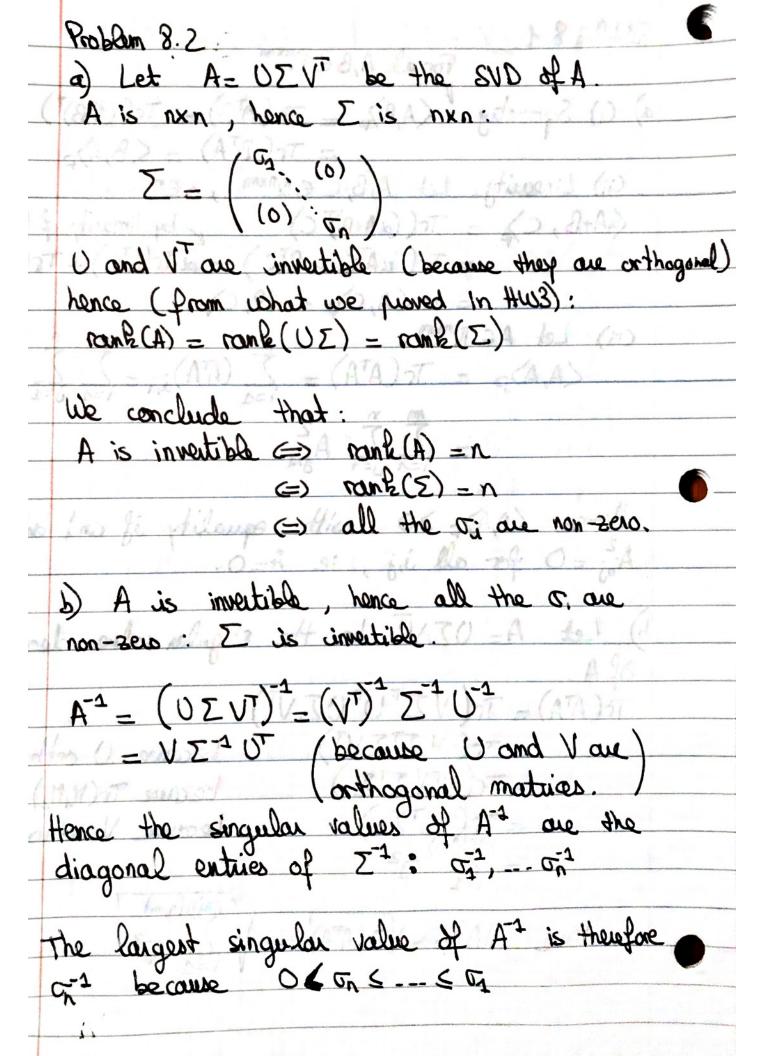
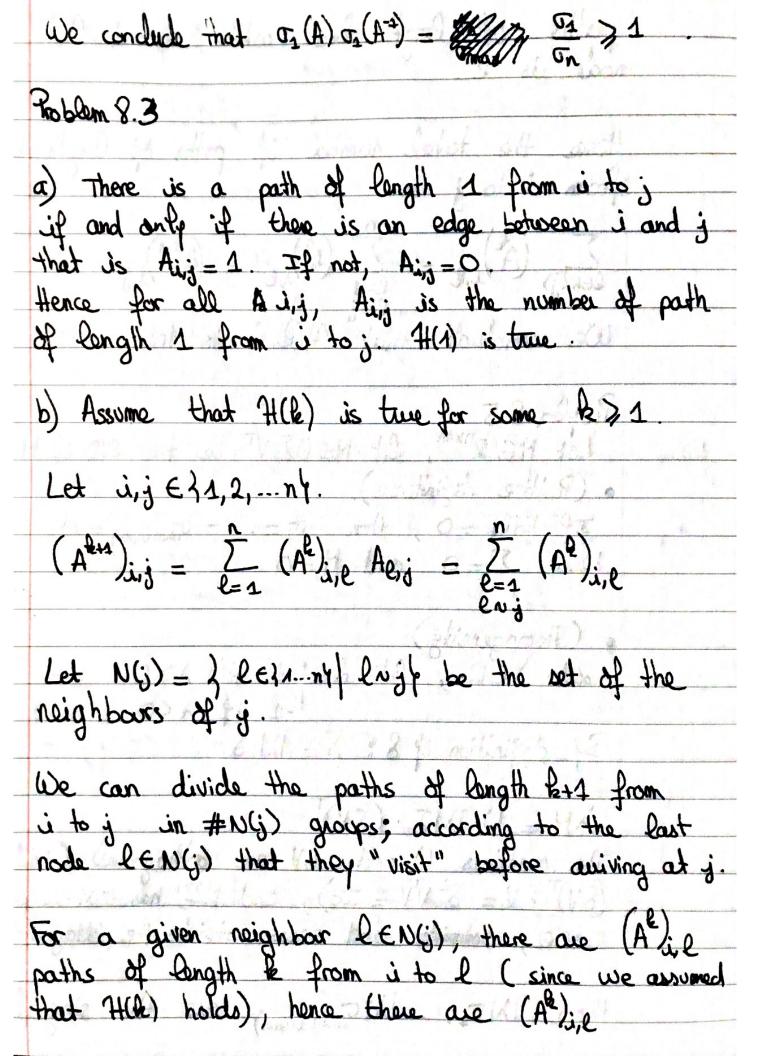
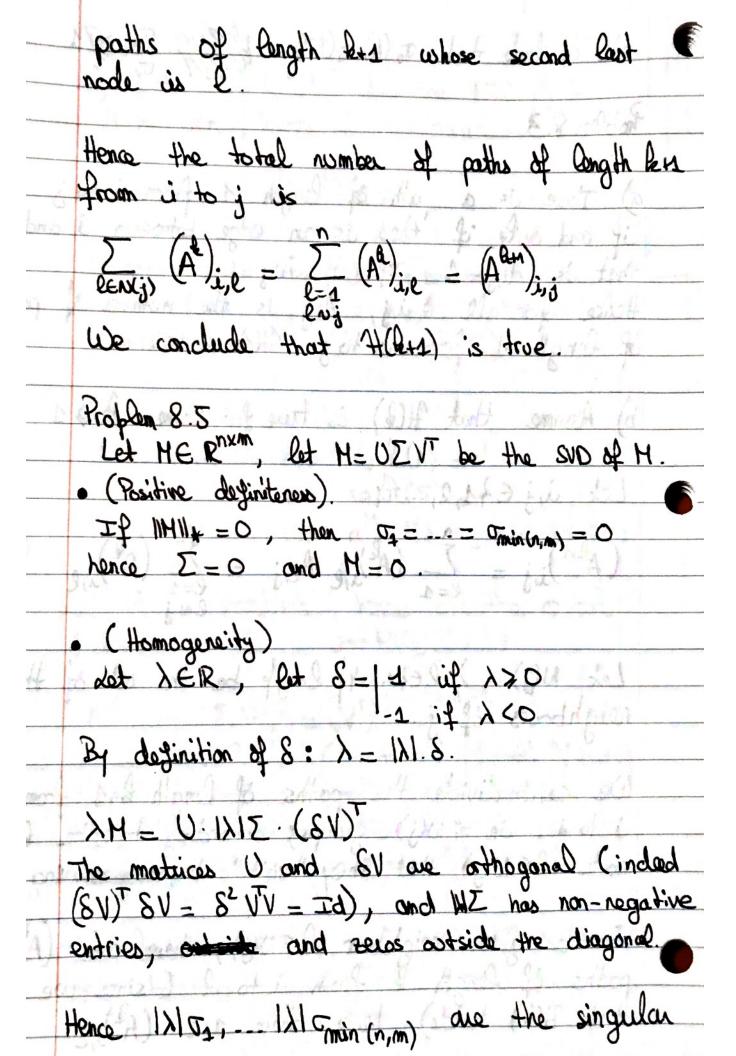
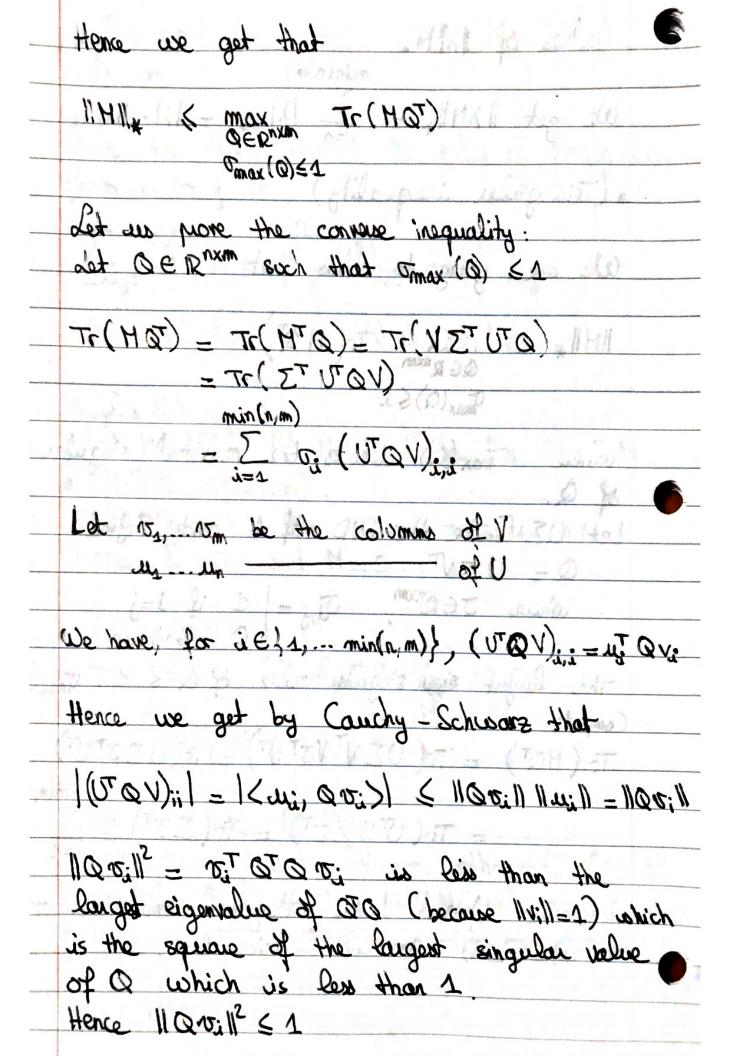
Robben 8.1 For all A, B ∈ Rnxm a) (i) Symmetry. (A,B)= Tr(ATB)= Tr((ATB)T) (ii) Linearity. Let A, B, C & Rnxm, a ER (aA+B, C) = Tr((aA+B) C) by linearity of the trace = Tr(aAC+BC) = aTr(AC)+Tr(BC) a (A, C) + (B, C)  $\langle A, A \rangle_{F} = Tr(A^{T}A) = \sum_{i=1}^{T} (A^{T}A)_{i,i} = \sum_{i=1}^{T} \sum_{j=1}^{A^{T}} A_{i,j}^{T} A_{j,i}$ Hence  $(A,A)_{c} \ge 0$  with equality if and only if  $A_{i,j}^2 = 0$  for all i, i.e. A = 0. Let A = UIV' be the singular value decomposition VITU OIVI) because U orthogonal because Tr(MM2)=Tr(M2HA) I Tr ( IT I because V orthogonal Honce MANG = VTr(ATA)







	values of 1/2 M. min(n,m)
	min(n, m)
	We get 11 x 11   = =
	· (Triangular inequality)
	We are going to show that
	IIMII * = max Tr(MQT) - CH) T
_	$\sigma_{\max}(Q) \leq 1$
	where max (a) denotes the maximal singular value
	of Q.
	Let UEVT be the SUD of A and define.
	0 = 021/2
	where $J \in \mathbb{R}^{n \times m}$ $J_{ij} =  1 \ i \neq i = j$ $0 \text{ otherwise}.$
	The largest eigen singular value of Q is 1: That (0)=1.
	Competer 5 12 12 12 12 12 12 12 12 12 12 12 12 12
	$T_{\Gamma}(HO^{T}) = T_{\Gamma}(U\Sigma V^{T}VJ^{T}U^{T}) = T_{\Gamma}(U\Sigma J^{T}U^{T})$
3	- A:41 1:001 2 (000; 11) - (000) min(n, m)
	$= Tr(U^T U \Sigma J^T) = Tr(\Sigma J^T) - \sum_{i=1}^{min(n,m)} C_i$
_	2000 parts 3000 300 10 60 10 60 10 10 10 10 10 10 10 10 10 10 10 10 10
No.	because U and V are orthogonal matrices and Tr(AB)=Tr(BA) for all matrices A, B
-	Tr(AB)=Tr(BA) for all matrices A, B
	the most policy of the contract of the



We conclude that ( UTOV) in & 1	and that
$Tr(HQ^T) \leq \sum_{i=1}^{min(n,m)} T_i = \ H\ _{\star}$	
This proves that max Tr(MQT) & 11H11+  GERNAM  TMAX(Q) & 1	
Consequently: 114114 = max ### Tr(HOT  GERENAM  GMAX(Q) S1	
Let now A, BERNAM	
$  A+B  _{\#} = \max_{\mathbf{C} \in \mathbb{R}^{n \times m}} \left( Tr(A\mathbf{G}^{T}) + Tr(B\mathbf{G}^{T}) \right)$ $Tr(\mathbf{A}\mathbf{G}^{T}) + Tr(\mathbf{B}\mathbf{G}^{T})$	
	= IIAN+ IICH+

```
In [2]: %matplotlib inline
import scipy
import matplotlib
import numpy as np
import matplotlib.pyplot as plot
from sklearn.cluster import KMeans
```

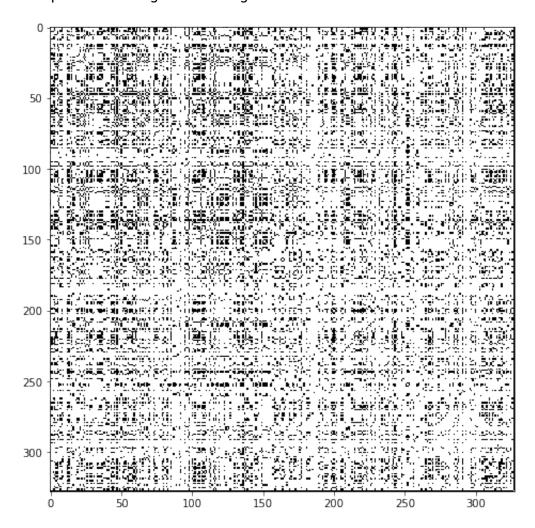
```
In [3]: # Reads the adjacency matrix from file
A=np.loadtxt('adjacency.txt')
print("There are", A.shape[0], "nodes.")
```

There are 328 nodes.

As you can see above, the adjacency matrix is relatively large (328x328): there are 328 persons in the graph. In order to visualize this adjacency matrix, it is convenient to use the 'imshow' function. This plots the 328x328 image where the pixel (i,j) is black if and only if A[i,j]=1.

```
In [4]: plot.figure(figsize=(8,8))
   plot.imshow(A,aspect='equal',cmap='Greys', interpolation='none')
```

## Out[4]: <matplotlib.image.AxesImage at 0x127ad6040>



a) Construct in the cell below the degree matrix:

$$D_{i,i} = \deg(i)$$
 and  $D_{i,j} = 0$  if  $i \neq j$ ,

the Laplacian matrix:

$$L = D - A$$

and the normalized Laplacian matrix:

$$L_{\text{norm}} = D^{-1/2} L D^{-1/2}.$$

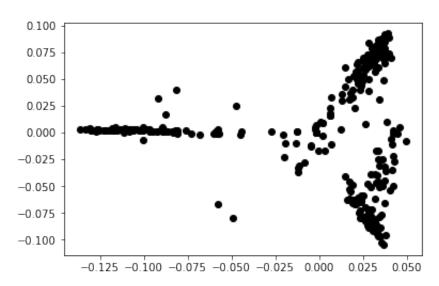
b) Using the command 'linalg.eigh' from numpy, compute the eigenvalues and the eigenvectors of  $L_{\rm norm}$  .

```
In [6]: v,w = np.linalg.eigh(L)
```

c) We would like to cluster the nodes (i.e. the users) in 3 groups. Using the eigenvectors of  $L_{\rm norm}$ , assign to each node a point in  $R^2$ , exactly as in 'Algorithm 1' of the notes where you replace L by  $L_{\rm norm}$ . Plot these points using the 'scatter' function of matplotlib.

```
In [7]: X=w[:,1:3]
plot.scatter(X[:,0],X[:,1], color='black')
```

Out[7]: <matplotlib.collections.PathCollection at 0x127dbed30>

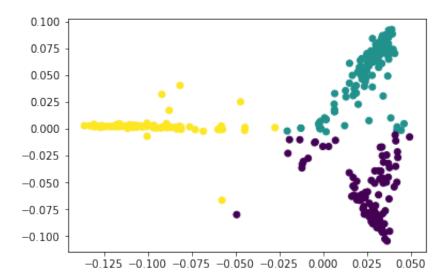


d) Using the K-means algorithm (use the built-in function from scikit-learn)

```
In [8]: # Replace ??? by the matrix of the points computed in (c)
kmeans = KMeans(n_clusters=3, random_state=0).fit(X)
labels=kmeans.labels_
# labels contains the membership of each node
```

```
In [9]: # Color the points according to their labels, to check that kmeans work
plot.scatter(X[:,0],X[:,1],c=labels)
```

Out[9]: <matplotlib.collections.PathCollection at 0x127b228b0>

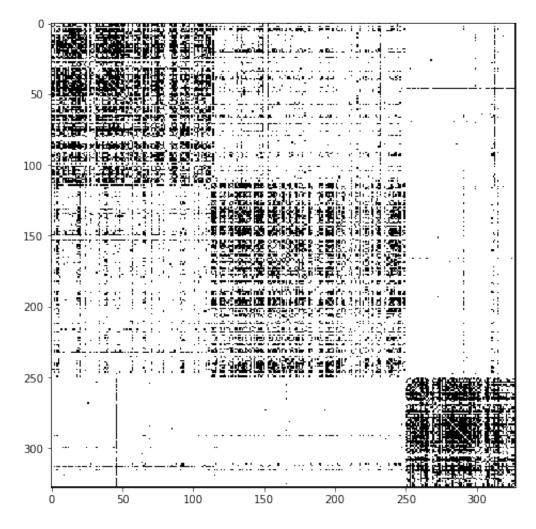


e) Re-order the adjacency matrix according to the clusters computed in the previous question. That is reorder the columns and rows of A to obtain a new adjacency matrix (that represent of course the same graph) such that the  $n_1$  nodes of the first cluster correspond to the first  $n_1$  rows/columns, the  $n_2$  nodes of the second cluster correspond to the next  $n_2$  rows/columns, and the  $n_3$  nodes of the third cluster correspond to the last  $n_3$  rows/columns. Plot the reordered adjacency matrix using 'imshow'.

```
In [10]: nodes = np.arange(N)
    cluster1 = nodes[labels==0]
    cluster2 = nodes[labels==1]
    cluster3 = nodes[labels==2]
    permutation = np.concatenate((cluster1, cluster2, cluster3))
    rA=np.zeros((N,N))
    for i in range(N):
        for j in range(N):
            rA[i,j]= A[permutation[i],permutation[j]]
```

```
In [11]: plot.figure(figsize=(8,8))
   plot.imshow(rA,aspect='equal',cmap='Greys', interpolation='none')
```

Out[11]: <matplotlib.image.AxesImage at 0x127ba3d60>



```
In [ ]:
```