

# Hierarchical Clustering Implementation

```
In [2]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn import datasets
```

```
In [3]: # Import IRIS Dataset
iris = datasets.load_iris()
```

```
In [4]: iris_data = pd.DataFrame(iris.data)
```

```
In [7]: iris_data.columns = iris.feature_names
```

```
In [8]: iris_data
```

```
Out[8]:
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
...	...	...	...	...
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns

```
In [9]: # Standard Scaler
from sklearn.preprocessing import StandardScaler
scaler = StandardScaler()
```

```
In [10]: X_scaled = scaler.fit_transform(iris_data)
```

```
In [11]: X_scaled
```

```
Out[11]: array([[ -9.00681170e-01,  1.01900435e+00, -1.34022653e+00,
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```

```
In [12]: X_scaled.shape
```

```
Out[12]: (150, 4)
```

```
In [14]: # apply the pca
from sklearn.decomposition import PCA
```

```
In [15]: pca = PCA(n_components=2)
```

```
In [16]: pca
```

```
Out[16]: PCA
Parameters
```

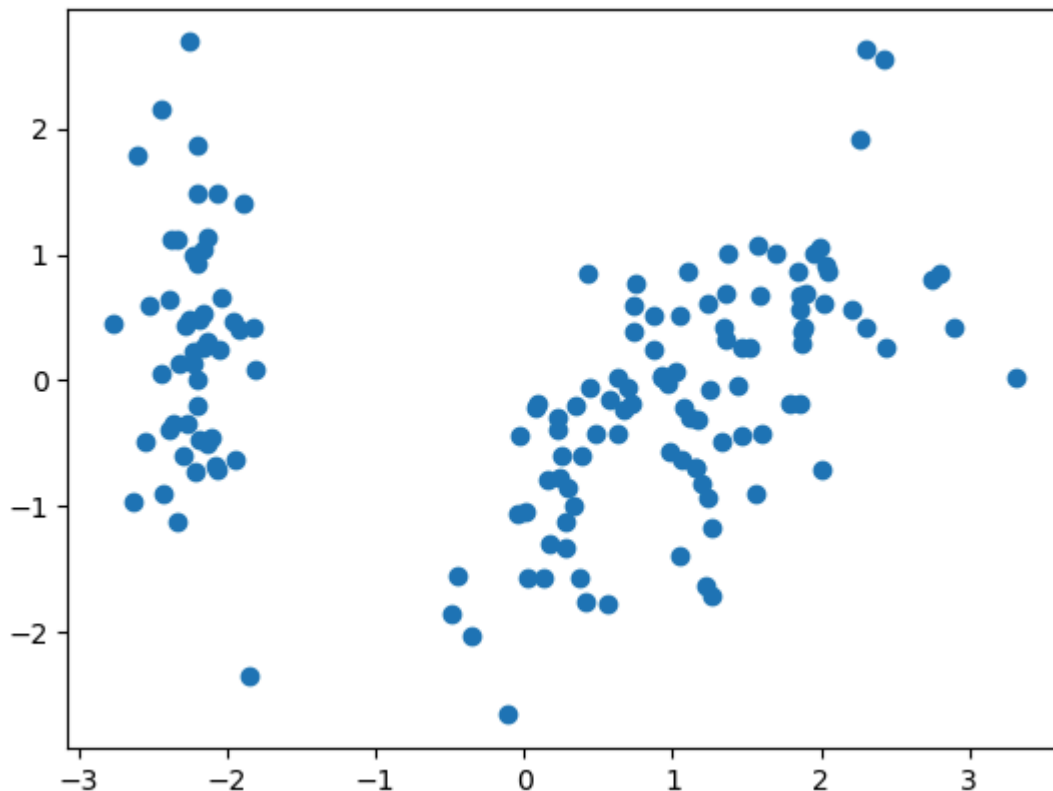
```
In [17]: pca_scaled = pca.fit_transform(X_scaled)
```

```
In [19]: pca_scaled.shape
```

```
Out[19]: (150, 2)
```

```
In [20]: plt.scatter(pca_scaled[:,0],pca_scaled[:,1])
```

```
Out[20]: <matplotlib.collections.PathCollection at 0x237e818fc90>
```

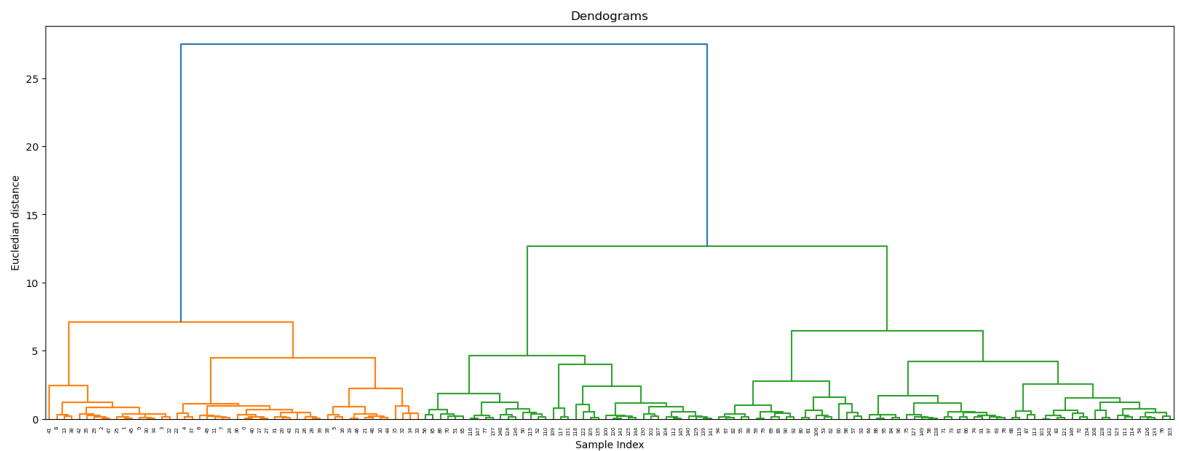


```
In [23]: ## Agglomerative Clustering
import scipy.cluster.hierarchy as sc

plt.figure(figsize=(20,7))
plt.title("Dendograms")
```

```
# craete dendograms
sc.dendrogram(sc.linkage(pca_scaled,method='ward'))
plt.title('Dendograms')
plt.xlabel('Sample Index')
plt.ylabel('Eucledian distance')
```

```
Out[23]: Text(0, 0.5, 'Euclidean distance')
```



```
In [26]: from sklearn.cluster import AgglomerativeClustering
cluster = AgglomerativeClustering(n_clusters=2, metric='euclidean', linkage='ward')
cluster.fit(pca_scaled)
```

```
Out[26]: ▼ AgglomerativeClustering
```

### ► Parameters

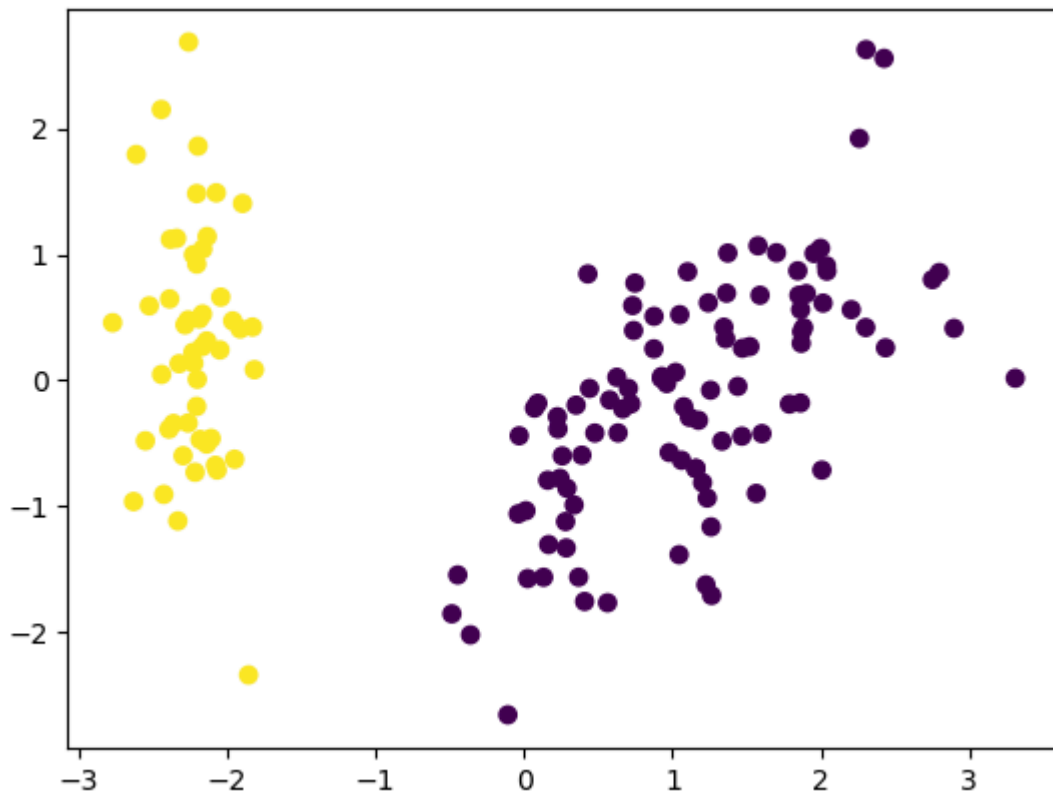
```
In [27]: cluster.labels_
```

[illegible]

```
In [28]: plt.scatter(pca_scaled[:,0],pca_scaled[:,1],c=cluster.labels_)
```

```
Out[28]: <matplotlib.collections.PathCollection at 0x237ea7616d0>
```





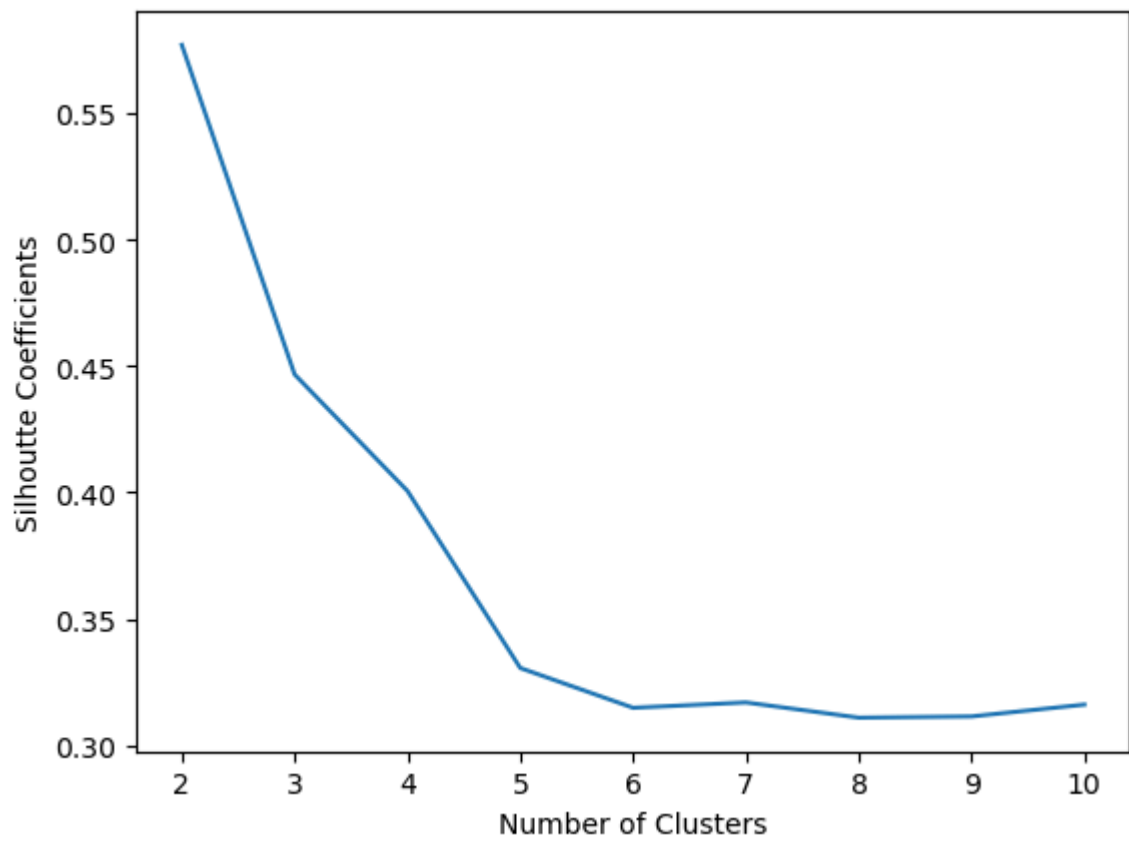
```
In [29]: # performance metrics
# silhoutte score
from sklearn.metrics import silhouette_score
```

```
In [32]: silhouette_coefficients = []
for k in range(2,11):
    agglo = AgglomerativeClustering(n_clusters=k,metric='euclidean',linkage='war
    agglo.fit(X_scaled)
    score = silhouette_score(X_scaled, agglo.labels_)
    silhouette_coefficients.append(score)
```

```
In [33]: silhouette_coefficients
```

```
Out[33]: [0.5770346019475988,
0.4466890410285909,
0.4006363159855973,
0.33058726295230545,
0.31485480100512825,
0.316969830299128,
0.310946529007258,
0.31143422475471655,
0.3161120375980681]
```

```
In [34]: # 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 [ ]: