

Configuration Model Toolkit

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1 Introduction

Extended phase graphs (EPG) are indispensable for the qualitative and quantitative description of echo generation in presence of gradients. In the most common interpretation [1], the associated EPG states are motivated by non-local dephasing patterns and formally defined via voxel-scale Fourier integrals. This approach, however, impedes the proper inclusion of susceptibility effects and the quantification of echo pathways in the reconstructed signal.

We present the closely related (albeit lesser known) configuration model (CM), which does not suffer from these limitations. Relying on a microscopic representation of phase graphs, the Bloch(-Torrey) equations are solved and the transition to the voxel scale is postponed. We present a multi-dimensional CM variant, which applies to arbitrary sequences.

CoMoTk is a Matlab class, which implements the configuration model as described in section 2.

The transition to the voxel scale is discussed in section 3.

2 Microscopic Scale

2.1 Configuration Model

We consider an arbitrary (periodic or non-periodic) sequence¹ of instantaneous RF pulses² $(\alpha_\nu(\mathbf{x}), \varphi_\nu(\mathbf{x}))$, separated by time intervals of duration $\tau_{\mu(\nu)}$ during which time-dependent gradients $\mathbf{G}_\nu(t)$ may be played out. The corresponding time-dependent zero-order gradient moment vector $\mathbf{p}_\nu(t)$ is defined as usual

$$\mathbf{p}_\nu(t) := \gamma \int_0^t d\tau \mathbf{G}_\nu(\tau) \quad \mathbf{p}_\nu(\tau_{\mu(\nu)}) =: \mathbf{p}_{\mu(\nu)} \quad (1)$$

If d denotes the number of different pairs³ $(\tau_{\mu(\nu)}, \mathbf{p}_{\mu(\nu)})$ in the sequence, the function

$$\mu : \mathbb{N} \rightarrow \mathbb{N} : \nu \mapsto \mu(\nu) \in \{1, \dots, d\} \quad (2)$$

assigns a unique identifier to them. Phase encoding gradients are a typical example, when different $\mathbf{G}_\nu(t)$ have the same zero-order net gradient moment $\mathbf{p}_{\mu(\nu)}$. More generally, all $\mathbf{G}_\nu(t)$ with the same $\mu(\nu)$ differ by a balanced gradient.

A *static* spin, located at \mathbf{x} with local off-resonance frequency $\omega(\mathbf{x})$, will accumulate the phase

$$\vartheta_\nu(\mathbf{x}, t) := \omega(\mathbf{x})t - \mathbf{p}_\nu(t) \mathbf{x} \quad \vartheta_\nu(\mathbf{x}, \tau_{\mu(\nu)}) =: \vartheta_{\mu(\nu)}(\mathbf{x}) \quad (3)$$

for $0 \leq t \leq \tau_{\mu(\nu)}$.

An example of a non-periodic sequence is shown in Figure 1.

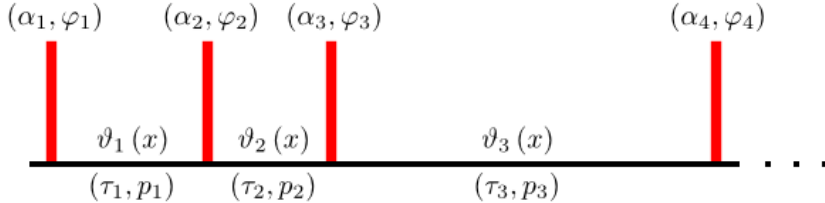


Figure 1: A non-periodic sequence

Within the configuration model of dimension d , we write the magnetization vector density $\mathbf{m}(\mathbf{x})$, immediately after any RF pulse or time interval, in the form⁴

$$\mathbf{m} =: \sum_{\mathbf{n} \in \mathbb{Z}^d} e^{i\mathbf{n}\vartheta} \mathbf{m}^{(\mathbf{n})} = \sum_{\mathbf{n} \in \mathbb{Z}^d} e^{i(\omega\tau_{\mathbf{n}} - \mathbf{p}_{\mathbf{n}}\mathbf{x})} \mathbf{m}^{(\mathbf{n})} \quad (4)$$

with

¹We use a running index $\nu = 1, 2, \dots$

²A dependence on \mathbf{x} could be caused by B_1^+ variations. Its relevance will be situational.

³ $(\tau_{\mu(\nu_1)}, \mathbf{p}_{\mu(\nu_1)}) = (\tau_{\mu(\nu_2)}, \mathbf{p}_{\mu(\nu_2)}) \Leftrightarrow \tau_{\mu(\nu_1)} = \tau_{\mu(\nu_2)} \wedge \mathbf{p}_{\mu(\nu_1)} = \mathbf{p}_{\mu(\nu_2)}$

⁴The d vector elements $\vartheta_{\mu}(\mathbf{x})$ are defined by Eq. (3). Keeping in mind that the configuration model is microscopic in nature, such that essentially everything possibly depends on the position \mathbf{x} , we will simplify the notation by dropping the argument \mathbf{x} , wherever it is not explicitly relevant.

$$\tau_{\mathbf{n}} := \sum_{\eta=1}^d n_{\eta} \tau_{\eta} \quad \text{and} \quad \mathbf{p}_{\mathbf{n}} := \sum_{\eta=1}^d n_{\eta} \mathbf{p}_{\eta} \quad (5)$$

We will refer to $\mathbf{m}^{(\mathbf{n})}(\mathbf{x})$ as *configuration vector* and \mathbf{n} as *configuration order*.

The configuration model does not rely on a specific convention for the magnetization vector. We will, however, also derive concrete expressions for actual calculations and therefore define the complex magnetization vector (density) via⁵

$$\mathbf{m} := \begin{pmatrix} (m_x + i m_y) / \sqrt{2} \\ m_z \\ (m_x - i m_y) / \sqrt{2} \end{pmatrix} =: \begin{pmatrix} m_1 \\ m_0 \\ m_{-1} \end{pmatrix} \quad (6)$$

It is related to the real magnetization density \mathbf{m}_r by a unitary transformation $\mathbf{m} = \mathbf{U} \mathbf{m}_r$ with

$$\mathbf{U} := \begin{pmatrix} 1/\sqrt{2} & i/\sqrt{2} & 0 \\ 0 & 0 & 1 \\ 1/\sqrt{2} & -i/\sqrt{2} & 0 \end{pmatrix} \quad (7)$$

A few comments on this arbitrary (and biased) choice:

- Dividing the transverse part by $\sqrt{2}$ (unlike in the common definition) prevents the magnetization from changing its magnitude under rotations like (9) below.
- The reordering and reindexing adapts to the intrinsic symmetries and simplifies the notation considerably.

2.2 Instantaneous RF pulses

We describe RF pulses as instantaneous rotations⁶

$$\mathbf{R}(\alpha_{\nu}, \varphi_{\nu}) := \mathbf{R}_z(\varphi_{\nu}) \mathbf{R}_x(\alpha_{\nu}) \mathbf{R}_z(-\varphi_{\nu}) \quad (8)$$

applied to the local magnetization density

$$\mathbf{m}_+ = \mathbf{R}(\alpha_{\nu}, \varphi_{\nu}) \mathbf{m}_- \quad (9)$$

This operation is trivially compatible with our ansatz (4), if we define

$$\mathbf{m}_+^{(\mathbf{n})} := \mathbf{R}(\alpha_{\nu}, \varphi_{\nu}) \mathbf{m}_-^{(\mathbf{n})} \quad (10)$$

In our chosen convention (6), we explicitly obtain the unitary matrices⁷

⁵CoMoTk uses this convention as well.

⁶To simplify the notation, we drop the possible dependence of flip angle and phase on the position. It can be added, if necessary.

⁷Here and in the following we use the abbreviations $c_{\beta} := \cos(\beta)$ and $s_{\beta} := \sin(\beta)$.

$$\mathbf{R}_x(\alpha) = \begin{pmatrix} (1+c_\alpha)/2 & -i s_\alpha/\sqrt{2} & (1-c_\alpha)/2 \\ -i s_\alpha/\sqrt{2} & c_\alpha & i s_\alpha/\sqrt{2} \\ (1-c_\alpha)/2 & i s_\alpha/\sqrt{2} & (1+c_\alpha)/2 \end{pmatrix} \quad (11)$$

and

$$\mathbf{R}_z(\varphi) = \begin{pmatrix} e^{i\varphi} & & \\ & 1 & \\ & & e^{-i\varphi} \end{pmatrix} \quad (12)$$

2.3 Time intervals

We distinguish cases without and with diffusion effects. For better readability, we use the abbreviation

$$\mu := \mu(\nu) \quad (13)$$

throughout this section.

Without diffusion

In this case, we have to solve the Bloch equations

$$\frac{\partial}{\partial t} \mathbf{m} = \left[i \frac{\partial \vartheta_\nu}{\partial t} \mathbf{P}_{xy} - \mathbf{T}^{-1} \right] \mathbf{m} + \mathbf{T}^{-1} \mathbf{m}_{eq} \quad (14)$$

with ϑ_ν defined as in Eq. (3), the proton density $\mathbf{m}_{eq} = m_{eq} \cdot \mathbf{e}_z$ and the definitions⁸

$$\mathbf{T} := \begin{pmatrix} T_2 & & \\ & T_1 & \\ & & T_2 \end{pmatrix} \quad \mathbf{P}_{xy} := \begin{pmatrix} 1 & & \\ & 0 & \\ & & -1 \end{pmatrix} \quad (15)$$

Since \mathbf{P}_{xy} and \mathbf{T}^{-1} commute, the formal solution can be written as

$$\mathbf{m}(t) = e^{i\vartheta_\nu(t)\mathbf{P}_{xy} - t\mathbf{T}^{-1}} \mathbf{m}(0) + (\mathbf{I}_3 - e^{-t\mathbf{T}^{-1}}) \mathbf{m}_{eq} \quad (16)$$

We now insert the ansatz (4) for $\mathbf{m}_- := \mathbf{m}(0)$ and get⁹

$$\mathbf{m}(t) = e^{i\vartheta_\nu(t)\mathbf{P}_{xy}} \sum_{\mathbf{n} \in \mathbb{Z}^d} e^{i\mathbf{n}\vartheta} \mathbf{E}(t) \mathbf{m}_-^{(\mathbf{n})} + (1 - E_1(t)) \mathbf{m}_{eq} \quad (17)$$

for $0 \leq t \leq \tau_\mu$. We introduced the common notation

⁸The specific form of the matrices depends on the notation. Here, we rely on Eq. (6).

⁹We could also write $\mathbf{R}_z(\vartheta_\mu)$ instead of $e^{i\vartheta_\mu\mathbf{P}_{xy}}$, but the latter form is better suited to equate powers in the derivation of the recursion (19).

$$\mathbf{E}(t) := e^{-t\mathbf{T}^{-1}} =: \begin{pmatrix} E_2(t) & & \\ & E_1(t) & \\ & & E_2(t) \end{pmatrix} \quad (18)$$

At the end of the interval, $\mathbf{m}_+ := \mathbf{m}(\tau_\mu)$ becomes compatible with the configuration model (4), since we then have $\vartheta_\nu(\tau_\mu) = \vartheta_\mu$, cf. Eq. (3).

We equate terms with equal¹⁰ powers $e^{i\mathbf{n}\boldsymbol{\vartheta}}$ and directly obtain the recursion for the ν^{th} interval

$$\mathbf{m}_+^{(\mathbf{n})} = \sum_{j=-1}^1 \mathbf{E}^{(j)}(\tau_\mu) \mathbf{m}_-^{(\mathbf{n}-j\mathbf{e}_\mu)} + \delta_{\mathbf{n}\mathbf{0}} \cdot (1 - E_1(\tau_\mu)) \mathbf{m}_{eq} \quad (19)$$

The longitudinal repolarization term $\propto \mathbf{m}_{eq}$ does not depend on $\boldsymbol{\vartheta}$ and therefore corresponds to $\mathbf{n} = \mathbf{0}$.

The matrix $\mathbf{E}^{(j)}$ is obtained from \mathbf{E} by setting everything but the j^{th} column¹¹ to zero. $\mathbf{e}_\mu \in \mathbb{Z}^d$ is defined by $(\mathbf{e}_\mu)_j = \delta_{j\mu}$.

With diffusion

We use the Bloch–Torrey equations

$$\frac{\partial}{\partial t} \mathbf{m}(t) = \left[i \frac{\partial \vartheta_\nu}{\partial t}(t) \mathbf{P}_{xy} - \mathbf{T}^{-1} + \nabla \mathbf{D} \nabla \right] \mathbf{m}(t) + \mathbf{T}^{-1} \mathbf{m}_{eq} \quad (20)$$

where \mathbf{D} denotes the diffusion tensor¹².

To solve the Bloch–Torrey equations (20), we try a generalization of the solution (17) of the Bloch equations

$$\mathbf{m}(t) = e^{i\vartheta_\nu(t)\mathbf{P}_{xy}} \sum_{\mathbf{n} \in \mathbb{Z}^d} e^{i\mathbf{n}\boldsymbol{\vartheta}} \mathbf{F}_\nu^{(\mathbf{n})}(t) \mathbf{E}(t) \mathbf{m}_-^{(\mathbf{n})} + (1 - E_1(t)) \mathbf{m}_{eq} \quad (21)$$

under the assumption that the only rapid spatial variations on the right hand side are caused by the gradients in $\vartheta_\nu(t)$

$$\nabla \vartheta_\nu(t) \approx -\mathbf{p}_\nu(t) \quad (22)$$

such that we only have to consider¹³

$$\begin{aligned} \nabla \mathbf{D} \nabla e^{i\vartheta_\nu(t)\mathbf{P}_{xy}} e^{i\mathbf{n}\boldsymbol{\vartheta}} &\approx \\ &- \left[\mathbf{p}_\nu^T \mathbf{D} \mathbf{p}_\nu \cdot \mathbf{I}_3 + 2 \mathbf{p}_\nu^T \mathbf{D} \mathbf{p}_\nu(t) \cdot \mathbf{P}_{xy} + \mathbf{p}_\nu^T(t) \mathbf{D} \mathbf{p}_\nu(t) \cdot \mathbf{P}_{xy}^2 \right] e^{i\vartheta_\nu(t)\mathbf{P}_{xy}} e^{i\mathbf{n}\boldsymbol{\vartheta}} \end{aligned} \quad (23)$$

We insert (21) into the Bloch–Torrey equations (20) and obtain a differential equation for the $\mathbf{F}_\nu^{(\mathbf{n})}(t)$:

¹⁰Since the equality must hold for arbitrary \mathbf{x} .

¹¹Now we benefit from the unconventional indexing, introduced in our convention (6).

¹²For isotropic diffusion, we have $\mathbf{D} = D \cdot \mathbf{I}_3$. Neglecting the spatial derivative of \mathbf{D} is consistent with Eq. (22), where we assume that gradients dominate all other spatial variations.

¹³Notation: $a := \|\mathbf{a}\|_2$

$$\frac{\partial}{\partial t} \mathbf{F}_\nu^{(\mathbf{n})}(t) = - [\mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\mathbf{n} \cdot \mathbf{I}_3 + 2 \mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\nu(t) \cdot \mathbf{P}_{xy} + \mathbf{p}_\nu^T(t) \mathbf{D} \mathbf{p}_\nu(t) \cdot \mathbf{P}_{xy}^2] \mathbf{F}_\nu^{(\mathbf{n})}(t) \quad (24)$$

The solution, which satisfies the boundary conditions $\mathbf{F}_\nu^{(\mathbf{n})}(0) \equiv \mathbf{I}_3$, reads

$$\mathbf{F}_\nu^{(\mathbf{n})}(t) = e^{-t[\mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\mathbf{n} \cdot \mathbf{I}_3 + \mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{s}_\nu(t) \cdot \mathbf{P}_{xy} + \text{tr}(\mathbf{D} \mathbf{S}_\nu(t)) \cdot \mathbf{P}_{xy}^2]} \quad (25)$$

with

$$\mathbf{s}_\nu(t) := 2t^{-1} \int_0^t d\tau \mathbf{p}_\nu(\tau) \quad \mathbf{S}_\nu(t) := t^{-1} \int_0^t d\tau \mathbf{p}_\nu(\tau) \mathbf{p}_\nu^T(\tau) \quad (26)$$

Because of (15), $\mathbf{F}_\nu^{(\mathbf{n})}(t)$ must be diagonal and we apply our indexing convention (6) to its elements

$$\mathbf{F}_\nu^{(\mathbf{n})}(t) =: \begin{pmatrix} F_{\nu,1}^{(\mathbf{n})}(t) & & \\ & F_{\nu,0}^{(\mathbf{n})}(t) & \\ & & F_{\nu,-1}^{(\mathbf{n})}(t) \end{pmatrix} \quad (27)$$

We conclude that diffusion effects modify the recursion (19) by additional damping factors

$$\mathbf{m}_+^{(\mathbf{n})} = \sum_{j=-1}^1 F_{\nu,j}^{(\mathbf{n}-j\mathbf{e}_\mu)}(\tau_\mu) \mathbf{E}^{(j)}(\tau_\mu) \mathbf{m}_-^{(\mathbf{n}-j\mathbf{e}_\mu)} + \delta_{\mathbf{n}\mathbf{0}} \cdot (1 - E_1(\tau_\mu)) \mathbf{m}_{eq} \quad (28)$$

which depend on the configuration order \mathbf{n} :¹⁴

$$F_{\nu,j}^{(\mathbf{n})}(\tau_\mu) = e^{-\tau_\mu[\mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\mathbf{n} + j \mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{s}_\nu + j^2 \text{tr}(\mathbf{D} \mathbf{S}_\nu)]} \quad (29)$$

The integrals in (29) depend on the shape of $\mathbf{G}_\nu(t)$. For the sake of simplicity¹⁵, a constant gradient is sometimes assumed, for which the moment depends linearly on time

$$\mathbf{p}_\nu(t) \equiv (t/\tau_\mu) \cdot \mathbf{p}_\mu \quad (30)$$

and the damping matrix (29) becomes

$$F_j^{(\mathbf{n})}(\tau_\mu) = e^{-\tau_\mu[\mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\mathbf{n} + j \mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\mu + j^2 \mathbf{p}_\mu^T \mathbf{D} \mathbf{p}_\mu/3]} \quad (31)$$

¹⁴ $\mathbf{s}_\nu := \mathbf{s}_\nu(\tau_\mu)$, $\mathbf{S}_\nu := \mathbf{S}_\nu(\tau_\mu)$ and $\text{tr}()$ denotes the trace.

¹⁵For large \mathbf{n} , the quadratic term, which does not depend on the gradient shape, can become dominant. The assumption (30), for which the steady-state of unbalanced SSFP has been solved analytically in [2], can be acceptable at $\text{TR} \ll T_2$, for example.

3 Voxel Scale

At first glance, the configuration model (4) is just a cumbersome and redundant way to rewrite the microscopic magnetization density $\mathbf{m}(\mathbf{x})$.

The situation changes beyond the microscopic scale, in particular for the reconstructed *voxel* m_ρ , when its true value becomes apparent.

Actually, two fundamental mechanisms are available for signal localization:

- Selective Excitation
- Spatial Encoding

Both rely on the application of gradients, effecting more or less complicated signal modulations in the vicinity of a given location \mathbf{x} . Under the assumption that other causes¹⁶ for these variations can be neglected on the voxel scale, the superposition of configurations in Eq. (4) encodes just this information via the phase factors $e^{i\mathbf{n}\boldsymbol{\vartheta}(\mathbf{x})}$.

As an example, how to apply this knowledge for both localization mechanisms, let us consider some 2D sequence with Cartesian sampling and slice selective excitation. The finite readout duration is neglected and we assume the coordinate axes to be aligned parallel to the reconstructed image, such that $x_{1,2}$ encode the locations along the orthonormal in-plane vectors $\mathbf{e}_{1,2}$ and x_3 the position along the slice normal \mathbf{e}_3 .

3.1 Spatial encoding

Due to finite sampling, the discrete reconstructed voxel signal m_ρ at position \mathbf{x}_ρ results from convolution of the transverse magnetization density $m(\mathbf{x})$ with a *point spread function* $\phi(\mathbf{x})$

$$m_\rho \propto \phi * m(\mathbf{x}_\rho) \quad (32)$$

which in our example is just a scaled sinc function

$$\phi(\mathbf{x}) := \prod_{j=1}^2 \text{sinc}\left(\frac{x_j}{\Delta x_j}\right) \quad (33)$$

where Δx_j denotes the pixel resolution in direction j .

We now insert the configuration model (4) into (32) with $f^{(\mathbf{n})} := e^{i\omega\tau_{\mathbf{n}}} m^{(\mathbf{n})}$ and get after short calculation

$$m_\rho \propto \sum_{\mathbf{n} \in \mathbb{Z}^d} \int d\mathbf{k} e^{i\mathbf{k}\mathbf{x}_\rho} \hat{f}^{(\mathbf{n})}(\mathbf{k}) \cdot \prod_{j=1}^2 u\left(\frac{\pi}{\Delta x_j} - |k_j + p_{\mathbf{n},j}|\right) \quad (34)$$

where u is the unit step function.

If the support of $\hat{f}^{(\mathbf{n})}$ in direction j is approximately¹⁷ bounded by $\pi/\Delta x_j$, only configurations with

$$|p_{\mathbf{n},j}| < \frac{2\pi}{\Delta x_j} \quad (35)$$

¹⁶Tissue properties, partial volume effects, B_0 inhomogeneity (beyond susceptibility variations), to name a few.

¹⁷This means that the reconstructed resolution recovers the essential fine structure of the sample. In a strict sense, $\hat{f}^{(\mathbf{n})}$ is not bounded at all, as it is the Fourier transform of a bounded function.

contribute to m_ρ .

Essentially, this matches the often encountered statement that crusher gradients should effect a 2π de-phasing over the voxel dimension, in order to be effective. But we also see that this is only approximately true and depends on the severity of partial volume effects.

To summarize, $\mathbf{p}_\mathbf{n}$ provides information, which configurations $m^{(\mathbf{n})}$ contribute to the reconstructed signal m_ρ .

3.2 Selective excitation

In real acquisitions, selective excitation is performed via application of a resonant \mathbf{B}_1^+ field in presence of slice encoding gradients. For example, an amplitude modulated RF pulse is usually split into a few hundred intervals of equal duration, within each of which the gradient and \mathbf{B}_1^+ are held constant¹⁸. Since rotations around different axes do not commute in general, RF pulse design treats RF pulses as a series of instantaneous small tip angle pulses interleaved by short intervals τ of constant gradient moments \mathbf{p} in direction of the slice normal \mathbf{e}_3 , cf. [3]. The accuracy of this approximate description is determined by the choice of τ .

In the configuration model, the small intervals generate a separate dimension with a unique index, say, $\mu = 1$ and the assignments $\tau_1 = \tau$ and $\mathbf{p}_1 = \mathbf{p}$.

For the reconstructed voxel m_ρ , Eq. (32) still applies in principle, since it also includes an unbounded integral along the slice normal \mathbf{e}_3 . Due to the approximate description of the RF pulse, however, the configuration model according to Eq. (4) becomes periodic in this direction

$$m(\mathbf{x}) \equiv m\left(\mathbf{x} + n \cdot \frac{2\pi}{p_1} \cdot \mathbf{e}_3\right) \quad n \in \mathbb{Z} \quad (36)$$

and the x_3 integral should therefore be restricted to the interval $[-\pi/p_1, \pi/p_1]$. To restore spin coherence across the slice, subsequent time intervals include a rephasing moment parallel to \mathbf{e}_3 . Since p_1 is rather small, it is not too restrictive, to assume that the third component of all other \mathbf{p}_μ is some integer multiple of p_1 . In this case, and in addition to restrictions due to spatial encoding, like in Eq. (35), only configuration orders \mathbf{n} with

$$p_{\mathbf{n},3} = 0 \quad (37)$$

contribute to m_ρ , since all other configurations are completely dephased in direction \mathbf{e}_3 .

If all non-zero in-plane gradient moments refer to (large enough) crusher gradients, the restrictions due to spatial encoding and selective excitation can be combined in a single statement:

$$\mathbf{p}_\mathbf{n} = \mathbf{0} \quad (38)$$

3.3 Susceptibility effects

Susceptibility related intra-voxel frequency variations $\omega_s(\mathbf{x})$, as part of the local resonance frequency $\omega(\mathbf{x})$, are typically considered as independent of gradient induced frequency modulations in the integral (32).

¹⁸VERSE pulses are an exception.

Under this assumption, the fluctuations are assumed to be distributed according to some zero-mean density $p(\omega_s)$ within the voxel. The associated damping factor depends on the configuration (via $\tau_{\mathbf{n}}$) and is given by

$$\hat{p}(\tau_{\mathbf{n}}) := \int d\omega_s e^{i\omega_s \tau_{\mathbf{n}}} \cdot p(\omega_s) \quad (39)$$

To calculate m_ρ , it has to be multiplied with $m^{(\mathbf{n})}$ in the configuration model (4) for all contributing \mathbf{n} . Most commonly, a Lorentzian distribution $p(\omega_s)$ is assumed and the damping factor $\hat{p}(\tau_{\mathbf{n}})$ takes the familiar form

$$\hat{p}(\tau_{\mathbf{n}}) = e^{-R'_2 \tau_{\mathbf{n}}} \quad (40)$$

4 Design Notes

The Matlab class `CoMoTk` implements the configuration model according to its definition in section 2. In addition to executing arbitrary sequences, various first order partial derivatives (with respect to sequence and tissue parameters) may be extracted as well.

4.1 Units

The toolkit assumes the following units throughout:

- **time:** [ms]
- **relaxation rate:** [1/ms]
- **angular frequency:** [rad/ms]
- **apparent diffusion coefficient:** [$\mu\text{m}^2/\text{ms}$]
- **angles:** [rad]
- **length, position:** [μm]
- **gradient moment, spatial frequency:** [$1/\mu\text{m}$]

4.2 Spoiler gradients

Spoiler and crusher gradients can be differentiated by their intended action. While the latter are used in sequences for de- and rephasing, the task of the former is to destroy transverse coherences reliably [4]. In principle, this goal can be (partly) reached by two effects:

- **Voxel scale:** Suppression of unwanted configurations via crusher gradients (included in $p_{\mathbf{n}}$), as outlined in section 3.1.
- **Microscopic scale:** By diffusion effects.

With respect to the second mechanism, it follows from Eq. (29) and the recursion (28) that ideal spoiling can be accomplished in the limit

$$\text{tr}(\mathbf{D} \mathbf{S}_\nu) \rightarrow \infty \quad (41)$$

since then we have

$$\mathbf{F}^{(\mathbf{n})}(\tau_\mu) \rightarrow \begin{pmatrix} 0 & & \\ & e^{-\tau_\mu \mathbf{p}_\mathbf{n}^T \mathbf{D} \mathbf{p}_\mathbf{n}} & \\ & & 0 \end{pmatrix} \quad (42)$$

and transverse magnetization is eliminated in all configuration orders.

Note that the condition (41) does not necessarily¹⁹ require $p_\mu \rightarrow \infty$ (or $p_\mathbf{n} \rightarrow \infty$). On the other hand, $p_\mu \rightarrow \infty$ will usually require²⁰ (41) as well.

4.3 General properties of occupied configurations

Proposition 1. *In absence of magnetization transfer and for $T_2 \leq T_1$, we always have*

$$\sum_{\mathbf{n}} \left\| \mathbf{m}^{(\mathbf{n})} \right\|_2^2 \leq m_{eq}^2 \quad (43)$$

Proof. The RF pulses do not affect $\left\| \mathbf{m}^{(\mathbf{n})} \right\|$. Therefore we only need to consider the time intervals and the proof can be done by induction. From (28), we derive

$$\left| m_{+,j}^{(\mathbf{n})} \right| = \begin{cases} \left| F_{\nu,0}^{(0)} E_{00} m_{-,0}^{(0)} + (1 - E_1) m_{eq} \right| & : \mathbf{n} = \mathbf{0} \wedge j = 0 \\ \left| F_{\nu,j}^{(\mathbf{n} - j\mathbf{e}_\mu)} E_{jj} \right| \left| m_{-,j}^{(\mathbf{n} - j\mathbf{e}_\mu)} \right| & : \text{else} \end{cases} \quad (44)$$

In combination with $T_2 \leq T_1$ and assuming that (43) holds for $\mathbf{m}_-^{(\mathbf{n})}$, we therefore obtain

$$\begin{aligned} \sum_{\mathbf{n}} \left\| \mathbf{m}_+^{(\mathbf{n})} \right\|_2^2 &\leq E_1^2 \cdot \sum_{\mathbf{n}} \left\| \mathbf{m}_-^{(\mathbf{n})} \right\|_2^2 + 2 E_1 (1 - E_1) \left| m_{-,0}^{(0)} \right| m_{eq} + (1 - E_1)^2 m_{eq}^2 \\ &\leq \left(E_1^2 + 2 E_1 (1 - E_1) + (1 - E_1)^2 \right) m_{eq}^2 \\ &= m_{eq}^2 \end{aligned} \quad (45)$$

which completes the proof. \square

4.4 Storage considerations

The actual state of the configuration is defined by the set $\{\mathbf{n}\}$ of occupied configurations and their associated configuration vectors $\mathbf{m}^{(\mathbf{n})}$.

The size of the set, n_c , will increase with every time interval

$$n_c \rightarrow \begin{cases} n_c + 2d & : d \rightarrow d \\ 3n_c & : d \rightarrow d + 1 \end{cases} \quad (46)$$

¹⁹Strong balanced gradients with $p_\mu = 0$ are a simple counterexample. In the specific case (30), however, the conditions (41) and $p_\mu \rightarrow \infty$ are equivalent.

²⁰More accurately, if we replace “ ∞ ” by “large”, this follows from the actual limitations of gradient hardware.

The first case applies, if the same time interval μ has been played out before²¹, whereas the second case applies for new intervals²².

Eq. (46) implies that n_c becomes particularly large for high dimensional configuration models. The most extreme scenario is a fingerprinting-type excitation pattern with distinct random intervals τ_μ . This pattern generates $n_c = 3^n$ occupied configurations after $n = d$ time intervals, easily exceeding the memory of actual computer hardware after less than about $n \approx 20$ intervals.

Fortunately, due to relaxation, magnetization forgets about its initial state with the consequence that many (if not most) of the occupied configuration vectors $\mathbf{m}^{(n)}$ may be safely ignored. Formally, this immediately follows from the upper bound on $\sum_n \|\mathbf{m}^{(n)}\|_2^2$, derived in Proposition 1.

Based on some given, user-defined limit ε , we will therefore repeatedly discard all configuration vectors with $\|\mathbf{m}^{(n)}\|_2 < \varepsilon$. To this end, we invoke the method `meltdown()` after every time interval.

For this to work properly, the actual set $\mathbf{m}^{(n)}$ is stored in a two-dimensional array `m` of size²³ $3 \times n_a$, where n_a specifies the allocated (not necessarily occupied) space. This storage concept requires a considerable amount of bookkeeping, which is described in more detail in section 4.5.

To improve performance further, reallocations²⁴ and redundant computations²⁵ are minimized as well.

4.5 Bookkeeping

The `n_conf` stored configuration orders \mathbf{n} of dimension `d` are collected in the $n_a \times \mu_a$ array `n`. The associated location in the allocated storage is determined by the logical $n_a \times 1$ array `b_n` and the $1 \times \mu_a$ array `b_mu`.²⁶

The `d` distinct dimensions μ are stored in the $1 \times \mu_a$ array `mu`.

Prior to and after the ν^{th} time interval, we denote the set of *stored* configurations \mathbf{n} by S_ν^- and S_ν^+ , respectively. To implement the time intervals properly, particularly recursions like (28), it is crucial to guarantee that S_ν^\pm does not contain any holes: If $\mathbf{n} \in S_\nu^\pm$ and $\mathbf{n} + c \cdot \mathbf{e}_\mu \in S_\nu^\pm$ for some $\mathbf{n} \in \mathbb{Z}^d$ and $c \in \mathbb{Z} > 0$, we must have $\mathbf{n} + b \cdot \mathbf{e}_\mu \in S_\nu^\pm$ for every $1 \leq b \in \mathbb{Z} < c$.

Since RF pulses do not change the set of occupied configurations, we set $S_{\nu+1}^- = S_\nu^+$.

For each $\mathbf{n} \in S_\nu^-$ and each dimension μ , the information whether $\mathbf{n} \pm \mathbf{e}_\mu \in S_\nu^-$ is fulfilled, is stored in a separate variable.²⁷ In case of $\mathbf{n} \pm \mathbf{e}_\mu \in S_\nu^-$, the corresponding location in memory must be available as well.²⁸

Due to the method `meltdown()`, the `n_conf` $\leq n_c$ stored configurations are some subset of \mathbb{Z}^d with²⁹ $d \leq d$. We demand that the stored set remains connected along each dimension, i.e. free of holes, as described above. We also demand that $\mathbf{n} = \mathbf{0}$ is included at all times.

²¹In this case, the dimension of the configuration model, d remains unchanged.

²²Here, the dimension d is increased by 1.

²³The first dimension refers to the three vector components.

²⁴Mainly achieved via a sufficiently large initial value of n_a .

²⁵Some quantities, which are repeatedly needed in calls of identical RF pulses or time intervals are stored for reuse. When possible, updates are limited to newly entering configurations.

²⁶Obviously, `sum(b_n)` and `sum(b_mu)` must be equal to `n_conf` and `d`, respectively.

²⁷We define two logical $n_a \times \mu_a$ arrays `b_up_free` and `b_do_free` as follows: For each stored configuration \mathbf{n} and dimension μ , the associated entry in `b_up_free` is `true`, if the direct neighbor $\mathbf{n} + \mathbf{e}_\mu$ is *not* stored and `false` otherwise. Similarly, `b_do_free` is `true`, if $\mathbf{n} - \mathbf{e}_\mu$ is *not* stored and `false` otherwise.

²⁸Similarly, we define two $n_a \times \mu_a$ arrays `idx_up` and `idx_do` as follows: For each stored configuration \mathbf{n} and dimension μ , with the associated entry in `b_up_free` equal to `false`, the corresponding value `idx_up` gives the index (with respect to the first dimension of size n_a) of the direct neighbor $\mathbf{n} + \mathbf{e}_\mu$. For the other direction, $\mathbf{n} - \mathbf{e}_\mu$, the array `idx_do` is defined in complete analogy.

²⁹A populated dimension μ , for which no configuration orders $n_\mu \neq 0$ are stored, is de facto nonexistent and can be discarded. This can, for example, happen, if the associated time period τ_μ has not been played out for a long time.

5 Usage

To familiarize with the toolkit, the scripts in the `test` and `examples` folders are recommended as a good starting point. Following, a brief overview of the basic functionality.

5.1 Initialize CoMoTk

First, we need to create an instance of `CoMoTk`:

```
cm = CoMoTk;
```

Next, we have to setup a few mandatory tissue parameters. In the simplest case of a 1-peak model without diffusion, this can look like this:

```
cm.R1 = 0.01;      % longitudinal relaxation rate
cm.R2 = 0.1;       % transverse relaxation rate
cm.D = 0;          % apparent diffusion coefficient
```

It is also possible, to specify more complex n-peak models with variable relative weighting (\propto proton density), chemical shift and diffusivity. Here, a 2-peak example:

```
cm.R1 = [ 0.01, 0.02 ];
cm.R2 = [ 0.1, 0.2 ];
cm.D = [ 3, 1.5 ];
cm.w = [ 0.6, 0.4 ]; % relative weight (does not need to add to 1)
cm.dom = [ 0, -0.1 ]; % chemical shift (angular frequency)
```

Setting `w` and `dom` is optional. If not set, they default to 1 and 0, respectively.

Another optional variable is the relative B_1^+ field (default = 1):

```
cm.B1 = 0.8;
```

There are further options, for which it is possible to modify the default settings. The most important one is the desired accuracy. A non-zero value of `epsilon` is set, if the number of stored configurations needs to be restricted, e.g. due to memory or performance restrictions.

```
options = cm.options; % get default options

options.alloc_d = 3; % allocated number of dimensions
options.alloc_n = 10000; % allocated number of configurations
options.epsilon = 0; % for maximal accuracy (enough memory needed)
options.verbose = true; % for more output
options.debug = true; % for debugging purposes (look into CoMoTk.m)

cm.options = options; % activate new options
```

Now we have to specify the initial configuration vector, corresponding to $\mathbf{n} = 0$. It is supplied as a real vector (row or column) in the usual convention (m_x, m_y, m_z):

```
cm.init_configuration ( [ 0; 0; 1 ] ); % longitudinal magnetization
```

If we want CoMoTk to calculate specific derivatives as well, we can do this now. A full example, involving every possible variable, looks like this:

```
cm.set_derivatives ( ...
    'R1', [ 1, 3 ], ... % tissue handles (n-peak model)
    'R2', 2, ... % for a 1-peak model the handle (= 1)
    'D', [ 2, 3 ], ... % must be supplied as well
    'B1', ... % only B1 has no handle
    'FlipAngle', [ 2, 3 ], ... % non-tissue handles can be chosen
    'Phase', [ 2, 1004, 12 ], ... % arbitrarily
    'tau', [ 3, 4, 6 ], ... % time interval (= \mu)
    'p', [ 1, 2, 4 ], ... % gradient moment (see below)
    's', [ 4, 7 ] ... % gradient shape (see below)
);
```

Of course, only the desired subset of parameter/handle combinations needs to be supplied.

After having initialized everything, we proceed with the actual sequence. Here, we apply instantaneous RF pulses and time intervals, typically in an alternate fashion.³⁰

5.2 RF pulse

Calling an RF pulse is as simple as:

```
cm.RF( flip , phase ); % RF pulse with flip angle and phase [rad]
```

If partial derivatives with respect to flip angle(s) and/or phase(s) have been set before, the associated handles can be additionally supplied according to the following format:³¹

```
cm.RF( flip , phase , 'FlipAngle', flip_handle , 'Phase', phase_handle );
```

5.3 Time interval

Whether the time derivatives are needed or not, executing the time interval always needs specification of the (otherwise arbitrary) index $\mu := \mu$, which is defined as in section 2. At least in the first call of each distinct interval, the duration $\tau_\mu := \tau_\mu$ must be supplied also:

³⁰There are no restrictions, though, i.e. multiple subsequent RF pulses or time intervals are allowed as well.

³¹If only the flip angle derivative is needed, the phase handle is not required (and vice versa). The user is responsible for the validity of value/handle combinations. Otherwise, the result is unpredicted.

```
cm.time( mu, 'tau', tau );
```

If we also require the gradient moment $\mathbf{p} := \mathbf{p}_\mu$ for the simulations (diffusion effects) or the results (e.g. slice profile), we have to add this parameter as well:³²

```
cm.time( mu, 'tau', tau, 'p', p );
```

If we include the influence of gradient shape(s) on diffusion damping, the shape $\mathbf{s} := (\mathbf{s}_\mu^{(1)}, \mathbf{s}_\mu^{(2)})$ must be added too:³³

```
cm.time( mu, 'tau', tau, 'p', p, 's', s );
```

In later calls, only the handle is required:³⁴

```
cm.time( mu );
```

5.4 Get results

After any RF pulse or time interval, we can extract the current state like this:

```
iso = cm.isochromat( om, x, b_n );
```

`om` and `x` respectively refer to the local angular off-resonance frequency $\omega(\mathbf{x})$ and the position \mathbf{x} according to the definition (3). If the position is not needed, e.g. in an idealized simulation without specifying gradient moments, just set `x = []`.

`b_n` selects a subset from the set of stored configuration orders, S_ν^\pm , to be included in the result. Setting `b_n = []` calculates the whole sum in Eq. (4). For a restriction, the `find` method can be used to generate `b_n`:

```
b_n = cm.find( mu, n );
```

If `mu` and `n` are scalars, we have $\mathbf{b}_n = \{\mathbf{n} \in S_\nu^\pm : n_{\text{mu}} = \mathbf{n}\}$. If no matching configurations are found, `b_n = []` is returned.

`mu` and `n` can also be 1d-arrays of equal length. This can be used as a shorthand for a combination with the `|` operator. For example, we obtain for arrays of length 3 a result equivalent to

³²`p` must be a 3×1 or 1×3 array. If we set an element of `p` equal to `Inf`, it is interpreted as an ideal spoiler, as defined in section 4.2.

³³`s` must be a 4×1 or 1×4 array.

³⁴The full version is also allowed. Of course, `s` still needs to be supplied, if the shape is variable.

```

b_n = ...
cm.find( mu( 1 ), n( 1 ) ) | ...
cm.find( mu( 2 ), n( 2 ) ) | ...
cm.find( mu( 3 ), n( 3 ) );

```

If we want to single out a specific configuration $\mathbf{n} \in S_\nu^\pm$, we have to use the `&` operator explicitly. For example, we get for $d = 3$:

```

b_n = ...
cm.find( cm.mu( 1 ), n( 1 ) ) & ...
cm.find( cm.mu( 2 ), n( 2 ) ) & ...
cm.find( cm.mu( 3 ), n( 3 ) );

```

It is also possible, to restrict the mask `b_n` directly. For example, to extract all stored configurations, which satisfy $\mathbf{p}_n = \mathbf{0}$, cf. Eq. (38), one could use

```

b_n = cm.b_n & reshape( ~any( cm.p_n ), size( cm.b_n ) );

```

References

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