Lab 5: Eigen-decomposition of Images

Course Title: Image Processing I (Spring 2022)

Course Number: ECE 63700

Instructor: Prof. Charles A. Bouman

Author: Zhankun Luo

Lab 5: Eigen-decomposition of Images

- 2. Multivariate Gaussian Distributions and Whitening
 - 2.1. Exercise: Generating Gaussian random vectors
 - 2.2. Exercise: Covariance Estimation and Whitening
- 4. Eigenimages, PCA, and Data Reduction
 - 4.1. Exercise
 - 4.1.1. the figure with the first 12 eigen-images
 - 4.1.2. plots of projection coef. vs. eigenvector number for first 4 images
 - 4.1.3. the 6 synthesized images for 1st image
- 5. Image Classification
 - 5.1. Exercise: Classification and PCA
 - 5.1.1. table for each mis-classified input image when $B_k=R_k$
 - 5.1.2. table for each mis-classified input image when $B_k = \Lambda_k \equiv \mathrm{diag}(R_k)$
 - 5.1.3. table for each mis-classified input image when $B_k = R_{wc} \equiv \frac{1}{K} \sum_k R_k$
 - 5.1.4. table for each mis-classified input image when $B_k = \Lambda \equiv {
 m diag}(R_{wc})$
 - 5.1.5. table for each mis-classified input image when $B_k = I$
 - 5.1.6. which of the above classifiers worked the best
 - 5.1.7. trade-off between accuracies of the data model and the estimates

Appendix

```
Python codes for functions
```

Python codes for solutions

```
solution to section 2.1: soln 2 1.py
```

solution to section 2.2: soln 2 2.py

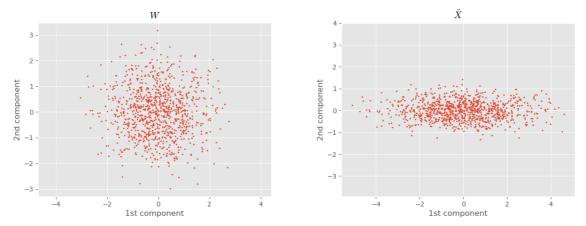
solution to section 4: soln 4.py

solution to section 5: soln 5.py

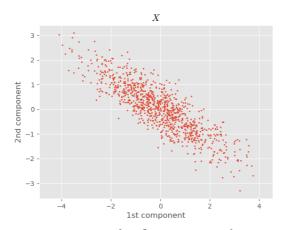
2. Multivariate Gaussian Distributions and Whitening

2.1. Exercise: Generating Gaussian random vectors solution

the scatter plots for W, \tilde{X} , and X



Scatter Plots for **W**(left), and Scaled Random Vectors (right)

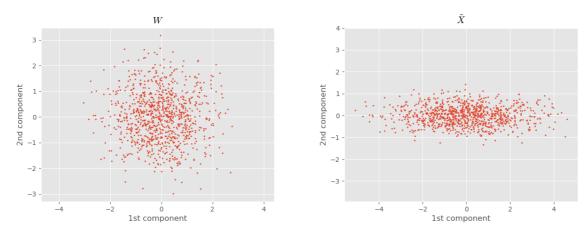


Scatter Plot for Generated X

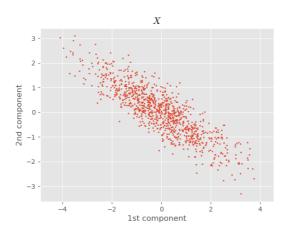
2.2. Exercise: Covariance Estimation and Whitening

solution

the scatter plots for W, \tilde{X} , and X



Scatter Plots for Whitened Random Vector **W**(left), and Uncorrelated Random Vectors (right)

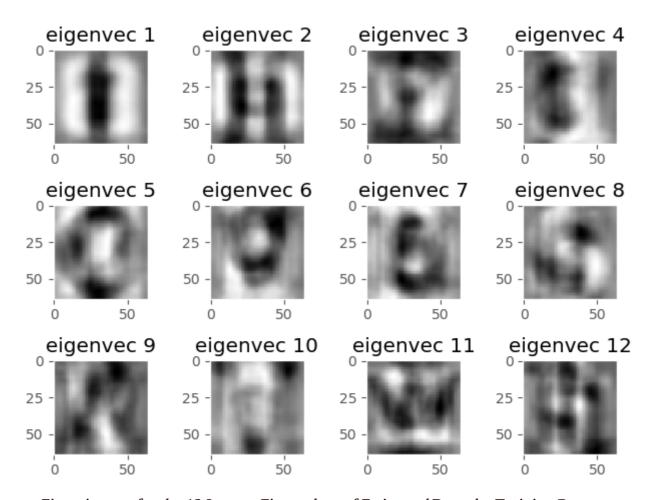


Scatter Plot for Original Gaussian Random Vector \mathbf{X}

4. Eigenimages, PCA, and Data Reduction

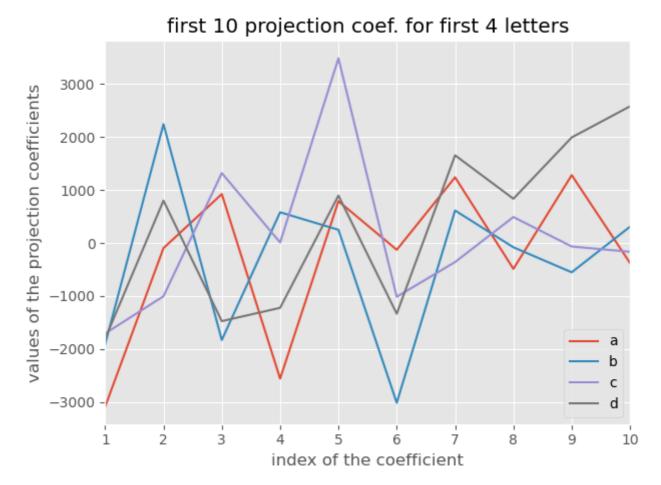
4.1. Exercise

4.1.1. the figure with the first 12 eigen-images



Eigen-images for the 12 Largest Eigenvalues of Estimated **R** on the Training Dataset

4.1.2. plots of projection coef. vs. eigenvector number for first 4 images

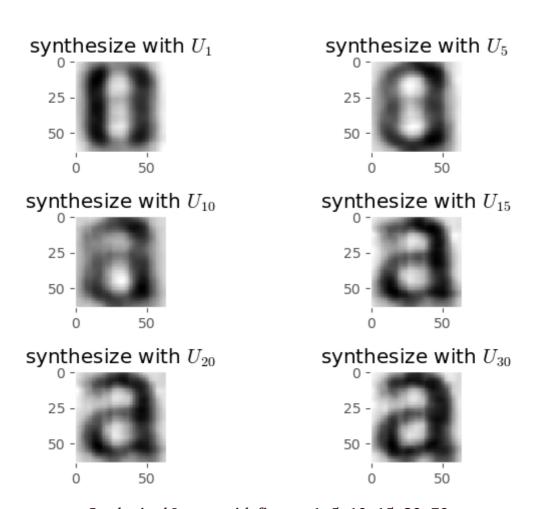


First 10 Projection Coefficients for First 4 Images in the Training Dataset

4.1.3. the 6 synthesized images for 1st image



Original Image for 1st Image in the Training Dataset



Synthesized Images with first m=1, 5, 10, 15, 20, 30 Eigenvectors for 1st Image in the Training Dataset

5. Image Classification

5.1. Exercise: Classification and PCA

5.1.1. table for each mis-classified input image when $B_k=R_k\,$

INPUT CHARACTER	OUTPUT FROM THE CLASSIFIER
d	a
j	у
1	i
n	V
p	e
q	a
u	a
У	V

5.1.2. table for each mis-classified input image when $B_k = \Lambda_k \equiv \mathrm{diag}(R_k)$

INPUT CHARACTER	OUTPUT FROM THE CLASSIFIER
i	1
у	V

5.1.3. table for each mis-classified input image when $B_k = R_{wc} \equiv \frac{1}{K} \sum_k R_k$

$$B_k = R_{wc} \equiv rac{1}{K} \sum_k R_k$$

INPUT CHARACTER	OUTPUT FROM THE CLASSIFIER
g	q
у	V

5.1.4. table for each mis-classified input image when $B_k = \Lambda \equiv { m diag}(R_{wc})$

INPUT CHARACTER	OUTPUT FROM THE CLASSIFIER
f	t
у	V

5.1.5. table for each mis-classified input image when $B_{\it k}=I$

INPUT CHARACTER	OUTPUT FROM THE CLASSIFIER
f	t
g	q
у	V

5.1.6. which of the above classifiers worked the best

when
$$B_k=\Lambda_k\equiv {
m diag}(R_k)$$
, , $B_k=R_{wc}\equiv rac{1}{K}\sum_k R_k$ or $B_k=\Lambda\equiv {
m diag}(R_{wc})$

the error rate is lowest $rac{2}{26} imes 100\%$, the accuracy reaches $rac{24}{26} imes 100\% pprox 92.3\%$

5.1.7. trade-off between accuracies of the data model and the estimates

Even thought the classifier for $B_k = R_k$ has the best estimate for covariances, it has the worst accuracy for the classifier.

Even
$$R_k=I$$
 is better than it, but $B_k=\Lambda_k\equiv {
m diag}(R_k)$, $B_k=R_{wc}\equiv \frac{1}{K}\sum_k R_k$ or $B_k=\Lambda\equiv {
m diag}(R_{wc})$ have the best accuracy.

We should balance the data parameter estimation and the model complexity.

Appendix

Python codes for functions

utils.py

```
from typing import Tuple, List
from string import ascii lowercase as alphabet
import os
from os.path import join
from math import sqrt, floor
from numpy import ndarray, zeros, array
from PIL import Image
import matplotlib.pyplot as plt
import matplotlib
matplotlib.rcParams['mathtext.fontset'] = 'cm'
plt.style.use('ggplot')
def plot data2d(data: ndarray, str title: str) -> None:
    plt.scatter(data[0], data[1], s=3)
    plt.grid(color='w')
    plt.xlabel("1st component")
    plt.ylabel("2nd component")
    plt.title(str title)
    plt.axis("equal")
def read data(dir data: str, height: int=64, width: int=64) ->
Tuple[ndarray, List[str]]:
    , folders, = list(os.walk(dir_data))[0]
    folders.sort()
    data = zeros((len(folders), len(alphabet), height, width))
    for i, folder in enumerate(folders):
        for j, char in enumerate(alphabet):
            data[i][j] = array(Image.open(join(dir data, folder,
char+'.tif')))
    return data, folders
def get divisor(x: int) -> Tuple[int, int]:
    t = floor(sqrt(x) + 1e - 6)
    while x%t != 0:
```

```
t -= 1
    return t, x//t
def display samples(data: ndarray, folders: List[str], char: str) ->
    index = ord(char) - ord(alphabet[0])
    rows, cols = get divisor(len(folders))
    fig, axs = plt.subplots(rows, cols)
    axs = [axs] if rows == 1 else axs
    axs = [[e] for e in axs] if cols == 1 else axs
    for i, folder in enumerate (folders):
        row, col = i//cols, i%cols
        axs[row][col].imshow(data[i][index],cmap=plt.cm.gray,
interpolation='none')
        axs[row][col].set title(folder)
        axs[row][col].grid(False)
    plt.tight layout()
def display scaled (eigenvecs: ndarray, height: int=64, width: int=64)
-> None:
    assert(height*width == eigenvecs.shape[0])
    num eigen = eigenvecs.shape[-1]
   rows, cols = get divisor(num eigen)
   fig, axs = plt.subplots(rows, cols)
    axs = [axs] if rows == 1 else axs
    axs = [[e] for e in axs] if cols == 1 else axs
    for i, vec in enumerate (eigenvecs.T):
        row, col = i//cols, i%cols
        v max, v min = vec.max(), vec.min()
        vec = (vec - v min) / (v max - v min)
        axs[row][col].imshow(vec.reshape((height, width)),
cmap=plt.cm.gray, interpolation='none')
        axs[row][col].set_title("eigenvec {}".format(i+1))
        axs[row][col].grid(False)
    plt.tight_layout()
def plot projection(Y: ndarray, range coef: int=10) -> None:
    num char = Y.shape[-1]
    for char, projection in list(zip(alphabet[:num char], Y.T)):
        plt.plot(range(1, range coef+1), projection[:range coef],
label=char)
    plt.legend()
    plt.xlim([1, range coef])
```

```
plt.title("first {} projection coef. for first {}
letters".format(range_coef, num char))
    plt.xlabel("index of the coefficient")
   plt.ylabel("values of the projection coefficients")
   plt.tight layout()
def display synthesized (vec img: ndarray, mu hat: ndarray,
    U: ndarray, list num eigen: list,
   height: int=64, width: int=64) -> None:
   assert(height*width == vec img.shape[0])
    cols, rows = get divisor(len(list num eigen))
   fig, axs = plt.subplots(rows, cols)
   axs = [axs] if rows == 1 else axs
   axs = [[e] for e in axs] if cols == 1 else axs
   for i, num eigen in enumerate(list num eigen):
        U top = U[:, :num eigen]
       row, col = i//cols, i%cols
        Y = U top.conj().T @ (vec img - mu hat)
       vec img syn = U top @ Y + mu hat
        axs[row][col].imshow(vec img syn.reshape((height, width)),
cmap=plt.cm.gray, interpolation='none')
        axs[row][col].set title(r"synthesize with
$U {{{}}}$".format(num eigen))
        axs[row][col].grid(False)
   plt.tight layout()
if name == " main ":
   print(len(alphabet))
    data train, folders train = read data("resource/training data")
   print(data train.shape)
    data_test, folders_test = read_data("resource/test_data")
   print(data test.shape)
   display samples(data train, folders train, char='a')
   plt.show()
    display_samples(data_test, folders_test, char='a')
    plt.show()
```

eigen_decompose.py

```
from typing import Union, Tuple
from string import ascii lowercase as alphabet
from numpy import sqrt, diag, ndarray, reshape, empty, zeros, log
from numpy.linalg import eig, svd, det, inv
from numpy.random import seed, randn
def generate data(R: ndarray, mu: Union[None, ndarray]=None,
    N: int=1000, se=0) -> ndarray:
    """Generate N data samples for a Guassian distribution cluster
    V[:,i] is the eigenvector corresponding to the eigenvalue D[i]
    if W \sim N(0, I), X = sqrt(D) W
       => X \sim N(0, D),
    if X \sim N(0, D), Y = V X
        => Y \sim N(0, V D V^T)
    thus Y = V \text{ sqrt}(D) W
        => Y \sim N(0, R)
        with decomposition: R = V D V^T
    11 11 11
    seed(se)
    dimension = R.shape[0]
    D, V = eig(R)
    if mu is None:
        return V @ diag(sqrt(D)) @ randn(dimension, N)
    return V @ diag(sqrt(D)) @ randn(dimension, N) + mu
def estimate param(X: ndarray) -> Tuple[ndarray, ndarray]:
    N, mu = X.shape[-1], X.mean(axis=-1, keepdims=True)
    R = ((X - mu) @ (X - mu).T) / (N-1)
    return R, mu
def decorrelate_data(X: ndarray, R: ndarray, mu: Union[None,
ndarray] = None) -> ndarray:
    if mu is not None:
        X -= mu
    D, V = eig(R)
    return V.T @ X
def whiten data(X: ndarray, R: ndarray, mu: Union[None,
ndarray] = None) -> ndarray:
    if mu is not None:
```

```
X -= mu
    D, V = eig(R)
    return diag(1./sqrt(D)) @ (V.T @ X)
def svd data(X: ndarray) -> Tuple[ndarray, list, ndarray]:
    """X = U \operatorname{diag}(D) V^H = U \operatorname{@np.diag}(D) \operatorname{@Vh} = (U * D) \operatorname{@Vh}
    note: The rows of Vh are the eigenvectors of A^H A
        the columns of U are the eigenvectors of A A^H
        Vector(s) with the singular values sorted in descending order
    If full matrices=True (default),
        U and Vh have the shapes (..., M, M) and (..., N, N)
    Otherwise, full matrices=False,
        the shapes are (..., M, K) and (..., K, N)
        where K = min(M, N)
    11 11 11
    U, D, Vh = svd(X, full matrices=False)
    return U, D, Vh.conj().T
class Classifier PCA:
    def init (self, num eigen: int=10):
        self.num folder = None
        self.alphabet = alphabet
        self.num char = None
        self.num eigen = num eigen
        self.height = None
        self.width = None
        self.mean global = None
        self.A = None
        self.params = None
        self.R wc = None
    def fit(self, data train: ndarray):
        self.num folder, self.num char, \
             self.height, self.width = data_train.shape
        X = reshape(data_train, (self.num_folder*self.num_char,
self.height*self.width)).T
        self.mean global = X.mean(axis=-1, keepdims=True)
        U, D, V = svd data((X - self.mean global) /
sqrt(self.num folder*self.num char-1))
        self.A = U[:, :self.num_eigen].conj().T
        self.params = [None] * self.num char
        self.R_wc = zeros((self.num_eigen, self.num eigen))
```

```
for ind, char in enumerate(self.alphabet[:self.num char]):
        x = data train[:, ind].reshape(self.num folder, -1).T
        y = self.A @ (x - self.mean global)
        R, mu = estimate param(y)
        self.params[ind] = {"mean": mu, "cov": R,
                            "invcov": inv(R),
                            "logdetcov": log(det(R))}
        self.R wc += R
    self.R wc /= self.num char
    self.invR wc = inv(self.R wc)
def predict(self, data test: ndarray,
            option: Union[str, None]=None):
    shape data = list(data test.shape)
    X = reshape(data test, (-1, self.height*self.width)).T
    Y = self.A @ (X - self.mean global)
    pred = empty((Y.shape[-1],), dtype='str')
    pred[:] = ' '
    if option is None or option == "default":
        f = lambda y, i: \
            (y-self.params[i]["mean"]).T \
            @ self.params[i]["invcov"] \
            @ (y-self.params[i]["mean"]) \
            + self.params[i]["logdetcov"]
    elif option == "diag":
        f = lambda y, i: \
            (y-self.params[i]["mean"]).T \
            @ inv(diag(diag(self.params[i]["cov"]))) \
            @ (y-self.params[i]["mean"]) \
            + log(det(diag(diag(self.params[i]["cov"]))))
    elif option == "global":
        f = lambda y, i: \
            (y-self.params[i]["mean"]).T \
            @ self.invR wc \
            @ (y-self.params[i]["mean"])
    elif option == "global diag":
        f = lambda y, i: \
            (y-self.params[i]["mean"]).T \
            @ inv(diag(diag(self.R wc))) \
            @ (y-self.params[i]["mean"])
    elif option == "identity":
        f = lambda y, i: \
            (y-self.params[i]["mean"]).T \
```

```
@ (y-self.params[i]["mean"])
else:
    print("error")
for ind, y in enumerate(Y.T):
    y = y.reshape((-1, 1))
    id_pred = min(range(self.num_char), key=lambda i: f(y,
i))
    pred[ind] = self.alphabet[id_pred]
    return pred.reshape(shape_data[:-2])
```

Python codes for solutions

solution to section 2.1: soln_2_1.py

```
import sys
from os.path import dirname
sys.path.insert(0, dirname(dirname(__file__)))
from numpy import array, diag, sqrt
from numpy.linalg import eig
from numpy.random import seed, randn
import matplotlib.pyplot as plt
from src.utils import plot data2d
if _ name__ == "__main__":
    N = 1000
    R = array([[2, -1.2]],
                [-1.2, 1]]
    dimension = R.shape[0]
    D, V = eig(R) # eigenvalues, eigenvectors
    W = randn(dimension, N)
    X tilde = diag(sqrt(D)) @ W
    X = V @ X tilde
    for data, str title, char \
        in list(zip([W, X_tilde, X],
                [r'$W$', r'$\tilde{X}$', r'$X$'],
                ['a', 'b', 'c'])):
        plot_data2d(data, str title)
        plt.savefig('result/fig_2_1' + char + '.png', Bbox='tight')
        plt.show()
```

solution to section 2.2: soln 2 2.py

```
import sys
from os.path import dirname
sys.path.insert(0, dirname(dirname( file )))
from numpy import array
from src.eigen decompose import generate data, estimate param, \
    decorrelate data, whiten data
from src.utils import plot data2d
import matplotlib.pyplot as plt
if name == " main ":
   N = 1000
   R = array([[2, -1.2],
                [-1.2, 1])
   X = generate data(R, N=N, se=0)
   R X hat, mu X hat = estimate param(X)
   print("R X\n", R X hat)
   print("mu X\n", mu X hat)
   X tilde = decorrelate data(X, R)
   W = whiten data(X, R)
   R W hat, mu W hat = estimate param(W)
   print("R W\n", R W hat)
   print("mu W\n", mu W hat)
    for data, str title, char \
        in list(zip([W, X_tilde, X],
                [r'$W$', r'$\tilde{X}$', r'$X$'],
                ['a', 'b', 'c'])):
        plot data2d(data, str title)
        plt.savefig('result/fig 2 2' + char + '.png', Bbox='tight')
        plt.show()
```

solution to section 4: soln 4.py

```
import sys
from os.path import dirname
sys.path.insert(0, dirname(dirname( file )))
from src.eigen decompose import svd data
from src.utils import read data, display scaled, plot projection,
display synthesized
from numpy import reshape, sqrt
import matplotlib.pyplot as plt
if name == " main ":
    data train, folders train = read data("resource/training data")
   num folder, num alphabet, height, width = data train.shape
    X = reshape(data train, (num folder*num alphabet,
height*width)).T
    mu hat = X.mean(axis=-1, keepdims=True)
    U, D, V = svd data((X - mu hat) / sqrt(num folder*num alphabet-
1))
   num top = 12
   U top = U[:, :num top]
    display scaled(U top)
    plt.savefig('result/fig 4 la.png', Bbox='tight')
   plt.show()
    num char = 4
   Y concat = U.conj().T @ (X[:, :num char] - mu hat)
    plot projection(Y concat)
    plt.savefig('result/fig 4 1b.png', Bbox='tight')
    plt.show()
    vec img = X[:, 0].reshape((-1, 1))
    list num eigen = [1, 5, 10, 15, 20, 30]
    display synthesized (vec img, mu hat, U, list num eigen)
    plt.savefig('result/fig_4_1c.png', Bbox='tight')
    plt.show()
```

solution to section 5: soln 5.py