K Means Clustering Algorithms Implementation

```
In [1]: import matplotlib.pyplot as plt
         from sklearn.datasets import make_blobs
         import pandas as pd
         import numpy as np
         %matplotlib inline
In [2]: X, y = make_blobs(n_samples=1000,centers=3, n_features=2, random_state=23)
In [3]: X.shape
Out[3]: (1000, 2)
         plt.scatter(X[:,0],X[:,1])
In [6]:
Out[6]: <matplotlib.collections.PathCollection at 0x1ca2588e750>
         12.5
         10.0
          7.5
          5.0
          2.5
          0.0
        -2.5
        -5.0
        -7.5
                                                                               7.5
                   -7.5
                             -5.0
                                       -2.5
                                                 0.0
                                                           2.5
                                                                     5.0
In [5]: from sklearn.model_selection import train_test_split
         X_train, y_train, X_test, y_test = train_test_split(X, y , test_size=0.33, rando
In [9]: from sklearn.cluster import KMeans
In [41]: # Manual process
         ## Elbow method to select the k value
         wcss = []
         for k in range(1,11):
```

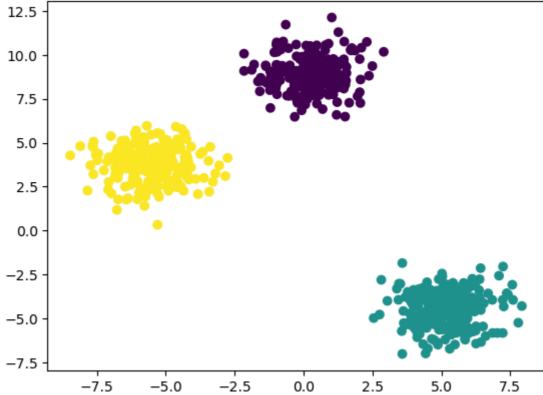
```
kmeans = KMeans(n_clusters=k,init='k-means++')
             kmeans.fit(X_train)
             wcss.append(kmeans.inertia_)
In [42]: wcss
Out[42]: [34827.57682552021,
           7935.437286145418,
           1319.2730531585605,
           1140.4677884655123,
           1006.8745739258457,
           902.3383256536542,
           770.7924368518035,
           709.6055857560772,
           644.8743127558693,
           532.3776756218431]
In [44]: ## plot the elbow curve
         plt.plot(range(1,11),wcss)
         plt.xticks(range(1,11))
         plt.xlabel("Number of Clusters")
         plt.ylabel("WCSS")
         plt.show()
           35000
           30000
           25000
           20000
           15000
           10000
            5000
                0
                            2
                                    3
                                                  5
                     1
                                                                              9
                                                                                     10
                                                                       8
                                            Number of Clusters
In [45]:
         kmeans = KMeans(n_clusters=3, init="k-means++")
In [46]: y_labels = kmeans.fit_predict(X_train)
In [59]: y_test_label = kmeans.predict(X_test)
```

```
ValueError
                                          Traceback (most recent call last)
Cell In[59], line 1
----> 1 y test label = kmeans.predict(X test)
File ~\anaconda3\Lib\site-packages\sklearn\cluster\_kmeans.py:1085, in _BaseKMean
s.predict(self, X)
  1067 """Predict the closest cluster each sample in X belongs to.
  1069 In the vector quantization literature, `cluster_centers_` is called
   (\ldots)
  1081
           Index of the cluster each sample belongs to.
  1082 """
  1083 check_is_fitted(self)
-> 1085 X = self._check_test_data(X)
  1087 # sample weights are not used by predict but cython helpers expect an arr
av
   1088 sample_weight = np.ones(X.shape[0], dtype=X.dtype)
File ~\anaconda3\Lib\site-packages\sklearn\cluster\_kmeans.py:944, in _BaseKMean
s._check_test_data(self, X)
   943 def _check_test_data(self, X):
--> 944
          X = validate_data(
   945
                self,
   946
                Χ,
   947
                accept_sparse="csr",
   948
               reset=False,
   949
               dtype=[np.float64, np.float32],
                order="C",
   950
   951
                accept_large_sparse=False,
   952
            )
   953
           return X
File ~\anaconda3\Lib\site-packages\sklearn\utils\validation.py:2954, in validate
data(_estimator, X, y, reset, validate_separately, skip_check_array, **check_para
ms)
  2952
                out = X, y
  2953 elif not no_val_X and no_val_y:
           out = check array(X, input name="X", **check params)
-> 2954
   2955 elif no val X and not no val y:
   2956
           out = _check_y(y, **check_params)
File ~\anaconda3\Lib\site-packages\sklearn\utils\validation.py:1091, in check_arr
ay(array, accept_sparse, accept_large_sparse, dtype, order, copy, force_writeabl
e, force_all_finite, ensure_all_finite, ensure_non_negative, ensure_2d, allow_nd,
ensure min samples, ensure min features, estimator, input name)
  1084
                else:
   1085
                    msg = (
  1086
                        f"Expected 2D array, got 1D array instead:\narray={arra
y}.\n"
   1087
                        "Reshape your data either using array.reshape(-1, 1) if "
  1088
                        "your data has a single feature or array.reshape(1, -1) "
  1089
                        "if it contains a single sample."
  1090
                    )
                raise ValueError(msg)
-> 1091
  1093 if dtype_numeric and hasattr(array.dtype, "kind") and array.dtype.kind in
"USV":
   1094
            raise ValueError(
   1095
                "dtype='numeric' is not compatible with arrays of bytes/strings."
                "Convert your data to numeric values explicitly instead."
   1096
```

```
1097
ValueError: Expected 2D array, got 1D array instead:
array=[1. 1. 1. 2. 2. 2. 1. 2. 1. 1. 0. 0. 1. 1. 0. 1. 0. 2. 0. 0. 2. 0. 0. 2.
 2. 1. 1. 2. 2. 1. 2. 2. 1. 2. 0. 0. 0. 2. 2. 1. 1. 2. 1. 1. 2. 0. 0. 1.
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 1. 0. 1. 0. 2. 2. 2. 2. 1. 1. 0. 1. 0. 0. 1. 0. 1. 0. 1. 0. 1. 2. 1. 1.
 2. 0. 0. 0. 0. 2. 0. 1. 2. 2. 0. 1. 0. 1. 2. 0. 1. 2. 2. 2. 2. 0. 1. 0.
 1. 1. 0. 2. 1. 2. 1. 0. 2. 2. 1. 2. 2. 2. 2. 2. 2. 2. 1. 1. 0. 0. 1. 2.
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 2. 1. 1. 0. 0. 1. 0. 0. 0. 0. 1. 2. 1. 1. 0. 2. 0. 0. 1. 2. 1. 2. 1. 2.
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 2. 2. 2. 1. 0. 2. 2. 2. 2. 1. 0. 1. 0. 1. 0. 1. 0. 2. 0. 1. 2. 1.].
Reshape your data either using array.reshape(-1, 1) if your data has a single fea
ture or array.reshape(1, -1) if it contains a single sample.
```

```
In [52]: plt.scatter(X_train[:,0],X_train[:,1],c=y_labels)
```

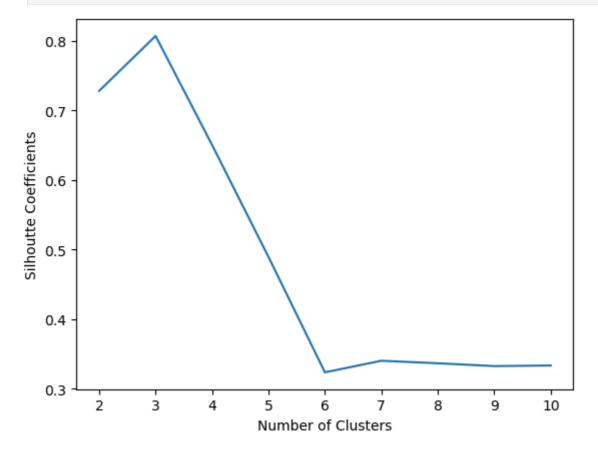
Out[52]: <matplotlib.collections.PathCollection at 0x1ca323ffa90>



```
In [53]: ## kneed Locatore
         from kneed import KneeLocator
         k1 = KneeLocator(range(1,11),wcss,curve='convex', direction='decreasing')
In [54]:
Out[54]: 3
In [55]: # performance metrics
         # silhoutte score
         from sklearn.metrics import silhouette_score
In [56]: silhouette_coefficients = []
         for k in range(2,11):
             kmeans = KMeans(n_clusters=k,init='k-means++')
             kmeans.fit(X_train)
             score = silhouette_score(X_train, kmeans.labels_)
             silhouette_coefficients.append(score)
In [57]: silhouette_coefficients
Out[57]: [0.7281443868598331,
           0.8071181203797672,
           0.6505454471731087,
           0.4895647834796006,
           0.3237480747005592,
           0.34040524166707997,
           0.33677299942813615,
```

0.3327061302389863,
0.3335585872729096]

```
In [60]: # plot the silhouette score
plt.plot(range(2,11), silhouette_coefficients)
plt.xticks(range(2,11))
plt.xlabel("Number of Clusters")
plt.ylabel("Silhoutte Coefficients")
plt.show()
```



In []: