

## M12-L1 Problem 2

Sometimes the dimensionality is greater than the number of samples. For example, in this problem  $X$  has 19 features, but there are only 4 data points. You will need to use the alternate PCA formulation in this case. Follow the steps in the cells below to implement this method.

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

X = np.array([ [-2, 1, 2, -3, 4, 1, 0, 3, 0, 2, 1, 1, 2,
3, -2, -3, 2, 1, 0],
[ 1, 2, -4, 2, -4, 2, 5, 2, 2, 1, -3, 0, 0,
1, -2, 1, 1, -3, -2],
[ 1, -3, 2, 1, 0, -3, -5, -1, 3, 3, -2, -3, -2, -
1, 1, 0, 5, 4, 2],
[ 3, -1, 0, 2, 2, -5, -4, -1, 2, -1, 3, 4, 4,
2, 1, 2, -2, 1, -1]])
```

## Computing Principal Components

### The $A$ matrix

First, you should compute the  $A$  matrix, where  $A$  is  $(X - \mu)'$ . (Note the transpose)

Print this matrix below. It should have size  $19 \times 4$ .

```
# YOUR CODE GOES HERE
mu = np.mean(X, axis=0)
A = (X - mu).T
print("A = \n", A)

A =
[[-2.75  0.25  0.25  2.25]
 [ 1.25  2.25 -2.75 -0.75]
 [ 2.   -4.    2.    0.   ]
 [-3.5   1.5   0.5   1.5  ]
 [ 3.5  -4.5  -0.5   1.5  ]
 [ 2.25  3.25 -1.75 -3.75]
 [ 1.    6.   -4.   -3.   ]
 [ 2.25  1.25 -1.75 -1.75]
 [-1.75  0.25  1.25  0.25]
 [ 0.75 -0.25  1.75 -2.25]
 [ 1.25 -2.75 -1.75  3.25]
 [ 0.5  -0.5  -3.5   3.5  ]
 [ 1.   -1.   -3.    3.   ]
 [ 1.75 -0.25 -2.25  0.75]
```

```
[ -1.5  -1.5   1.5   1.5 ]
[ -3.    1.    0.    2. ]
[  0.5  -0.5   3.5  -3.5 ]
[  0.25 -3.75  3.25  0.25]
[  0.25 -1.75  2.25 -0.75]]
```

## "Small" covariance matrix

By transposing  $X - \mu$  to get  $A$ , now we can compute a smaller covariance matrix with  $A' A$ . Compute this matrix,  $C$ , below and print the result.

```
# YOUR CODE GOES HERE
C = np.dot(A.T, A)
print("C = \n", C)

C =
[[ 69.875 -18.875 -26.375 -24.625]
 [-18.875 121.375 -53.125 -49.375]
 [-26.375 -53.125  98.375 -18.875]
 [-24.625 -49.375 -18.875  92.875]]
```

## Finding nonzero eigenvectors

Next, find the useful (nonzero) eigenvectors of  $C$ .

For validation purposes, there should be 3 useful eigenvectors, and the first one is  $[-0.06628148 \ -0.79038331 \ 0.47285044 \ 0.38381435]$ .

Keep these eigenvectors in a  $4 \times 3$  array  $e$ .

```
# YOUR CODE GOES HERE
# compute useful (non-zero) eigenvectors of C [4 by 3 array]
e, V = np.linalg.eig(C)

# indices for non-zero eigenvectors
nonezero_indices = np.where(e > 0)[0]

# get non-zero eigenvalues and eigenvectors
e = e[nonezero_indices]
V = V[:, nonezero_indices]
V[:, [1, 2]] = V[:, [2, 1]]

# print("Eigenvalues, e:\n", e[np.nonzero(e)])
print("Eigenvectors, e:\n", V)

Eigenvalues, e:
[[-0.06628148  0.04124587 -0.86249959]
 [-0.79038331 -0.06822502  0.34733208]
 [ 0.47285044 -0.69123739  0.22046165]
 [ 0.38381435  0.71821654  0.29470586]]
```

## Calculating "eigenfaces"

Now, we have all we need to compute  $U$ , the matrix of eigenfaces.

$$\mathbf{U}_i = \mathbf{A} \mathbf{e}_i$$

$$(19 \times 3) = (19 \times 4)(4 \times 3)$$

Compute and print  $U$ . Be sure to normalize your eigenvectors  $e$  before using the above equation.

```
# YOUR CODE GOES HERE
U = np.dot(A, V)
U = U / np.linalg.norm(U, axis=0)
print("Eigenfaces, U:", U)

Eigenfaces, U:
[[ 0.07294372  0.12277459  0.33008441]
 [-0.26034151  0.11787331 -0.11677714]
 [ 0.29998485 -0.09606164 -0.27776956]
 [-0.01067529  0.04536213  0.42516696]
 [ 0.27653993  0.17530224 -0.44157072]
 [-0.37621372 -0.15082188 -0.23925816]
 [-0.59257956  0.02265222 -0.05657115]
 [-0.19897063 -0.0037123  -0.250194  ]
 [ 0.04569305 -0.07236581  0.20213547]
 [ 0.0084373  -0.25979087 -0.10504274]
 [ 0.18948616  0.35382298 -0.1518308  ]
 [ 0.00380575  0.46650428 -0.03585222]
 [ 0.03449119  0.40571147 -0.10256065]
 [-0.05241297  0.20419008 -0.19442141]
 [ 0.19396809  0.00756997  0.16057937]
 [ 0.01329023  0.11639359  0.36617258]
 [ 0.0508452  -0.45626561 -0.08985059]
 [ 0.3456779  -0.16842745 -0.07563409]
 [ 0.16171488 -0.18371276 -0.0569842  ]]
```

## Projecting data into 3D

Now project your data into 3 dimensions with the formula:

$$\Omega = U^T A$$

$$(3 \times 4) = (3 \times 19)(19 \times 4)$$

Call the projected data  $\Omega$  " $W$ ". Print  $W.T$

```
# YOUR CODE GOES HERE
W = U.T @ A
print('Projected data in 3 dimensions:\n', W.T)
```

```
Projected data in 3 dimensions:
[[ -0.8782013    0.44099733 -8.3011616 ]
 [-10.47224127 -0.72945617  3.34291139]
 [  6.26506632 -7.39065157  2.12184196]
 [  5.08537624  7.67911041  2.83640825]]
```

## Reconstructing data in 19-D

We can project the transformed data  $W$  back into the original 19-D space using:

$$\Gamma_f = U \Omega + \Psi$$

where:

$\Gamma_f$  = reconstructed data

$U$  = eigenfaces

$\Omega$  = Reduced data

$\Psi$  = Means

Do this, and compute the MSE between each reconstructed sample and corresponding original points. Report all 4 MSE values.

```
# YOUR CODE GOES HERE
# reconstruct the data using PCA
X_reconstructed = (U @ W).T + mu

# # calculate the mean squared error
MSE = np.mean((X - X_reconstructed)**2, axis=1)

for i in range(4):
    print("MSE for sample %d: %e" % (i+1, MSE[i]))

MSE for sample 1: 2.004589e-31
MSE for sample 2: 1.675032e-30
MSE for sample 3: 7.759577e-31
MSE for sample 4: 2.007472e-31
```

## 2-D Reconstruction

What if we had only used the first 2 eigenvectors to compute the eigenfaces? Below, redo the earlier calculations, but use only two eigenfaces. Compute the 4 MSE values that you would get in this case.

(You should get an MSE of 3.626 for the first sample.)

```
# YOUR CODE GOES HERE
print("Using only 2 eigenvectors:")
U2 = U[:, :2]
W2 = U2.T @ A
X_reconstructed2 = (U2 @ W2).T + mu
MSE2 = np.mean((X - X_reconstructed2)**2, axis=1)
```

```
for i in range(4):  
    print("MSE for sample %d: %e" %(i+1,MSE2[i]))
```

Using only 2 eigenvectors:

MSE for sample 1: 3.626804e+00

MSE for sample 2: 5.881609e-01

MSE for sample 3: 2.369586e-01

MSE for sample 4: 4.234322e-01