

PROXIMAL ALGORITHM FOR L_1 COMBINED WITH MAXENT PRIOR: A NEW APPROACH FOR DOSY SIGNAL PROCESSING

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DOSY NMR

Diffusion Ordered SpectroscopY is a Nuclear Magnetic Resonance experiment using PFG (Pulsed Field Gradient), used to identify species in a complex chemical mixture by measuring its diffusion distribution.

• DOSY NMR Signal

$$I(q) = \int_{D_{min}}^{D_{max}} X(D) \exp(-D\Delta q^2) dD$$

with $X(D)$ is the diffusion distribution of the molecular species, I is the signal intensity, $q = \gamma\delta g$ presents the measure of the phase dispersion created by the PFG. γ is the gyromagnetic ratio of the studied spin, and δ and g are the duration and intensity of the PFG respectively.

• Goal

Find the chemical species from its diffusion profile by inverting the Laplace equation \rightsquigarrow Ill posed problem.

Theory

• Lagrangian formulation

$$\underset{X \in \mathbb{R}^N}{\text{minimise}} \|HX - Y\|^2 + \mu\Psi(X) \quad (1)$$

• Proposition : Hybrid regularisation

$$\Psi(X) = \lambda \text{ent}(X, a) + (1 - \lambda)\ell_1(X) \quad (2)$$

$$\text{ent}(X, a) = \begin{cases} \sum_{n=1}^N \frac{x_n}{a} \log\left(\frac{x_n}{a}\right) & \text{if } x_n > 0 \\ 0 & \text{if } x_n = 0 \\ +\infty & \text{elsewhere.} \end{cases} \quad \ell_1(X) = \sum_{n=1}^N |x_n|$$

Proximal solution

$$\text{prox}_{\Psi}(X) = (p(x_n))_{1 \leq n \leq N} \quad (3)$$

with

$$p(x_n) = \begin{cases} \frac{\lambda W}{a} \left[\frac{a}{\lambda} \exp\left(\frac{ax_n - a(1-\lambda)}{\lambda}\right) + \log(a) - 1 \right] & \text{if } \lambda \in]0, 1] \\ \text{sign}(x_n) \max(|x_n| - (1 - \lambda), 0) & \text{if } \lambda = 0 \end{cases} \quad (4)$$

W states for the Lambert function, defined as the inverse function of $f : z \rightarrow ze^z$ for all $z \in \mathbb{C}$ i.e :

$$z = we^w \Leftrightarrow w = W(z).$$

PALMA algorithm

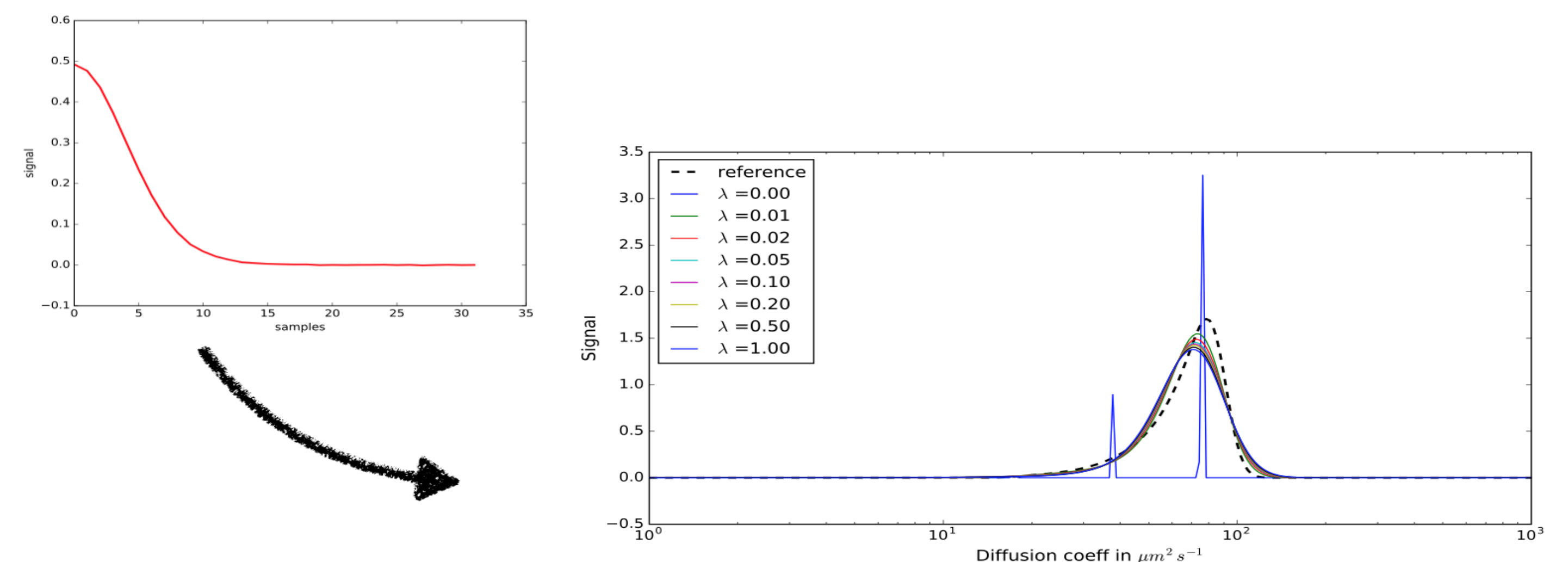
Proximal Algorithm for L_1 combined with MA_{xent} prior, available on the web site <http://palma.labo.igbmc.fr>, allows the analyse of your DOSY experiment in simple, efficient and easy way!

Simulated tests

• Polydisperse species

PALMA reconstruction for various values of λ of the simulated asymmetric distribution built from 15 log-normal components, ranging from 18 to $85\mu\text{m}^2/\text{s}$, with intensities ranging from 0.1 to 10. The intensities are chosen as $10/(1.4^{i-1})$, $i = 1, \dots, 15$.

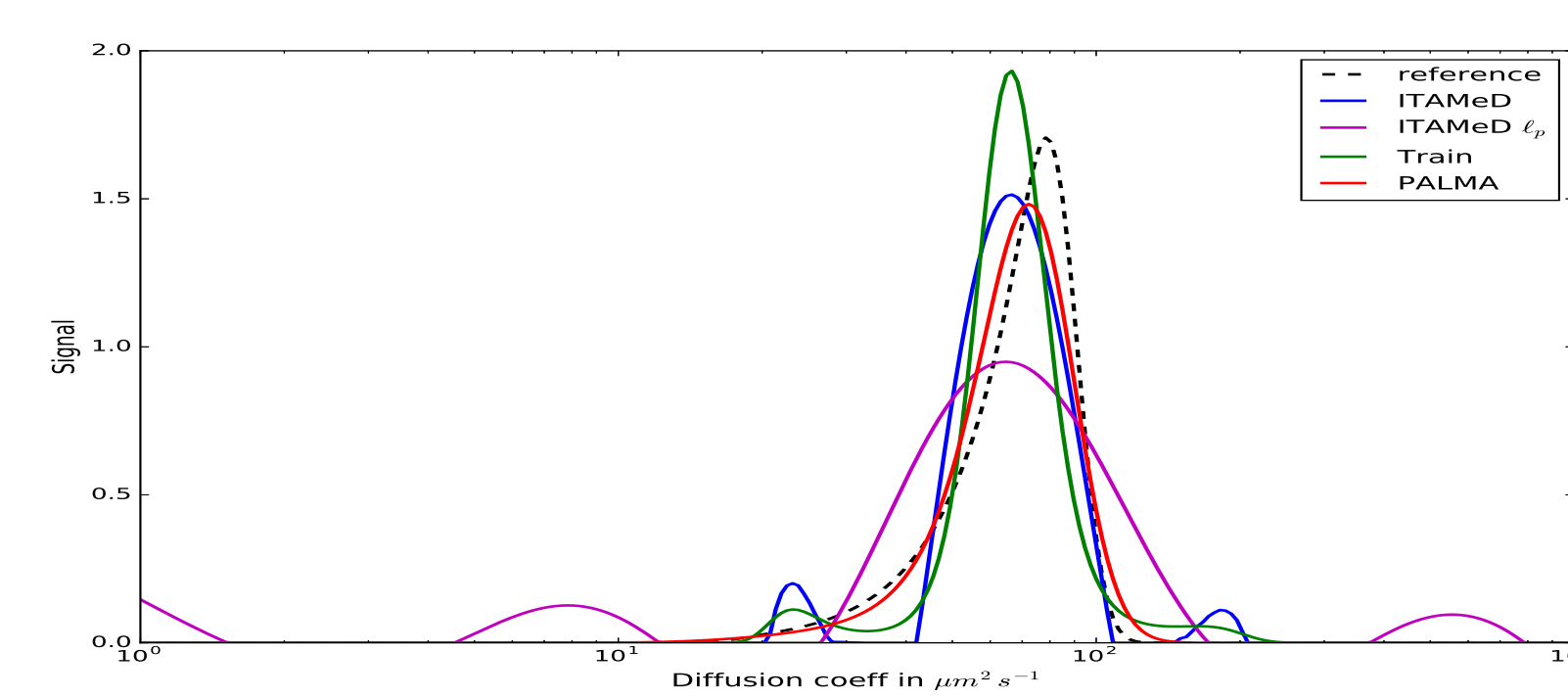
$M = 64$, $N = 256$, with noise level 0.1%.



\rightsquigarrow Best choice of $\lambda = 0.01$

• Comparaison with other algorithms

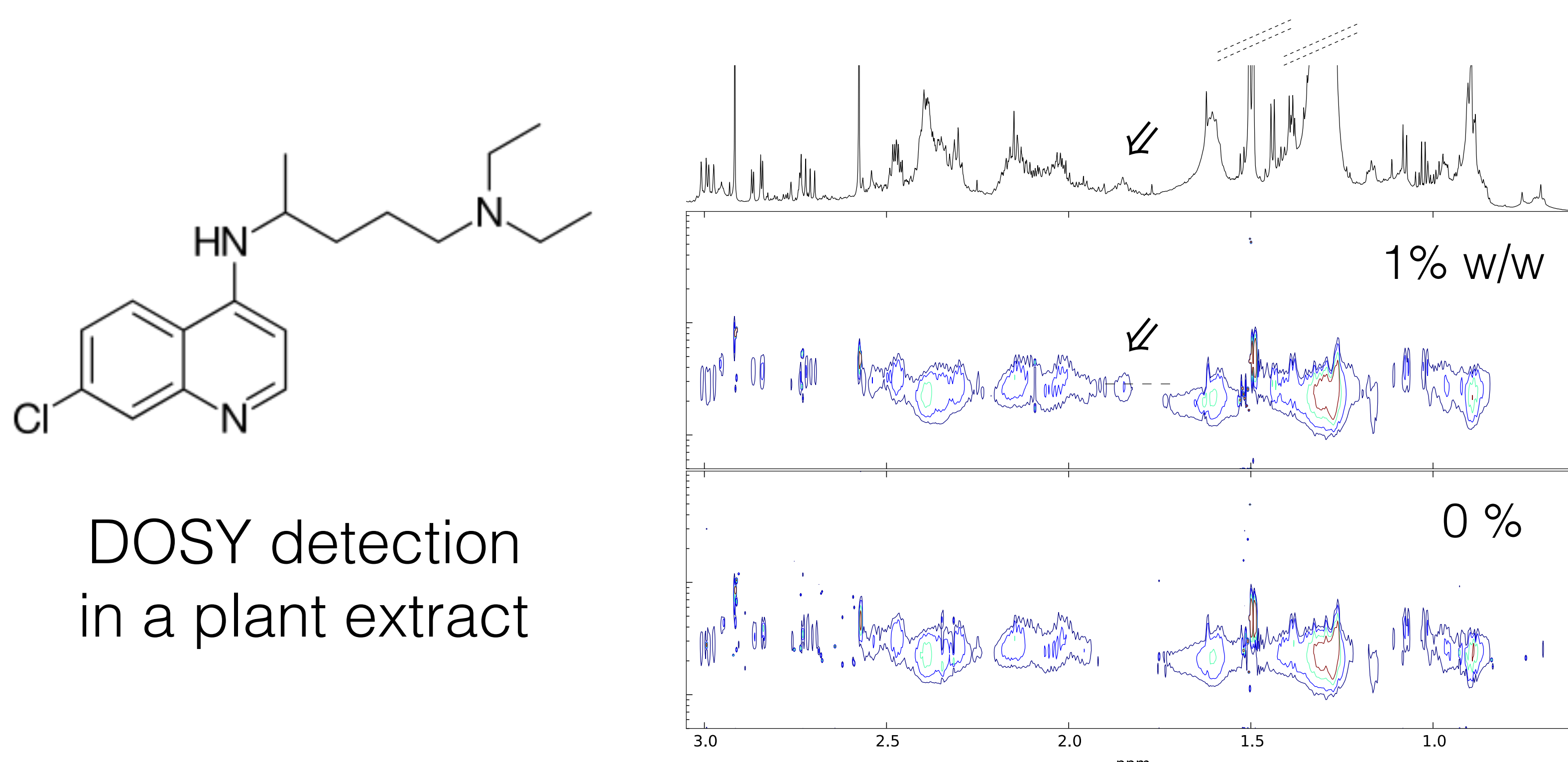
Comparison between our approach and other algorithms proposed to solve the ill-posed problem in DOSY NMR by recovering the signal defined in the above.



Algorithm	Qlty reconstruction ($-10 \log_{10} \left(\frac{\ x_{\text{rec}} - x_{\text{orig}}\ }{\ x_{\text{orig}}\ } \right)$) <i>Higher is better</i>
ITAMeD	08.89
ITAMeD with ℓ_p	06.18
TRAIIn	06.60
PALMA with $\lambda = 0.01$	14.60

\rightsquigarrow Best reconstruction with PALMA algorithm

Application polydisperse polymers



Comparison of two DOSY experiments on a brown algae extracts, one with 1% w/w 0.16 mg mL^{-1} Chloroquine added to the sample. The aliphatic region is shown, with the CH_2 signal of the Chloroquine highlighted with an arrow. The dashed horizontal label is at $285\mu\text{m}^2/\text{s}$.

Conclusion

• PALMA algorithm is a good tool to recover the DOSY NMR signal. It can be run in a fully automatic manner, where users may submit their datasets for automatic processing on the web site <http://palma.labo.igbmc.fr>

Perspectives

- Test with complex data in spectrometry field.
- Hybrid regularization in blind signal restoration problems