# CS540 Spring 2022 Homework 3

### 1 Assignment Goals

- Explore Principal Component Analysis (PCA) and the related Python packages (numpy, scipy, and matplotlib)
- Make pretty pictures :)

# 2 Summary

In this project, you'll be implementing a facial analysis program using PCA, using the skills from **the linear** algebra + PCA lecture. You'll also continue to build your Python skills.

We'll walk you through the process step-by-step (at a high level).

# 3 Packages Needed for this Project

You'll use Python 3. In this project, you'll need the following packages (these instructions apply to scipy >= 1.5.0):

```
>>> from scipy.linalg import eigh
>>> import numpy as np
>>> import matplotlib.pyplot as plt
```

#### 4 Dataset

You will be using part of the Yale face dataset (processed). You can find the dataset in the same zip archive you got this PDF from.

The dataset contains 2414 sample images, each of size  $32 \times 32$ . We will use n to refer to the number of images (so n = 2414) and d to refer to the number of features for each sample image (so  $d = 1024 = 32 \times 32$ ). We will test your code only using the provided data set. Note, we'll use  $x_i$  to refer to the ith sample image which would be a d-dimensional feature vector.

# 5 Program Specification

Implement these six Python functions to do PCA on our provided dataset in a file called pca.py:

- 1. load\_and\_center\_dataset(filename): load the dataset from a provided .npy file, re-center it around the origin, and return it as a numpy array of floats.
- 2. get\_covariance(dataset): calculate and return the covariance matrix of the dataset as a numpy matrix  $(d \times d \text{ array})$ .
- 3. get\_eig(S, m): perform eigendecomposition on the covariance matrix S and return a diagonal matrix (numpy array) with the largest m eigenvalues on the diagonal in descending order, and a matrix (numpy array) with the corresponding eigenvectors as columns.

- 4. get\_eig\_prop(S, prop): similar to get\_eig, but instead of returning the first m, return all eigenvalues and corresponding eigenvectors in a similar format that explain more than a prop proportion of the variance (specifically, please make sure the eigenvalues are returned in descending order).
- 5. project\_image(image, U): project each  $d \times 1$  image into your m-dimensional subspace (spanned by m vectors of size  $d \times 1$ ) and return the new representation as a  $d \times 1$  number array.
- 6. display\_image(orig, proj): use matplotlib to display a visual representation of the original image and the projected image side-by-side.

#### 5.1 Load and Center the Dataset ([20] points)

You'll want to use the the numpy function load() to load the YaleB\_32x32.npy file into Python (You may need to install numpy first).

```
>>> x = np.load(filename)
```

This should give you a  $n \times d$  dataset (recall that n = 2414 is the number of images in the dataset and d = 1024 is the number of dimensions of each image). In other words, each row represents an image feature vector.

Your next step is to center this dataset around the origin. Recall the purpose of this step from lecture — it is a technical condition that makes it easier to perform PCA, but it does not lose any important information.

To center the dataset is simply to subtract the mean  $\mu_x$  from each data point  $x_i$  (image, in our case), i.e.,  $x_i^{\text{cent}} = x_i - u_x$ , where

$$\mu_x = \frac{1}{n} \sum_{i=1}^n x_i.$$

You can take advantage of the fact that  $\mathbf{x}$  (as defined above) is a **numpy** array and, as such, has this convenient behavior:

After you've implemented this function, it should work like this:

```
>>> x = load_and_center_dataset('YaleB_32x32.npy')
>>> len(x)
2414
>>> len(x[0])
1024
>>> np.average(x)
-8.315174931741023e-17
```

(Its center isn't *exactly* zero, but taking into account precision errors over 2414 arrays of 1024 floats, it's what we call "close enough.")

From now on, we will use  $x_i$  to refer to  $x_i^{\text{cent}}$ .

### 5.2 Find the Covariance Matrix ([15] points)

Recall, from lecture, that one of the interpretations of PCA is that it is the eigendecomposition of the sample covariance matrix. We will rely on this interpretation in this assignment, with all of the information you need below.

The covariance matrix is defined as

$$S = \frac{1}{n-1} \sum_{i=1}^{n} x_i x_i^{\mathsf{T}}.$$

Note that  $x_i$  is one of the *n* images in the (centered) dataset and is considered to be a column vector of size  $d \times 1$ .

To calculate S, you'll need a couple of tools from numpy again:

The result of this function for our sample dataset should be a  $d \times d$  (or  $1024 \times 1024$ ) matrix.

#### 5.3 Get the m Largest Eigenvalues and their Eigenvectors ([17] points)

Again, recall from lecture that eigenvalues and eigenvectors are useful objects that characterize matrices. Better yet, PCA can be performed by doing an eigendecomposition and taking the eigenvectors corresponding to the largest eigenvalues. This replaces the recursive deflation step we discussed in class.

You may find scipy.linalg.eigh from the scipy library very helpful when writing this function. The optional parameter called subset\_by\_value might be of particular use.

Return the *largest* m eigenvalues of S as a diagonal matrix, in descending order, and the corresponding eigenvectors as columns in a matrix.

To return more than one thing from a function in Python, you can do this:

```
def multi_return():
    return "a string", 5
my_string, my_int = multi_return()
```

Make sure to return the diagonal matrix of eigenvalues *first*, then the eigenvectors in corresponding columns. You may have to rearrange the output of **eigh** to get the eigenvalues in decreasing order and *make* sure to keep the eigenvectors in the corresponding columns after that rearrangement.

# 5.4 Get all Eigenvalues/Eigenvectors that Explain More than a Certain Proportion of the Variance ([8] points)

We want all the eigenvalues that explain more than a certain proportion of the variance.

Let  $\lambda_i$  be an eigenvalue of the covariance matrix S. Then the proportion of variance explained is

$$\frac{\lambda_i}{\sum_{j=1}^n \lambda_j}.$$

Return the eigenvalues as a diagonal matrix, in descending order, and the corresponding eigenvectors as columns in a matrix. Hint: **subset\_by\_value** was useful for the previous function, so perhaps something similar could come in handy here. What is the trace of a matrix?

Again, make sure to return the diagonal matrix of eigenvalues *first*, then the eigenvectors in corresponding columns. You may have to rearrange the output of **eigh** to get the eigenvalues in decreasing order and *make sure to keep the eigenvectors in the corresponding columns* after that rearrangement.

## 5.5 Project the Images ([15] points)

Given one of the images from your dataset and the results of your get\_eig (or get\_eig\_prop) function, create and return the *projection* of that image.

Let  $u_j$  represent the jth column of U. In other words, each  $u_j$  is an eigenvector of S and has size  $d \times 1$ . If U has m columns, then for any image  $x_i$ , we project it into the m dimensional subspace as  $x_i^{\text{pro}} = \sum_{j=1}^m \alpha_{ij} u_j$ , where  $\alpha_{ij} = u_j^{\mathsf{T}} x_i$ . Note that  $\alpha_i \in \mathbb{R}^m$  and  $x_i^{\text{pro}} \in \mathbb{R}^d$ .

Find the  $\alpha_{ij}$ s for your image, then use them together with the eigenvectors to create your projection.

```
>>> projection = project_image(x[0], U)
>>> print(projection)
[6.84122225 4.83901287 1.41736694 ... 8.75796534 7.45916035 5.4548656 ]
```

### 5.6 Visualize ([25] points)

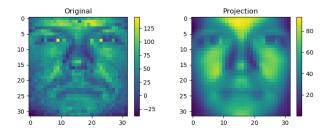
We'll be using matplotlib's imshow. First, make sure you have matplotlib installed. Follow these steps to visualize your images:

- 1. Reshape the images to be  $32 \times 32$  (before this, they were being thought of as 1 dimensional vectors in  $\mathbb{R}^{1024}$ ).
- 2. Create a figure with one row of two subplots.
- 3. Title the first subplot (the one on the left) as "Original" (without the quotes) and the second (the one on the right) as "Projection" (also without the quotes).
- 4. Use imshow with the optional argument aspect='equal'

- 5. Use the return value of **imshow** to create a **colorbar** for each image.
- 6. Render your plots!

Below is a simple snippet of code for you to test your functions. Do **not** include it in your submission!

```
>>> x = load_and_center_dataset('YaleB_32x32.npy')
>>> S = get_covariance(x)
>>> Lambda, U = get_eig(S, 2)
>>> projection = project_image(x[0], U)
>>> display_image(x[0], projection)
```



#### 6 Submission Notes

Please submit your files in a .zip archive named hw3\_<netid>.zip, where you replace <netid> with your netID (i.e., your wisc.edu login). Inside your zip file, there should be only one file named hw3.py. Do not submit a Jupyter notebook .ipynb file.

Be sure to **remove all debugging output** before submission; failure to do so will be **penalized** ([10] **points**):

- Your functions should run silently (except for the image rendering window in the last function).
- No code should be put outside the function definitions (except for import statements; helper functions are allowed).

ALL THE BEST!