Supporting information for Protein Dynamics from Nuclear Magnetic Relaxation

Cyril Charlier, $\ddagger a, b, c$ Samuel F. Cousin, $\ddagger a, b, c$ and Fabien Ferrage $\ast a, b, c$

1 Correlation function and spectral density function

We chose a model-free approach to describe motions in liquid and solid phases.

1.1 Correlation function and spectral density function found in liquid

The correlation function used for the global tumbling $C_0(t)$ is classically taken as:

$$\mathcal{C}_0(t) = e^{-\frac{t}{\tau_c}} \tag{S. 1}$$

The correlation function, for the model-free (MF) and the extended model-free (EMF) approaches, has been taken as:

$$C_{liq}^{MF}(t) = C_0(t)C_f(t)$$
(S. 2)

$$C_{liq}^{EMF}(t) = C_0(t)C_f(t)C_s(t)$$
(S. 3)

with:

$$C_i(t) = S_i^2 + (1 - S_i^2)e^{-\frac{t}{\tau_i}}$$
(S. 4)

where i stands for f (fast motions) or s (slow motions). The corresponding spectral density function for the MF and EMF approaches are:

$$J_{liq}^{MF}(\omega) = \frac{2}{5} \left((1 - S_f^2) \frac{\tau_{fe}}{1 + (\omega \tau_{fe})^2} + S_f^2 \frac{\tau_c}{1 + (\omega \tau_c)^2} \right)$$
(S. 5)

$$J_{liq}^{EMF}(\omega) = \frac{2}{5} \left(S^2 \frac{\tau_c}{1 + (\omega \tau_c)^2} + (1 - S_f^2) \frac{\tau_{fe}}{1 + (\omega \tau_{fe})^2} + (S_f^2 - S^2) \frac{\tau_{se}}{1 + (\omega \tau_{se})^2} \right)$$
(S. 6)

with $\tau_{ie}^{-1} = \tau_i^{-1} + \tau_c^{-1}$, the effective correlation time for the motion, with the order of parameter S_i^2 and

$$S^2 = S_f^2 S_s^2$$
 (S. 7)

^a École Normale Supérieure-PSL Research University, Département de Chimie, 24, rue Lhomond, 75005 Paris, France. Fabien.Ferrage@ens.fr

^b Sorbonne Universités, UPMC Univ Paris 06, LBM, 4 place Jussieu, F-75005, Paris, France

^c CNRS, UMR 7203 LBM, F-75005, Paris, France.

1.2 Correlation function and spectral density function found in solids

Correlations functions in solids are characterized by the absence of global tumbling

$$C_{solid}^{MF}(t) = C_f(t) \tag{S.8}$$

$$C_{solid}^{EMF}(t) = C_f(t)C_s(t)$$
(S. 9)

Only the fluctuating part of the interaction contributes to relaxation (the non-fluctuating part corresponds to a coherent interaction). The spectral density function is defined as the Fourier transform of the decaying part of the correlation function.

$$J_{solid}^{MF}(\boldsymbol{\omega}) = \frac{2}{5} \left((1 - S_f^2) \frac{\tau_f}{1 + (\boldsymbol{\omega}\tau_f)^2} \right)$$
(S. 10)

$$J_{solid}^{EMF}(\omega) = \frac{2}{5} \left((S_s^2 - S^2) \frac{\tau_f}{1 + (\omega \tau_f)^2} + (S_f^2 - S^2) \frac{\tau_s}{1 + (\omega \tau_s)^2} (1 - S_f^2 - S_s^2 + S^2) \frac{\tau_p}{1 + (\omega \tau_p)^2} \right)$$
(S. 11)

in which S^2 is defined as Eq. S. 7 and,

$$\tau_p^{-1} = \tau_s^{-1} + \tau_f^{-1} \tag{S. 12}$$

In the case of methyl groups, we used a two correlation times model with angular constraint already present in the literature [1]:

$$J_{Met}^{MF}(\omega, \theta_{i,j}) = \frac{1}{5} \left((1 - \alpha S_f^2) \frac{\tau_{fe}}{1 + (\omega \tau_{fe})^2} + \alpha S_f^2 \frac{\tau_c}{1 + (\omega \tau_c)^2} \right)$$
(S. 13)

where α depends on the angle $\theta_{i,j}$ between the rotation axis of the methyl group and the principal axis of the interaction as:

$$\alpha = \left(\frac{3\cos^2[\theta_{i,j}] - 1}{2}\right)^2 \tag{S. 14}$$

Relaxation rates for a selection of spin systems 2

2.1 Amplitudes of the interactions

We first define the dipolar constant δ_{ij} , the chemical shift anisotropy (CSA) constant Δ_i and the quadrupolar constant Ξ by:

$$\delta_{ij} = \frac{\mu_0 \hbar \gamma_i \gamma_j}{4\pi r_{i,j}^3} \tag{S. 15a}$$

$$\Delta_i = \frac{\Delta \sigma_i \gamma_i B_0}{3} \tag{S. 15b}$$

[1] Kay, JACS Vol 124 NO 22, 2002

1–7

2 |

$$\Xi = \frac{e^2 q Q}{\hbar}$$
(S. 15c)

where μ_0 is the permeability of free space, \hbar is the reduced Planck constant, γ_i is the gyromagnetic ratio of the nucleus i, $r_{i,j}$ is the distance between the two nucleis i and j, B_0 is the magnetic field, $\Delta\sigma$ the chemical shift anisotropy (we assume a axially symmetrical tensor), eQ is the nuclear quadrupole moment, and eq the electric field gradient along its principal axis.

2.2 Nitrogen-15 relaxation rates in an ¹⁵N¹H group

The longitudinal relaxation rate of nitrogen-15 in a NH group is:

$$\mathcal{R}_{I}(\boldsymbol{\omega}) = \frac{1}{4} \delta_{NH}^{2} \left(3J[\boldsymbol{\omega}_{N}] + J[\boldsymbol{\omega}_{N} - \boldsymbol{\omega}_{H}] + 6J[\boldsymbol{\omega}_{N} + \boldsymbol{\omega}_{H}] \right) + \Delta_{N}^{2} J[\boldsymbol{\omega}_{N}]$$
(S. 16)

The transverse relaxation rate is:

$$\mathcal{R}_{2}(\omega) = \frac{1}{8} \delta_{NH}^{2} \left(4J[0] + 3J[\omega_{N}] + J[\omega_{N} - \omega_{H}] + 6J[\omega_{H}] + 6J[\omega_{N} + \omega_{H}] \right)$$
(S. 17)
+ $\frac{1}{2} \Delta_{N}^{2} \left(3J[0] + 2J[\omega_{N}] \right)$

In our simulation we took $J(\omega) = J_{liq}^{MF}(\omega)$ (See Eq. S. 5) and the other parameters are given in Table 1.

σ_N	$r_{N,H}$	τ_c	S	$ au_f$
175 ppm	1.02 Å	5 or 10 ns	0.8	100 ps

Table 1 : Parameters used to calculate nitrogen-15 for relaxation rates in ¹⁵N¹H group.

2.3 Carbon-13 relaxation rates in a ¹³C¹H₃ group:

To calculate the relaxation rates of ${}^{13}C$ in a ${}^{13}C^{1}H_{3}$ group, four interactions have been considered: dipolar interactions with all three protons and the CSA of the carbon-13 nucleus. The angle ϕ lies between the rotation axis of the methyl group and each CH bond. The longitudinal relaxation rate is:

$$\mathcal{R}_{I}(\omega) = \frac{3}{4} \delta_{CH}^{2} \left(3J[\omega_{C}, \phi] + J[\omega_{C} - \omega_{H}, \phi] + 6J[\omega_{C} + \omega_{H}, \phi] \right) + \Delta_{N}^{2} J[\omega_{C}, 0]$$
(S. 18)

The transverse relaxation rate is:

$$\mathcal{R}_{2}(\omega) = \frac{3}{24} \delta_{CH}^{2} (4J[0,\phi] + 3J[\omega_{C},\phi] + J[\omega_{C} - \omega_{H},\phi] + 6J[\omega_{H},\phi] + 6J[\omega_{C} + \omega_{H},\phi])$$
(S. 19)
+ $\frac{1}{4} \Delta_{N}^{2} (4J[0,0] + 3J[\omega_{C},0])$

In this equation, the spectral density function is J_{Met}^{MF} (see Eq S. 13). We considered an ideal geometry for the methyl group, and we oriented the CSA vector along the CC axis.

τ_c	S_f^2	$ au_f$	σ_C	$r_{C,H}$	¢
5 or 10 ns	0.6	50 ns	25 ppm	1.115 Å	109.47°

Table 2 : Parameters used to calculate carbon-13 relaxation rates in a ${}^{13}C^{1}H_{3}$ group

2.4 Deuterium relaxation rates in a ${}^{13}C^2H^1H_1$ group

Three dipolar interactions and one quadrupolar interaction (CSA negligeable) contribute to the relaxation rate of deuterium in ${}^{13}C^{1}H_{2}{}^{2}H$ group. The longitudinal relaxation rate is:

$$\mathcal{R}_{J}(\omega) = \frac{1}{4} (\delta_{DC}^{2} (3J[\omega_{D}, \theta_{C,D}] + J[\omega_{C} - \omega_{D}, \theta_{C,D}] + 6J[\omega_{C} + \omega_{D}, \theta_{C,D}]) + 2\delta_{DH}^{2} (3J[\omega_{D}, \theta_{H,D}] + J[\omega_{H} - \omega_{D}, \theta_{H,D}] + 6J[\omega_{H} + \omega_{D}, \theta_{H,D}]))$$
(S. 20)
$$+ \frac{1}{40} \Xi^{2} (12J[\omega_{D}, \theta_{Quad}]) + 3J[\omega_{D}, \theta_{Quad}])$$

The transverse relaxation rate is:

$$\mathcal{R}_{2}(\omega) = \frac{2}{8} \delta_{CD}^{2} (4J[0, \theta_{C,D}] + 3J[\omega_{D}, \theta_{C,D}] + J[\omega_{C} - \omega_{D}, \theta_{C,D}] + 6J[\omega_{C}, \theta_{C,D}] + 6J[\omega_{C} + \omega_{D}, \theta_{C,D}]) + \frac{1}{8} \delta_{HD}^{2} (4J[0, \theta_{H,D}] + 3J[\omega_{D}, \theta_{H,D}] + J[\omega_{D} - \omega_{H}, \theta_{H,D}] + 6J[\omega_{H}, \theta_{H,D}] + 6J[\omega_{D} + \omega_{H}, \theta_{H,D}]) + \frac{1}{80} \Xi (9J[0, \theta_{Quad}] + 15J[\omega_{D}, \theta_{Quad}] + 6J[2\omega_{D}, \theta_{Quad}])$$
(S. 21)

 $J(\omega, \theta_{i,j})$ is depending between the angle between the principal axis of the interaction and the rotation axis of the methyl group (see Eq. S. 13). The principal axis of the quadrupolar interaction is considered aligned with the CC bond.

τ_c	S_f^2	$ au_f$	σ_C	$\Xi/2\pi$	$r_{C,D}$	$r_{D,H}$	θ_{Quad}	$\theta_{H,D}$	$\theta_{C,D}$
5 or 10 ns	0.6	100 ps	25 ppm	2167.10 ³ Hz	1.110 Å	1.115 Å	0°	90°	109.47°

Table 3 : Parameters used to calculate deuterium relaxation rates in a ${}^{13}C^2H^1H_1$ group

2.5 Carbonyl carbon-13 relaxation rates:

We considered only the CSA of the carbonyl and the dipolar interactions with ${}^{13}C_{\alpha}$, ${}^{1}H_{\alpha}$, ${}^{15}N$ and the ${}^{1}H^{N}$ nuclei.

The longitudinal relaxation rate is:

$$\mathcal{R}_{I}(\omega) = \frac{\delta_{CC}}{4} (J[0] + 3J[\omega_{C}] + 6J[2\omega_{C}]) + \frac{\delta_{CH_{\alpha}}}{4} (J[\omega_{C} - \omega_{H}] + 3J[\omega_{C}] + 6J[\omega_{H} - \omega_{C}]) + \frac{\delta_{CN}}{4} (J[\omega_{C} - \omega_{N}] + 3J[\omega_{C}] + 6J[\omega_{N} - \omega_{C}]) + \frac{\delta_{CH^{N}}}{4} (J[\omega_{C} - \omega_{H}] + 3J[\omega_{C}] + 6J[\omega_{H} - \omega_{C}]) + \Delta_{C} J[\omega_{C}]$$
(S. 22)

The transverse relaxation rate is:

$$\begin{aligned} \mathcal{R}_{2}(\omega) &= \frac{\delta_{CC}}{8} (5J[0] + 9J[\omega_{C}] + 6J[2\omega_{C}]) \\ &+ \frac{\delta_{CH_{\alpha}}}{8} (4J[0] + J[\omega_{C} - \omega_{H}] + 3J[\omega_{C}] + 3J[\omega_{H}] + 6J[\omega_{H} - \omega_{C}]) \\ &+ \frac{\delta_{CN}}{8} (4J[0] + J[\omega_{C} - \omega_{N}] + 3J[\omega_{C}] + 3J[\omega_{N}] + 6J[\omega_{N} - \omega_{C}]) \\ &+ \frac{\delta_{CH^{N}}}{8} (4J[0] + J[\omega_{C} - \omega_{H}] + 3J[\omega_{C}] + 3J[\omega_{H}] + 6J[\omega_{H} - \omega_{C}]) \\ &+ \frac{\Delta_{C}}{6} (4J[0] + 3J[\omega_{C}]) \end{aligned}$$
(S. 23)

A model-free spectral density function has been chosen, without angular constraint and the following parameters have been used to give evaluate this relaxation rate.

τ_c	S_f^2	τ_f	$\Delta \sigma_C$	$r_{C,N}$	r_{C,H_N}	$r_{C,C_{\alpha}}$	$r_{C,H_{\alpha}}$
5 or 10 ns	0.85	50 ps	140 ppm	1.33 Å	2.0 Å	1.52 Å	2.1 Å

Table 4 : Parameters used to calculate carbonyl carbon-13 relaxation rates

3 Chemical exchange simulations

Spectra in the presence of chemical exchange (Figure 7.d-m) were simulated with a simple integration of Bloch-McConnell equations. Chemical exchange saturation transfer (CEST) and dark-state exchange saturation transfer (DEST) saturation profiles (Figure 8.d-f) were obtained using an integration of Bloch-McConnell equations in the presence of a weak radio-frequency field. The exchange constant was $k_{ex} = 200 \text{ s}^{-1}$; with $p_B = 2\%$ (Figure 8-d) or $p_B = 10\%$ (Figure 8-f) and a static magnetic field strength of 18.7 T. CEST Profiles were simulated with an amplitude of radiofrequency fields $v_1 =$ 25 Hz (d-e) while DEST profiles were simulated with three different radiofrequency fields.

A classical model-free spectral density function was used (see Equ. 5) to describe intrinsic relaxation rates. Simulations of CEST profiles (Figure 8.d-e) were carried out with an overall correlation time τ_c = 7 ns for both ground and excited states, an internal correlation time $\tau_f = 100$ ps and an order parameter $S_f^2 = 0.9$. DEST profiles were simulated with $\tau_c = 7$ ns for the ground state, $\tau_c = 10 \ \mu$ s in the excited state, parameters of internal dynamics were $\tau_f = 100$ ps and $S^2 = 0.9$ in both states. Proton decoupling in simulations was achieved with an on-resonance continuous wave radiofrequency irradiation with an amplitude vH = 1 kHz.

4 Further readings

Relaxation theory

- Goldman, M. Quantum Description of High Resolution NMR in Liquids Chap. 8. Clarendon Press, 1988, Oxford.
- Goldman, M. J. Magn. Reson., 2001, 149, 160-187.
- Luginbuhl, P., and K. Wuthrich. Prog. Nucl. Magn. Reson. Spectrosc., 2002, 40, 199-247.
- Cavanagh, J., W. J. Fairbrother, A. G. Palmer III, M. Rance, and N. J. Skelton. Protein NMR Spectroscopy: Principles and practice. Academic Press, 2006, San Diego.
- Nicholas, M. P., E. Eryilmaz, F. Ferrage, D. Cowburn, and R. Ghose. Prog. Nucl. Magn. Reson. Spectrosc., 2010, 57,111-158.

Reorientational motions in water and biomolecules

- L. D. Favro, Phys. Rev., **1960**, *119*, 53-62.
- J. R. Brainard and A. Szabo, Biochemistry, **1981**, 20, 4618-4628.
- F. Ochsenbein, J. M. Neumann, E. Guittet and C. Van Heijenoort, Protein Sci., 2002, 11, 957-964.
- D. Laage and J. T. Hynes, Science, **2006**, *311*, 832-835.

Dynamic interpretation of high-field measurements of relaxation

- R. Ishima and K. Nagayama, J. Magn. Reson., Ser B, 1995, 108, 73-76.
- D. Fushman, S. Cahill and D. Cowburn, J. Mol. Biol., 1997, 266, 173194.
- M. Bieri, E. J. d'Auvergne and P. R. Gooley, J. Biomol. NMR, **2011**, *50*, 147-155.

Relaxometry

• C. Luchinat and G. Parigi, J. Am. Chem. Soc., 2007, 129, 1055-1064.

1–7

6 |

Relaxation in the solid state

• J. M. Lamley, M. J. Lougher, H. J. Sass, M. Rogowski, S. Grzesiek and J. R. Lewandowski, PCCP, 2015, *17*, 21997-22008.

Translational dynamics: introduction

• P. H. Fries and G. N. Patey, J. Chem. Phys., 1984, 80, 6253-6266

Highlighting transient proximity

- M. M. Dedmon, K. Lindorff-Larsen, J. Christodoulou, M. Vendruscolo and C. M. Dobson, J. Am. Chem. Soc., 2005, *127*, 476-477.
- C. Tang, J. Iwahara and G. M. Clore, Nature, 2006, 444, 383-386.
- J. Iwahara and G. M. Clore, Nature, 2006, 440, 1227-1230.
- C. Tang, C. D. Schwieters and G. M. Clore, Nature, 2007, 449, 1078-U1012.

Chemical reactions at equilibrium: the example of proton exchange

• J. Iwahara, Y.-S. Jung and G. M. Clore, J. Am. Chem. Soc., 2007, 129, 2971-2980.

CEST

• S. Forsen and R. A. Hoffman, J. Chem. Phys., 1963, 39, 2892.

Carr-Purcell-Meiboom-Gill Dispersion (historic references)

- H. Y. Carr and E. M. Purcell, Physical Review, 1954, 94, 630-638.
- S. Meiboom and D. Gill, Rev. Sci. Instrum., 1958, 29, 688-691.

Chemical exchange by relaxation dispersion: applications

- M. Akke and A. G. Palmer, J. Am. Chem. Soc., 1996, 118, 911-912.
- R. B. Hill, C. Bracken, W. F. DeGrado and A. G. Palmer, J. Am. Chem. Soc., **2000**, *122*, 11610-11619.
- E. Z. Eisenmesser, D. A. Bosco, M. Akke and D. Kern, Science, 2002, 295, 1520-1523.
- D. M. Korzhnev, T. L. Religa, W. Banachewicz, A. R. Fersht and L. E. Kay, Science, **2010**, *329*, 1312-1316.
- Korzhnev, D. M., V. Y. Orekhov, and L. E. Kay, J. Am. Chem. Soc. 2005, 127, 713-721.