Dendrimeric calcium-sensitive MRI probes: the first low-field relaxometric study

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Different classes of small- or nano-sized calcium-sensitive probes for magnetic resonance imaging (MRI) have been proposed in the last two decades. These compounds have been developed mainly for functional MRI purposes and tested in vivo in different animal models. Most of them are paramagnetic systems that change their relaxivity in the presence of the divalent ion calcium, resulting in increased $T_1$ or $T_2$ contrast. In this work, we report the investigation of their relaxometric behavior at low magnetic fields, specifically the comparison of the monomeric Ca-sensitive probe and the corresponding dendrimer conjugates of generations 0, 1 and 2 (G0, G1 and G2, respectively). As a result, a relaxivity hump between 10 and 100 MHz of the Larmor frequency progressively appeared with an increase in the size of the investigated contrast agent, indicative of a restricted rotational motion of the complexes as long as the size of the molecule increases. The same trend with a more pronounced effect was detectable in the presence of calcium. The relaxivity enhancement for the Ca$^{2+}$ adducts, primarily caused by an increase of the hydration state of Gd$^{3+}$, went from ca. 130% for the monomeric probe to ca. 310% for the G2 dendrimer conjugate at 0.5 T and 25 °C. $T_1$ weighted magnetic resonance images acquired at 1 T displayed the strong ability of these systems to change their contrast according to the presence of calcium at this field, thus laying the basis for promising future in vivo applications.

Introduction

Molecular imaging is gaining increasing attention in clinical and preclinical research due to its ability to non-invasively visualize biological processes at the cellular and molecular levels. Among the various imaging techniques that can be employed to this end, magnetic resonance imaging (MRI) stands out among the rest because of its superlative spatial resolution (μm), ability to distinguish between hard and soft tissues, and the absence of ionizing radiation. However, the low sensitivity associated with MRI requires the use of specific contrast agents able to report on various pathophysiological processes. These agents generally act through contrast-mediated alteration of tissue relaxation-times for the detection and localization of molecular disease markers, cells and therapeutic drugs. Recently, many so-called smart contrast agents (SCAs), which generate contrast as a response to a change in their chemical or physical environment, have been developed. The mechanism of action of SCAs generally resides in either (1) reversible conformational changes that modulate water accessibility to the agents’ metal ions (e.g. changes in the number of coordinated water molecules in the inner-coordination sphere or in water exchange time between the inner-sphere and bulk water molecules) or (2) reversible intermolecular associations that change the agents’ rotational correlation time. It must be taken into account that if these variations are faster than the relatively slow MRI acquisition timeframe, an averaging effect can be obtained, further limiting the sensitivity of the agent’s response. SCAs can be sensitive to pH-variations, enzymatic activity, glucose uptake, lactate or nitric oxide concentrations, partial oxygen pressure, the redox status and metal ion coordination states or concentrations. Among SCAs responsive to metal ions, compounds sensitive to copper, zinc and calcium have been mainly developed. Monitoring metal ions by MRI, in fact, requires at least an intracellular metal ion concentration in the micromolar range, due to the intrinsic low sensitivity of the technique. Calcium and zinc, the first and second most abundant metal ions in the body, respectively, are the most attractive examples due to their sufficient concentration and irreplaceable role in neuronal signalling, muscle contraction, and enzymatic reactions (extracellular magnesium concentration is also in the micromolar range; however, this metal
has been less studied, as its concentration remains constant during neuronal activity. Calcium imaging, in particular, attracts considerable interest for functional MRI applications as it can aid tracking the neuronal activity in vivo. SCAs designed for calcium real-time MRI are generally composed of a calcium chelator linked to one or two MR reporting moieties. The first paramagnetic calcium sensitive compound developed was based on the common BAPTA (1,2-bis(o-aminophenoxo)ethyl-N,N,N’,N’-tetraacetic acid) calcium chelator attached to two units of the highly paramagnetic Gd3+ complex of 1,4,7,10 tetraazacyclododecane-1,4,7-tricarboxylic acid (Gd-DOTA).28 Subsequently, SCAs composed of a modified EGTA chelator linked to two macrocyclic DOTA-derived moieties that bear a pair of Gd3+ ions were synthesized, resulting in excellent relaxivity performance. A recent study with this smart probe demonstrated the capability to detect and monitor the time course of cerebral ischemia, thus announcing the exciting prospects of this class of functional biomarkers for future advancements in molecular imaging.

Development of even more potent SCAs with appropriate biocompatibility is desired in order to progress further in functional MRI. To this end, existing probes may undergo synthetic modifications that lead to improved properties for their future application. In such cases, the EGTA-derived chelator remains a standard building block for the preparation of Ca-sensitive SCAs, due to its very high selectivity towards calcium. On the other hand, the existence of two macrocyclic chelators provides sufficient flexibility for various structural alterations in SCAs, which may result in different types of multimeric and nanosized systems. Following the initial preparation of the potent SCA mentioned above, this bifunctional chelator was synthetically modified and coupled to the poly(amidoamine) (PAMAM) dendrimer, aiming to modulate the pharmacokinetic properties of the SCA. The obtained system retained calcium sensitivity and displayed a slower diffusion in rat cerebral cortex in comparison to the monomeric equivalent, thus suggesting that dendrimeric systems can have great advantages over monomeric ones. On this wave, a calcium bioresponsive generation 4 (G4) dendrimer was developed and tested at high field (7 T). Thanks to the variations of the \( r_2/r_1 \) ratio at different calcium concentrations, an almost four times greater signal gain per unit of time as compared to conventional \( T_1 \) weighted imaging with small sized contrast agents was obtained. Recently, a series of novel dendrimeric calcium sensitive MRI probes were designed and examined in terms of size change, electrostatic behavior and relaxometric performance at high field (7 T). The study was carried out on different G4 dendrimers based on D3A-, D3A-monoamide or DOTA (1,4,7,10 tetraazacyclododecane-1,4,7,10-tetraacetic acid) complexes and indicated that a combination of changes in size, hydration number, and rigidity of the system is required to maximize an increase in the \( r_2/r_1 \) ratio change.

Interestingly, all these macromolecular systems reported so far were assessed at high magnetic field (7 T). Although a specific set of properties should be expected due to their large molecular size, almost none of them were studied at low fields, except for the liposome-based SCA. In the present work, we aimed to add more insights into the relaxometric behaviour of calcium sensitive dendrimeric contrast agents at magnetic fields that are closer to the ones currently more diffuse in clinics (1.5 and 3 T). More specifically, we coupled monomeric SCAs to three different generations of dendrimers, G0, G1 and G2, and compared their properties in terms of calcium responsivity at 0.5 and 1 T.

**Materials and methods**

**Synthesis of DCAs**

Dendrimeric contrast agents (DCAs) were prepared following the previously reported procedure for G1 dendrimers. Specifically, a protected monomeric ligand L was coupled via an alkylisothiocyanate group to the amine surface Starburst® PAMAM G0 (4 amine surface groups), G1 (8 amine surface groups) or G2 (16 amine surface groups) dendrimers to yield the dendrimer conjugates D0, D1 and D2, respectively. Their hydrolysis in formic acid provided the dendrimeric conjugate ligands D0b, D1b and D2b, which were complexed with Gd3+ to result in dendrimeric contrast agents DCA0, DCA1 and DCA2, respectively. Dendrimers D0-2 and D0-2b were purified by size-exclusion chromatography using Sephadex® LH-20 (bead size: 25–100 μm) and methanol or Sephadex® G-15 (bead size: 40–120 μm) and water as an eluent, respectively. DCA0-2 samples were treated with ethylenediaminetetraacetic acid (EDTA) to remove the excess of Gd3+, while formed GdEDTA and remaining EDTA were removed from DCA0-2 by centrifugation using centrifugal filter units (3 kDa molecular weight cut-off). Following the same literature procedure, the monomeric Gd3+ complex GdL was also prepared and its properties were compared to those of DCA0-2. Purity of GdL and DCA0-2 dendrimers was assessed by the Evans method on a Bruker Avance 600 spectrometer equipped with a 5 mm probe and a standard temperature control unit.

**MALDI analysis**

MALDI-ToF spectra were recorded at the Center for Metabolomics and Mass Spectrometry of The Scripps Research Institute, La Jolla, USA. All DCA0-2 samples were obtained as a mixture of dendrimeric products with different numbers of monomeric SCA units coupled to the dendrimer surface. Analytical data for DCA1 were reported previously. DCA0: a mixture of products with 3 and 4 SCA monomeric units. With 3 SCA units: \([M + 3Na + 3H_2O]^-\) celled for C142H228Gd3N37Na3O46S3(H2O)3 8880, found 8878. With 4 SCA units: \([M + 4Na]^-\) celled for C185H336Gd4N40Na4O60S4 4929, found 4930.

DCA2: the spectrum with broad signals indicated the presence of multiple species, and the average number of SCA monomeric units was 9. \([M + 14Na + 3H_2O]^-\) celled for C502H828Gd9N139Na14O154S9(H2O)34 13911, found 13911.

**r2 titrations with CaCl2**

GdL, DCA0, DCA1 and DCA2 were dissolved in HEPES buffer at 0.70 mM Gd3+ concentration and the pH was adjusted to 7.0 ± 0.1. Starting longitudinal relaxivity \( r_{1g} \) was measured at 21.5 MHz (0.5 T) and 25 °C on a Stelar Spinmaster (Stelar S. R. L., Mede (PV), Italy). Stepwise additions of 10 mM CaCl2 solution were
then carried out for each sample, until a Ca\(^{2+}\):Gd\(^{3+}\) ratio of 2.5 was reached. Variations in longitudinal relaxivity \(r_{1,\text{obs}}\) at 21.5 MHz were monitored 15 min after each addition to let the system equilibrate. The pH was maintained constant at 7.0 ± 0.1 before each measurement and addition. Nuclear magnetic relaxation dispersion (NMRD) profiles were acquired before and after the addition of CaCl\(_2\). \(^1\)H NMRD profiles were recorded at 25 °C on a Stolar Field Cycling relaxometer (FFC2000), operating at a magnetic field strength ranging from 0.01 to 20 MHz (2.4 × 10\(^{-4}\) to 0.47 T). The additional points measured in the range 20–70 MHz (0.47–1.65 T) were acquired on a tunable magnet (Stolar). The uncertainty of \(T_1\) measurements was ±1%. For each measurement, the longitudinal relaxation \(r_{1,p}\) was calculated according to eqn (1).

\[
r_{1,p} = \frac{R_{1,\text{obs}} - R_{1,0}}{[\text{Gd}^{3+}\text{]}^n}
\]  

\(r_1\) titration with EDTA

GdL, DCA0, DCA1 and DCA2 were dissolved in HEPES buffer at 0.70 mM Gd\(^{3+}\) concentration and the pH was adjusted to 6.0 ± 0.1. Starting longitudinal relaxivity was measured at 21.5 MHz (25 °C), and then stepwise additions of 20 mM EDTA solution were carried out for each sample. Variations in longitudinal relaxivity at 21.5 MHz were monitored 15 min after each addition. The pH was maintained constant at 6.0 ± 0.1 before each measurement and addition. When a molar concentration ratio of EDTA : Gd\(^{3+}\) of 3 : 1 was reached, samples were exhaustively dialysed against isotonic HEPES/NaCl buffer (pH 7.3) to remove Gd-EDTA and the excess of EDTA ligand. Then the amount of residual Gd\(^{3+}\) was relaxometrically determined through the glass vial test. Briefly, aliquots of each dialysed sample were diluted 1 : 1 in volume with HCl 37%, transferred to pre-scored glass ampoules and centrifuged for 3 min at 2000 rpm. Then the glass vials were sealed and placed at 120 °C for 16 hours to achieve the complete mineralization of the samples. Longitudinal relaxivity \(R_{1,\text{obs}}\) was then measured and the amount of residual gadolinium was determined according to eqn (2), where \(R_{1,\text{dia}}\) is the diamagnetic contribution of water \((R_{1,\text{dia}} = 0.5 \text{s}^{-1})\). Finally, titration with CaCl\(_2\) was performed as reported in the previous paragraph.

\[
[\text{Gd}^{3+}\text{]} = \frac{R_{1,\text{obs}} - R_{1,\text{dia}}}{13.7} \times 2
\]  

\(\text{Variable-temperature } ^{17}\text{O NMR measurements}\)

Variable-temperature \(^{17}\text{O NMR measurements of GdL were recorded on a Bruker Avance 600 spectrometer equipped with a 5 mm probe and a standard temperature control unit. The GdL sample contained 5\% of } ^{17}\text{O water (Cambridge Isotope) and 12.76 mM Gd}^{3+}\text{. The paramagnetic contribution was calculated using HEPES buffer containing 5\% of } ^{17}\text{O water. } ^{17}\text{O NMR measurements were also performed after calcium addition (Gd:Ca ratio 1.5 : 1) both on the GdL sample and on the control suspension (19.14 mM CaCl}_2\text{). The observed transverse relaxation rates were calculated from the signal width at half-height. Measurements were performed in a temperature interval ranging from 278 to 343 K.}\)

\textbf{Analysis of relaxometric data}

NMRD profiles were analyzed using a single relaxation model based on Solomon–Bloembergen–Morgan and Freed theories.\(^{36}\)

The model considered the relaxivity as the sum of three contributions arising from (i) the water protons in the inner-sphere (is) of Gd\(^{3+}\), (ii) the water protons in the second-coordination sphere (ss) of Gd\(^{3+}\), and (iii) the water protons that diffuse around the paramagnetic species (outer sphere, os) and do not interact with any molecular site of the SCA.

\[
r_1 = r_1^{\text{is}} + r_1^{\text{ss}} + r_1^{\text{os}}
\]  

The inner sphere contribution can be described as

\[
r_1^{\text{is}} = 1.8 \times 10^{-5} q^6 \frac{1}{T_1^M + \tau_M} T_1^{M(\omega^2,\nu_C,\nu_R)}
\]  

where \(q\) is the number of water protons coordinated to Gd\(^{3+}\), \(\tau_M\) is the residence lifetime of the coordinated water protons, \(1.8 \times 10^{-5}\) is the molar fraction of water protons in the inner sphere for a solution containing 1 mM of Gd\(^{3+}\), and \(T_1^M\) is the relaxation time of the water protons when they are bound at the metal site. \(T_1^M\) is defined by several dynamic and structural factors like the proton–metal distance \(r\), the relaxation rate of the unpaired electrons of the metal described by the parameters \(A^2\) and \(\tau_V\), and the reorientational correlation time \(\tau_R\).

The second-sphere contribution is supposed to follow a similar model:

\[
r_1^{\text{ss}} = 1.8 \times 10^{-5} q^6 \frac{1}{T_1^M} T_1^{M(\omega^2,\nu_C,\nu_R)}
\]

where the meaning of most of the parameters is unvaried with respect to the is contribution, but represents an average of the behavior of the water protons in the second-coordination sphere of the metal.

The contribution from the outer-sphere water protons can be modeled as

\[
r_1^{\text{os}} = C^{\text{os}} \times \left(1 \times 10^{-3}\right)^a \frac{1}{D} \times \left[7J_{(os)} + 3J_{(ad)}\right]
\]

where \(a\) is the distance of maximum approach of the diffusing water protons to the metal ion, \(D\) is the relative solute–solvent diffusion constant, and \(C^{\text{os}}\) is a physical constant, whereas the spectral density functions include the terms controlling the electronic relaxation rate.

The analysis of the NMRD data for the calcium-free agents was carried out putting \(q = 0\) and, therefore, only second- and outer-sphere contributions were considered. The fitting was performed keeping constant the values of \(a\) and \(D\) (3.8 Å and 2.24 × 10\(^{-5}\), respectively, as determined for most of the Gd-based agents reported in the literature).

For the calcium-adducts, \(q\) was switched to 1 and the inner-sphere contribution was considered. This mechanism introduced \(\tau_R\) and \(\tau_M\), which were varied, and \(r\), which was kept fixed at 3 Å.
Magnetic resonance imaging

Magnetic resonance imaging (MRI) is a non-invasive imaging technique that uses strong magnetic fields and radio waves to create detailed images of internal structures. It is widely used in medical diagnostics and research. The primary journal article discusses the use of PAMAM dendrimers in MRI imaging.

In the study, PAMAM dendrimers were used as contrast agents to enhance the visibility of specific regions in the body. The authors report that these dendrimers can be tailored to target specific biological markers and improve the sensitivity and specificity of MRI images.

The research highlights the potential of dendrimers as multifunctional contrast agents, offering improved tissue contrast and potentially revolutionizing medical imaging. The findings suggest that PAMAM dendrimers could play a significant role in the development of more effective and targeted diagnostic tools in the future.
Interestingly, the \( r_1 \) value measured for DCA2, 28.8 \( \text{mM}^{-1} \text{s}^{-1} \), was not too far from the corresponding value reported under the same experimental conditions using Ca-sensitive liposomes (37.0 \( \text{mM}^{-1} \text{s}^{-1} \)). However, as it has been widely reported previously, the relaxivity obtained at 0.5 T for a slowly tumbling agent is substantially higher than that measured for the same system at high field, e.g., at 7 T. Furthermore, a drop in relaxivity at high field can be quite prominent to make this agent less efficient than a fast tumbling agent. Representative examples are agents GdL and DCA1, whose longitudinal relaxivities for the Ca-adducts were measured at 7 T; in turn, the relaxivity of the monomer was slightly higher than that of the dendrimer (7.4 \( \text{mM}^{-1} \text{s}^{-1} \) vs. 6.5 \( \text{mM}^{-1} \text{s}^{-1} \)). In this work, the relaxivity values for the two Ca-adducts were measured at 0.5 T, and, as expected, DCA1 displayed a relaxivity much higher than that of the GdL (20 \( \text{mM}^{-1} \text{s}^{-1} \) vs. 6.5 \( \text{mM}^{-1} \text{s}^{-1} \)). Hence, these observations highlight the relevant role played by the magnetic field of the MRI scanner in defining the detection sensitivity of these calcium-sensitive CAs. Apart from pharmacokinetic considerations, large-sized agents may be superior at lower fields, whereas small-sized agents may be superior at higher fields.

The stability of the Gd\(^{3+}\) complexation for the monomer and dendrimers was monitored by measuring the relaxivity upon addition of EDTA. Almost no variations in pH values were detected and no decrease in relaxivity was observed, thus suggesting that EDTA, as the competing ligand, did not cause destruction of the initially prepared Gd-complexes, i.e., GdL and DCA0-2 (Fig. 3).

In the next step, we dialysed DCA0-2 (note: GdL cannot be collected due to the large pore size of the membrane) and titration with Ca\(^{2+}\) was carried out again. As expected, no significant differences in the DCA0-2 response towards Ca\(^{2+}\) were detected before and after treatment with EDTA (Fig. 4). These results confirm that DCA0-2 remained intact upon the treatment with EDTA, thereby also excluding the transmetallation of Gd\(^{3+}\) and the presence of GdEDTA species.

### Variable temperature \(^{17}\text{O\! NMR and NMRD experiments\!}

The ditopic ligand L, characteristic of the Ca-sensitive SCAs investigated in this work, contains a macrocyclic cage designed for hosting the Gd\(^{3+}\) ion responsible for the generation of the MRI contrast, which is linked to a bis-amide EGTA-derived ligand for the successive coordination of the Ca\(^{2+}\) ion. For the GdL, the coordination of Gd\(^{3+}\) (coordination number 9) is expected to be

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**Fig. 1** Structures of the monomeric complex GdL and illustrations of the dendrimeric conjugates DCA0, DCA1 and DCA2 studied in this work.

**Fig. 2** Relaxometric titrations of GdL, DCA0, DCA1 and DCA2 with CaCl\(_2\) at 0.5 T (25 °C), [Gd\(^{3+}\) ] = 0.7 mM. Dashed lines connect the experimentally observed values and provide better visualization of the observed \( r_1 \) changes.

**Fig. 3** Relaxometric titrations of the monomer and GdL, DCA0, DCA1 and DCA2 with EDTA, at 0.5 T (25 °C), pH 6.0 ± 0.1, [Gd\(^{3+}\) ] = 0.7 mM. Dashed lines connect the experimentally observed values and provide better visualization of the observed \( r_1 \) changes.
saturated by the 4 nitrogens of the macrocycle, the 3 oxygens of the pendant carboxylates, and the 2 oxygens coming from one amide and one carboxylate group on the proximal end (with respect to the macrocyclic cage) of the EGTA-derived ligand. Such coordination behavior was already observed for different mono- and bismacroyclic complexes that bear the same type of the EGTA-derived chelator.29,46

The confirmation of the GdL not having a water molecule coordinated to the metal (i.e. it acts as a non-hydrated, \( q = 0 \) complex) was gained from the acquisition of the temperature-dependent transverse relaxation rates of water \( ^{17}\text{O} \) nuclei in a solution of the GdL (Fig. 5, red circles). This technique is the first choice to get information about the number and the dynamics of the water molecules coordinated to a paramagnetic center. In fact, the \( ^{17}\text{O} \) transverse relaxation rate of water is proportional to the fraction of the water molecule bound to the metal, and to the number \( q \) of metal coordinated water. Furthermore, it is modulated by several variables such as the \( T_2 \) of the water molecule at the metal site \( T_{2M} \), the chemical shift of the water oxygen induced by the metal \( \Delta \omega_{M}^{2} \), and the residence lifetime of the water molecule in the inner coordination sphere \( \tau_{M} \). However, due to the very small chemical shift effect induced by Gd(m), the relaxation model simplifies in a form already seen for the inner-sphere contribution to the \(^1\text{H} \) paramagnetic relaxation (eqn (4)), though here the predominant relaxation mechanism arises from contact interaction:

\[
R_{2p} = \frac{[\text{GdL}]}{55.56} \frac{1}{T_{2M} + \tau_{M}} T_{2M(c=4,8)} \quad (8)
\]

where \( \tau_{M} \) and \( A/h \) refer to the electronic relaxation time of the unpaired electrons of the metal and the Gd–\(^{17}\text{O} \) coupling constant, respectively.

Hence, the occurrence of very low \( ^{17}\text{O}-R_{2p} \) values (as measured for GdL, Fig. 5) indicates the absence of the metal bound water molecule (\( q = 0 \)). However, the paramagnetic contribution is not null due to the presence of small dipolar effects from outer- and second-sphere contributions.

On the other hand, for \( q > 0 \), the temperature dependence of \( ^{17}\text{O}-R_{2p} \) values reflects the balance between \( T_{2M} \) and \( \tau_{M} \) values, which show an opposite temperature dependence.

Fig. 5 (blue diamonds) indicates that the complexation of calcium causes a significant increase in the \( ^{17}\text{O} \) water transverse relaxation rate, which is fully consistent with an increase in the hydration state of Gd\(^{3+} \) from 0 to 1. To verify this hypothesis, the NMRD profiles were analyzed with the relaxation model that includes the second-sphere and outer-sphere water protons for the calcium-free compounds, whereas for the samples where Ca\(^{2+} \) was added, the contribution from inner-sphere water protons was also considered (see below).

The values of the initial longitudinal relaxivities (i.e. in the absence of Ca\(^{2+} \), Table 1) for GdL and DCA0-2 show that there is a direct correlation between the relaxivity and the molecular size of the SCAs as previously reported in other studies.47,48 Furthermore, the value measured for GdL appears to be quite lower with respect to Gd(m) complexes with a similar size, but \( q = 1 \), thus confirming the qualitative result (\( q = 0 \)) obtained from the analysis of the \( ^{17}\text{O} \) data reported above. Since even the relaxivities of the dendrimer-based complexes are consistent with that of GdL (considering the differences in molecular size), it is reasonable to assume that also DCA0-2 species lack inner-sphere water molecules.

Based on these observations, the NMRD profile of GdL (Fig. 6, red circles) was analyzed using a relaxation model lacking the contribution from inner-sphere water protons. However, as the relaxivity of the complex was significantly higher than the numbers reported for “pure” outer-sphere Gd\(^{3+} \) complexes (3.9 mM\(^{-1} \) s\(^{-1} \) vs. 2.2–2.3 mM\(^{-1} \) s\(^{-1} \) at 20 MHz and 25 °C),\(^{49} \) the NMRD data were analyzed adding a contribution arising from the so-called “second-sphere” water protons. These are exchangeable protons relatively close to the metal center (typically within 5 Å), which intensively interact with polar moieties of the ligand.\(^{50} \)

**Table 1** Values of longitudinal relaxivity determined at 25 °C and 21.5 MHz. The values were normalized to the Gd\(^{3+} \) concentration

<table>
<thead>
<tr>
<th>Sample</th>
<th>( r_1 ) (mM(^{-1} ) s(^{-1} ))</th>
</tr>
</thead>
<tbody>
<tr>
<td>GdL</td>
<td>3.9</td>
</tr>
<tr>
<td>DCA0</td>
<td>5.1</td>
</tr>
<tr>
<td>DCA1</td>
<td>6.7</td>
</tr>
<tr>
<td>DCA2</td>
<td>7.3</td>
</tr>
</tbody>
</table>
The obtained results supported previous observations that GdL and its analogues are indeed $q = 0$ complexes, with substantial contribution of the second-sphere water to the overall relaxivity value.

This contribution can be modeled using a set of equations very similar to those used for describing the inner-sphere contribution (see the Experimental section), with an average number of contributing water protons ($q_{\text{2sf}}$), their average distance from Gd$^{3+}$ ($r_{\text{2sf}}$), and an overall correlation time ($\tau_{\text{C}}$) that modulates the dipolar interaction and encloses the role of the rotation and exchange dynamics of the second sphere protons. As the same ditopic ligand was used to get the other three dendrimeric SCAs investigated in this work, an identical relaxation model was applied for the analysis of all the NMRD data.

Besides the three parameters that describe the second-sphere contribution, the fitting was carried out varying the parameters describing the electronic relaxation rate of the unpaired electrons of Gd$^{3+}$ according to the Bloembergen–Morgan theory, such as the mean energy of the transient zero field splitting $\Delta$ and the corresponding time modulation $\tau_{\text{r}}$. Conversely, the other two parameters defining the outer-sphere contribution according to Freed’s model, i.e., the distance of maximum approach $a$ and the relative diffusion constant $D$, were kept fixed to the typical values for this type of Gd$^{3+}$ chelate, i.e. 3.8 Å and 2.24 cm$^2$ s$^{-1}$, respectively.

The quantitative analysis of the NMRD profiles according to the above-described relaxation model allowed the estimation of the relevant parameters controlling the relaxivity of the investigated systems (Table 2). The results obtained highlight the difference in the set of relaxation parameters between the monomeric complex GdL and the dendrimeric agent DCA0-2. This difference is mainly observed for the parameters determining the electronic relaxation rate, the number of second sphere protons, and their correlation times. On the other side, the main difference within the group of the three dendrimers relies on the number of the second-sphere protons, which progressively increases from G0 to G2.

To summarize, the coupling of the GdL monomer to PAMAM dendrimers to form calcium-sensitive paramagnetic dendrimer conjugates leads to a small (ca. 20%, from 190 ps to 230 ps) increase in the low-field limit ($\tau_{\text{500}}$) for the electronic relaxation time for the unpaired electrons of Gd$^{3+}$. In addition, dendrimers display a longer second-sphere correlation time that is likely mainly responsible for the relaxivity hump characteristic of the DCA0-2 profiles. Likely, the elongation of $\tau_{\text{C}}$ originates from the slow down of the rotational motion of the complexes when they are part of the dendrimers.

However, the increase of relaxivity going from DCA0 to DCA2 does not seem attributable to $\tau_{\text{C}}$ that is almost unvaried along the series. Likely, the rotation of the chelates is only very weakly coupled to the tumbling motion of the dendrimeric core, and therefore, $\tau_{\text{C}}$ results are rather uncorrelated to the whole size of the system.

Hence, what appears to be mainly responsible for the relaxation enhancement within the dendrimeric series is the number of second-sphere proton protons that increases from 1.9 to 2.6.

However, the Gd$^{3+}$ chelates have the same structure and they are located at the terminal end of the dendrimer branches, thus making it difficult to find an explanation for the increase in $q_{\text{2sf}}$ values. One might speculate that the increase in the dendrimer generation (and consequently in the number of branches) may somehow affect the structure/dynamics of the water around the terminal end of the branches, thus leading to a little increase of the number of water protons in the second sphere of the paramagnetic ion.

Another observation that can be noted from the data presented in Table 1 and the NMRD profiles in Fig. 6 deals with the relaxivity jump among the investigated SCAs: $r_1$ increases for 1.2 mM$^{-1}$ s$^{-1}$ (~30%) between the monomer and DCA0, 1.6 mM$^{-1}$ s$^{-1}$ (~30%) between DCA0 and DCA1, but only 0.6 mM$^{-1}$ s$^{-1}$ (~10%) between DCA1 and DCA2. Interestingly, a similar result was reported in a previous study carried out by Bryant et al., where a plateau in relaxivity values was reached by

![Fig. 6](image)
G7 dendrimers and was maintained for G9 and G10 dendrimers. In principle, a reduced relaxivity enhancement upon increasing the size could be the consequence of a long residence lifetime of the inner-sphere water molecule, but if this hypothesis could explain the data cited above, it cannot be applied to this work due to the lack of inner-sphere water in the agents investigated here. As the relaxivity of the calcium-free SCAs examined in this work is primarily determined by $g^{2nd}$, the observed changes within the series are nicely correlated with the number of second-sphere water protons reported in Table 2: the increase in $g^{2nd}$ is $\sim 27\%$ from GdL to DCA0, $\sim 26\%$ from DCA0 to DCA1, and $\sim 8\%$ from DCA1 to DCA2.

To understand the relaxation mechanisms associated with the calcium coordination of the investigated SCAs, the NMRD profiles at 25 °C for all the examined species after addition of CaCl$_2$ were measured (Fig. 7) and integrated with the temperature-dependent experiments of water $^{17}$O-R$_{2p}$ performed with GdL (Fig. 5, blue diamonds).

As anticipated in the discussion of the $^{17}$O-water data shown in Fig. 5, the significant relaxation enhancement observed when calcium was added to GdL is attributable to a $q = 0 \rightarrow q = 1$ hydration switch. Since the ditopic structure of the chelates in the dendrimers is equal to that of GdL, the NMRD data of the Ca$^{2+}$-adducts were analyzed with the same relaxation model used for the calcium-free systems, but implemented with a model describing the contribution from inner-sphere water protons.

The NMRD data confirmed the significant relaxivity enhancement observed for all the Gd$^{3+}$ complexes upon Ca$^{2+}$ coordination over the entire magnetic field window examined. Particularly clear is the increase of relaxivity in the low-to-intermediate field range, where $r_1$ is mostly dependent on the rotational motion of the system, especially for the Gd$^{3+}$ complexes having at least one inner-sphere water coordinated to the metal center. Actually, the relaxivity followed the order of the molecular size of the SCAs, i.e. GdL < DCA0 < DCA1 < DCA2. The good quality of the quantitative analysis of the profiles confirmed the goodness of the selected relaxation model, and, consequently, provided a support to the occurrence of the $q = 0/q = 1$ switch at the Gd$^{3+}$ site triggered by the coordination of Ca$^{2+}$ ions.

As expected, the correlation time associated with the rotational motion, $\tau_B$, was found to be proportional to the overall size of the SCAs, with limit values of 80 ps for the Ca-GdL and 9.1 ns (i.e. two orders of magnitude larger) for Ca-DCA2. By considering that the $^{17}$O-R$_{2p}$ profile reported in Fig. 5 for the Ca$^{2+}$ adduct is completely consistent with a short $\tau_M$ value and that the relaxivity of all the Ca$^{2+}$ adducts examined in this work decreases from 25 °C to 37 °C (data not shown), the NMRD profiles of the Gd-complexes coordinated with Ca$^{2+}$ have been analyzed fixing the $\tau_M$ value to 50 ns.

**MRI on tube phantoms at low magnetic field**

$T_{1w}$ and $T_{2w}$ MR images were acquired at 1T with an Aspect M2 MRI System (Fig. 8a and b). To this end, GdL, DCA0, DCA1 and DCA2 were suspended in HEPES buffer at a Gd$^{3+}$ concentration of 0.57 mM. The same amount of complexes was also separately incubated with CaCl$_2$ (Ca:Gd 1.5:1) and then suspended in HEPES buffer. $T_1$ contrast observed by MRI matched the NMRD profile results, displaying increasing signals from GdL to DCA2, both in the absence and presence of calcium. Moreover, the difference in intensity of the $T_{1w}$ signal between the two states...
(without and with Ca$^{2+}$) was clearly the highest for DCA2. This strong $T_1$ contrast observed for the investigated dendrimeric systems after interaction with Ca$^{2+}$ could be considered extremely promising for future in vitro and in vivo applications. Similarly, changes in the $T_{2w}$ contrast expectedly led to darkening of the recorded MR images (i.e. $r_2$ relaxivity also increases along with the increase in $r_1$).$^{31,33}$ Hence, a $T_2$ contrast enhancement was also observed for the dendrimer systems along with increasing generations upon addition of Ca$^{2+}$. However, changes in $T_{1w}$ contrast completely followed the increase in the size of the investigated SCAs; on the other hand, the change in $T_{2w}$ contrast was pronounced between the GdL and already the first dendrimer conjugate DCA0, while the remaining dendrimer conjugates displayed similar properties and resulted in constant $T_{2w}$ contrast irrespective of the DCA0-2 size (Fig. 8c).

Conclusions

In this work, we prepared a series of four SCAs with different sizes, consisting of a typical monomacroyclic, DO3A-derived SCA and a triplet of dendrimeric conjugates of different generation (G0–2) bearing the same monomeric SCA unit. We thoroughly investigated the effect of the molecular size on the relaxometric properties of these systems at low magnetic fields. Combined NMRD and $^{17}$O NMR studies indicated the existence of non-hydrated species with substantial contribution from second-sphere and outer-sphere water for these complexes. Relaxometric titrations in the presence of Ca$^{2+}$ showed strong response of the investigated systems towards this cation. Moreover, the magnitude of enhancement in longitudinal relaxivity was strongly correlated with the size of complexes, with the biggest G2 dendrimer SCA conjugate exhibiting the strongest $r_1$ response in the presence of Ca$^{2+}$. The results reported in this study confirm once again the similarity properties and SCAs smart contrast agents.

List of abbreviations

<table>
<thead>
<tr>
<th>Abbreviation</th>
<th>Description</th>
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<tbody>
<tr>
<td>BAPTA</td>
<td>1,2-Bis(o-aminophenoxy)ethane-N,N,N',N'-tetraacetic acid</td>
</tr>
<tr>
<td>CAs</td>
<td>Contrast agents</td>
</tr>
<tr>
<td>CNR</td>
<td>Contrast to noise ratio</td>
</tr>
<tr>
<td>DCAs</td>
<td>Dendrimeric contrast agents</td>
</tr>
<tr>
<td>DCA0</td>
<td>G0 dendrimeric contrast agent</td>
</tr>
<tr>
<td>DCA1</td>
<td>G1 dendrimeric contrast agent</td>
</tr>
<tr>
<td>DCA2</td>
<td>G2 dendrimeric contrast agent</td>
</tr>
<tr>
<td>EDTA</td>
<td>Ethylenediaminetetraacetic acid</td>
</tr>
<tr>
<td>EGTA</td>
<td>Ethylene glycol-bis(β-aminoethyl ether)-N,N,N',N'-tetraacetic acid</td>
</tr>
<tr>
<td>FOV</td>
<td>Field of view</td>
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<td>Gd-DOTA</td>
<td>Gd$^{3+}$-1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid</td>
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<td>GdL</td>
<td>Monomeric Gd$^{3+}$ complex</td>
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<tr>
<td>HEPES</td>
<td>4-(2-Hydroxyethyl)-1-piperazineethanesulfonic acid</td>
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<tr>
<td>is</td>
<td>Inner sphere</td>
</tr>
<tr>
<td>MRI</td>
<td>Magnetic resonance imaging</td>
</tr>
<tr>
<td>NAV</td>
<td>Number of averages</td>
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<tr>
<td>NMR</td>
<td>Nuclear magnetic resonance</td>
</tr>
<tr>
<td>NMRD</td>
<td>Nuclear magnetic relaxation dispersion</td>
</tr>
<tr>
<td>os</td>
<td>Outer sphere</td>
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<tr>
<td>PAMAM</td>
<td>Poly(amidoamine)</td>
</tr>
<tr>
<td>r$_1$</td>
<td>Longitudinal relaxivity</td>
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<td>SCAs</td>
<td>Smart contrast agents</td>
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<tr>
<td>SD</td>
<td>Standard deviation</td>
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<tr>
<td>SI</td>
<td>Signal intensity</td>
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<tr>
<td>ss</td>
<td>Second-coordination sphere</td>
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<td>$T_{1w}$</td>
<td>$T_1$ weighted</td>
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<td>TE</td>
<td>Echo time</td>
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<td>TR</td>
<td>Repetition time</td>
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Conflicts of interest

There are no conflicts to declare.

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Notes and references


