



PCA

▼ Manual PCA

Imports

```
1 import numpy as np
2 import pandas as pd
3 import matplotlib.pyplot as plt
4 import seaborn as sns
```

✓ 0.2s

```
1 df = pd.read_csv('cancer_tumor_data_features.csv')
```

✓ 0.1s

```
1 df
```

✓ 0.2s

| | mean radius | mean texture | mean perimeter | mean area | mean smoothness | mean compactness | mean concavity | mean concave points | mean symmetry | mean fractal dimension |
|---|----------------|-----------------|-------------------|--------------|--------------------|---------------------|-------------------|---------------------------|------------------|------------------------------|
| 0 | 17.99 | 10.38 | 122.80 | 1001.0 | 0.11840 | 0.27760 | 0.30010 | 0.14710 | 0.2419 | 0.07871 |
| 1 | 20.57 | 17.77 | 132.90 | 1326.0 | 0.08474 | 0.07864 | 0.08690 | 0.07017 | 0.1812 | 0.05667 |
| 2 | 19.69 | 21.25 | 130.00 | 1203.0 | 0.10960 | 0.15990 | 0.19740 | 0.12790 | 0.2069 | 0.05999 |
| 3 | 11.42 | 20.38 | 77.58 | 386.1 | 0.14250 | 0.28390 | 0.24140 | 0.10520 | 0.2597 | 0.09744 |
| 4 | 20.29 | 14.34 | 135.10 | 1297.0 | 0.10030 | 0.13280 | 0.19800 | 0.10430 | 0.1809 | 0.05883 |

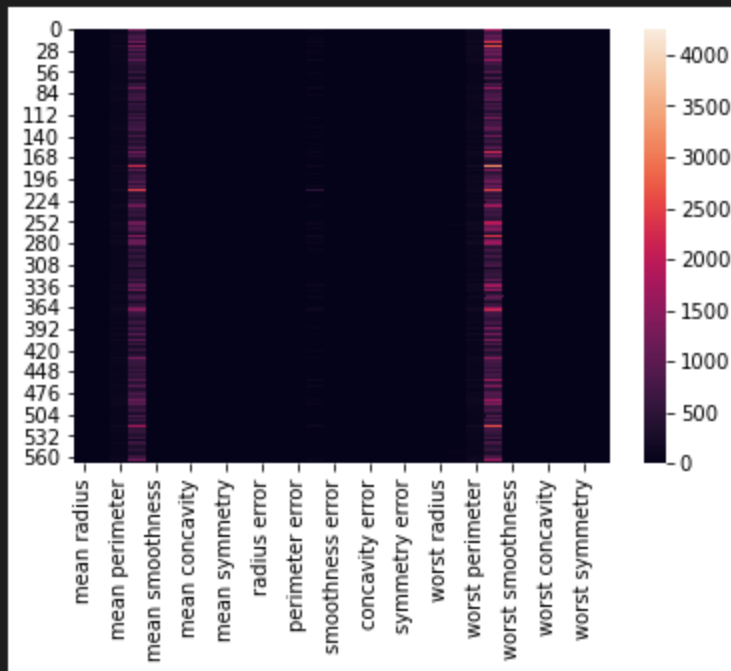
Manual Construction of PCA

Scaling Data

```
1 sns.heatmap(df)
2 # difficult to understand
```

✓ 1.2s

<AxesSubplot:>



```
1 from sklearn.preprocessing import StandardScaler
```

✓ 0.1s

```
1 scaler = StandardScaler()
```

✓ 0.1s

```
1 scaled_X = scaler.fit_transform(df)
```

✓ 0.1s

```
1 scaled_X.mean()
```

✓ 0.1s

-6.826538293184326e-17

```
1 # Grab Covariance Matrix
```

```
2 covariance_matrix = np.cov(scaled_X, rowvar=False)
```

✓ 0.1s

```
1 eigen_values, eigen_vectors = np.linalg.eig(covariance_matrix)
```

✓ 0.1s

```
1 # Choose some number of components
```

```
2 num_components=2
```

✓ 0.8s

```
1 # index [0,1,2]
```

```
2 np.argsort([2,1,3])
```

```
3 # sorted the index not the values
```

✓ 0.1s

array([1, 0, 2], dtype=int64)

```
1 sorted_key = np.argsort(eigen_values)[::-1][:num_components]
```

✓ 0.1s

```
1 eigen_values, eigen_vectors = eigen_vectors[sorted_key], eigen_vectors[:,sorted_key]
```

✓ 0.1s

```
1 principal_components = np.dot(scaled_X, eigen_vectors)
2 principal_components
```

✓ 0.1s

```
array([[ 9.19283683,  1.94858307],
       [ 2.3878018 , -3.76817174],
       [ 5.73389628, -1.0751738 ],
       ...,
       [ 1.25617928, -1.90229671],
       [10.37479406,  1.67201011],
       [-5.4752433 , -0.67063679]])
```

```

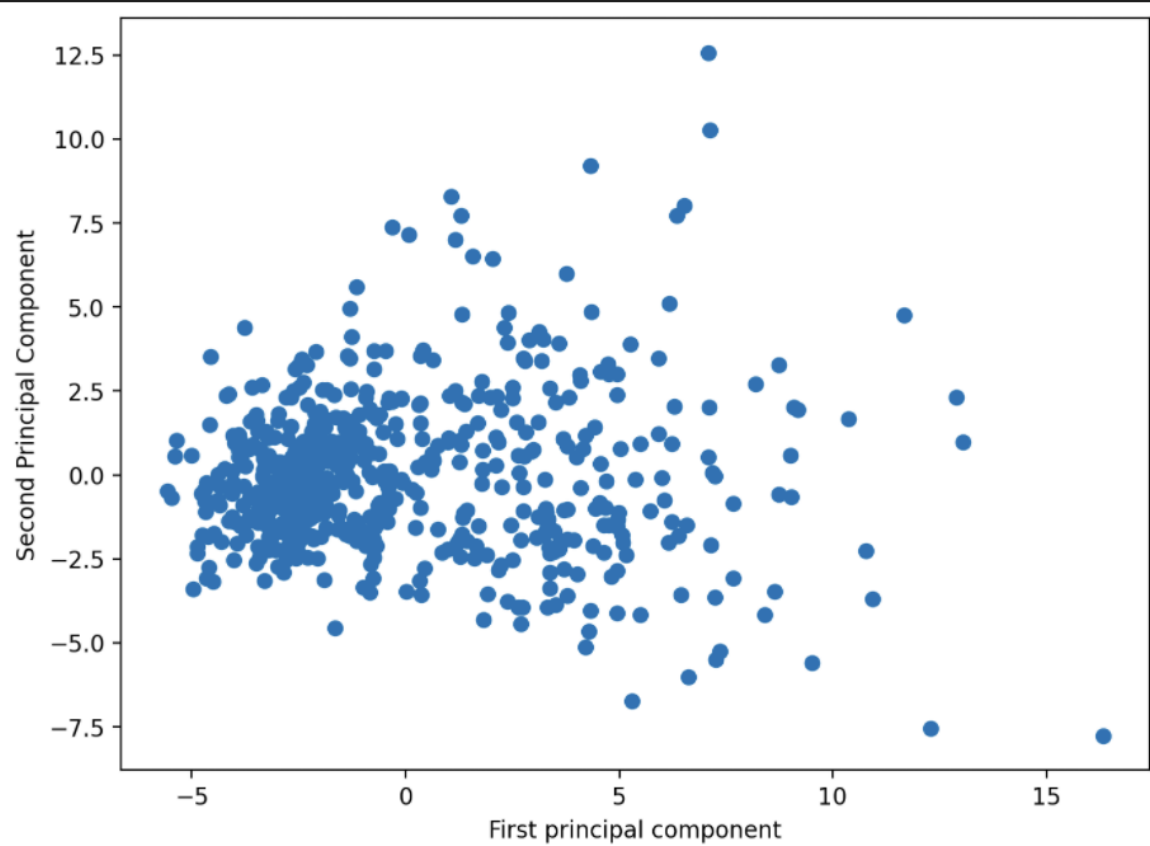
1 plt.figure(figsize=(8,6), dpi=200)
2 plt.scatter(principal_components[:,0],principal_components[:,1])
3 plt.xlabel('First principal component')
4 plt.ylabel('Second Principal Component')

```

✓ 0.4s

Python

Text(0, 0.5, 'Second Principal Component')



```

1 from sklearn.datasets import load_breast_cancer

```

✓ 0.8s

[+ Code](#) [+ Markdown](#)

```

1 cancer_dictionary = load_breast_cancer()

```

✓ 0.1s

```

1 cancer_dictionary.keys()

```

✓ 0.1s

```
dict_keys(['data', 'target', 'frame', 'target_names', 'DESCR', 'feature_names', 'filename'])
```

```
1 print(cancer_dictionary["DESCR"])
```

✓ 0.1s

Output exceeds the [size limit](#). Open the full output data [in a text editor](#)

.. _breast_cancer_dataset:

Breast cancer wisconsin (diagnostic) dataset

****Data Set Characteristics:****

:Number of Instances: 569

:Number of Attributes: 30 numeric, predictive attributes and the class

:Attribute Information:

- radius (mean of distances from center to points on the perimeter)
- texture (standard deviation of gray-scale values)
- perimeter

...

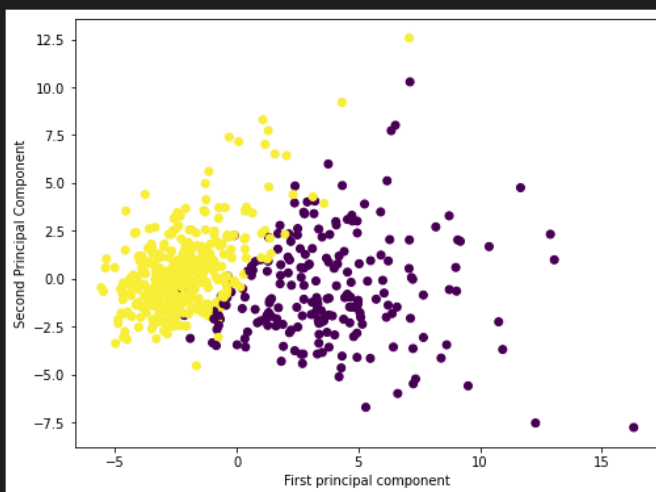
July-August 1995.

- W.H. Wolberg, W.N. Street, and O.L. Mangasarian. Machine learning techniques to diagnose breast cancer from fine-needle aspirates. Cancer Letters 77 (1994) 163-171.

```
1 plt.figure(figsize=(8,6))
2 plt.scatter(principal_components[:,0],principal_components[:,1],c=cancer_dictionary['target'])
3 plt.xlabel('First principal component')
4 plt.ylabel('Second Principal Component')
```

✓ 0.7s

Text(0, 0.5, 'Second Principal Component')



▼ SKLearn PCA

```
1 import numpy as np
2 import pandas as pd
3 import matplotlib.pyplot as plt
4 import seaborn as sns
```

✓ 7.1s

```
1 df = pd.read_csv('cancer_tumor_data_features.csv')
```

✓ 0.1s

```
1 from sklearn.preprocessing import StandardScaler
```

✓ 0.1s

```
1 scaler = StandardScaler()
2 scaled_X = scaler.fit_transform(df)
3 scaled_X
```

✓ 0.2s

```
array([[ 1.09706398, -2.07333501,  1.26993369, ...,  2.29607613,
         2.75062224,  1.93701461],
       [ 1.82982061, -0.35363241,  1.68595471, ...,  1.0870843 ,
        -0.24388967,  0.28118999],
       [ 1.57988811,  0.45618695,  1.56650313, ...,  1.95500035,
        1.152255  ,  0.20139121],
```

```
1 from sklearn.decomposition import PCA
✓ 0.1s

1 pca_model = PCA(n_components=2)
✓ 0.1s

1 pca_model.fit(scaled_X)
✓ 0.2s

PCA(n_components=2)

1 pca_model.transform(scaled_X)
✓ 0.1s

array([[ 9.19283683,  1.94858307],
       [ 2.3878018 , -3.76817174],
       [ 5.73389628, -1.0751738 ],
       ...,
       [ 1.25617928, -1.90229671],
       [10.37479406,  1.67201011],
       [-5.4752433 , -0.67063679]])

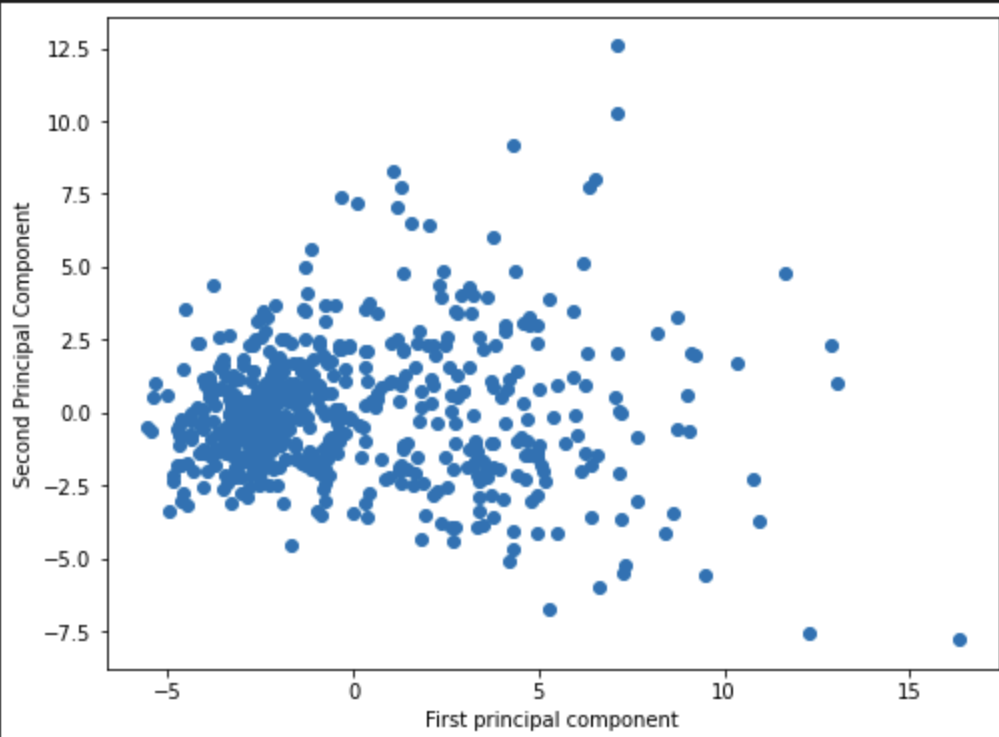
1 pc_results = pca_model.fit_transform(scaled_X)
✓ 0.1s
```



```
1 plt.figure(figsize=(8,6))
2 plt.scatter(pc_results[:,0],pc_results[:,1])
3 plt.xlabel('First principal component')
4 plt.ylabel('Second Principal Component')
```

✓ 0.3s

```
Text(0, 0.5, 'Second Principal Component')
```



```
1 from sklearn.datasets import load_breast_cancer
```

✓ 0.1s

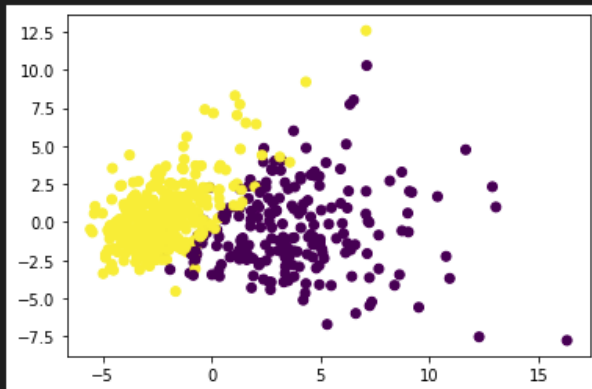
```
1 cancer_dictionary = load_breast_cancer()
```

✓ 0.8s

```
1 plt.scatter(pc_results[:,0],pc_results[:,1], c=cancer_dictionary["target"])
```

✓ 0.3s

<matplotlib.collections.PathCollection at 0x2ceb3b2aee0>



```
1 pca_model.n_components
✓ 0.9s

2

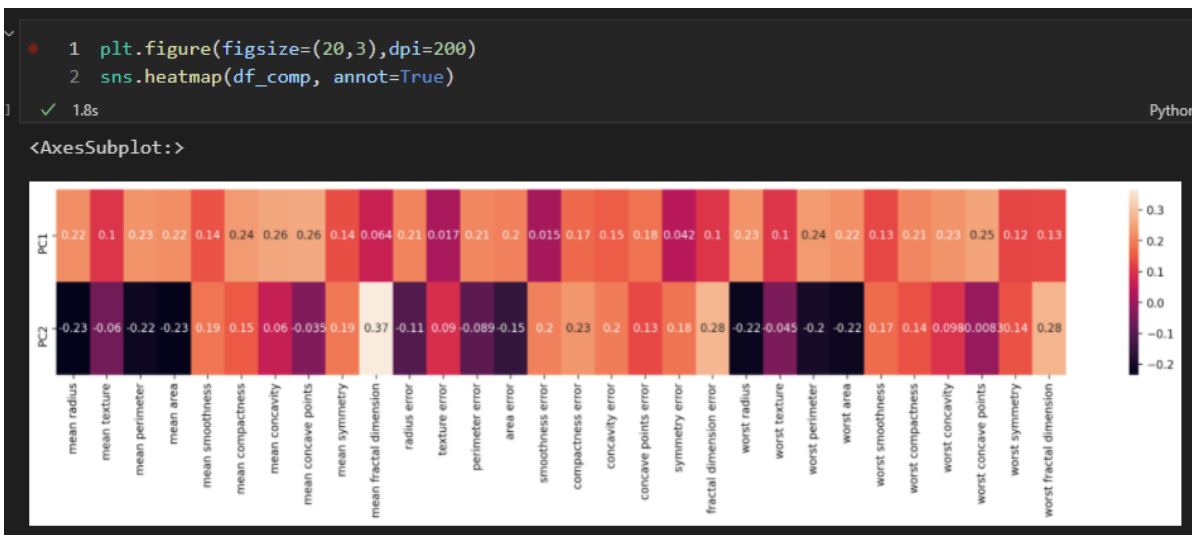
1 pca_model.components_
✓ 0.1s

array([[ 0.21890244,  0.10372458,  0.22753729,  0.22099499,  0.14258969,
         0.23928535,  0.25840048,  0.26085376,  0.13816696,  0.06436335,
         0.20597878,  0.01742803,  0.21132592,  0.20286964,  0.01453145,
         0.17039345,  0.15358979,  0.1834174 ,  0.04249842,  0.10256832,
         0.22799663,  0.10446933,  0.23663968,  0.22487053,  0.12795256,
         0.21009588,  0.22876753,  0.25088597,  0.12290456,  0.13178394],
        [-0.23385713, -0.05970609, -0.21518136, -0.23107671,  0.18611302,
         0.15189161,  0.06016536, -0.0347675 ,  0.19034877,  0.36657547,
        -0.10555215,  0.08997968, -0.08945723, -0.15229263,  0.20443045,
         0.2327159 ,  0.19720728,  0.13032156,  0.183848 ,  0.28009203,
        -0.21986638, -0.0454673 , -0.19987843, -0.21935186,  0.17230435,
         0.14359317,  0.09796411, -0.00825724,  0.14188335,  0.27533947]])

1 df_comp = pd.DataFrame(pca_model.components_, index=['PC1','PC2'],columns=df.columns)
✓ 0.8s

1 df_comp
✓ 0.1s
```

| | mean radius | mean texture | mean perimeter | mean area | mean smoothness | mean compactness | mean concavity | mean concave points | mean symmetry |
|-----|----------------|-----------------|-------------------|--------------|--------------------|---------------------|-------------------|---------------------------|------------------|
| PC1 | 0.218902 | 0.103725 | 0.227537 | 0.220995 | 0.142590 | 0.239285 | 0.258400 | 0.260854 | 0.138167 |
| PC2 | -0.233857 | -0.059706 | -0.215181 | -0.231077 | 0.186113 | 0.151892 | 0.060165 | -0.034768 | 0.190349 |



```
1  pca_model.explained_variance_ratio_
✓ 0.1s
array([0.44272026, 0.18971182])

1  np.sum(pca_model.explained_variance_ratio_)
✓ 0.9s
0.6324320765155946

1  pca_30 = PCA(n_components=30)
✓ 0.1s

1  pca_30.fit(scaled_X)
✓ 0.1s
PCA(n_components=30)

1  pca_30.explained_variance_ratio_
✓ 0.1s
array([4.42720256e-01, 1.89711820e-01, 9.39316326e-02, 6.60213492e-02,
       5.49576849e-02, 4.02452204e-02, 2.25073371e-02, 1.58872380e-02,
       1.38964937e-02, 1.16897819e-02, 9.79718988e-03, 8.70537901e-03,
       8.04524987e-03, 5.23365745e-03, 3.13783217e-03, 2.66209337e-03,
       1.97996793e-03, 1.75395945e-03, 1.64925306e-03, 1.03864675e-03,
       9.99096464e-04, 9.14646751e-04, 8.11361259e-04, 6.01833567e-04,
       5.16042379e-04, 2.72587995e-04, 2.30015463e-04, 5.29779290e-05,
       2.49601032e-05, 4.43482743e-06])
```

```
1 np.sum(pca_30.explained_variance_ratio_)
```

✓ 0.1s

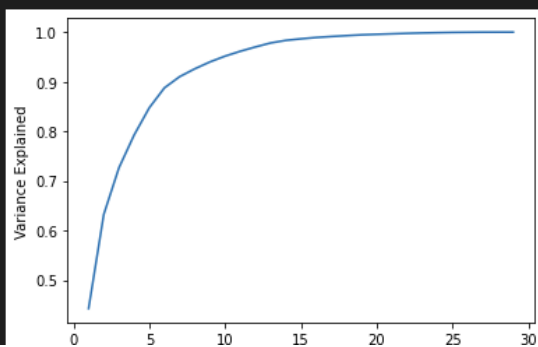
1.0

```
1 explained_variance = []
2
3 for n in range(1,30):
4     pca = PCA(n_components=n)
5     pca.fit(scaled_X)
6
7     explained_variance.append(np.sum(pca.explained_variance_ratio_))
```

✓ 0.4s

```
1 plt.plot(range(1,30),explained_variance)
2 plt.xlabel("Number of Components")
3 plt.ylabel("Variance Explained");
```

✓ 0.4s



▼ Handwritten Digits

Imports

```
1 import numpy as np
2 import pandas as pd
3 import matplotlib.pyplot as plt
4 import seaborn as sns
```

✓ 7.4s

```
1 digits = pd.read_csv('digits.csv')
```

✓ 0.1s

```
1 digits
```

✓ 0.1s

| | pixel_0_0 | pixel_0_1 | pixel_0_2 | pixel_0_3 | pixel_0_4 | pixel_0_5 | pixel_0_6 | pixel_0_7 | pixel_1_0 | pixel_1_1 |
|---|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| 0 | 0.0 | 0.0 | 5.0 | 13.0 | 9.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 1 | 0.0 | 0.0 | 0.0 | 12.0 | 13.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 2 | 0.0 | 0.0 | 0.0 | 4.0 | 15.0 | 12.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 3 | 0.0 | 0.0 | 7.0 | 15.0 | 13.0 | 1.0 | 0.0 | 0.0 | 0.0 | 8.0 |
| 4 | 0.0 | 0.0 | 0.0 | 1.0 | 11.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |

```
1 pixels = digits.drop('number_label',axis=1)
2 pixels
```

✓ 0.1s

| | pixel_0_0 | pixel_0_1 | pixel_0_2 | pixel_0_3 | pixel_0_4 | pixel_0_5 | pixel_0_6 |
|-----|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| 0 | 0.0 | 0.0 | 5.0 | 13.0 | 9.0 | 1.0 | 0.0 |
| 1 | 0.0 | 0.0 | 0.0 | 12.0 | 13.0 | 5.0 | 0.0 |
| 2 | 0.0 | 0.0 | 0.0 | 4.0 | 15.0 | 12.0 | 0.0 |
| 3 | 0.0 | 0.0 | 7.0 | 15.0 | 13.0 | 1.0 | 0.0 |
| 4 | 0.0 | 0.0 | 0.0 | 1.0 | 11.0 | 0.0 | 0.0 |
| ... | ... | ... | ... | ... | ... | ... | ... |

```
1 single_image = pixels.iloc[0]
2 single_image
✓ 0.1s

pixel_0_0    0.0
pixel_0_1    0.0
pixel_0_2     5.0
pixel_0_3    13.0
pixel_0_4     9.0
...
pixel_7_3    13.0
pixel_7_4    10.0
pixel_7_5     0.0
pixel_7_6     0.0
pixel_7_7     0.0
Name: 0, Length: 64, dtype: float64

1 single_image.to_numpy()
✓ 0.1s

array([ 0.,  0.,  5., 13.,  9.,  1.,  0.,  0.,  0.,  0., 13., 15., 10.,
        15.,  5.,  0.,  0.,  3., 15.,  2.,  0., 11.,  8.,  0.,  0.,  4.,
        12.,  0.,  0.,  8.,  8.,  0.,  0.,  5.,  8.,  0.,  0.,  9.,  8.,
         0.,  0.,  4., 11.,  0.,  1., 12.,  7.,  0.,  0.,  2., 14.,  5.,
        10., 12.,  0.,  0.,  0.,  0.,  6., 13., 10.,  0.,  0.,  0.])

1 single_image.to_numpy().shape
✓ 0.1s

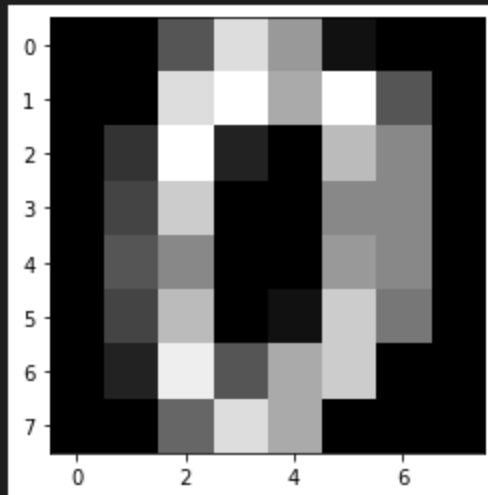
(64,)
```

```
1 num = single_image.to_numpy().reshape(8,8)
✓ 0.1s
```

```
1 plt.imshow(num, cmap="gray")
```

✓ 0.5s

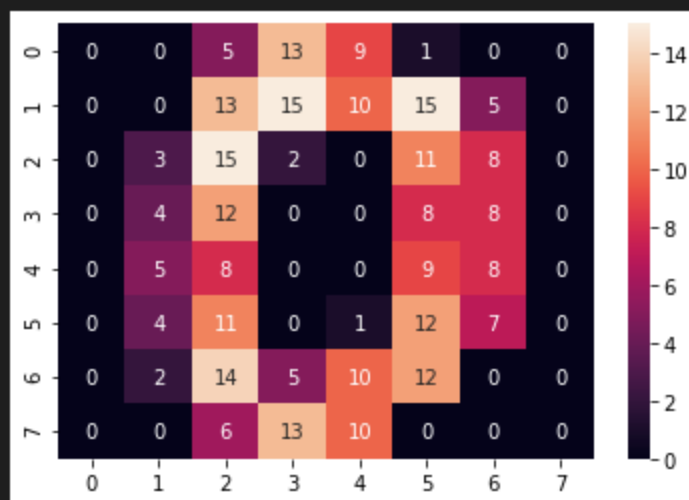
<matplotlib.image.AxesImage at 0x12a7769ac40>



```
1 sns.heatmap(num, annot=True)
```

✓ 1.1s

<AxesSubplot:>



Scaling Data

```
1 from sklearn.preprocessing import StandardScaler
```

✓ 0.5s

```
1 scaler = StandardScaler()
```

✓ 0.5s

```
1 scaled_pixels = scaler.fit_transform(pixels)
```

✓ 0.8s

```
1 pixels
```

✓ 0.2s

| | pixel_0_0 | pixel_0_1 | pixel_0_2 | pixel_0_3 | pixel_0_4 | pixel_0_5 | pixel_0_6 | pixel_0_7 |
|------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| 0 | 0.0 | 0.0 | 5.0 | 13.0 | 9.0 | 1.0 | 0.0 | 0.0 |
| 1 | 0.0 | 0.0 | 0.0 | 12.0 | 13.0 | 5.0 | 0.0 | 0.0 |
| 2 | 0.0 | 0.0 | 0.0 | 4.0 | 15.0 | 12.0 | 0.0 | 0.0 |
| 3 | 0.0 | 0.0 | 7.0 | 15.0 | 13.0 | 1.0 | 0.0 | 0.0 |
| 4 | 0.0 | 0.0 | 0.0 | 1.0 | 11.0 | 0.0 | 0.0 | 0.0 |
| ... | ... | ... | ... | ... | ... | ... | ... | ... |
| 1792 | 0.0 | 0.0 | 4.0 | 10.0 | 13.0 | 6.0 | 0.0 | 0.0 |
| 1793 | 0.0 | 0.0 | 6.0 | 16.0 | 13.0 | 11.0 | 1.0 | 0.0 |
| 1794 | 0.0 | 0.0 | 1.0 | 11.0 | 15.0 | 1.0 | 0.0 | 0.0 |
| 1795 | 0.0 | 0.0 | 2.0 | 10.0 | 7.0 | 0.0 | 0.0 | 0.0 |
| 1796 | 0.0 | 0.0 | 10.0 | 14.0 | 8.0 | 1.0 | 0.0 | 0.0 |

1797 rows × 64 columns

PCA

```
1 from sklearn.decomposition import PCA
```

✓ 0.1s

```
1 pca_model = PCA(n_components=2)
```

✓ 0.8s

```
1 pca_pixels = pca_model.fit_transform(scaled_pixels)
```

✓ 0.1s

```
1 pca_model.explained_variance_ratio_
```

✓ 0.1s

```
array([0.12033916, 0.09561054])
```

```
1 np.sum(pca_model.explained_variance_ratio_)
```

✓ 0.1s

```
0.2159497049303386
```

```
1 pca_pixels[:,1]
```

✓ 0.1s

```
array([-0.95450277,  0.92463286, -0.31734493, ..., -0.14782995,  
       -0.38088582, -2.22747391])
```

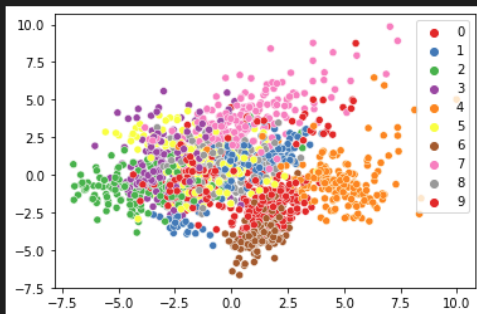
```
1 s.scatterplot(pca_pixels[:,0] ,pca_pixels[:,1], hue=digits["number_label"].values, palette="Set1")
```

✓ 1.6s

Python

C:\Users\mbatu\anaconda3\lib\site-packages\seaborn\decorators.py:36: FutureWarning: Pass the following variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.
warnings.warn(

<AxesSubplot:>



```
1 pca_model = PCA(n_components=3)
```

✓ 0.1s

```
1 pca_pixels = pca_model.fit_transform(scaled_pixels)
```

✓ 0.1s

```
1 from mpl_toolkits import mplot3d
```

✓ 0.1s

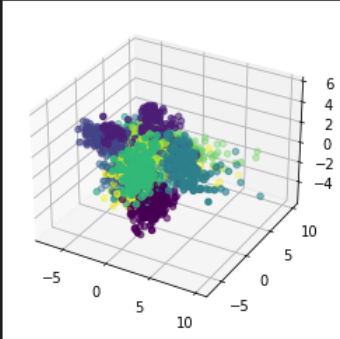
```

1
2 ax = plt.axes(projection='3d')
3 ax.scatter3D(pca_pixels[:,0],pca_pixels[:,1],pca_pixels[:,2],c=digits['number_label']);

```

✓ 0.4s

Pytho



```

1 %matplotlib notebook
2 toadece jupyter notebookta çalışılıyo
3 plt.figure(figsize=(8,8),dpi=150)
4 ax = plt.axes(projection='3d')
5 ax.scatter3D(pca_pixels[:,0],pca_pixels[:,1],pca_pixels[:,2],c=digits['number_label']);

```