(ist1100032, ist1100070)

I. Pen-and-paper

1)



E-step: posterior probabilities for each observation

 \mathcal{H}_1 : (1,0.6,0.1)

likelihoods:
$$p(\frac{1}{2}n3 \mid K=1) = 0.3^{1} \cdot (1-0.3)^{1-1} = 0.3$$

bernoulli
$$p(\frac{1}{2}n3 \mid K=2) = 0.7^{1} \cdot (1-0.7)^{1-1} = 0.7$$

multiraride
$$\begin{cases} P(\{y_1,y_3\} \mid N=1) = (0.6,0.1) \sim N_1(y_1, \Xi_1) = 0.06657 \\ P(\{y_1,y_3\} \mid N=2) = (0.6,0.1) \sim N_2(y_2, \Xi_2) = 0.11961 \end{cases}$$

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fosteriors:

$$P(N=1|x_1) = \frac{P(x_1|N=1) \times \tilde{11}_1}{\sum_{i=1}^{2} \tilde{11}_{i}^{i} \cdot P(x_1|N=1)} = 0.1826$$

$$0.05185$$

$$P(x_{1}|x_{1}) = \frac{P(x_{1}|x_{2}) \times \tilde{x}_{2}}{\tilde{z}} = 0.8074$$

$$\frac{\text{likelihoods:}}{\text{barnoulli}} \begin{cases} P(\frac{1}{2}1\frac{1}{3}) | N=1 \\ P(\frac{1}{2}1\frac{1}{3}) | N=2 \\ P(\frac{1}{2}1\frac$$

multirative
$$\begin{cases} P(\{y_1,y_3\} | y_{=1}) = (-0.4,0.8) \sim N_1(y_1, \Xi_1) = 0.050048 \\ P(\{y_1,y_3\} | y_{=2}) = (-0.4,0.8) \sim N_2(y_2, \Xi_2) = 0.06819 \end{cases}$$

$$P(\chi_2 | K=1) = P(\{y_1\} | K=1) \times P(\{y_2, y_3\} | K=1) = 0.035034$$



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Posteriors:

$$\frac{P(N=1|x_2) = \frac{P(x_2|N=1) \times 1111}{\sum_{i=1}^{2} 11i} = 0.63134$$

$$P(K=2|X_2) = \frac{P(X_2|K_2) \times \tilde{I}Z}{\sum_{i=1}^{2} \tilde{I}_i^2 \cdot P(X_2|K=i)} = 0.36865$$

K3: (0,0.2,05)

likelihoods:
$$p(\frac{1}{2} + \frac{1}{3} + \frac{1}{4} + \frac{1}{4}) = 0.3^{\circ} \cdot (1 - 0.3)^{1 - \circ} = 0.7$$

becould
$$p(\frac{1}{2} + \frac{1}{3} + \frac{1}{4} + \frac{1}{4}) = 0.7^{\circ} \cdot (1 - 0.7)^{1 - \circ} = 0.3$$



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Posteriors:

$$P(N=1|x_3) = \frac{P(x_3|N=1) \times \tilde{11}_1}{\sum_{i=1}^{2} \tilde{11}_{i}^{i} \cdot P(x_3|N=1)} = 0.55181$$

$$P(x=1)x_3) = \frac{P(x_3|K_2) \times \tilde{x}_2}{\sum_{i=1}^{2} \tilde{x}_i, P(x_3|K=i)} = 0.44819$$

likelihoods:
$$P(\frac{1}{2},\frac{1}{3},\frac{1}{4},\frac{1}{4}) = 0.3^{1} \cdot (1-0.3)^{1-1} = 0.3$$

$$P(\frac{1}{2},\frac{1}{3},\frac{1}{4},\frac{1}{4}) = 0.7^{1} \cdot (1-0.7)^{1-1} = 0.7$$



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$$P(N=1|x_4) = \frac{P(x_4|N=1) \times \tilde{11}_1}{\sum_{i=1}^{2} \tilde{11}_i, P(x_4|N=1)} = 0.1689$$

Cluster 1

$$\frac{\frac{4}{2} \operatorname{P}(N=1) = \frac{\frac{4}{2} \operatorname{P}(N=1) \times 1}{\frac{1}{4}} = 0.38617$$

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Bernoulli Parameteus:

$$\rho_{1} = \rho(y_{1}=1 \mid X_{1}=1) = \frac{\sum_{i=1}^{4} \rho(x_{1}=1 \mid y_{1i}) \cdot y_{1i}}{\sum_{i=1}^{4} \rho(x_{1}=1 \mid X_{1i})} = \frac{0.3615}{1.5447} = 0.2340$$

Gaussim Parmeters:

$$\mu_{1} = \frac{\sum_{i=1}^{4} P(N=1|X_{i}) \cdot \{y_{1}, y_{3}\};}{\sum_{i=1}^{4} P(N=1|X_{i})} = \begin{pmatrix} 0.0409 \\ 0.7833 \end{pmatrix} - 1.5447 = \begin{bmatrix} 0.026509 \\ 0.307129 \end{bmatrix}$$

$$\begin{bmatrix} 0.21836 & -0.16282 \\ -0.16282 & 0.14837 \end{bmatrix} - 1.5447 - \begin{bmatrix} 0.14136 & -0.10541 \\ -0.10541 & 0.09605 \end{bmatrix}$$

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Cluster 2

$$\frac{2 \operatorname{rior}:}{2 \operatorname{rior}:} \widehat{I}_{1} = P(N=2) = \frac{\frac{4}{2} \operatorname{P}(N=2|X_{i})}{4} = 0.61383$$

Barnoulli Parameters

$$\rho_{2} = \rho(y_{1}=1 \mid N=2) = \frac{\sum_{i=1}^{4} \rho(N=2 \mid y_{1i}) \cdot y_{1i}}{\sum_{i=1}^{4} \rho(N=2 \mid X_{1i})} = \frac{1.6385}{2.4553} = 0.66731$$

Gaussian Parameters:

$$\mu_{2} = \frac{\frac{4}{5} P(N=2|X_{1}) \cdot \{N_{1},N_{3}\}}{\sum_{i=1}^{3} P(N=2|X_{1})} = \begin{pmatrix} 0.75905 \\ 0.051665 \end{pmatrix} - 2.41553 = \begin{pmatrix} 0.30614 \\ 0.21642 \end{pmatrix}$$

$$= \frac{\sum_{k=1}^{4} P(k=2|X_{k}) \cdot (\xi_{k},y_{3}\xi_{1} - M_{k}) \cdot (\xi_{k},y_{3}\xi_{1} - M_{k})}{\sum_{k=1}^{4} P(k=2|X_{k})} =$$

$$= \begin{bmatrix} 0.26589 & -0.217669 \\ -0.217669 & 0.233657 \end{bmatrix} - 2.41533 - \begin{bmatrix} 0.10829 & -0.08265 \\ -0.08265 & 0.10412 \end{bmatrix}$$

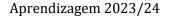


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Código auxiliar:

```
import numpy as np
from scipy.stats import bernoulli, multivariate normal
data = np.array([
    [1, 0.6, 0.1],
    [0, -0.4, 0.8],
    [0, 0.2, 0.5],
    [1, 0.4, -0.1]
1)
# Given parameters
pi = np.array([0.5, 0.5])
p = np.array([0.3, 0.7])
N1 = {
    'mean': np.array([1, 1]),
    'cov': np.array([[2, 0.5], [0.5, 2]])
N2 = {
    'mean': np.array([0, 0]),
    'cov': np.array([[1.5, 1], [1, 1.5]])
```

```
posterior_probs = np.zeros((len(data), 2))
for i in range(len(data)):
   print(f"\n\nX{ i }\n")
    likelihood_bernoulli_1 = p[0]**data[i, 0] * (1-p[0])**(1-data[i, 0])
    likelihood_bernoulli_2 = p[1]**data[i, 0] * (1-p[1])**(1-data[i, 0])
   print(f"Bernoulli Likelihood (Cluster 1): {likelihood_bernoulli_1}")
   print(f"Bernoulli Likelihood (Cluster 2): {likelihood_bernoulli_2}")
    likelihood_gaussian_1 = multivariate_normal.pdf(data[i, 1:], mean=N1['mean'], cov=N1['cov'])
    likelihood_gaussian_2 = multivariate_normal.pdf(data[i, 1:], mean=N2['mean'], cov=N2['cov'])
   print(f"Gaussian Likelihood (Cluster 1): {likelihood_gaussian_1}")
   print(f"Gaussian Likelihood (Cluster 2): {likelihood_gaussian_2}'
    likelihood_1 = likelihood_bernoulli_1 * likelihood_gaussian_1
    likelihood_2 = likelihood_bernoulli_2 * likelihood_gaussian_2
   print(f"likelihood (Cluster 1): {likelihood_1}")
   print(f"likelihood (Cluster 2): {likelihood_2}")
   denominator = pi[0] * likelihood_1 + pi[1] * likelihood_2
   print(f"Denominator): {denominator}")
   posterior_probs[i, 0] = (pi[0] * likelihood_bernoulli_1 * likelihood_gaussian_1) / denominator
    posterior_probs[i, 1] = (pi[1] * likelihood_bernoulli_2 * likelihood_gaussian_2) / denominator
print("Posterior Probabilities:\n", posterior_probs)
```





X0

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Gaussian Likelihood (Cluster 1): 0.06657529920303393 Gaussian Likelihood (Cluster 2): 0.11961837142058572

likelihood (Cluster 1): 0.019972589760910178 likelihood (Cluster 2): 0.08373285999441

Bernoulli Likelihood (Cluster 1): 0.3 Bernoulli Likelihood (Cluster 2): 0.7

```
Denominator): 0.05185272487766009
X1
Bernoulli Likelihood (Cluster 1): 0.7
Bernoulli Likelihood (Cluster 2): 0.300000000000000004
Gaussian Likelihood (Cluster 1): 0.05004888824270901
Gaussian Likelihood (Cluster 2): 0.0681905803254947
likelihood (Cluster 1): 0.035034221769896304
likelihood (Cluster 2): 0.020457174097648412
Denominator): 0.027745697933772358
X2
Bernoulli Likelihood (Cluster 1): 0.7
Bernoulli Likelihood (Cluster 2): 0.300000000000000004
Gaussian Likelihood (Cluster 1): 0.06837452355368487
Gaussian Likelihood (Cluster 2): 0.12958103481626038
likelihood (Cluster 1): 0.047862166487579405
likelihood (Cluster 2): 0.038874310444878116
Denominator): 0.04336823846622876
ΧЗ
Bernoulli Likelihood (Cluster 1): 0.3
Bernoulli Likelihood (Cluster 2): 0.7
Gaussian Likelihood (Cluster 1): 0.059046993443730274
Gaussian Likelihood (Cluster 2): 0.12450008976589248
likelihood (Cluster 1): 0.01771409803311908
likelihood (Cluster 2): 0.08715006283612474
Denominator): 0.052432080434621914
Posterior Probabilities:
 [[0.19258959 0.80741041]
 [0.63134512 0.36865488]
 [0.55181128 0.44818872]
 [0.16892423 0.83107577]]
```



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```
# M-step: Update model parameters
new_pi = np.mean(posterior_probs, axis=0)
print("PI:", new pi)
print("\n\n")
print("BERNOULLI")
new_p = np.zeros(2)
for j in range(2): # 2 clusters
   print("cluster:", j)
    p_sum = 0
    for i in range(len(data)):
        p_sum += data[i, 0] * posterior_probs[i, j]
    print("p_sum:", p_sum)
    denominator = np.sum(posterior_probs[:, j])
    print("denominator:", denominator)
    new_p[j] = p_sum / denominator
print("p:", new_p)
print("\n\n")
```

```
new_N1_mean = np.zeros(2)
new_N2_mean = np.zeros(2)
new_N1_cov = np.zeros((2, 2))
new_N2_cov = np.zeros((2, 2))
print("\nGAUSS")
for j in range(2): # 2 clusters
   print("\ncluster:", j)
   mean_sum = np.zeros(2)
    cov_sum = np.zeros((2, 2))
    for i in range(len(data)):
       mean_sum += posterior_probs[i, j] * data[i, 1:]
       cov_sum += posterior_probs[i, j] * np.outer(data[i, 1:] - [N1['mean'], N2['mean']][j], data[i, 1:] - [N1['mean'], N2['mean']][j])
    print("mean_sum:", mean_sum)
    print("cov_sum:", cov_sum)
    if j == 0:
       new N1 mean = mean sum / np.sum(posterior probs[:, j])
        new_N1_cov = cov_sum / np.sum(posterior_probs[:, j])
       new_N2_mean = mean_sum / np.sum(posterior_probs[:, j])
       new_N2_cov = cov_sum / np.sum(posterior_probs[:, j])
```



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```
# Update the model parameters
pi = new_pi
p = new_p

N1['mean'] = new_N1_mean
N2['mean'] = new_N2_mean
N1['cov'] = new_N1_cov
N2['cov'] = new_N2_cov

print("\n")
print("N1 mean:", N1['mean'])
print("N1 cov:\n", N1['cov'])
print("\n")
print("N2 mean:", N2['mean'])
print("N2 cov:\n", N2['cov'])
```

PI: [0.38616755 0.61383245]

```
BERNOULLI
```

cluster: 0

p_sum: 0.36151382107026986

denominator: 1.5446702187154808

cluster: 1

p_sum: 1.63848617892973

denominator: 2.455329781284519 p: [0.23403948 0.66731817]



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```
GAUSS
cluster: 0
mean_sum: [0.04094766 0.78334827]
cov_sum: [[ 0.21836232 -0.16281668]
[-0.16281668 0.1483696 ]]
cluster: 1
mean_sum: [0.75905234 0.51665173]
cov_sum: [[ 0.26589514 -0.21766927]
[-0.21766927 0.25565705]]
N1 mean: [0.026509 0.50712978]
N1 cov:
 [[ 0.14136501 -0.10540546]
 [-0.10540546 0.0960526 ]]
N2 mean: [0.30914476 0.2104205 ]
N2 cov:
[[ 0.10829305 -0.08865175]
 [-0.08865175 0.1041233 ]]
```

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2)
$$\times_{MU} = \begin{pmatrix} 0.3 \\ 0.7 \end{pmatrix}$$

After EM update

 $\Pi_1 = 0.38617$
 $\Lambda_2 = 0.61383$
 $\Lambda_1 = 0.23404$
 $\Lambda_2 = 0.66732$
 $\Lambda_1 = 0.50713$
 $\Lambda_1 = 0.50713$
 $\Lambda_2 = 0.14137$
 $\Lambda_1 = 0.14137$
 $\Lambda_1 = 0.50713$
 $\Lambda_2 = 0.14137$
 $\Lambda_1 = 0.14137$
 $\Lambda_2 = 0.10841$
 $\Lambda_2 = 0.108865$
 $\Lambda_2 = 0.108865$
 $\Lambda_2 = 0.108865$
 $\Lambda_2 = 0.108865$

likelihood cluster
$$1 = \int (\{y_1\} | k=1) \times \int (\{y_2,y_3\} | k=1) =$$

$$= 0,006337$$
likelihood cluster $2 = \int (\{y_1\} | k=2) \times \int (\{y_2,y_3\} | k=2) =$

$$= 0,04567$$

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Código auxiliar:

```
import numpy as np
from scipy.stats import multivariate_normal

# Given initial values
pi1 = new_pi[0]
pi2 = new_pi[1]
p1 = new_p[0]
p2 = new_p[1]
mu1 = new_N1_mean
sigma1 = new_N1_cov
mu2 = new_N2_mean
sigma2 = new_N2_cov
x_new = np.array([1, 0.3, 0.7])
```



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```
# Likelihood of x_new under each cluster
likelihood_c1 = (p1**x_new[0]) * ((1 - p1)**(1 - x_new[0])) * multivariate_normal.pdf(x_new[1:], mean=mu1, cov=sigma1)
likelihood_c2 = (p2**x_new[0]) * ((1 - p2)**(1 - x_new[0])) * multivariate_normal.pdf(x_new[1:], mean=mu2, cov=sigma2)

print("Likelihood for Cluster 1: ",likelihood_c1)
print("Likelihood for Cluster 2: ",likelihood_c2)

# Posterior probabilities
posterior_c1 = (likelihood_c1 * pi1) / (likelihood_c1 * pi1 + likelihood_c2 * pi2)
posterior_c2 = (likelihood_c2 * pi2) / (likelihood_c1 * pi1 + likelihood_c2 * pi2)

print("Posterior Probability for Cluster 1:", posterior_c1)
print("Posterior Probability for Cluster 2:", posterior_c2)
```

```
Likelihood for Cluster 1: 0.006336753293816409
Likelihood for Cluster 2: 0.045665170414993406
Posterior Probability for Cluster 1: 0.08028950846197516
Posterior Probability for Cluster 2: 0.9197104915380249
```



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Homework I V- Group 35

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II. Programming and critical analysis

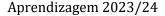
```
from scipy.io import arff
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn import metrics
from sklearn.preprocessing import MinMaxScaler
from sklearn.cluster import KMeans
from sklearn.decomposition import PCA
import warnings
warnings.filterwarnings("ignore", category=FutureWarning) # Ignore FutureWarnings
data = arff.loadarff('column_diagnosis.arff')
df = pd.DataFrame(data[0])
# Features
X = df.drop(columns='class').values
# Normalization
scaler = MinMaxScaler()
X normalized = scaler.fit transform(X)
```



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```
def purity_score(y_true, y_pred):
    confusion_matrix = metrics.cluster.contingency_matrix(y_true, y_pred)
    return np.sum(np.amax(confusion_matrix, axis=0)) / np.sum(confusion_matrix)
k_{values} = [2, 3, 4, 5]
silhouette_scores = {}
purity_scores = {}
for k in k_values:
    kmeans = KMeans(n_clusters=k, random_state=0)
    cluster_labels = kmeans.fit_predict(X_normalized)
    silhouette = metrics.silhouette_score(X_normalized, cluster_labels)
    silhouette_scores[k] = silhouette
    purity = purity_score(df['class'], cluster_labels)
    purity_scores[k] = purity
for k, silhouette in silhouette_scores.items():
   print(f"K = {k}, Silhouette Score: {silhouette}")
for k, purity in purity_scores.items():
    print(f"K = {k}, Purity Score: {purity}")
```

```
K = 2, Silhouette Score: 0.36044124340441114
K = 3, Silhouette Score: 0.29579055730002257
K = 4, Silhouette Score: 0.27442402122340176
K = 5, Silhouette Score: 0.23823928397844843
K = 2, Purity Score: 0.632258064516129
K = 3, Purity Score: 0.667741935483871
K = 4, Purity Score: 0.6612903225806451
K = 5, Purity Score: 0.6774193548387096
```





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```
n_components = 2

pca = PCA(n_components=n_components)
principal_components = pca.fit_transform(X_normalized)

explained_variance = pca.explained_variance_ratio_
print("Explained Variance for Top Two Components:", explained_variance)

vfor i, component in enumerate(range(1, n_components + 1)):
    print(f"\nTop Variables for Principal Component {component} (sorted by relevance):")
    component_weights = pca.components_[i]
    sorted_indices = np.argsort(np.abs(component_weights))[::-1]

v for idx in sorted_indices:
    print(f"Variable: {df.columns[idx]}, Weight: {component_weights[idx]}")
```

```
Explained Variance for Top Two Components: [0.56181445 0.20955953]

Top Variables for Principal Component 1 (sorted by relevance):

Variable: pelvic_incidence, Weight: 0.5916206177372234

Variable: lumbar_lordosis_angle, Weight: 0.5150847620730926

Variable: pelvic_tilt, Weight: 0.46703943896727157

Variable: sacral_slope, Weight: 0.3256888625569196

Variable: degree_spondylolisthesis, Weight: 0.21692963450485403

Variable: pelvic_radius, Weight: -0.11582397626328887

Top Variables for Principal Component 2 (sorted by relevance):

Variable: pelvic_tilt, Weight: -0.6703727595553627

Variable: pelvic_radius, Weight: -0.5810738370953603

Variable: sacral_slope, Weight: 0.4433029949470745

Variable: pelvic_incidence, Weight: 0.10003707489152235

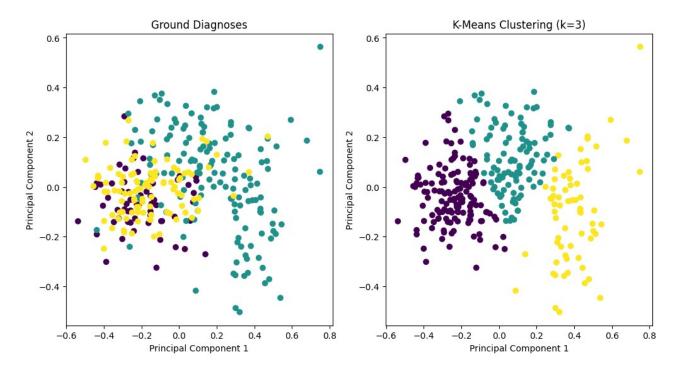
Variable: lumbar_lordosis_angle, Weight: 0.08004745059088263

Variable: degree_spondylolisthesis, Weight: 0.0045829097093999325
```



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```
kmeans = KMeans(n clusters=3, random state=0)
cluster_labels = kmeans.fit_predict(X_normalized)
# Mapping of class labels to numerical values
y = df['class']
y = y.astype(str)
class_mapping = {'Hernia': 0, 'Spondylolisthesis': 1, 'Normal': 2}
class_numerical = y.map(class_mapping)
class_numerical
# ground diagnoses
plt.figure(figsize=(12, 6))
plt.subplot(1, 2, 1)
plt.scatter(principal_components[:, 0], principal_components[:, 1], c=class_numerical)
plt.title("Ground Diagnoses")
plt.xlabel("Principal Component 1")
plt.ylabel("Principal Component 2")
# k-means clustering
plt.subplot(1, 2, 2)
plt.scatter(principal_components[:, 0], principal_components[:, 1], c=cluster_labels)
plt.title("K-Means Clustering (k=3)")
plt.xlabel("Principal Component 1")
plt.ylabel("Principal Component 2")
plt.show()
```





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END