```
import numpy as np
import pandas as pd
import io
import matplotlib.pyplot as plt
import scipy
from google.colab import files
uploaded = files.upload()
    Choose Files train_set.mat
    • train_set.mat(n/a) - 3298808 bytes, last modified: 4/19/2023 - 100% done
    Saving train_set.mat to train_set.mat
train_set = scipy.io.loadmat("train_set", mdict = None, appendmat = True)
train_set
'data': array([[ 39, 44, 53, ..., 29, 26, 29],
           [ 63, 53, 35, ..., 41, 10, 24],
[ 64, 76, 80, ..., 35, 37, 39],
           [111, 114, 112, ..., 88, 86, 92],
           [110, 112, 113, ..., 92, 87, 90],
           [111, 111, 110, ..., 88, 79, 90]], dtype=uint8),
     'labels': array([[ 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 1,
             2, 2, 2, 2, 2, 2, 2, 3, 3, 3, 3, 3, 3, 3, 3, 4, 4, 4, 4, 4, 4, 4, 4, 5, 5, 5, 5, 5, 5, 5, 5, 5, 5,
                                                       3, 3, 3, 3,
             12, 12, 12, 12, 12, 12, 12, 13, 13, 13, 13, 13, 13, 13,
            14, 14, 14, 14, 14, 14, 14, 15, 15, 15, 15, 15, 15, 15, 15,
            16, 16, 16, 16, 16, 16, 16, 17, 17, 17, 17, 17, 17, 17, 17,
            18, 18, 18, 18, 18, 18, 18, 19, 19, 19, 19, 19, 19, 19, 19,
            20, 20, 20, 20, 20, 20, 20, 20, 21, 21, 21, 21, 21, 21, 21,
            24, 24, 24, 24, 24, 24, 24, 25, 25, 25, 25, 25, 25, 25,
            26, 26, 26, 26, 26, 26, 26, 27, 27, 27, 27, 27, 27, 27,
            34, 34, 34, 34, 34, 34, 34, 35, 35, 35, 35, 35, 35, 35,
            36, 36, 36, 36, 36, 36, 36, 37, 37, 37, 37, 37, 37, 37, 37,
            38, 38, 38, 38, 38, 38, 38, 38, 39, 39, 39, 39, 39, 39, 39, 39]],
          dtype=int32)}
from google.colab import files
uploaded = files.upload()
    Choose Files test_set.mat

    test_set.mat(n/a) - 824888 bytes, last modified: 4/19/2023 - 100% done

    Saving test set.mat to test set.mat
test_set = scipy.io.loadmat("test_set", mdict = None, appendmat = True)
test set
    {'_header__': b'MATLAB 5.0 MAT-file Platform: nt, Created on: Mon Apr 10 23:29:46 2023',
    '_version__': '1.0',
    '_globals__': [],
     'data': array([[ 48, 49, 45, ..., 47, 46, 46],
           [ 60, 60, 62, ..., 32, 34, 34],
           [140, 134, 135, ..., 25, 26, 49],
           [131, 128, 126, ..., 41, 40, 35],
[105, 102, 106, ..., 72, 62, 75],
           [100, 110, 107, ..., 64, 71, 59]], dtype=uint8),
     'labels': array([[ 0, 0, 1, 1, 2, 2, 3, 3, 4, 4, 5, 5,
             8, 8, 9, 9, 10, 10, 11, 11, 12, 12, 13, 13, 14, 14, 15, 15,
            16, 16, 17, 17, 18, 18, 19, 19, 20, 20, 21, 21, 22, 22, 23, 23,
            24, 24, 25, 25, 26, 26, 27, 27, 28, 28, 29, 29, 30, 30, 31, 31,
            32, 32, 33, 33, 34, 34, 35, 35, 36, 36, 37, 37, 38, 38, 39, 39]],
          dtvpe=int32)}
```

Visualizing one image for each label/identity in the training set

```
# create figure
fig = plt.figure(figsize=(10, 7))

# setting values to rows and column variables
rows = 5
columns = 8
for i in range(0,320,8):
    t = np.reshape(train_set['data'][i],(112,92))
    # Adds a subplot at the 1st position
    fig.add_subplot(rows, columns, ((int)(i/8) + 1))
    # showing image
    plt.imshow(t, cmap = 'gray')
    plt.title("label " + str((int)(i/8)), size = 6)
    plt.axis('off')
```



Visualising each image in the test_set

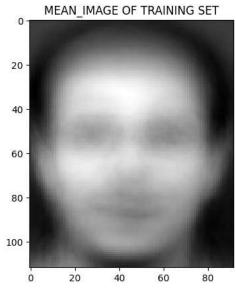
```
# create figure
fig = plt.figure(figsize=(10, 7))

# setting values to rows and column variables
rows = 5
columns = 8
for i in range(0,80,2):
    t = np.reshape(test_set['data'][i],(112,92))
    # Adds a subplot at the 1st position
    fig.add_subplot(rows, columns, ((int)(i/2) + 1))
    # showing image
    plt.imshow(t, cmap = 'gray')
    plt.title("label " + str((int)(i/2)), size = 6)
    plt.axis('off')
```



```
means = np.mean(train_set['data'],axis = 0)
stdev = np.std(train_set['data'],axis = 0)
mean_image = np.reshape(means, (112,92))
plt.imshow(mean_image,cmap ='gray')
plt.title("MEAN_IMAGE OF TRAINING SET")
```

Text(0.5, 1.0, 'MEAN_IMAGE OF TRAINING SET')

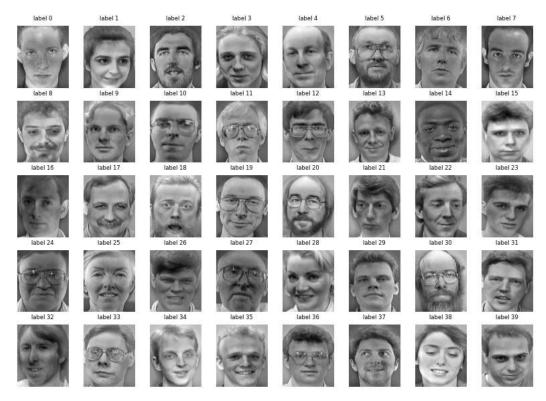


Centering the whole data set

```
centered_train_data = np.zeros((320,10304))
centered_test_data = np.zeros((80,10304))
for i in range(320):
 centered_train_data[i] += (train_set['data'][i] - means)
for i in range(80):
 centered_test_data[i] += (test_set['data'][i] - means)
```

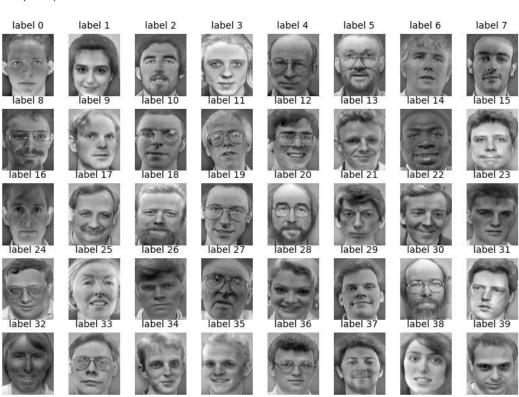
Visualsing the centered training set

```
# create figure
fig = plt.figure(figsize=(10, 7))
\ensuremath{\text{\#}} setting values to rows and column variables
rows = 5
columns = 8
for i in range(0,320,8):
  t = np.reshape(centered_train_data[i],(112,92))
  \mbox{\#} Adds a subplot at the 1st position
  fig.add_subplot(rows, columns, ((int)(i/8) + 1))
  # showing image
  plt.imshow(t, cmap = "gray")
plt.title("label " + str((int)(i/8)), size = 6)
  plt.axis('off')
```



Visualising the centered test set

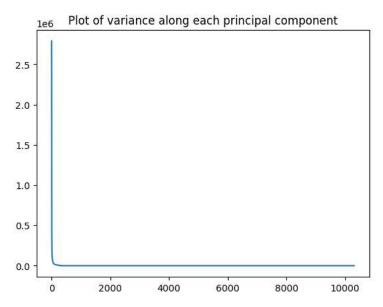
```
# create figure
fig = plt.figure(figsize=(10, 7))
# setting values to rows and column variables
rows = 5
columns = 8
for i in range(0,80,2):
    t = np.reshape(centered_test_data[i],(112,92))
    # Adds a subplot at the 1st position
    fig.add_subplot(rows, columns, ((int)(i/2) + 1))
    # showing image
    plt.imshow(t, cmap = "gray")
    plt.title("label " + str((int)(i/2)), size = 10)
    plt.axis('off')
```



Let $X \in R$ N×D be a centered data matrix with N data points. Let vj be the j - th eigenvector of XXT . The j - th principal component is along XT vj . This approach for finding the principal components is preferable over eigen decomposition of the covariance matrix when D > N

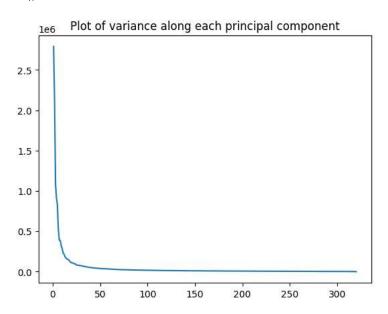
```
covar = (centered_train_data @ centered_train_data.T)/320
eigen_values , eigen_vectors = np.linalg.eigh(covariance)
# eigen vectors in columns in ascending order
eigen_vectors_order = eigen_vectors[:,::-1] # flipping in descending order
eigen_values_order = eigen_values[::-1]
#eigen_faces = (np.dot(centered_train_data.T,eigen_vectors)).T  # principle_components
#print(eigen_faces.shape)
#normalized_eigen_faces = eigen_faces/ np.linalg.norm(eigen_faces, axis = 0)
#eigen_vectors = eigen_vectors.T
eigen_vectors_order[:5]
     array([[ 0.00133785, -0.01462058, -0.01818825, ..., 0.
           [\ 0.00138743,\ -0.01450587,\ -0.01807086,\ \dots,\ -0.36739404,
              0.33220772, 0.40898063],
            [\ 0.00146925,\ -0.01458946,\ -0.01828916,\ \ldots,\ 0.36538352,
            -0.28295005, 0.32860456],
[ 0.00134757, -0.01457466, -0.01815938, ..., -0.25378774,
             -0.36623836, -0.34596835]])
# create figure
fig = plt.figure(figsize=(10, 7))
# setting values to rows and column variables
rows = 5
columns = 5
for i in range(0,25):
 t = np.reshape(eigen_vectors_order[:,i],(112,92))
 # Adds a subplot at the 1st position
 fig.add_subplot(rows, columns,i + 1)
  # showing image
  plt.imshow(t, cmap = "gray")
 plt.title("label " + str(i + 1), size = 6)
  plt.axis('off')
```

```
x = np.linspace(1,10304,10304)
plt.plot(x.reshape(-1,1), eigen_values_order.reshape(-1,1))
plt.title("Plot of variance along each principal component")
plt.show()
```



Since we cannot deduce anything from this, I'm drawing the plot only for 1000 eigen values

```
x = np.linspace(1,320,320)
plt.plot(x.reshape(-1,1), eigen_values_order[:320].reshape(-1,1))
plt.title("Plot of variance along each principal component")
plt.show()
```



• Sort the variance along each principal component in descending order and plot them. How many principal components do you need to capture 95% of the total variance? Let's call this number 'k'. If the variance along i – th principal component is given by λi , then k is the smallest number for which (sum of first k eigen values)/(sum of all the eigen values) ≥ 0.95 . Here k is the maximum number of principal components

```
eigen_total_sum = np.sum(eigen_values_order)
eigen_sum = 0
k = 0
for i in range(10304):
    eigen_sum += eigen_values_order[i]
    if ( eigen_sum >= 0.95* eigen_total_sum):
        k = i + 1
        break
```

d = 10304 # since d is the number of eigen values of covariance matrix, it will be the maximum number of principal components

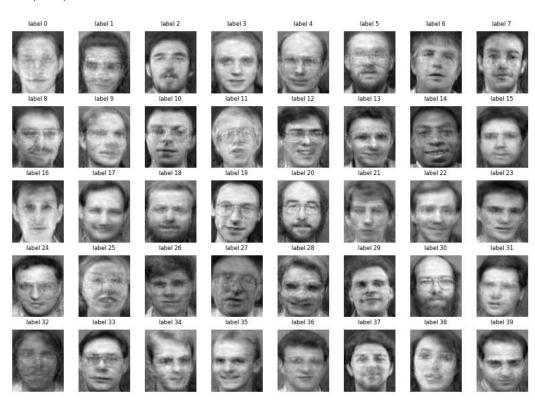
```
print(d) # total data points
print(k) # no of eigen vectors that contribute to 95 % of variance
     10304
     164
Reconstruction of data
print(centered_train_data.shape)
     (320, 10304)
z_train = (centered_train_data @ eigen_vectors_order[:,:k] )
print(z_train.shape)
z_test = (centered_test_data @ eigen_vectors_order[:,:k])
print(z_test.shape)
reconstructed\_train\_data = (z\_train \ @ \ eigen\_vectors\_order[:,:k].T) \ + \ means
print(reconstructed_train_data.shape)
reconstructed_test_data = (z_test @ eigen_vectors_order[:,:k].T) + means
print(reconstructed_test_data.shape)
     (320, 164)
     (80, 164)
     (320, 10304)
     (80, 10304)
Reconstructed training data
# create figure
fig = plt.figure(figsize=(10, 7))
# setting values to rows and column variables
rows = 5
columns = 8
for i in range(0,320,8):
  t = np.reshape(reconstructed_train_data[i],(112,92))
  # Adds a subplot at the 1st position
  \label{fig.add_subplot} \verb|fig.add_subplot(rows, columns, ((int)(i/8) + 1))| \\
  # showing image
  plt.imshow(t, cmap ='gray')
  plt.title("label " + str((int)(i/8)), size = 6)
  plt.axis('off')
          label 0
                       lahel 1
```



Reconstructed test data

```
# create figure
fig = plt.figure(figsize=(10, 7))

# setting values to rows and column variables
rows = 5
columns = 8
for i in range(0,80,2):
    t = np.reshape(reconstructed_test_data[i],(112,92))
    # Adds a subplot at the 1st position
    fig.add_subplot(rows, columns, ((int)(i/2) + 1))
    # showing image
    plt.imshow(t, cmap = 'gray')
    plt.title("label " + str((int)(i/2)), size = 6)
    plt.axis('off')
```



```
test_nearer = [np.argmin(np.linalg.norm(i - z_train, axis = 1)) for i in z_test]
test_nearer = np.array(test_nearer)
```

Does this accuracy change if you increase or decrease k?

If we decrease k, underfitting will occur, since principal components will take less than 95 % variance, there is a chance of underfitting the data If we increase k, overfitting will occur since principal components will take lot more than 95 % variance, there is a chance of overfitting the data k = 164 explains > 95 % variance, hence it is a good measure

```
correct_pred = 0

for i in range(0,80):
    if ( test_set["labels"].T[i] == train_set["labels"].T[test_nearer[i]]):
        correct_pred += 1

Accuracy = correct_pred/80
print(Accuracy*100)
        98.75

for i in range(0,80):
    plt.figure(figsize=(2,2))
    t = np.reshape(test_set['data'][i],(112,92))
    plt.subplot(1,2,1)
    plt.imshow(t, cmap = 'gray')
    plt.title("Test_original", size = 6)
    plt.axis('off')
```

```
m = np.reshape(train_set['data'][test_nearer[i]],(112,92))
plt.subplot(1,2,2)
plt.imshow(m, cmap = 'gray')
plt.title("Nearest_trained_data",size = 6)
plt.axis('off')
```

<ipython-input-124-1d625be41838>:2: RuntimeWarning: More than 20 figures have been opened. Figures crea plt.figure(figsize=(2,2))

Test_original Nearest_trained_data

















































If we decrease k, underfitting will occur, since principal components will take less than 95 % variance, there is a chance of underfitting the data If we increase k, overfitting will occur since principal components will take lot more than 95 % variance, there is a chance of overfitting the data k = 164 explains > 95 % variance, hence it is a good measure































































































































Test_original Nearest_trained_data



















































































Test_original Nearest_trained_data The same of the sa