

roozara_hw3

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- ECE 657A: Data and Knowledge Modelling and Analysis
- Winter 2019
- WATIAM:roozara ID: 20801583
- Homework 3:Eigenvector Decomposition

Reference used :<https://hadrienj.github.io/posts/Preprocessing-for-deep-learning/>
<https://archive.ics.uci.edu/ml/machine-learning-databases/communities/>

```
In [1]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
#for eigen value
from numpy import cov
from numpy import linalg as LA
#pd.options.display.max_columns = None
pd.options.display.max_rows = None

def readFromFile(name):
    with open(name) as f:
        features= [line.split(' ')[1] for line in f.readlines()
                    if line.startswith('@attr')]
    return features
```

1 Importing the crime dataset and storing in a matrix

The crime dataset was loaded into a matrix ,its observed that the data is already normalized but so many missing values. About 1675 out of 1993 missing values each in columns 101 to 126 were replaced by the mean of those features. the first 5 columns including two (county & community of which contained about 1174 missing values each were not included in creating the matrix.These first five non predictive attributes were left out of the analysis. 128 scaled to 123 features being analysed.

```

In [2]: #importing the communities dataset into variable cdata
        cdata = pd.read_csv('data/communities.data',sep= ',', header = None, na_values=["?"])

        #print(Acdata)
        #examining the data to correct for missing valieue

        #IDENTIFYING COLUMNS WITH null value and filling with the mean
        #nan_col=pd.DataFrame(cdata.isnull().sum(axis=0))
        #nan_col.index = list(range(128))

        cdata.iloc[:,4:] = cdata.iloc[:,4:].apply(lambda x: x.fillna(x.mean()),axis=0)
        cdata.columns = readFromFile('data/communities.names')
        print('Crime dataset dimension', cdata.shape)
        cdata.head(5)

```

Crime dataset dimension (1994, 128)

```

Out[2]:

```

	state	county	community	communityname	fold	population	\
0	8	NaN	NaN	Lakewoodcity	1	0.19	
1	53	NaN	NaN	Tukwilacity	1	0.00	
2	24	NaN	NaN	Aberdeentown	1	0.00	
3	34	5.0	81440.0	Willingborotownship	1	0.04	
4	42	95.0	6096.0	Bethlehemtownship	1	0.01	

	householdsize	racepctblack	racePctWhite	racePctAsian	\
0	0.33	0.02	0.90	0.12	
1	0.16	0.12	0.74	0.45	
2	0.42	0.49	0.56	0.17	
3	0.77	1.00	0.08	0.12	
4	0.55	0.02	0.95	0.09	

	...	LandArea	PopDens	PctUsePubTrans	PolicCars	\
0	...	0.12	0.26	0.20	0.060000	
1	...	0.02	0.12	0.45	0.163103	
2	...	0.01	0.21	0.02	0.163103	
3	...	0.02	0.39	0.28	0.163103	
4	...	0.04	0.09	0.02	0.163103	

	PolicOperBudg	LemasPctPolicOnPatr	LemasGangUnitDeploy	\
0	0.040000	0.900000	0.500000	
1	0.076708	0.698589	0.440439	
2	0.076708	0.698589	0.440439	
3	0.076708	0.698589	0.440439	
4	0.076708	0.698589	0.440439	

	LemasPctOfficDrugUn	PolicBudgPerPop	ViolentCrimesPerPop
0	0.32	0.140000	0.20
1	0.00	0.195078	0.67
2	0.00	0.195078	0.43
3	0.00	0.195078	0.12
4	0.00	0.195078	0.03

[5 rows x 128 columns]

```
In [3]: #matrix creating with numpy
        cdata_matrix = np.matrix(cdata.iloc[:,5:])
        print('Matrix created' , '(dimension' , cdata_matrix.shape, ')', '\n', cdata_matrix)
```

Matrix created (dimension (1994, 123))

```
[[0.19      0.33      0.02      ... 0.32      0.14      0.2       ]
 [0.       0.16      0.12      ... 0.       0.19507837 0.67      ]
 [0.       0.42      0.49      ... 0.       0.19507837 0.43      ]
 ...
 [0.16      0.37      0.25      ... 0.91      0.28      0.23      ]
 [0.08      0.51      0.06      ... 0.22      0.18      0.19      ]
 [0.2       0.78      0.14      ... 1.       0.13      0.48      ]]
```

2 Compute the eigenvectors and eigenvalue and Reporting the top 20 eigenvalues

We compute the eigenvectors and thus eigen value by first calculating the covaraince matrix. We project any data onto the principal subspace that is spanned by the eigenvectors that belong to the largest eigenvalues.

```
In [4]: cov_matrix = np.cov(cdata_matrix, rowvar=False, bias=True)
        eigenvalues, eigenVector = LA.eig(cov_matrix)
        x = np.arange(1, 124,1)
        eig_valTable = pd.DataFrame(eigenvalues, index = x, columns = ['Eigenvalues'])
        eig_valTable.sort_values(by='Eigenvalues', ascending=False, inplace=True)
        eig_valTable.head(20)
```

```
Out[4]:      Eigenvalues
1      1.091241
2      0.760521
3      0.333826
4      0.287335
5      0.187330
6      0.161874
7      0.133454
8      0.111856
9      0.088807
10     0.076775
```

11	0.057316
12	0.055735
13	0.053821
14	0.048017
15	0.045117
16	0.039349
17	0.033284
18	0.029946
19	0.028341
20	0.027238

As it can be seen from the first plot (left plot) below, it is hard to have a clear cut off since the curve is more shallow. The first 20 eigenvalues count for ~85% of the variance. The 95% were calculated below and it turned out that we need approximately 39 eigenvalues to process 95% of the data.

```
In [5]: i = 1
        j = 0
        y = np.zeros(shape=(123,1))

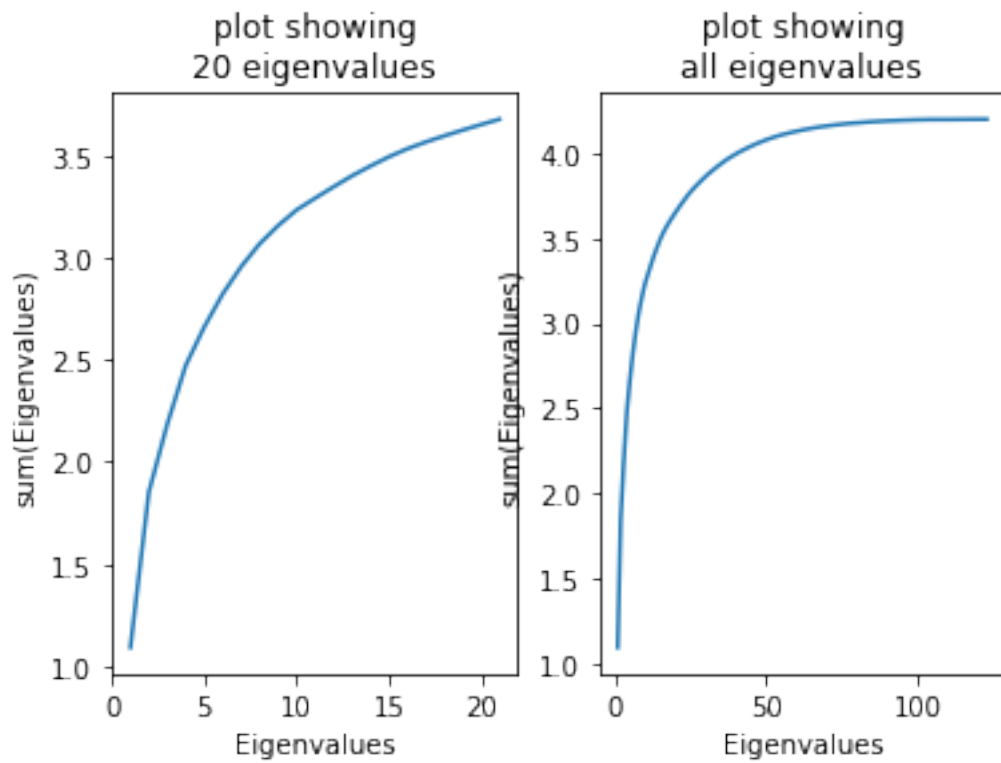
        for index, row in eig_valTable.iterrows():
            y[j] = eig_valTable['Eigenvalues'][i] + y[j-1]
            i += 1
            j += 1

        sum_eigen = pd.DataFrame(y, index = x)
        sum_eigen['Eigenvalue No'] = x

        f, (ax1, ax2) = plt.subplots(1, 2)
        ax1.plot(x[:21], y[:21])
        ax1.set_title('plot showing\n20 eigenvalues')
        ax1.set_xlabel('Eigenvalues')
        ax1.set_ylabel('sum(Eigenvalues)')

        ax2.plot(x, y)
        ax2.set_title(' plot showing \nall eigenvalues')
        ax2.set_xlabel('Eigenvalues')
        ax2.set_ylabel('sum(Eigenvalues)')

Out[5]: Text(0, 0.5, 'sum(Eigenvalues)')
```



```
In [6]: print(sum_eigen[(sum_eigen[0] > 3.98) & (sum_eigen[0] < 3.992268253580597)])
```

```
0 Eigenvalue No
39 3.986718 39
```