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An automatic L_1 -based regularization method for the analysis of FFC dispersion profiles with quadrupolar peaks

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Abstract

Fast Field-Cycling Nuclear Magnetic Resonance relaxometry is a non-destructive technique to investigate molecular dynamics and structure of systems having a wide range of applications such as environment, biology, and food. Besides a considerable amount of literature about modeling and application of such technique in specific areas, an algorithmic approach to the related parameter identification problem is still lacking. We believe that a robust algorithmic approach will allow a unified treatment of different samples in several application areas. In this paper, we model the parameters identification problem as a constrained L_1 -regularized non-linear least squares problem. Following the approach proposed in [*Analytical Chemistry* **2021** 93 (24)], the non-linear least squares term imposes data consistency by decomposing the acquired relaxation profiles into relaxation contributions associated with $^1H - ^1H$ and $^1H - ^{14}N$ dipole-dipole interactions. The data

fitting and the L_1 -based regularization terms are balanced by the so-called regularization parameter.

For the parameters identification, we propose an algorithm that computes, at each iteration, both the regularization parameter and the model parameters. In particular, the regularization parameter value is updated according to a Balancing Principle and the model parameters values are obtained by solving the corresponding L_1 -regularized non-linear least squares problem by means of the non-linear Gauss-Seidel method. We analyse the convergence properties of the proposed algorithm and run extensive testing on synthetic and real data. A Matlab software, implementing the presented algorithm, is available upon request to the authors.

Keywords: parameter identification, L_1 regularization, non-linear Gauss-Seidel method, Fast Field Cycling NMR relaxation, Free-model, quadrupole relaxation enhancement.

1. Introduction

Fast Field-Cycling (FFC) Nuclear Magnetic Resonance (NMR) relaxometry is a non-destructive magnetic resonance technique particularly useful in revealing information on slow molecular dynamics, carried out at very low magnetic field strengths. Standard NMR relaxation experiments are only performed in a relatively large fixed magnetic field that determines the resonance frequency of the molecules under investigation. Conversely, FFC-NMR relaxometry [1, 2] provides relaxation studies in a remarkably wide frequency range from approximately 1 kHz to 40 MHz. The FFC technique allows one to evaluate how the rate R_1 (also referred to as longitudinal re-

laxation rate) of a sample varies by changing the strength of an applied magnetic field, so forming the NMR Dispersion (NMRD) profiles. Therefore, FFC-NMR relaxometry measurements can detect the motion across a wide range of timescales (from millisecond to picoseconds) within an experiment. In addition, frequency-dependent relaxation studies have the exceptional potential to reveal the underlying mechanisms of molecular motion (not just its timescale). Provided that the proper constraints are fulfilled [3], spin relaxation can be described as a linear combination of spectral density functions (Fourier transform of the time correlation function) characterising the motional frequencies and their intensities present in the correlation function.

However, complex spins dynamical interactions may occur such as the Quadrupole Relaxation Enhancement (QRE) due to their intramolecular magnetic dipolar coupling with quadrupole nuclei of arbitrary spins $S \geq 1$ [4, 5]. In the case of nitrogen-containing systems, for instance, the presence of QRE is represented by local maxima or peaks of the R_1 profiles due to $^1H - ^{14}N$ interactions. The positions of the peaks depend on the quadrupole parameters which are determined by the electric field gradient tensor at the ^{14}N position. Consequently, even subtle changes in the electronic structure around ^{14}N reflect in changes of the position and shape of the quadrupole peaks. The QRE is a very sensitive fingerprint of molecular arrangement which has a wide range of applications ranging from environmental science [6], the study of ionic liquids, proteins [5] and food [7, 8].

Despite the consistent literature about the modeling of relaxation rate R_1 of protons fluids within a confined environment (see for instance [9, 10, 11])

and applications of FFC-NMR (see for example [12] and references therein), the study of a computational framework for the automatization of the FFC-NMR analysis is still missing. To the authors’ best knowledge, only P. Lo Meo et al. [13] propose a computational approach where, following the “model-free” approach introduced by [14, 15], the relaxation rate R_1 is represented as the sum of a constant term (offset) accounting for very “fast” molecular motion, a term describing proper $^1H - ^1H$ relaxation as an integral function of the correlated time distribution function, and a non-linear term depending on several characteristic parameters related to the QRE occurrence.

Therefore, the analysis of the NMRD profiles requires the solution of a parameter identification problem dealing with the estimation of the offset term, the correlation time distribution and the QRE parameters. In the present contribution, we formulate the parameter identification problem as a regularized non-linear least squares problem with box constraints and we propose a completely automatic strategy for its solution. In particular, the objective function contains a non-linear least squares term, imposing data consistency, and a L_1 -based regularization term. An L_1 -based regularization term, added to the L_2 -data fitting term, promotes sparse solutions since it forces only some components to be non-null while pushing all the other components to zero. Since the correlation time distribution function is known to be sparse, i.e. to have only a few non-null values, L_1 -based regularization is a suitable choice compared to L_2 -based regularization, also known as Tikhonov regularization, which indeed promotes smooth solutions. For a deeper discussion on regularization approaches based on L_1 and L_2 -norms we refer the reader to [16, 17, 18, 19]. The data-fitting and regularization terms are balanced by

the so-called regularization parameter. Physical constraints on the unknown parameters lead to bound constraints in the optimization problem.

The parameter identification problem crucially depends on the regularization parameter whose value has to be properly identified in order to perform a meaningful NMRD analysis. Therefore, our mathematical model depends on several parameters: the NMRD parameters (i.e. the offset, the correlation time distribution), the QRE parameters, and the regularization parameter. The estimation of all these parameters is carried out by an iterative process where, at each iteration, the regularization parameter is computed according to a balancing principle [20]. The NMRD and QRE parameters are estimated solving the corresponding constrained optimization problem by the constrained two-blocks non-linear Gauss-Seidel (GS) method [21, 22], since the unknown NMRD and QRE parameters can be naturally partitioned into two blocks. In the GS method, the objective function is iteratively minimized with respect to the offset and the correlation time distribution while the QRE parameters are held fixed; then, fixed the updated values for the offset and the correlation time distribution, the objective is minimized with respect to the QRE parameters. The first subproblem involves solving a constrained linear least squares problem, obtained by the *model-free* approach [13], with an L_1 regularization term. The second subproblem requires the solution of a constrained non-linear least squares problem.

This computational approach, separating the contribution due to the offset and the relaxation distributions from the parameters of the quadrupolar relaxation, is able to provide a very accurate fit not only of the overall NMRD profile, but also of the local maxima due to the QRE.

Besides analysing the convergence of the proposed approach, we tested it on synthetic and real data aiming to illustrate the algorithm efficiency and its robustness to data noise.

The main contributions of the present paper can be summarized as follows.

- We formulate the problem of identifying the offset, the correlation time distribution and the QRE parameters from the NMRD profiles as a L_1 -regularized non-linear least squares problem with box constraints related to physical properties of the parameters.
- We derive an automatic procedure, named AURORA (AUtomatic L_1 -Regularized mOdel fRee Analysis) for the identification of *all* the parameters of mathematical model, i.e, the NMRD parameters (the offset term, the correlation time distribution), the QRE parameters and the regularization parameter and we analyse its convergence properties.
- We prove the robustness of the proposed approach to data noise by testing it on synthetic and real NMRD profiles.

The remainder of this paper is organised as follows. In section 2 we present the parameter identification problem; in section 3 we introduce the solution method, analyse its properties and present the AURORA algorithm. The results from several numerical experiments are reported and discussed in section 4. Finally, in section 5, we draw some conclusions.

2. The parameter identification problem

In the following, we first describe the continuous model for NMRD profiles, then we derive its discretization and, finally, we present the parameter identification problem.

2.1. The continuous model of NMRD profiles

Following the *model-free* approach, [13] proposes a model for the NMRD profiles R_1 made of three components:

$$R_1(\omega) = R_0 + R^{HH}(\omega) + R^{NH}(\omega) \quad (1)$$

where R_0 is nonnegative offset keeping into account very *fast* molecular motions, the term $R^{HH}(\omega)$ describes the correlation distribution function $f(\tau)$ as:

$$R^{HH}(\omega) = \int_0^\infty \left[\frac{\tau}{(1 + (\omega\tau)^2)} + \frac{4\tau}{(1 + 4(\omega\tau)^2)} \right] f(\tau) d\tau \quad (2)$$

where τ is the correlation time, i.e., the average time required by a molecule to rotate one radian or to move for a distance as large as its radius of gyration.

Noticeably, the typical approach to data analysis in FFC NMR relaxometry relies on using an ad-hoc mathematical model containing information about both the number and meaning of the correlation times that describe a given system. Conversely, the approach described by equation (2) reverts the philosophy of the elaboration process. In fact, the integral form of equation (2) unconstrainedly retrieves only the number of possible correlation times describing the dynamics of the overall physical system. In other words, with-

out any ad-hoc pre-defined physical model, the proposed free-model analysis affords a fingerprint of the possible motion regimes regardless of their physical-chemical interpretation. The latter, in turn, can be reasonably attempted ex-post by interpreting the chemistry of the investigated system. The term $R^{HN}(\omega)$ describes the occurrence of the *quadrupolar peaks* [12]:

$$R^{HN}(\omega) = C^{HN} \left(\frac{1}{3} + \sin^2(\Theta) \cos^2(\Phi), \frac{1}{3} + \sin^2(\Theta) \sin^2(\Phi), \frac{1}{3} + \cos^2(\Theta) \right) \cdot \left(\begin{array}{c} \frac{\tau_Q}{1 + (\omega - \omega_-)^2 \tau_Q^2} + \frac{\tau_Q}{1 + (\omega + \omega_-)^2 \tau_Q^2} \\ \frac{\tau_Q}{1 + (\omega - \omega_+)^2 \tau_Q^2} + \frac{\tau_Q}{1 + (\omega + \omega_+)^2 \tau_Q^2} \\ \frac{\tau_Q}{1 + (\omega - (\omega_+ - \omega_-))^2 \tau_Q^2} + \frac{\tau_Q}{1 + (\omega + (\omega_+ - \omega_-))^2 \tau_Q^2} \end{array} \right) \quad (3)$$

where

- i) C^{HN} refers to “the gyromagnetic ratios and the average interaction distance of the nuclei”;
- ii) Θ and Φ are two angles accounting for “ the orientation of the $^1H - ^{14}N$ dipole-dipole axis with respect to the principal axis system of the electric field gradient at the position of ^{14}N ”;
- iii) τ_Q is the correlation time for the $^1H - ^{14}N$ quadrupolar interaction;
- iv) ω_- and ω_+ are the angular frequency position of the peaks on the NMRD profiles.

(We remark that in (3) the \cdot operator denotes the scalar product of two vectors.)

Simply speaking, the interpolation of the experimental data by applying equation (1) can be considered as a combination of the free model approach given by equation (2) for the homonuclear R^{HH} term, with the non-linear function developed in reference [4] and described by equation (3) for the quadrupolar R^{NH} term.

2.2. The discrete model for NMRD profiles

Before describing the discretization of the continuous model (1), let us introduce the following notation. Let $\boldsymbol{\omega} \in \mathbb{R}^m$ be the vector of the m Larmor angular frequency values ($\omega = 2\pi\nu$, ν in Mhz) at which R_1 is evaluated, and let $\mathbf{y} \in \mathbb{R}^m$ be the corresponding observations vector, i.e., $y_i = R_1(\omega_i)$, $i = 1, \dots, m$. Let $\mathbf{f} \in \mathbb{R}^n$ be the vector obtained by sampling $f(\tau)$ in n logarithmically equispaced values τ_1, \dots, τ_n . Finally, let $\boldsymbol{\psi} \in \mathbb{R}^6$ be such that $\psi_1 \equiv C^{HN}$, $\psi_2 \equiv \sin^2(\Theta)$, $\psi_3 \equiv \sin^2(\Phi)$, $\psi_4 \equiv \tau_Q$, $\psi_5 \equiv \omega_-$, $\psi_6 \equiv \omega_+$. The discrete model, obtained by discretizing the equations (2) and (3), is

$$\mathbf{y} = \mathcal{F}(\mathbf{f}, \boldsymbol{\psi}, R_0) \equiv \mathcal{F}_1(\mathbf{f}) + \mathcal{F}_2(\boldsymbol{\psi}) + R_0 \quad (4)$$

where $\mathcal{F} : \mathbb{R}^{n+6+1} \rightarrow \mathbb{R}^m$. The first term $\mathcal{F}_1 : \mathbb{R}^n \rightarrow \mathbb{R}^m$, only depending on \mathbf{f} , is a linear function of \mathbf{f} deriving from the discretization of the integral for R^{HH} in (2):

$$\mathcal{F}_1(\mathbf{f}) \equiv \mathbf{K}\mathbf{f}, \text{ with } \mathbf{K} \in \mathbb{R}^{m \times n}, \quad \mathbf{f} \in \mathbb{R}^n \quad (5)$$

and

$$\mathbf{K}_{i,j} = \frac{\tau_j}{(1 + (\omega_i \tau_j)^2)} + \frac{4\tau_j}{(1 + 4(\omega_i \tau_j)^2)}, \quad i = 1, \dots, m, \quad j = 1, \dots, n.$$

In a typical FFC-NMR experiment, $m \ll n$.

The second term $\mathcal{F}_2(\boldsymbol{\psi}) : \mathbb{R}^6 \rightarrow \mathbb{R}^m$ represents the quadrupolar component R^{HN} (3) and only depends on the parameters ψ_j , $j = 1, \dots, 6$:

$$(\mathcal{F}_2(\boldsymbol{\psi}))_i = \psi_1 \left(\frac{1}{3} + \psi_2(1 - \psi_3), \frac{1}{3} + \psi_2 \cdot \psi_3, \frac{1}{3} + (1 - \psi_2) \right) \cdot \left(\begin{array}{c} \frac{\psi_4}{1 + (\omega_i - \psi_5)^2 \psi_4^2} + \frac{\psi_4}{1 + (\omega_i + \psi_5)^2 \psi_4^2} \\ \frac{\psi_4}{1 + (\omega_i - \psi_6)^2 \psi_4^2} + \frac{\psi_4}{1 + (\omega_i + \psi_6)^2 \psi_4^2} \\ \frac{\psi_4}{1 + (\omega_i - (\psi_6 - \psi_5))^2 \psi_4^2} + \frac{\psi_4}{1 + (\omega_i + (\psi_6 - \psi_5))^2 \psi_4^2} \end{array} \right) \quad (6)$$

for $i = 1, \dots, m$.

The last term in \mathcal{F} is the constant parameter $R_0 \geq 0$ representing the offset in the NMRD curve.

2.3. The parameter identification problem

Mathematically, the problem of identifying the parameters \mathbf{f} , $\boldsymbol{\psi}$ and R_0 from the observations \mathbf{y} is an ill-conditioned non linear inverse problem (4). In order to stabilize the parameter identification procedure, we use a regularization approach adding some *a priori* information on the unknown parameters. In particular, we use L_1 regularization to induce sparsity of \mathbf{f} since the distribution $f(\tau)$ is known to be a sparse function with only a few non-null terms. Therefore, the parameter identification problem is reformulated as

the following optimization problem

$$\begin{aligned}
& \min_{\mathbf{f}, \boldsymbol{\psi}, R_0} \|\mathbf{y} - (\mathcal{F}_1(\mathbf{f}) + \mathcal{F}_2(\boldsymbol{\psi}) + R_0)\|_2^2 + \lambda \|\mathbf{f}\|_1 \\
& \text{s.t.} \quad \mathbf{f} \geq \mathbf{0}, \\
& \quad \boldsymbol{\psi} \in \mathcal{B}_\psi, \\
& \quad R_0 \geq 0,
\end{aligned} \tag{7}$$

where the set \mathcal{B}_ψ defines the box constraints on $\boldsymbol{\psi}$:

$$\mathcal{B}_\psi = \{\boldsymbol{\psi} : \psi_1 \in [0, \bar{C}]; \psi_2, \psi_3 \in [0, 1]; \psi_4 \in [0, \bar{\tau}]; \psi_5, \psi_6 \in [\omega_\ell, \omega_u]\}. \tag{8}$$

The bounds on the parameters ψ_i , $i = 1, \dots, 6$, can be derived from the physical properties of the system and from the data \mathbf{y} ; a deeper discussion on this topic will be given in the Section 4.

The regularization parameter $\lambda > 0$ weights the contribution of the L_1 regularization term; the parameters $(\mathbf{f}, \boldsymbol{\psi}, R_0)$ obtained by solving (7) depend critically on the value of λ .

3. The solution method

The presented parameter identification method is an iterative procedure where, at each iteration, a value of the regularization parameter λ is provided and the corresponding parameters $(\mathbf{f}_\lambda, \boldsymbol{\psi}_\lambda, R_{0,\lambda})$ are computed by solving problem (7). The constrained two-blocks non-linear Gauss-Seidel (GS) method [21, 22] is used for its solution. In the following, we firstly describe the GS method and recall its convergence properties, then we introduce the iterative procedure for the regularization parameter computation, and, fi-

nally, we draw the overall parameter identification procedure.

3.1. The constrained two-blocks Gauss-Seidel method

In this subsection, we describe the GS method used for the solution of the constrained optimization problem (7) for a fixed value of the regularization parameter λ . To this end, we partition the unknowns of (7) into two blocks such that the data fitting term is linear with respect to the first block and non-linear with respect to the second block. Therefore, we reformulate problem (7) as follows:

$$\begin{aligned} \min_{\mathbf{x}_1, \mathbf{x}_2} \quad & g(\mathbf{x}_1, \mathbf{x}_2) = \|\mathbf{y} - \mathbf{K}_e \mathbf{x}_1 - \mathcal{F}_2(\mathbf{x}_2)\|_2^2 + \lambda \|\mathbf{x}_1\|_1 + \eta \|\mathbf{x}_1\|_2^2 \\ \text{s.t.} \quad & \mathbf{x}_1 \in X_1, \\ & \mathbf{x}_2 \in X_2, \end{aligned} \tag{9}$$

where

$$\mathbf{x}_1 \equiv (\mathbf{f}, R_0), \quad \mathbf{x}_2 \equiv \boldsymbol{\psi} \tag{10}$$

$$X_1 = \{\mathbf{x}_1 \geq \mathbf{0}\}, \quad X_2 \equiv \mathcal{B}_\psi \tag{11}$$

and

$$\mathbf{K}_e = \begin{bmatrix} \mathbf{K} & \mathbf{1} \end{bmatrix} \in \mathbb{R}^{m \times (n+1)}. \tag{12}$$

The last L_2 -based penalty term $\eta \|\mathbf{x}_1\|_2^2$ in the objective function has been introduced to ensure that $\mathbf{K}_e^T \mathbf{K}_e + \eta \mathbf{I}$ is a definite positive matrix; to this end, a small value for η , as $\eta = 10^{-10}$ for example, can be fixed. Moreover, observe that in (9), the parameter R_0 has been included in the L_1 -based penalty term.

The closed subsets $X_1 \subseteq \mathbb{R}^{n+1}$ and $X_2 \subseteq \mathbb{R}^6$ are both convex; the objective function $g(\mathbf{x}_1, \mathbf{x}_2)$ is continuous and it is convex with respect to \mathbf{x}_1 for fixed \mathbf{x}_2 , but it is not convex with respect to \mathbf{x}_2 for fixed \mathbf{x}_1 . However, since $\mathbf{K}_e^T \mathbf{K}_e + \eta \mathbf{I}$ is definite positive and X_2 is bounded, it is easy to show that g is coercive on $X_1 \times X_2$.

Definition 3.1. A function $g : \mathbb{R}^q \rightarrow \mathbb{R}$ is called coercive in X if, for every sequence $\{\mathbf{x}^{(k)}\} \in X$ such that $\|\mathbf{x}^{(k)}\| \rightarrow \infty$, we have

$$\lim_{k \rightarrow \infty} g(\mathbf{x}^{(k)}) = +\infty$$

Proposition 3.1. *The function $g : \mathbb{R}^{n+1+6} \rightarrow \mathbb{R}$ such that*

$$g(\mathbf{x}_1, \mathbf{x}_2) = \|\mathbf{y} - \mathbf{K}_e \mathbf{x}_1 - \mathcal{F}_2(\mathbf{x}_2)\|_2^2 + \lambda \|\mathbf{x}_1\|_1 + \eta \|\mathbf{x}_1\|_2^2$$

is coercive in $X_1 \times X_2$.

Proof. The function g can be rewritten as

$$g(\mathbf{x}_1, \mathbf{x}_2) = \mathbf{x}_1^T (\mathbf{K}_e^T \mathbf{K}_e + \eta \mathbf{I}) \mathbf{x}_1 + 2\mathbf{x}_1^T \mathbf{K}_e^T (\mathcal{F}_2(\mathbf{x}_2) - \mathbf{y}) + \|\mathcal{F}_2(\mathbf{x}_2) - \mathbf{y}\|_2^2 + \lambda \|\mathbf{x}_1\|_1$$

where $\mathbf{K}_e^T \mathbf{K}_e + \eta \mathbf{I}$ is positive definite. Let $\{(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})\}$ be a sequence in $X_1 \times X_2$ such that $\lim_{k \rightarrow \infty} \|(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})\| = \infty$. Since X_2 is bounded, we have

$$\lim_{k \rightarrow \infty} \|\mathbf{x}_1^{(k)}\| = \infty \quad \text{and} \quad \lim_{k \rightarrow \infty} \|\mathbf{x}_2^{(k)}\| < \infty. \quad (13)$$

Let $\mu > 0$ be the smallest eigenvalue of $\mathbf{K}_e^T \mathbf{K}_e + \eta \mathbf{I}$. It holds

$$\begin{aligned} g(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)}) &\geq \mu \|\mathbf{x}_1^{(k)}\|^2 - 2\|\mathbf{K}_e^T(\mathcal{F}_2(\mathbf{x}_2^{(k)}) - \mathbf{y})\| \|\mathbf{x}_1^{(k)}\| + \lambda \|\mathbf{x}_1^{(k)}\| + \\ &\quad + \|\mathbf{K}_e^T(\mathcal{F}_2(\mathbf{x}_2^{(k)}) - \mathbf{y})\|^2 \\ &\geq \left(\mu \|\mathbf{x}_1^{(k)}\| - 2\|\mathbf{K}_e^T(\mathcal{F}_2(\mathbf{x}_2^{(k)}) - \mathbf{y})\| + \lambda \right) \|\mathbf{x}_1^{(k)}\| \end{aligned}$$

From (13) we have $\mu \|\mathbf{x}_1^{(k)}\| - 2\|\mathbf{K}_e^T(\mathcal{F}_2(\mathbf{x}_2^{(k)}) - \mathbf{y})\| + \lambda > 0$ for sufficiently large k , then

$$\lim_{k \rightarrow \infty} g(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)}) = +\infty$$

□

Continuity and coerciveness ensure the existence of at least one global minimizer of $g(\mathbf{x}_1, \mathbf{x}_2)$ in $X_1 \times X_2$ [23].

In the constrained two-blocks Gauss-Seidel method, at each iteration, the objective function is minimized with respect to each of the block coordinate vectors \mathbf{x}_i over the subsets X_i , $i = 1, 2$, as summarized in Algorithm 1, where the convergence condition is:

$$|g(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)}) - g(\mathbf{x}_1^{(k-1)}, \mathbf{x}_2^{(k-1)})| \leq Tol_{GS} |g(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})|. \quad (14)$$

We observe that the GS method is well defined since each subproblem has solutions. Indeed, the function g is strictly convex with respect of \mathbf{x}_1 and hence there exist at most one global minimum of f over X_1 for fixed \mathbf{x}_2 . On the other hand, Weierstrass's theorem guarantees the existence of at least one global minimum of g over X_2 for fixed \mathbf{x}_1 since g is continuous and X_2

Algorithm 1 Constrained two-blocks non-linear Gauss-Seidel method

```
1: function GS( $\mathbf{x}_1^{(0)}, \mathbf{x}_2^{(0)}$ )
2:   Set  $k = 0$  and  $\mathbf{x}^{(0)} = (\mathbf{x}_1^{(0)}, \mathbf{x}_2^{(0)})$ .
3:   repeat
4:      $k = k + 1$ 
5:     Set  $\mathbf{x}_1^{(k)} \in \arg \min_{\mathbf{z} \in X_1} g(\mathbf{z}, \mathbf{x}_2^{(k-1)})$ 
6:     Set  $\mathbf{x}_2^{(k)} \in \arg \min_{\mathbf{z} \in X_2} g(\mathbf{x}_1^{(k)}, \mathbf{z})$ 
7:   until convergence condition (14)
8:   return  $(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})$ 
9: end function
```

is a closed and bounded set.

For general nonconvex, constrained problems, convergence of sequences generated by the GS method to critical points has been proved in [22]. For the reader's convenience, we report here the main convergence result for the GS method and we refer to [22] for its proof.

Theorem 3.2. *Consider the problem*

$$\begin{aligned} & \min_{\mathbf{x}_1, \mathbf{x}_2} g(\mathbf{x}_1, \mathbf{x}_2) \\ & s.t. \quad \mathbf{x}_1 \in X_1, \\ & \quad \quad \mathbf{x}_2 \in X_2, \end{aligned} \tag{15}$$

where g is a continuously differentiable function and the subsets X_i are closed, nonempty and convex for $i = 1, 2$. Suppose that the sequence $\{(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})\}$ generated by the two-blocks GS method has limit points. Then, every limit point of $\{(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})\}$ is a critical point of the problem.

We have already observed that the objective function g of (9) is coercive; since the level sets of continuous coercive functions are compact, the sequence

$\{(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)})\}$ generated by the GS method has limit points (eventually, it has a convergent subsequence); hence, the GS method converges to critical points of (9).

We conclude this subsection with a remark on the solution of the two constrained subproblems to be solved at each iteration of algorithm 1. The first subproblem at step 3 is a L_1 -regularized least squares problem with nonnegativity constraints:

$$\begin{aligned} \min_{\mathbf{z}} \quad & \|\mathbf{w} - \mathbf{K}_e \mathbf{z}\|_2^2 + \lambda \sum_{i=1}^{m+1} z_i \\ \text{s.t.} \quad & z_i \geq 0, \quad i = 1, \dots, m+1 \end{aligned} \tag{16}$$

where $\mathbf{w} = \mathbf{y} - \mathcal{F}_2(\mathbf{x}_2^{(k)})$. For its solution, we use the truncated Newton interior-point method described in [24].

The second subproblem at step 4 is a bound constrained non-linear least squares problem:

$$\begin{aligned} \min_{\mathbf{z}} \quad & \|\mathcal{F}_2(\mathbf{z}) - \mathbf{w}\|^2 \\ \text{s.t.} \quad & \mathbf{z} \in X_2 \end{aligned} \tag{17}$$

where $\mathbf{w} = \mathbf{K}_e \mathbf{x}_1^{(k+1)} - \mathbf{y}$. For its solution, we use the Newton Projection method [25, 26] where the Hessian matrix is approximated as in the Levenberg-Marquardt method [27] since the Jacobian of \mathcal{F}_2 is ill-conditioned.

3.2. Computation of the regularization parameter λ

In order to correctly analyse the NMRD profiles, it is necessary to choose an appropriate value for the regularization parameter λ . Even if several parameter selection rules have been proposed in the literature for L_2 -regularized

minimization problems (see [28, 29, 30] for a theoretical discussion of such rules), the case of L_1 -based regularization still remains largely unexplored. In [31, 32], the discrepancy principle has been investigated for nonsmooth regularization. This principle is difficult to be realized since it requires the prior knowledge of the noise norm and a solution of the discrepancy equation is not guaranteed to exist. In [20], the Balancing Principle (BP) has been proposed where the regularization parameter is selected by balancing, up to a multiplicative factor γ , the data fidelity and the regularization term, i.e.:

$$\gamma\lambda\|\mathbf{x}_1\|_1 = \|\mathbf{y} - \mathbf{K}_e\mathbf{x}_1 - \mathcal{F}_2(\mathbf{x}_2)\|_2^2 + \eta\|\mathbf{x}_1\|_2^2 \quad (18)$$

The regularization properties of the BP has been deeply investigated and a convergent fixed-point iterative scheme for its realization has been proposed in [20]. We set $\gamma = 1$ which gives the following rule for the regularization parameter selection:

$$\lambda = \frac{\|\mathbf{y} - \mathbf{K}_e\mathbf{x}_1 - \mathcal{F}_2(\mathbf{x}_2)\|_2^2 + \eta\|\mathbf{x}_1\|_2^2}{\|\mathbf{x}_1\|_1}.$$

3.3. The parameter identification method

The proposed iterative method for the identification of both the NMRD parameters \mathbf{f} , $\boldsymbol{\psi}$ and R_0 and the regularization parameter λ is outlined in algorithm 2 where, given an initial guess for λ , at each iteration, the NMRD parameters are computed by solving problem (7) by the GS method, and the regularization parameter value is updated by the BP until the following

convergence condition is met:

$$|\lambda^{(k+1)} - \lambda^{(k)}| \leq Tol_\lambda |\lambda^{(k)}|, \quad Tol_\lambda > 0. \quad (19)$$

We refer to this method as AURORA (AUtomatic L_1 -Regularized mOdel fRee Analysis).

Algorithm 2 AURORA

- 1: Set $k = 0$, $\eta = 10^{-10}$ and choose a starting guess $\lambda^{(0)}$.
- 2: **repeat**
- 3: $k = k + 1$
- 4: **NMRD and QRE parameters update**

By algorithm 1 compute $(\mathbf{x}_2^{(k)}, \mathbf{x}_2^{(k)}) = GS(\mathbf{x}_1^{(k-1)}, \mathbf{x}_2^{(k-1)})$ i.e.

$$(\mathbf{x}_1^{(k)}, \mathbf{x}_2^{(k)}) \in \arg \min_{\substack{\mathbf{x}_1 \in X_1 \\ \mathbf{x}_2 \in X_2}} \|\mathbf{y} - \mathbf{K}_e \mathbf{x}_1 - \mathcal{F}_2(\mathbf{x}_2)\|_2^2 + \lambda^{(k)} \|\mathbf{x}_1\|_1 + \eta \|\mathbf{x}_2\|_2^2$$

- 5: **Regularization parameter update**

$$\lambda^{(k+1)} = \frac{\|\mathbf{y} - \mathbf{K}_e \mathbf{x}_1^{(k)} - \mathcal{F}_2(\mathbf{x}_2^{(k)})\|_2^2 + \eta \|\mathbf{x}_2^{(k)}\|_2^2}{\|\mathbf{x}_1^{(k)}\|_1}$$

- 6: **until** convergence condition (19)
 - 7: **return** $(\mathbf{f}, R_0) = \mathbf{x}_1^{(k)}$ and $\boldsymbol{\psi} = \mathbf{x}_2^{(k)}$ \triangleright Result $(\mathbf{f}, R_0, \boldsymbol{\psi})$
-

Following the analysis of the BP performed in [20], algorithm AURORA can be viewed as a fixed point-like scheme for the problem

$$\begin{aligned} (\mathbf{x}_1^*, \mathbf{x}_2^*) &= \arg \min_{\substack{\mathbf{x}_1 \in X_1 \\ \mathbf{x}_2 \in X_2}} \|\mathbf{y} - \mathbf{K}_e \mathbf{x}_1 - \mathcal{F}_2(\mathbf{x}_2)\|_2^2 + \lambda^* \|\mathbf{x}_1\|_1 + \eta \|\mathbf{x}_1\|_2^2, \\ \lambda^* &= \frac{\|\mathbf{y} - \mathbf{K}_e \mathbf{x}_1^* - \mathcal{F}_2(\mathbf{x}_2^*)\|_2^2 + \eta \|\mathbf{x}_1^*\|_2^2}{\|\mathbf{x}_1^*\|_1}. \end{aligned} \quad (20)$$

The monotone convergence of the sequence $\{\lambda^{(k)}\}$ generated by the fixed point scheme has been proved in [20] when $\lambda^{(0)}$ is chosen in an interval containing only one solution of equation (18).

4. Results and Discussion

In this section, we present and discuss the results obtained by a set of numerical experiments to assess the accuracy, robustness and efficiency of the proposed algorithm. In paragraph 4.1, we describe the experimental setting. In paragraph 4.2, we test AURORA on a synthetic NMRD profile R_1 computed by the model (1) with assigned values of the parameters ψ , \mathbf{f} and R_0 . We evaluate computational efficiency and accuracy of AURORA comparing it with some algorithms available in the Matlab optimization Toolbox. Moreover, we investigate the algorithm robustness in presence of data noise. Then, in paragraph 4.3, we report the results of the analysis of NMRD profiles from two different samples: Dry Nanosponge (DN) and Parmigiano-Reggiano (PR) cheese.

4.1. Numerical Experimental setting

All numerical computations are carried out using Matlab R2022a on a laptop equipped with Apple M1 processor and 16 GB 2133 MHz RAM. For all tests, the values \bar{C} and $\bar{\tau}$ in the constraints set \mathcal{B}_ψ (8) are set equal to a value large enough so that the intermediate solutions $\psi_1^{(k)}$ and $\psi_4^{(k)}$ never reach such bounds. In our tests $\bar{C} = \bar{\tau} = 100$ are suitable values. The interval $[\omega_\ell, \omega_u]$ in (8), representing the region where R_1 interrupts its decaying behaviour due to QRE, is defined by inspection of the NMRD profile.

The starting guess for the parameter $\psi_1^{(0)} \equiv C^{HN}$ is obtained by the literature [5]:

$$C^{HN} = \frac{2}{3} \left(\frac{\mu_0 \gamma_H \gamma_N \hbar}{4\pi r_{NH}^3} \right)^2 \approx 0.18 \left[\frac{\mu S}{s^2} \right] \quad (21)$$

where the value of the physical constants is reported in table 1. Concerning

Constant	Description	Value
μ_0	permeability of vacuum	$10^{-7} T^2 J^{-1} m^3$
γ_H	1H gyromagnetic factor	$2.577 \cdot 10^6 T^{-1} s^{-1}$
γ_N	^{14}N gyromagnetic factor	$3.078 \cdot 10^6 T^{-1} s^{-1}$
\hbar	reduced Planck's constant	$1.05472 \cdot 10^{-34} J s$
r_{HN}	$^1H - ^{14}N$ inter-spin distance	$1.4 \cdot 10^{-10} m$

Table 1: Characteristic constants for C^{HN} in (21).

the quadrupolar parameters, $\psi_2^{(0)} \equiv \sin^2 \Theta^{(0)}$, $\psi_3^{(0)} \equiv \sin^2 \Phi^{(0)}$, the initial values are equal to the mean of the corresponding upper and lower bounds in \mathcal{B}_ψ , i.e. 1/2. The initial value of $\psi_4^{(0)} \equiv \tau_Q$, is set to 1, while $\psi_5^{(0)} \equiv \omega_-^{(0)}$, and $\psi_6^{(0)} \equiv \omega_+^{(0)}$ are defined as follows:

$$\psi_5^{(0)} = \omega_\ell + \frac{1}{4} |\omega_u - \omega_\ell|, \quad \psi_6^{(0)} = \omega_u - \frac{1}{4} |\omega_u - \omega_\ell|.$$

The computed results are evaluated by the Mean Squared Error (MSE)

$$\text{MSE} = \frac{\|R_1 - \mathcal{F}(\mathbf{f}, \boldsymbol{\psi}, R_0)\|^2}{m},$$

and the Parameter Relative Error (PRE):

$$\text{PRE}(x) = \frac{\|x^{exact} - x^{computed}\|^2}{\|x^{exact}\|^2} \quad (22)$$

with x representing either the vector \mathbf{f} or the scalars $R_0, \psi_i, i = 1, \dots, 6$.

The components of the vector $\boldsymbol{\psi}$ are referenced by the name in the physical model (3), according to the mapping introduced in section 2.2, and reported in table (2) for convenience. All the tests apply algorithm 2 with $Tol_\lambda = 10^{-2}$

C^{HN}	Φ	Θ	τ_Q	ω_-	ω_+
ψ_1	$asin(\sqrt{\psi_2})$	$asin(\sqrt{\psi_3})$	ψ_4	ψ_5	ψ_6

Table 2: Quadrupolar parameters mapping.

in (19) and algorithm 1 with $Tol_{GS} = 10^{-6}$ in (14).

The computational cost is evaluated in terms of execution time and iterations number.

4.2. Synthetic test Problem

To investigate the properties of AURORA, we first test it on the synthetic NMRD profile R_1 represented in figure 1(a), and obtained by setting the parameters of model (1) as in the second column of table 3, with the distribution function \mathbf{f}^* represented in red in figure 2(a). Throughout the paragraph we use the frequencies ν instead of the angular frequencies ω , i.e. $\nu_- \equiv \omega_-/(2\pi)$ and $\nu_+ \equiv \omega_+/(2\pi)$.

The accuracy of the computed results can be appreciated in the correlation distribution \mathbf{f} and R_1 curves shown in figure 2. To test the convergence behaviour we evaluate the PRE and MSE at each step of the GS method in algorithm 1. Figure 3(a) shows the the behaviour of the relative errors for each parameter ($\mathbf{f}, R_0, C^{HN}, \Phi, \Theta, \tau_Q, \nu_-, \nu_+$) compared to their reference values. The convergence to reference parameters values is initially non monotonic for most parameters with the exception of τ_Q and ν_- . On the contrary, MSE has

	reference	computed	PRE
R_0	3.69	3.6868	$7.0267 \cdot 10^{-4}$
C^{HN}	18.84	18.8453	$6.1449 \cdot 10^{-5}$
τ_Q	0.96	0.9554	$8.5033 \cdot 10^{-6}$
Θ	1.09	1.0901	$6.1449 \cdot 10^{-5}$
Φ	0.57	0.5696	$6.9199 \cdot 10^{-4}$
ν_-	2.15	2.1502	$5.7363 \cdot 10^{-6}$
ν_+	2.87	2.8696	$1.1316 \cdot 10^{-6}$

Table 3: Model parameters: reference (second column), AURORA computed values (third column) and PRE (fourth column).

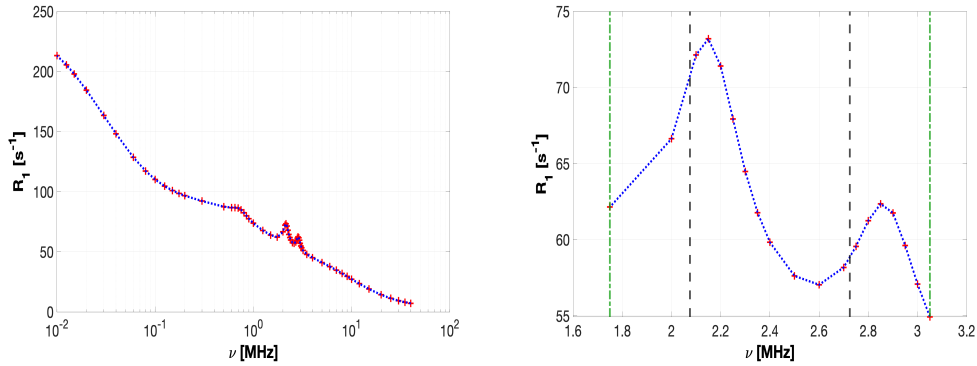


Figure 1: Synthetic sample. (a) Full NMRD profile (b) Zoom of NMRD profile in the reference interval $[\nu_\ell, \nu_u]$ represented by the left and right green vertical lines. Left and right black vertical lines represent the values $\psi_5^{(0)}/(2\pi)$, $\psi_6^{(0)}/(2\pi)$ respectively.

monotonic decrease as reported in figure 3(b). The values of the computed parameters and relative errors reported in the third and fourth columns of table 3 confirm the excellent accuracy obtained by the proposed algorithm. The computed value of the regularization parameter is $\lambda^* = 1.216 \cdot 10^{-9}$ with computation time of 90.44 ± 0.3 s.

The computation cost, evaluated in terms of inner iterations of the two Gauss-Seidel blocks in algorithm 1, consists of 147 iterations of the Newton Projection method and 74258 iterations of the truncated Newton interior

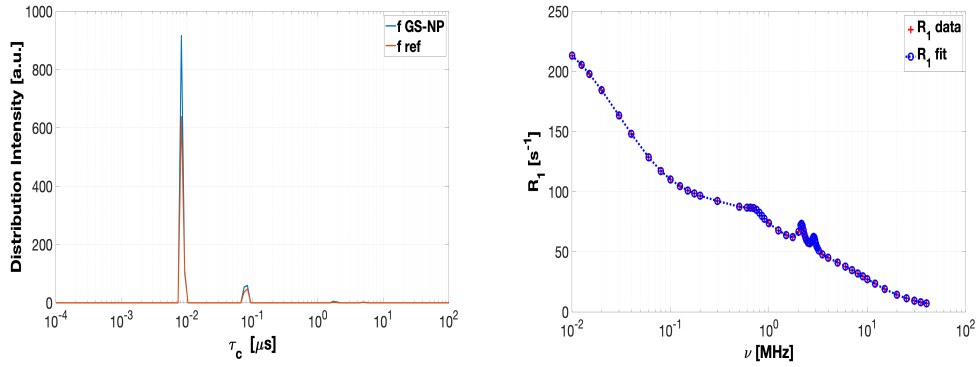


Figure 2: Synthetic sample. (a) Reference (red) and computed correlation distribution (blue). (b) Reference and computed R_1 profiles.

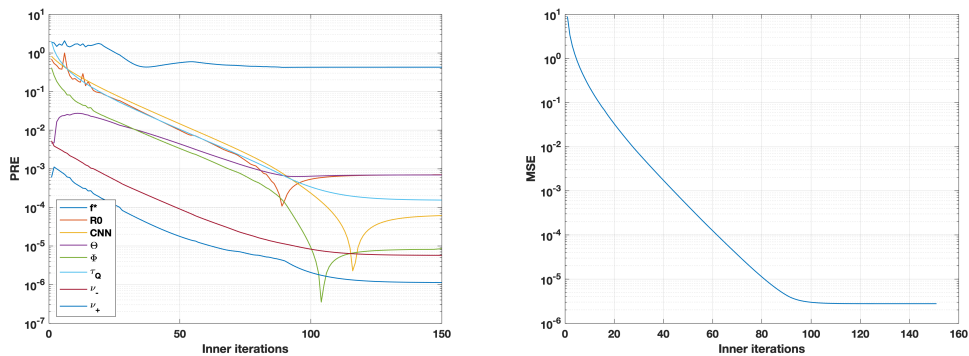


Figure 3: Synthetic R_1 . (a) PRE values per iteration (b) MSE values per iteration.

point method. Note that the greater computational weight lies in truncated Newton interior point iterations due to the larger size of problem (16) compared to (17).

Although the convergence of the update formula (18) depends on the initial guess $\lambda^{(0)}$, we experimentally found convergence for $\lambda^{(0)}$ in a quite large interval ($[10^{-16}, 10^0]$). In figure 4 we represent the sequences $\lambda^{(k)}$, $k = 0, \dots, 15$ obtained by algorithm 2 with $\lambda^{(0)} \in \{10^{-16}, 10^{-6}, 10^{-4}, 10^{-2}, 10^0\}$. Optimal convergence ($k = 1$) is obtained for $10^{-16} \leq \lambda^{(0)} \leq 10^{-4}$ while $\lambda^{(0)} > 10^{-4}$

causes a slight increase of the iterations number, still preserving the convergence up to $\lambda^{(0)} = 1$, which is usually considered as a standard starting guess. Therefore, to keep computations efficient, $\lambda^{(0)} = 10^{-6}$ is used throughout the numerical experiments of this section.

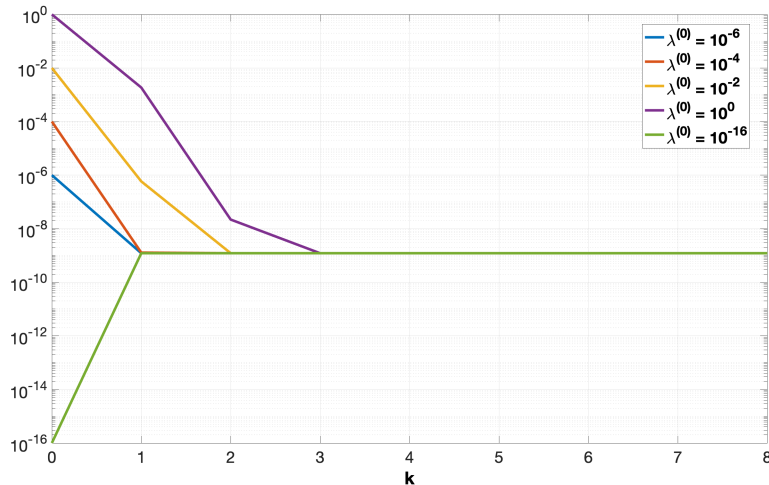


Figure 4: Synthetic R_1 . Sequence $\{\lambda^{(k)}\}$, obtained by AURORA with $\lambda^{(0)} \in \{10^{-16}, 10^{-6}, 10^{-4}, 10^{-2}, 10^0\}$.

Comparison with Matlab solvers. With this test problem, we aim to compare AURORA with several methods implemented by the Matlab function `fmincon`: such as interior-point (`ip`), the active-set (`as`), the sequential quadratic programming (`sqp`) and trust-region-reflective (`trr`) methods. We highlight that AURORA automatically computes the value of the regularization parameter λ while the Matlab function `fmincon` solves the optimization problem (7) for a fixed value of λ . Therefore, we compare the GS algorithm 1 with `ip`, `as`, `sqp` `trr` for the same fixed value $\lambda = 1 \cdot 10^{-8}$, which we heuristically found to be a good value for all the methods.

Besides the automatic computation of the regularization parameter λ , AURORA splits the unknown parameters in two blocks and alternatively minimizes the objective function for (R_0, \mathbf{f}) , the offset and correlation distribution, and for the quadrupolar parameters $\boldsymbol{\psi}$. Two different methods are used for the solution of the corresponding sub-problems. On the contrary, `fmincon` computes all the parameters applying the same method.

Table 4 shows the PRE and MSE values (last row) obtained by AURORA (second column) and by the Matlab solvers, highlighting the smallest values. The distribution \mathbf{f} computed by `sqp` is shown in figure 5.

Parameter	PRE				
	AURORA	ip	active-set	sqp	trr
\mathbf{f}	4.2834 10^{-1}	1.5509	1.4497	1.3020	8.5279 10^{-1}
R_0	7.0032 10^{-4}	9.9629 10^{-1}	1.0000	2.7930 10^{-1}	1.3671 10^{-1}
C^{HN}	5.8238 10^{-5}	4.2908	9.6353 10^{-1}	1.5045 10^{-5}	1.1591 10^{-2}
Θ	6.9108 10^{-4}	6.5929 10^{-2}	7.7862 10^{-2}	7.2072 10^{-4}	1.5758 10^{-2}
Φ	8.7093 10^{-6}	5.5535 10^{-1}	2.1372	2.6548 10^{-5}	7.2619 10^{-3}
τ_Q	1.5660 10^{-4}	9.9228 10^{-1}	3.1584 10^1	1.8903 10^{-4}	1.1033 10^{-2}
ν_-	5.7679 10^{-6}	4.0856 10^{-1}	2.2756 10^{-1}	5.6228 10^{-6}	5.9438 10^{-5}
ν_+	1.1391 10^{-6}	5.5197 10^{-2}	3.3362 10^{-2}	1.2084 10^{-6}	1.8516 10^{-5}
<i>MSE</i>	2.8131 10^{-6}	9.1906	9.0766	3.1658 10^{-6}	2.8289 10^{-3}

Table 4: Parameter relative errors and MSE of AURORA and methods implemented by the Matlab function `fmincon`.

We observe that AURORA has globally superior accuracy both in data fitting and parameter estimation. Only `sqp` has MSE value similar to AURORA ($3.1658e - 06$ compared to $2.8131e - 06$), and a slightly better PRE for parameters C^{HN} and ν_- , but the amplitude distribution in figure 5 shows too many spurious peaks.

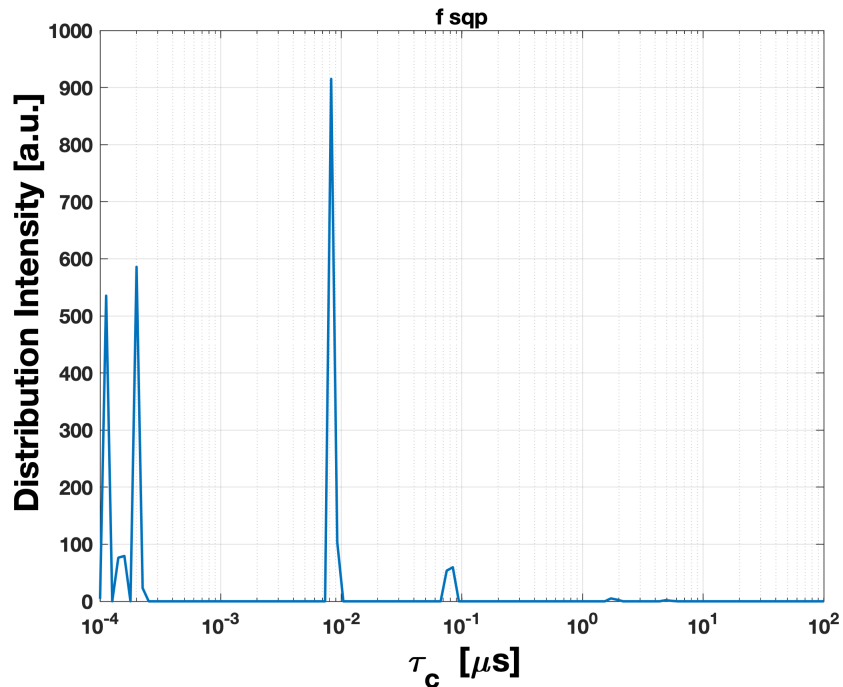


Figure 5: Correlation time distribution \mathbf{f} computed by `sqp` method.

Test with noisy data. In this paragraph we test the algorithm robustness to data perturbations by computing noisy data $\mathbf{y}^\delta \in \mathbb{R}^m$ from a random uniformly distributed vector $\mathbf{v} \in \mathbb{R}^m$ with values in the interval $[-1, 1]$ s.t.

$$y_i^\delta = y_i(1 + \delta v_i), \quad i = 1, \dots, m$$

and consider the cases $\delta = 1\%, 5\%, 10\%$. The computation times (averaged over 10 runs) and the iteration numbers, reported in table 5, show that both iteration numbers and computation times decrease with increasing noise percentage. This is due to the effect of increasing noise values on the computed regularisation parameters, as shown in the third column of table 5. The re-

δ	Time [s]	Total iterations Subproblem (16)	Total iterations Subproblem (17)	λ
$1 \cdot 10^{-2}$	38.39 ± 0.28	52217	89	$5.8405 \cdot 10^{-5}$
$5 \cdot 10^{-2}$	14.39 ± 0.16	44414	72	$1.8803 \cdot 10^{-3}$
$1 \cdot 10^{-1}$	4.90 ± 0.10	16846	28	$9.3561 \cdot 10^{-3}$

Table 5: Computation times and iteration numbers with noise $\delta = 1\%, 5\%, 10\%$.

lations between the execution times are not found precisely in the number of iterations because they do not take into account the backtracking steps internal to the Newton method.

Computing 500 noisy samples \mathbf{y}_j^δ , we run AURORA and compare the errors on the estimated parameters as well as reconstructed NMRD profiles.

For the noise values $\delta = 1\%, 5\%, 10\%$, we compute the mean PRE for each parameter and represent the mean values in the bar plot shown in figure 6 together with the product $C^{HN} \cdot \tau_Q$. The mean PRE and MSE are reported

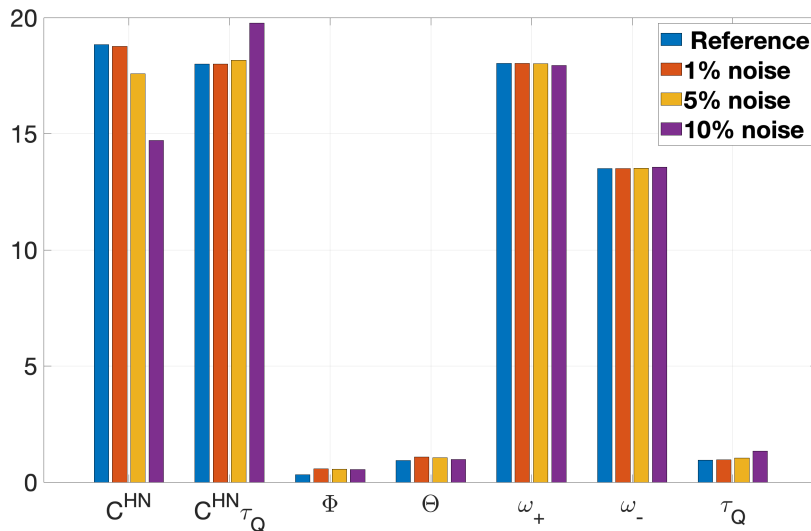


Figure 6: Mean parameter values computed by 500 noisy NMRD profiles with noise $\delta = 1\%, 5\%, 10\%$.

	PRE		
	1%	5%	10%
\mathbf{f}	$5.9019 \cdot 10^{-1}$	1.1816	1.4509
R_0	$3.6393 \cdot 10^{-2}$	$1.6726 \cdot 10^{-1}$	$1.8099 \cdot 10^{-1}$
C^{HN}	$3.3625 \cdot 10^{-2}$	$2.7021 \cdot 10^{-1}$	$4.7742 \cdot 10^{-1}$
Θ	$2.3023 \cdot 10^{-2}$	$1.0678 \cdot 10^{-1}$	$2.1726 \cdot 10^{-1}$
Φ	$3.5151 \cdot 10^{-2}$	$4.0280 \cdot 10^{-1}$	$6.5910 \cdot 10^{-1}$
τ_Q	$4.4998 \cdot 10^{-2}$	1.8862	$1.1095 \cdot 10^1$
ν_-	$4.3917 \cdot 10^{-3}$	$4.8712 \cdot 10^{-2}$	$7.2441 \cdot 10^{-2}$
ν_+	$3.0889 \cdot 10^{-3}$	$3.8712 \cdot 10^{-2}$	$5.6856 \cdot 10^{-2}$
MSE	$1.5980 \cdot 10^{-1}$	3.1441	$1.0055 \cdot 10^1$

Table 6: Mean PRE and MSE on 500 noisy NMRD profiles with $\delta = 1\%, 5\%, 10\%$.

in table 6. The computed R_1 curves and the zoom in the QRE interval are shown in figures 7,8 and 9 for $\delta = 1\%, 5\%, 10\%$ respectively.

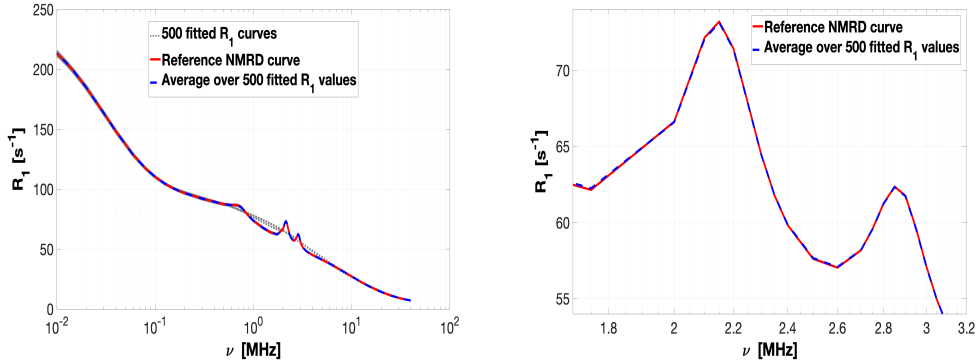


Figure 7: Fit of NMRD obtained from 500 noisy Synthetic NMRD curves with noise $\delta = 1\%$. (a) Light gray: 500 fitted R_1 curves, Red line: Reference NMRD curve. Blue line: average over 500 fitted R_1 values. (b) zoom in QRE interval.

In figure 6, we observe that data noise affects mainly C^{HN} , τ_Q and Φ values. However, considering the value of the product $C^{HN}\tau_Q$, represented by the second group in figure 6, we see that the value is preserved when $\delta = 1\%, 5\%$. This feature is a physical characteristic and allows us to consider

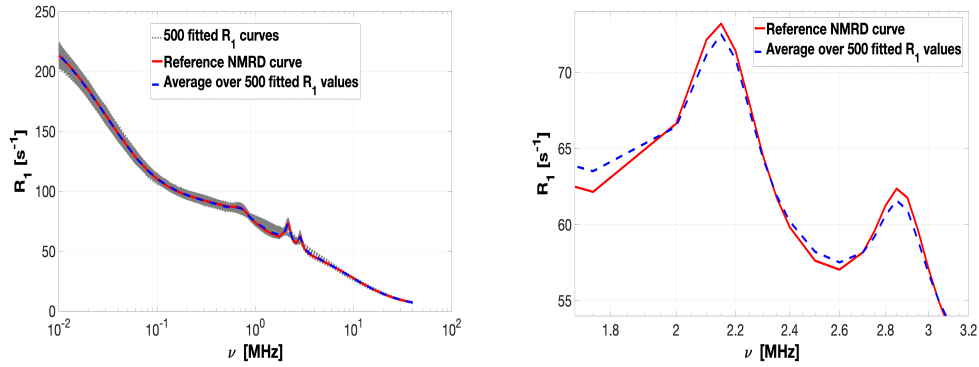


Figure 8: Fit of NMRD obtained from 500 noisy Synthetic NMRD curves with noise $\delta = 5\%$. (a) Light gray: 500 fitted R_1 curves, Red line: Reference NMRD curve. Blue line: average over 500 fitted R_1 values. (b) zoom in QRE interval.

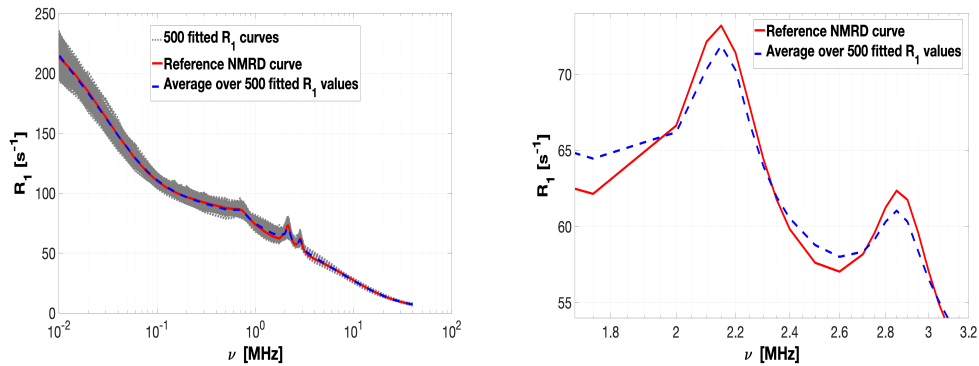


Figure 9: Fit of R_1 obtained from 500 noisy synthetic NMRD profiles with noise $\delta = 10\%$. (a) Light gray: 500 fitted R_1 curves, Red line: Reference NMRD curve. Blue line: average over 500 fitted R_1 values. (b) zoom in QRE interval.

accurate the related parameters.

Although the average MSE increase with data noise, the computed average R_1 curves show a very good agreement to the reference NMRD profiles (figures 7, 8 and 9). The QRE is well reproduced even with high noise (figures 7(b), 8(b) and 9(b)).

4.3. NMRD profiles from FFC measures

In this paragraph we consider the NMRD profiles obtained from two different materials described in [13].

- A sample of 24-month aged Parmigiano-Reggiano (PR) cheese. The NMRD profile represented in figure 10(a) has $m = 48$ values with confidence intervals ranging from $\pm 0.35\%$ to $\pm 3.07\%$ of the value. The quadrupolar peaks, represented in figure 11(a), correspond to frequency values $\nu_- = 2.1$ and $\nu_+ = 2.8$ of values $R1_- = 32.2 \text{ s}^{-1}$ and $R1_+ = 30.7 \text{ s}^{-1}$ respectively.
- Dry nanosponge (DN). In this case the NMRD profile represented in figure 10(b) has $m = 44$ values with confidence intervals ranging from $\pm 0.47\%$ to $\pm 1.54\%$ of the value. The quadrupolar peaks, represented in figure 10(b), correspond to frequency values $\nu_- = 2.4991 \text{ MHz}$ and $\nu_+ = 3.1488 \text{ MHz}$ of values $R1_- = 104.85 \text{ s}^{-1}$ and $R1_+ = 104.85 \text{ s}^{-1}$ respectively.

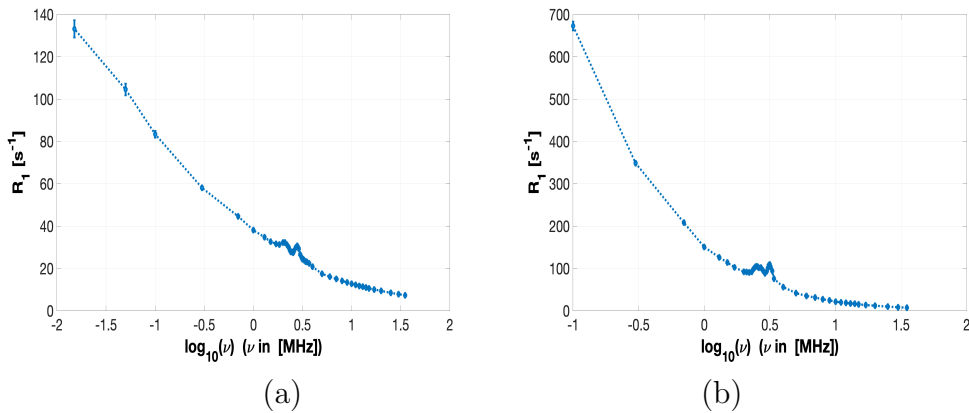


Figure 10: NMRD profiles. (a) Parmigiano Reggiano sample. (b) Dry nanosponge sample.

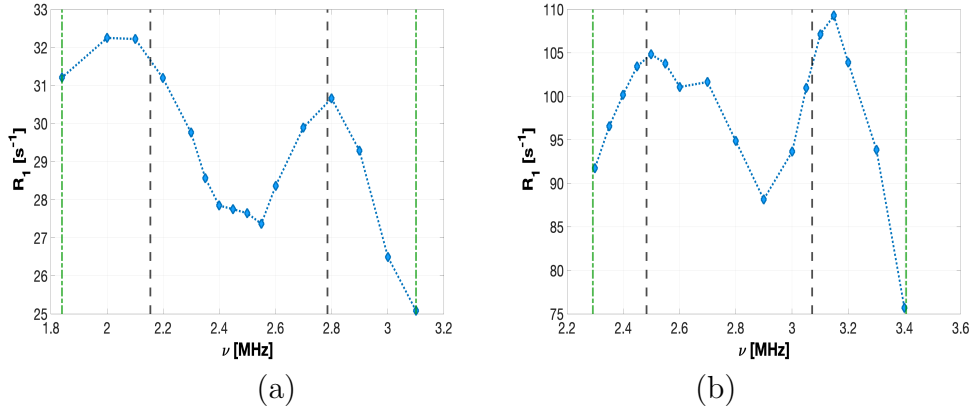


Figure 11: Zoom of quadrupolar dips. (a) Parmigiano Reggiano Cheese. (b) Dry nanosponge sample.

The proposed AURORA method has been used to compute the model parameters reported in table 7. The obtained correlation distributions are

	Parameter values	
	PR	DN
R_0	3.23	2.73
C^{NH}	5.66	69.00
Θ	1.25	0.91
Φ	0.86	0.87
τ_Q	1.02	0.74
ν_-	2.1	2.56
ν_+	2.8	3.17
MSE	$7.8887 \cdot 10^{-2}$	2.7853

Table 7: Values of the parameters fitted by AURORA and MSE in the last row.

represented in figure 12 in dark green line.

Concerning the fit of the NMRD profiles we measured the MSE reported in the last row of table 7. The fitted NMRD profiles, represented in figure 13, show in blue line the data and error bars while the fitted curves are represented in red line for both samples.

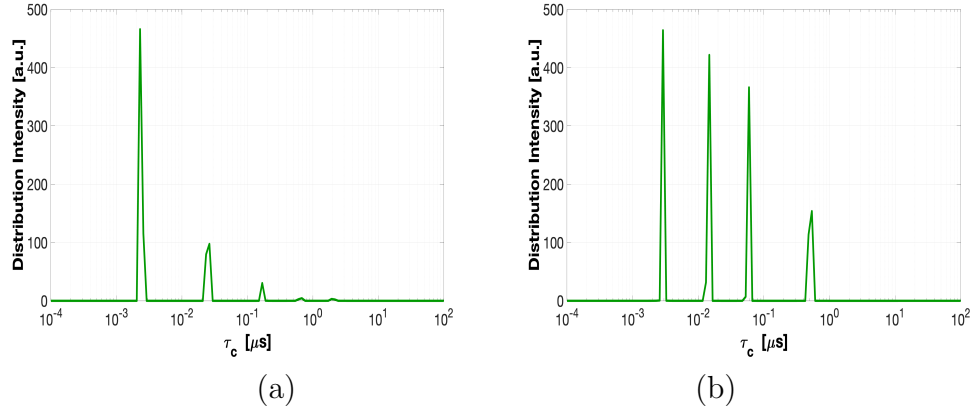


Figure 12: Correlation distribution (dark green lines). (a) Parmigiano Reggiano sample. (b) Dry nanosponge sample.

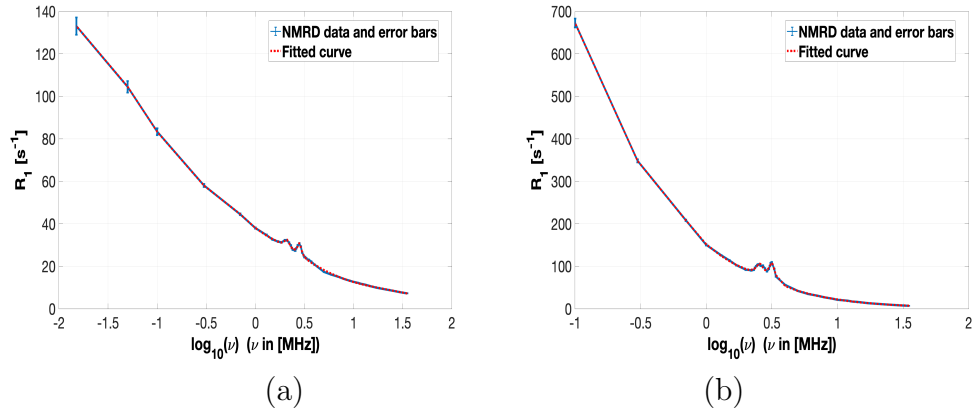


Figure 13: NMRD data and error bars (blue lines) and fitted curve (red lines). (a) Parmigiano Reggiano sample. (b) Dry nanosponge sample.

The zoom in the frequencies of QRE interval is shown in figure 14. The results confirm the excellent fit to the NMRD profile (figure 13) also in the QRE interval (figure 14).

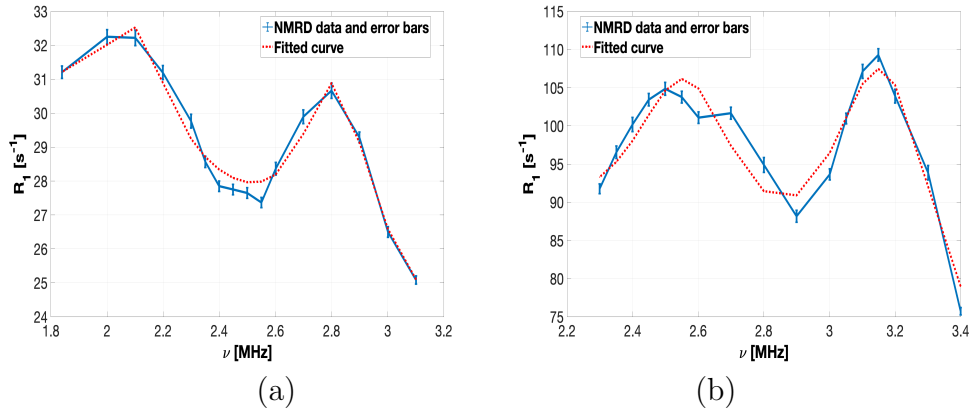


Figure 14: Zoom of data and fitted curves in the QRE intervals. NMRD data and error bars (blue lines) and fitted curve (red lines). (a) PR sample. (b) DN sample.

5. Conclusion

The present contribution investigates an automatic approach for analyzing the NMRD profiles in the presence of the quadrupolar relaxation enhancement. This feature yields a non-linear model whose parameters require the solution of a constrained non-linear least squares problem. Coupling the model-free approach and L_1 regularization, we tackle the constrained problem by a two-blocks non-linear Gauss-Seidel method. We assess the well-posedness of the optimization problem (existence of a minimum) and the convergence of the GS iterations to a critical point. Finally, we introduce an automatic convergent update rule of the regularization parameter based on the Balancing Principle.

The proposed algorithm is investigated both with synthetic and real data and the results state that it is a robust, fast approach to obtain accurate estimates of the correlation times distributions as well as modeling the quadrupolar function.

Moreover, we highlight that AURORA can be viewed as a reference framework to construct parameter estimation procedures when the model parameters can be split into independent blocks allowing the use of different computational approaches for each block. In this regard, future work will include the extension of such a framework to different models of NMRD profiles where the number of correlation times τ in (2) is assigned, and their values are to be estimated together with the corresponding component $\mathbf{f}(\tau)$.

Given the very accurate and promising results, AURORA will be included in the Matlab software tool `FreeModelFFC Tool` for the inversion of NMRD profiles with QRE (available in <https://site.unibo.it/softwareedicam/en/software>).

Acknowledgement

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