HW2 Solution, CSE 569 Fall 2018

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Question Attempting: Q (II)

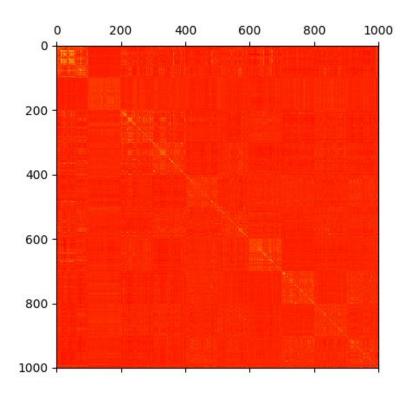
Q(II)

(a) Perform PCA and reduce the dimensions of training images from 28 × 28 = 784 to 100. Output the percentage of variance retained in the data after dimensionality reduction. Plot the covariance matrix for the training data after dimensionality reduction and analyze. Plot one example of each of the digits after dimensionality reduction and analyze.

Solution (a):

Percentage Variance Retained after dimensionality reduction is **84.76**% (with Standardized Scaling of the training images) and **92.84** % (without Standardized scaling). The plots below depict results with Standardized Scaling.

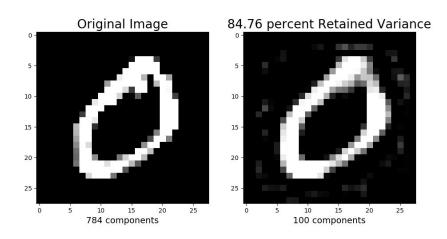
Plot of the covariance matrix for the training data after dimensionality reduction:

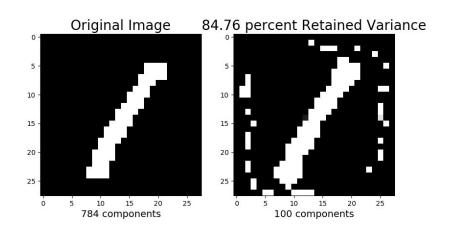


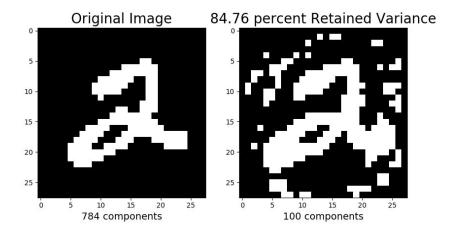
The covariance matrix gives us the relation between points present in a space, and how they change with respect to the change in the other. In this case, the variables are the 1000 training images with reduced dimensions(100), and the matrix tells us how they vary with each other. Here, the covariance matrix is a diagonal matrix, which tells us that that the images do not vary with each other, but only with itself.

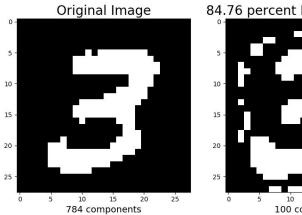
Plots of each digit after dimensionality reduction:

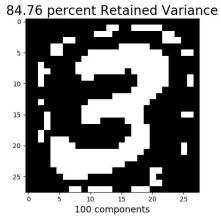
After reducing the principal components from 784 to 100, we observe that 84.76 % of the variance is retained. In the reduced dimensional space, we observe that there is loss because of the removal of principal components which correspond to dimensions with low variation.

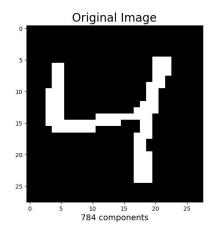


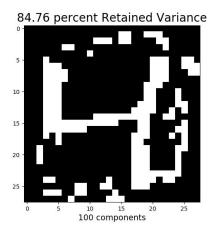


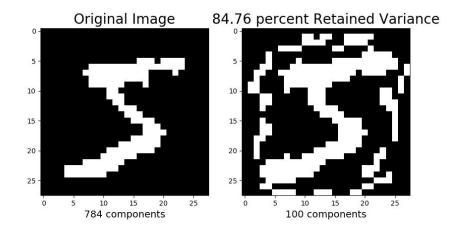


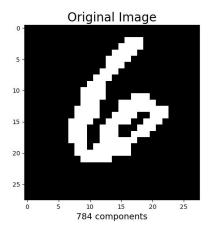


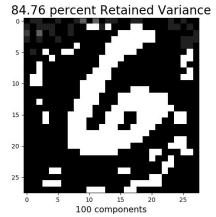


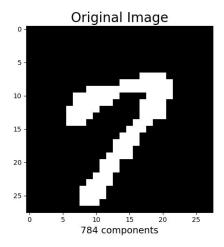


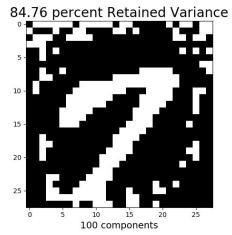


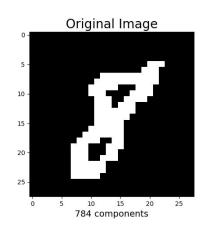


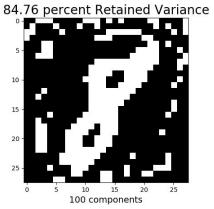


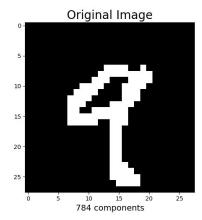


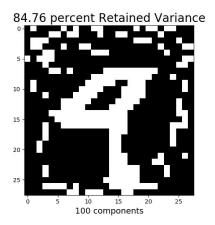








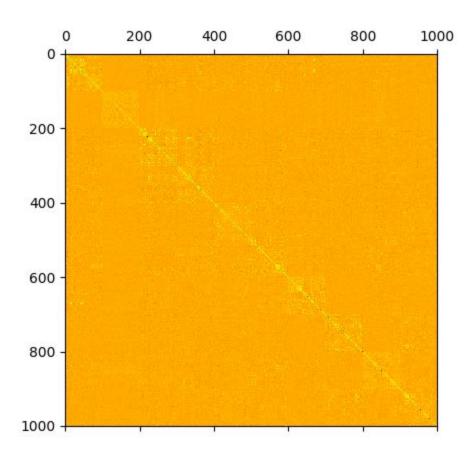




b) Perform PCA whitening on the reduced dimensions. Plot the covariance matrix after whitening and analyze. Plot one example of each of the digits after PCA whitening and analyze. Implement ZCA whitening and plot the covariance matrix. Plot one example of each of the digits after ZCA whitening and analyze.

Solution (b):

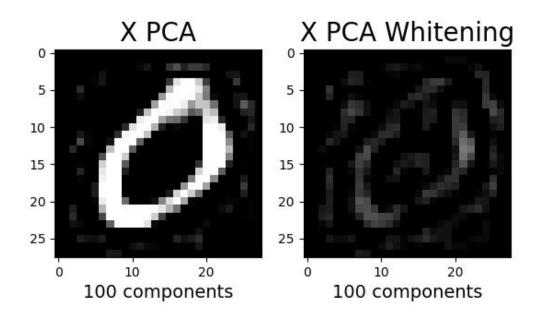
Covariance Matrix Plot after performing PCA Whitening:

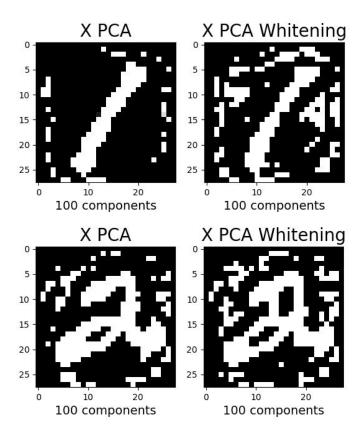


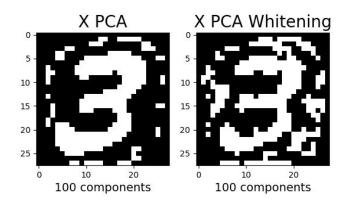
Similar to the covariance after using PCA alone, PCA with whitening also results in a diagonal matrix, which explains the same reason of variation in data with each other. Here, we are whitening with regularization to avoid numerical instability. Due to this, some diagonal values will be lesser than 1, which would have been 1 for all diagonal values in case of $PCA_{Whitening}$ without regularization. In the plot above, we observe a bright yellow diagonal, with some points of different colors and intensities. These points illustrate the diagonal values lesser than 1 due to regularization.

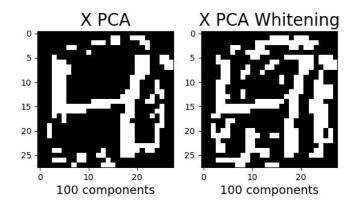
Plots of each digit after PCA Whitening:

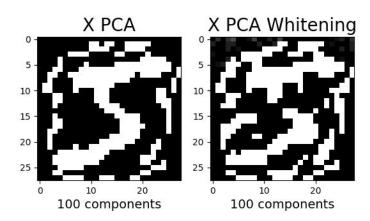
With the aid of PCA Whitening, we aim to remove redundancies of the raw data, in which the adjacent pixels are highly correlated. Thus, it makes the input less redundant. The following plots illustrate the effect of PCA whitening with regularization.

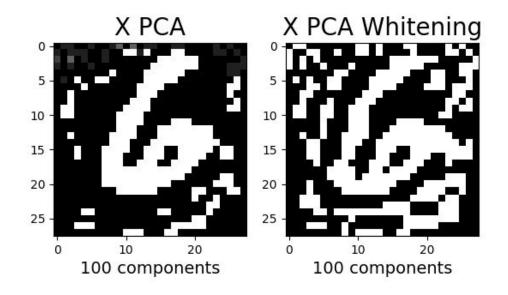


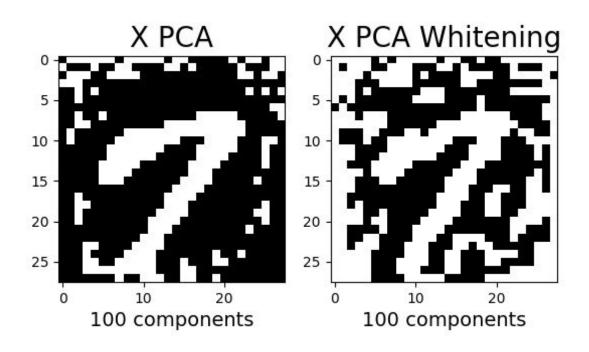


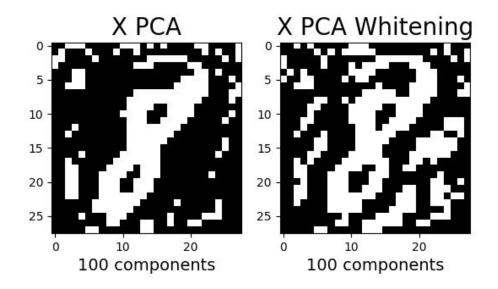


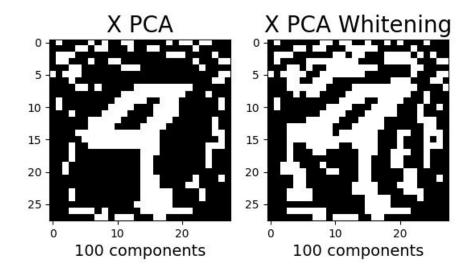






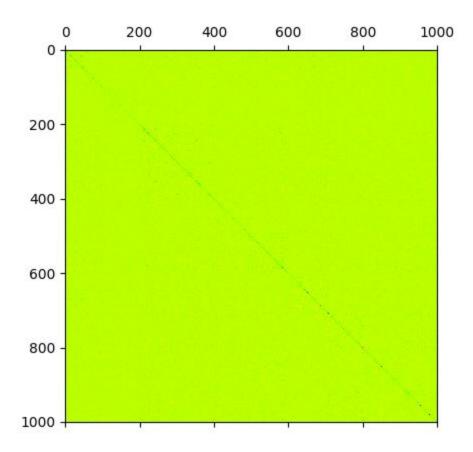






ZCA Whitening:

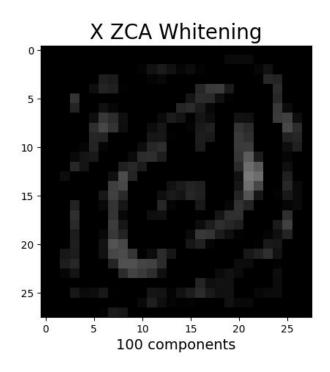
• Covariance Matrix Plot:

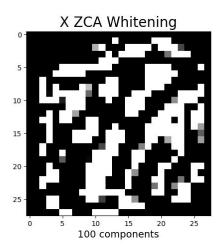


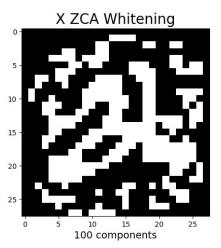
We again get a similar diagonal matrix, which illustrates the lack of variation in data with respect to each other other than itself. With ZCA whitening, we attempt to project the PCA-whitened data back in the original space.

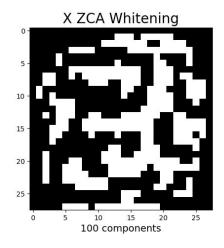
• Plots of each digit after ZCA Whitening:

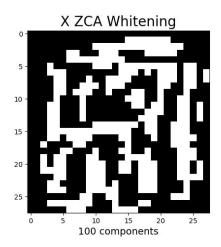
With ZCA Whitening, we attempt to project the reduced-dimensional PCA-whitened data back in the original space. It is a rotation. We achieve it by taking the product of the $X_{PCAWhitened}$ with the eigen vector E. If we compare the ZCA-whitened results with the original data, we observe more enhanced edges in the results with ZCA-whitening.

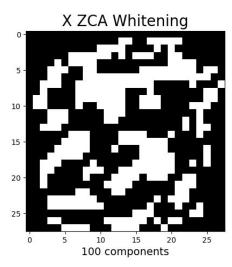


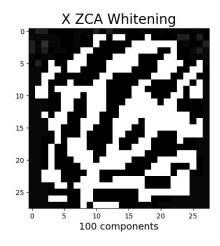


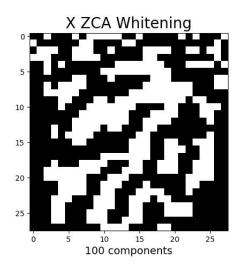


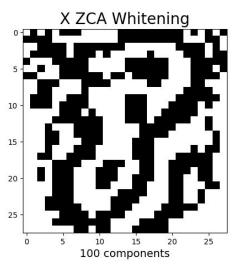


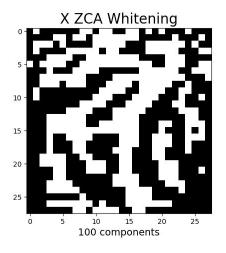












• Code Snippet for Part (a) and (b):

```
import numpy as np
import os
import pdb
import matplotlib.pyplot as plt
import cv2
import math
from load_mnist_updated_40ct2018 import mnist
from load_mnist_updated_40ct2018 import one_hot
from sklearn.decomposition import PCA
from sklearn.preprocessing import StandardScaler
datasets_dir = '/home/kunal/Desktop/CSE569_HW2/data/'
def calculate_percentVarianceRetained(pca1,pca2):
      return (sum(pca1.explained_variance_)/sum(pca2.explained_variance_))*100;
def main():
   trX, trY, tsX, tsY = mnist(noTrSamples=1000,
                               noTsSamples=100, digit_range=[0, 10],
                               noTrPerClass=100, noTsPerClass=10)
   trX=trX.T
    tsX=tsX.T
    pca=PCA(n_components=100)
    #pca object having 784 components
    pca2=PCA(n components=784)
    scalar= StandardScaler()
    trX=scalar.fit_transform(trX)
    eTx=pca.fit_transform(trX)
    #Transforming reduced data back to its original space
    #for plotting and comparison with original
    apX=pca.inverse transform(eTx)
```

```
ld2=pca2.fit_transform(trX)
    #Transforming reduced data back to its original space
    apX2=pca2.inverse_transform(ld2)
    #calculating retained % variance
    percVretained=calculate_percentVarianceRetained(pca,pca2);
    percVretained=round(percVretained,2)
    print percVretained
    cov=getCovariance(eTx)
    #Performing PCA Whitening
   Xpca_w=perform_PCA_Whitening(pca,eTx,cov)
    #for plotting and comparison with original
    apX3=pca.inverse_transform(Xpca_w)
    cov2=getCovariance(Xpca w)
    #Performing ZCA Whitening
   Xzca_w=perform_ZCA_Whitening(pca,Xpca_w,cov)
    cov3=getCovariance(Xzca w)
    displayImgs(apX3,Xzca_w,percVretained)
def perform_ZCA_Whitening(pca,Xpca_w,cov):
       eigen_vectors=pca.components_
       return np.dot(Xpca_w,eigen_vectors)
def perform_PCA_Whitening(pca,eTx,cov):
      eigen_vectors=pca.components_
      eigen_values=pca.explained_variance_
      x=np.dot(eTx,eigen_vectors)
      for i in range(0,100):
             if(eigen_values[i]==0):
                   eigen_vectors[i]=eigen_vectors[i]/math.sqrt(eigen_values[i]+1e-3);
                   eigen_vectors[i]=eigen_vectors[i]/math.sqrt(eigen_values[i]);
      return np.dot(x,eigen_vectors.T)
```

```
def getCovariance(apX):
      covarianceMat=np.cov(apX)
      plt.matshow(covarianceMat*255,cmap=plt.cm.hsv) # hsv, RdBu
      plt.show()
def displayImgs(trX,apX,percVretained):
      i=0
      while i < 999:
          plt.subplot(1, 2, 1);
          trX=trX*255;
          plt.imshow(trX[i].reshape(28,28),cmap = plt.cm.gray,
             interpolation='nearest',
                           clim=(0, 255));
          plt.xlabel('100 components', fontsize = 14)
          plt.title('X PCA Whitening', fontsize = 20);
          apX=apX*255;
          plt.subplot(1, 2, 2);
          plt.imshow(apX[i].reshape(28, 28),cmap = plt.cm.gray,
             interpolation='nearest',
                           clim=(0, 255));
          plt.xlabel('100 components', fontsize = 14)
          plt.title('X ZCA Whitening', fontsize = 20);
          plt.show()
          i+=100
if __name__ == "__main__":
    main()
```

c) Implement Fisher's Linear discriminant for digits 0 and 1 in the training dataset. Use reduced PCA dimensions (100). Report the accuracy on the test dataset having only digits 0 and 1.

Solution (c):

Fitting Fisher's Linear Discriminant with 200 images of 0's and 1's in total, and testing them on 20 images of 0s and 1s from a testing set of 100 images, we get an accuracy of **100 percent** on the test dataset having only digits 0 and 1.

Code Snippet for this Problem: On the next Page

```
import numpy as np
import os
import pdb
import matplotlib.pyplot as plt
import cv2
import math
from load_mnist_updated_40ct2018 import mnist
from load_mnist_updated_40ct2018 import one_hot
from sklearn.decomposition import PCA
from sklearn.preprocessing import StandardScaler
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
import matplotlib as mpl
from matplotlib import colors
datasets_dir = '/home/kunal/Desktop/CSE569_HW2/data/'
def main():
    trX, trY, tsX, tsY = mnist(noTrSamples=1000,
                               noTsSamples=100, digit_range=[0, 10],
                               noTrPerClass=100, noTsPerClass=10)
    trX=trX.T
    tsX=tsX.T
    tsY=tsY.T
    trX_01=trX[0:200]
    trY_01=trY[0][0:200]
    trX_01=np.array(trX_01)
    trY_01=np.array(trY_01)
    tsX_01=[]
    tsY 01=[]
    pca=PCA(n_components=100)
    scalar= StandardScaler()
    trX_01=scalar.fit_transform(trX_01)
    tsX=scalar.fit_transform(tsX)
```

```
eTx=pca.fit_transform(trX_01)
   # Linear Discriminant space
   fLD= LinearDiscriminantAnalysis()
   fLD.fit(eTx,trY_01)
   #reducing test set to 100 components
   reduced_tsX=pca.fit_transform(tsX)
   for i in range(0,100):
      if tsY[i]==0 or tsY[i]==1:
             tsX_01.append(reduced_tsX[i])
             tsY_01.append(tsY[i])
   tsX_01=np.array(tsX_01)
   tsY_01=np.array(tsY_01)
   results=fLD.predict(tsX_01)
   print results
   print fLD.score(tsX_01,tsY_01)*100
if __name__ == "__main__":
   main()
```