# **Progress in Bayesian DOSY and ROSY Transforms**



MESTRELAB RESEARCH Chemistry Software Solutions Carlos Cobas<sup>1</sup>, Stan Sykora<sup>2</sup>



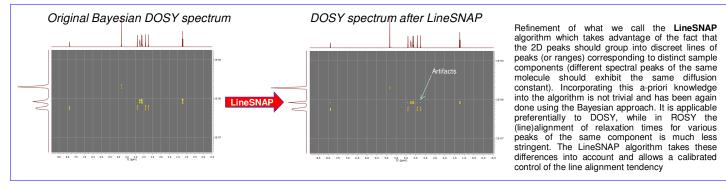
<sup>1</sup>Mestrelab Research, Santiago de Compostela, Spain, <u>www.mestrelab.com</u>

<sup>2</sup>Extra Byte, Castano Primo, Italy, <u>www.ebyte.it</u>

# Introduction

An efficient Bayesian approach (nicknamed BDT or **Bayesian Data Transform**) to the evaluation and graphic representation of multi-array NMR data sets characteristic of the diffusion (**DOSY**) and relaxation (**ROSY**) spectroscopies was first presented at ENC in 2008. Since than, it was beta tested and a considerable progress has been made which includes (a) a refinement of the LineSNAP section of the algorithm, (b) spectral alignment of the arrayed spectra using the novel **Global Spectrum Deconvolution (GSD**) algorithm and (c) combination of the BDT algorithm with a Bayesian handling of bi-exponentiality of overlapping spectral peaks. We will be happy to discuss with you the details of these developments.

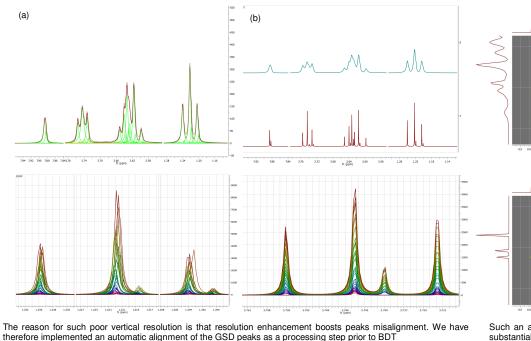


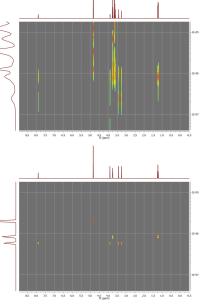


### (b) Spectral Alignment

Spectral overlap is an important problem in any DOSY processing: where peaks overlap, a monoexponential hypothesis yields compromise apparent diffusion coefficient. Overlapping can be minimized by exploiting the resolution power of GSD, as illustrated in the two figures below: (a) shows deconvolved peaks corresponding to the first trace of an PFG experiment whilst (b) shows the result of creating a synthetic GSD spectrum with a uniform line with of 0.2 Hz compared to the original, untreated spectrum

However, when BDT is applied to the GSD resolution enhanced spectra, it yields a DOSY map with long vertical strips





ve Such an automatic alignment algorithm results in a substantially higher quality of the final DOSY spectra

## (c) handling of bi-exponentiality

The Bayesian algorithm has been now combined with a Bayesian handling of bi-exponentiality. Considering the very limited number of spectra available in typical arrayed data sets (anything beyond 32 is considered impractical and 16 is a very popular choice), a full multi-exponential analysis is hardly ever feasible, but a separation of two decay components in cases of overlapping spectral peaks can be done with profit. However, since it affects negatively the final z-resolution, it must be applied only to those spectral regions where it is really needed and this, in addition to the incorporation of the bi-exponential hypothesis into the BDT algorithm, implies the need for an automatic procedure capable of selecting the non-exponential regions automatically on the basis of objective statistical criteria.

#### References

1 C.Cobas, M.Sordo, N.Larin, S.Sykora, Novel Data Evaluation Algorithms: Bayesian DOSY and ROSY Transforms, DOI: 10.3247/SL2Nmr08.009 2. C.Cobas, S.Sykora, Bayesian DOSY and ROSY transforms, J.Magn.Reson., submitted