```
In [114]: import numpy as np
           import matplotlib.pyplot as plt
           import pandas as pd
           import random
           %matplotlib inline
           from sklearn.decomposition import PCA
           from sklearn.preprocessing import StandardScaler,RobustScaler
           import seaborn as sns
           from sklearn.metrics import mean_squared_error as mse
           from scipy.stats import mode
           from sklearn.cluster import KMeans
           from sklearn.datasets import load_digits
           import warnings
           warnings.filterwarnings('ignore')
  In [ ]: def find_dunn_index(c,d,df):
               numerator=np.inf
               denominator=-1*np.inf
               #minimize the intercluster distance
               for i in range(len(c)):
                   for k in range(len(c)):
                     if k!=i:
                        ans=0
                        for j in range(len(c[0])):
                            ans+=(c[i][j]-c[k][j])**2
                        numerator=min(numerator, ans)
               #maximize the
               return ans
  In [4]: def find_centroids(k,n):
               random.seed(5)
               idx=[]
               for i in range(k):
                   idx.append(random.randint(0,n-1))
               return idx
  In [5]: def find_clusters(df,c,k,n):
               dist=[[] for _ in range(k)]
               for i in range(n):
                   temp=[]
                   for j in range(k):
                       dist_temp = np.sum((df[i]-c[j])**2)
                        temp.append(dist_temp)
                   location=temp.index(min(temp))
                   dist[location].append(i)
               return dist
  In [6]: def find_new_centroids(df, dist, c, k, m):
               for i in range(k):
               temp_mean=[0 for _ in range(m)]
               for j in range(len(dist[i])):
                 id=dist[i][j]
                   temp=0
                   for kk in range(m):
                       temp_mean[kk]+=df[id][kk]/len(dist[i])
               for j in range(m):
                   c[i][j]=temp_mean[j]
               return c
 In [26]: def plot_2D(dist,df,c,k):
               plt.figure(figsize=[10,6])
               distx=[[] for i in range(k)]
               disty=[[] for i in range(k)]
               for i in range(k):
                   for j in range(len(dist[i])):
                       id=dist[i][j]
                        distx[i].append(df[id][0])
                        disty[i].append(df[id][1])
               cx=[];cy=[]
               for i in range(k):
                   cx.append(c[i][0])
                   cy.append(c[i][1])
                   sns.scatterplot(distx[i], disty[i], markers='o', s=200)#, c=next(clr))
               sns.scatterplot(cx,cy,color='.2',marker='*',s=400)
               plt.grid()
               plt.xlabel('X', fontsize=15)
               plt.ylabel('Y', fontsize=15)
In [108]: def k_means(df, k, plot):
               # clr= iter(['b', 'r','c', 'm', 'g', 'tab:orange', 'tab:brown', 'tab:pink'])
               n=df.shape[0]
               m=df.shape[1]
               #indices of centroid
               idx=find_centroids(k,n)
               #to store the centroids kxm
               c=[[] for _ in range(k)]
               cprev=[[] for _ in range(k)]
               for i in range(k):
                   for j in range(m):
                        c[i].append(df[idx[i]][j])
                        cprev[i].append(df[idx[i]][j])
               #to get the points in a given cluster kxdim
               #just store the indices in the original array
               dist=[[] for _ in range(k)]
               flag=False
               while flag==False and itr!=1000:
                   #k clusters with elements in the cluster
                   dist=find_clusters(df,c,k,n)
                   #store previous centroids
                   cprev=np.copy(c)
                   #finding the new centroids
                   c=find_new_centroids(df,dist,c,k,m)
                   count=0
                   for i in range(k):
                        for j in range(len(c[0])):
                            if c[i][j]==cprev[i][j]:
                                count+=1
                   if count==k*len(c[0]):
                       break
                   itr+=1
               if plot==True:
                   plot_2D(dist,df,c,k)
               return c, dist
  In [9]: def find_inertia(c,d,df):
               ans=0
               for i in range(len(c)):
                   indices=d[i]
                   for kk in range(len(indices)):
                       for j in range(len(c[0])):
                            ans+=(c[i][j]-df[indices[kk]][j])**2
               return ans
          Q1
 In [28]: x = [10, 14, 8, 12, 15, 12, 15, 17, 5, 18, 22, 25, 35, 21, 39, 27, 25, 33, 30, 36]
          y = [8, 25, 10, 30, 35, 12, 14, 15, 22, 32, 2, 21, 35, 7, 15, 29, 33, 23, 17, 11]
 In [29]: df=[]
           df.append(x)
           df.append(y)
           df=np.array(df).transpose()
           scaler = StandardScaler()
           df_scaled = scaler.fit_transform(df)
 In [30]: SSE_q1=[]
           for i in range(1,n):
               c_q1, dist_q1=k_means(df, i, False)
               SSE_q1.append(find_inertia(c_q1, dist_q1, df))
 In [31]: x_axis=np.arange(1,20)
           plt.figure(figsize=[10,6])
           plt.plot(x_axis, SSE_q1, 'k-o')
           plt.grid()
           plt.ylabel('SSE', fontsize=15)
           plt.xlabel('K(number of clusters)', fontsize=15)
          plt.title('Elbow Curve', fontsize=15)
 Out[31]: Text(0.5, 1.0, 'Elbow Curve')
                                               Elbow Curve
              4000
              3500
              3000
              2500
           2000
2000
              1500
              1000
               500
                          2.5
                                                                              17.5
                                  5.0
                                                             12.5
                                                                      15.0
                                           K(number of clusters)
 In [32]: diff_x=np.arange(2,20)
           \label{eq:diff_sse_q1} \mbox{diff\_sse\_q1[i-1]-SSE\_q1[i]} \mbox{ for } \mbox{i in } \mbox{range(1,len(SSE\_q1))]}
           plt.figure(figsize=[10,6])
           plt.plot(diff_x, diff_sse_q1, 'k-o')
           plt.grid()
           plt.ylabel('SSE[i-1] - SSE[i]', fontsize=15)
           plt.xlabel('i', fontsize=15)
           plt.title('Optimal cluster size', fontsize=15)
 Out[32]: Text(0.5, 1.0, 'Optimal cluster size')
                                            Optimal cluster size
              1400
              1200
              1000
           SSE[i-1] - SSE[i]
               800
               600
               400
               200
                      2.5
                                         7.5
                                                  10.0
                                                            12.5
                                                                     15.0
                                                                              17.5
          We can see that at i=6 we get almost zero difference and after that we get such a case for i>10 for which the computation time increases significantly hence, K=6 is a good choice for number of clusters.
 In [33]: c_q1, dist_q1=k_means(df, 6, True)
              35
              30
              25
              15
              10
               5
                                                                  30
                                      15
                                                        25
                                                   Χ
           Q2
 In [17]: data_digits=load_digits().data
           data_labels=load_digits().target
 In [82]: data_digits.shape
 Out[82]: (1797, 64)
 In [18]: | scaler = StandardScaler()
           data_scaled = scaler.fit_transform(data_digits)
          In built function
 In [19]: kmeans = KMeans(n_clusters = 10, init='k-means++')
           kmeans.fit(data_scaled)
           SSE_kmeans=kmeans.inertia_
          print('SSE using inbuilt function:',SSE_kmeans)
          SSE using inbuilt function: 69426.25372371473
 In [76]: fig, ax = plt.subplots(2, 5, figsize=(8, 3))
           centers = kmeans.cluster_centers_.reshape(10, 8, 8)
           for axi, center in zip(ax.flat, centers):
               axi.set(xticks=[], yticks=[])
               axi.imshow(center, interpolation='nearest', cmap=plt.cm.binary)
          My function
In [109]: c_q2, dist_q2=k_means(data_scaled, 10, False)
           SSE_q2_myfunc = find_inertia(c_q2, dist_q2, data_scaled)
           print('SSE using my function(k_means):', SSE_q2_myfunc)
          SSE using my function(k_means): 69476.70340699745
In [110]: fig, ax = plt.subplots(2, 5, figsize=(8, 3))
           centers = np.array(c_q2).reshape(10, 8, 8)
           for axi, center in zip(ax.flat, centers):
               axi.set(xticks=[], yticks=[])
               axi.imshow(center, interpolation='nearest', cmap=plt.cm.binary)
          We can see that above 10 centers/centroids identified can be associated to actual digits/labels as:
          8/3 7 6 2 9
          8/3 5 4 0 1
          Although they are not very clear the SSE of the inbuilt and my function is comparable.
          To visualize the clustering we can use PCA for 2 components and then apply k means
          clustering on the obtained data
In [111]: estimator = PCA(n_components=2)
           x_pca = estimator.fit_transform(data_scaled)
In [112]: kmeans = KMeans(n_clusters = 10, init='k-means++')
           kmeans.fit(x_pca)
           SSE_kmeans_pca=kmeans.inertia_
           print('SSE using inbuilt function:',SSE_kmeans_pca)
          SSE using inbuilt function: 2815.3317395769977
In [113]: c_pca, dist_pca=k_means(x_pca, 10, True)
           SSE_pca_myfunc = find_inertia(c_pca, dist_pca, x_pca)
          print('SSE using my function(k_means):',SSE_pca_myfunc)
          SSE using my function(k_means): 2818.2310148725037
              10.0
               7.5
               5.0
               2.5
               0.0
              -2.5
              -5.0
              -7.5
                                                                           7.5
                            -5.0
                                     -2.5
                                                    Χ
          For the above plot X, Y represents the two principal components.
          To verify the correctness of my function elbow curve is constructed for 1-10 clusters to
          show the similarity in SSE between the inbuilt and my function
 In [54]: | SSE_inbuilt = []
           for cluster in range(1,11):
               kmeans = KMeans(n_clusters = cluster, init='k-means++')
               kmeans.fit(data_scaled)
               SSE_inbuilt.append(kmeans.inertia_)
In [116]: | SSE_q2=[]
           n=11
           for i in range(1,n):
             c_q2, dist_q2=k_means(data_scaled, i, False)
             SSE_q2.append(find_inertia(c_q2, dist_q2, data_scaled))
In [117]: | x_axis=[i for i in range(1,11)]
           plt.figure(figsize=[10,6])
           plt.plot(x_axis, SSE_q2, 'k-o', label='My Function')
           plt.plot(x_axis, SSE_inbuilt, 'r-o', label='Inbuilt')
           plt.grid()
          plt.ylabel('SSE', fontsize=15)
           plt.xlabel('K(number of clusters)', fontsize=15)
           plt.title('Elbow Curve', fontsize=15)
          plt.legend(fontsize=12)
Out[117]: <matplotlib.legend.Legend at 0x7fdf074c6190>
                                                 Elbow Curve
              110000
                                                                            My Function
                                                                            → Inbuilt
              105000
              100000
               95000
               90000
               85000
               80000
               75000
               70000
                                             K(number of clusters)
```

CS306: DATA ANALYSIS AND VISUALIZATION

LAB 8: K-Means Clustering

STUDENT ID: 201801407

NAME: PRATVI SHAH

between the sum of square errors of the two implementation.

SSE shows that the k_means() function and the inbuilt function are comparable as there is very little difference

In []: