

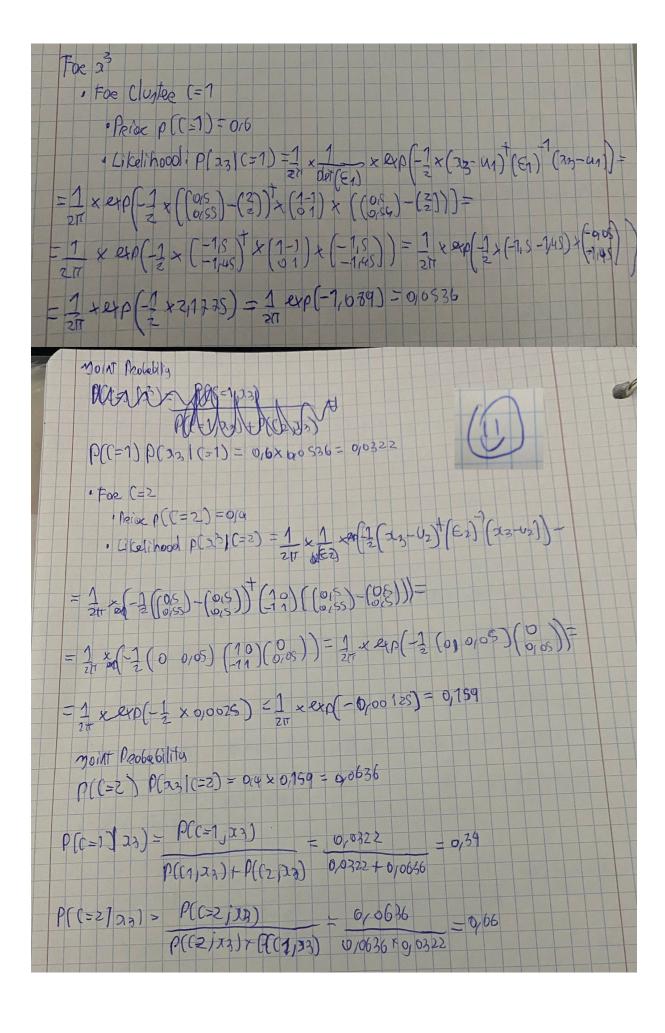
## LEIC-T 2023/2024 Aprendizagem - Machine Learning

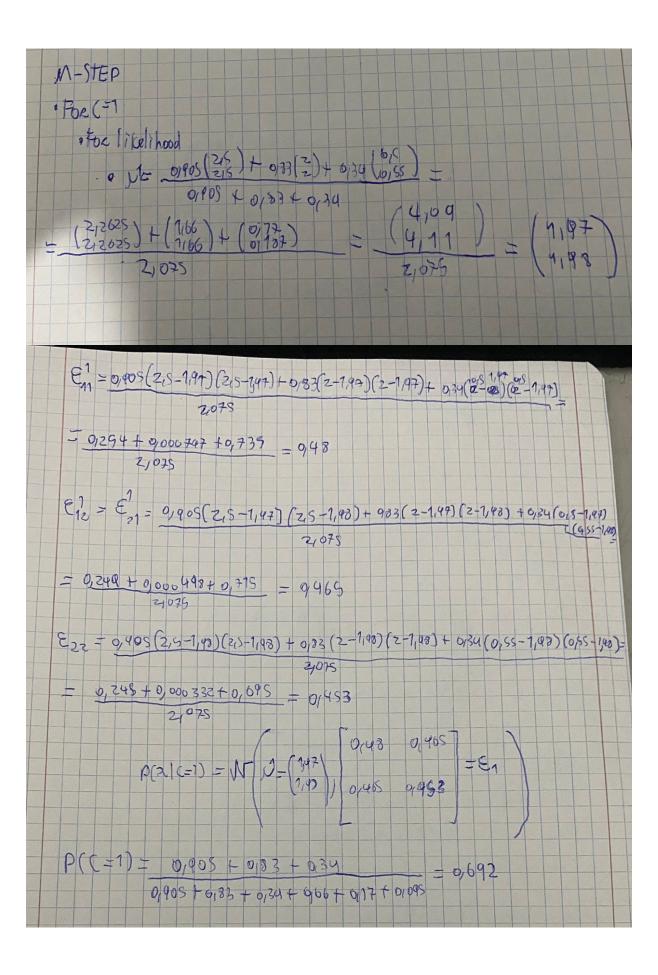
Homework IV - Group 007

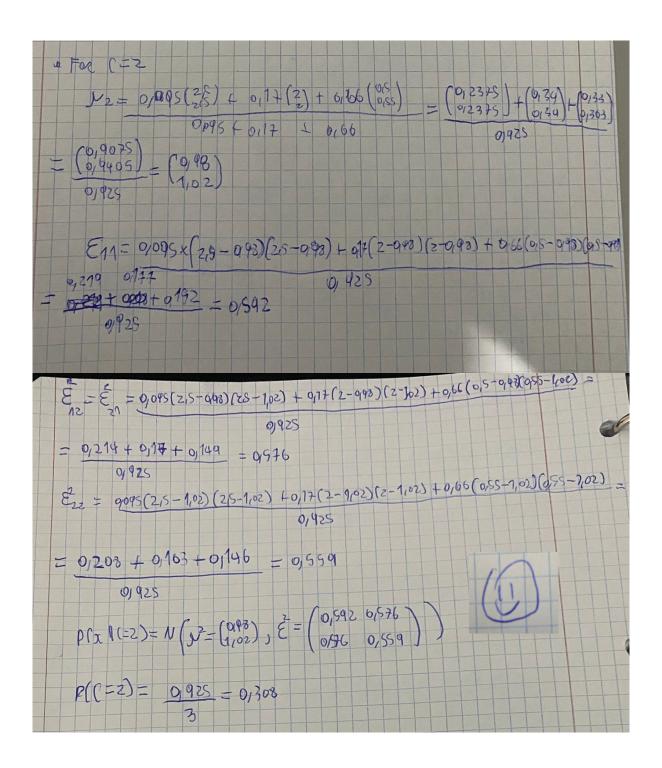
Miguel Teixeira - 103449

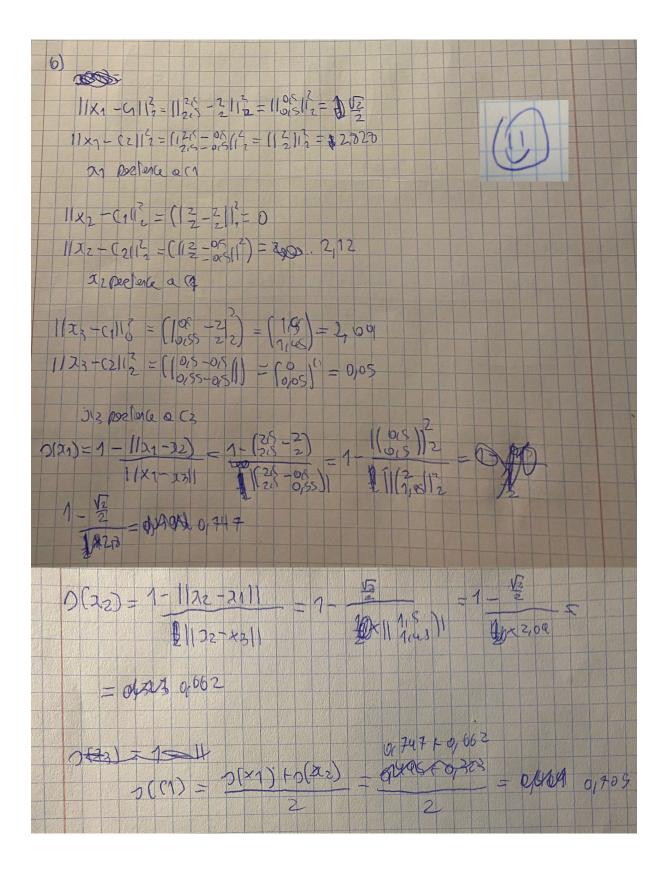
Rodrigo Alves - 103299

```
· FOR Cluster C=2
                                                                          Peior p((=2)=014
                                                                   1 Like (ihax): P(x1 (=1) = 1 7 P(-1 (x1-42) (E2) (x1-42) =
                 =\frac{1}{2\pi} \times \frac{1}{1} \exp\left(-\frac{1}{2}\left(\frac{2}{2}\right) - \frac{0}{0}\right) \left(\frac{1}{1}\right) \left(\frac{2}{2}\right) - \frac{0}{0}\right) + \frac{1}{2}
      = \frac{1}{2\pi} \times \exp\left(-\frac{1}{2}(2)^2\right) \left(\frac{10}{10}(2)\right) = \frac{1}{2\pi} \times \exp\left(-\frac{1}{2}(2)^2\right) \left(\frac{2}{0}\right) = \frac{1}{2\pi} \times \exp\left(-\frac{1}{2}(2)^2\right) = \frac{1}{2\pi} \times \exp\left(-\frac{1}{2}(2)^2\right)
        = 0,022
                                                     · Doint peobolility: P(C=2) P(x11C=2) = 0,4 ×0,022 = 0,00 23
     P((=1))+P((=2))+P((=2))+P((=2))
P((=2/21) = P((=2/21)) = 5,000 0,0000 = 9,095
                                                                                                                                                                                                                                                                                                                                                                                  6/039 4 0/0633
                                                   Toe 22
                                                                  · Foe ( lugloe C= 1
                                                                                       · Peioe p((=1)=0,6
                                                                                        · Likelihood: p(22 (=1) = 1 1 x0xp(-7 x((3)-(3))x(1)) x((3)-(3))
                                               =\frac{1}{2\pi}\times\exp(-\frac{1}{2}(00)(\frac{1-1}{2})(0))=\frac{1}{2\pi}\times\exp(-\frac{1}{2}x_0)=\frac{1}{2\pi}\times1-\frac{1}{2\pi}
                                                                                  · Doint DROBABILITY : P(C=4) P(221C=1) = 016 × 1 = 0,099
                                                      · For Cluster (=2
                                                                                · Price p((-2)-014
                                                                              ·Licelinood: P(xz) (=z)=1x1 x2xp(-1x(jz-uz) (Ez) (zz-uz)=
                              =\frac{1}{2\pi} \times exp(-\frac{1}{2} \times ((\frac{2}{2}) - (\frac{0}{15})) + (\frac{1}{15})(\frac{1}{15})) = \frac{1}{2\pi} \times exp(-\frac{1}{2} \times (\frac{1}{15} \times \frac{1}{15})(\frac{1}{15})) = \frac{1}{2\pi} \times exp(-\frac{1}{2} \times (\frac{1}{15} \times \frac{1}{15})(\frac{1}{15})(\frac{1}{15} \times \frac{1}{15})(\frac{1}{15})
                             = \frac{1}{2\pi} \exp(-\frac{1}{3} \times z_1 z_5) = \frac{1}{2\pi} \times \exp(-\frac{1}{125}) = 0,0517
                                                                 · Moint Poo bebility: P(C=1) P(r=1(=2) = 09 x0,051= 0,02
                                    P(C=1) \times 2) = P(C_1, x_2) + P(C_2, x_2) = 0.000 0.000 0.000 = 0.000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 = 0.0000 0.0000 =
                          P((=2/x2) = P(cz/x2)
                                                                                                                                P((1/22) + P(22/22) = 0/18 0/02 = 0/17
```









## **II Software Experiments**

(a)

```
import matplotlib.pyplot as plt
from sklearn import metrics, datasets, cluster, mixture
from sklearn.decomposition import PCA
# Load the wine dataset
data = datasets.load wine()
X, y = data.data, data.target
# Initialize a list to store silhouette scores for k-means and EM
kmeans_silhouettes = []
em_silhouettes = []
k_{values} = [2, 3, 4]
for k in k_values:
    # K-means clustering
    kmeans_algo = cluster.KMeans(n_clusters=k, algorithm='elkan', n_init=10)
    kmeans_model = kmeans_algo.fit(X)
    kmeans_labels = kmeans_model.labels_
    kmeans_silhouette = metrics.silhouette_score(X, kmeans_labels, metric='euclidean')
    kmeans_silhouettes.append(kmeans_silhouette)
    em_algo = mixture.GaussianMixture(n_components=k, covariance_type='full', n_init=10)
    em_model = em_algo.fit(X)
    em_labels = em_model.predict(X)
    em_silhouette = metrics.silhouette_score(X, em_labels, metric='euclidean')
    em_silhouettes.append(em_silhouette)
for i, k in enumerate(k_values):
    print(f'K-Means with {k} clusters - Silhouette: {kmeans_silhouettes[i]}')
    print(f'EM with {k} clusters - Silhouette: {em_silhouettes[i]}')
# Now, perform PCA with two components and repeat the clustering experiments
pca = PCA(n_components=2)
X_pca = pca.fit(X).transform(X)
# Initialize new lists for silhouette scores with PCA
kmeans_silhouettes_pca = []
em_silhouettes_pca = []
```

```
for k in k_values:
    # K-means clustering with PCA
    kmeans_algo_pca = cluster.KMeans(n_clusters=k, algorithm='elkan', n_init=10)
    kmeans_model_pca = kmeans_algo_pca.fit(X_pca)
    kmeans_labels_pca = kmeans_model_pca.labels_
    kmeans_silhouette_pca = metrics.silhouette_score(X_pca, kmeans_labels_pca, metric='euclidean')
    kmeans_silhouettes_pca.append(kmeans_silhouette_pca)
    # EM clustering with PCA
    em_algo_pca = mixture.GaussianMixture(n_components=k, covariance_type='full', n_init=10)
    em_model_pca = em_algo_pca.fit(X_pca)
    em_labels_pca = em_model_pca.predict(X_pca)
    em_silhouette_pca = metrics.silhouette_score(X_pca, em_labels_pca, metric='euclidean')
    em_silhouettes_pca.append(em_silhouette_pca)
for i, k in enumerate(k_values):
    print(f'K-Means with PCA and {k} clusters - Silhouette: {kmeans_silhouettes_pca[i]}')
    print(f'EM with PCA and {k} clusters - Silhouette: {em_silhouettes_pca[i]}')
```

K-Means with 2 clusters - Silhouette: 0.6568536504294317 EM with 2 clusters - Silhouette: 0.5510515269549274

K-Means with 3 clusters - Silhouette: 0.5711381937868838 EM with 3 clusters - Silhouette: 0.34726590057721557

K-Means with 4 clusters - Silhouette: 0.5620323449580341 EM with 4 clusters - Silhouette: 0.32712767339444565

K-Means with PCA and 2 clusters - Silhouette: 0.6572176888364498 EM with PCA and 2 clusters - Silhouette: 0.6419971749179373

K-Means with PCA and 3 clusters - Silhouette: 0.5722554756855063 EM with PCA and 3 clusters - Silhouette: 0.2623333079949891

K-Means with PCA and 4 clusters - Silhouette: 0.5633930017441461 EM with PCA and 4 clusters - Silhouette: 0.16208459426663982

\_\_\_\_\_

## The ideal k-values for K-Means and EM clustering are as follows: Without PCA:

- For K-Means, the ideal k-value is 2, with a silhouette score of 0.657.
- For EM, the ideal k-value is 2, with a silhouette score of 0.551.

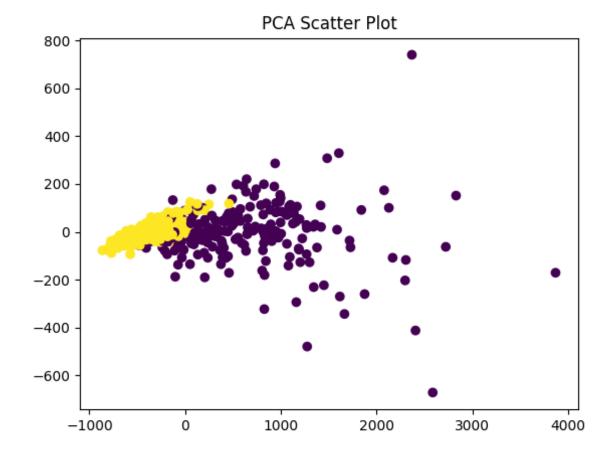
## With PCA (2 components):

- For K-Means with PCA, the ideal k-value is 2, with a silhouette score of 0.657.
- For EM with PCA, the ideal k-value is 2, with a silhouette score of 0.642.

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The ideal k-values for K-Means and EM are the same when using PCA and when not using PCA in this specific case. However, it's important to note that the ideal k-value can vary depending on the dataset and the specific problem we are trying to solve. In this case, k=2 seems to provide the best clustering results based on silhouette scores.

```
import matplotlib.pyplot as plt
from sklearn import metrics, datasets, cluster, mixture
from sklearn.decomposition import PCA
# Load the breast_cancer dataset
data = datasets.load_breast_cancer()
X, y = data.data, data.target
# K-means clustering with 2 clusters
kmeans_algo = cluster.KMeans(n_clusters=2, algorithm='elkan', n_init=10)
kmeans_model = kmeans_algo.fit(X)
kmeans_labels = kmeans_model.labels_
kmeans_silhouette = metrics.silhouette_score(X, kmeans_labels, metric='euclidean')
# EM clustering with 2 clusters
em_algo = mixture.GaussianMixture(n_components=2, covariance_type='full', n_init=10)
em_model = em_algo.fit(X)
em_labels = em_model.predict(X)
em_silhouette = metrics.silhouette_score(X, em_labels, metric='euclidean')
# Perform PCA with two components
pca = PCA(n_components=2)
X_pca = pca.fit(X).transform(X)
# Scatter plot of PCA mapped data
plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y)
plt.title("PCA Scatter Plot")
plt.show()
print("K-Means Silhouette Score: ", kmeans_silhouette)
print("EM Silhouette Score: ", em_silhouette)
```



K-Means Silhouette Score: 0.6972646156059464 EM Silhouette Score: 0.5315172918032405

Based on the silhouette values and the scatter plot of the PCA-mapped data, K-Means clustering with 2 clusters appears to be better, as it has a higher silhouette score (0.697) compared to EM clustering (0.532).