데이터 다운로드 및 불러오기

```
!wget https://bit.ly/fruits_300_data -0 fruits_300.npy
import numpy as np
fruits = np.load('fruits_300.npy')
fruits_2d = fruits.reshape(-1, 100*100)
```

K-평균 기능 불러오기

```
from sklearn.cluster import KMeans

km = KMeans(n_clusters=3, random_state=42)
km.fit(fruits_2d)
```

K-평균 클러스터 확인

```
print(km.labels_)
print(np.unique(km.labels_, return_counts=True))
```

군집으로 형성된 각 클러스터별 확인

```
import matplotlib.pyplot as plt
def draw_fruits(arr, ratio=1):
   n = len(arr) # n은 샘플 개수입니다
   rows = int(np.ceil(n/10))
   cols = n if rows < 2 else 10
   fig, axs = plt.subplots(rows, cols,
                          figsize=(cols*ratio, rows*ratio), squeeze=False)
    for i in range(rows):
       for j in range(cols):
           if i*10 + j < n: # n 개까지만 그립니다.
               axs[i, j].imshow(arr[i*10 + j], cmap='gray_r')
           axs[i, j].axis('off')
    plt.show()
 draw_fruits(fruits[km.labels_==0])
```

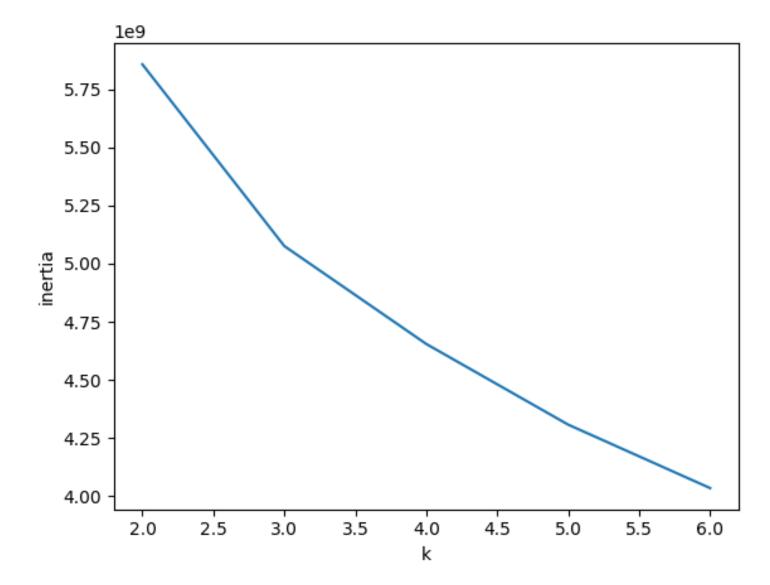
K-평균 클러스터 중심 확인(시각화)

```
draw_fruits(km.cluster_centers_.reshape(-1, 100, 100), ratio=3)
```

최적의 K 찾기

```
inertia = []
for k in range(2, 7):
    km = KMeans(n_clusters=k, n_init='auto', random_state=42)
    km.fit(fruits_2d)
    inertia.append(km.inertia_)

plt.plot(range(2, 7), inertia)
plt.xlabel('k')
plt.ylabel('inertia')
plt.show()
```



PCA 구동 및 차원 확인

```
from sklearn.decomposition import PCA

pca = PCA(n_components=50)
pca.fit(fruits_2d)

print(pca.components_.shape)
```

주성분 분석 결과 이미지 출력

```
import matplotlib.pyplot as plt
def draw_fruits(arr, ratio=1):
   n = len(arr) # n은 샘플 개수입니다
   rows = int(np.ceil(n/10))
   cols = n if rows < 2 else 10
   fig, axs = plt.subplots(rows, cols,
                          figsize=(cols*ratio, rows*ratio), squeeze=False)
    for i in range(rows):
       for j in range(cols):
           if i*10 + j < n: # n 개까지만 그립니다.
               axs[i, j].imshow(arr[i*10 + j], cmap='gray_r')
           axs[i, j].axis('off')
    plt.show()
draw_fruits(pca.components_.reshape(-1, 100, 100))
```

원본 이미지 -> PCA 변환 값

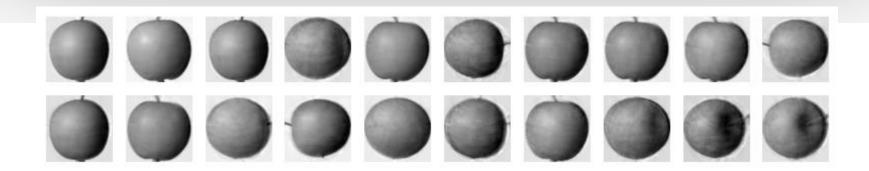
```
fruits_pca = pca.transform(fruits_2d)

fruits_inverse = pca.inverse_transform(fruits_pca)
print(fruits_inverse.shape)

fruits_reconstruct = fruits_inverse.reshape(-1, 100, 100)
```

이미지 출력

```
for start in [0, 100, 200]:
    draw_fruits(fruits_reconstruct[start:start+100])
    print("\n")
```



PCA 분산 확인 및 그래프 확인

```
print(np.sum(pca.explained_variance_ratio_))
plt.plot(pca.explained_variance_ratio_)
```

