

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

```
df=pd.read_csv('iris.csv')
```

```
df.head()
```

	sepalen	sepalwid	petallen	petalwid	label
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0

```
df.drop(['label'],axis=1,inplace=True)
```

```
df.head()
```

	sepalen	sepalwid	petallen	petalwid
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
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2

```
from sklearn.cluster import KMeans
```

```
km=KMeans(n_clusters=3)
```

```
km.fit(df)
```

```
ym=km.predict(df)
```

```
km.labels_
```

```
array([1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
1, 1, 1, 1, 1, 1, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2,
2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 0, 0, 0, 0, 2, 0, 0,
0,
0, 0, 0, 2, 2, 0, 0, 0, 0, 2, 0, 2, 0, 2, 0, 0, 2, 2, 0, 0, 0,
0,
0, 2, 0, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 2])
```

```
ym
```

```
array([1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
1, 1, 1, 1, 1, 1, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2,
2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 0, 0, 0, 0, 2, 0, 0,
0,
0, 0, 0, 2, 2, 0, 0, 0, 0, 2, 0, 2, 0, 2, 0, 0, 2, 2, 0, 0, 0,
0,
0, 2, 0, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 2])
```

```

1,      1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,      1, 1, 1, 1, 1, 1, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,      2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2,
2,      2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 0, 0, 0, 0, 2, 0, 0,
0,      0, 0, 0, 2, 2, 0, 0, 0, 0, 2, 0, 2, 0, 2, 0, 0, 2, 2, 0, 0, 0,
0,      0, 2, 0, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 2, 0, 0, 2])

```

```

from scipy.spatial.distance import cdist
distortions = []
K = range(1,20)
for k in K:
    km = KMeans(n_clusters=k).fit(df)
    km.fit(df)
    distortions.append(sum(np.min(cdist(df, km.cluster_centers_,
'euclidean'), axis=1)) / df.shape[0])

```

```

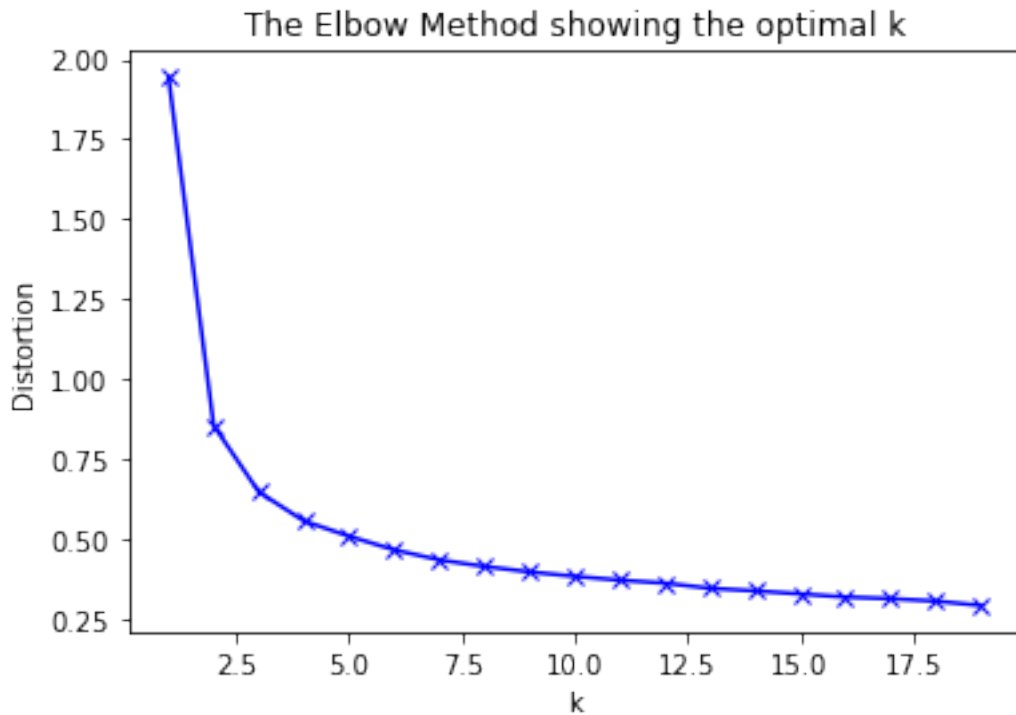
# Plot the elbow
plt.plot(K, distortions, 'bx-')
plt.xlabel('k')
plt.ylabel('Distortion')
plt.title('The Elbow Method showing the optimal k')
plt.show()

```

```

C:\Users\user\anaconda3\lib\site-packages\sklearn\cluster\
_kmeans.py:1036: UserWarning: KMeans is known to have a memory leak on
Windows with MKL, when there are less chunks than available threads.
You can avoid it by setting the environment variable
OMP_NUM_THREADS=1.
  warnings.warn(
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You can avoid it by setting the environment variable
OMP_NUM_THREADS=1.
  warnings.warn(

```



```

km=KMeans(n_clusters=3)
km.fit(df)
ym=km.predict(df)
center=km.cluster_centers_
km.labels_

array([[1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
      1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
      1, 1, 1, 1, 1, 1, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,
      2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,
      2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 0, 0, 0, 0, 2, 0, 0,
0,
      0, 0, 0, 2, 2, 0, 0, 0, 0, 2, 0, 2, 0, 2, 0, 0, 2, 2, 0, 0, 0,
0,
      0, 2, 0, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 0, 2, 0, 0, 2])

df.columns

Index(['sepalen', 'sepalwid', 'petallen', 'petalwid'],
      dtype='object')

tdf=pd.DataFrame(data=center,columns=df.columns)

tdf

```

```

    sepallen  sepalwid  petallen  petalwid
0  6.850000   3.073684   5.742105   2.071053
1  5.006000   3.428000   1.462000   0.246000
2  5.901613   2.748387   4.393548   1.433871

```

```
com_df=pd.concat([df,tdf])
```

```

from sklearn.preprocessing import StandardScaler
df_norm=StandardScaler().fit_transform(com_df)

```

```
df_norm[:10]
```

```

array([[ -0.90389781,   1.02375167,  -1.34096881,  -1.31681816],
       [ -1.14661325,  -0.13392882,  -1.34096881,  -1.31681816],
       [ -1.3893287 ,   0.32914338,  -1.39778675,  -1.31681816],
       [ -1.51068642,   0.09760728,  -1.28415088,  -1.31681816],
       [ -1.02525553,   1.25528777,  -1.34096881,  -1.31681816],
       [ -0.53982464,   1.94989607,  -1.17051501,  -1.05354215],
       [ -1.51068642,   0.79221557,  -1.34096881,  -1.18518015],
       [ -1.02525553,   0.79221557,  -1.28415088,  -1.31681816],
       [ -1.75340187,  -0.36546492,  -1.34096881,  -1.31681816],
       [ -1.14661325,   0.09760728,  -1.28415088,  -1.44845616]])

```

```

from sklearn.decomposition import PCA
pca = PCA(n_components=2)
trans_pc = pca.fit_transform(df_norm)

```

```
trans_pc[:10]
```

```

array([[ -2.26957633,   0.47908601],
       [ -2.08213805,  -0.68122811],
       [ -2.36712663,  -0.34682497],
       [ -2.30168974,  -0.60342672],
       [ -2.39559757,   0.64702343],
       [ -2.08352464,   1.49389065],
       [ -2.44866953,   0.04533184],
       [ -2.23708554,   0.22101257],
       [ -2.33554646,  -1.1239887 ],
       [ -2.18629012,  -0.47502155]])

```

```
trans_pc[: -3].shape
```

```
(150, 2)
```

```
trans_pc[-3:]
```

```

array([[ 1.92577832,   0.6018583 ],
       [-2.22920499,   0.28719207],
       [ 0.61244603,  -0.60506777]])

```

```
pc_df=pd.DataFrame(data = trans_pc[: -3], columns = ['pc1', 'pc2'])
```

```
pc_df.shape
```

```
plt.scatter(x=first.pc1,y=first.pc2,color='purple',label='Cluster 0')
plt.scatter(x=second.pc1,y=second.pc2,color='green',label='Cluster 1')
plt.scatter(x=third.pc1,y=third.pc2,color='blue',label='Cluster 2')
plt.scatter(x=center_df.pc1,y=center_df.pc2,color='red',label='Centers
')
plt.legend()
plt.show()
```

