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Synthesis and evaluation of MR probes for targeted-reporter imaging†

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Visualizing disease heterogeneity remains a challenging task since most imaging agents are targeted to a single receptor. We describe the development of an MR platform able to report on multiple molecular events. Enzyme activation and enhanced cellular uptake of this modular probe make it suitable for subsequent targeted-reporter imaging applications.

Extracellular receptors enable cells to interact with and adapt to their surrounding environment through a series of interwoven signalling cascades.¹ Receptor expression levels therefore provide valuable insight with regard to cellular health. For instance, given the increased nutrient requirements necessary for uncontrolled cellular proliferation, cancerous cells are often characterized by the concerted overexpression of multiple cell surface receptors.² Specifically, overexpression and amplification of epidermal growth factor receptor (EGFR) has been reported as the primary gain-of-function mutation observed in brain,^{3,4} breast,⁵ lung,⁶ ovarian,⁷ and skin cancers.⁸ Moreover, transferrin receptor (TfR) has been implicated in several cancers given the involvement of the iron-dependent enzyme ribonucleotide reductase during rapid DNA synthesis.^{9,10} Thus, targeted molecular imaging strategies able to report on the status of multiple biomarkers have the potential to function as innovative diagnostic agents during drug development.

The Basilion lab has demonstrated the ability to visualize multiple biomarkers using complementing β -galactosidase (β -gal) fragments targeted to EGFR and TfR.¹¹ This approach relied on colocalization of complementing enzyme fragments at extracellular surfaces overexpressing the target receptors to restore enzyme activity, thus permitting their detection following colorimetric staining with X-gal *in vitro*. While this approach represents a first attempt at multimarker visualization, the colorimetric output is restricted to translucent specimens, thereby precluding longitudinal studies *in vivo*. More recently, Grimm and coworkers achieved *in vivo* imaging of multiple

biomarkers by utilizing Cerenkov luminescence produced by targeted positron-emitting radionuclides to excite nearby bio-activatable fluorescent nanoparticles.¹² However, the use of radionuclides make this imaging platform unsuitable for longitudinal studies given prolonged exposure to ionizing radiation. To overcome these limitations, we designed a complementary approach employing magnetic resonance (MR) imaging.

MR imaging has greatly impacted diagnostic medicine due to its high spatiotemporal resolution and unlimited penetration depth.^{13,14} To improve image contrast, paramagnetic agents are frequently administered to shorten the longitudinal (T_1) or transverse (T_2) relaxation times of nearby water protons. This increase in the observed water relaxation rate ($1/T_{i,obs}$, $i = 1, 2$) yields a change in image contrast commensurate with the rate in the presence ($1/T_{i,p}$) and absence ($1/T_{i,d}$) of the paramagnetic species. With seven unpaired electrons and high magnetic moment, Gd(III) chelates are the most commonly employed T_1 agents. The efficiency with which these agents shorten T_1 is termed relaxivity (r_1), and is influenced by the number of bound waters (q), their residency lifetime (τ_m), and the rotational correlation time of the complex (τ_R). Modulating any one of the above variables upon interaction with a stimulus generates responsive MR probes.^{15–23}

Here, we present a modular platform for the noninvasive visualization of enzyme activity and receptor expression. To achieve this aim we designed a Gd(III) chelate that incorporates a proximal glycoside and a distal biotin that allow signal transduction and receptor targeting, respectively. The corresponding galactose residue is covalently tethered to the Gd(III) core through a self-immolative linker capable of modulating the coordination chemistry about the metal center in a manner similar to that previously reported by our group.²⁴ Biotinylation of the probe enables facile complexation with avidin, a tetrameric protein capable of binding four biotin molecules ($K_d = 10^{-15}$ M).²⁵ Complexation of three MR agents to one avidin was predicted to enable facile targeting to the desired receptor through subsequent complexation between the remaining

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subunit and the desired biotinylated ligand (*e.g.*, Tf). β -gal could then similarly be targeted to a separate receptor with the appropriate biotinylated ligand (*e.g.*, EGF). Overexpression of both receptors is therefore expected to result in colocalization, receptor-mediated internalization, and subsequent activation through targeted-reporter complex formation (Fig. 1).

Briefly, 1,4,7,10-tetraazacyclododecane was protected at the 1 and 7 positions with Cbz-Cl²⁶ prior to alkylation with ethyl bromoacetate; subsequent deprotection of the Cbz groups *via* catalytic hydrogenation yielded *N-trans* symmetrical material (Scheme S1†). Alkylation of this material with half an equivalent of 1-(*tert*-butyl)6-ethyl 2-bromohexanedioate²⁷ (**3**) followed by acid-catalyzed cleavage of the *t*-butyl ester gave trisubstituted macrocycle in 34% over five steps. Reaction with **11**, followed by global deprotection and subsequent metalation yielded untargeted complex **Gd14** in 5% after preparative HPLC. Peptide coupling with norbiotinamine (**15**)²⁸ and **16**²⁹ gave the desired biotinylated MR agents **Gd2** and **Gd1**, respectively (Table 1 and Scheme S3†). The corresponding Tb(III) complexes were similarly generated to quantify *q* following luminescence lifetime decay measurements.

Gd1 and **Gd2** were assessed regarding their ability to influence the $1/T_1$ of nearby water protons at pH 7.4 in 100 mM MOPS buffer. T_1 measurements made at 1.5 T and 37 °C yielded r_1 values for **Gd1** and **Gd2** of 7.8 and 8.6 mM⁻¹ s⁻¹, respectively. Exposing each agent to β -gal afforded a measurable change in absorbance and $1/T_1$ (Fig. S19†). The apparent decrease in the latter is consistent with findings previously reported by our lab,²⁴ and is likely due to the decrease in molecular weight upon

enzyme activation.³⁰ Both agents exhibited complete activation within three hours.

We next examined the ability of **Gd1** and **Gd2** to form host-guest complexes with avidin. This was confirmed by changes in $1/T_1$ and $1/T_2$ of nearby water protons upon successive additions of avidin. Measurements were performed identically (*vide supra*), and converted to r_1 and r_2 values following ICP-MS. As anticipated, **Gd2** exhibited an increase in both r_1 (149%) and r_2 (382%) upon complexation with avidin (Fig. S20†). This finding is expected given an increase in τ_R following complexation with biomacromolecules. Moreover, r_1 and r_2 values appeared to saturate upon binding of approximately three **Gd2** molecules to a single avidin tetramer, as has been seen before.³¹ While **Gd1** produced a similar increase in r_2 (40%) that became saturated at the desired 4 : 1 stoichiometry, r_1 remained largely unchanged throughout the experiment (Fig. 2). This response is important for subsequent targeted-reporter imaging and was later attributed to a concurrent decrease in *q* (*vide infra*). We therefore aimed to further establish **Gd1** as a viable candidate for such imaging applications by demonstrating its ability to drastically alter relaxation rates of nearby water protons upon activation while complexed with avidin.

Attempts to activate **Gd2** complexed to avidin failed to alter r_1 , suggesting the proximity of avidin to the glycoside was sufficient to obviate β -gal recognition, and hence MR activation. However, **Gd1** readily underwent enzyme activation while complexed with avidin owing to the increased linker length. However, unlike before, activation of this macromolecular complex by β -gal afforded a desirable increase in r_1 from 7.8 to 17.2 mM⁻¹ s⁻¹.

To understand these changes in r_1 prior to, and following complexation with avidin and/or β -gal activation, luminescence lifetime measurements were performed with **Tb1** and **Tb2** (Table 1). As noted above, the constant r_1 value for **Gd1** may be rationalized given a concurrent decrease in *q* and increase in τ_R , the latter of which dominates at clinical field strengths.¹³ These findings indicate that avidin, like carbonate,^{24,27,32} is capable of efficiently masking the open coordination sites in **Gd1** to afford an MR agent exhibiting a desirable increase in r_1 (130%) following activation by β -gal.

We determined the binding stoichiometry between this agent and avidin *via* a colorimetric assay using 2-(4-hydroxyphenylazo)benzoic acid (HABA). To a reconstituted HABA/avidin solution was titrated in a solution of **Gd1** and the change in HABA absorption at 500 nm monitored. Plotting the difference in absorption between each addition against the corresponding ratio of **Gd1** to avidin indicated the number of probes bound to each avidin (Fig. S21†). Upon the addition of four equivalents of **Gd1**, no observable change in the absorption of HABA was noted. These results confirm complete and efficient displacement of HABA from avidin in the presence of four equivalents of **Gd1**, suggesting negligible influence of the macrocyclic chelate on binding of the distal biotin to avidin.

Having established the desired 4 : 1 binding stoichiometry, we investigated the ability to target **Gd1** to cells overexpressing TfR. The corresponding **Gd1** : avidin : Tf (3 : 1 : 1) macromolecular construct was generated and cellular uptake

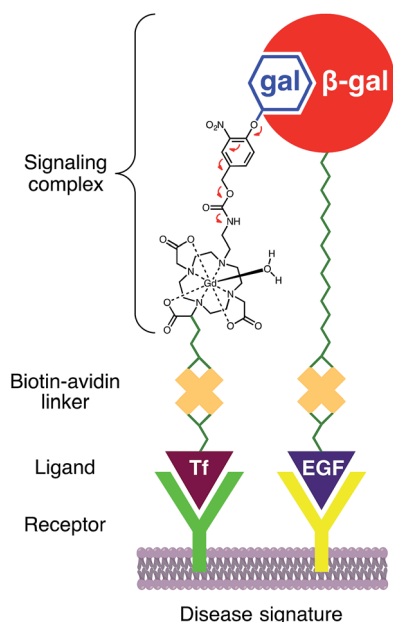
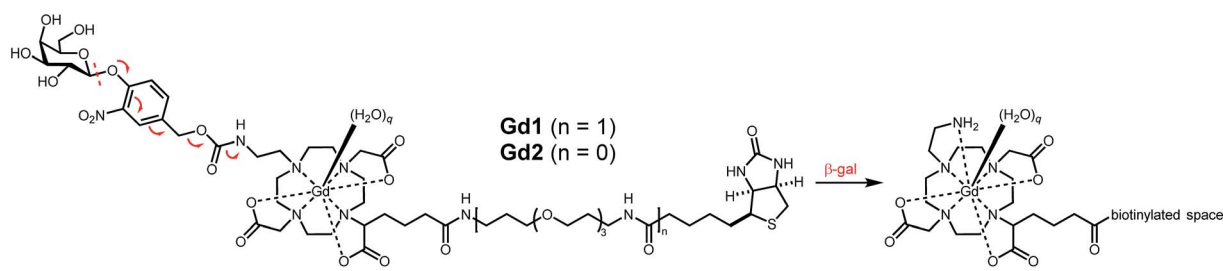


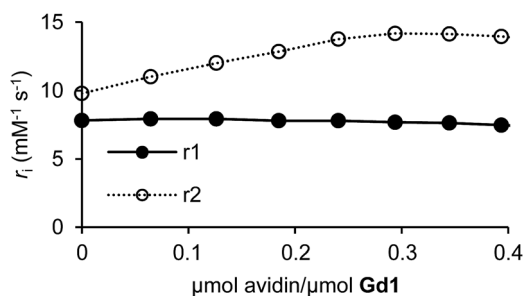
Fig. 1 Formation of a targeted-reporter complex is achieved by conjugating a targeting ligand to an enzyme or its corresponding substrate. Thus, multiple cell surface receptors may be targeted simply by varying the ligands used. Cells overexpressing the receptors of interest enable positioning of the targeted enzyme and substrate close to one another, resulting in activation of the MR probe *via* a self-immolative cascade.



Table 1 Relaxivities and q measurements at 1.41 T and 37 °C in 100 mM MOPS, pH = 7.4


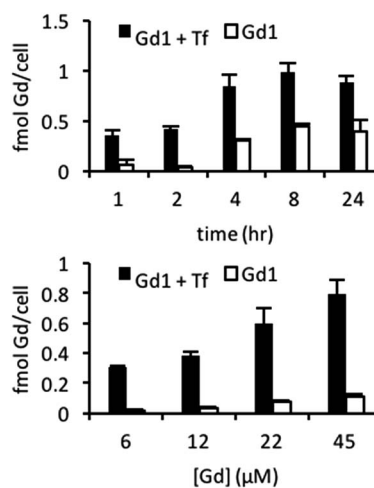
	Relaxivity at 1.41 T (60 MHz)		q	
	(–) Avidin ($\text{mM}^{-1} \text{s}^{-1}$)	(+) Avidin ($\text{mM}^{-1} \text{s}^{-1}$)	(–) Avidin	(+) Avidin
Gd1	7.8	7.8	0.95	0.14
Gd1 + β -gal	4.2	17.2	1.3	1
Gd2	8.6	19.3	1.0	0.9
Gd2 + β -gal	4.2	n.a.	1.2	n.a.

subsequently examined in 9L cells known to overexpress TfR.¹¹ Cells incubated with targeted agent (20 μM Gd(III), 6.6 μM Tf) and untargeted agent (20 μM Gd(III), 0 μM Tf) for 1, 2, 4, 8, or 24 hours show higher labelling for the targeted agent of approximately 0.3–0.4 fmol Gd(III) per cell at all time points (Fig. 3). Additionally, at short incubation times of 1 and 2 hours, the magnitude of labelling by the untargeted agent is less than 0.1 fmol Gd(III) per cell indicating that short incubation times are necessary to minimize the magnitude of non-specific binding. We therefore examined uptake in cells incubated with increasing concentrations of targeted and untargeted contrast agents for 2 hours (Fig. 3). At each concentration, there is significant preferential uptake of the targeted contrast agent. However, there is no significant increase in uptake of targeted contrast agent between incubations of 22 and 45 μM Gd(III) indicating that cells have been saturated with the agent. These findings demonstrate that we are well within the limit of detection necessary for bioactivatable Gd(III)-based agents (*vide infra*).^{33–35} However, attempts to visualize combinatorial biomarkers through colocalization of TfR-targeted **Gd1** and EGFR-targeted β -gal and subsequent activation of the former *in vitro* were unsuccessful. We conjectured that this finding was a result sluggish MR activation.

Fig. 2 r_1 and r_2 as a function of avidin equivalents for **Gd1** (0.15 mM).

We investigated the kinetics associated with biotinylated β -gal activation required for subsequent targeted-reporter imaging. Biotinylated β -gal activity was established using *ortho*-nitrophenyl- β -galactoside (ONPG) through spectrophotometric measurements (Fig. S22†). Initial rates were determined and the reciprocal initial velocities ($1/V_0$) plotted against the corresponding inverse concentrations ($1/[\text{ONPG}]$) to afford a Lineweaver–Burk plot (Fig. S23†). We examined ONPG activation by this enzyme while complexed with 0.25 equivalents of avidin (Fig. S24 and S25†). The corresponding host–guest interaction was found to exert little influence on the observed k_{cat} , indicating this platform's suitability for subsequent targeted-reporter imaging applications.

Spectrophotometric measurements were repeated in the presence of **Gd1** (Fig. S26 and S27†). However, complexation of

Fig. 3 Time- (top) and concentration-dependent uptake (bottom) of TfR-targeted and untargeted **Gd1** illustrate significant uptake when complexed with biotinylated transferrin.

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