Ex. No. 06

06.07.2024

AGGLOMERATIVE CLUSTERING ALGORITHM

Aim

Develop agglomerative clustering models to cluster the given dataset using the scikit-learn.

Description

Hierarchical clustering is a type of clustering algorithm used in unsupervised machine learning to group similar data points together in a hierarchical tree-like structure, also known as a dendrogram. The algorithm starts by treating each data point as a separate cluster, and then iteratively merges the closest pair of clusters until a stopping criterion is met.

Hierarchical clustering can be performed using two approaches:

Agglomerative hierarchical clustering: In this approach, each data point is initially considered as a separate cluster and then, at each step, the two closest clusters are merged into a larger cluster. This process is continued until all data points belong to a single cluster. This is a bottom-up approach.

Divisive hierarchical clustering: In this approach, all data points are initially considered as belonging to a single cluster, and then at each step, the cluster is recursively split into smaller clusters based on some distance metric, until each cluster contains only one data point. This is a top-down approach.

Dendrogram: A Useful Tool for Summarizing Similarity Measurements

- Min Distance: Minimum distance of two points.
- Max Distance: Maximum distance of two points.
- Group Average: Average of distance between every two points of the cluster.
- Ward's method: Similarity is based on the increase in square when two clusters are merged.

Source Code

1. Pre-process the data and fill the missing values and apply normalization

```
import pandas as pd
import numpy as np
from sklearn.preprocessing import StandardScale
data=pd.read_csv(hopping-data.csv')
data.fillna(data.mean(), inplace=True)
scaler = StandardScaler()

data_normalized = scaler.fit_transform(data.select_dtypes(include=[np.number]))
data_normalized_df = pd.DataFrame(data_normalized,
columns=data.select_dtypes(include=[np.number]).columns)
non_numeric_cols = data.select_dtypes(exclude=[np.number]).columns
data_normalized_df = pd.concat([data_normalized_df,
data[non_numeric_cols].reset_index(drop=True)], axis=1)
print(data_normalized_df.head())
```

2. Apply label encoding to convert the categorical values to numerical values

```
from sklearn.preprocessing import LabelEncoder
encoders = {}
for column in data.select_dtypes(include=['object']).columns:
    encoders[column] = LabelEncoder()
    data[column] = encoders[column].fit transform(data[column])
```

OUTPUT:

3 4 23 16 77 4 5 31 17 40	0 1 2 3 4	1	19 21 20 23	15 15 16	Spending Score (1-100) 39 81 6 77 40	
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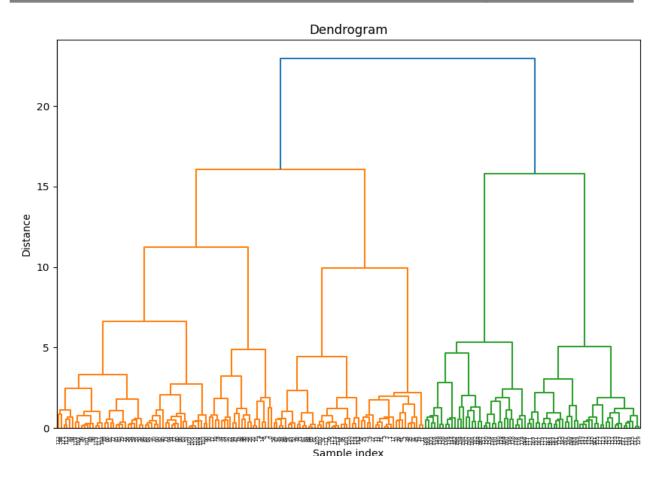
3. Plot the dendrogram

import matplotlib.pyplot as plt

```
from scipy.cluster.hierarchy import dendrogram, linkage
def plot_dendrogram(data):
    linked = linkage(data, method='ward') # You can change the method here
    plt.figure(figsize=(10, 7))
    dendrogram(linked, orientation='top', distance_sort='descending', show_leaf_counts=True)
    plt.title('Dendrogram')
    plt.xlabel('Sample index')
    plt.ylabel('Distance')
    plt.show()

plot_dendrogram(data_normalized)
```

OUTPUT:



4. Implement agglomerative clustering algorithms to cluster the given data for different distance metrics (Euclidean, Manhattan, Cosine, L1, L2) and linkage functions (single, complete, average, wards).

from sklearn.cluster import AgglomerativeClustering distance_metrics = ['euclidean', 'manhattan', 'cosine'] linkage_methods = ['ward', 'single', 'complete', 'average'] results = {} for metric in distance metrics:

for linkage_method in linkage_methods:

Error occurred for metric: manhattan, linkage: ward manhattan was provided as metric. Ward can only work with euclidean distances.

5. Try with the whole dataset (except the label) and with any 2 attributes

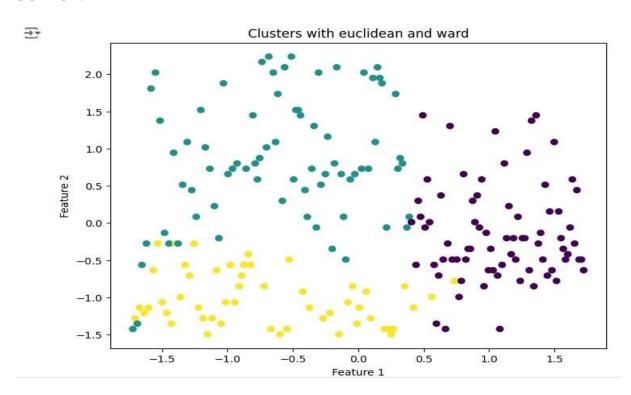
```
def plot_scatter(data, cluster_labels, title):
    plt.figure(figsize=(8, 6))
    plt.scatter(data[:, 0], data[:, 1], c=cluster_labels, cmap='viridis', marker='o')
    plt.title(title)
    plt.xlabel('Feature 1')

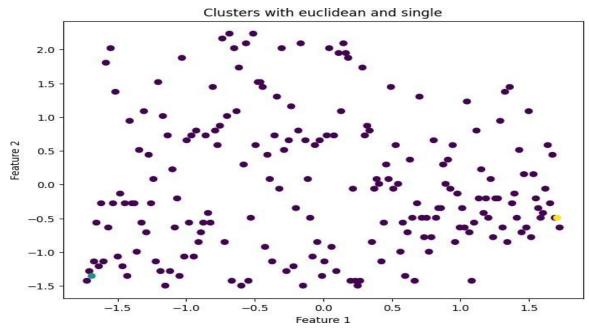
plt.ylabel('Feature 2')
    plt.show()

for (metric, linkage method), cluster labels in results.items():
```

plot_scatter(data_normalized[:, :2], cluster_labels, f'Clusters with {metric} and {linkage_method}')

OUTPUT:





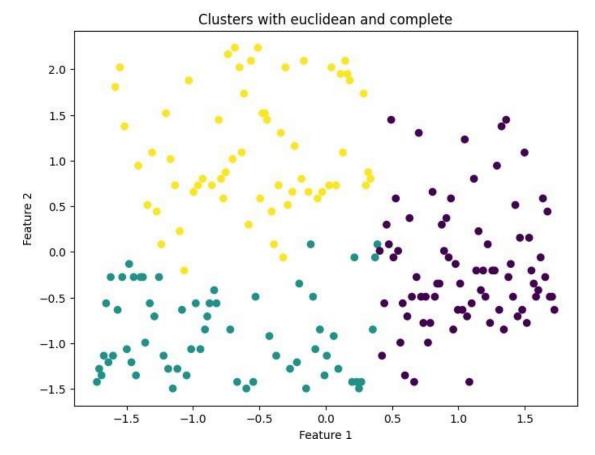
6. Analyse the results using scatter plot

```
def plot_scatter(data, cluster_labels, title):
    plt.figure(figsize=(8, 6))
    plt.scatter(data[:, 0], data[:, 1], c=cluster_labels, cmap='viridis', marker='o')
    plt.title(title)
    plt.xlabel('Feature 1')

plt.ylabel('Feature 2')
    plt.show()

OUTPUT:
```

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7. Compare the results using Mutual information, Silhouette Score (Silhouette Coefficient), Davies-Bouldin Index.

```
From sklearn.metrics import silhouette_score, davies_bouldin_score, mutual_info_score true_labels = data['true_label_column_name']

for (metric, linkage_method), cluster_labels in results.items():
    silhouette = silhouette_score(data_normalized, cluster_labels)
    davies_bouldin = davies_bouldin_score(data_normalized, cluster_labels)
    mutual_info = mutual_info_score(true_labels, cluster_labels)
    print(f''Metric: {metric}, Linkage: {linkage_method}'')
    print(f''Silhouette Score: {silhouette:.4f}'')
    print(f''Davies-Bouldin Index: {davies_bouldin:.4f}'')

Print(f''Mutual Information: {mutual_info:.4f}'') # Uncomment if true labels are available print("-" * 50)

OUTPUT:
```

```
Metric: euclidean, Linkage: ward
Silhouette Score: 0.3087
Davies-Bouldin Index: 1.1303

Metric: euclidean, Linkage: single
Silhouette Score: 0.0992
Davies-Bouldin Index: 0.5491

Metric: euclidean, Linkage: complete
Silhouette Score: 0.3260
Davies-Bouldin Index: 1.0617

Metric: euclidean, Linkage: average
Silhouette Score: 0.3335
Davies-Bouldin Index: 0.9122
```

Result

Agglomerative Clustering Algorithm are executed and verified successfully.