## q3-Copy1

November 6, 2018

## 1 Principal Component Analysis

The goal of this question is to build a conceptual understanding of dimensionality reduction using PCA and implement it on a toy dataset. Only have to use numpy and matplotlib for this question.

```
In [1]: import numpy as np
        import matplotlib
        import matplotlib.pyplot as plt
In [2]: # (a) Load data (features)
        def load_data():
            data = np.load('q3-data/features.npy')
            labels = np.load('q3-data/labels.npy')
            mean = np.mean(data, axis=0)
            mean = np.array(mean, ndmin=2)
            data = np.subtract(data, mean)
            mean2 = np.mean(data, axis=0)
            stdvn = np.std(data, axis=0)
            stdvn = np.array(stdvn, ndmin=2)
            print(stdvn.shape, stdvn)
            data = data/stdvn
            print(np.std(data, axis=0))
            return data, labels
        datafile, labels = load_data()
(1, 8) [[0.82530129 0.43214658 1.75852918 0.76061262 1.29681839 1.18787014
  1.93816328 1.22044697]]
[1. 1. 1. 1. 1. 1. 1. 1.]
In [3]: labels = labels.astype(int)
        print(labels.shape, labels.dtype)
```

```
(150,) int 64
```

```
In [4]: # (b) Perform eigen decomposition and return eigen pairs in desecending order of eigen
        def eigendecomp(X):
            # covariance matrix:
            \# cov_mat = (X - np.mean(X, axis=0)).T.dot((X - np.mean(X, axis=0))) / (X.shape[0])
            cov mat = np.cov(X.T)
            print(cov_mat.shape, type(cov_mat), cov_mat)
            # Eigendecomposition on coveriance matrix, cov_mat:
            eig_vals, eig_vecs = np.linalg.eig(cov_mat)
            print('Eigenvectors \n%s' %eig_vecs)
            print('\nEigenvalues \n%s' %eig_vals)
            # softing eigenvalues in decreasing order and corresponding eigenvectors:
            idx = eig_vals.argsort()[::-1]
            sorted_eig_vals = eig_vals[idx]
            sorted_eig_vecs = eig_vecs[:,idx]
            return (sorted_eig_vals, sorted_eig_vecs)
        sorted_evalues, sorted_evectors = eigendecomp(datafile)
        print(sorted_evalues)
        print(sorted_evectors)
(8, 8) <class 'numpy.ndarray'> [[ 1.00671141 -0.11010327 0.87760486 0.82344326 0.61123001 -
   0.776564
               0.48190166]
 [-0.11010327 1.00671141 -0.42333835 -0.358937
                                                     0.07459068 0.52416995
 -0.28731011 -0.06998508]
  \begin{bmatrix} 0.87760486 & -0.42333835 & 1.00671141 & 0.96921855 & 0.4704054 & -0.25019762 \end{bmatrix} 
   0.85445501 0.51042104]
                         0.96921855 1.00671141 0.44634059 -0.21541033
 [ 0.82344326 -0.358937
   0.8273983 0.54484244]
 [ 0.61123001  0.07459068  0.4704054
                                        0.44634059 1.00671141 0.70151359
   0.84012249 0.93454714]
  \begin{bmatrix} -0.07220205 & 0.52416995 & -0.25019762 & -0.21541033 & 0.70151359 & 1.00671141 \end{bmatrix} 
   0.2731165 0.659988 ]
 [ 0.776564   -0.28731011   0.85445501   0.8273983   0.84012249   0.2731165
   1.00671141 0.88321462]
```

```
[ 0.48190166 -0.06998508 \ 0.51042104 \ 0.54484244 \ 0.93454714 \ 0.659988 
   0.88321462 1.00671141]]
Eigenvectors
[[-0.39124937 0.13884872 -0.46160937 0.58034539 0.24934936 0.21747713
  -0.38816186 0.1118572 ]
 [ 0.11687696 -0.4391715 -0.78711289 -0.2905579 -0.12725786  0.02806659
   0.21689908 0.15922802]
  \begin{bmatrix} -0.40655289 & 0.29080021 & -0.13961871 & -0.12636707 & -0.54994554 & -0.61159334 \end{bmatrix} 
  -0.17223991 0.13591601]
 [-0.39944906 \quad 0.26454833 \quad -0.16206048 \quad -0.54404218 \quad 0.49904279 \quad 0.01470075
   0.05047577 -0.44213114]
  \begin{bmatrix} -0.3778555 & -0.35426671 & 0.07790627 & 0.42060984 & 0.12822569 & -0.34172774 \\ \end{bmatrix} 
   0.6099293 -0.17576426]
 [-0.09816172 -0.64299795 \ 0.11941452 -0.04972667 -0.0795516 \ -0.07714852
  -0.59620498 -0.43768069]
 [-0.45509399 -0.03231459 0.12200908 -0.08034689 -0.51935676 0.67406772
   0.18983425 -0.14979985]
  \begin{bmatrix} -0.38587285 & -0.30545597 & 0.29393481 & -0.28457653 & 0.27864817 & -0.02358821 \end{bmatrix} 
  -0.08099129 0.70942501]]
Eigenvalues
[ 4.74298961e+00 2.29585309e+00 7.76910512e-01 2.04172901e-01
  3.37651661e-02 -8.56766184e-16 -1.16776912e-16 5.26888208e-16
[ 4.74298961e+00 2.29585309e+00 7.76910512e-01 2.04172901e-01
  3.37651661e-02 5.26888208e-16 -1.16776912e-16 -8.56766184e-16]
 \begin{bmatrix} [-0.39124937 & 0.13884872 & -0.46160937 & 0.58034539 & 0.24934936 & 0.1118572 \end{bmatrix} 
  -0.38816186 0.21747713]
  \begin{bmatrix} 0.11687696 & -0.4391715 & -0.78711289 & -0.2905579 & -0.12725786 & 0.15922802 \end{bmatrix} 
   0.21689908 0.02806659]
  \begin{bmatrix} -0.40655289 & 0.29080021 & -0.13961871 & -0.12636707 & -0.54994554 & 0.13591601 \end{bmatrix} 
  -0.17223991 -0.61159334]
  \begin{bmatrix} -0.39944906 & 0.26454833 & -0.16206048 & -0.54404218 & 0.49904279 & -0.44213114 \end{bmatrix} 
   0.05047577 0.01470075]
 \begin{bmatrix} -0.3778555 & -0.35426671 & 0.07790627 & 0.42060984 & 0.12822569 & -0.17576426 \end{bmatrix}
   0.6099293 -0.34172774]
 [-0.09816172 -0.64299795 \ 0.11941452 -0.04972667 -0.0795516 \ -0.43768069
  -0.59620498 -0.07714852]
 [-0.45509399 -0.03231459 \ 0.12200908 -0.08034689 -0.51935676 -0.14979985
   0.18983425 0.67406772]
 [-0.38587285 - 0.30545597 \ 0.29393481 - 0.28457653 \ 0.27864817 \ 0.70942501
  -0.08099129 -0.02358821]]
```

After sorting the eigenpairs, the next question is "how many principal components are we going to choose for our new feature subspace?" A useful measure is the so-called "explained variance," which can be calculated from the eigenvalues.

• The explained variance tells us how much information (variance) can be attributed to each of the principal components.

```
tot = sum(sorted_evalues)
           var_exp = [(i / tot)*100 for i in sorted(sorted_evalues, reverse=True)]
           cum_var_exp = np.cumsum(var_exp)
           var_exp = np.array(var_exp, ndmin=2)
           cum_var_exp = np.array(cum_var_exp, ndmin=2)
           sorted_evalues = np.array(sorted_evalues, ndmin=2)
           np.set_printoptions(suppress=True)
           eval_var = np.concatenate((np.array(sorted_evalues.T, ndmin=2), np.array(var_exp.T
           cum_eval_var = np.concatenate((np.array(eval_var, ndmin=2), np.array(cum_var_exp.T
           print("[EigenValue
                                               Cumulative_Variance]")
                                  Variance
           print(cum_eval_var)
           return var_exp, cum_var_exp
       var, cum_var = eval(sorted_evalues, sorted_evectors)
[EigenValue
               Variance
                            Cumulative_Variance]
[[ 4.74298961 58.89212098 58.89212098]
[ 2.29585309 28.50684249 87.39896347]
[ 0.77691051  9.64663886  97.04560233]
[ 0.2041729  2.53514686  99.58074919]
[ 0.03376517 0.41925081 100.
                    100.
Γ 0.
                0.
                                       1
[ -0.
               -0.
                           100.
                                       ]
[ -0.
               -0.
                           100.
                                       11
```

I would pick k = 3 for this problem, as it covers 97% of the variance. Remaining of the values are quite insignificant and thus can be left out without much loss of information.

## In []:

The construction of the projection matrix that will be used to transform the Iris data onto the new feature subspace. Although, the name "projection matrix" has a nice ring to it, it is basically just a matrix of our concatenated top k eigenvectors.

 Here, we are reducing the 8-dimensional feature space to a 2-dimensional feature subspace, by choosing the "top 2" eigenvectors with the highest eigenvalues to construct our -dimensional eigenvector matrix.

```
Y = datafile.dot(matrix_w)

Matrix W:

[[-0.39124937  0.13884872]

[ 0.11687696 -0.4391715 ]

[-0.40655289  0.29080021]

[-0.39944906  0.26454833]

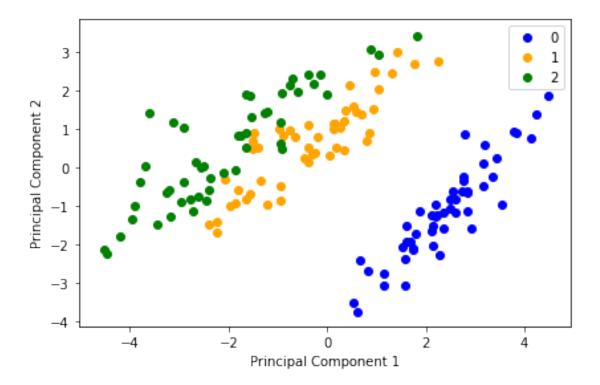
[-0.3778555  -0.35426671]

[-0.09816172 -0.64299795]

[-0.45509399  -0.03231459]

[-0.38587285  -0.30545597]]
```

In this last step we will use the 8Œ2-dimensional projection matrix W to transform our samples onto the new subspace via the equation Y=XŒW, where Y is a 150Œ2 matrix of our transformed samples.



```
eval(sorted_evalues, sorted_evectors)
           viz(Y, labels, datafile)
       if __name__ == "__main__":
           main()
[EigenValue
               Variance
                            Cumulative_Variance]
[[ 4.74298961
               58.89212098 58.89212098]
[ 2.29585309
               28.50684249 87.39896347]
[ 0.77691051
                9.64663886 97.04560233]
[ 0.2041729
                2.53514686 99.58074919]
[ 0.03376517
                0.41925081 100.
[ 0.
                0.
                            100.
                                       ]
                            100.
                                        ]
[ -0.
               -0.
[ -0.
               -0.
                            100.
                                       ]]
```

In [8]: def main():