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# A Cell Permeable Esterase-Activated Ca(II)-Sensitive MRI Contrast Agent

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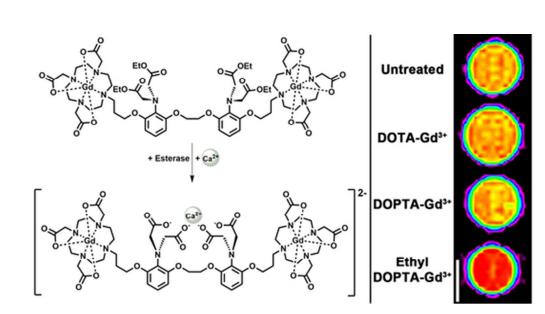
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Despite the importance of Ca(II) in physiology, visualizing Ca(II) flux in the central nervous system (CNS) in vivo remains challenging. To address these challenges we have developed a new generation, cell-permeable Ca(II)-activated MRI contrast probe that utilizes ethyl esters to increase cell permeability and prevent extracellular divalent Ca(II) binding. Upon cellular internalization, the ethyl esters are cleaved allowing the agent to bind intracellular Ca(II) increasing relaxivity and resulting in enhanced positive MR image contrast. 44x23mm (300 x 300 DPI)

## A Cell Permeable Esterase-Activated Ca(II)-Sensitive MRI Contrast Agent

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#### Abstract

Calcium [Ca(II)] is a fundamental transducer of electrical activity in the central nervous system (CNS). Influx of Ca(II) into the cytosol is responsible for action potential initiation and propagation, and initiates inter-neuronal communication via release of neurotransmitters and activation of gene expression. Despite the importance of Ca(II) in physiology, it remains a challenge to visualize Ca(II) flux in the central nervous system (CNS) in vivo. To address these challenges, we have developed a new generation, Ca(II)-activated MRI contrast agent that utilizes ethyl esters to increase cell labeling and prevent extracellular divalent Ca(II) binding. Following labelling, the ethyl esters can be cleaved thus allowing the agent to bind Ca(II), increasing relaxivity and resulting in enhanced positive MR image contrast. The ability of this probe to discriminate between extra- and intracellular Ca(II) may allow for spatiotemporal in vivo imaging of Ca(II) flux during seizures or ischemia where large Ca(II) fluxes  $(1 - 10 \ \mu M)$  can result in cell death.

# Introduction

Gradients and fluctuations in Ca(II) concentration play critical roles in virtually all cellular events including membrane potential maintenance, action potential propagation, signal transduction cascades, and apoptosis.<sup>1-4</sup> In the central nervous system (CNS), Ca(II) flux is

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critical to neuronal function where aberrant fluctuations of intracellular Ca(II) cause a number of pathologies affecting brain activity resulting in seizures, ischemia, and hypoglycemia.<sup>1, 5-10</sup>

The pivotal role of Ca(II) flux in physiology has led to the development of Ca(II) sensitive imaging techniques to investigate the fundamental mechanisms of this important cation. The most common technique for Ca(II) visualization has been the use of optical imaging which relies on the use of Ca(II)-sensitive fluorescent dyes.<sup>11-13</sup> The major challenge for optical techniques is limited penetration depth (~5-10 mm) due to light scattering and tissue absorption. As a result, non-invasive in vivo longitudinal studies using optical techniques such as bioluminescence and two-photon confocal microscopy of Ca(II) flux are challenging.<sup>11</sup> More recent methods have combined electrophysiology or fiber optic Ca(II) imaging with MRI to record neural activity with Ca(II) flux simultaneously but are technically challenging and have limited clinical potential.<sup>14</sup> Additionally, the acquired data is in the form of maps that are not in real time and are fundamentally static representations of dynamic activity.

An alternative Ca(II) imaging strategy has been the use of blood-oxygen-level-dependent (BOLD) contrast generated in functional magnetic resonance imaging (fMRI).<sup>15-19</sup> Further, while fMRI is not limited by tissue depth, the alterations in BOLD contrast are used to measure brain activity and not necessarily Ca(II) flux and are indirect, as they arise from changes in cerebral blood flow, blood volume, and oxygenation.<sup>17-18</sup> Although optical imaging and BOLD fMRI approaches have yielded insights regarding Ca(II) signaling, transport and trafficking, these techniques are unable to provide high resolution, real-time in vivo imaging of Ca(II) flux.

Contrast-enhanced magnetic resonance imaging (MRI) is an attractive alternative to both optical imaging and BOLD fMRI due to its unlimited penetration depth, soft-tissue contrast, high resolution tomographic (3D) information, and the ability to provide anatomical and physiological information about surrounding tissues.<sup>20</sup> In order to visualize Ca(II) via contrast-enhanced MR imaging we developed bioactivated contrast agents that are conditionally activated by the presence of Ca(II).<sup>21-22</sup> Since these initial studies, several agents have been prepared that detect Ca(II) by a similar mechanism.<sup>23-27</sup> However, these agents are either confined to the extracellular space or cannot discern between intra- and extracellular Ca(II) limiting their effectiveness to report on intracellular Ca(II) flux.

Our work has focused on conjugating two paramagnetic Gd(III) complexes (1,4,7,10tetraazacycloddodecane-1,4,7,10-tetraacetic acid, Gd(III)-DOTA) to the well-known Ca(II) chelator 1,2-bis-(O-aminophenoxy) ethane-N,N,N',N'-tetraacetic acid (BAPTA). Complex **2** [**DOPTA-Gd(III**)] relies on the four carboxylates of the BAPTA domain to block water coordination to the Gd(III) ions (*q*) in the absence of Ca(II) (low relaxivity or 'off' state). In the presence of 1-100  $\mu$ M Ca(II), the four carboxylates of BAPTA preferentially bind Ca(II) allowing water access to the Gd(III) ions and resulting in an 80% increase in *T*<sub>1</sub> relaxivity allowing Ca(II) fluctuations to be visualized via small molecule contrast-enhanced MR imaging. However, while the relaxivity enhancement of complex **2** (**Figure 1**) is adequate to detect Ca(II) binding at low concentrations (~10  $\mu$ M), the overall negative charge of the complex could limit cell permeability. Further, complex **2** is not capable of distinguishing intracellular versus extracellular Ca(II) (1-2 mM) in vivo.

To overcome these limitations we have designed and synthesized complex 1 [DOPTA-Ethyl-Gd(III)] where the negatively-charged imino acetates of BAPTA are protected via

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esterification (**Figure 1**).<sup>28-30</sup> The addition of the esters to the BAPTA ligand address the two issues cited above simultaneously: i. increases the lipophilicity and cell permeability of the agent and more importantly ii. prevents extracellular Ca(II) binding maintaining the contrast agent in a low relaxivity ('off') state prior to cell internalization. Upon cell uptake, endogenous cytosolic esterases cleave the esters of complex 1 deprotecting the aromatic imino acetates (resulting in complex 2) and allowing intracellular Ca(II) binding with a concomitant increase in relaxivity (**Figure 1**).

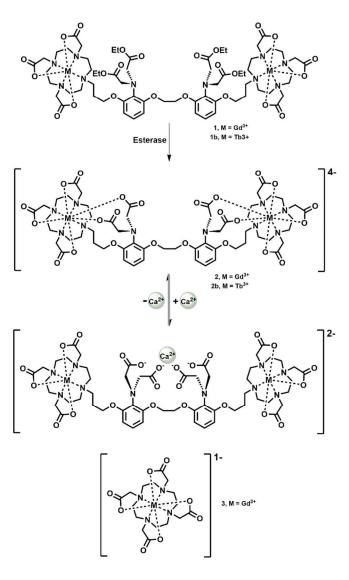


Figure 1. Schematic of complex 1 activation after ethyl ester hydrolysis and subsequent Ca(II) binding. Esterases cleave the ethyl esters on the BAPTA moiety unmasking four carboxylates

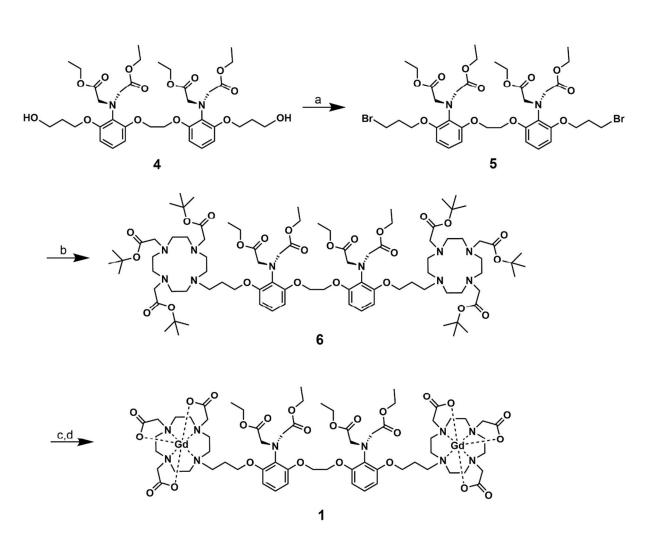
that coordinate to the metal center blocking water access to the macrocyclic chelates maintaining the agent in a low relaxivity state. In the presence of Ca(II) a conformational change occurs whereby the free carboxylates bind Ca(II) providing water access to the Gd(III) centers thereby increasing contrast agent relaxivity. Complex **3** was used in a parallel experiment as a control.

#### **Results and Discussion**

#### *Synthesis*

Complex 1 was synthesized in five steps according to modified literature procedures<sup>31</sup> using 1,2-bis(o-aminophenoxy)-ethane-N,N,N',N'-tetraacetic acid (BAPTA) as a Ca(II)-binding domain due to its  $10^5$ -fold selectivity for Ca(II) over other divalent metal cations such as Mg(II) and protons.<sup>14, 21-22, 32</sup> The intermediate diol, **4** was brominated to the dibromide **5** using bromine and triphenylphosphine. Attachment of the 2,2',2"-(1,4,7,10-tetraazacyclododecane-1,4,7-triyl triacetate (DO3A) was accomplished using excess *tris-t*-butyl-DO3A in order to minimize the formation of the monoalkylated product. The cyclen imino *tris-t*-butyl acetates as part of **6** were selectively hydrolyzed with excess formic acid at 50 °C (the aromatic imino ethyl diacetates on the BAPTA ligand remain). Metalation using GdCl<sub>3</sub> was performed under weakly acidic conditions (pH = 5.5) and after purification by semi-preparative HPLC the final product **1** was isolated (**Figure 2**).

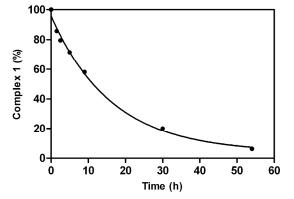
The design of the agent incorporates a propyl linker to couple the aromatic iminoacetates of BAPTA to the macrocycle. We have designed the linker to maximize acetate back-binding (after ester cleavage) to Gd(III) when the Ca(II) concentration is low (< 1  $\mu$ M). As a result, the relaxivity of the complex is minimized. An internal rearrangement allows water to bind to Gd(III) when the Ca(II) concentration is between 5 and 10  $\mu$ M. Utilizing the stable ethyl bromoacetates in place of acetoxymethyl esters reduced synthetic challenges while decreasing the rate of ethyl ester hydrolysis and as a result minimized cleavage by extracellular esterases.



**Figure 2.** Synthetic scheme of DOPTA-Ethyl-Gd(III) . (a)  $Br_2$ , PPh<sub>3</sub>, TEA, DCM, 60%. (b) *tris-t*-butyl-DO3A,  $K_2CO_3$ , MeCN, 65%. (c) formic acid, 55°C. (d) GdCl<sub>3</sub>, NaOH, pH 5-6, 44%.

# Effect of Ester Hydrolysis and Ca(II) Binding on Relaxivity

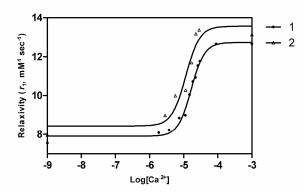
Ethyl ester hydrolysis of **1** was investigated using HPLC-MS equipped with an evaporative light scattering detector (ELSD) to monitor the reaction with porcine esterase.<sup>28, 30, 33-35</sup> The kinetics of ester hydrolysis were determined by following the disappearance of the HPLC peak of **1** post-incubation with porcine esterase at 37 °C in HEPES buffer at pH 7.4 (**Figure 3**, and **Supporting Figure 6**). After 50 hours, 92.4% of **1** was hydrolyzed resulting in a rate constant of  $1.69 \times 10^{-5}$  s<sup>-1</sup> (k<sub>obs</sub>, 37 °C). Employing pseudo first-order kinetics this result corresponds to an 11.4 hour half-life.



**Figure 3.** Plot of ethyl ester hydrolysis of **1** by porcine esterase at 37 °C (in pH 7.4 HEPES buffer) monitored by HPLC-MS using an evaporative light scattering detector (ELSD). Individual data points represent the integration of the HPLC-MS peak of **1** after exposure to porcine esterase divided by the HPLC-MS peak of **1** prior to porcine esterase exposure (multiplied by 100 to give percent complex hydrolysis). The line corresponds to a pseudo first-order kinetic fit of the data with an 11.4 h half-life for **1**.

Relaxivities ( $r_1$ ) of 1, 2, and 1,4,7,10-tetraazacyclododecane-N,N',N'',N'''-tetraacetic acid, 3 [**DOTA-Gd(III**)] were measured. The  $r_1$  of 1, 2, and 3 were 10.6 mM<sup>-1</sup>s<sup>-1</sup>, 7.4 mM<sup>-1</sup>s<sup>-1</sup>, and 3.1 mM<sup>-1</sup>s<sup>-1</sup>, respectively prior to activation and Ca(II) binding (**Table 1**). Following ester hydrolysis (in pH 7.4 HEPES buffer and 37 °C for 24 hours), the  $r_1$  of 1 is 7.6 mM<sup>-1</sup>sec<sup>-1</sup> in the absence of Ca(II) and reaches a maximum of 12.6 mM<sup>-1</sup>s<sup>-1</sup> in the presence of 1 mM Ca(II) with a dynamic relaxivity range between 1 and 100  $\mu$ M Ca(II) (**Table 1, Figure 4**). To confirm that ester hydrolysis of 1 produces the same relaxivity changes as (2), this complex was independently synthesized and the  $r_1$  measured in the presence of increasing Ca(II) concentration. The increase in  $r_1$  of 2 post Ca(II) binding is 64% suggesting the product of the ester hydrolysis of 1 is complex 2 (**Figure 4**). Additionally, these same relaxivity samples were then run on a 7 T imaging spectrometer to determine high field relaxivity. As shown in supporting figure 10 the high field relaxivity of 2 is 6.21 mM<sup>-1</sup>s<sup>-1</sup> and 9.14 mM<sup>-1</sup>s<sup>-1</sup> for 1. This is very promising since the relaxivity decrease due to increased field strength is less than 20% for both molecules.





**Figure 4.** Relaxivity ( $r_1$ , mM<sup>-1</sup>s<sup>-1</sup>) changes of complex **1** (after ethyl ester hydrolysis via porcine esterase) and **2** upon addition of Ca(II) at 1.41 T and 37° C (in pH 7.4 0.1 M HEPES buffer wih 0.1 M KCl and 1% DMSO). Both synthesized **2** and ester hydrolyzed **1** show a > 64% change in  $r_1$  upon Ca(II) binding. Lines represent non-linear sigmoidal dose-response fits to the data. EC50 value for compound **1** is 16.6  $\mu$ M Ca<sup>2+</sup> and for compound **2** is 11.5  $\mu$ M Ca<sup>2+</sup>.

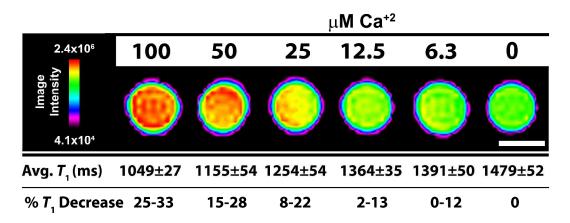
To determine q of **1**, fluorescence lifetime of the Tb(III) analog (**1b**) was measured in D<sub>2</sub>O and H<sub>2</sub>O. Tb(III) was chosen over Eu(III) due to its increased luminescence and longer luminescent lifetime in both H<sub>2</sub>O and D<sub>2</sub>O.<sup>36-38</sup> The difference in luminescence lifetimes in H<sub>2</sub>O versus D<sub>2</sub>O reflects the number of inner-sphere water molecules, q, and can be calculated using Equation (**1**).<sup>38</sup>

$$q = 4.2 \left( \frac{1}{\tau_{H_2O}} - \frac{1}{\tau_{D_2O}} - 0.06 \right) \tag{1}$$

The fluorescence lifetimes of the complexes were measured and fit to an exponential curve with observed decay times of 1.41 ms and 2.42 ms in H<sub>2</sub>O and D<sub>2</sub>O, respectively where q is 0.5 for **1b** and 0.3 for **2b** (per metal). This result indicates that the ethyl esters of **1b** prevent water from coordinating to Gd(III) as efficiently as the acetates of **2b**. Furthermore, q was measured for **1b** and **2b** after the addition of porcine esterase and Ca(II) to determine whether the changes in the relaxivities of the complexes are due to changes in the inner-sphere water coordination of the Gd(III) chelates. As shown in **Table 1**, q for **2b** increases from 0.3 to 1 in the presence of 1 mM

Ca(II) whereas q for 1c (prior to ester hydrolysis) remains constant at 0.5 even in the presence of 1 mM Ca(II). This data shows the ethyl ester protected iminoacetates likely do not participate in Ca(II) binding prior to enzymatic hydrolysis. Following hydrolysis, the q value of 1 increases from 0.5 to 0.9 confirming that the ester hydrolysis of 1 results in 2.

To investigate whether changes in low field relaxivity can be visualized via high field MRI (7 T, 300 MHz) phantoms of a 100  $\mu$ M solution of **2** [200  $\mu$ M Gd(III)] were prepared with different stoichiometric concentrations of CaCl<sub>2</sub> with concentrations of 100, 50, 25, 12.5, and 6.25, and 0  $\mu$ M Ca(II) (**Figure 5**). There is a 32% decrease in *T*<sub>1</sub> from 0  $\mu$ M Ca(II) to 100  $\mu$ M Ca(II) which is visible via high field MRI. Additionally, Ca(II) concentrations as low as 6.25  $\mu$ M produce a *T*<sub>1</sub> decrease of at least 10% which is required to visualize large Ca(II) fluxes of approximately 10  $\mu$ M.



**Figure 5.** MR images and  $T_1$  analysis of 100  $\mu$ M solution phantoms of **2** binding with increasing amounts of Ca(II) corresponding to 100, 50, 25, 12.5, and 6.25  $\mu$ M Ca(II) at 7T and 25 °C. Scale bar (white) represents 1 mm.  $T_1$  values are calculated as the average of 5, 1 mm slices  $\pm$  1 standard deviation.  $\%T_1$  decrease is calculated versus the 0  $\mu$ M Ca(II) sample.

Table 1 - Relaxivity, octanol-water partition coefficients and hydration numbers of complexes 1, 2, and 3

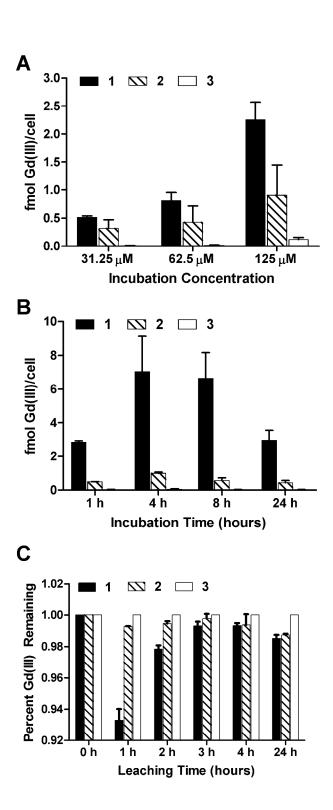
	- Ca(II)		+ 1 mM Ca(II)		+ Esterase + 1 mM Ca(II)	
log P	$r_1$ (mM <sup>-1</sup> s <sup>-1</sup> )	q	$r_1$ (mM <sup>-1</sup> s <sup>-1</sup> )	q	$r_1$ (mM <sup>-1</sup> s <sup>-1</sup> )	q

1	-2.44	$10.6 \pm 0.3$	$0.5 \pm 0.1$	$10.2 \pm 0.2$	$0.5 \pm 0.1$	$12.6\pm0.3$	$0.9 \pm 0.1$
2	-3.69	$7.4 \pm 0.2$	$0.3 \pm 0.1$	$12.5 \pm 0.4$	$1.0 \pm 0.1$	-	-
3	-3.42	$3.1 \pm 0.3$	$1 \pm 0.1$	$3.1 \pm 0.2$	-	-	-

#### In vitro Cellular Uptake and Retention

The previous data showed that Ca(II) can only bind to complex **1** following ester hydrolysis resulting in its detectable form (complex **2**) which can then bind Ca(II) (5-10  $\mu$ M) leading to a decrease in *T*<sub>1</sub> providing positive MR image contrast. This provides a novel bio-activated probe for determining intracellular Ca(II) fluxes but requires cellular uptake of the contrast agent. One factor that has been used to predict the ability of a molecule or drug to diffuse into biomembranes is lipophilicity where a balance in amphilicity is desired for ease of aqueous administration and cell uptake.<sup>39-40</sup> In order to determine the lipophilicity of Ca(II)-sensitive MRI contrast agents, octanol–water partition coefficients (log *P*) were determined for each complex where **1** was one order of magnitude more hydrophobic than **2** (-2.44 versus -3.69) which should lead to an increase in cell uptake.<sup>41-42</sup>

To determine the cell labelling properties of **1**, **2**, and **3** a hippocampal neuronal cell line (HT-22 cells were chosen as they have frequently been used to investigate neuronal activity).<sup>43-44</sup> Concentration-dependent uptake of **1**, **2** and **3** was determined by incubating HT-22 cells with 31.25, 62.5, and 125  $\mu$ M of the complexes for 48 hours. After incubation, cells were washed, trypsinized, counted, and digested with nitric acid. Gd(III) content was measured using ICP-MS of the acid-digested samples (**Figure 6A, Supporting Table 1**). As shown, cellular labelling is concentration dependent and directly correlated to the octanol–water partition coefficients of the complexes where the more hydrophobic **1** exhibited higher uptake than **2** at incubation concentrations of 125  $\mu$ M. At lower incubation concentrations, uptake of **1** and **2** were the same.



**Figure 6**. Concentration-dependent (A), time-dependent (B), and cellular retention studies (C) of **1**, **2**, and **3** in the hippocampal neuronal cell line, HT-22. Cellular uptake of **1** was significantly higher than that of **2** and **3**. A small amount of contrast agent (<7%) leached out of the cells in 24

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hours. All samples were washed 3x with DPBS, trpsyinized, counted, and digested in nitric acid to determine Gd(III) content via ICP-MS. Data are mean  $\pm$  standard deviation of triplicate runs

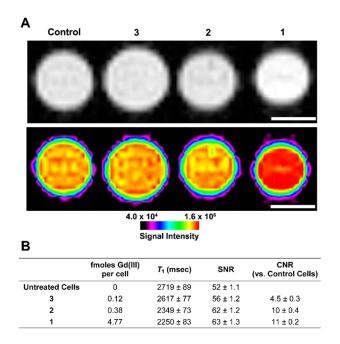
To determine the time-dependent cell labelling, HT-22 cells were incubated with 125  $\mu$ M of the complexes for 1, 4, 8, 24 hours. After incubation, cells were washed, trypsinized, counted, and digested with nitric acid. Finally, Gd(III) content was measured using ICP-MS of the acid-digested samples. As shown in **Figure 6B** (and **Supporting Table 2**) maximum Gd(III) labelling occurs after a 4 or 8 hour incubation followed by a significant decrease after 24 hours.

To elucidate the relationship between cellular labelling and cellular retention HT-22 cells were incubated with 125  $\mu$ M of each complex for 2 hours. After incubation, cells were washed with DPBS (2x) followed by the addition of fresh media. Cells were allowed to efflux (leach) for various time points (0, 1, 2, 3, 4, and 24 hrs) in fresh media followed by media removal, cell washing with DPBS (2x), cell harvesting and Gd(III) determination via ICP-MS. At each time point cell samples were counted and analyzed by ICP-MS.

Figure 6C shows that, despite multiple washes, greater than 90% of the Gd(III) is retained. The cells treated with 1 lost ~ 5% more of the agent compared to cells treated with the other two agents. However, because the cellular uptake of 1 is more than 200% higher than complex 2 (and more than 500% higher than complex 3), the overall amount of cell associated 1 remains higher than 2 and 3 prior to and after leaching. The absolute quantities were divided by their respective initial concentrations (those collected at time = 0) in order to quantify the relative amount of cellular retention for each agent.

To determine whether the cell labelling and molecular relaxivity of 1 and 2 leads to detectable image contrast, HT-22 cells were treated with 125  $\mu$ M of 1, 2, and 3 complexes for 4 hours and centrifuged into capillaries (~ 1.0 mm diameter) prior to MR imaging. The greyscale

and color intensity MR images of the cell pellets show increases in image intensity for cells treated with 1 or 2 compared with cells treated with 3 or untreated cells where the contrast to noise ratio (CNR) between 1 and 3 is 11 (Figure 7A). Lastly, The amount of Gd(III) per cell was determined by ICP-MS for each cell pellet and as expected,  $T_1$  was found to be inversely proportional to Gd(III) content (Figure 7B).



**Figure 7.** (A)  $T_1$ -weighted MR image with corresponding image-intensity color map of HT-22 cells incubated with **1**, **2**, **3** and media alone for 4 h (scale bar = 1 mm). 9.4 T (400 MHz) and 25°C. (B) Calculated values corresponding to the MR image in (A). Values represent the average of 4, 1 mm slices and the error is plus or minus one standard deviation of the mean.

The findings from these *in vitro* studies show that **1** labels cells up to 2-fold more efficiently than **2** or **3**, which is directly correlated to the octanol-water partition coefficients. The results are in agreement with our hypothesis that esterification enhances lipophilicity and consequently cell labelling.

# Conclusion

We have synthesized a cell-permeable version of a previously reported Ca(II)-sensitive MRI contrast agent, **1**, for visualizing intracellular Ca(II) signaling. By incorporating ethyl esters in the BAPTA binding ligand the charge of the complex is neutralized resulting in increased lipophilicity and concurrent cellular labelling. In addition, the ethyl esters maintain the contrast agent in a low relaxivity or 'off' state until ethyl ester hydrolysis resulting in the active form, **2**, which is capable of binding Ca(II) producing a high relaxivity or 'on' state. This probe design will allow for the discrimination between extra- and intracellular Ca(II) by MR imaging and thereby providing the ability to monitor intracellular Ca(II) signaling.

#### **Materials and Methods**

#### **General Methods**

Unless otherwise noted, materials and solvents were purchased from Sigma-Aldrich Chemical Co. (St. Louis, MO) and used without further purification. Common precursor **4** (2-{2-[3-(3-hydroxypropoxy)-2-di(ethoxycarbonylmethyl)aminophenoxy]ethoxy} -6-(3hydroxypropoxy)-N,N-di(ethoxycarbonylmethyl)benzenamine), *tris-t*-butyl-DO3A (*Tris-tert*butyl 2,2',2"-(1,4,7,10-tetraazacyclododecane-1,4,7-triyl) triacetate) were prepared according to literature procedures.<sup>3e, 20</sup> Unless noted, all reactions were performed under a nitrogen or argon atmosphere. Acetonitrile was purified using a Glass Contour Solvent system (Pure Process Technology, Nashua, NH, USA). Deionized water (18.2 M $\Omega$ ·cm) was obtained from a Millipore Q-Guard System (Billerica, MA). Thin-layer chromatography (TLC) was performed on EMD 60 F254 silica gel plates. Standard grade 60 Å 230 – 400 mesh silica gel (Sorbent Technologies, Norcross, GA, USA) was used for flash column chromatography. <sup>1</sup>H and <sup>13</sup>C NMR spectra were obtained on a Bruker 500 MHz Avance III NMR spectrometer (Bruker Biospin, Billerica, MA, USA) or a Varian Inova 400 MHz NMR spectrometer (Agilent Technologies, Santa Clara, CA, USA) with deuterated solvents as noted. Electrospray ionization mass spectrometry (ESI-MS) was carried out using a Varian 1200L single-quadrupole mass spectrometer (Agilent Technologies, Santa Clara, CA, USA). Matrix-assisted laser desportion ionization time-of-flight mass spectra (MALDI-TOF-MS) were recorded on a Bruker AutFlex III (Bruker, Billerica, MA, USA), using 2, 5-dihydroxybenzoic acid as the matrix. Analytical reverse-phase HPLC-MS was performed on a Varian Prostar 500 system (Agilent Technologies, Santa Clara, CA, USA) using either a Waters (Milford, MA, USA) Atlantis C18 column ( $4.6 \times 250, 5 \mu m$ ) or an XBridge C18 column (4.6  $\times$  150, 5 µm). This system is equipped with a Varian 380 LC ELSD system, a Varian 363 fluorescence detector, a Varian 335 UVvis detector, and a Varian 1200L quadrupole MS detector (Agilent Technologies, Santa Clara, CA, USA). Preparative runs were performed on a Waters (Milford, MA, USA) Atlantis C18 column ( $19 \times 250$ ,  $10 \mu m$ ). The mobile phase consisted of water or water with  $NH_3 \cdot H_2O$  (pH = 10.4) (solvent A) and HPLC-grade acetonitrile (solvent B).

# 2-{2-[3-(3-bromopropoxy)-2-di(ethoxycarbonylmethyl)aminophenoxy]ethoxy}-6-(3-

**bromopropoxy**)-*N*,*N*-**di**(ethoxycarbonylmethyl)benzenamine, 5. 0.37 mL of TEA (2.66 mmol) was added via syringe to 0.70 g of triphenylphosphine (2.66 mmol) dissolved in 10 mL of dry dichloromethane at 0 °C. 0.14 mL of bromine (2.66 mmol), diluted in 15 mL of dry dichloromethane, was added to the reaction mixture. Stirring was continued at 0 °C for 30 min. 0.89 g of compound 4 (1.21 mmol), in 10 mL of dichloromethane, was added via syringe and the reaction mixture was stirred at room temperature. Absence of starting material was confirmed by TLC after 24 h. The crude material was purified by flash column chromatography over silica gel

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with hexanes: ethyl acetate (3:1) to afford a yellow oil (0.63g, 60%,  $R_f = 0.45$ ). <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>):  $\delta$  6.93 (t, 2H, J = 8.3 Hz), 6.60 (d, 2H, J = 8.2 Hz), 6.56 (d, 2H, J = 8.2 Hz), 4.30 (s, 4H), 4.09 (t, 4H, J = 5.8 Hz), 4.08 (s, 8H), 4.05 (q, 8H, J = 7.15 Hz), 3.63 (t, 4H, J = 6.5 Hz), 2.30 (m, 4H), 1.16 (t, 12H, J = 7.15 Hz); <sup>13</sup>C-{H} NMR (126 MHz, CDCl<sub>3</sub>):  $\delta$  172.2, 154.2, 128.3, 123.6, 106.9, 106.7, 67.5, 66.5, 60.2, 53.9, 32.5, 30.4, 14.2

### 2-{2-[3-(3-(tris-t-butyl-DO3A)propoxy)-2-di(ethoxycarbonylmethyl)aminophenoxy]

# ethoxy}-6-(3-(tris-t-butyl-DO3A)propoxy)-N,N-di(ethoxycarbonylmethyl) benzenamine,

**DOPTA-Ethyl, 6**. 131 mg (0.26 mmol) of tris-t-butyl-DO3A in 50 mL of acetonitrile and 49.7 mg (0.36 mmol) of K<sub>2</sub>CO<sub>3</sub> was added to a 100 mL round-bottom flask. The mixture was sealed, stirred and placed under nitrogen atmosphere for 30 min before 100 mg (0.12 mmol) of **5** in 10 mL of acetonitrile was added. The mixture was heated to 75 °C and stirred for 48 h. The reaction was monitored by mass spectrometry to follow the disappearance of the starting materials. The reaction was filtered and evaporated to yield a yellow oil. The crude material was purified by flash column chromatography over silica gel with dichloromethane: methanol (50:1) to afford a yellow oil (135 mg, 65% R<sub>f</sub> = 0.90). <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>, 70 °C):  $\delta$  6.92 (2H, br), 6.64 (4H, br), 4.28 (s, 4H), 3.97 (m, 20H,), 3.67-1.79 (m, 52H), 1.43 (s, 54H), 1.11 (t, 12H, *J* = 6.92 Hz); MALDI-MS *m/z* observed: 1730.33, calculated: 1730.17 [M + H]<sup>+</sup>.

# 2-{2-[3-(3-(4,7,10-tris(carboxymethyl)-1,4,7,10-tetraazacyclododecyl)propoxy)-2-

di(ethoxycarbonylmethyl)aminophenoxy]ethoxy}-6-(3-(4,7,10-tris(carboxymethyl)-1,4,7,10-tetraazacyclododecyl)propoxy)-N,N-di(ethoxycarbonylmethyl)benzenamine-gadolinium(III), DOPTA-Ethyl-Gd(III), 1. 200 mg (0.12 mmol) of compound 6 wasdissolved in 50 mL of formic acid and heated to 55 °C for 18 hours. After formic acid wasevaporated on a rotary evaporator, the resulting oil was redissolved in methanol 3 × 10 mL and

evaporated again to remove most of the formic acid. The resulting glassy solid 7 was dissolved in 20 mL of water and the pH of the solution was adjusted to 5. 107 mg (0.28 mmol) of GdCl<sub>3</sub>·  $6H_2O$  was added and the pH of the reaction was adjusted back to 6 every 4-6 h until no further change occurred. (Typically the reaction was completed in 1-2 days). The reaction was evaporated and purified by reverse phase HPLC according to method 1, retention time of 44.5 min, and 99% purity, followed by lyophilization to yield 87 mg of **1** as a white powder in 44%. MALDI-TOF-MS m/z observed: 1703.2 (M + H<sup>+</sup>) calcd for C<sub>64</sub>H<sub>94</sub>Gd<sub>2</sub>N<sub>10</sub>O<sub>24</sub> 1702.0.

The terbium(III) analog of **1** were prepared in a similar manner for luminescence lifetime measurements. HPLC and MALDI-TOF-MS were used to determine the purity. DOPTA-Ethyl-Tb(III), retention time: 47.0 min, yield: 42%, m/z observed: 1705.1, calcd for  $C_{64}H_{94}Yb_2N_{10}O_{24}$  1705.3

**Relaxation Time Measurements. 1** was dissolved in 500  $\mu$ L of filtered, 0.1 M HEPES, 0.1 M KCl containing 1% DMSO for  $T_1$  acquisition. Relaxation times were measured on a Bruker mq60 NMR analyzer equipped with Minispec v2.51 Rev.00/NT software (Bruker Biospin, Billerica, MA, USA) operating at 1.41 T (60 MHz) and 37 °C  $T_1$  relaxation times were measured using an inversion recovery pulse sequence ( $T_1$ \_ir\_mb) using the following parameters: 4 scans per point, 10 data points for fitting, mono-exponential curve fitting, phase cycling, 10 ms first pulse separation, and a recycle delay and final pulse separation  $\geq$  5  $T_1$ . The Gd(III) concentrations of each solution were determined using ICP-MS. The relaxivity  $r_1$  was calculated from Equation (2). The inverse of the longitudinal relaxation time ( $1/T_1$ , s<sup>-1</sup>) was plotted against Gd(III) concentration (mM) and fitted to a straight line with an R<sup>2</sup> > 0.99. The slope of the fitted line was recorded as the relaxivity,  $r_1$ .

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$$\frac{1}{T_{1,obs}} = \frac{1}{T_{1,d}} + r_1[Gd]$$
(2)

**Octanol–Water Partition Coefficients.** 0.5 mg of **1**, **2** and **3** were dissolved in 1 mL of a 1:1 mixture of water:1-octanol. After shaking for 30 s, tubes containing the mixtures were placed on a rotator for gentle mixing to equilibrate for 48 h. After mixing, samples were allowed to settle at room temperature for 24 hours. Gd(III) concentrations of each layer were determined by ICP-MS. Partition coefficients were calculated from the Equation log  $P = log(C_o/C_w)$ , where log P is the logarithm of the partition coefficient,  $C_o$  is the concentration of Gd in the 1-octanol layer, and  $C_w$  is the concentration of Gd in the water layer.

Luminescence Lifetime Measurements of 1C. The luminescence lifetime of a 1 mM solution of 1c was measured in H<sub>2</sub>O and D<sub>2</sub>O on a Hitachi (San Francisco, CA) F4500 fluorometer, using a  $\lambda_{ex} = 232$  nm and  $\lambda_{em} = 544$  nm. Twenty-five scans were acquired, averaged and fit to a mono-exponential decay function.

Kinetics of Ethyl Ester Hydrolysis. Porcine liver esterase was diluted 10 times with 0.1 M HEPES, 0.1 M KCl and 1% DMSO buffer and used to hydrolyze 1. The solution of 1 was prepared by dissolving the agent in in the aforementioned buffer to give a final concentration of 1 mM. 10 U/ml of porcine esterase was added to the solution and was maintained at 37 °C. 200  $\mu$ L aliquots of the reaction was taken at different time points for 72 h and centrifuged at 4000 RPM for 5 mins and filtered through microporous (0.2  $\mu$ m) membrane filter. The filtrate was analyzed by HPLC-MS with an evaporative light scattering detector (ELSD). Pseudo-first-order rate constants for enzymatic hydrolysis were determined based on the Equation,  $y = A_0 \times e^{-x/t}$  (where  $A_0 = 1$  and  $t^{-1} = k_{obs}$ ), using Origin 7 (OriginLab, Northampton, MA).

General Cell Culture. Dulbecco's modified phosphate buffered saline (DPBS), media, and TrypLE were purchased from Invitrogen (Carlsbad, CA). Serum is from VWR (Corning, Mediatech). Cell culture consumables (flasks, plates, etc.) were purchased from VWR (Radnor, PA). Charcoal dextran stripped FBS was purchased from Atlanta Biologicals (Lawrenceville, GA). The HT-22 hippocampal neuronal cell line obtained from Professor Schubert at Salt Institute (La Jolla, CA) and were cultured using phenol red free DMEM supplemented with 10% FBS. Experiments and incubations were done in a 5.0% CO<sub>2</sub> incubator at 37 °C unless otherwise noted. Cells were harvested by incubation with 0.25% TrypLE for 10 min at 37 °C in a 5.0% CO<sub>2</sub> incubator.

**Cellular Uptake Studies**. *Concentration-dependent uptake*. Cell studies were performed to determine the ability of **1**, **2** and **3** to label cells. HT-22 hippocampal neuronal cells were grown to 60% confluence in a 24 well cell culture plate at 350  $\mu$ L per well. Media was then replaced with media containing 1% DMSO and increasing concentrations (31.25  $\mu$ M, 62.5  $\mu$ M and 125  $\mu$ M) of the three agents in duplicate. The plate was incubated for 48 h and washed three times with sterile phosphate buffered saline. Each sample was then trypsinized and tested for cell count and viability on a Guava PCA system using the Guava viacount protocol according to the manufacturer's instructions. The remaining volume in each sample was dissolved in concentrated nitric acid and diluted for analysis by ICP-MS to determine the amount of Gd(III) associated with the cells. Viability analysis revealed no change in viability of treated cells.

*Time-dependent uptake.* The cells were prepared in a similar fashion to previous assays. HT-22 hippocampal neuronal cells were grown to 60% confluence and incubated with 125  $\mu$ M of the three agents for 1, 4, 8, and 24 h in duplicate. After incubation, the medium was removed, and the cells were rinsed three times with DPBS. These cells were trypsinized, counted, and analyzed for Gd(III) content using Guava EasyCyte Mini PCA System and ICP-MS.

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*Cellular Retention Studies.* The cells were prepared in a similar fashion to previous assays. HT-22 hippocampal neuronal cells were grown to 60% confluence and incubated with 125  $\mu$ M of the three agents for 2 h. After incubation, the cells were washed with DPBS (time = 0, initial amount), placed in fresh media, and allowed leach for 0, 1, 2, 3, 4, and 24 hours. After each time point, cells were harvested, counted, and analysed for Gd(III) content by ICP-MS.

Cell Counting Using A Guava EasyCyte Mini Personal Cell Analyzer (PCA) System. After cell harvesting, an aliquot of the cell suspensions was mixed with Guava ViaCount reagent and allowed to stain at room temperature for at least 5 min (no longer than 20 min). Stained samples were vortexed for 10 s and percent cell viability was determined via manual analysis using a Guava EasyCyte Mini Personal Cell Analyzer (PCA) and ViaCount software module. For each sample, 1000 events were acquired with dilution factors that were determined based upon optimum machine performance (~25-100 cells/ $\mu$ L). Instrument reproducibility was assessed daily using GuavaCheck Beads and following the manufacturer's suggested protocol using the Daily Check software module.

**MR Imaging and**  $T_1$  **analysis of Cell Pellets at 9.4 T.** 9.4 T MR imaging and  $T_1$  measurements were performed on a Bruker Biospec 9.4 T imaging spectrometer fitted with shielded gradient coils at 25 °C. For cell pellet images, ~ 2.0 × 10<sup>6</sup> HT-22 cells were incubated in 25-cm<sup>2</sup> T-flasks with 125  $\mu$ M of **1**, **2** and **3** for 4 h. The cells were rinsed with DPBS (3 × 5 mL/flask), and harvested with 500  $\mu$ L of a 0.25-vol% trypLE Express solution. After addition of 500  $\mu$ L of the appropriate fresh media (1.0 mL total volume/flask), cells were transferred to 1.5-mL microcentrifuge tubes and centrifuged at 200 g for 5 min. The supernatant was removed, and cell pellets were re-suspended in 1 mL of complete media, added to 5 3/4″ flame-sealed Pasteur pipets, and centrifuged at 200 g and 4.0 °C for 5 min. The bottom sections of the flame-sealed pipets were scored with a glass scribe, broken into small capillaries, and imaged using a RF RES 400 1H 089/038 quadrature transmit receive 23-mm volume coil (Bruker BioSpin, Billerica, MA, USA).

Spin-lattice relaxation times ( $T_1$ ) were measured using a rapid-acquisition rapid-echo (RARE-VTR) T1-map pulse sequence, with static TE (10 ms) and variable TR (100, 200, 400, 500, 750, 1000, 2500, 5000, 7500, and 10000 ms) values. Imaging parameters were as follows: field of view (FOV) =  $25 \times 25$  mm<sup>2</sup>, matrix size (MTX) =  $256 \times 256$ , number of axial slices = 3, slice thickness (SI) = 1.0 mm, and averages (NEX) = 4 (total scan time = 2 hrs 58 min).  $T_1$  analysis was carried out using the image sequence analysis tool in Paravision 5.0 pl3 software (Bruker, Billerica, MA, USA) with mono-exponential curve-fitting of image intensities of selected regions of interest (ROIs) for each axial slice.

Quantification of Gd(III) with Inductively Coupled Plasma Mass Spectrometry. Quantification of Gd(III) was accomplished using ICP-MS of acid digested samples. Specifically, solution samples were digested in ACS reagent grade nitric acid (70%, Sigma, St. Louis, MO, USA) and incubated in a water bath at 70 °C for at least 2 h to allow for complete sample digestion. A portion of the digested sample was added to a 15 mL conical tube along with 5 ng/mL of multi-element internal standard containing Bi, Ho, In, Li, Sc, Tb, and Y (Inorganic Ventures, Christiansburg, VA, USA) in filtered, de-ionized H<sub>2</sub>O (18.2 M $\Omega$ ·cm). Instrument calibration was accomplished by preparing individual-element Gd standard (Inorganic Ventures, Christiansburg, VA, USA) using concentrations of 1.000, 5.000, 10.00, 20.00, 50.00, 100.0, and 200.0 ng/mL containing 3.0% nitric acid (v/v) and 5.0 ng/mL of multi-element internal standard.

ICP-MS was performed on a computer-controlled (Plasmalab software) Thermo X series II ICP-MS (Thermo Fisher Scientific, Waltham, MA, USA) operating in standard mode

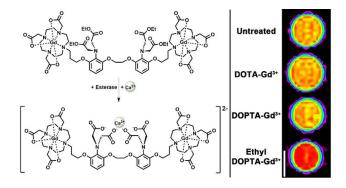
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equipped with an ESI SC-2 autosampler (Omaha, NE, USA). Each sample was acquired using 1 survey run (10 sweeps) and 3 main (peak jumping) runs (100 sweeps). The isotopes selected for analysis were <sup>157,158</sup>Gd with <sup>115</sup>In and <sup>165</sup>Ho isotopes selected as internal standards for data interpolation. Instrument performance is optimized daily by an autotune followed by verification via a performance report (passing manufacturer specifications). Addition of all reagents for all samples and standards were weighed using a Mettler Toledo (Columbus, OH, USA) XS205 DeltaRange analytical micro balance (with 0.01 mg resolution for up to 81 g of sample).

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#### For Table of Contents Only



# TOC Figure Synopsis:

Despite the importance of Ca(II) in physiology, visualizing Ca(II) flux in the central nervous system (CNS) in vivo remains challenging. To address these challenges we have developed a new generation, cell-permeable Ca(II)-activated MRI contrast probe that utilizes ethyl esters to increase cell permeability and prevent extracellular divalent Ca(II) binding. Upon cellular internalization, the ethyl esters are cleaved allowing the agent to bind intracellular Ca(II) increasing relaxivity and resulting in enhanced positive MR image contrast.

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