

Principal Component Analysis and Hierarchical Clustering

Data Boot Camp

Lesson 18.2



The Big Picture



This Week: Unsupervised Machine Learning

By the end of this week, you'll know how to:



To reduce features or dimensions using principal component analysis (PCA)



Perform PCA on datasets



Use hierarchical clustering as alternative to K-means



This Week's Challenge

Using the skills learned throughout the week, you will analyze cryptocurrency data, reduce data dimensions with PCA, cluster with K-means, and visualize the results.

Today's Agenda

By completing today's activities, you'll learn the following skills:



Reducing features or dimensions using PCA



Performing PCA on datasets



Using hierarchical clustering as an alternative to K-means



Make sure you've downloaded any relevant class files!



Principal component analysis is a statistical technique to speed up machine learning algorithms.



It does this by reducing the number of input features (or dimensions).



This allows us to transform large sets of variables into smaller ones that contain most of the information.

Before using PCA, we need to standardize the data using scikit-learn's StandardScaler module.

Then, the fit_transform() method combines training and transforming data into a single step.

```
# Standardize data with StandardScaler
Iris_scaled = StandardScaler().fit_transform(df_iris)
print(iris_scaled[0.5])≈
ΓΓ-0.90068117
               1.03205722 -1.34127244
                                           -1.321976737
 Γ-1.14301791
               -0.1249576 -1.34127244
                                           -1.321976737
 Γ-1.38535265
                0.33784833 -1.39813811
                                           -1.321976737
 \lceil -1.50652052 \qquad 0.10644536 \quad -1.2844067
                                           -1.321976737
 \Gamma-1.021849041
                 1.26346019 -1.3412724
                                           -1.3219767377
```

Once the features are standardized, PCA can be used to reduce the number of features.

The n_components parameter specifies the final number of features.

PROA reduces the ataset grates igner to be to find in the majorest called the different postents. The two main dimensions of variations that contain most of the information in the original dataset.

```
# Transform PCA data to a DataFrame
df_iris_pca = pd.DataFrame(
    data=iris_pca, columns=["principal component 1", "principal component 2"]
df_iris_pca.head()
       principal component 1
                                        principal component 2
                    -2.264542
                                                       0.50574
                    -2.086426
                                                     -0.655405
                    -2.367950
                                                     -0.318477
3
                                                     -0.575368
                    -2.304197
                                                      0.674767
                    -2.388777
```

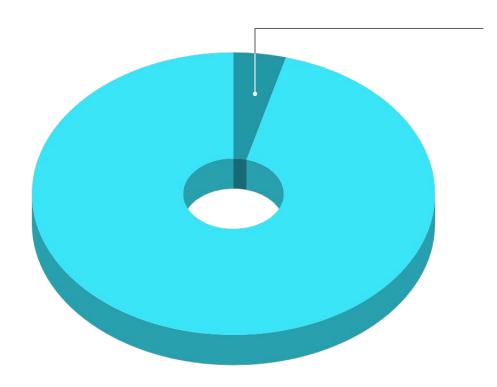
The principal component values bear little resemblance to the original dataset. They can be seen as a reduced representation of the original data. The explained_variance_ratio attribute is used to assess the amount of information that has been preserved in the PCA dimensionality reduction.

Here, the first principal component is responsible for 72.77% of the variance, and the second contains 23.03% of the variance.

```
# Fetch the explained variance
pca.explained_variance_ratio_
```

arry([0.33690046, 0.26230645, 0.232606390])

Together, the principal components preserve 95.80% of the information.



In other words...

A little over 4% of the useful information was lost in the dimensionality reduction performed by PCA.



Instructor Demonstration

Speed up Machine Learning with PCA



Activity: PCA in Action

In this activity, you will use PCA to reduce the dimensions of a consumer shopping dataset.

Suggested Time:

20 minutes







Activity: PCA

In this activity, you will perform PCA on the Boston Marathon dataset.

Suggested Time:

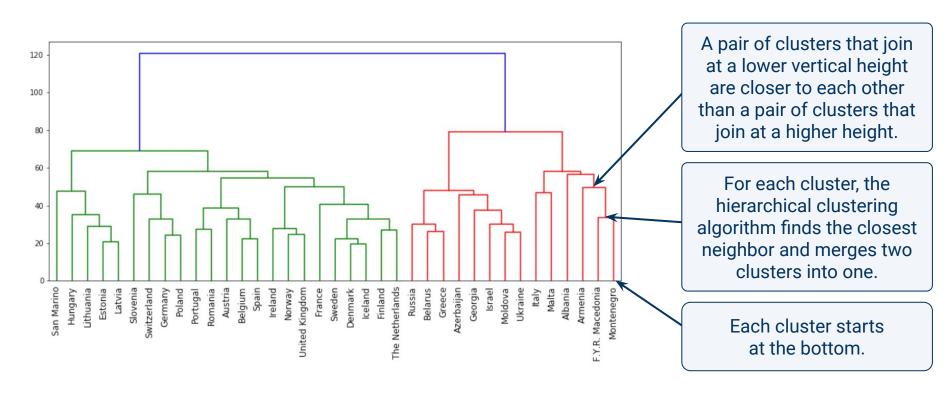
25 minutes





Dendrograms

This tree-like structure is called a dendrogram. Each cluster starts at the bottom.



Dendrograms

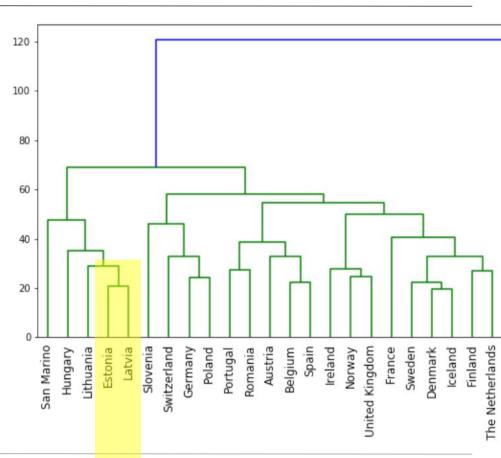
Each country is paired with its closest neighboring cluster based on voting patterns.

For example

Estonia's closest neighbor is Latvia, and they merge into a single cluster before merging with Lithuania.

There appear to be two large clusters:

- **1.** The cluster on the left, in which eastern European countries are more heavily represented
- **2.** The cluster on the right, in which eastern European countries are more heavily represented



Dendrograms

Because the mergings are based on the distances between samples, datasets should be standardized or normalized.

Here, we use scikit-learn's normalize method. The actual clustering is performed here using SciPy's linkage method.

```
normalized = normalize(df)
mergings = linkage(normalized, method='ward')
```

We're using the ward method to compute distances between clusters.

Hierarchical Clustering

To generate the dendrogram, SciPy's dendrogram method is used.

The first argument, mergings, is the linkage matrix that we just generated.

The next two arguments, leaf_rotation and leaf_font_size, refer to the text label of each sample. Here, the text is turned vertically, and its font size is set at 5.

Hierarchical Clustering

Hierarchical clustering has different types of linkage methods:

Single

The difference between two clusters is defined by the closest distance between two clusters.

Complete

The difference between two clusters is defined by the farthest distance between two clusters.

Ward

This method is based on the squared euclidean distance between clusters. It's the method used in our example, and it is often used as a default.





Activity: Hierarchical Customer Data

In this activity, you will use hierarchical clustering to group and plot customer data.

Suggested Time:

20 minutes



