2D CORRELATION SPECTROSCOPY AND ITS APPLICATION IN VIBRATIONAL SPECTROSCOPY USING MATLAB

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Abstract

Two dimensional correlation spectroscopy is a powerfull tool for spectral analysis. It is able to reveal correlations between spectral changes and to deconvolve overlapping peaks. It is easily applicable in a study of biomolecules. Here presented program was created for easy accessibility of all necessary operations. Presented algorithm was deeply tested on regular spectra as well as on a spectra of vibrational optical activity. Basic mathematics needed for understanding and performing two-dimensional correlation spectroscopy was also presented.

1 Two dimensional correlation spectroscopy

Two dimensional correlation spectroscopy (2DCoS) originated in a NMR spectroscopy. There were efforts to use it in different methods of optical spectroscopy since its development. The main reason is that 2DCos has an ability to determine an order of bands and also has superior deconvolution abilities. This effort has run into troubles with a time scale because very short light pulses are needed for excitation. That wasn't possible until construction of femtosecond lasers and therefore existed an effort for creating a generalised form of 2DCoS. Practical use of so called Generalised 2DCoS was enabled in 1986 when Noda [1] utilised its basic theory. In Generalised 2DCos external impulses (pH, temperature, concentration etc.) are used to stimulate the system instead of short excitation pulses. Excited system is then analysed by one or more spectroscopic techniques. This spectra are then processed by a correlation function into a 2D spectra.

This method allows us to study the response of our system on different physical-chemical impulses with use of conventional spectrometers and without need of femtosecond lasers and optical stimulation [2]. It also provides information about order of spectral bands in dependency on external impulse and has very strong deconvoluting abilities.

The first part of this spectra (so called synchronous) is representing simultaneous or coincidental changes of measured spectral series. This spectrum is allways symetrical along the diagonal and has peaks on the diagonal. Intensity of these peaks (also called autocorrelation) is representing a strength of the band. The peaks off the diagonal are called cross-peaks and are representing a degree of correlation. When this peak is positive then both peaks are changing in the same direction (both increasing or both decreasing). When negative, peaks are changing in the different way (one is decreasing and the other is increasing). These rules are reversed when these peaks have different signs.

Second part of this spectra (asynchronous) is representing sequential or successive changes of measured spectral series. It is always antisymetrical along the diagonal and there are no peaks on the diagonal. When the cross-peak is positive then a band from the first spectra is growing earlier or more intensive then a band from second spectra and vice versa.

2DCoS can be performed in two ways. First of them is called homospectral correlation and there are two identical assemblies of data (spectra) on the input. This method can be used to deconvolve and to determine correlations between bands in the spectra. The second one is called heterospectral correlation and there are two different sets of data on the input. They are obtained by measuring the system with different spectroscopic techniques. This is the most powerfull variant because when we know the explanation of some peaks in one type of spectra and we see that this band is strongly correlated with another band in the second spectra it is most probable that these two bands have the same origin.



Figure 1: Synchronous (A) and asynchronous (B) spectra with marked correlation squares and peaks.

Some basic properties of correlation spectra are marked on figure 1. We can see diagonal peaks in the synchronous part and a symetrical positive cross-peak which indicates that both bands are changing in the same way. There are no diagonal peaks in the asynchronous part and from negative peak in the lower right corner we can say that band on 70 is changing slower than band on 30 (positive peak in the upper left corner is saying the same). In both cases we can draw so called correlation square which is connecting all peaks belonging to two correlating bands.

2 Computation of 2D spectra

First of all we need a set of spectra measured on a system which was induced by some external impulse. Spectral intensities of this set can be expressed as $I(\nu, t)$ where ν is a spectral characteristic (wavenumber, wavelength or Raman shift) and t is a parameter of an external impulse. That can be an evolution in time, temperature, pH, concentration etc. Only certain range of t can be measured and therefore a dynamical spectrum is implemented as [3]:

$$\widetilde{y}(\nu, t) = \begin{cases} y(\nu, t) - \widetilde{y}(\nu) & \text{if } T_{min} \le t \le T_{max}, \\ 0 & \text{otherwise} \end{cases}$$
(1)

where $\tilde{y}(\nu)$ is a reference spectrum of the system. An average spectrum is usually picked as a reference spectrum but any reasonable spectrum can be chosen. Correlation spectrum is then defined as:

$$\chi(\nu_1, \nu_2) = \langle \widetilde{y}(\nu_1, t) \cdot \widetilde{y}(\nu_2, t) \rangle \tag{2}$$

where the symbol $\langle \rangle$ is a crosscorrelation function defined as:

$$C(t,\tau) = \langle \Phi(\tau) | \Psi(t) \rangle \tag{3}$$

This spectra expresses a functional dependency between ν_1 and ν_2 during the interval of external variable t. We will treat this function as a complex one for further simplifying and divide it into a synchronous and asynchronous part [4]:

$$\chi(\nu_1, \nu_2) = \Phi(\nu_1, \nu_2) + i\Psi(\nu_1, \nu_2) \tag{4}$$

and formal definition of 2D correlation spectra is given by:

$$\Phi(\nu_1,\nu_2) + i\Psi(\nu_1,\nu_2) = \frac{1}{\pi(T_{max} - T_{min})} \times \int_0^\infty \widetilde{Y}_1(\omega) \cdot \widetilde{Y}_2(\omega) d\omega$$
(5)

The expression $\widetilde{Y}_1(\omega)$ is a Fourier transformation of spectral intensity variations $\widetilde{y}_1(\nu_1, t)$ measured in certain spectral variable ν_1 during an external impulse t. It can be written as:

$$\widetilde{Y}_1(\omega) = \int_{-\infty}^{\infty} \widetilde{y}(\nu_1, t) e^{-i\omega t} dt$$
(6)

As it is clear from Eq. 1, this Fourier transformation is related with a strict interval of an external impulse between T_{min} and T_{max} . $\tilde{Y}_2(\omega)$ is defined in a similar way but it is an inverse Fourier transform.

It is possible, with use of classical timeseries analysis and Wiener-Khintchinsky theorem [5], to express synchronous and asynchronous part:

$$\Phi(\nu_1, \nu_2) = \frac{1}{T_{max} - T_{min}} \int_{T_{min}}^{T_{max}} \widetilde{y}(\nu_1, t) \cdot \widetilde{y}(\nu_2, t) dt$$
(7)

$$\Psi(\nu_1,\nu_2) = \frac{1}{T_{max} - T_{min}} \int_{T_{min}}^{T_{max}} \widetilde{y}(\nu_1,t) \cdot \widetilde{z}(\nu_2,t) dt$$
(8)

where $\tilde{z}(\nu_2, t)$ is defined as follows:

$$\widetilde{z}(\nu_2, t) = \frac{1}{\pi} \int_{-\infty}^{\infty} \frac{\widetilde{y}(\nu_2, t')}{t' - t} dt$$
(9)

But this direct computation is very time consuming and therefore is used very rarely. Some simplification can be made when the data are equidistantly spaced.

For synchronous part we will start from the equation (7) and rewrite it as follows [6]:

$$\Phi(\nu_1, \nu_2) = \frac{1}{m-1} \sum_{j=1}^m \widetilde{y}(\nu_1) \cdot \widetilde{y}(\nu_2)$$
(10)

where m is a number of spectra equidistantly measured between T_{min} and T_{max} with growth defined by:

$$\Delta t = \frac{T_{max} - T_{min}}{m - 1} \tag{11}$$

It is even more simple when written in a Matlab notation:

Computation of asynchronous part is a little bit more complicated. First of all we need so called Hilbert–Noda matrix defined as:

$$N_{jk} = \begin{cases} 0 & \text{for } j = k \\ \frac{1}{\pi(k-j)} & \text{otherwise} \end{cases}$$
(12)

written in a Matlab notation:

```
N = zeros(n-1);
for i=2:(n-1)
    N(1,i) = 1/(pi * (i-1));
end
for i=2:(n-2)
    for j=i:(n-1)
        N(i,j) = N(i-1,j-1);
    end;
end
for i=2:(n-1)
    for j=1:i
        N(i,j) = -N(j,i);
    end;
end
```

Hilbert–Noda matrix is a slightly modified Hilbert matrix. This matrix is bringing into a time delay between two changes in a spectrum.

With use of Hilbert–Noda matrix we can compute asynchronous spectra using this prescription:

$$\Psi(\nu_1, \nu_2) = \frac{1}{m-1} \widetilde{y}(\nu_1)^T N \widetilde{y}(\nu_2)$$
(13)

and in a Matlab notation:

asynch = data(:,2:n) * N * data(:,2:n)'/(n-2);

3 Graphics User Interface

Here presented program has all necessary controls and abilities to perform 2DCoS on a custom set of data and to present and save the output results. It is based on the algorithms presented in chapter 2. It's design is focused on maximum lucidity and effectivity. GUI is pictured in figure 2 with all important places marked. Main plot of 2D spectra is placed in (1) and one of the spectra from each spectral set is placed in (2) (both spectra are identical in case of homospectral correlation). Colorbar (when active) is placed in (3) and all necessary controls are grouped in (4).

All controls are integrated into menus according to their functions. They are arranged in an assumed order of use. Particular items become active only after all prerequisites for their save execution are met. All functions are described in the included help file. Details are in figure 3.

4 Testing of the program

Functionality of these algorithms and program itself was tested on series of different spectra. They are simulating a lot of situations which can occur in spectra. The first testing spectra contains two Gaussian peaks which were described by function:

$$G = k \cdot e^{-\frac{(x - pos)^2}{20}} \tag{14}$$

where pos is a position of a peak and k is changing with an order of spectra (5i and $\frac{10}{i^2}$).

The whole set of spectra can be seen on figure 4 and the output of 2DCoS can be seen on figure 5.



Figure 2: GUI window with marked main controls.



Figure 3: GUI with some of menus and submenus.

As the second testing spectra we selected Lorentzian peaks (fig 6) which are described by the equation:

$$L = k \frac{1}{1 + \frac{(x - pos)^2}{20}} \tag{15}$$

As can be seen from fig 7 their correlation peaks are sharper and has slightly different shape but same rules described previously can be applied. We will focus on Lorentzian peaks further because they are more related to the real bands in spectra.

All performed tests (data not shown) were with a very good agreement with previously published results [7] and therefore we can assume that our algorithm is working correctly.

5 Problems with vibrational optical activity

Currently exists the methodology for performing and scoring results 2DCos for optical spectroscopy which contains bands of only one sign. Rules for scoring this correlation spectra are sometimes called Noda rules [6]. An attempt to extend this methodology also to spectra of Raman optical activity (ROA) was conducted recently [8] and first use of 2DCoS in ROA was in a last year [8] but only simple polyaminoacid chains were measured [9, 10] without working



Figure 4: Set of spectra representing 2 growing Gaussian peaks.



Figure 5: Synchronous and asynchronous 2D spectra of 2 growing Gaussian peaks.



Figure 6: Set of spectra representing 2 growing Lorentzian peaks.



Figure 7: Synchronous and asynchronous 2D spectra of 2 growing Lorentzian peaks.

out a detailed methodology. ROA and vibrational circular dichroism (VCD) are different from other spectroscopic techniques because they are differential methods and so they have bands of both polarity in spectra. The final methodology is a little bit different from the original rules and is listed in [8].

Ashton et al. have also conducted some theoretical test for veryfing this methodology but these tests were very simple. Used peaks were constructed from only three points and only growing or decreasing in magnitude were tested. So we have decided to make a proper testing which can be also applied on VCD. We picked the most simple ROA spectra as a test subject. It contains two bands of oposite polarity, one of them placed on 40 and the other one on 60.

The first set of spectra (fig 8) has both bands growing as we can see from the synchronous part (fig 9). Cross-peaks in asynchronous part of the spectra tell us that the band on 60 is growing more rapidly than the one on 40.



Figure 8: Set of spectra representing 2 growing ROA/VCD Lorentzian peaks.



Figure 9: Synchronous and asynchronous 2D spectra of 2 growing ROA/VCD Lorentzian peaks.

The second set of spectra (fig 10) has bands changing in an oposite direction because the cross-peaks in a synchronous part are negative. From the asynchronous part we can say that the band on 40 is changing more rapidly than the band on 60.



Figure 10: Set of spectra representing 2 ROA/VCD Lorentzian peaks changing in an oposite direction.



Figure 11: Synchronous and asynchronous 2D spectra of 2 ROA/VCD Lorentzian peaks changing in an oposite direction.

Another set of spectra (fig 12) has bands which are moving towards each other without changing magnitude. Both 2D spectra (fig. 13) are becoming more complicated.

The biggest disadvantage of ROA 2DCoS is its high sensitivity for spectral background, even small change has a great effect on an output 2D spectrum and so precise data treatment is needed.



Figure 12: Set of spectra representing 2 ROA/VCD Lorentzian peaks moving towards each other.



Figure 13: Synchronous and asynchronous 2D spectra of 2 ROA/VCD Lorentzian peaks moving towards each other.

6 Conclusions and future work

Two-dimensional correlation spectroscopy is a usefull tool in studying dynamic changes in spectra or for deconvolving overlapping peaks. It can analyze a spectral set under some perturbation and find any correlations or coincidences between spectral changes. It can also analyze different spectra under the same type of perturbation and then transfer explanation from one type of spectra to another type of spectra. This method is applicable on study of biomolecules and their interactions.

Future work will be focused on optimalizing the code and on expanding capabilities of this program to cover even some simple data treatment methods and to enhance its current abilities. We will also focus on practical use of this program in studying protein dynamics.

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