

# M12-L2-P1

December 4, 2023

## 1 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
from sklearn.metrics import mean_squared_error

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]])
```

### 1.1 Computing Principal Components

#### 1.1.1 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  $19 \times 4$ .

```
[ ]: A = (X - np.mean(X,axis=0)).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]]
```

### 1.1.2 “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.

```
[ ]: C = 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]]
```

### 1.1.3 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  $e$ .

```
[ ]: w,v = np.linalg.eig(C)
      w,v = np.real(w), np.real(v)
      idx = np.argsort(-w)
      w,v = w[idx], v[:,idx]
      e = v[:,0:3]
      print(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]]
```

### 1.1.4 Calculating “eigenfaces”

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

$$U_i = A e_i$$

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

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

```
[ ]: U = A @ e
      U /= np.linalg.norm(U, axis=0)
      print("Eigenfaces, U:\n",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  ]]
```

## 1.2 Projecting data into 3D

Now project your data into 3 dimensions with the formula:

$$W = U^T A$$

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

Call the projected data  $\Omega$  “W”. Print W.T

```
[ ]: 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]]
```

## 1.3 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:

\$ \_f = \$ reconstructed data

\$U = \$ eigenfaces

\$ = \$ Reduced data

\$ = \$ Means

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

```
[ ]: MSE = []
reconstructed = (U@W) + np.mean(X,axis=1)
mse = np.mean((X - reconstructed.T) **2, axis=1)
for i in range(4):
    print(f"MSE for sample {i}: {mse[i]}")
```

MSE for sample 0: 0.7468836565096959

MSE for sample 1: 0.716412742382272

MSE for sample 2: 0.7164127423822705

MSE for sample 3: 0.691481994459834

## 1.4 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.)

```
[ ]: MSE2 = []
U2 = A @ e[:, :2]
U2 /= np.linalg.norm(U2, axis=0)
W2 = U2.T@A
reconstructed = (U2@W2) + np.mean(X,axis=1)
print("Using only 2 eigenvectors:")
mse = np.mean((X - reconstructed.T) ** 2, axis=1)
for i in range(4):
    print(f"MSE for sample {i}: {mse[i]}")
```

Using only 2 eigenvectors:

MSE for sample 0: 3.882784888090766

MSE for sample 1: 1.3575824909878655

MSE for sample 2: 0.9870175957576055

MSE for sample 3: 1.260330232133847