

For IRIS Dataset,

K-Means (when init is 'random'):

The K-means (when init is 'random') cluster analysis are:

[00
1121111111111111111111111111111121111111111111111111111111111111
21222212222221122221212122112222212222122221222122212221]

Cluster Centres for k-means (when init is 'random') are:

[[5.006 3.428 1.462 0.246]]

[5.9016129 2.7483871 4.39354839 1.43387097]

[6.85 3.07368421 5.74210526 2.07105263]]

Distortion for k-means (when init is 'random'): 26.59

K-Means (when init is 'k-means++'):

K-Means Cluster analysis for k-means (when init is 'k-means++') are:

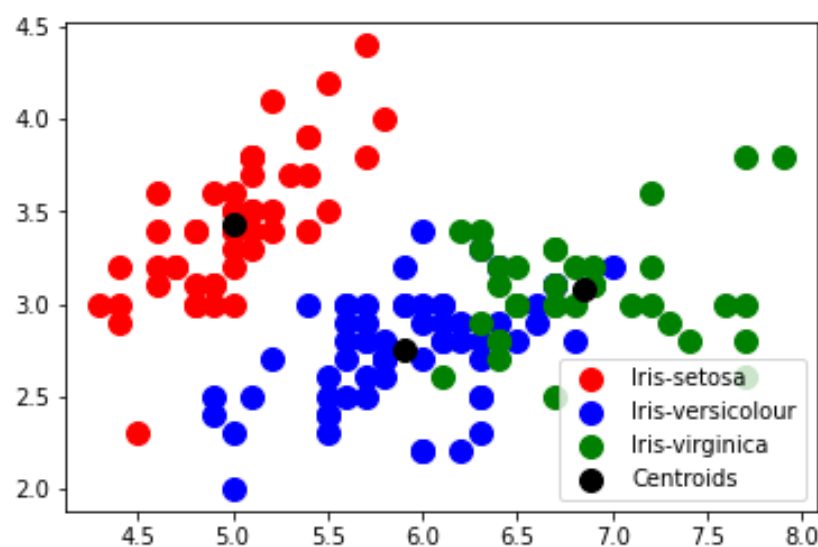
```
[111111111111111111111111111111111111111111111111111111111111111111  
0020000000000000000000000000000020000000000000000000000000000000  
202222022222200222202020220022222022220222202220222022202220]
```

Cluster Centres for k-means (when init is 'k-means++') are:

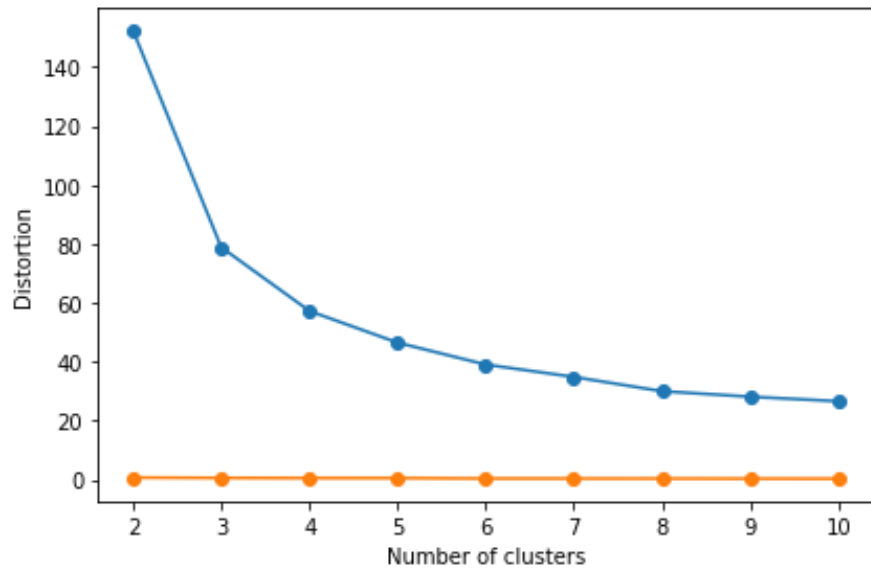
[[5.9016129 2.7483871 4.39354839 1.43387097]]

[5.006 3.428 1.462 0.246]

```
[6.85 3.07368421 5.74210526 2.07105263]]
```

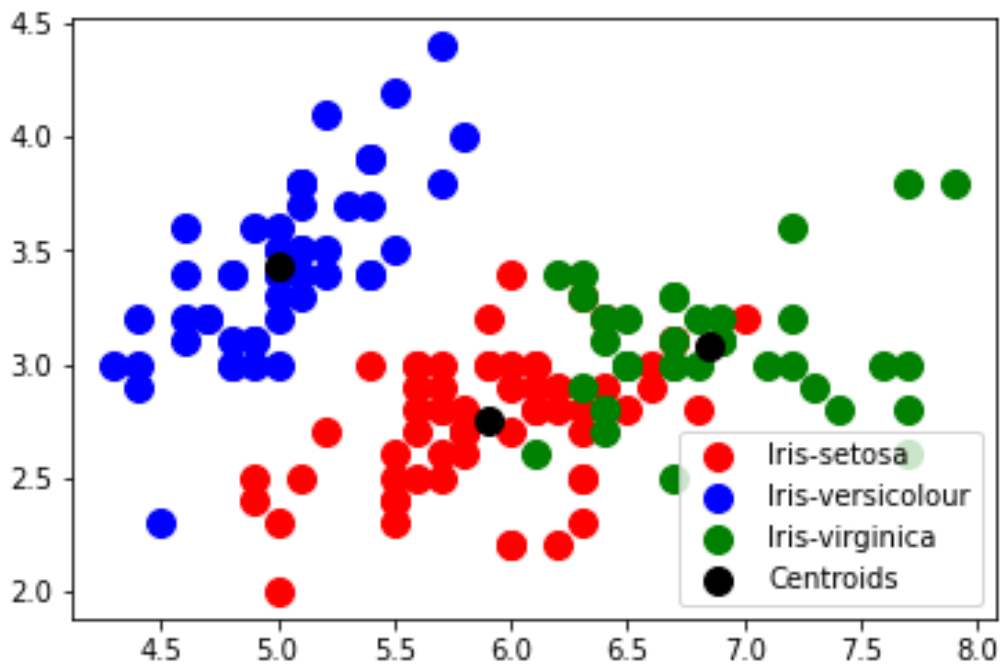
Distortion for k-means (when init is 'k-means++'): 26.09

k-means plot (when init is 'random')

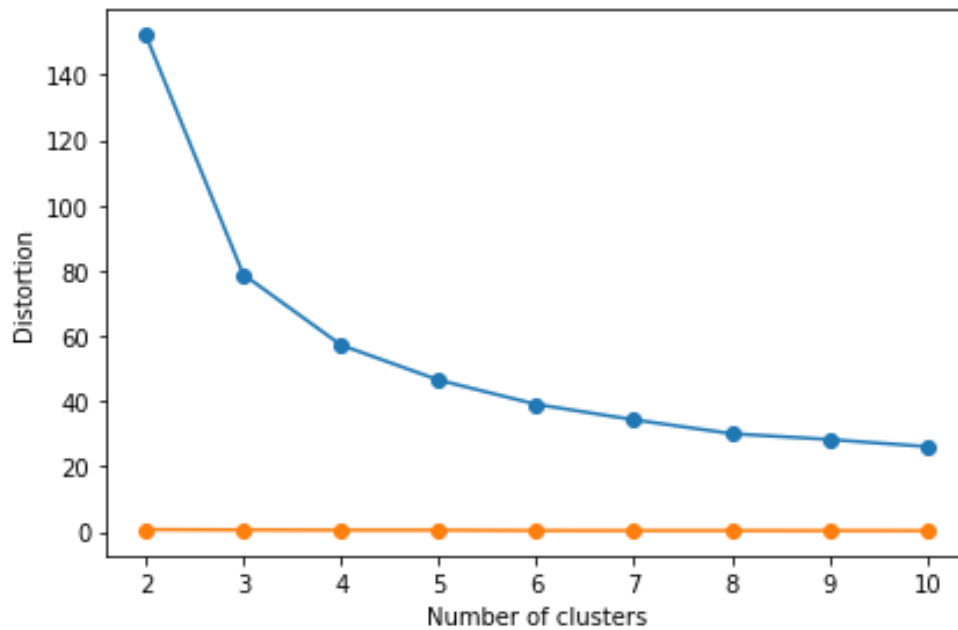


Elbow approach and the silhouette (when init is 'random')

The Blue colour line is the Elbow approach for the k-means (when init is 'random') as we can see from the plot the K value for the elbow method is unsure what to take so I compared the elbow with that of the silhouette score (orange colour line) so that the K value can be fixated on what to take.

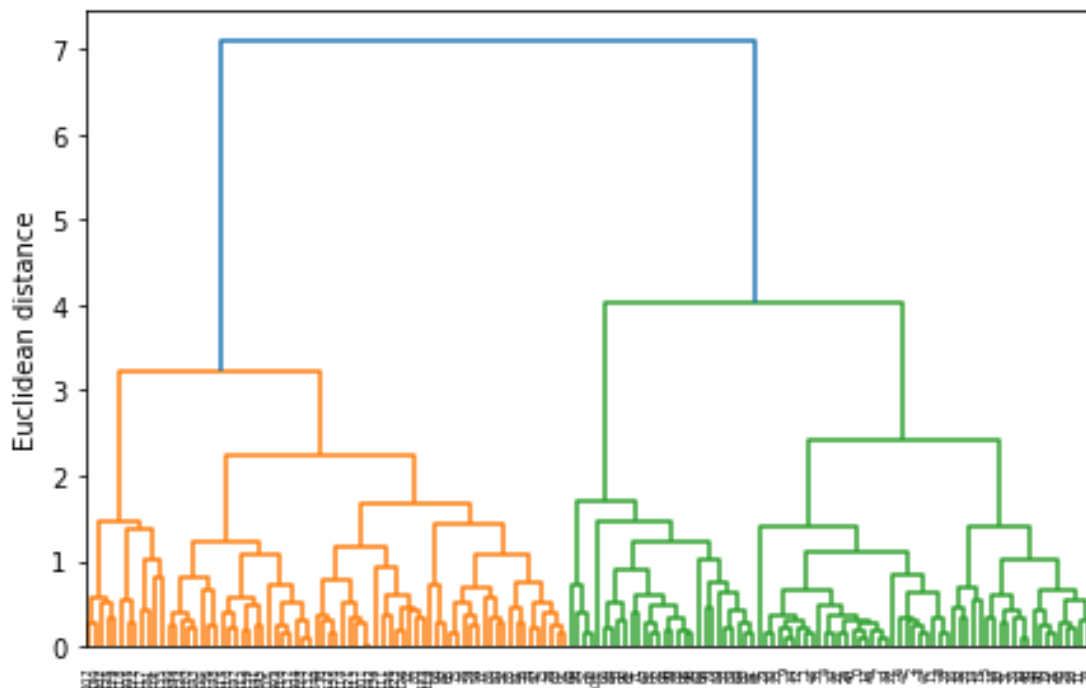


k-means plot (when init is 'k-means++')

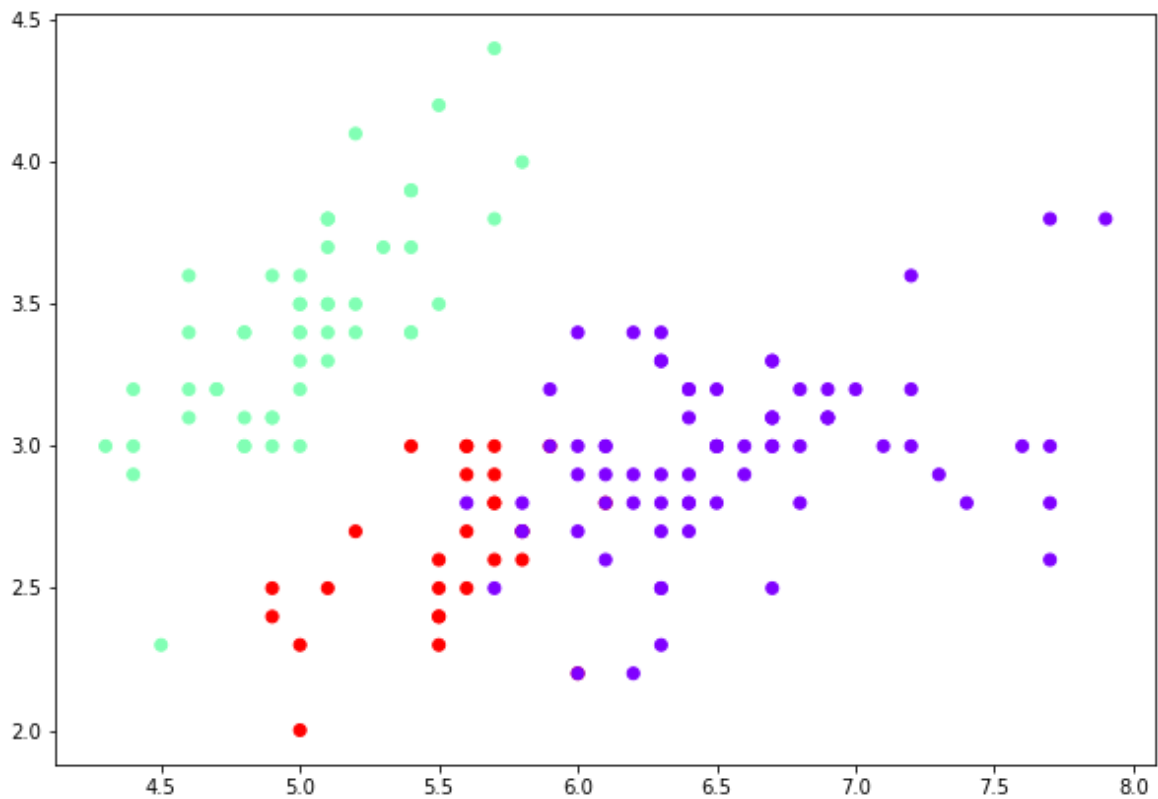


Elbow approach and the silhouette (when init is 'k-means++')

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Hierarchical Clustering (using SciPy library) Dendrograms



Hierarchical clustering (using Scikit library) Dendrograms

Algorithms	Sum of Squared Errors	Silhouette Scores	Time Taken
K-means (when init is 'random')	78.851	0.5528	0.2076
Elbow Method (when init is 'random')	--	0.3044	0.1713
K-means (when init is 'k-means++')	78.851	0.5528	0.0255
Elbow Method (when init is 'k-means++')	--	0.3312	0.3763
Hierarchical approach (using SciPy library)	--	Cluster1 (when max_d is taken) = 0.4998 Cluster2 (when k is taken) = 0.5159	0.2689
Hierarchical approach (using scikit library)	--	0.5135	0.0025

For MNIST Dataset,

I took a subset of test - size '0.90'.

K-Means (when init is 'random'):

The K-means (when init is 'random') cluster analysis are:

[2 2 1 ... 2 0 1]

Cluster Centres for k-means (when init is 'random') are:

[[0. 0. 0. ... 0. 0. 0.]

[0. 0. 0. ... 0. 0. 0.]

[0. 0. 0. ... 0. 0. 0.]]

Distortion for k-means (when init is 'random'): 17839831688.70

K-Means (when init is 'k-means++'):

K-Means Cluster analysis for k-means (when init is 'k-means++') are:

[0 0 1 ... 0 2 1]

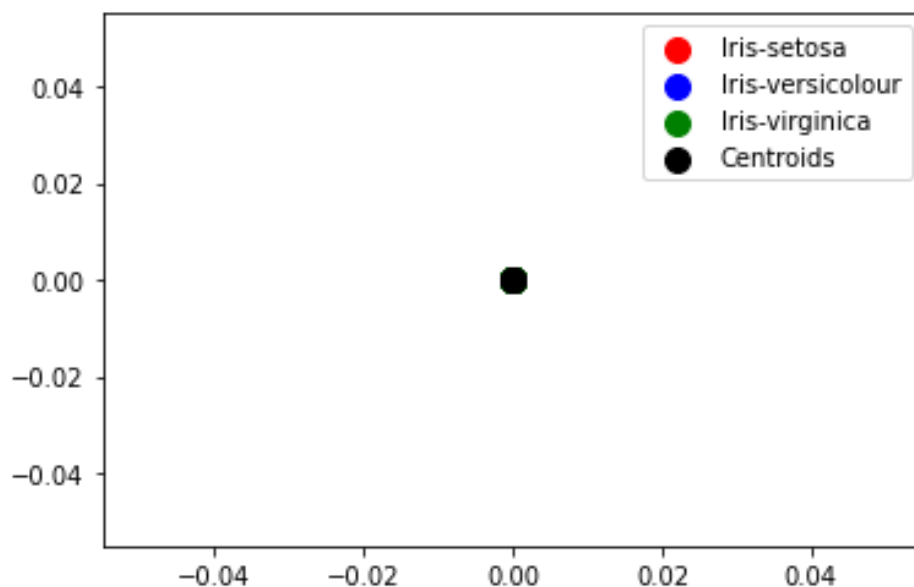
Cluster Centres for k-means (when init is 'k-means++') are:

[[0. 0. 0. ... 0. 0. 0.]

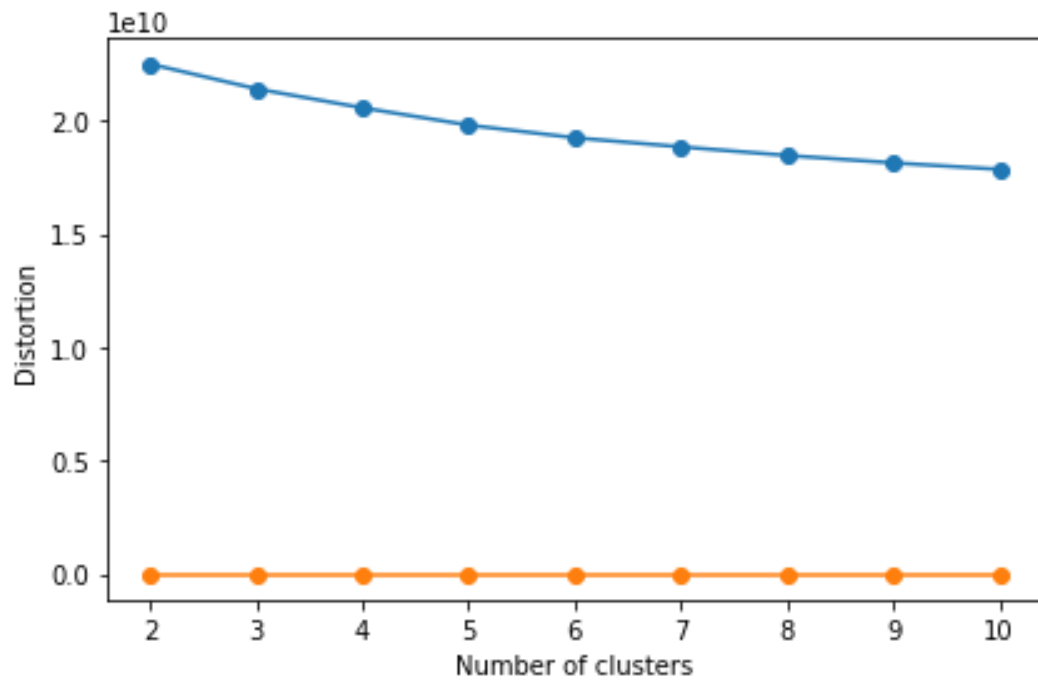
[0. 0. 0. ... 0. 0. 0.]

[0. 0. 0. ... 0. 0. 0.]]

Distortion for k-means (when init is 'k-means++'): 17838960304.95

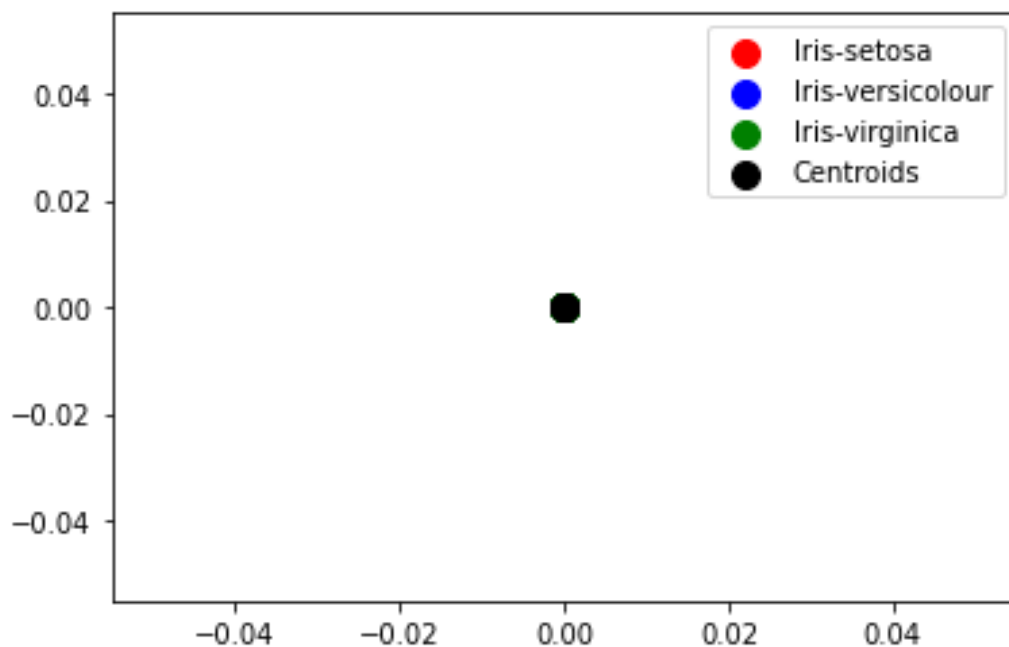


k-means plot (when init is 'random')

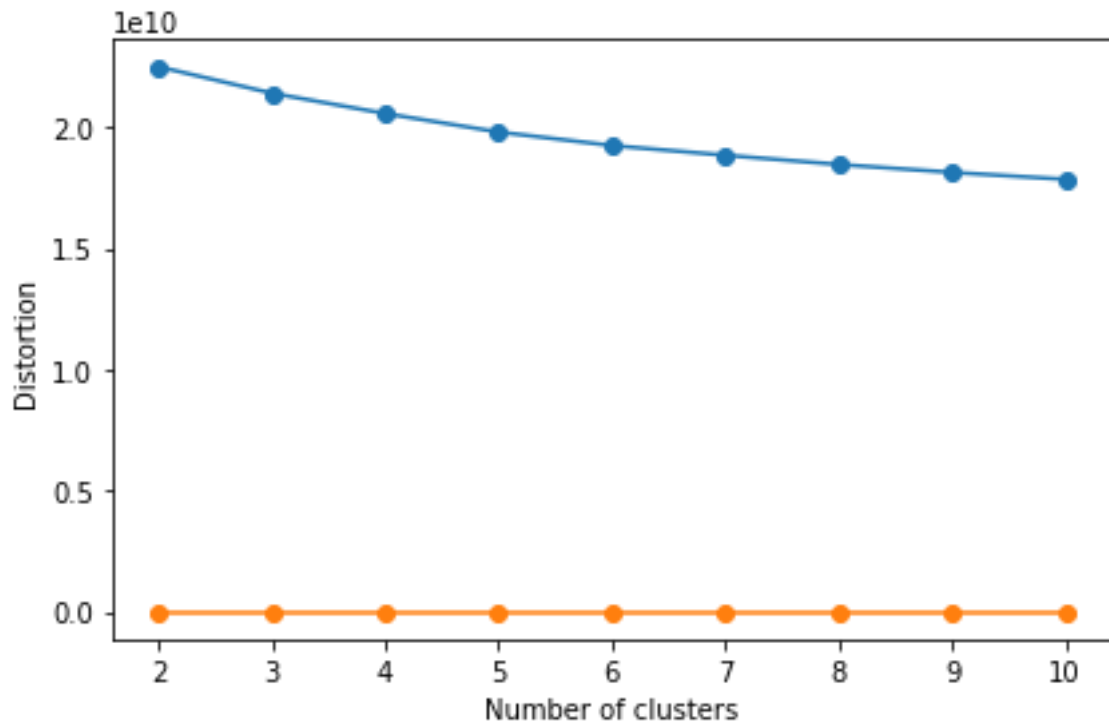


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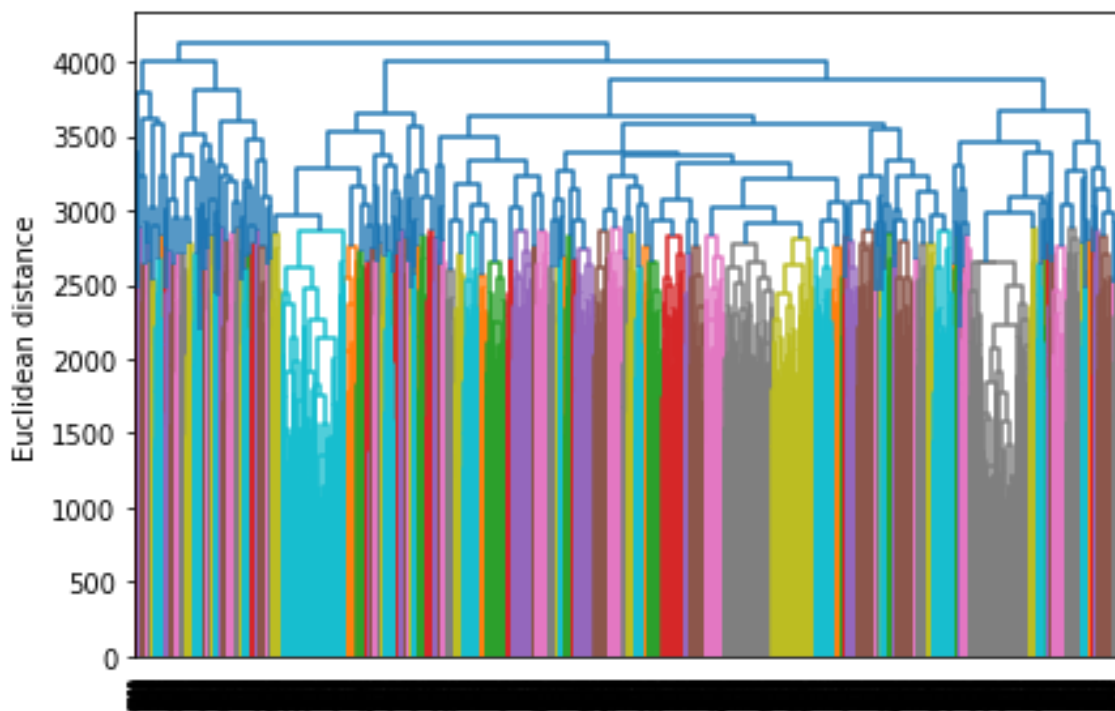


k-means plot (when init is 'k-means++')

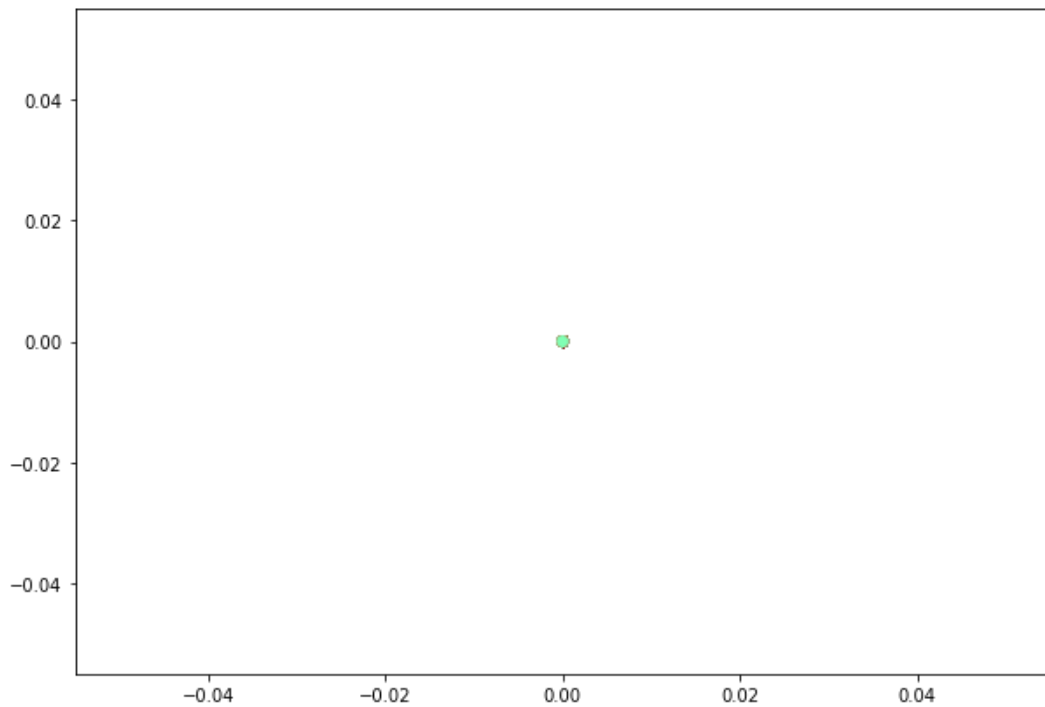


Elbow approach and the silhouette (when init is 'k-means++')

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Hierarchical Clustering (using SciPy library) Dendrograms



Hierarchical clustering (using Scikit library) Dendrograms

Algorithms	Sum of Squared Errors	Silhouette Scores	Time Taken
K-means (when init is 'random')	21392925381.585	0.0538	2.9420
Elbow Method (when init is 'random')	--	0.0572	33.8220
K-means (when init is 'k-means++')	21392906205.685	0.0539	3.9344
Elbow Method (when init is 'k-means++')	--	0.0577	40.6300
Hierarchical approach (using SciPy library)	--	--	102.7345
Hierarchical approach (using scikit library)	--	0.0399	22.7918