K-Means Clustering with Scikit-learn

This example implements K-means clustering with Scikit-learn. Since this is an unsupervised learning use case, we use just the features of the Iris dataset to cluster the observations into labels.

```
# import packages
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn import datasets
from sklearn.model_selection import train_test_split
# load dataset
data = datasets.load iris()
# get the dataset features
X = data.data
# create the model. Since we know that the Iris dataset has 3 classes, we
set n clusters = 3
kmeans = KMeans(n clusters=3, random state=0)
# fit the model on the training set
kmeans.fit(X)
'Output':
KMeans(algorithm='auto', copy_x=True, init='k-means++', max iter=300,
    n clusters=3, n init=10, n jobs=1, precompute_distances='auto',
    random state=0, tol=0.0001, verbose=0)
# predict the closest cluster each sample in X belongs to.
y kmeans = kmeans.predict(X)
# plot clustered labels
plt.scatter(X[:, 0], X[:, 1], c=y kmeans, cmap='viridis')
# plot cluster centers
centers = kmeans.cluster centers
plt.scatter(centers[:, 0], centers[:, 1], c='black', s=200, alpha=0.7);
plt.show()
```

The code to plot the clustered labels and the cluster centers should be executed in the same notebook. The plot of clusters made by the K-means algorithm is shown in Figure 25-4.

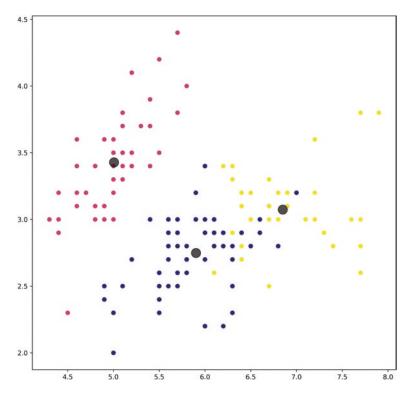


Figure 25-4. Plot of K-means clusters and their cluster centers

Hierarchical Clustering

Hierarchical clustering is another clustering algorithm for finding homogeneous sub-groups or classes within a dataset. However, as opposed to k-means, we do not need to make an a priori assumption of the number of clusters in the dataset before running the algorithm.

The two main techniques for performing hierarchical clustering are

- · Bottom-up or agglomerative
- Top-down or divisive