l\cdot P(D)}{(k+m\cdot P(D))^2} = rac{k\cdot l}{(k+m\cdot P(D))^2}$$

The derivative is poitive, so as P(D) decrease, P(D|P) decrease, Thus the disease is rare is a good news since P(D|P) is lower.

Where,
$$P(P|D)=0.99$$
, $P(ar{P}|ar{D})=0.99$, $P(D)=1/10,000$
$$P(D|P)=0.0098$$

b)

According to the Bayes' rule:

$$P(A|B,E)P(B|E) = rac{P(A,B,E)}{P(B,E)} imes rac{P(B,E)}{P(E)}$$

$$= rac{P(A,B,E)}{P(E)}$$

$$= P(A,B|E)$$

$$\frac{P(B|A, E)P(A|E)}{P(B|E)} = \frac{\frac{P(A, B, E)}{P(A, E)} \frac{P(A, E)}{P(E)}}{\frac{P(B, E)}{P(E)}}$$
$$= \frac{P(A, B, E)}{P(B, E)}$$
$$= P(A|B, E)$$

2.

a)

Denote 1 as up, 0 as down.

x2, x3	x1 up	x1 down
1, 1	$ heta_{X1 1,1}$	1 - $ heta_{X1 1,1}$
1, 0	$ heta_{X1 1,0}$	1 - $ heta_{X1 1,0}$
0, 1	$ heta_{X1 0,1}$	1 - $ heta_{X1 0,1}$
0, 0	$ heta_{X1 0,0}$	1 - $ heta_{X1 0,0}$

```
110:0)=P(D10)
        = T. PLD:10)
        = T. P(X17, X2;, X3) 28)
        = [[ P(x, c; ] | x, c; ] x, c; ] ) [[ P(x, c; ] ) [[ P(x, c; ] ]
C). Organiz (0:0) = Organiz In (L(0:0)): organize let).
In order to argmax (10), we take derivative w.r.t. each Bxx/Pax and set to zero.
 of Oxipak, we have, \frac{M_{x_{K}|Pak}}{\Theta_{K|Pak}} = \frac{M_{pak} - M_{x_{Apak}}}{1 - \Theta_{K|Pak}} = 0
So, Oxulpak: M[th.phr)
       Penote Gal80: A, Gal4: B, Gal2: C,
 In model 1: ÂA, ÎBIA=1, ÎBIA=0, ÊCIB=1, ÎCIB=0.
 In mode( 2: ÔA, ÔB, ÔCIA=1,B=1, ÔCIA=1,B=0, ÔCIA=0,B=1, ÔCIA=0,B=0.
b). For model 1:
2(θ: D), = TCP(A, ) PCB: 1A: ) PCC7113; )

- MAINT MANA MBIA=1 MAINTABIA, MBIA=0 MAIO) - MBIA=0 MC1B=1 MB=1-MC1B=0 MBIOMAIN
- ΘΑ (1-ΘΑ) . ΘΒΙΑ=1 (1-ΘΒΙΑ=1) ΘΒΙΑ=0 (1-ΘΒΙΑ=0) . ΘC1B=1(1-ΘC1B=1) . ΘC1B=0(1-ΘC1B=0)
   For model 2:
1 (0:17) = TOP(A:) PC131) PCC-11A1131)
         C. Mathematically we should take derivative u.r.t. each & and set to zero, Computationally, we
 can compute the cpd for each node to get wh the \theta.
d.
```

2.

Codes are attached in the end.

CPDs for Model 1

Gal80(0)	0.5179
Gal80(1)	0.4821

Gal80	0	1
Gal4(0)	0.3103	0.6667
Gal4(1)	0.6897	0.3333

Gal4	0	1
Gal2(0)	0.6296	0.2931
Gal2(11)	0.3704	0.7069

CPDs for Model 2

Gal80(0)	0.5179
Gal80(1)	0.4821

Gal4(0)	0.4821
Gal4(1)	0.5179

Gal4	0	0	1	1
Gal80	0	1	0	1
Gal2(0)	0.6667	0.6111	0.2	0.5
Gal2(1)	0.3333	0.3889	0.8	0.5

Likelihood for two models.

Model	Likelihood Score
1	-218.535478586
2	-223.129027335

e.

Model 1 has higher score, so it's selected.

Code for 3.d

```
from __future__ import division, absolute_import, print_function
import numpy as np
import pandas as pd
from pgmpy.models import BayesianModel
from pgmpy.factors.discrete import TabularCPD
from pgmpy.estimators import K2Score
#%%
#Method 1
# Read data
data = pd.read table('disc-gal80-gal4-gal2.txt', index col=0).values
# Get params for Model 1
M = len(data[0])
Ma1 = np.count nonzero(data[0])
Ma0 = M - Ma1
Mb1 = np.count_nonzero(data[1])
Mb0 = M - Mb1
Mb_a1 = np.count_nonzero(data[1][np.where(data[0]==1)])
Mb_a0 = np.count_nonzero(data[1][np.where(data[0]==0)])
Mc_b1 = np.count_nonzero(data[2][np.where(data[1]==1)])
Mc b0 = np.count nonzero(data[2][np.where(data[1]==0)])
theta_a = Ma1 / M
theta_b_1 = Mb_a1 / Ma1
theta_b_0 = Mb_a = 0 / Ma0
theta c 1 = Mc b1 / Mb1
theta_c_0 = Mc_b0 / Mb0
#Compute log Likelihood
def local 1(theta, M, M ):
    return M_* np.log(theta) + (M - M_) * np.log(1 - theta)
L1 = np.sum(local_l(np.array([theta_a, theta_b_1, theta_b_0, theta_c_1, theta_c_0]),
                        np.array([M, Ma1, Ma0, Mb1, Mb0]),
                        np.array([Ma1, Mb_a1, Mb_a0, Mc_b1, Mc_b0])))
print ('log likelilhood for model 1 is ' + str(L1))
#%%
# Get params for Model 2
M = len(data[0])
Ma1 = np.count nonzero(data[0])
\#Ma0 = M - Ma1
Mb1 = np.count_nonzero(data[1])
\#Mb0 = M - Mb1
M11 = len(np.intersect1d(np.where(data[0]==1)[0], np.where(data[1]==1)[0]))
M10 = len(np.intersect1d(np.where(data[0]==1)[0], np.where(data[1]==0)[0]))
M01 = len(np.intersect1d(np.where(data[0]==0)[0], np.where(data[1]==1)[0]))
M00 = len(np.intersect1d(np.where(data[0]==0)[0], np.where(data[1]==0)[0]))
Mc_11 = np.count_nonzero(data[2][np.intersect1d(np.where(data[0]==1)[0], np.where(data[1]==1)
[0])])
Mc_10 = np.count_nonzero(data[2][np.intersect1d(np.where(data[0]==1)[0], np.where(data[1]==0)
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[0])])
Mc_01 = np.count_nonzero(data[2][np.intersect1d(np.where(data[0]==0)[0], np.where(data[1]==1)
Mc_00 = np.count\_nonzero(data[2][np.intersect1d(np.where(data[0]==0)[0], np.where(data[1]==0)]
[0])])
theta_a = Ma1 / M
theta b = Mb1 / M
theta c 11 = Mc 11 / M11
theta c 10 = Mc 10 / M10
theta_c_01 = Mc_01 / M01
theta c 00 = Mc 00 / M00
#Compute Likelihood
L2 = np.sum(local_l(np.array([theta_a, theta_b, theta_c_11, theta_c_10, theta_c_01,
theta c 00]),
                       np.array([M, M, M11, M10, M01, M00]),
                       np.array([Ma1, Mb1, Mc_11, Mc_10, Mc_01, Mc_00])))
print ('log likelilhood for model 2 is ' + str(L2))
#%%
#Method 2, Using pgmpy
# Read Data
train data = pd.DataFrame(pd.read table('disc-gal80-gal4-gal2.txt', index col=0).values.T,
columns=['Gal80', 'Gal4', 'Gal2'])
# Define Model
model1 = BayesianModel([('Gal80', 'Gal4'), ('Gal4', 'Gal2')])
model2 = BayesianModel([('Gal80', 'Gal2'), ('Gal4', 'Gal2')])
# Fit the data
model1.fit(train data)
model2.fit(train_data)
# Get CPDs
print('For Model 1')
print(model1.get_cpds('Gal80'))
print(model1.get_cpds('Gal4'))
print(model1.get_cpds('Gal2'))
print('For Model 2')
print(model2.get_cpds('Gal80'))
print(model2.get_cpds('Gal4'))
print(model2.get_cpds('Gal2'))
#Calculate K2 Score
print('K2Score of Model1 and 2')
print(K2Score(train_data).score(model1))
print(K2Score(train_data).score(model2))
```