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# The reliability of parameters obtained by fitting of $^1\text{H}$ NMRD profiles and $^{17}\text{O}$ NMR data of potential $\text{Gd}^{3+}$ -based MRI contrast agents

Joop A. Peters\*



Synthetic variable temperature  $^1\text{H}$  NMRD profiles and  $^{17}\text{O}$  NMR relaxation and shift data were generated with a model based on the Solomon–Bloembergen–Morgan and Freed theories and then fitted simultaneously or individually. The effects of the fitting procedure and of experimental uncertainties on the resulting best-fit parameters were investigated. The most reliable best-fit parameters were obtained when all data were included in a simultaneous fitting procedure. Fitting of only NMRD and/or  $^{17}\text{O}$  NMR data provided considerably less accurate best-fit parameters. Very large deviations from the values of the parameters used for the construction of the datasets were obtained due to the combined effects of uncertainties resulting from the fitting and from the data. For these fittings, the accuracy of the best-fit parameters appeared to be strongly dependent on the magnitude of synthetic parameters applied. For example, the accuracy of  $\tau_{\text{M}}^{\text{C}}$  was low around  $\tau_{\text{M}}^{\text{S}} = 10^{-8}$  s. The parameters  $\tau_{\text{V}}$  and  $\Delta^2$  are strongly correlated in fittings of only  $^{17}\text{O}$  NMR data. Consequently, only the ratio of these parameters can be evaluated in this way. The observations underline the need to reduce the number of adjustable parameters by constraining as many as possible of them at values obtained by independent techniques. The inaccuracies observed in these simulations come in addition to those caused by the inadequacy of the Solomon–Bloembergen–Morgan theory, particularly at low magnetic field strengths. Copyright © 2015 John Wiley & Sons, Ltd.

**Keywords:** MRI contrast agents; relaxivity; water exchange rate; rotational correlation times; electronic relaxation times

## 1. INTRODUCTION

Insight into the parameters that influence the efficacy of contrast agents (CAs) for MRI is essential for the rational design of novel more efficient agents (1–4). The most frequently applied CAs are  $\text{Gd}^{3+}$ -based compounds, which improve the contrast by enhancing the longitudinal relaxation rates of water protons in their proximity. Their efficacy is usually expressed as the relaxivity ( $r_1$ ), the longitudinal relaxation rate enhancement normalized to 1 mmol of  $\text{Gd}^{3+}$ , which is determined by (i) experimental parameters: magnetic field strength ( $B$ ) and temperature ( $T$ ), (ii) structural and dynamic parameters of the CA and (iii) physical parameters associated with the electronic relaxation of the  $\text{Gd}^{3+}$  ion. The relaxivity as a function of the magnetic field strength (the  $^1\text{H}$  NMRD profile) can be evaluated with models based on the Solomon–Bloembergen–Morgan (SBM) (5–8) and the Freed theory (9) for the inner and outer sphere contributions, respectively. Various refinements of the SBM model have been proposed, which more accurately describe the electronic relaxation (10) and the mobility of the system (11). It should be noted that the SBM model is an approximate theory based on many assumptions and is not generally valid (10,12). However, many groups use this model to evaluate the parameters governing the relaxivity of CAs by fitting their NMRD profiles. Such a fitting is a cumbersome procedure because at least 10 adjustable parameters are required for each temperature. Therefore, constraints are usually introduced by fixing as many adjustable parameters as possible at the most likely values or better at values that have been determined independently. For instance, fitting of the variable temperature  $^{17}\text{O}$  NMR relaxation rates ( $T_1$  and  $T_2$ ) and chemical shifts induced by the CA with a model based on

the equations of Swift and Connick (8,13,14) affords values for the residence time of water in the first coordination sphere of  $\text{Gd}^{3+}$  ( $\tau_{\text{M}}$ ) and electronic relaxation time parameters. The fitting of the NMRD profiles and the  $^{17}\text{O}$  data can be performed either separately or simultaneously (8). Further constraints can be applied by including multiple frequency EPR linewidth data in the fitting procedure (8).

A complete physicochemical characterization of a potential CA through the parameters that govern its relaxivity requires collection and analysis of a large amount of data, and therefore is very time consuming. The attention of researchers in the field is shifting from low molecular weight Gd-chelates for non-selective contrast enhancement to complex materials, which allow combination of highly sensitive MRI probes with those for other diagnostic imaging techniques in multiple probes or with therapeutic agents in so-called theranostics. NMRD profiles can be measured with samples having a  $\text{Gd}^{3+}$  concentration of about 1 mM, but accurate  $^{17}\text{O}$  NMR measurements require concentrations that are at least an order of magnitude higher. Unfortunately, such concentrations often cannot be reached for complex materials such as nanoparticles, and variable temperature measurements may also be difficult with these materials.

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Usually, the inaccuracies of best-fit parameters of experimental relaxation and shift data of CAs are reported as standard deviations as obtained by the fitting algorithm. This may be misleading because these procedures do not take into account inaccuracies due to (i) the invalidity of the assumptions on which the of the SBM model is based and (ii) the presence of multiple local minima in the optimization. The former source of inaccuracies has been studied extensively by the groups of Fries, Merbach and Helm (1,10,12,15). The present study focusses on aspect (ii), particularly on the reliability of best-fit values obtained by simultaneous fittings of various synthetic NMRD profiles and variable temperature  $^{17}\text{O}$  NMR relaxation and shift data, compared with those obtained by fitting of exclusively NMRD or  $^{17}\text{O}$  NMR data.

## 2. RESULTS AND DISCUSSION

### 2.1. Construction of the synthetic datasets

All synthetic data were generated for five  $\tau_{\text{M}}^{298}$  values in the range typical for potential CAs ( $10^{-5}$ – $10^{-9}$  s) and for  $\tau_{\text{R}}^{298} = 10^{-10}$  and  $10^{-8}$  s, typical values for low and high molecular weight compounds, respectively.  $^1\text{H}$  NMRD data were calculated for 28 Larmor frequencies (LFs) between 0.01 and 400 MHz at 5, 15, 25, 37 and 50 °C and  $^{17}\text{O}$  NMR data for 10 temperatures in the range 5–95 °C at a magnetic field strength of 7.05 T. Other parameters used to calculate the data were arbitrarily chosen in the range usually reported for CAs (1) and are tabulated in Table 1. To simulate the effect of inaccuracies in the experimental data, additional synthetic datasets were constructed with random errors of 3% added to the  $^1\text{H}$  and  $^{17}\text{O}$  relaxation rates and 5% added to the  $^{17}\text{O}$  chemical shifts.

#### 2.1.1. Synthetic $^1\text{H}$ NMRD data

The study is limited to the relatively simple situation where second sphere waters are absent and local motions are unimportant. Two contributions are considered to influence the longitudinal

proton relaxation rate profile, inner sphere ( $r_{1,\text{IS}}$ ) and outer sphere relaxivity ( $r_{1,\text{OS}}$ ):

$$r_1 = r_{1,\text{IS}} + r_{1,\text{OS}} \quad (1)$$

The dipolar interaction dominates  $r_{1,\text{IS}}$  and is described by eqn (2) (13) and the Solomon–Bloembergen equation (eqn (3)) (5–7), where  $q$  is the number of inner sphere water molecules (assumed to be unity in the present calculations) and  $T_{1\text{M}}$  is the longitudinal relaxation time of a  $\text{Gd}^{3+}$ -bound water molecule. The distance between  $\text{Gd}^{3+}$  and the H atom of a bound water molecule ( $r_{\text{GdH}}$ ) was fixed at 3.10 Å, unless stated otherwise.

$$r_{1,\text{IS}} = \frac{q}{55556(T_{1\text{M}} + \tau_{\text{M}})} \quad (2)$$

$$\frac{1}{T_{1\text{M}}} = \frac{2}{15} \left( \frac{\mu_0}{4\pi} \right)^2 \frac{\hbar^2 \gamma_{\text{S}}^2 \gamma_{\text{I}}^2}{r_{\text{GdH}}^6} S(S+1) \left( \frac{3\tau_{\text{d1}}}{1 + \omega_{\text{I}}^2 \tau_{\text{d1}}^2} + \frac{7\tau_{\text{d2}}}{1 + \omega_{\text{S}}^2 \tau_{\text{d2}}^2} \right) \quad (3)$$

Here  $(\mu_0/4\pi)$  is the magnetic permeability in vacuum,  $S$  is the electron spin ( $S = 7/2$  for  $\text{Gd}^{3+}$ ),  $\gamma_{\text{I}}$  is the nuclear gyromagnetic ratio,  $\gamma_{\text{S}}$  is the electron gyromagnetic ratio,  $\omega_{\text{I}}$  and  $\omega_{\text{S}}$  are the Larmor frequencies of the proton and electron spin, respectively, and  $\tau_{\text{di}}^{-1} = \tau_{\text{M}}^{-1} + \tau_{\text{R}}^{-1} + T_{\text{ie}}^{-1}$ , where  $T_{\text{ie}}$  ( $i = 1, 2$ ) stands for electronic relaxation time. The latter are mainly due to the zero field splitting (ZFS) interaction and can be expressed by eqns (4) and (5) (16):

$$\frac{1}{T_{1\text{e}}} = \frac{1}{25} \Delta^2 \tau_{\text{v}} [4S(S+1) - 3] \left( \frac{1}{1 + \omega_{\text{S}}^2 \tau_{\text{v}}^2} + \frac{1}{1 + 4\omega_{\text{S}}^2 \tau_{\text{v}}^2} \right) \quad (4)$$

$$\frac{1}{T_{2\text{e}}} = \Delta^2 \tau_{\text{v}} \left( \frac{5.26}{1 + 0.372\omega_{\text{S}}^2 \tau_{\text{v}}^2} + \frac{7.18}{1 + 1.24\omega_{\text{S}}^2 \tau_{\text{v}}^2} \right) \quad (5)$$

In these equations,  $\Delta^2$  represents the mean-squared fluctuation of the ZFS and  $\tau_{\text{v}}$  is the correlation time for the instantaneous distortion of the coordination polyhedron of  $\text{Gd}^{3+}$ .

The outer sphere contribution to the relaxivity ( $r_{1,\text{OS}}$ ) is described by eqs (6) and (7) (9):

$$r_{1,\text{OS}} = \left( \frac{32\pi}{405} \right) \left( \frac{\mu_0}{4\pi} \right)^2 \gamma_{\text{I}}^2 \gamma_{\text{S}}^2 \hbar^2 S(S+1) \frac{N_{\text{A}}}{a_{\text{GdH}} D_{\text{GdH}}} [3J_{\text{OS}}(\omega_{\text{I}}, T_{1\text{e}}) + 7J_{\text{OS}}(\omega_{\text{S}}, T_{1\text{e}})] \quad (6)$$

$$J_{\text{OS}}(\omega, T_{\text{je}}) = \text{Re} \left\{ \frac{1 + \frac{1}{4} \left[ i\omega\tau_{\text{GdH}} + \left( \frac{\tau_{\text{GdH}}}{T_{\text{je}}} \right) \right]^2}{1 + \left[ i\omega\tau_{\text{GdH}} + \left( \frac{\tau_{\text{GdH}}}{T_{\text{je}}} \right) \right]^2 + \frac{4}{9} \left[ i\omega\tau_{\text{GdH}} + \left( \frac{\tau_{\text{GdH}}}{T_{\text{je}}} \right) \right]^2 + \frac{1}{9} \left[ i\omega\tau_{\text{GdH}} + \left( \frac{\tau_{\text{GdH}}}{T_{\text{je}}} \right) \right]^2} \right\} \quad (7)$$

Here  $N_{\text{A}}$  is Avogadro's number,  $a_{\text{GdH}}$  is the distance of closest approach of a diffusing water molecule to  $\text{Gd}^{3+}$ , which was fixed at 3.5 Å in all calculations,  $D_{\text{GdH}}$  stands for the diffusion coefficient and  $J_{\text{OS}}(\omega, T_{\text{je}})$  ( $j = 1, 2$ ) are spin density functions. Generally,  $D_{\text{GdH}}$  can be well estimated by an empirical equation proposed by Vander Elst *et al.* (17), and consequently this parameter can be fixed in the fitting procedure. In the present calculations  $D_{\text{GdH}}^{298}$  was arbitrarily set at  $2.2 \times 10^{-9} \text{ m}^2 \text{ s}^{-1}$ . The diffusion correlation time ( $\tau_{\text{GdH}}$ ) is given by  $a_{\text{GdH}}^2/D_{\text{GdH}}$ . All other correlation times are assumed to obey an exponential temperature dependence (eqn (8)), where  $x = \text{R, M or V}$ ,  $\tau_x^T$  and  $\tau_x^{298}$  are the values of the parameter concerned at temperature  $T$  and 298.15 K,

**Table 1.** Values of variables used in calculations of synthetic  $^1\text{H}$  NMRD and  $^{17}\text{O}$  NMR data. The values that were fixed during the fittings are in italic

$q$	1.0
$r_{\text{GdH}}$ (Å)	3.10
$\tau_{\text{M}}^{298}$ (s)	$10^{-5}$ – $10^{-9}$
$\tau_{\text{R}}^{298}$ (s)	$10^{-8}$ , $10^{-10}$
$\tau_{\text{V}}^{298}$ (ps)	15.0
$\Delta^2$ ( $10^{19} \text{ s}^{-2}$ )	1.60
$D_{\text{GdH}}^{298}$ ( $10^{-9} \text{ m}^2 \text{ s}^{-1}$ )	2.20
$a_{\text{GdH}}^2$ (Å)	3.50
$A/\hbar$ ( $10^6 \text{ rad s}^{-1}$ )	3.60
$\chi(1 + \eta^2/3)^{1/2}$ (MHz)	7.58
$r_{\text{GdO}}$ (Å)	2.50
$C_{\text{OS}}$	0.1
$E_{\text{M}}$ ( $\text{kJ mol}^{-1}$ )	47.6
$E_{\text{R}}$ ( $\text{kJ mol}^{-1}$ )	21.9
$E_{\text{V}}$ ( $\text{kJ mol}^{-1}$ )	1.0
$E_{\text{D}}$ ( $\text{kJ mol}^{-1}$ )	18.2

respectively,  $E_x$  is the associated activation energy and  $R$  is the gas constant.

$$\tau_x^T = \tau_x^{298} \exp \left[ \frac{E_x}{R} \left( \frac{1}{T} - \frac{1}{298.15} \right) \right] \quad (8)$$

The temperature dependence of  $D_{\text{GdH}}$  is assumed to obey

$$D_{\text{GdH}}^T = D_{\text{GdH}}^{298} \exp \left[ \frac{E_D}{R} \left( \frac{1}{298.15} - \frac{1}{T} \right) \right] \quad (9)$$

$E_v$  was fixed at  $1.0 \text{ kJ mol}^{-1}$  and  $E_D$  at  $18.2 \text{ kJ mol}^{-1}$ .

## 2.1.2. Synthetic $^{17}\text{O}$ NMR data

Synthetic datasets consisting of reduced  $\text{Gd}^{3+}$ -induced water  $^{17}\text{O}$  longitudinal and transverse relaxation rates ( $1/T_{1r}$  and  $1/T_{2r}$ ) and angular frequencies ( $\Delta\omega_r$ ) were generated using eqns (10–12):

$$\frac{1}{T_{1r}} = \frac{1}{T_{1M} + \tau_M} \quad (10)$$

$$\frac{1}{T_{2r}} = \frac{1}{\tau_M} \frac{T_{2M}^{-2} + \tau_M^{-1} T_{2M}^{-1} + \Delta\omega_M^2}{(\tau_M^{-1} + T_{2M}^{-1})^2 + \Delta\omega_M^2} \quad (11)$$

$$\Delta\omega_r = \frac{\Delta\omega_M}{(1 + \tau_M T_{2M}^{-1})^2 + \tau_M^2 \Delta\omega_M^2} + C_{OS} \Delta\omega_M \quad (12)$$

The outer sphere contributions to the relaxation rates have been shown to be negligible (18), and therefore are not included in eqns (10) and (11). The outer sphere contribution to  $\Delta\omega_r$  is assumed to be proportional to  $\Delta\omega_M$  and is represented by the last term in eqn (12). The constant  $C_{OS}$  was fixed at 0.1 in the present calculations. The  $^{17}\text{O}$  longitudinal relaxation rates in eqn (10) are dominated by contributions of the dipole–dipole and quadrupolar mechanisms as expressed by eqns (13–16) (16):

$$\frac{1}{T_{1M}} = \frac{1}{T_{1dd}} + \frac{1}{T_{1q}} \quad (13)$$

$$\frac{1}{T_{1dd}} = \frac{1}{15} \frac{\gamma_I^2 \gamma_S^2 \hbar^2}{r_{\text{GdO}}^6} S(S+1) \left( \frac{\mu_0}{4\pi} \right)^2 \left[ 6\tau_{d1} + 14 \frac{\tau_{d2}}{1 + \omega_S^2 \tau_{d2}^2} \right] \quad (14)$$

$$\frac{1}{T_{1q}} = \frac{1}{T_{2q}} = \frac{3\pi^2(2I+3)}{10I^2(2I-1)} \chi^2 (1 + \eta^2/3) \tau_{RO} \quad (15)$$

$$\tau_{di}^{-1} = \tau_M^{-1} + \tau_{RO}^{-1} + T_{ie}^{-1} \quad i = 1, 2 \quad (16)$$

Here,  $r_{\text{GdO}}$  is the distance between the electron charge and the  $^{17}\text{O}$  nucleus, which was fixed at  $2.50 \text{ \AA}$  in the present calculations. The rotational correlation time for the Gd–O vector ( $\tau_{RO}$ ) is not necessarily equal that of the Gd–water H vector,  $\tau_{RH}$  (19). In the present calculations, the ratio  $\tau_{RO}/\tau_{RH}$  was fixed at 1.54. In eqn (15),  $\chi$  is the quadrupolar coupling constant and  $\eta$  an asymmetry parameter. The value of the quadrupole coupling parameter  $\chi\sqrt{(1 + \eta^2/3)}$  was set at the value of pure water,  $7.58 \text{ MHz}$  (20,21).

$1/T_{2q}$  is equal to  $1/T_{1q}$  (see eqn (15)) and the scalar and dipolar contributions to the transverse  $^{17}\text{O}$  relaxation rate are given by eqns (17) and (18), (19), respectively:

$$\frac{1}{T_{2SC}} = \frac{S(S+1)}{3} \left( \frac{A}{\hbar} \right)^2 \tau_{e1} \quad (17)$$

$$\frac{1}{T_{2dd}} = \frac{1}{15} \frac{\gamma_I^2 \gamma_S^2 \hbar^2}{r_{\text{GdO}}^6} S(S+1) \left( \frac{\mu_0}{4\pi} \right)^2 \left[ 7\tau_{d1} + 13 \frac{\tau_{d2}}{1 + \omega_S^2 \tau_{d2}^2} \right] \quad (18)$$

$$\tau_{e1}^{-1} = \tau_M^{-1} + T_{1e}^{-1} \quad (19)$$

$$\frac{1}{T_{2M}} = \frac{1}{T_{2dd}} + \frac{1}{T_{2q}} + \frac{1}{T_{2SC}} \quad (20)$$

Micskei *et al.* have determined that, for  $[\text{Gd}(\text{DTPA})(\text{H}_2\text{O})]^{2-}$  and for  $[\text{Gd}(\text{DOTA})(\text{H}_2\text{O})]^-$ , the scalar contribution makes up 95% of  $1/T_{2M}$ , and therefore, in their calculations, the contributions of the dipolar and the quadrupolar relaxation mechanisms could be neglected (18). Generally, in fittings of  $^{17}\text{O}$  NMR relaxation rates, the dipolar and quadrupolar contributions are neglected as well (1). However, simulations show that for systems with smaller  $\tau_M$  and/or larger  $\tau_R$  this assumption is no longer valid. For example, for  $LF = 40 \text{ MHz}$ ,  $\tau_M = 1 \text{ ns}$ ,  $\tau_R = 1 \text{ ns}$  and with the other parameters the same as for  $[\text{Gd}(\text{DTPA})(\text{H}_2\text{O})]^{2-}$ , the dipolar, scalar and quadrupolar contributions to  $1/T_{2M}$  are 27, 54 and 19%, respectively. Therefore, in the present calculations, all these contributions are included in  $1/T_{2M}$  (see eqn (20)).

## 2.2. Fittings of the synthetic datasets

### 2.2.1. Procedures

The fittings were carried out by minimization of  $S^2$ , the sum of the squared differences between calculated and synthetic data (see eqn (21)) using the evolutionary algorithm of the Microsoft Excel Solver (22). This algorithm has the advantage that it results usually in a global minimum. Initial estimates of the fitting parameters were randomly generated numbers within the ranges commonly observed for these parameters (see Table S1). The minimization procedure was continued until no further significant decrease of  $S^2$  occurred after restarting. For display purposes, the goodness of the fits was expressed as the agreement factor ( $AF$ ; see eqn (22)). The final values of  $S^2$  and  $AF$  were always below those calculated for the synthetic data after adding the random errors with respect to synthetic data without added errors ( $AF \leq 0.02$ ). Standard deviations of the best-fit parameters were calculated by repeating a minimization five times with different randomly generated initial estimates.

$$S^2 = \sum_i (y_i^c - y_i^s)^2 \quad (21)$$

$$AF = \sqrt{\frac{\sum_i (y_i^c - y_i^s)^2}{\sum_i (y_i^s)^2}} \quad (22)$$

The fittings of the synthetic data were carried out with four different methods:

(A) variable temperature NMRD data and  $^{17}\text{O}$  NMR data simultaneously;

**Table 2.** Comparison of parameters used for the construction of synthetic variable temperature  $^1\text{H}$  NMRD data with best-fit parameters obtained from this dataset with Method B

	Synthetic <sup>1</sup>	Best-fit parameters		
		No errors <sup>2,3</sup>		Errors included <sup>2</sup>
			Minimum 1	Minimum 2
$\tau_M^{298}$ (s)	$10^{-9}$	$(2.1 \pm 0.1) \times 10^{-9}$	$(1.5 \pm 0.5) \times 10^{-7}$	$(3 \pm 2) \times 10^{-9}$
$\tau_R^{298}$ (ps)	100.0	$95.0 \pm 0.3$	$90.0 \pm 0.8$	$93 \pm 3$
$\tau_V^{298}$ (ps)	15.0	$15.0 \pm 0.0$	$14.1 \pm 0.4$	$15.5 \pm 0.1$
$\Delta^2$ ( $10^{19} \text{ s}^{-2}$ )	1.60	$1.59 \pm 0.00$	$1.38 \pm 0.05$	$1.50 \pm 0.01$
$E_M$ ( $\text{kJ mol}^{-1}$ )	47.6	$56.9 \pm 0.9$	$7 \pm 1$	$53 \pm 5$
$E_R$ ( $\text{kJ mol}^{-1}$ )	21.9	$22.9 \pm 0.1$	$26.0 \pm 0.4$	$23.3 \pm 0.8$
AF	0.0	$2.4 \times 10^{-4}$	$2.1 \times 10^{-2}$	$2.2 \times 10^{-2}$

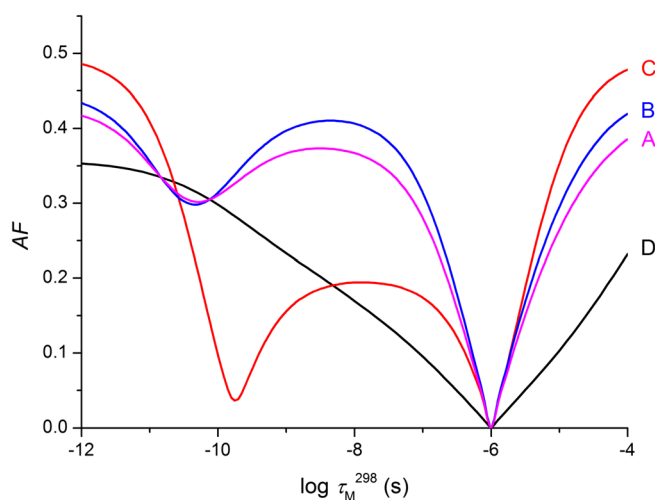
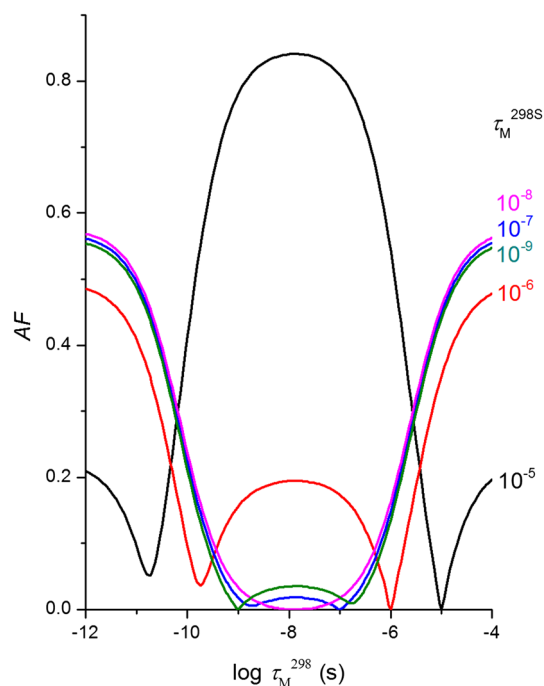
<sup>1</sup>All other parameters were fixed at the values given in Table 1. The best-fit values are averages of results of five fittings with different initial parameters.  
<sup>2</sup>In synthetic data.  
<sup>3</sup>A single minimum was found.

- (B) exclusively the variable temperature NMRD data;  
 (C) exclusively an NMRD profile for 25 °C;  
 (D) exclusively variable temperature  $^{17}\text{O}$  NMR data.

In all cases excellent fits were obtained with  $AF \leq 0.02$ . Below, the parameters employed for the construction of the datasets, the “synthetic parameters”, will be denoted with a superscript S and those obtained after fitting of these data with a superscript C.

### 2.2.2. Method A: simultaneous fitting of variable temperature $^1\text{H}$ NMRD and $^{17}\text{O}$ NMR data

Fittings of synthetic data for  $\tau_M^{298S} = 10^{-5}$ – $10^{-9}$  s and  $\tau_R^{298S} = 10^{-10}$  s afforded best-fit parameters that were always within 0.1% of the values employed for their construction (see Table S2). The standard deviations were negligibly small.

**Figure 1.** Plot of AF versus  $\log \tau_M^{298}$  for the various fitting procedures of synthetic datasets constructed for  $\tau_M^{298S} = 10^{-6}$  s,  $\tau_R^{298S} = 10^{-10}$  s and the other parameters fixed at the values mentioned in Table 1.**Figure 2.** Plot of AF versus  $\log \tau_M^{298}$  for fitting of synthetic NMRD datasets (Procedure C) constructed for various values of  $\tau_M^{298S}$ , with the other parameters fixed at the values mentioned in Table 1.

### 2.2.3. Method B: fitting of variable temperature $^1\text{H}$ NMRD data

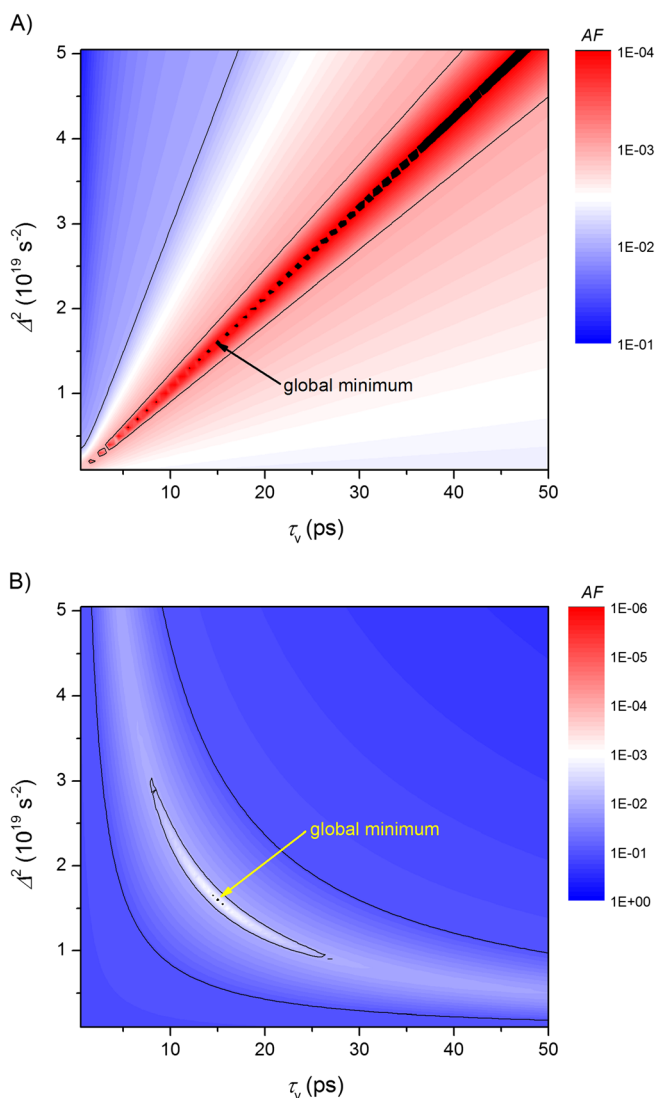
Fittings of synthetic NMRD data for  $\tau_M^{298S} = 10^{-5}$ – $10^{-8}$  s and  $\tau_R^{298S} = 10^{-10}$  s again all resulted in excellent fits ( $AF < 10^{-4}$ ), and produced best-fit parameters that were identical to the corresponding synthetic values and that had negligible standard deviations (see Table S4). However, for  $\tau_M^{298S} = 10^{-9}$  s,  $\tau_M^{298C}$  deviated by a factor of about 2 from the corresponding synthetic values and had relatively large standard errors (see Table 2), although the fit was still perfect ( $AF = 2 \times 10^{-4}$ ). The deviations in  $\tau_M$  were mainly compensated by small changes in  $\tau_R$  and  $E_M$ .

**Table 3.** Comparison of parameters used for the construction of synthetic variable temperature  $^{17}\text{O}$  NMR data with best-fit parameters obtained from this dataset with Method D. Standard deviations are given in parentheses<sup>1</sup>

	Synthetic	Best-fit parameters	
		No errors <sup>2</sup>	Errors included <sup>2</sup>
$\tau_M^{298}$ ( $\mu\text{s}$ )	1.00	1.00 (0.00)	0.99 (0.00)
$\tau_R^{298}$ (ps)	100.0	100.0 (0.0)	100.9 (0.0)
$\tau_V^{298}$ (ps)	15.0	70 (13)	58 (13)
$\Delta^2$ ( $10^{19} \text{ s}^{-2}$ )	1.60	7.5 (1.4)	5 (1)
$E_M$ ( $\text{kJ mol}^{-1}$ )	47.6	47.6 (0.0)	47.9 (0.0)
$E_R$ ( $\text{kJ mol}^{-1}$ )	21.9	21.9 (0.0)	22.3 (0.0)
$\Delta^2/\tau_V^{298}$ ( $10^{30} \text{ s}^{-3}$ )	1.0667	1.06 (0.00)	1.02 (0.00)
AF	0.0	$7.4 \times 10^{-7}$	$1.6 \times 10^{-3}$

<sup>1</sup>All other parameters were fixed at the values given in Table 1. The best-fit values are averages of results of five fittings with different initial parameters.  
<sup>2</sup>In synthetic data.





**Figure 3.** Contour plot of AF for a synthetic dataset with  $\tau_M^{298S} = 1 \mu s$ , as a function of  $\tau_v^{298}$  and  $\Delta^2$  for (A) fitting of  $^{17}O$  data (Procedure D) and (B) fitting variable temperature NMRD data (Procedure B).

This decrease in accuracy of  $\tau_M^{298C}$  can be ascribed to the transition of the water exchange rate from the slow to the fast exchange limit. For  $\tau_M^{298} = 10^{-9}$  s, the fast exchange limit is reached and  $\tau_M$  is so small relative to  $T_{1M}$  that it can be neglected in eqn (2). Consequently,  $\tau_M$  influences  $r_1$  only through the correlation times that govern  $T_{1M}$ , namely  $\tau_{d1}$  and  $\tau_{d2}$ , which are given

by  $\tau_{di}^{-1} = \tau_M^{-1} + \tau_R^{-1} + T_{ie}^{-1}$ . In the NMRD profiles concerned, the relaxivity contribution due to electronic relaxation dominates up to  $LF \approx 1$  MHz, whereas the effects of  $\tau_M$  and  $\tau_R$  dominate at 1–100 MHz. Thus in the latter  $LF$  region, deviations of  $\tau_M^S$  from  $\tau_M^C$  are mainly reflected in variations in the shape of the NMRD profile at 1–100 MHz, which can be compensated optimally by variations in  $\tau_R^C$  and vice versa. The sum of the reciprocals of these correlation times of each fitting was the same as that of the corresponding synthetic values  $\tau_M^S$  and  $\tau_R^S$ .

#### 2.2.4. Method C: fitting of only $^1H$ NMRD data for 25 °C

An inspection of AF curves for fittings of synthetic NMRD datasets for a single temperature (Procedure C) as a function of  $\tau_M^{298}$  shows that next to the global minimum a local minimum is present with a slightly higher AF (see Fig. 1). This can be explained by the shape of curves of the relaxivity  $r_1$  as a function of  $\tau_M$ . From eqns (2–5) it can be deduced that at a particular  $LF$  such a curve presents a maximum when  $T_{1M}^{-1} < \tau_M^{-1} < \tau_R^{-1}$ ;  $T_{1e}^{-1}$  (1). Hence, below this maximum, a given value of  $r_1$  corresponds to two values of  $\tau_M$ , which are located opposite each other with respect to the maximum. For the datasets without included random errors, the fitting procedures always ended always in the proper minimum for AF, when the evolutionary algorithm was applied for the minimization. Near the value of  $\tau_M$ , where the maxima in  $r_1$  occur ( $\tau_M^{298} \approx 10^{-8}$  s), the two minima in the curve for AF overlap (see Fig. 2), and consequently  $\tau_M^C$  had larger standard deviations and differed substantially from  $\tau_M^S$  (see also Table S6).

For  $\tau_M^{298S} = 10^{-5}$  s, the best-fit value  $\tau_R^{298C}$  appeared to be a factor of 10 higher than its synthetic counterpart. This can be rationalized by eqn (2), where  $\tau_M$  is larger than  $T_{1M}$  under these conditions, whereas at the same time it is of minor importance in  $\tau_{di}$  (see eqn (3)). As a result, small deviations in  $\tau_M$  have to be compensated by relatively large deviations in the other fitting parameters to reach a good fit.

#### 2.2.5. Method D: fitting of only variable temperature $^{17}O$ NMR data

If only  $^{17}O$  NMR data were considered, the evaluated best-fit values for the parameters governing the electronic relaxation ( $\tau_v^{298}$  and  $\Delta^2$ ) deviated substantially from the corresponding synthetic values, whereas the best fits of all other parameters were perfectly in agreement with the corresponding synthetic values. This is illustrated in Table 3 for  $\tau_M^{298S} = 10^{-6}$  s (see also Table S8). Although the fittings did not afford the correct values for  $\tau_v^{298}$  and  $\Delta^2$ , the qualities of the fits were excellent, as reflected in very low values for AF. A contour plot of AF as a function of  $\tau_v^{298}$  and

**Table 4.** Comparison of best-fit parameters for synthetic data constructed with  $\tau_M^{298S} = 10^{-7}$  s and  $\tau_R^{298S} = 10^{-10}$  s as obtained with Procedures A–D. Random errors were included in the data

	Synthetic	A	B	C	D
$\tau_M^{298}$ (s)	$1.00 \times 10^{-7}$	$(1.04 \pm 0.00) \times 10^{-7}$	$(2.9 \pm 0.7) \times 10^{-7}$	$(1.3 \pm 1.0) \times 10^{-7}$	$(0.99 \pm 0.00) \times 10^{-7}$
$\tau_R^{298}$ (ps)	100.0	$100.1 \pm 0.0$	$104 \pm 2$	$106 \pm 3$	$98.4 \pm 0.0$
$\tau_v^{298}$ (ps)	15.0	$16.2 \pm 0.0$	$14.6 \pm 0.9$	$10 \pm 4$	$57 \pm 16$
$\Delta^2$ ( $10^{19} s^{-2}$ )	1.60	$1.49 \pm 0.00$	$1.36 \pm 0.02$	$2.9 \pm 0.8$	$6 \pm 2$
$E_M$ (kJ mol $^{-1}$ )	47.6	$48.5 \pm 0.0$	$28 \pm 5$	—	$47.6 \pm 0.0$
$E_R$ (kJ mol $^{-1}$ )	21.9	$21.9 \pm 0.0$	$23.8 \pm 0.7$	—	$21.7 \pm 0.0$
AF	0.0	0.017	0.019	0.025	0.002

$\Delta^2$  (see Fig. 3) shows a valley with local minima along the line  $\Delta^2 \approx 1.1 \times 10^{30} \tau_V^{298}$  (see Fig. 3(A)). Inside this valley a global minimum occurs at  $\tau_V^{298} = 15$  ps,  $\Delta^2 = 1.6 \times 10^{19} \text{ s}^{-2}$  ( $AF = 7.5 \times 10^{-6}$ ), which was missed in the fitting procedure, due to the shallowness of the curve along the minima. The strong correlation between  $\tau_V^{298}$  and  $\Delta^2$  can be explained by consideration of eqns (4) and (5). At the magnetic field strengths usually applied for  $^{17}\text{O}$  NMR (in the present example  $B = 7.04$  T), the term  $\omega_s \tau_V$  is larger than unity. After filling in all the physical constants, it can be seen that both  $1/T_{1e}$  and  $1/T_{2e}$  are approximately proportional to  $\Delta^2/\tau_V$ . Consequently, fitting of the  $^{17}\text{O}$  NMR data can be used to get an estimate only of the ratio  $\Delta^2/\tau_V$  rather than of the individual parameters (see Table 3). By contrast, the part of the NMRD profiles for  $B < 1$  T is mainly determined by the electronic relaxation rates, and at these magnetic field strengths the approximations mentioned above are no longer valid. Accordingly, there is no correlation between  $\tau_V^{298}$  and  $\Delta^2$  and thus a good estimate of both parameters can be obtained from fitting of  $^1\text{H}$  NMRD (Method B or C) data alone or simultaneously with the  $^{17}\text{O}$  NMR data (Method A; see Fig. 3(B)).

#### 2.2.6. Effects of inclusion of random errors in the synthetic data

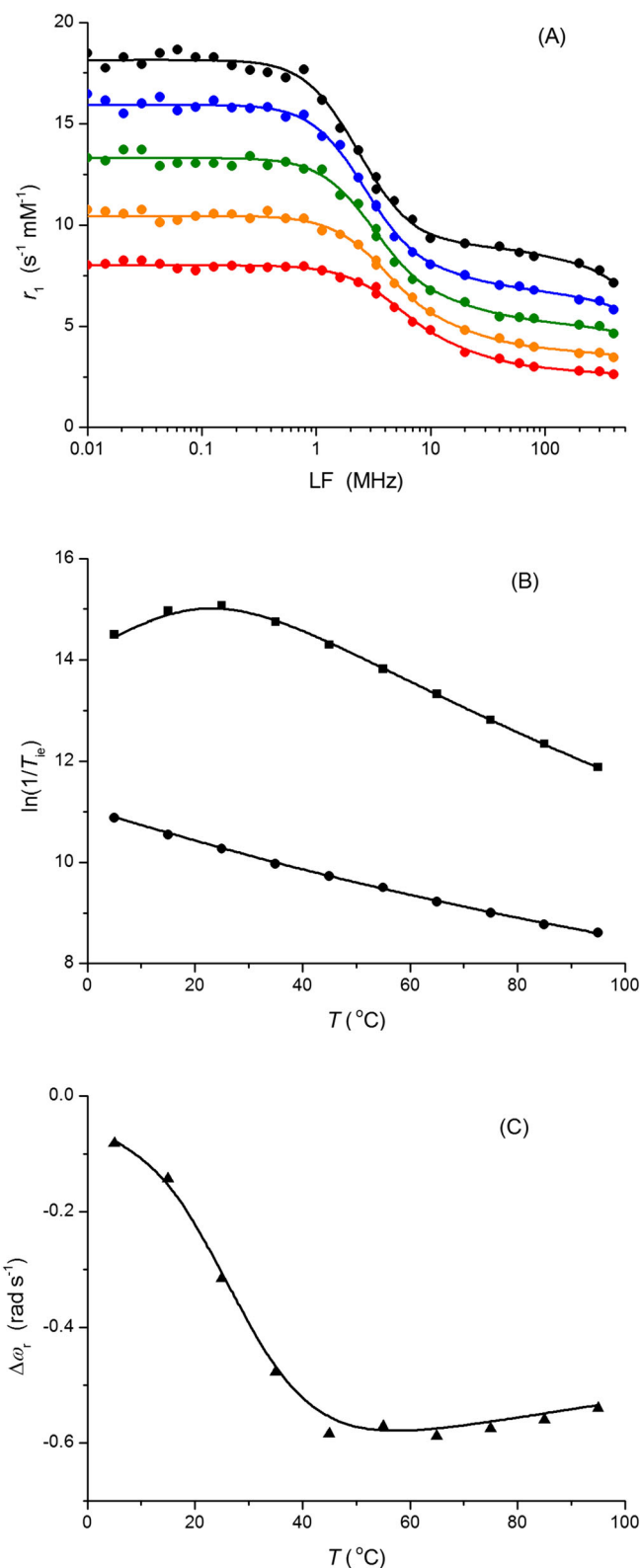
Inclusion of random errors of 3% in the relaxation rates and of 5% in the chemical shifts had a relatively small magnifying effect on the inaccuracy of the best-fit data obtained with a simultaneous fit (Method A): all best-fit parameters were within 10% of their synthetic values. However, smaller datasets (Methods B–D) afforded best-fit values that deviated substantially and had relatively large standard deviations, although excellent fits were always obtained irrespective of the procedure employed (see Tables S2–S9). This is illustrated for the best-fit parameters for the datasets for  $\tau_M = 10^{-7}$  s in Table 4, whereas in Fig. 4 the curves calculated with the values obtained with Procedure A are displayed.

As expected based on the results mentioned in the previous section, the fittings of only  $^{17}\text{O}$  NMR data gave unreliable best-fit values for  $\tau_V^{298}$  and  $\Delta^2$ , but values close to the synthetic ones were obtained for  $\tau_M^{298}$ ,  $\tau_R^{298}$  and the activation energies. The opposite holds for the fittings of NMRD data only. Fittings of the NMRD dataset constructed for  $\tau_M^{298} = 10^{-9}$  s ended up in one of the two  $AF$  minima. In this case the minimizations were repeated until five of each of the minima were obtained. In most cases, the global minimum could be identified by a careful inspection of a plot of  $AF$  as a function of  $\tau_M^{298}$ . However, great care is needed, since the differences between the  $AF$  values of the two minima are always very small. If possible, either the temperature dependence of NMRD data or  $^{17}\text{O}$  NMR data should be used to discriminate between the two minima.

The plots of  $AF$  as function of  $\tau_M^{298}$  almost coincided with the corresponding plot shown in Fig. 2; only the lower part of the curve was shifted upward by about 0.02  $AF$  units. Consequently, the separation of the two local minima was less than prior to the inclusion of random errors in the synthetic dataset. This flattening of the minima leads to a larger spreading of the best-fit values for the results of individual fittings, as reflected in the relatively large standard deviations for the best-fit values of  $\tau_M^{298C}$  for datasets constructed with  $\tau_M^{298S} = 10^{-7}$ – $10^{-9}$  s (see Table S5).

#### 2.2.7. Synthetic datasets for $\tau_R^{298S} = 10^{-8}$ s

Upon increase of  $\tau_R^{298S}$ , the valleys in the profiles of  $AF$  as a function of  $\tau_R$  become narrower. Consequently, the values of the



**Figure 4.** Results of simultaneous fitting of synthetic datasets using the best-fit values given in Table 4, Procedure A. (A)  $^1\text{H}$  NMRD data for 5 (black), 15 (blue), 25 (green), 37 (orange) and 50 °C (red); (B) reduced  $^{17}\text{O}$  NMR relaxation rates ( $\blacksquare$ ,  $1/T_{2r}$ ;  $\bullet$ ,  $1/T_{1r}$ ); (C) reduced  $^{17}\text{O}$  NMR chemical shifts of water.

best-fit parameters are closer to those of the parameters used to construct the synthetic datasets, and their standard deviation generally decreases (compare Tables 5 and 4). When Procedure C is

**Table 5.** Comparison of best-fit parameters for synthetic data constructed with  $\tau_M^{298} = 10^{-7}$  s and  $\tau_R^{298} = 10^{-8}$  s as obtained with Procedures A–D

	Synthetic	A	B	C	D
$\tau_M^{298}$ (s)	$1.00 \times 10^{-7}$	$(1.01 \pm 0.00) \times 10^{-7}$	$(1.01 \pm 0.00) \times 10^{-7}$	$(1.03 \pm 0.00) \times 10^{-7}$	$(1.00 \pm 0.00) \times 10^{-7}$
$\tau_R^{298}$ (s)	$10^{-8}$	$(0.98 \pm 0.01) \times 10^{-8}$	$(0.98 \pm 0.02) \times 10^{-8}$	$(0.94 \pm 0.01) \times 10^{-8}$	$(1.00 \pm 0.00) \times 10^{-8}$
$\tau_V^{298}$ (ps)	15.0	$15.0 \pm 0.0$	$15.0 \pm 0.0$	$15.0 \pm 0.0$	$43 \pm 18$
$\Delta^2$ ( $10^{19}$ s $^{-2}$ )	1.60	$1.60 \pm 0.00$	$1.59 \pm 0.01$	$1.59 \pm 0.01$	$4 \pm 2$
$E_M$ (kJ mol $^{-1}$ )	47.6	$48.0 \pm 0.1$	$47.9 \pm 0.2$	—	$47.7 \pm 0.0$
$E_R$ (kJ mol $^{-1}$ )	21.9	$22.0 \pm 0.0$	$22.2 \pm 0.4$	—	$21.9 \pm 0.0$
AF	0.0	0.020	0.021	0.021	0.0015

applied, the separation between the two local minima is better. Then, the global minimum can be located more easily. This is illustrated in Fig. 5, which displays the profiles for  $\tau_R = 10^{-8}$  and  $10^{-10}$  s.

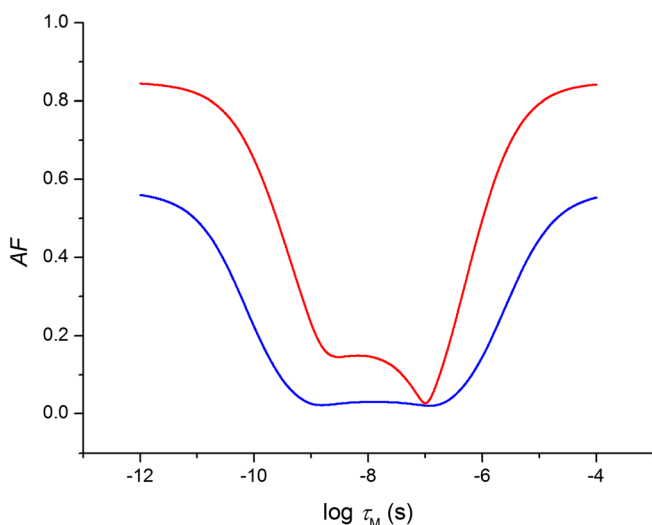
### 2.2.8. Variation of $q$ and $r_{\text{GdH}}$

From eqn (2), it is evident that evaluation of NMRD data can provide the value of the ratio  $q/r_{\text{GdH}}^6$ , but not that of either  $q$  or  $r_{\text{GdH}}$ . The parameters  $q$  and  $r_{\text{GdH}}$  are often both fixed in fitting procedures of NMRD data. Determination of either of these parameters from NMRD data requires an independent determination of the other parameter by a different technique.

The value of  $q$  can be estimated from the  $\text{Gd}^{3+}$  induced shift of the water  $^{17}\text{O}$  nucleus, provided that the exchange between bound and bulk water is fast on the  $^{17}\text{O}$  NMR time scale. This shift is purely of contact origin and is, to a first approximation, linearly proportional to  $q$  (23,24). A widely used alternative method to estimate  $q$  values of  $\text{Gd}^{3+}$  complexes is from  $q$  values of corresponding  $\text{Eu}^{3+}$  and/or  $\text{Tb}^{3+}$  complexes as determined by their luminescence decay rates in  $\text{H}_2\text{O}$  and  $\text{D}_2\text{O}$  (25). A double check of the correctness of a  $q$  value can be obtained by fitting NMRD and/or  $^{17}\text{O}$  NMR data according to Procedures A or D, by including  $A/\hbar$  in the set of adjustable fitting parameters. A

best-fit value obtained for  $A/\hbar$  in the range from  $-4.1 \times 10^{-6}$  to  $-3.6 \times 10^{-6}$  rad s $^{-1}$  is an indication that the assumed  $q$  value is correct, since the value of  $A/\hbar$  is generally almost independent of the structure of the lanthanide complex. Sometimes non-integral numbers have been obtained for  $q$ , which may be due to, for example, (i) occurrence of equilibria of Gd-complexes with different hydration states, (ii) neglect of second hydration sphere effects when  $q$  was determined by luminescence or (iii)  $A/\hbar$  values outside the common range when the  $^{17}\text{O}$  NMR method was used. When non-integral  $q$  values occur, reliable fittings of NMRD data can only be obtained if the thermodynamic parameters of the hydration equilibria are known. A good estimate can usually be obtained by high resolution UV-vis spectrometric measurements on the corresponding  $\text{Eu}^{3+}$  system. The intensity of the absorption band for the  $^7\text{F}_0 \rightarrow ^5\text{D}_0$  transition is low, but generally so narrow that deconvolution into bands for the different hydration states is possible (26–29).

Since the number of required adjustable parameters in fittings is very large, the value of  $r_{\text{GdH}}$  is often fixed in fittings of NMRD data reported in the literature. Caravan *et al.* have determined the value of  $r_{\text{GdH}}$  for several Gd-based CAs from anisotropic hyperfine constants as determined by pulsed ENDOR spectroscopy on glassy methanol–water samples (30,31). These distances were all found to be in the narrow range of 3.0–3.2 Å. This seems to justify fixing of  $r_{\text{GdH}}$  at 3.10 Å. However, some caution is needed, because  $r_1$  is very sensitive to variations of  $r_{\text{GdH}}$  due to the proportionality of the inner sphere relaxation rate to  $r_{\text{GdH}}^{-6}$  (see eqn (3)). Table 6 shows that fixing of  $r_{\text{GdH}}$  at an incorrect value



**Figure 5.** Profiles of AF as function of  $\log \tau_M$  at 25 °C for fittings (Procedure C) of a synthetic NMRD dataset constructed with  $\tau_M^{298} = 10^{-7}$  s and  $\tau_R = 10^{-8}$  s (red) and  $10^{-10}$  s (blue). All other parameters are as in Table 1. Random errors were included in the synthetic datasets.

**Table 6.** Best-fit parameters (Procedure D) for synthetic NMRD data constructed with  $r_{\text{GdH}}^S = 3.10$  Å obtained for various fixed values of  $r_{\text{GdH}}^1$ 

	Synthetic $r_{\text{GdH}} = 3.10$ Å	Best-fit parameters for $r_{\text{GdH}}$ fixed at		
		3.00 Å	3.10 Å	3.20 Å
$\tau_M^{298}$ ( $10^{-8}$ s)	1.00	$0.08 \pm 0.05$	$0.42 \pm 0.40$	$77.4 \pm 0.2$
$\tau_R^{298}$ ( $10^{-10}$ s)	$10^{-10}$	$138 \pm 305$	$1.03 \pm 0.02$	$0.93 \pm 0.00$
$\tau_V^{298}$ (ps)	15.0	$21.9 \pm 0.0$	$15.0 \pm 0.0$	$6.6 \pm 0.1$
$\Delta^2$ ( $10^{19}$ s $^{-2}$ )	1.60	$1.00 \pm 0.00^2$	$1.61 \pm 0.00$	$2.04 \pm 0.04$
AF		0.017	0.0055	0.0071

<sup>1</sup>No random errors were included in the synthetic dataset.

<sup>2</sup>Equal to lower boundary imposed on  $\Delta^2$  during fitting.



has rather dramatic consequences for the resulting best-fit parameters. On the other hand the  $-6$  power relationship also means that fitting of NMRD data with inclusion of  $r_{\text{GdH}}$  in the adjustable parameters while  $q$  is fixed at a known value reproduces the value of  $r_{\text{GdH}}$  exactly (see Table S10).

A complicated situation occurs for systems with very fast water exchange rates. For both dissociative and associative water exchange mechanisms, minimal  $\tau_{\text{M}}$  values and hence minimal activation energies may be expected when the energy difference between the two hydration states involved is small (1). Then, both hydration states may be populated, resulting in non-integral  $q$  values. In this situation the Gd–water bond is weakened and accordingly the Gd–O bond may be longer than usual (32), which in turn will have an effect on the value for  $A/\hbar$  for the Gd-bound water O atom. When this parameter is not known,  $^{17}\text{O}$  NMR cannot provide a value for  $q$ .

### 2.2.9. Beyond the approximate SBM theory

The above data were all generated and fitted with the simplest form of the SBM theory, which is the only theory leading to analytical equations. In reality it has been shown that the SBM theory is not generally valid (1,10,12,15). Especially at low frequencies at which transverse electronic relaxation is important, the SBM approximation in general does not lead to correct results, particularly not for the electronic relaxation parameters. Therefore, many researchers now fit  $^1\text{H}$  NMRD data only for  $LF \geq 10$  MHz. Some fittings were performed on the datasets discussed above after deleting the  $^1\text{H}$  NMRD data for  $LF < 10$  MHz (see Table 7). The best-fit parameters obtained for the dataset for  $\tau_{\text{M}}^{298\text{S}} = 10^{-7}\text{ s}$  and  $\tau_{\text{R}}^{298\text{S}} = 10^{-10}\text{ s}$  deviate substantially from the corresponding synthetic parameters. An inspection of the profiles of  $AF$  as a function of the various parameters shows that no minima occur in the profiles for the electronic parameters ( $\tau_{\text{V}}^{298}$  and  $\Delta^2$ ) within the boundaries set on these parameters during the fitting procedure ( $10^{-12}\text{ s} < \tau_{\text{V}}^{298} < 10^{-10}\text{ s}$ ;  $10^{19}\text{ s}^{-2} < \Delta^2 < 2 \times 10^{20}\text{ s}^{-2}$ ). This is due to the low contribution of the electronic relaxation time to the correlation time  $\tau_{\text{di}}$  for the selected rotational correlation time ( $\tau_{\text{R}}^{298\text{S}} = 10^{-10}\text{ s}$ ). The inaccuracy is obviously enhanced by the

small number of  $^1\text{H}$  NMRD data points available at  $LF < 10$  MHz. By contrast, the dataset generated for  $\tau_{\text{M}}^{298\text{S}} = 10^{-7}\text{ s}$  and  $\tau_{\text{R}}^{298\text{S}} = 10^{-8}\text{ s}$  resulted in best-fit parameters that are in good agreement with the corresponding synthetic parameters. This can be rationalized by the relative importance of  $T_{\text{ie}}$  in  $\tau_{\text{di}}$  at slow rotation.

## 3. CONCLUSIONS

The fitting of synthetic datasets described demonstrates that simultaneous fitting of NMRD and  $^{17}\text{O}$  data results in the most reliable best-fit parameters. When only  $^{17}\text{O}$  NMR data are fitted, a strong correlation between  $\tau_{\text{V}}$  and  $\Delta^2$  prevents determination of these parameters individually, although an accurate estimation of the ratio  $\tau_{\text{V}}/\Delta^2$  is still possible. The quality of the best-fit parameters depends strongly on the magnitudes of the various parameters, particularly on  $\tau_{\text{M}}$ . For example, fitting of NMRD data for a single temperature (Method C) gives two local minima of  $AF$  as function of  $\tau_{\text{M}}$ , located symmetrically with respect to the value that gives rise to the  $\tau_{\text{M}}$  value for which  $r_1$  as function  $\tau_{\text{M}}$  of would be maximal. The separation of the two local minima decreases upon approaching the latter maximum, and therefore the accuracy of the best-fit parameter and its deviation of  $\tau_{\text{M}}^{\text{S}}$  decrease as well. For  $\tau_{\text{R}} > 10^{-8}\text{ s}$ ,  $r_1$  is rather insensitive for changes in this parameter, and therefore a reliable determination by fitting is impossible. Plots of  $AF$  as a function of the various adjustable parameters may be very helpful for the identification of the global minimum and to get an impression of the accuracy of the best-fit parameters. In general, the accuracy of the fitting can be improved by reduction of the number of adjustable parameters by fixing them at values independently determined by other techniques. DFT calculations are emerging as an important tool to accurately predict the various parameters governing the relaxivity of CAs, including  $q$ ,  $\tau_{\text{M}}$ ,  $r_{\text{GdH}}$  and  $A/\hbar$  (32). Ultimately, combination of these calculations with fitting of experimental data may be an attractive approach to obtain more insight into the relation between structure and relaxivity.

The synthetic datasets described above were calculated for a very simple model. Local mobility and the presence of second sphere water molecules were not taken into account. If these

**Table 7.** Comparison of best-fit parameters for synthetic data constructed with  $\tau_{\text{M}}^{298\text{S}} = 10^{-7}\text{ s}$  and  $\tau_{\text{R}}^{298\text{S}} = 10^{-10}$  or  $10^{-8}\text{ s}$  as obtained with Procedures A–D. The NMRD data for  $LF < 10$  MHz were excluded during the fittings. Random errors were included in the data

	Synthetic	A	B	C	D
$\tau_{\text{R}}^{298}$ (ps)	100.0	$99.9 \pm 0.0$	$101.5 \pm 0.2$	$274 \pm 3$	$98.4 \pm 0.0$
$\tau_{\text{M}}^{298}$ (s)	$1.00 \times 10^{-7}$	$(1.00 \pm 0.00) \times 10^{-7}$	$(1.76 \pm 0.09) \times 10^{-7}$	$(3.24 \pm 0.01) \times 10^{-7}$	$(0.99 \pm 0.00) \times 10^{-7}$
$\tau_{\text{V}}^{298}$ (ps)	15.0	$73.1 \pm 0.3$	$69 \pm 2$	$1.49 \pm 0.03$	$57 \pm 16$
$\Delta^2$ ( $10^{19}\text{ s}^{-2}$ )	1.60	$4.95 \pm 0.02$	$4.5 \pm 0.2$	$4.2 \pm 0.1$	$5 \pm 2$
$E_{\text{M}}$ ( $\text{kJ mol}^{-1}$ )	47.6	$48.1 \pm 0.0$	$36 \pm 1$	—	$47.6 \pm 0.0$
$E_{\text{R}}$ ( $\text{kJ mol}^{-1}$ )	21.9	$21.5 \pm 0.0$	$22.1 \pm 0.0$	—	$21.7 \pm 0.0$
$AF$	0.0	0.008	0.013	0.014	0.002
$\tau_{\text{R}}^{298}$ (s)	$1.00 \times 10^{-8}$	$(0.95 \pm 0.00) \times 10^{-8}$	$(0.95 \pm 0.00) \times 10^{-8}$	$(0.91 \pm 0.00) \times 10^{-8}$	$(1.00 \pm 0.00) \times 10^{-8}$
$\tau_{\text{M}}^{298}$ (s)	$1.00 \times 10^{-7}$	$(1.02 \pm 0.00) \times 10^{-7}$	$(1.02 \pm 0.00) \times 10^{-7}$	$(1.06 \pm 0.00) \times 10^{-7}$	$(1.00 \pm 0.00) \times 10^{-7}$
$\tau_{\text{V}}^{298}$ (ps)	15.0	$15.6 \pm 0.0$	$15.6 \pm 0.0$	$16.1 \pm 0.0$	$43 \pm 18$
$\Delta^2$ ( $10^{19}\text{ s}^{-2}$ )	1.60	$1.60 \pm 0.00$	$1.60 \pm 0.00$	$1.62 \pm 0.0$	$4 \pm 2$
$E_{\text{M}}$ ( $\text{kJ mol}^{-1}$ )	47.6	$48.5 \pm 0.0$	$48.6 \pm 0.0$	—	$47.7 \pm 0.0$
$E_{\text{R}}$ ( $\text{kJ mol}^{-1}$ )	21.9	$21.6 \pm 0.0$	$21.6 \pm 0.0$	—	$21.9 \pm 0.0$
$AF$	0.0	0.019	0.020	0.014	0.002

phenomena have to be considered, additional adjustable parameters are needed in the fitting procedure, which further reduce the reliability of the best-fit parameters.

Furthermore, it should be noted that the best-fit values of  $\tau_v$  and  $\Delta^2$  often are physically not very meaningful due to (i) the strong correlation between these parameters in  $^{17}\text{O}$  relaxation data and (ii) the inadequacy of the SBM theory for the description of the electronic relaxation, particularly at low magnetic field strengths. Physically meaningful values for these parameters can only be evaluated if EPR data are included and if more advanced models are used to describe the electronic relaxation (15). However, a wealth of data has been published in the literature, which has been evaluated by applying the SBM theory on NMR data only. The trends in these data are generally consistent and have been appeared to be very useful for the rational design of more efficient CAs. The values of the electronic parameters obtained in this way should however be considered as effective rather than as physically meaningful.

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