Solutions - Assignment 5

1. Non-negative matrix factorization (NMF) to extract pure component spectra

The absorbance data considered from 'Inorfull.mat'. In Part (a), the absorbance data of the first replicate of the mixtures collectively forms the data matrix $Z_{26\times176}$. The number of species is known a priori to be 3. The NMF code provided by *Prof. Haesun Park, GATech* is used for solving the problem. The input to the *NMF* function ('nmf.m') are the data matrix, number of species and the initial guesses are the absolute values of the loadings and the scores matrix calculated from svd(Z).

$$\mathbf{B}_{26\times 3} = abs(\mathbf{U}(:,1:Ns)) P_{3\times 176} = abs(\mathbf{S}(1:Ns,1:Ns)\mathbf{V}(:,1:Ns)')$$

In part (b), the average of the 5 replicates of every mixture is considered for the data set *Z*. The correlation matrices computed between the actual pure component spectra and the extracted ones is presented below.

-0.5629	0.2143	0.3433
0.6284	0.6342	-0.4021
0.8738	-0.0757	0.2442

Figure 1: Correlation matrix for part (a)

-0.5209	-0.1247	0.7194
0.5826	0.8541	-0.4682
0.9023	-0.0483	0.0583

Figure 2: Correlation matrix for part (b)

Observations: Pure species spectra are unambiguously extracted using average values (since average data has less noise), the sources are not separated well using non-averaged measurements. The first extracted spectra has a high correlation with all pure species spectra. The second and third extracted spectra have maximum correlation with the same (second) pure species spectra.

2. A)
$$A_{struct}$$
 of the given network $\begin{bmatrix} X & 0 & X \\ X & 0 & X \\ 0 & X & X \\ X & 0 & X \\ X & X & 0 \\ X & 0 & X \\ 0 & X & X \end{bmatrix}$

For the network to be NCA compliant, the number of zeros in each column has to be $\geq m-1$. Here m=3, therefore the given network is NCA compliant.

B) Assuming the number of species is known to be 3, the rotation matrix M can be calculated by solving system of linear equations

$$\textbf{\textit{U}}_1\textbf{\textit{M}} = \textbf{\textit{A}}_{struct}$$
 Thus the calculated rotation matrix is given by $M = \begin{bmatrix} 1 & 2.596 & 9.325 \\ 0.27 & 1 & -11.721 \\ 1.556 & -0.211 & 1 \end{bmatrix}$

The correlation matrix is given in Figure 3

-0.9425	0.4291	0.4343
0.3662	-0.9968	-0.3815
0.4940	-0.4074	-0.9967

Figure 3: Correlation matrix for part (b)

Observations: The pure species spectra of the three species are extracted very well (correlations greater than 0.9).

C) Solution using the NCA toolbox

0.9425	-0.4293	-0.4342
-0.3659	0.9968	0.3815
-0.4942	0.4075	0.9967

Figure 4: Correlation matrix from NCA algorithm

