INORGANIC CHEMISTRY

FRONTIERS

RESEARCH ARTICLE

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Cite this: Inorg. Chem. Front., 2022, 9, 2271

The critical role of ligand topology: strikingly different properties of Gd(III) complexes with regioisomeric AAZTA derivatives†

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The substitution of an acetate pendant arm on the endocyclic or exocyclic nitrogen atoms of AAZTA with a hydroxybenzyl group results in two regioisomeric Gd(III) complexes with different hydration numbers, thermodynamic stabilities differing by $5.5 \log K$ units and remarkably different kinetic inertness. The ligand functionalized with the phenol group on the exocyclic N atom (AAZ3A-exoHB) forms a Gd(III) complex with remarkably high stability (log K_{Grll} = 25.06) thanks to the tight coordination of the phenol group, which presents a rather low protonation constant (log K_{GdHL} = 3.22). Conversely, the complex formed with the ligand bearing a phenol unit attached to an endocyclic N atom (AAZ3A-endoHB) is considerably less stable (log K_{GdL} = 19.57) and more prone to protonation (log K_{GdHL} = 6.22). Transmetallation kinetics studies in the presence of Cu(II) evidence that the Gd(III) complexes dissociate via the proton- and metal-assisted dissociation pathways, with the AAZ3A-exoHB derivative being considerably more inert. A detailed ¹H nuclear magnetic relaxation dispersion (NMRD) study coupled with ¹⁷O NMR measurements demonstrates that the complex with AAZ3A-exoHB contains a single water molecule in the inner coordination sphere, while the AAZ3A-endoHB analogue has two water molecules coordinated to the metal ion endowed with significantly different water exchange rates. Finally, a binding study of the two complexes with human serum albumin showed a stronger interaction and higher relaxivity (r_1^b = 36.5 mM⁻¹ s⁻¹ at 30 MHz and 298 K) for Gd (AAZ3A-endoHB) than for Gd(AAZ3A-exoHB). Overall, this study highlights the importance that ligand topology has in the properties of Gd(III) complexes relevant in the field of magnetic resonance imaging (MRI).

Received 28th February 2022, Accepted 28th March 2022 DOI: **10.1039/d2gi00451h**

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Introduction

Among the macrocyclic or mesocyclic polydentate ligands used for imaging applications, AAZTA (6-amino-6-methylperhydro-1,4diazepine tetraacetic acid, Scheme 1) is one of the most versatile platforms with remarkable affinity for both di- and trivalent metal ions.¹⁻⁶ The first and most investigated use of this heptadentate chelator is the preparation of the bis-hydrated Gd(m) complex, which displays excellent properties for its application as a magnetic resonance imaging (MRI) contrast agent (CA).¹ Furthermore, the same ligand has been used to complex efficiently radiometals such as ⁶⁸Ga^{2,3} and ⁴⁴Sc^{4,5} for Positron Emission Tomography (PET) and other lanthanide ions like Yb(m) or Tm(m) for chemical exchange saturation transfer (CEST) studies.^{7,8} It has also been recently reported that, passing from Gd(m) to Yb(m), Ln(AAZTA) complexes change the number of coordinated water molecules from two (q = 2) to one (q = 1) as their



Scheme 1 Structures of the ligands discussed or mentioned in this work.

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[†]Electronic supplementary information (ESI) available: Characterizations of the ligands; supplementary figures and tables. See DOI: https://doi.org/10.1039/ d2qi00451h

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exchange rate decreases by several orders of magnitude.⁸ Moreover, X-ray structural studies and DFT calculations have shown that the complexes adopt a monocapped square antiprismatic structure when q = 2 and a dodecahedral structure when $q = 1.^{7,8}$ Finally, it has been recently highlighted by ¹H relaxometric, ¹⁷O NMR and DFT studies that [Gd(AAZTA)(H₂O)₂]⁻ presents one Gd-bound water molecule with an exchange rate 6 times faster than the other, due to a longer metal–water distance.⁹

Notably, another heptadentate ligand that forms a very inert Gd(m) complex is the functionalized tris-serinolamido PCTA derivative (Gadopiclenol) that shows remarkable relaxivity values due to its large molecular weight and the presence of hydrophilic polyalcohol pendants that favour the additional contribution of second sphere water molecules.¹⁰

The modification of the ligand AAZTA has been pursued with the intent to form more rigid ligands to increase the kinetic inertness of the resulting Gd(III) or Ga(III) complexes^{11,12} and bifunctional chelates suitable for building multimeric structures,¹³ for conjugation to targeting vectors,^{2–5} nanoparticles or hydrophobic moieties.^{14,15} However, the modification of the coordination cage with different pendant arms received limited attention.^{16,17} Only a derivative in which the two acetate groups on the diazepine ring were replaced by methylphosphinate pendant arms¹⁶ and mono- and bis-phosphonate derivatives¹⁷ have been reported. In these examples, however, the presence of a more hindering phosphonate or phosphinate group reduces the space available for the coordination of one water molecule, thus reducing the relaxivity of the final Gd(m) complex. Moreover, two diastereoisomeric di-glutarate AAZTA derivatives were reported forming Gd(III) complexes with different water exchange rates, in both cases lower than that of GdAAZTA.⁵⁹

The importance of the donor group in the coordination of the metal ion and in the properties of the resulting metal complexes requires special attention in the choice of the functional group. Phenolate anions are more basic than carboxylates and thus can be strong donor groups for hard metal ions, although they are easier to protonate in acidic media. In fact, macrocyclic and acyclic chelates functionalized with hydroxybenzyl units are mainly reported in the literature as ligands for hard metal ions such as Fe(m) and Ga(m).¹⁸

In this study, we replaced one acetate pendant arm from the AAZTA structure with hydroxybenzyl moieties and synthesized two regioisomeric ligands shown in Scheme 1 (namely AAZ3A-endoHB and AAZ3A-exoHB) with the aim to study the influence of the phenolate donor on the solution properties of the resulting Gd(m) complexes. The different topologies of the two ligands result in Gd(m) complexes astonishingly diverse in terms of thermodynamic stability and kinetic inertness, key properties for the design of chelates for biomedical application.

Results and discussion

Synthesis of the ligands

The synthesis of the two ligands started from an appropriately substituted 6-aminodiazepine¹⁹ (1 and 3 in Scheme 2) and pro-



Scheme 2 Synthesis of ligands AAZ3A-endoHB and AAZ3A-exoHB: (i) H_2 , 10% Pd/C, rt, overnight; (ii) BMPA, K_2CO_3 , MeCN, reflux, overnight; (iii) 1 M NaOH, 1:1 H₂O/THF, 60 °C, 3 h; (iv) 1:1 DCM/TFA, rt, overnight; (v) BrCH₂COOtBu, K_2CO_3 , MeCN, reflux, overnight.

ceeded *via* a series of alkylations, hydrogenation and deprotection steps. In particular, AAZ3A-endoHB was obtained by removal of the benzyl group of **1** through catalytic hydrogenation, introduction of the protected hydroxybenzyl moiety *via* substitution with 2-(bromomethyl)phenyl acetate (BMPA) and final deprotection of the functional groups through basic hydrolysis followed by acidic treatment. On the other hand, AAZ3A-exoHB was synthesized by alkylation of the secondary amine of **3** with *tert*-butyl bromoacetate followed by deprotection of the *tert*-butyl ester moieties.

Thermodynamic properties

Considering the key importance of the thermodynamic properties of any metal-complex proposed for *in vivo* applications, the equilibrium properties of the two regioisomeric ligands and those of the Gd(π) complexes were investigated in detail. First of all, the protonation constants of both ligands, defined by eqn (1), were determined by pH-potentiometry and the log $K_i^{\rm H}$ values are listed in Table 1.

$$K_i^{\rm H} = \frac{[{\rm H}_i {\rm L}]}{[{\rm H}_{i-1} {\rm L}][{\rm H}^+]} \tag{1}$$

Comparing the log K_i^{H} values of the two ligands with those of AAZTA, it is clear that the main difference is observed for log K_2^{H} , which can be ascribed to the phenolate group. To confirm this, the protonation sequence of AAZ3A-endoHB was also determined by recording the absorption spectra of the ligand in the pH range 2.0–11.5 (Fig. S1†). According to the spectral changes, it can be assumed that the log $K_i^{\text{H}} = 9.2$ (1) value characterizes the protonation of the phenolate group of the AAZ3A-endoHB ligand, as it is also very similar to the p K_a of phenol (log $K^{\text{H}} = 10.0, 0.1 \text{ M NaClO}_4, 25 \text{ °C}).^{20}$

A comparison of the protonation constants (Table 1) indicates that the $\log K_1^{\rm H}$ value of AAZ3A-exoHB is slightly higher than that of AAZ3A-endoHB, which can be explained by the H-bonding between the phenolate group and the protonated exocyclic N-atom which is less likely to occur in case of AAZ3A- **Table 1** Protonation constants (log K_i^{H}) of AAZ3A-endoHB and AAZ3A-exoHB compared with AAZTA and an AAZTA-like ligand (CyAAZTA); stability (log K_{GdL}) and protonation constants (log $K_{Gd(HiL)}$) of Gd(AAZ3A-endoHB), Gd(AAZ3A-exoHB), Gd(CyAAZTA) and Gd(AAZTA) complexes (25 °C). Standard deviations are shown in parentheses

Ι			11.10	AAZTA	
	AAZ3A-exoHB 0.15	AAZ3A-exoHB AAZ3A-endoHB 0.15 M NaCl		0.15 M NaCl ^{21,22}	0.1 M KCl ²³
$\log K_1^{\rm H}$	12.14 (1)	11.47 (4)	10.48	10.06	11.23
$\log K_2^{\rm H}$	10.66 (2)	9.18 (2)	6.43	6.50	6.52
$\log K_3^{\text{H}}$	5.82(4)	6.17 (3)	4.23	3.77	3.78
$\log K_{4}^{H}$	3.84 (4)	3.76 (3)	2.76	2.33	2.24
$\log K_5^{\tilde{H}}$	2.88 (4)	2.69 (3)	1.68	1.51	1.56
$\log K_6^{\rm H}$	1.34 (3)	1.43 (2)	_	_	_
$\Sigma \log K_i^{\rm H}$	36.69	32.14	25.58	24.16	25.33
$\log K_{GdL}$	25.06(4)	19.57 (5)	18.26	18.93	20.24
$\log K_{GdHL}^{a}$	3.22 (3)	6.22 (3)	3.79	2.18	1.89
log K _{GdH-L}	2.51 (6)	3.44 (4)	_	_	_
$\log K_{GdLH-1}$	_ ``	10.51 (4)	_	_	_
pGd ^b	18.00	14.67	16.10	17.17	17.31

^{*a*} Protonation constants of the phenol group obtained by spectrophotometry: Gd(AAZ3A-endoHB), log $K_{GdHL} = 5.82$ (2); Gd(AAZ3A-exoHB, log $K_{GdHL} = 3.99$ (4), 0.15 M NaCl, 25 °C. ^{*b*} p $G_d = -\log[Gd(m)]_{free}$, $[Gd(m)]_{tot} = 1 \mu M$, $[L]_{tot} = 10 \mu M$, pH = 7.4.²⁴

endoHB. On the other hand, $\log K_1^{\rm H}$ of AAZTA (in 0.15 M NaCl) is significantly lower due to the formation of a Na⁺ complex competing with the first protonation process.^{21,22} The lower affinity of AAZ3A-endoHB and AAZ3A-exoHB towards Na⁺ maybe related to the formation of a H-bond between the protonated nitrogen and the basic phenolate group, which can hinder the interaction of the ligands with the Na⁺ ion.

The stability and protonation constants of the Gd(III)-complexes, calculated from the titration curves obtained at 1:1 metal to ligand concentration ratios, are reported in Table 1 and defined by eqn (2)–(4):

$$K_{\rm GdL} = \frac{[\rm GdL]}{[\rm Gd^{3+}][\rm L]} \tag{2}$$

$$K_{\rm GdH_{iL}} = \frac{[\rm Gd(H_{iL})]}{[\rm Gd(H_{i-1}L)][\rm H^{+}]}$$
(3)

$$K_{\rm GdLH_{-1}} = \frac{[\rm GdL]}{[\rm Gd(L)OH^{-}][\rm H^{+}]}$$
(4)

where i = 1-3. The best fitting was obtained by using a model that includes the formation of GdL, GdHL and GdH₂L species in equilibrium. The titration data of AAZ3A-endoHB in the presence of Gd(m) indicate base consuming processes at pH > 9. These processes can be interpreted by assuming the hydrolysis of the metal ion; the coordination of an OH⁻ ion results in the formation of Gd(L)OH species characterized by the equilibrium constant K_{GdLH-1} (eqn (4)).

The deprotonation of the phenol group in the Gd(\mathfrak{m}) complexes was studied by spectrophotometry following the absorbance values at 238, 275 and 291 nm of the aromatic group of the ligand (Fig. S2 and S3†). The protonation constants of the phenol group in the Gd(\mathfrak{m}) complexes were calculated by fitting the absorbance–pH data pairs to eqn (S3).†

While the stability constant of Gd(AAZ3A-endoHB) is comparable with those of AAZTA and CyAAZTA complexes, the data in Table 1 evidence a $\Delta \log K_{GdL}$ between the two regioisomers of 5.49 units in favour of Gd(AAZ3A-exoHB), due to the stronger interaction between the Gd(m) ion and the basic exocyclic phenolate-O⁻ donor than with the endocyclic phenolate. This stronger interaction can be further highlighted by the log K_{GdHL} values related to the protonation of the phenolate donor atom, which is about 3 log *K* unit smaller in Gd(AAZ3AexoHB) than for Gd(AAZ3A-endoHB). Interestingly, the stability constants of Gd(AAZ3A-exoHB) and Gd(DOTA) are comparable and significantly higher (*ca.* 3 log *K* units) than that of Gd(DOTA): log $K_{GdL} = 24.7$; Gd(DTPA): log $K_{GdL} = 22.03$).^{25,26} To the best of our knowledge Gd(AAZ3A-exoHB) is characterized by the highest thermodynamic stability among the Gd(m) complexes formed with heptadentate ligands.

Furthermore, a second lower protonation constant could be determined by pH-potentiometric studies for both Gd(m) complexes, probably due to one more weakly coordinated donor atom (a carboxylate-O), which can be protonated in the pH range 2–5. By taking into account the equilibrium data in Table 1, the species distribution diagrams of the Gd(m)-AAZ3A-endoHB and Gd(m)-AAZ3A-exoHB systems were calculated as a function of pH and shown in Fig. S4 and S5.†

Kinetic inertness

The kinetic inertness of Gd(AAZ3A-endoHB) and Gd(AAZ3A-exoHB) was determined by following the transmetallation reactions between the Gd(m) complexes and the Cu(n) ion *via* spectrophotometry. These experiments were monitored on the absorption band of the resulting Cu(n) complexes in the pH range 2.8–5.0 in the presence of a Cu(n) excess to guarantee the pseudo-first order kinetic condition ([GdL] = 0.2 mM, [Cu(n)] = 2.0–8.0 mM, 0.15 M NaCl, 25 °C). The proposed mechanism for the transmetallation of Gd(m)-complexes is shown in Scheme 3, whereas the definitions and equations used for the evaluation of the kinetic data are reported in the ESI.†

Scheme 3 Proton- and metal-assisted dissociation of Gd(AAZ3AendoHB) and Gd(AAZ3A-exoHB) complexes.

The kinetic data (Fig. S8 and S9^{\dagger}) show that the k_d values increase with increasing $[H^+]$ (particularly at lower $[Cu(\pi)]$) and decrease with increasing $[Cu(\pi)]$ at pH < 4.0. By taking into account the species distribution of the Gd(m)-L systems, the increase in the k_d values with increasing $[H^+]$ can be interpreted in terms of the formation (K_{GdH_2L} , eqn (3)) and the slow dissociation of protonated $Gd(H_2L)$ intermediate (k_{GdH_2L} , eqn $(S6)^{\dagger}$), which is followed by a fast reaction between the free ligands and the exchanging $Cu(\pi)$ metal ions. The transmetallation reaction can also occur via direct attack of the exchanging metal ion on the Gd(III) complex (k_{GdLCu} , eqn (S8)†), through the formation of a dinuclear intermediate (K_{GdLCu}) eqn (S7)[†]). The formation of the dinuclear [Lu(AAZTA)]Cu complex was also observed by ¹H NMR spectroscopy in the presence of a large excess of the Cu(II) ion.²³ It can be assumed that in the dinuclear intermediate Gd(L)Cu, the functional groups of the ligand are slowly transferred from Gd(III) to the attacking Cu(II) step by step (eqn (S7)[†]). The $k_1 = k_{\text{GdH},\text{L}} \times$ $K_{\text{GdH}_{2L}}$ and $k_3 = k_{\text{GdLCu}} \times K_{\text{GdLCu}}$ rate constants, as well as the protonation and stability constants K_{GdH,L} and K_{GdLCu} characterizing the proton- and metal-assisted dissociation for the two complexes, are shown in Table 2 and compared with the corresponding values of Gd(AAZTA) and Gd(CyAAZTA).

The k_1 rate constant characterizing the proton-assisted dissociation of Gd(AAZ3A-exoHB) is about 60 times smaller than that of Gd(AAZ3A-endoHB). Based on the mechanism of the proton-assisted dissociation, the protonated intermediates facilitate the proton transfer from the carboxylic acid/phenol groups to the N-atom of the ligand backbone *via* the formation of a relatively labile protonated intermediate. The proton of this intermediate can displace the Gd(m) ion from the coordination cage resulting in the dissociation of the GdL complex. The stronger interaction between the Gd(m) ion and the phenol group in Gd(AAZ3A-exoHB) results in a less probable proton transfer and slower rate of de-coordination of the Gd(m) ion from the N-donor atoms. Moreover, the transfer of the phenol-OH proton to the endocyclic N atom is more likely for Gd(H₂AAZ3A-endoHB) than for Gd(H₂AAZ3A-exoHB), which can cause the faster H⁺-assisted dissociation of the former Gd(III) complex. However, the proton-assisted dissociation of both Gd(III) complexes is faster than for Gd(AAZTA) and Gd (CvAAZTA), which might be interpreted by the electrostatic repulsion between the protonated phenol-OH group and the metal centre of $Gd(H_2L)$ intermediates, which might weaken the interaction between the ring N donor atoms and the Gd(m) ion in Gd(AAZ3A-endoHB) and Gd(AAZ3A-exoHB) complexes. The k_3 rate constants characterizing the Cu(II) assisted dissociation of Gd(AAZ3A-exoHB) is 75 times slower than that of Gd(AAZ3A-endoHB). These findings suggest that the coordination cage around the Gd(m) ion in the Gd(AAZ3A-exoHB) complex is more compact, hindering the transformation of the dinuclear intermediate to the final $Cu(\pi)$ complex. The stability constants of the dinuclear intermediates formed by Gd(AAZ3AexoHB) and Gd(AAZ3A-endoHB) with Cu(II) are very similar and somewhat higher than that of Gd(AAZTA), as a consequence of the more basic phenolate pendant group. Finally, the 60 times slower dissociation rate constant (k_d) and higher half-lives ($t_{1/2} = \ln 2/k_d$, pH = 7.4, 25 °C) for Gd(AAZ3A-exoHB) indicate that this complex is remarkably more inert than Gd (AAZ3A-endoHB).

Structural properties

The relaxivity values (r_1) for Gd(AAZ3A-endoHB) and Gd (AAZ3A-exoHB) at 30 MHz, 25 °C and pH 7.0 (Table 3) are 7.3 and 5.4 mM⁻¹ s⁻¹, respectively, and reflect different hydration states for the two complexes, consistent with the presence of two water molecules in the inner coordination sphere of the metal ion for Gd(AAZ3A-endoHB) (q = 2) and one for Gd (AAZ3A-exoHB) (q = 1).

Density Functional Theory (DFT) calculations were performed to rationalise the strikingly different behaviour of the regioisomeric Gd(III) complexes. We performed calculations on the $[Gd(L)(H_2O)_2]^{-}\cdot 4H_2O$ systems (L = AAZ3A-endoHB or AAZ3A-exoHB) at the wB97XD/LCRECP/6-311G(d,p) level,^{27,28} including an explicit second-sphere solvation shell²⁹ and a polarized continuum to model bulk solvent effects (see Computational details below). Calculations performed for the

Table 2 Rate (k_i) and equilibrium (K_i) constant characterizing the transmetallation reaction of Gd(III) complexes formed with AAZTA derivatives (25 °C)

Ι	Gd(AAZ3A-exoHB) 0.15 M NaCl	Gd(AAZ3A-endoHB)	Gd(CyAAZTA) ¹² 0.1 M KCl	Gd(AAZTA) ²³ 1.0 M KCl
$k_1 (M^{-1} s^{-1})$	5.1 ± 0.2	300 ± 30	$6.0 imes 10^{-3}$	1.05
$k_2 (M^{-2} s^{-1})$	_	_	53	_
k_{3}^{Cu} (M ⁻¹ s ⁻¹)	$(4 \pm 1) \times 10^{-3}$	0.30 ± 0.05	_	$1.9 imes 10^{-4}$
$K_{\rm GdH_{2L}}({\rm M}^{-1})$	238 ± 37	2400 ± 600	6166 (GdHL)	233 (GdHL)
$K_{\rm GdLCu}$ (M ⁻¹)	27 ± 7	37 ± 10	_ ``	9
$k_{\rm d} ({\rm s}^{-1}) {\rm pH} = 5.0$	$5.1 imes 10^{-5}$	2.9×10^{-3}	6.2×10^{-8}	$1.0 imes 10^{-5}$
$t_{1/2}$ (h) pH = 5.0	3.76	0.07	3106	18.4
$k_{\rm d} ({\rm s}^{-1}) {\rm pH} = 7.4$	2.0×10^{-7}	$1.1 imes 10^{-5}$	$2.4 imes 10^{-10}$	$4.0 imes10^{-8}$
$t_{1/2}$ (h) pH = 7.4	939	15.8	$8.0 imes 10^5$	$4.3 imes 10^3$

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[Gd(AAZ3A-endoHB)(H₂O)₂]⁻·4H₂O system predicted a nine-coordinated structure with two inner-sphere water molecules (Fig. 1), which is more stable than the octacoordinated species by $\Delta G^{\circ} = 6.6 \text{ kJ mol}^{-1}$. The reverse situation is observed for the [Gd(AAZ3A-exoHB)(H₂O)₂]⁻·4H₂O complex, the octacoordinated species being favoured by $\Delta G^{\circ} = 12.9 \text{ kJ mol}^{-1}$. This is in nice agreement with the experimental ¹⁷O NMR data (see below). The analysis of the thermodynamic contributions shows that the hydration reactions [Gd(L)(H₂O)₂]⁻ \rightleftharpoons [Gd(L) (H₂O)]⁻ + H₂O are characterised by similar negative reaction entropies with the enthalpy contribution being the main



Fig. 1 Structures of the $[Gd(AAZ3A-endoHB)(H_2O)_2]^-$ (bottom) and $[Gd(AAZ3A-exoHB)(H_2O)]^-$ (top) systems obtained with DFT calculations. Second-sphere water molecules are omitted for simplicity. Bond distances (Å), $[Gd(AAZ3A-endoHB)(H_2O)_2]^-$: Gd–N1, 2.734; Gd–N2, 2.725; Gd–N3, 2.701; Gd–O1, 2.391; Gd–O2, 2.426; Gd–O3, 2.402; Gd–O4, 2.452; Gd–O1w, 2.515; Gd–O2w, 2.518. $[Gd(AAZ3A-exoHB)(H_2O)_2]^-$: Gd–N1, 2.661; Gd–N2, 2.649; Gd–N3, 2.681; Gd–O1, 2.403; Gd–O2, 2.421; Gd–O3, 2.360; Gd–O4, 2.325; Gd–O1w, 2.470.

responsible for the different hydration numbers (Tables S1–S4, ESI†).

The coordination polyhedron around the Gd(III) ions in [Gd $(AAZ3A-endoHB)(H_2O)_2$ can be described as a capped square antiprism, where the capping position is occupied by an oxygen atom of one of the carboxylate groups. Conversely, the coordination polyhedron in [Gd(AAZ3A-exoHB)(H₂O)]⁻ is a biaugmented trigonal prism, with the exocyclic N atom and the coordinated water molecule occupying the capping positions (Fig. 1). The most important difference in bond distances is observed for the Gd(III)-O_{phenol} bond, which is considerably shorter for $[Gd(AAZ3A-exoHB)(H_2O)]^-$ (Gd-O4 = 2.325 Å) than for $[Gd(AAZ3A-endoHB)(H_2O)_2]^-$ (Gd–O1 = 2.391 Å). This indicates that the phenolate group in the latter complex occupies a sterically demanding position, resulting in a weaker interaction with the metal ion. As a result, the phenolate group in the $[Gd(AAZ3A-endoHB)(H_2O)_2]^-$ complex is characterised by a higher protonation constant ($\log K_{GdHL} = 6.22$) than the [Gd $(AAZ3A-exoHB)(H_2O)$ regionsomer $(\log K_{GdHL} = 3.22)$. Thus, the presence of the bulky phenolate group attached to the endocyclic N atom introduces an important steric hindrance for the coordination to the Gd(m) ion, an effect that is clearly reflected in the impressive difference in stability of the two complexes. A weak coordination of the phenolate group in [Gd $(AAZ3A-endoHB)(H_2O)_2$ ⁻ likely facilitates the protonation of the complex, and thus its dissociation following the acid-catalysed mechanism.

Relaxation properties

The relaxometric characterization of the Gd(III) complexes started by measuring the pH dependence of r_1 for the two Gd^{III} complexes to assess possible variations of hydration state or coordination environment with pH (Fig. S13 and S14†). In particular, r_1 of Gd(AAZ3A-endoHB) is almost constant from pH = 3 to pH = 11. This indicates that over this broad range of pH values the complex does not hydrolyze and does not vary the hydration number *q*. In case of Gd(AAZ3A-exoHB), a slight and constant decrease of r_1 from pH 4 to pH 7 is observed

Table 3 Parameters obtained from the fits of ¹⁷O NMR and ¹H NMRD data for Gd(AAZ3A-endoHB) and Gd(AAZ3A-exoHB)^{a,b}

Parameter	Gd(AAZ3A-endoHB)	Gd(AAZ3A-exoHB)	Gd(AAZTA) ^{1,8}
$^{298}r_1$ (30 MHz, pH 7) (mM ⁻¹ s ⁻¹)	7.3 ± 0.1	5.4 ± 0.1	7.1
$\Delta^2/10^{19} \mathrm{s}^{-2}$	8.6 ± 0.3	2.0 ± 0.2	2.2
$^{298}\tau_{\rm V}/{\rm ps}$	16.2 ± 1.1	29.1 ± 1.6	31
$^{298}\tau_{\rm M}^{\rm A}/\rm{ns}$	35 ± 3	20 ± 1	29
$\Delta H_{M}^{A'}/kJ \text{ mol}^{-1}$	15.0 ± 0.5	15.9 ± 0.4	20
r_{M}^{298} r_{M}^{B}/ns	290 ± 10	_	169
$\Delta H_{M}^{B}/kJ \text{ mol}^{-1}$	20.0 ± 0.3	_	29.5
$A_{0}^{A}/\hbar/10^{6} \text{ rad s}^{-1}$	-3.3 ± 0.1	-3.3 ± 0.1	-3.8
$A_{\rm O}^{\rm B}/\hbar/10^{\rm 6} {\rm ~rad~s^{-1}}$	-3.6 ± 0.1	_	-3.9
$298\tau_{\rm R}/{\rm ps}$	79 ± 2	80 ± 1	74
q	2	1	2

^{*a*} Some of the parameters affecting r_1 and the ¹⁷O NMR data were fixed to the values reported in literature for GdAAZTA; the water proton-gadolinium distance (r = 3.05 Å), the outer sphere relaxation parameters (a = 4.0 Å, ²⁹⁸ $D = 2.24 \times 10^5$ cm² s⁻¹) and the activation energy for the diffusion coefficient ($E_D = 20$ kJ mol⁻¹). Then, the activation energies of $\tau_V (E_V)$ and $\tau_R (E_R)$ were set to 1.0 and 18.0 kJ mol⁻¹, respectively, and the distance r_{GdO} was fixed at 2.5 Å. ^{*b*} Information on q and on the scalar coupling A_O/h are derived from the temperature dependence of $\Delta \omega_r$.



Fig. 2 ¹H nuclear magnetic relaxation dispersion (NMRD) profiles at pH 7.0 of Gd(AAZ3A-endoHB) (A) and Gd(AAZ3A-exoHB) (B) recorded at different temperatures 283 K (\blacksquare), 298 K (\bullet) and 310 K (\blacklozenge). The solid lines correspond to the fits of the data as described in the text.

(from about 7 to 5.5 mM⁻¹ s⁻¹) and then it remains almost constant going to basic pH. The slightly higher r_1 at acidic pH maybe due to several reasons: (1) the presence of an equilibrium between q = 1 and q = 2 species occurring after protonation of the phenolate oxygen as can be seen from the species distribution diagram in Fig. S5;† (2) a contribution to the relaxivity of acid catalyzed proton exchange or (3) second sphere water molecules. Finally, for both complexes, at pH < 3 r_1 increases likely because of proton-assisted metal release.

Then, the ¹H nuclear magnetic relaxation dispersion (NMRD) profiles (Fig. 2) and the temperature dependence of the transverse ¹⁷O NMR relaxation rate (R_{2r}) and chemical shifts $(\Delta \omega_r)$ were recorded for both complexes (Fig. 3 and S10, S11[†]). Notably, the shape of the variable-temperature ¹⁷O NMR R_2 profile of Gd(AAZ3A-endoHB) (Fig. 3A) evidences the presence of two water molecules with significantly different water exchange dynamics; in fact, the curve is distant from the simple pseudo-exponential decay expected for systems containing one (or two) coordinated water molecule(s) in exchange with bulk solvent, as shown, conversely, by the ${}^{17}OR_2$ profile of Gd(AAZ3A-exoHB) (Fig. 3B). ¹H NMR spectra recorded for the Eu(AAZ3A-endoHB) complex (273-313 K, Fig. S15†) exclude that the shape of the ¹⁷O NMR R_2 profile is related to the presence of different coordination isomers with water molecules exchanging at different rates, as observed in the case of some GdDOTA-like complexes.³⁰ Another possibility considers the



Fig. 3 ¹⁷O NMR transverse relaxation rates as a function of temperature measured at pH 7.0 and 500 MHz (11.74 T) for Gd(AAZ3A-endoHB) (A, 2.6 mM) and Gd(AAZ3A-exoHB) (B, 12.3 mM). The solid lines correspond to the fits of the data as described in the text, while the blue and green lines in (A) represent the calculated contributions of the two different water species, in faster and slower exchange with the bulk, respectively.

presence of a mixture of a protonated and a non-protonated species at about 1:1 ratio at pH 7 with substantially different water exchange rates. Thus, ¹⁷O NMR data were recorded also at pH 5 and 8 for Gd(AAZ3A-endoHB) (Fig. S12†) where the species with the protonated and deprotonated phenol, respectively, are predominating (see Fig. S10†). The qualitative analysis of the profiles highlights that at acidic pH both water molecules exchange faster, whereas at pH 8 there is no difference with the profile at pH 7. Therefore, we can exclude that the two species exchanging differently are the protonated and deprotonated and deprotonated and deprotonated and deprotonated that the two species exchanging differently are the protonated and deprotonated complexes.

Thus, the ¹⁷O NMR and NMRD data for Gd(AAZ3A-endoHB) were analysed by considering the presence in solution of two water molecules subject to different exchange rates, whereas for Gd(AAZ3A-exoHB) the analysis considered the presence of a q = 1 complex. The ¹H NMRD and ¹⁷O NMR data (Fig. 2 and 3), were fitted simultaneously according to the established theory of paramagnetic relaxation.³¹ The data for Gd(AAZ3A-endoHB) were well reproduced by considering that the two Gd(m)-bound water molecules are characterized by significantly different residence lifetimes ($\tau_M^A = 35$ ns and $\tau_M^B = 290$ ns) and enthalpies associated to the exchange process ($\Delta H_M^A = 15$ kJ mol⁻¹ and $\Delta H_M^B = 20$ kJ mol⁻¹). The water molecule residing for shorter times on the metal center (A) exchanges more than 8 times faster than the other with a reaction enthalpy 1.3 times lower.

As observed for Gd(AAZTA),⁸ it is reasonable to assume that breaking the Gd–O bond of the water molecule less strongly bound to the metal centre and with a longer Gd–O distance requires the lowest energy. In case of Gd(AAZ3A-exoHB) the coordinated water molecule exchanges very fast with the bulk (²⁹⁸ $\tau_{\rm M}$ = 20 ns), similar to the faster water molecule of Gd (AAZ3A-endoHB) and more than four times faster than that measured for other GdAAZTA-like complexes.^{1,12} The higher water exchange rate may be ascribed to the increased steric compression near the water coordination site imposed by the phenolate donor group, as shown already in cases where a sixmembered chelate ring is formed.^{32,33}

Moreover, $\tau_{\rm R}$ values around 80 ps and the electronic parameters characterizing the two complexes are in line with the corresponding values obtained for GdAAZTA-like complexes.^{1,12} The fitted ¹⁷O hyperfine coupling constants for Gd(AAZ3A-endoHB) show a higher value for the more strongly bound and more slowly exchanging water molecule ($A_O/\hbar^A =$ -3.3×10^6 rad s⁻¹, $A_O/\hbar^B = -3.6 \times 10^6$ rad s⁻¹) whereas for Gd (AAZ3A-exoHB) the value is similar to that of the faster exchanging molecule (-3.3×10^6 rad s⁻¹).

Relativistic DFT calculations (TPSSh/SARC2-DKH-QZVP/ DKH-def2-TZVPP³⁴⁻³⁷ see below) provide calculated values in excellent agreement with the experiment $(A_O/\hbar^A = -3.0 \times 10^6 \text{ rad s}^{-1} \text{ and } A_O/\hbar^B = -3.6 \times 10^6 \text{ rad s}^{-1}$ for Gd(AAZ3A-endoHB), $A_O/\hbar^B = -3.7 \times 10^6 \text{ rad s}^{-1}$ for Gd(AAZ3A-exoHB), confirming that the hydration numbers assumed for the analysis of the ¹H NMRD and ¹⁷O NMR data are correct. The zero field splitting (ZFS) parameters obtained using NEVPT2 calculations (Table S5, ESI†) provide Δ^2 values of 8.1×10^{19} and 6.5×10^{19} rad² s⁻² for [Gd(AAZ3A-endoHB)(H₂O)₂]⁻ and [Gd(AAZ3A-

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exoHB)(H₂O)]⁻, respectively, again in reasonable agreement with the experiment. The lower value of Δ^2 calculated for the latter complex appears to be related to the smaller splitting of the four Kramer doublets associated to the ⁸S_{7/2} ground state.

Interaction with human serum albumin (HSA)

We argued that the presence of a phenol ring in these Gd(m)complexes would enable non-covalent interaction with hydrophobic binding sites on HSA, the most abundant serum protein in blood and a primary biological target for MRI contrast agents. In fact, the formation of supramolecular adducts between a paramagnetic probe and HSA is accompanied by a strong enhancement of the relaxivity because of the decreased tumbling rate of the system. This is generally exploited for increasing the lifetime of the probe in the vascular system for angiographic MRI applications. Notably, in the case of Gd (AAZ3A-endoHB) and Gd(AAZ3A-exoHB) the phenolate group can act both as donor group for the lanthanide ion and as hydrophobic group for HSA interaction. The longitudinal ¹H relaxation rate (R_1 ; 30 MHz and 298 K) of diluted aqueous solutions of the two complexes were measured following the



Fig. 4 (A) Water proton relaxation rate of aqueous solutions of Gd (AAZ3A-endoHB) (0.1 mM, circles) and Gd(AAZ3A-exoHB) (0.1 mM, squares) as a function of increasing amounts of HSA, measured at 20 MHz and 298 K. (B) $1/T_1$ NMRD profiles of the Gd(AAZ3A-endoHB)-HSA (circles) and Gd(AAZ3A-exoHB)-HSA adducts (squares), collected at 298 K. The solid lines through the data points are calculated using the parameters reported in Table 4. The data were analyzed only in the magnetic field region >1 T because of the known limitations of the SBM theory to completely account for the behaviour of slowly rotating systems at low magnetic field strengths.

addition of increasing amount of HSA (PRE method, Fig. 4A). The analysis of the relaxometric binding isotherm allows estimating the affinity constant, K_A , for the protein and the relaxivity of the bound complex, $r_1^{\rm b}$. Both Gd(m)-chelates interact with the protein, although to a different extent. The fit of the experimental data, performed considering two independent binding sites (n = 2), gave quite different values for the two complexes: K_A for Gd(AAZ3A-endoHB)-HSA is more than one order of magnitude higher than for Gd(AAZ3A-exoHB)-HSA (Table 4) probably due to the higher availability of the aromatic moiety to interact with the hydrophobic pocket of the protein. The stronger interaction of Gd(AAZ3A-endoHB) parallels to the longer distance and weaker interaction between Gd(m) and the phenolic oxygen with respect to Gd(AAZ3A-endoHB). Also $r_1^{\rm b}$ resulted to be three times greater for the Gd(AAZ3A-endoHB)-HSA adduct as a result of the two coordinated water molecules present in Gd(AAZ3A-endoHB).

The NMRD profiles of the two supramolecular adducts were recorded at neutral pH and 298 K under conditions ensuring that more than 98% of the chelate was bound to the protein (Fig. 4B). The high field region of the profiles (>1 T) was fitted to the SBM theory including the Lipari-Szabo approach for the description of the rotational dynamics.^{38,39} This model considers a local rotation of the complex superimposed on the global reorientation of the supramolecular adduct. These two types of motion are characterized by different correlation times: $\tau_{\rm RL}$ and $\tau_{\rm RG}$, respectively, related by the parameter S^2 , which takes values between 0 (completely independent motions) and 1 (entirely correlated motions). A four-parameter $(\Delta^2, \tau_{\rm V}, \tau_{\rm RL} \text{ and } S^2)$ least-squares fit of the data was performed and the best-fit parameters are listed in Table 4 and compared with the supramolecular system obtained by interacting an amphiphilic Gd(AAZTA)-like complex with HSA.40 The analysis of the parameters remarks again the better interaction of Gd (AAZ3A-endoHB) with HSA than Gd(AAZ3A-exoHB), highlighted by the higher values of $\tau_{\rm RL}$ and S^2 , translated into a higher relaxivity (although caused also by the q = 2 of Gd (AAZ3A-endoHB)). Moreover, the comparison with the GdAAZTA-C12-HSA adduct shows that the C12 alkyl chain strongly interacts with the hydrophobic pocket of HSA result-

Table 4 Selected best-fit parameters obtained from the analysis of the $1/T_1$ NMRD profiles of Gd(AAZ3A-endoHB)-HSA and Gd(AAZ3A-exoHB)-HSA adducts collected at 298 K compared to the analogous data obtained for Gd(AAZTA-C₁₂)^{40 a}

Parameter	Gd(AAZ3A-endoHB)-HSA	Gd(AAZ3A-exoHB)-HSA	Gd(AAZTA C ₁₂)-HSA
r_{1} (30 MHz) (mM ⁻¹ s ⁻¹)	36.5	11.4	52.5
$\Delta^2/10^{19} \mathrm{s}^{-2}$	2.4 ± 0.2	3.4 ± 0.2	1.5
$^{298}\tau_{\rm V}/{\rm ps}$	10 ± 2	42 ± 4	10
$^{298}\tau_{\rm M}/\rm{ns}$	160^b	21.3	90
$^{298}\tau_{\rm RG}/\rm ns$	40	40	80
$^{298}\tau_{\rm RL}/{\rm ps}$	355 ± 20	98 ± 8	587
S^2	0.23 ± 0.01	0.07 ± 0.01	0.18
q	2	1	2
\bar{K}_{A}/M^{-1}	$7.5 imes 10^4$	$1.5 imes 10^3$	$8.5 imes 10^4$

^{*a*} The distance of the hydrogen atoms of the coordinated water molecule from the metal ion (r_{GdH}) was fixed to 3.0 Å. The outer-sphere component of the relaxivity was estimated by using standard values for the distance of closest approach a (4 Å) and the relative diffusion coefficient of solute and solvent D (2.25 × 10⁻⁵ cm² s⁻¹). ^{*b*} The average between $\tau_{\text{M}}^{\text{A}}$ and $\tau_{\text{B}}^{\text{B}}$ for Gd(AAZ3A-endoHB) was used.

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ing also into a slower local rotation of the complex around the alkyl chain (higher $\tau_{\rm RL}$ value).

Conclusions

In conclusion we have highlighted the importance of ligand topology in two AAZ3A-HB chelators, which has a huge effect on the hydration number, stability and dissociation kinetics of two regioisomeric Gd(III) complexes. In fact, the replacement of an endocyclic acetate group of AAZTA with a 2-hydroxybenzyl group results in a q = 2 Gd-complex with higher relaxivity (5.9 mM⁻¹ s⁻¹ at 20 MHz and 310 K), but lower thermodynamic stability (log K_{GdL} = 19.57) and kinetic inertness ($t_{1/2}$ = 15.8 h at pH 7.4). Contrarily, the insertion of the phenolic pendant on the exocyclic nitrogen allows the formation of a more compact octadentate q = 1 Gd(III) complex with lower relaxivity (4.0 mM⁻¹ s⁻¹ at 20 MHz and 310 K), but remarkable stability (log K_{GdL} = 25.06) and good kinetic inertness ($t_{1/2}$ = 960 h at pH 7.4). The relevance of ligand topology has also nicely recently been demonstrated for a highly inert constrained pentaazamacrocyclic Cu(II) complex.⁴¹ The different properties of metal complexes with regioisomeric ligands have been highlighted previously in other examples, *i.e.* the $Mn(\pi)$ complexes of 1,4- and 1,7-DO2A which showed different hydration, and thus different relaxivities,⁴² but similar thermodynamic stability and kinetic inertness.43 However, the remarkable $\Delta \log K_{GdL}$ of 5.5 between AAZ3A-exo and endoHB Gd(m) complexes shows that in this case the ligand topology plays a remarkable role in metal ion coordination properties.

Experimental section

Chemicals and materials

All chemicals were purchased from commercial sources and were used without further purification unless otherwise stated. NMR spectra were recorded on Bruker Avance III spectrometers operating at 9.39 and 11.74 T, corresponding to protonic resonance frequencies of 399.8 and 499.8 MHz. ¹H and ¹³C NMR chemical shifts are reported relative to TMS and are referenced using the residual proton solvent resonances. Splitting patterns are described as singlet (s), broad singlet (bs), doublet (d), triplet (t) or multiplet (m). ESI mass spectra were recorded on a Waters SQD 3100 (Waters Corporation, Milford, MA, USA). Analytical HPLC-MS were carried out on Waters modular system equipped with Waters 1525 binary pump, Waters 2487 UV/Vis and Waters SQD 3100 (ESCI ionization mode) detectors, using an XBridge[™] Phenyl 3.5 µm 4.6 × 150 mm column (Waters). Semi-preparative HPLC purifications were performed with an XBridge[™] Prep Phenyl 5 µm OBD[™] 19 × 100 mm column (Waters). The HPLC methods are reported in the ESI.†

Synthesis

N,N,1-Tris(tert-butylacetate)-4-(2-acetoxybenzyl)-6-methyl-6aminoperhydro-1,4-diazepine (2). Compound 1 (0.14 g,

0.25 mmol) was dissolved in MeOH. 10 w/w% Pd/C (0.014 g) was added and the suspension was stirred under H2-atmosphere (balloon) at room temperature overnight. After filtration through Celite and evaporation of the filtrate, the obtained debenzylated intermediate was dissolved in MeCN (10 mL). K₂CO₃ (0.066 g, 0.48 mmol) was added followed by 2-(bromomethyl)phenyl acetate (0.11 g, 0.48 mmol), and the suspension was refluxed overnight. After evaporation of the solvent, the residue was suspended in EtOAc (20 mL) and washed with H_2O (2 × 10 mL) and brine (10 mL). The organic phase was dried over anhydrous Na₂SO₄, filtered and evaporated. The crude product was purified by flash chromatography (SiO₂, PE/ EtOAc 80:20, $R_{\rm f}$ = 0.20), leading to the aimed product as a yellow oil (0.048 g, 0.077 mmol, 32%).

¹H NMR (500 MHz, 25 °C, CDCl₃), δ (ppm): 7.55 (d, ³J_{HH} = 7.0 Hz, 1H, CH^{Ar}), 7.21 (m, 2H, $2 \times CH^{Ar}$), 6.97 (d, ${}^{3}J_{HH} = 7.5$ Hz, 1H, CH^{Ar}), 3.72 (s, 4H, $N(CH_2CO)_2$), 3.54 (d, ${}^2J_{HH}$ = 13.9 Hz, 1H, NCHH'Ar), 3.42 (d, ²*J*_{HH} = 13.9 Hz, 1H, NCHH'Ar), 3.25 (s, 2H, NCH₂CO), 3.05 (d, ${}^{2}J_{HH}$ = 14.3 Hz, 1H, NCHH'CN), 2.99 (d, ${}^{2}J_{HH}$ = 13.6 Hz, 1H, NCH"H"'CN), 2.73 + 2.63 + 2.47 + 2.35 (m, 4H, NCH₂CH₂N), 2.67 (d, ${}^{2}J_{HH}$ = 14.3 Hz, 1H, NCHH'CN), 2.53 (d, ${}^{2}J_{HH}$ = 13.6 Hz, 1H, NCH"<u>H</u>"'CN), 2.30 (s, 3H, CH₃CO), 1.42 (s, 27H, CH₃^{tBu}), 1.04 (s, 3H, CH₃). ¹³C NMR (125 MHz, 25 °C, CDCl₃), δ (ppm): 172.8 (N(CH₂CO)₂), 170.9 (NCH₂CO), 169.2 (CH₃<u>C</u>O), 149.1 (OC^{Ar}), 131.8 (CH₂<u>C</u>^{Ar}), 130.3 (CH^{Ar}), 127.7 (CH^{Ar}), 126.0 (CH^{Ar}), 122.1 (CH^{Ar}), 80.4 (C^{tBu}), 67.7 (NCH₂CN), 66.4 (NC'H₂CN), 62.1 (NCH₂CO), 60.7 (NCCH₃), 59.2 (NCH₂CH₂N), 58.1 (NCH₂Ar), 51.6 (N(CH₂CO)₂), 28.2 (CH_3^{tBu}) , 24.2 (NCCH₃), 20.9 (CH₃CO). ESI⁺ MS: m/z 620.6 [M + H^+], calc. for $[C_{33}H_{54}N_3O_8]^+ = 620.80 \text{ g mol}^{-1}$.

AAZ3A-endoHB. Intermediate 2 (0.38 g, 0.61 mmol) was dissolved in THF (40 mL), 1 M NaOH (40 mL) was added and the biphasic system was stirred vigorously at 60 °C for 3 h. The solvents were evaporated and the residue was suspended in EtOAc (100 mL) and washed with H_2O (2 × 50 mL) and brine (50 mL). The organic phase was dried over anhydrous MgSO₄, filtered and evaporated. The obtained residue was dissolved in DCM (10 mL), triisopropylsilane (2 drops) and TFA (10 mL) were added and the solution was stirred at room temperature overnight. After removal of the volatiles under reduced pressure, the remaining solid was re-dissolved in TFA (1 mL) and precipitated in Et₂O (10 mL). The suspension was centrifuged (4000 rpm, 15 min, 10 °C) and the precipitate was washed/centrifuged with Et_2O (3 × 10 mL) and dried under vacuum. The product was purified by semi-preparative HPLC (see ESI[†]), re-dissolved in 1 M HCl (2 mL) and stirred for 2 h. The solution was evaporated, the residue dissolved in H₂O (2 mL) and lyophilized, and the product (mono hydrochloride salt) was obtained as a white powder (0.12 g, 0.26 mmol, 43%).

¹H NMR (500 MHz, 25 °C, D_2O), δ (ppm): 7.4–7.3 (m, 2H, $2 \times CH^{Ar}$), 6.94 (m, 2H, $2 \times CH^{Ar}$), 4.36 (d, ²J_{HH} = 13.1 Hz, 2H, NCHH'Ar), 4.32 (d, ²*J*_{HH} = 13.1 Hz, 2H, NCHH'Ar), 3.87 (m, 2H, $NCH_2CH'_2N$), 3.3–3.7 (m, 12H, 3 × $NCH_2CO + 2 × NCH_2C +$ NCH₂CH'₂N), 1.12 (CH₃). ¹³C NMR (125 MHz, 25 °C, D₂O), δ (ppm): 174.3 (N(CH₂CO)₂), 172.7 (NCH₂CO), 155.7 (COH^{Ar}), $133.1 + 132.3 + 120.9 + 116.1 (4 \times CH^{Ar}), 115.6 (CH_2C^{Ar}), 62.8$ *N*,1,4-Tris(*tert*-butylacetate)-*N*-(2-acetoxybenzyl)-6-methyl-6aminoperhydro-1,4-diazepine (4). Compound 3 (789 mg, 1.70 mmol) was dissolved in DMF (30 mL). K₂CO₃ (470 mg, 3.40 mmol) was added, followed by *tert*-butyl bromoacetate (0.30 mL, 2.04 mmol), and the mixture was stirred at 80 °C overnight. The solvent was evaporated under reduced pressure, the residue was suspended in EtOAc (40 mL) and washed with H₂O (2 × 20 mL) and brine (20 mL). The organic phase was dried over anhydrous Na₂SO₄, filtered and evaporated under vacuum. The crude product was purified by flash chromatography (SiO₂, DCM/MeOH 99:1 to 95:5, $R_{\rm f}$ (95:5) = 0.31), leading to the aimed compound as a yellow-colored oil (642 mg, 65%).

¹H NMR (500 MHz, 25 °C, CDCl₃), δ (ppm): 7.69 (m, 1H, CH^{Ar}), 7.17 (m, 1H, CH^{Ar}), 6.93 (m, 1H, CH^{Ar}), 4.03 (s, 2H, CH₂Ar), 3.38 (s, 2H, CH₂CO), 3.26 (s, 4H, 2 × CH₂CO), 3.13 (d, ²J_{HH} = 13.9 Hz, 2H, NC<u>H</u>H'CN), 2.8–2.6 (m, 6H, NCH<u>H</u>'CN + NCH₂CH₂N), 1.42 + 1.33 (s, 27H, CH₃^{tBu}), 1.11 (s, 3H, CH₃). ¹³C NMR (125 MHz, 25 °C, CDCl₃), δ (ppm): 172.8 + 170.9 (3 × CO0tBu), 148.8 (C^{Ar}), 133.2 (C^{Ar}), 130.1 (CH^{Ar}), 127.2 (CH^{Ar}), 126.2 (CH^{Ar}), 121.2 (CH^{Ar}), 80.7 + 80.0 (2 × C^{tBu}), 65.2 (N<u>C</u>H₂CN), 62.4 (2 × NCH₂CO), 59.1 (NCH₂CH₂N), 52.2 (N<u>C</u>H₂CO), 47.2 (CH₂Ar), 28.2 (CH₃^{tBu}), 20.8 (CH₃). ESI⁺ MS: m/z 578.9 [M + H⁺] (calc. for [C₃₁H₅₂N₃O₇]⁺: 578.66 g mol⁻¹).

AAZ3A-exoHB. Intermediate 4 (0.24 mg, 0.41 mmol) was dissolved in DCM (2 mL), triisopropylsilane (3 drops) and TFA (2 mL) were added and the solution was stirred at room temperature for 5 h. After removal of the volatiles under reduced pressure, the residue was dissolved again in TFA (1 mL) and precipitated in Et₂O (10 mL). The suspension was centrifuged (4000 rpm, 15 min, 10 °C) and the precipitate was washed/centrifuged with Et₂O (3 × 10 mL) and dried under vacuum. The product was obtained as a white powder (57 mg, 65%).

¹H NMR (500 MHz, 25 °C, D₂O), δ (ppm): 7.4–7.3 (m, 2H, 2 × CH^{Ar}), 6.96 (m, 2H, 2 × CH^{Ar}), 4.38 (d, ²J_{HH} = 13.1 Hz, 1H, C<u>H</u>H'Ar), 4.34 (d, ²J_{HH} = 13.1 Hz, 1H, CL<u>H</u>'Ar), 3.6–3.4 (m, 10H, 3 × CH₂CO + 2 × NCH₂CN), 3.4–3.3 (m, 4H, NCH₂CH₂N), 1.11 (s, 3H, CH₃). ¹³C NMR (125 MHz, 25 °C, D₂O), δ (ppm): 174.7 + 172.7 (3 × COOH), 155.7 (C^{Ar}), 133.1 (CH^{Ar}), 132.4 (CH^{Ar}), 120.9 (CH^{Ar}), 116.1 (CH^{Ar}), 115.6 (C^{Ar}), 62.3 (NCH₂CN), 58.9 + 58.3 + 58.1 (3 × NCH₂CO), 57.3 (CH₂Ar), 53.5 (NCH₂CN), 49.8 (NCH₂CH₂N), 17.7 (CH₃). Elemental analysis: found C, 54.96; H, 6.77; N, 10.09; calc. for C₁₉H₂₇N₃O₇ C, 55.74; H, 6.65; N, 10.26. ESI⁺ MS: *m/z* 410.6 [M + H⁺] (calc. for [C₁₉H₂₈N₃O₇]⁺: 410.45 g mol⁻¹).

Thermodynamic studies

The chemicals used for the experiments were of the highest analytical grade. The concentration of the GdCl₃ and EuCl₃ solutions was determined by complexometric titration with standardized Na₂H₂EDTA and xylenol orange as indicators. The concentration of the H_4AAZ3A -endoHB and H_4AAZ3A -exoHB was determined by pH-potentiometric titration in the presence and absence of a large (40-fold) excess of CaCl₂. The pH-potentiometric titrations were made with standardized 0.2 M NaOH.

The stability and protonation constants of the Gd(III)-complexes of AAZ3A-endoHB and AAZ3A-exoHB ligands were determined by pH-potentiometric titration. The metal-to-ligand concentration ratio was 1:1 (the concentration of the ligand was generally 0.002 M). For the pH measurements and titrations, a Metrohm 888 Titrando automatic titration workstation with a Metrohm-6.0234.110 combined electrode was used. Equilibrium measurements were carried out at a constant ionic strength (0.15 M NaCl) in 6 mL samples at 25 °C under magnetic stirring and N₂ bubbling. The titrations were made in the pH range 1.7-12.0. KH-phthalate (pH = 4.005) and borax (pH = 9.177) buffers were used to calibrate the pH meter. For the calculation of [H⁺] from the measured pH values, the method proposed by Irving et al. was used.44 A 0.01 M HCl solution was titrated with the standardized NaOH solution in the presence of 0.15 M NaCl ionic strength. The differences (A) between the measured (pH_{read}) and calculated $pH(-log[H^+])$ values were used to obtain the equilibrium H⁺ concentration from the pH values measured in the titration experiments (A =0.035). The waiting time between two pH measurements was 60 s. For the equilibrium calculations, the stoichiometric water ionic product (pKw) was also needed to calculate [OH⁻] values under basic conditions. The V_{NaOH}-pH_{read} data pairs of the HCl-NaOH titration obtained in the pH range 10.5-12.0 were used to calculate the pK_w value ($pK_w = 13.75$).

The protonation constants of the phenolate group of both ligands and those of Gd(m)-complexes were also determined by spectrophotometry at the absorption band of the aromatic group. The absorption spectra of 0.1 mM solutions of AAZ3A-endoHB, Gd(AAZ3A-endoHB) and Gd(AAZ3A-exoHB) were recorded in the pH range 2–12 and in the wavelength range of 210–350 nm in 0.15 M NaCl. The pH was adjusted by stepwise addition of concentrated NaOH or HCl. The spectrophotometric measurements were made with the use of PerkinElmer Lambda 365 UV-Vis spectrophotometer at 25 °C, using 1.0 cm cells. The protonation and stability constants were calculated with the PSEQUAD program.⁴⁵

Transmetallation reactions with Cu(II)

The kinetic inertness of Gd(AAZ3A-endoHB) and Gd(AAZ3A-exoHB) was determined by the rates of the Cu(π) mediated metal exchange reactions at different pH values. The exchange reactions with Cu(π) were studied by spectrophotometry, following the formation of the CuL complexes at 275 nm with PerkinElmer Lambda 365 UV-Vis spectrophotometer. The concentration of the Gd(AAZ3A-endoHB) complex was 0.2 mM, while the concentration of Cu(π) was 10 to 40 times higher, in order to guarantee pseudo-first-order conditions. The temperature was maintained at 25 °C and the ionic strength of the solutions was kept constant using 0.15 M NaCl. The exchange rates were studied in the pH range about 3.0–5.0. For keeping

the pH values constant, 1,4-dimethylpiperazine (pH range 3.0–4.1) and *N*-methylpiperazine (pH range 4.1–5.2) buffers (0.01 M) were used. The pseudo-first-order rate constants (k_d) were calculated by fitting the absorbance data to eqn (5)

$$A_t = (A_0 - A_p)e^{-k_{\rm d}t} + A_p \tag{5}$$

where A_t , A_0 and A_p are the absorbance values at time t, the start of the reaction and at equilibrium, respectively. The calculation of the kinetic parameters were performed by the fitting of the absorbance-time and relaxation rate-time data pairs with the Micromath Scientist computer program (version 2.0, Salt Lake City, UT, USA).

Relaxometric measurements

The Gd(III) complexes were prepared by mixing solutions of GdCl₃ hexahydrate and the ligand (in *ca*. 5% molar excess) and adjusting the pH to about 6.5 with aqueous NaOH. The exact concentration of the aqueous solutions for ¹H NMRD and ¹⁷O NMR measurements was determined by measuring the bulk magnetic susceptibility shifts of the tert-BuOH ¹H NMR signal at 11.7 T.⁴⁶ Proton relaxation measurements $(1/T_1)$ and the resulting $1/T_1$ NMRD profiles were measured on a Fast-Field Cycling (FFC) Stelar SmarTracer Relaxometer (Stelar s.r.l., Mede (PV), Italy) over a continuum of magnetic field strengths from 0.00024 to 0.25 T (corresponding to 0.01-10 MHz proton Larmor frequencies). The relaxometer operates under computer control with an absolute uncertainty in $1/T_1$ of $\pm 1\%$. A precise control of the temperature was operated during the measurements by means of a Stelar VTC-91 airflow heater equipped with a calibrated copper constantan thermocouple (uncertainty of ±0.1 °C). Furthermore, the real temperature inside the probe head was additionally monitored by a Fluke 52 k/j digital thermometer (Fluke, Zürich, Switzerland). Additional data in the 20-120 MHz frequency range were obtained with a High Field Relaxometer (Stelar) equipped with the HTS-110 3T Metrology Cryogen-free Superconducting Magnet. The data were collected using the standard inversion recovery sequence (20 experiments, 2 scans) with a typical 90° pulse width of 3.5 ms and the reproducibility of the data was within ±0.5%. ¹⁷O NMR spectra were recorded on a Bruker Avance III spectrometer (11.7 T) equipped with a 5 mm probe and a standard temperature control unit. Aqueous solutions of the complexes (ca. 6-10 mM) containing 2.0% of the ¹⁷O isotope (Cambridge Isotope Laboratories Inc., Tewksbury, MA, USA) were used. The observed transverse relaxation rates were calculated from the signal width at half-height as a function of temperature in the 278-350 K range.

Theoretical calculations

Optimizations of the $[Gd(L)(H_2O)_2]^-\cdot 4H_2O$ systems (L = AAZ3AendoHB or AAZ3A-exoHB) were performed with the Gaussian 16 program package (revision B.01), using the pseudopotential approximation and the wB97XD functional,²⁷ which includes empirical dispersion. We selected the quasirelativistic largecore pseudopotential of the Stuttgart/Cologne group (46 + 4f7

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electrons in the core for Gd) and the related [5s4p3d] valence basis set,²⁸ while the standard 6-311G(d,p) basis set was used for all other atoms. No restraints were imposed for geometry optimization and frequency calculations were used to confirm the nature of the optimized geometries as local energy minima. The output of frequency calculations provided the thermal contributions and zero-point energies required to calculate the enthalpy and entropy of the systems. Bulk solvent effects (water) were considered with integral equation formalism of the polarized continuum model (IEFPCM), using the default parameters implemented in Gaussian 16.⁴⁷ The size of the integration grid was increased with the integral = ultrafine keyword.

Scalar relativistic calculations were performed with the ORCA program package³⁴ using the second-order Douglas-Kroll-Hess method.^{36,48} Solvent effects (water) were included with the SMD solvation model.49 DFT calculations used the TPSSh functional,³⁵ in combination with the SARC2-DKH-OZVP³⁷ basis set for Gd and the DKH-def2-TZVPP^{50,51} basis set for all other atoms. The resolution of identity and chain of spheres RIJCOSX approximation⁵² was used to speed up the calculations using the Autoaux⁵³ procedure to generate auxiliary basis sets for Gd and the SARC/J basis set for other atoms. The quasi-restricted orbitals generated from these calculations were used as starting orbitals for CASSCF/ NEVPT2^{54,55} calculations, in which the active space included 7 electrons distributed in the seven 4f-based orbitals (1 octet root and 49 sextets were considered). The RIJCOSX approximation⁵⁶ was used also in these calculations, with the SARC2-DKH-QZVP/JK³⁷ auxiliary basis set for Gd and the Autoaux⁵³ procedure to generate auxiliary basis sets for all other atoms. Spin-orbit coupling effects were included by using quasidegenerate perturbation theory (QDPT).57 All ORCA calculations used a Gaussian finite nucleus model.⁵⁸

Conflicts of interest

There are no conflicts to declare.

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