▼ K-means Clustering - 군집분석

```
import warnings
warnings.filterwarnings('ignore')
```

I. Import Packages and Lead Dataset

→ 1) Import Packages

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

→ 2) Load Dataset

· Load iris Dataset

```
from sklearn.datasets import load_iris
iris = load_iris()
```

• iris: Dictionary

X:iris.datay:iris.target

iris

```
[4.9, 2.5, 4.5, 1.7],

[7.3, 2.9, 6.3, 1.8],

[6.7, 2.5, 5.8, 1.8],

[7.2, 3.6, 6.1, 2.5],

[6.5, 3.2, 5.1, 2.],

[6.4, 2.7, 5.3, 1.9],

[6.8, 3., 5.5, 2.1],

[5.7, 2.5, 5., 2.],

[5.8, 2.8, 5.1, 2.4],

[6.4, 3.2, 5.3, 2.3],

[6.5, 3., 5.5, 1.8],

[7.7, 3.8, 6.7, 2.2],

[7.7, 2.6, 6.9, 2.3],
```

```
[0. , 2.2, 5. , 1.5],
     [6.9, 3.2, 5.7, 2.3],
     [5.6, 2.8, 4.9, 2.],
     [7.7, 2.8, 6.7, 2.],
     [6.3, 2.7, 4.9, 1.8],
     [6.7, 3.3, 5.7, 2.1],
     [7.2, 3.2, 6., 1.8],
     [6.2, 2.8, 4.8, 1.8],
     [6.1, 3., 4.9, 1.8],
     [6.4, 2.8, 5.6, 2.1],
     [7.2, 3., 5.8, 1.6],
     [7.4. 2.8. 6.1. 1.9].
     [7.9, 3.8, 6.4, 2.],
     [6.4, 2.8, 5.6, 2.2],
     [6.3, 2.8, 5.1, 1.5],
     [6.1, 2.6, 5.6, 1.4],
     [7.7, 3., 6.1, 2.3],
     [6.3, 3.4, 5.6, 2.4],
     [6.4, 3.1, 5.5, 1.8],
     [6., 3., 4.8, 1.8],
     [6.9, 3.1, 5.4, 2.1],
     [6.7, 3.1, 5.6, 2.4],
     [6.9, 3.1, 5.1, 2.3],
     [5.8, 2.7, 5.1, 1.9],
     [6.8, 3.2, 5.9, 2.3],
     [6.7, 3.3, 5.7, 2.5],
     [6.7, 3., 5.2, 2.3],
     [6.3, 2.5, 5., 1.9],
     [6.5, 3., 5.2, 2.],
     [6.2, 3.4, 5.4, 2.3],
[5.9, 3. , 5.1, 1.8]]), 'feature_names': ['sepal length (cm)',
'sepal width (cm)',
'petal length (cm)'
'petal width (cm)'],
filename': '/usr/local/lib/python3.7/dist-packages/sklearn/datasets/data/iris.csv',
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
    'target_names': array(['setosa', 'versicolor', 'virginica'], dtype='<U10')}
```

pandas DataFrame

DF.head(3)

| | sepal_length | sepal_width | petal_length | petal_width |
|---|--------------|-------------|--------------|-------------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 |

▼ II. K-means Clustering

→ 1) Modeling

• n_clusters : 군집 개수 지정

• init : 초기 중심 설정 방식(기본값)

• max_iter: 최대 반복 횟수

→ 2) Clustering Results

• 반복 횟수

```
kmeans_3.n_iter_
```

3

• 군집별 중심점

```
kmeans_3.cluster_centers_
```

```
array([[5.9016129 , 2.7483871 , 4.39354839, 1.43387097],
```

```
[5.006 , 3.428 , 1.462 , 0.246 ],
[6.85 , 3.07368421, 5.74210526, 2.07105263]])
```

• 군집 결과 레이블

```
kmeans_3.labels_
```

• 군집 중심까지의 제곱 거리의 합

```
kmeans_3.inertia_
```

78.85144142614601

→ III. Scree Plot

→ 1) DataFrame

| | sepal_length | sepal_width | petal_length | petal_width |
|---|--------------|-------------|--------------|-------------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 |

▼ 2) K(1~10) 군집분석

```
inertia = []
```

```
K = range(1,10)

for k in K:
    kmeanModel = KMeans(n_clusters = k).fit(Z)
    kmeanModel.fit(Z)
    inertia.append(kmeanModel.inertia_)
```

▼ 3) 군집 중심까지의 제곱 거리의 합

inertia

```
[681.3706,
152.34795176035792,
78.85144142614601,
57.25600931571815,
46.44618205128205,
39.03998724608725,
34.29822966507177,
30.06459307359308,
28.3473708513708541
```

→ 4) Plot the elbow

```
plt.figure(figsize = (9, 7))
plt.plot(K, inertia, 'bx-')
plt.xlabel('k')
plt.ylabel('inertia')
plt.title('Scree Plot with Kink')
plt.show()
```



▼ IV. Visualization with PCA(Principal Component Analysis)

▼ 1) target 및 cluster 추가

```
DF['cluster'] = kmeans_3.labels_
DF['target'] = iris.target

DF.head(3)
```

| | sepal_length | sepal_width | petal_length | petal_width | cluster | target |
|---|--------------|-------------|--------------|-------------|---------|--------|
| (| 5.1 | 3.5 | 1.4 | 0.2 | 1 | 0 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 | 1 | 0 |
| 2 | 2 4.7 | 3.2 | 1.3 | 0.2 | 1 | 0 |

▼ 2) 군집 결과 확인

DF.groupby('target')['cluster'].value_counts()

```
target cluster
0 1 50
1 0 48
2 2 2
2 2
2 36
0 14
Name: cluster, dtype: int64
```

▼ 3) PCA 차원 축소(4차원 -> 2차원)

```
from sklearn.decomposition import PCA

pca = PCA(n_components = 2)
pca_transformed = pca.fit_transform(iris.data)
```

pca_transformed[:5]

▼ 4) pca_x와 pca_y 추가

```
DF['pca_x'] = pca_transformed[:, 0]
DF['pca_y'] = pca_transformed[:, 1]
```

DF.head(5)

| | sepal_length | sepal_width | petal_length | petal_width | cluster | target | pca_x |
|---|--------------|-------------|--------------|-------------|---------|--------|-----------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 | 1 | 0 | -2.684126 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 | 1 | 0 | -2.714142 |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 | 1 | 0 | -2.888991 |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 | 1 | 0 | -2.745343 |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 | 1 | 0 | -2.728717 |

▼ 5) 2차원 시각화

• 군집 값 0, 1, 2 인덱스 추출

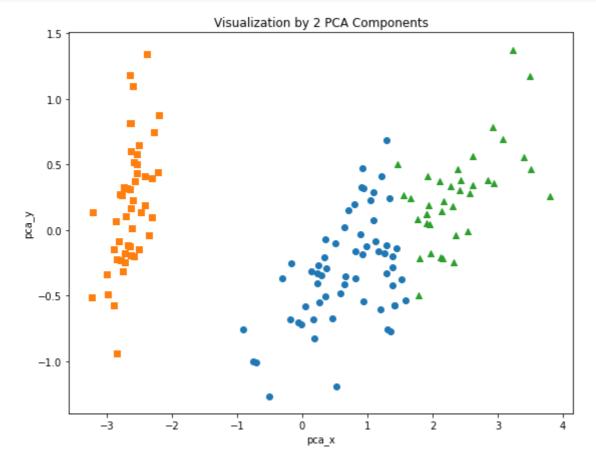
```
idx_0 = DF[DF['cluster'] == 0].index
idx_1 = DF[DF['cluster'] == 1].index
idx_2 = DF[DF['cluster'] == 2].index
```

idx_0 , idx_1 , idx_2

```
(Int64Index([ 50, 51, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 101, 106, 113, 114, 119, 121, 123, 126, 127, 133, 138, 142, 146, 149], dtype='int64'),
Int64Index([ 0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49], dtype='int64'),
Int64Index([ 52, 77, 100, 102, 103, 104, 105, 107, 108, 109, 110, 111, 112,
```

```
115, 116, 117, 118, 120, 122, 124, 125, 128, 129, 130, 131, 132, 134, 135, 136, 137, 139, 140, 141, 143, 144, 145, 147, 148], dtype='int64'))
```

• 0, 1, 2 인덱스 시각화



#

#

#

The End

#

#

#