

Q2

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1 Question 2

1.1 Part 2

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[ ]: # Import useful libraries. Feel free to use sklearn.
     from sklearn.datasets import make_blobs

     # Construct a 2D toy dataset for clustering.
     X, _ = make_blobs(n_samples=1000,
                       centers=[[0, 0], [1, 1], [-1, 1], [-1, -1], [1, -1]],
                       cluster_std=[0.2, 0.3, 0.3, 0.3, 0.3],
                       random_state=26)

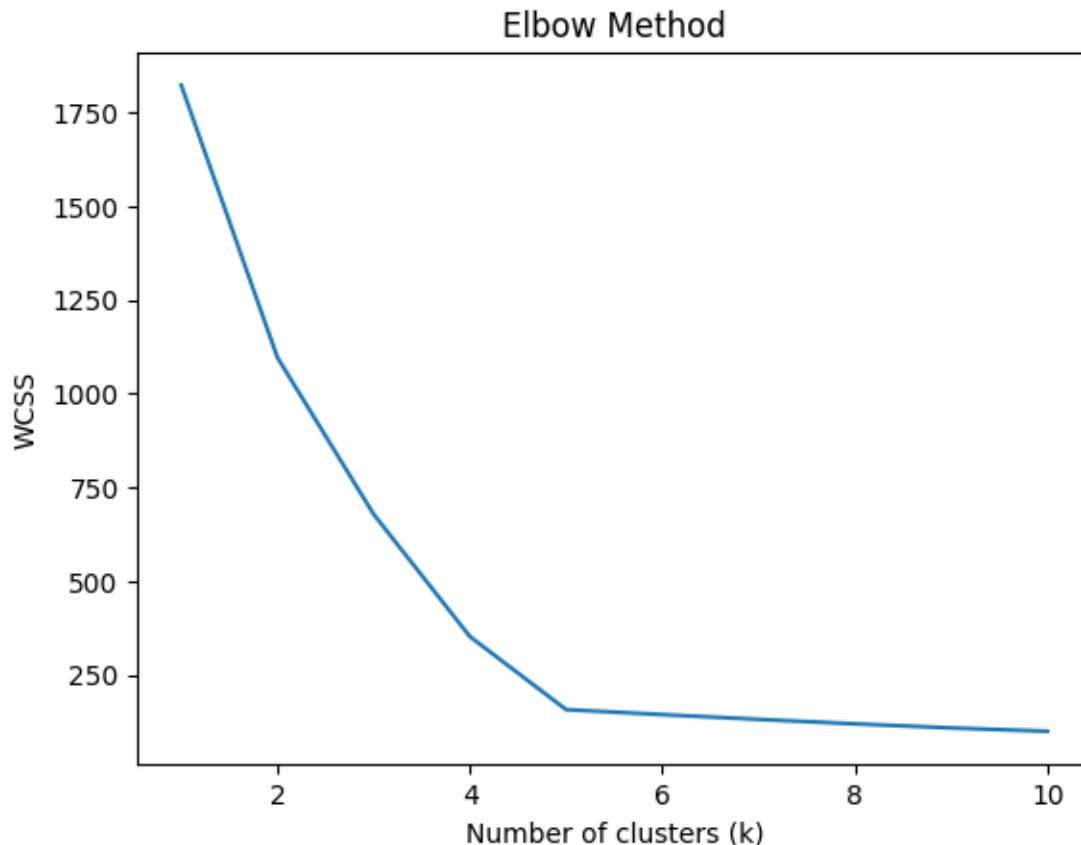
     # Conduct clustering on X using k-Means, and determine the best k with the
     ↪ elbow method.

[ ]: import matplotlib.pyplot as plt
     from sklearn.cluster import KMeans
     import numpy as np

     distances = []

     for i in range(1, 11):
         kmeans = KMeans(n_clusters=i, init='k-means++', max_iter=300, n_init=10,
         ↪ random_state=0)
         kmeans.fit(X)
         distances.append(kmeans.inertia_)

     plt.plot(range(1, 11), distances)
     plt.title('Elbow Method')
     plt.xlabel('Number of clusters (k)')
     plt.ylabel('WCSS')
     plt.show()
```



As can be seen in the graph, for the first values of k , the improvement of the WCSS value calculated by the elbow method is considerable. For values of k greater than 5, the resulting value continues to decrease, but at a much slower rate than in the previous cases.

In this method, we are looking for the right point to go from a big improvement to a not so big improvement. This is because although the result continues to improve, for each larger value of k , the computational cost increases, and a balance is sought between this computational cost and a k -means result that is as accurate as possible.

Therefore, as can be seen in the graph, the best value is $k=5$. The values of the centroids of each cluster are shown in the following graph, with all points related with each cluster (coloured for each one with a different color):

```
[ ]: k = 5
kmeans = KMeans(n_clusters=k, init='k-means++', max_iter=300, n_init=10,
↳ random_state=0)
pred_y = kmeans.fit_predict(X)

# Plotting the clusters
plt.scatter(X[:,0], X[:,1], c=pred_y, cmap='viridis')
```

```
plt.scatter(kmeans.cluster_centers_[ :, 0], kmeans.cluster_centers_[ :, 1],  
            s=300, c='red')  
plt.title('Clustering Result with k={}'.format(k))  
plt.xlabel('Feature 1')  
plt.ylabel('Feature 2')  
plt.show()
```

