

How can I perform multidimensional scaling in Python?

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Multidimensional scaling is a statistical technique used to analyze and visualize high-dimensional data by reducing its dimensionality to two or three dimensions. In Python, this technique can be performed using the scikit-learn library, specifically the "MDS" function in the "manifold" module. This function takes in a distance or dissimilarity matrix as input and outputs the coordinates of the data points in a lower-dimensional space. The resulting coordinates can then be plotted to visualize the similarities and differences between the data points. Overall, performing multidimensional scaling in Python allows for a better understanding and interpretation of complex data sets.

Perform Multidimensional Scaling in Python

In statistics, multidimensional scaling is a way to visualize the similarity of observations in a dataset in an abstract cartesian space (usually a 2-D space).

The easiest way to perform multidimensional scaling in Python is by using the MDS() function from the sklearn.manifold sub-module.

The following example shows how to use this function in practice.

Example: Multidimensional Scaling in Python

Suppose we have the following pandas DataFrame that contains information about various basketball players:

```
import pandas as pd
```

```
#create DataFrame
```

```
df = pd.DataFrame({'player': ,  
'points': ,  
'assists': ,  
'blocks': ,  
'rebounds': })
```

```
#set player column as index column
```

```
df = df.set_index('player')
```

```
#view Dataframe
```

```
print(df)
```

```
points assists blocks rebounds
```

```
player
```

```
A 4 3 7 4
```

```
B 4 2 3 5
```

```
C 6 2 6 5
```

```
D 7 5 7 6
```

```
E 8 4 5 5
```

```
F 14 8 8 8
```

```
G 16 7 8 10
```

```
H 19 6 4 4
```

```
I 25 8 2 3
```

```
J 25 10 2 2
```

K 28 11 1 2

We can use the following code to perform multidimensional scaling with the `MDS()` function from the `sklearn.manifold` module:

```
from sklearn.manifold import MDS

#perform multi-dimensional scaling
mds = MDS(random_state=0)
scaled_df = mds.fit_transform(df)

#view results of multi-dimensional scaling
print(scaled_df)

]
```

Each row from the original DataFrame has been reduced to an (x, y) coordinate.

We can use the following code to visualize these coordinates in a 2-D space:

```
import matplotlib.pyplot as plt

#create scatterplot
```

```
plt.scatter(scaled_df, scaled_df)
```

```
#add axis labels
```

```
plt.xlabel('Coordinate 1')
```

```
plt.ylabel('Coordinate 2')
```

```
#add labels to each point
```

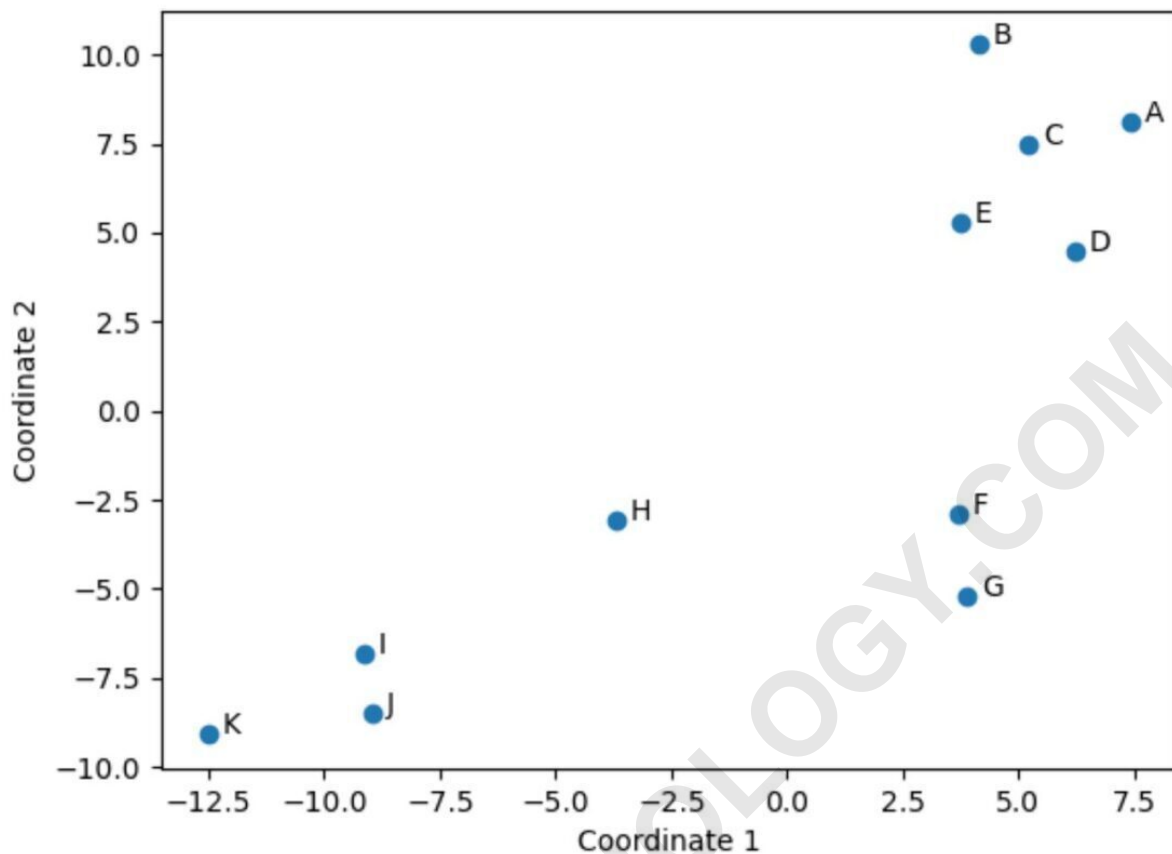
```
for i, txt in enumerate(df.index):
```

```
plt.annotate(txt, (scaled_df+.3, scaled_df))
```

```
#display scatterplot
```

```
plt.show()
```

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Players from the original DataFrame who have similar values across the original four columns (points, assists, blocks, and rebounds) are located close to each other in the plot.

For example, players F and G are located close to each other. Here are their values from the original DataFrame:

```
#select rows with index labels 'F' and 'G'  
df.loc]
```

points assists blocks rebounds

player

F 14 8 8 8

G 16 7 8 10

Their values for points, assists, blocks, and rebounds are all fairly similar, which explains why they're located so close together in the 2-D plot.

By contrast, consider players B and K who are located far apart in the plot.

```
#select rows with index labels 'B' and 'K'
```

```
df.loc]
```

points assists blocks rebounds

player

B 4 2 3 5

K 28 11 1 2

Thus, the 2-D plot is a nice way to visualize how similar each players are across all of the variables in the DataFrame.

Players who have similar stats are grouped close

together while players who have very different stats are located far apart from each other in the plot.

The following tutorials explain how to perform other common tasks in Python:

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