# 40.319 Statistical and Machine Learning Spring 2020 Homework 3 Solutions

### Problem 1 (10 points)

In this problem, we will implement the EM algorithm for clustering. Start by importing the required packages and preparing the dataset.

```
import numpy as np
import matplotlib.pyplot as plt
from numpy import linalg as LA
from matplotlib.patches import Ellipse
from sklearn.datasets.samples_generator import make_blobs
from scipy.stats import multivariate_normal
NUMDATAPTS = 150
X, y = make_blobs(n_samples=NUM.DATAPTS, centers=K, shuffle=False,
                                random_state=0, cluster_std=0.6)
g1 = np.asarray([[2.0, 0], [-0.9, 1]])
g2 = np.asarray([[1.4, 0], [0.5, 0.7]])
mean1 = np.mean(X[:int(NUM.DATAPTS/K)])
mean2 = np.mean(X[int(NUM_DATAPTS/K):2*int(NUM_DATAPTS/K)])
X[:int(NUM_DATAPTS/K)] = np.einsum('nj,ij->ni',
                X[:int(NUM_DATAPTS/K)] - mean1, g1) + mean1
X[int(NUM.DATAPTS/K):2*int(NUM.DATAPTS/K)] = np.einsum('nj,ij->ni',
                X[int(NUM.DATAPTS/K):2*int(NUM.DATAPTS/K)] - mean2, g2) + mean2
X[:,1] = 4
```

- (a) Randomly initialize a numpy array mu of shape (K, 2) to represent the mean of the clusters, and initialize an array cov of shape (K, 2, 2) such that cov[k] is the identity matrix for each k. cov will be used to represent the covariance matrices of the clusters. Finally, set  $\pi$  to be the uniform distribution at the start of the program.
- (b) Write a function to perform the E-step:

```
def E_step():
    gamma = np.zeros((NUMLDATAPTS, K))
    ...
    return gamma
```

(c) Write a function to perform the M-step:

```
def M_step(gamma):
...
```

(d) Now write a loop that iterates through the E and M steps, and terminates after the change in log-likelihood is below some threshold. At each iteration, print out the log-likelihood, and use the following function to plot the progress of the algorithm:

(e) Use sklearn's KMeans module

https://scikit-learn.org/stable/modules/generated/sklearn.cluster.KMeans.html

to perform K-means clustering on the dataset, and compare both clustering results.

```
Solution. import numpy as np
import matplotlib.pyplot as plt
from numpy import linalg as LA
from matplotlib.patches import Ellipse
from sklearn.datasets.samples_generator import make_blobs
from scipy.stats import multivariate_normal
K = 3
NUM.DATAPTS = 150
X, y = make\_blobs(n\_samples=NUM\_DATAPTS, centers=K, shuffle=False,
                                         random_state=0, cluster_std=0.6)
g1 = np.asarray([[2.0, 0], [-0.9, 1]])
g2 = np.asarray([[1.4, 0], [0.5, 0.7]])
mean1 = np.mean(X[:int(NUM.DATAPTS/K)])
mean2 = np.mean(X[int(NUM_DATAPTS/K):2*int(NUM_DATAPTS/K)])
X[: int(NUM.DATAPTS/K)] = np.einsum('nj,ij->ni',
                X[:int(NUM.DATAPTS/K)] - mean1, g1) + mean1
X[int(NUM.DATAPTS/K):2*int(NUM.DATAPTS/K)] = np.einsum('nj,ij->ni',
                X[int(NUM.DATAPTS/K):2*int(NUM.DATAPTS/K)] - mean2, g2) + mean2
X[:,1] -= 4
#For 1(a)
pi = np.ones(K)
```

```
pi /= np.sum(pi)
mu = 10 * np.random.rand(K, 2) - 5
cov = np.zeros((K, 2, 2))
for i in range (K):
          cov[i] = np.eye(2)
#For 1(b)
def E_step():
          gamma = np.zeros((NUM.DATAPTS, K))
          for k in range(K):
                     numer = \; pi \, [\, k\,] \; * \; multivariate\_normal.pdf (X, \; mu[\, k\,] \;, \; cov \, [\, k\,] \,)
                     denom = 0
                     for j in range(K):
                               denom += pi[j] * multivariate_normal.pdf(X, mu[j], cov[j])
                     gamma[:,k] = numer / denom
          return gamma
#For 1(c)
def M_step(gamma):
          for k in range(K):
                     pi[k] = np.mean(gamma[:, k])
                     denom = np.sum(gamma[:,k])
                     mu\left[\,k\;,0\,\right] \;=\; np\;. \textbf{sum}\left(X\left[\,:\;,0\,\right] \;\;*\;\; gamma\left[\,:\;,\;\;k\,\right]\,\right) \;\;/\;\; denom
                     mu\,[\,k\,,1\,] \ = \ np\,. \\ \text{sum}(X\,[\,:\,,1\,] \ * \ gamma\,[\,:\,,\ k\,]\,) \ / \ denom
                     sample\_cov = np.reshape(np.einsum('ni,nj->nij', X - mu[k], X - mu[k]),
                                          (NUM.DATAPTS, 4)
                     cov[k] = np.reshape(np.sum(sample_cov * np.reshape(gamma[:, k],
                                          (NUM_DATAPTS, 1)), axis=0) / denom, (2, 2))
\#For 1(d)
def main():
           fig = plt.figure()
           plot_result()
          t = 0
           prev_ll = -5000
          {\rm c}\,{\rm u}\,{\rm r}\,{}_{\text{-}}{\rm l}\,{\rm l}\ =\ -4000
          while abs(prev_ll - cur_ll) > 0.1:
                     prev_ll = cur_ll
                     t += 1
                     \mathbf{print}\,(\,\text{"Time:}\,\,\,\,\,\,\,\,\,\,\,\,t\,\,)
                     gamma = E_step()
                     M_step(gamma)
                     cur_{-}11 = 0
                     for i in range(NUM_DATAPTS):
                               likelihood = 0
                                for k in range (K):
```

Comparison: Both clustering results look similar with slight differences (2 data points).

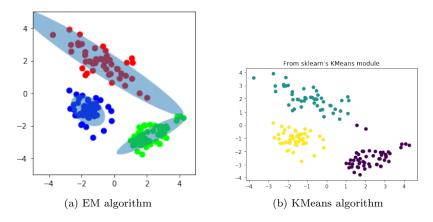


Figure 1: Comparison between both clustering results

# Problem 2 (5 points)

Let p and q be distributions on  $\{1,2,3,4,5\}$  such that  $p_1=\frac{1}{8},\ p_2=\frac{1}{2},\ p_3=p_4=p_5=\frac{1}{8},\$ and  $q_1=\frac{1}{4},\ q_2=q_3=\frac{1}{8},\ q_4=q_5=\frac{1}{4}.$ 

- (a) Compute the cross-entropy H(p,q) in bits. Is H(q,p) = H(p,q)?
- (b) Compute the entropies H(p) and H(q) in bits.
- (c) Compute the KL-divergence  $D_{KL}(p \mid q)$  in bits.

Show all working and leave your answers in fractions.

Solution.

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(a) We have

$$\begin{split} H(p,q) &= \frac{1}{8} \log_2 4 + \frac{1}{2} \log_2 8 + \frac{1}{8} \log_2 8 + \frac{1}{8} \log_2 4 + \frac{1}{8} \log_2 4 \\ &= \frac{2}{8} + \frac{3}{2} + \frac{3}{8} + \frac{2}{8} + \frac{2}{8} = \frac{21}{8}. \end{split}$$

On the other hand,

$$\begin{split} H(q,p) &= \frac{1}{4} \log_2 8 + \frac{1}{8} \log_2 2 + \frac{1}{8} \log_2 8 + \frac{1}{4} \log_2 8 + \frac{1}{4} \log_2 8 \\ &= \frac{3}{4} + \frac{1}{8} + \frac{3}{8} + \frac{3}{4} + \frac{3}{4} = \frac{22}{8} \neq H(p,q). \end{split}$$

(b) We have

$$H(p) = \frac{1}{8}\log_2 8 + \frac{1}{2}\log_2 2 + \frac{1}{8}\log_2 8 + \frac{1}{8}\log_2 8 + \frac{1}{8}\log_2 8 + \frac{1}{8}\log_2 8$$
$$= \frac{3}{8} + \frac{1}{2} + \frac{3}{8} + \frac{3}{8} + \frac{3}{8} = 2$$

and

$$\begin{split} H(q) &= \frac{1}{4} \log_2 4 + \frac{1}{8} \log_2 8 + \frac{1}{8} \log_2 8 + \frac{1}{4} \log_2 4 + \frac{1}{4} \log_2 4 \\ &= \frac{2}{4} + \frac{3}{8} + \frac{3}{8} + \frac{2}{4} + \frac{2}{4} = \frac{9}{4}. \end{split}$$

(c)  $D_{KL}(p | q) = \frac{21}{8} - 2 = \frac{5}{8}$ .

## Problem 3 (10 pts)

(a) Perform singular value decomposition (SVD) on the following matrix

$$X = \begin{bmatrix} 1 & 1 \\ 1 & 0 \\ 0 & 1 \end{bmatrix} = U \Sigma V^T.$$

(b) For a general design matrix X, why are the columns of the transformed matrix T = XV orthogonal?

Solution. (a) We compute  $X^TX$  and get

$$X^T X = \begin{bmatrix} 2 & 1 \\ 1 & 2 \end{bmatrix}.$$

The eigenvalues, in order of decreasing magnitude, are  $\lambda_1 = 3$  and  $\lambda_2 = 1$ , and the corresponding singular values are  $\sigma_1 = \sqrt{3}$ ,  $\sigma_2 = 1$ . This means that

$$\Sigma = \begin{bmatrix} \sqrt{3} & 0 \\ 0 & 1 \\ 0 & 0 \end{bmatrix}.$$

The corresponding orthonormal eigenvector of  $\lambda_1$  is  $v_1 = \left[\frac{1}{\sqrt{2}}, \frac{1}{\sqrt{2}}\right]^T$ , and that of  $\lambda_2$  is  $v_2 = \left[\frac{1}{\sqrt{2}}, \frac{-1}{\sqrt{2}}\right]^T$ . Thus

$$V = \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \\ \frac{1}{\sqrt{2}} & \frac{-1}{\sqrt{2}} \end{bmatrix}.$$

To obtain U, we compute

$$XX^T = \begin{bmatrix} 2 & 1 & 1 \\ 1 & 1 & 0 \\ 1 & 0 & 1 \end{bmatrix},$$

which we know will have eigenvalues 3, 1 and 0. The corresponding orthonormal eigenvectors are  $u_1 = \frac{1}{\sqrt{6}} \left[2,1,1\right]^T$ ,  $u_2 = \frac{1}{\sqrt{2}} \left[0,1,-1\right]^T$  and  $u_3 = \frac{1}{\sqrt{3}} \left[-1,1,1\right]^T$  respectively. Hence

$$X = \begin{bmatrix} \frac{2}{\sqrt{6}} & 0 & \frac{-1}{\sqrt{3}} \\ \frac{1}{\sqrt{6}} & \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{3}} \\ \frac{1}{\sqrt{6}} & \frac{-1}{\sqrt{2}} & \frac{1}{\sqrt{3}} \end{bmatrix} \begin{bmatrix} \sqrt{3} & 0 \\ 0 & 1 \\ 0 & 0 \end{bmatrix} \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \\ \frac{1}{\sqrt{2}} & \frac{-1}{\sqrt{2}} \end{bmatrix}^{T}.$$

(b) We have

$$T^TT = V^TX^TXV = V^T(V\Sigma^TU^T)(U\Sigma V^T)V = \Sigma^T\Sigma,$$

which is a diagonal matrix. This means that  $\langle t_i, t_j \rangle = 0$  for all  $i \neq j$  as this quantity is equal to the  $ij^{th}$  entry of the matrix.

# Problem 4 (4 points)

In this problem, we will perform principal component analysis (PCA) on sklearn's diabetes dataset. Start by importing the required packages and load the dataset.

import numpy as np
from sklearn import decomposition
from sklearn import datasets

X = datasets.load\_diabetes().data

You can find out more on how to use sklearn's PCA module from: https://scikit-learn.org/stable/modules/generated/sklearn.decomposition.PCA.html

For this problem, make sure the design matrix is first normalized to have zero mean and unit standard deviation for each column.

- (a) Write code to print the matrix V that will be used to transform the dataset, and print all the singular values.
- (b) Now perform PCA on the dataset and print out the 3 most important components for the first 10 data-points.

As this problem is short, print out your script and results, and include it with your hardcopy submission. You do not have to upload the solution.

```
Solution. import numpy as np
from sklearn import decomposition
from sklearn import datasets

X = datasets.load_diabetes().data

#For 4(a)

X = preprocessing.scale(X)
pca = decomposition.PCA(n_components=10)
pca.fit(X)

V = pca.transform(np.eye(10))
print("Transformation_matrix_V:_", V)
print("Singular_values:_", pca.singular_values_)

#For 4(b)

X = pca.transform(X)
print(X[:10, :3])
```

### Problem 5 (5 points)

An AR(2) model assumes the form

$$r_t = \phi_0 + \phi_1 r_{t-1} + \phi_2 r_{t-2} + a_t,$$

where  $a_t$  is a white noise sequence. Show that if the model is stationary, then

- (a)  $E(r_t) = \frac{\phi_0}{1 \phi_1 \phi_2}$  (assume  $\phi_1 + \phi_2 \neq 1$ );
- (b) the ACF is given by

$$\rho(1) = \frac{\phi_1}{1 - \phi_2}, \qquad \rho(s) = \phi_1 \rho(s - 1) + \phi_2 \rho(s - 2), \quad \forall s \ge 2.$$

Solution. (a) Applying expectation on both sides, we have

$$\mathbb{E}\left[r_t\right] = \mu = \phi_0 + \phi_1 \mu + \phi_2 \mu,$$

and thus  $\mu = \frac{\phi_0}{1 - \phi_1 - \phi_2}$ .

(b) Using  $\phi_0 = \mu(1 - \phi_1 - \phi_2)$ , we can rewrite the AR(2) equation as

$$r_t - \mu = \phi_1(r_{t-1} - \mu) + \phi_2(r_{t-2} - \mu) + a_t.$$

Now multiplying both sides by  $r_{t-s} - \mu$  and taking expectations, and using the fact that  $Cov(r_{t-s}, a_t) = 0$ , we get

$$Cov(r_t, r_{t-s}) = \phi_1 Cov(r_{t-1}, r_{t-s}) + \phi_2 Cov(r_{t-2}, r_{t-s}).$$

Dividing everything by  $Var(r_t)$ , we obtain

$$\rho(s) = \phi_1 \rho(s-1) + \phi_2 \rho(s-2), \quad \forall s \ge 2,$$

 $\quad \text{and} \quad$ 

$$\rho(1) = \phi_1 + \phi_2 \rho(-1) = \phi_1 + \phi_2 \rho(1) \implies \rho(1) = \frac{\phi_1}{1 - \phi_2}.$$

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