

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
```

```
from sklearn.datasets import load_wine
```

```
data=load_wine()
df=pd.DataFrame(data=data.data, columns=data.feature_names)
```

df

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	flav
<b>0</b>	14.23	1.71	2.43	15.6	127.0	2.80	
<b>1</b>	13.20	1.78	2.14	11.2	100.0	2.65	
<b>2</b>	13.16	2.36	2.67	18.6	101.0	2.80	
<b>3</b>	14.37	1.95	2.50	16.8	113.0	3.85	
<b>4</b>	13.24	2.59	2.87	21.0	118.0	2.80	
...	...	...	...	...	...	...	...
<b>173</b>	13.71	5.65	2.45	20.5	95.0	1.68	
<b>174</b>	13.40	3.91	2.48	23.0	102.0	1.80	
<b>175</b>	13.27	4.28	2.26	20.0	120.0	1.59	
<b>176</b>	13.17	2.59	2.37	20.0	120.0	1.65	
<b>177</b>	14.13	4.10	2.74	24.5	96.0	2.05	

178 rows × 13 columns

Next steps:

[Generate code with df](#)

[New interactive sheet](#)

```
from sklearn.preprocessing import StandardScaler
scaler = StandardScaler()
```

```
df_scaled= scaler.fit_transform(df)
```

df\_scaled

```
array([[ 1.51861254, -0.5622498 ,  0.23205254, ...,  0.36217728,
         1.84791957,  1.01300893],
       [ 0.24628963, -0.49941331,  0.82799632, ...,  0.40605066,
         1.1134493 ,  0.96524152],
```

```
[ 0.19687903,  0.02123125,  1.10933436, ...,  0.31830389,
  0.78858745,  1.39514818],
...,
[ 0.33275817,  1.74474449, -0.38935541, ..., -1.61212515,
 -1.48544548,  0.28057537],
[ 0.20923168,  0.22769377,  0.01273209, ..., -1.56825176,
 -1.40069891,  0.29649784],
[ 1.39508604,  1.58316512,  1.36520822, ..., -1.52437837,
 -1.42894777, -0.59516041]])
```

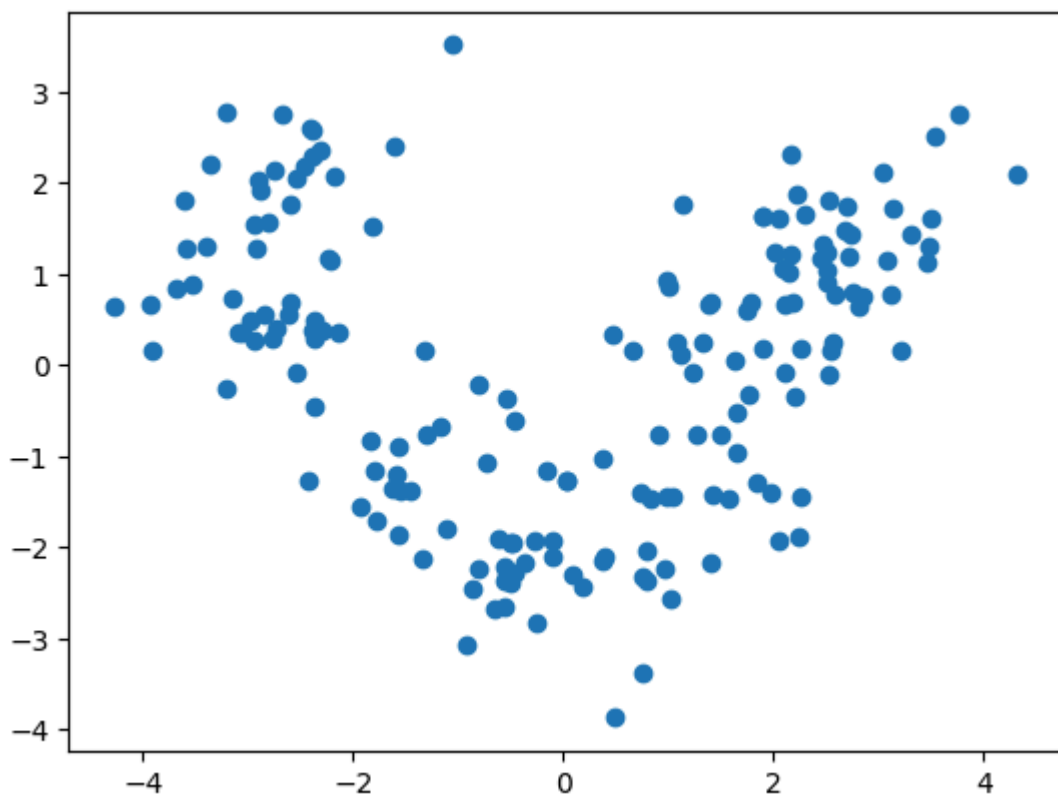
```
from sklearn.decomposition import PCA
pca=PCA(n_components=2)
pca_scaled = pca.fit_transform(df_scaled)
```

```
pca_scaled.shape
```

```
(178, 2)
```

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

```
<matplotlib.collections.PathCollection at 0x79418172b830>
```



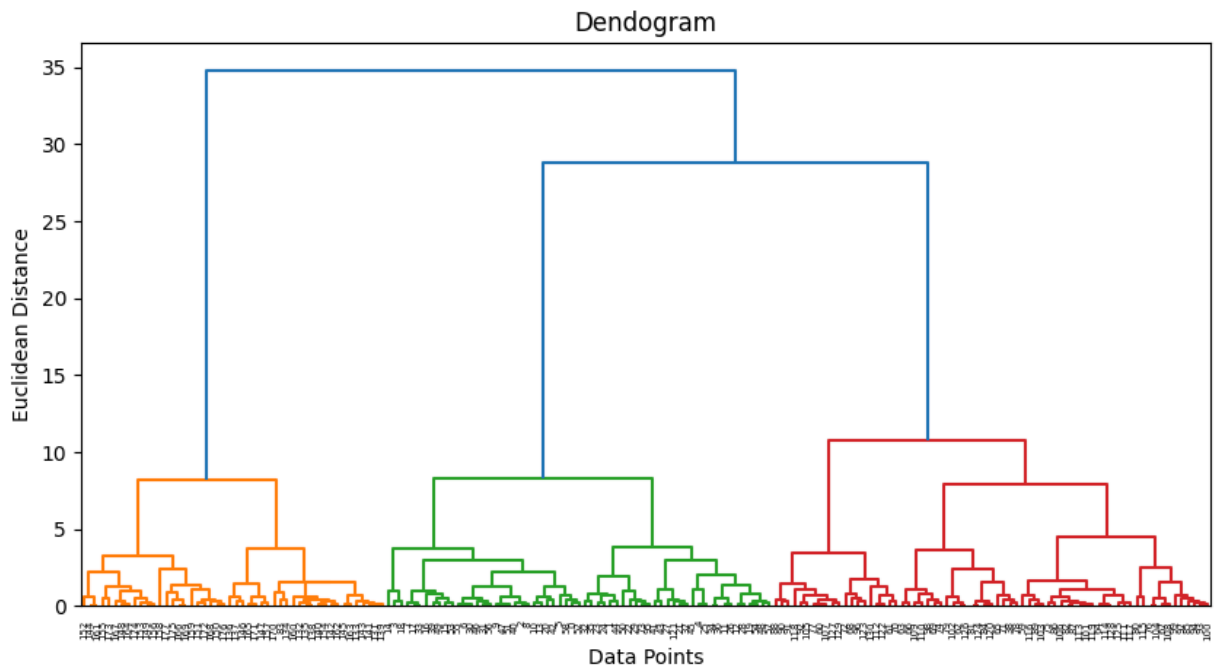
```
import scipy.cluster.hierarchy as sc
```

```
plt.figure(figsize=(10,5))
plt.title('Dendrogram')
```

```
# create a Dendrogram
sc.dendrogram(sc.linkage(pca_scaled, method='ward'))
```

```
plt.title('Dendrogram')
plt.xlabel('Data Points')
plt.ylabel('Euclidean Distance')
```

```
Text(0, 0.5, 'Euclidean Distance')
```



Hence the `n_cluster` value is 3

```
# silhouette score
from sklearn.cluster import KMeans
from sklearn.metrics import silhouette_score

for k in range(2,11):
    kmeans = KMeans(n_clusters=k, init='k-means++')
    labels = kmeans.fit_predict(pca_scaled)
    score = silhouette_score(pca_scaled, labels)
    print(f"for k={k} silhouette score is {score}")
```

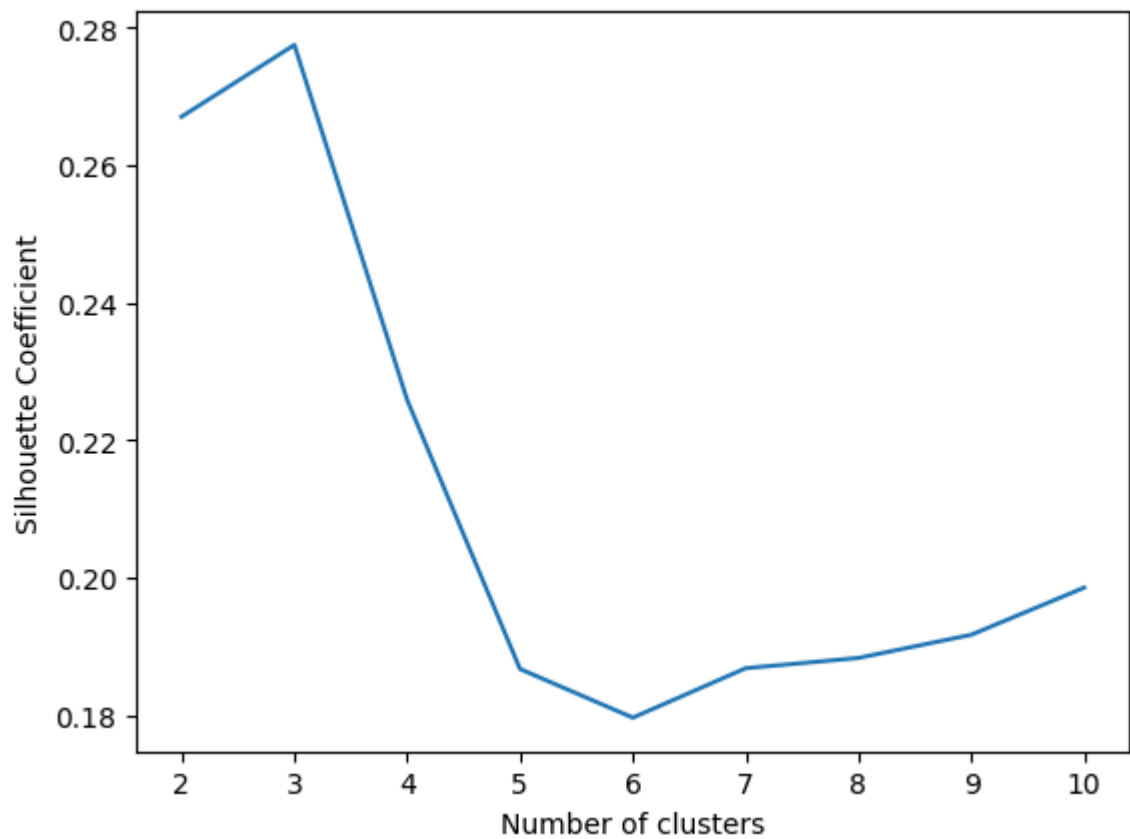
```
for k=2 silhouette score is 0.4750102141135033
for k=3 silhouette score is 0.5609305721582905
for k=4 silhouette score is 0.4855976780800213
for k=5 silhouette score is 0.4336819174782484
for k=6 silhouette score is 0.4244059902675462
for k=7 silhouette score is 0.42251924243479017
for k=8 silhouette score is 0.3777792659691172
for k=9 silhouette score is 0.380704142674008
for k=10 silhouette score is 0.40269911475418657
```

```
# Agglomerative Clustering
from sklearn.cluster import AgglomerativeClustering
```

```
cluster=AgglomerativeClustering(n_clusters=3, metric='euclidean', linkage='ward')
```



```
plt.xlabel('Number of clusters')  
plt.ylabel('Silhouette Coefficient')  
plt.show()
```



Start coding or [generate](#) with AI.