

Electronic Supplementary Material

for

Supramolecular assemblies based on amphiphilic Mn²⁺-Complexes as High Relaxivity MRI probes

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1. Dynamic Light scattering measurements:

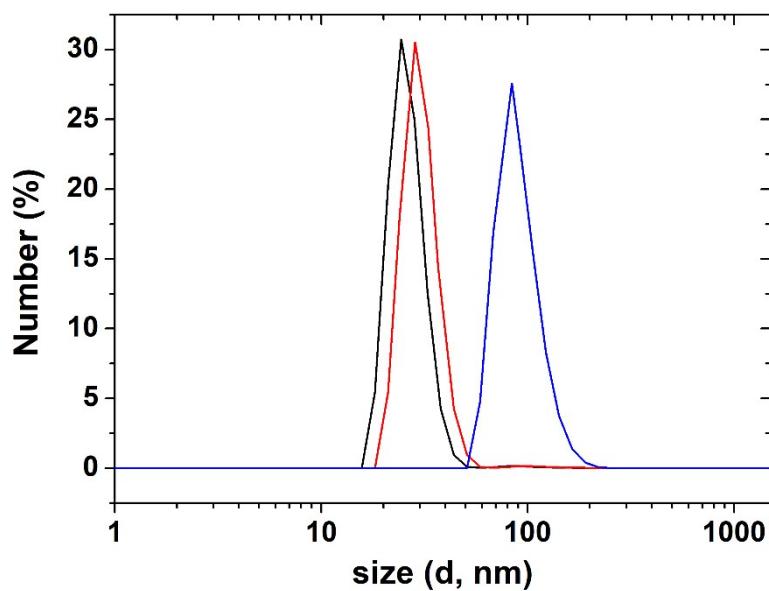


Figure S1: DLS analysis of **MnTDDTA** (black), **MnODDTA** (red) and **MnHCDTA** (blue) in aggregated forms: **MnTDDTA**: 26.2 nm (Pdl = 0.26); **MnODDTA**: 28.8 nm (Pdl = 0.22); **MnHCDTA**: 83.8 nm (Pdl = 0.33).

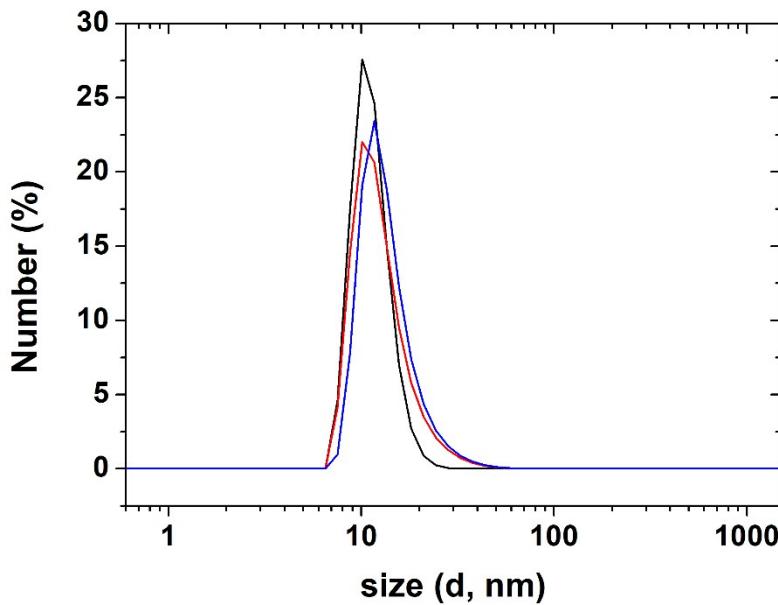


Figure S2: DLS analysis of mixed micelles made of 50% of DSPE-PEG₂₀₀₀-OMe and **MnHCDTA** (blue), **MnDD-DO2A** (red) and **MnDH-DO2A** (black): **MnHCDTA**: 13.9 nm (Pdl = 0.26); **MnDD-DO2A**: 13.0 nm (Pdl = 0.26); **MnDH-DO2A**: 11.4 nm (Pdl = 0.38).

2. Relaxometry

Table S1. Parameters obtained from the simultaneous analysis of ^{17}O NMR and NMRD data for MnTDDTA (in the aggregated form) and MnDB-DO2A. In case of MnODDTA (in the aggregated form) and of mixed micelles made by DSPE-PEG2000 and MnHCDTA, MnDD-DO2A or MnDH-DO2A at 1:1 molar ratio the best-fit was carried out by analysis of the ^1H NMRD profiles at 298 K.

Parameter	Mn-TDDTA	Mn-ODDTA	Mn-HCDTA	MnDB-DO2A	MnDD-DO2A	MnDH-DO2A
$A^2 / 10^{19} \text{ s}^{-2}$	10.7 ± 0.7	8.7 ± 0.7	1.8 ± 0.1	28 ± 10	3.5 ± 0.8	1.6 ± 0.8
$^{298}\tau_V / \text{ps}$	23 ± 2	26 ± 2	33 ± 2	10 ± 3	36 ± 8	36 ± 5
$k_{\text{ex}}^{298} / 10^6 \text{ s}^{-1}$	310 ± 15	$310^{[a]}$	$310^{[a]}$	1650 ± 50	340 ± 30	340 ± 30
$\Delta H^\ddagger / \text{kJ mol}^{-1}$	29.5 ± 2.4	--	--	36.8 ± 1.3	--	--
$^{298}\tau_{\text{RG}} / \text{ns}$	0.117 ± 0.003	0.170 ± 0.005	3.2 ± 0.3	0.038 ± 0.006	9 ± 2	9 ± 2
$^{298}\tau_{\text{RL}} / \text{ps}$	--	--	102 ± 12	--	78 ± 8	81 ± 5
S^2	--	--	0.38 ± 0.01	--	0.25 ± 0.01	0.28 ± 0.02
$E_R / \text{kJ mol}^{-1}$	18 ± 1	--	--	17 ± 2	--	--
$E_v / \text{kJ mol}^{-1}^{[a]}$	1	--	--	1	--	--
q	$1^{[a]}$	$1^{[a]}$	$1^{[a]}$	0.87 ± 0.01	$0.87^{[a]}$	$0.87^{[a]}$
$r_{\text{MnH}} / \text{\AA}^{[a]}$	2.83	2.83	2.83	2.83	2.83	2.83
$^{298}D / 10^{-5} \text{ cm}^2 \text{ s}^{-1}^{[a]}$	2.3	2.3	2.3	2.3	2.3	2.3
$a / \text{\AA}^{[a]}$	3.6	3.6	3.6	3.6	3.6	3.6

[a] fixed in the fitting procedure.

2.1 Variable temperature experiments

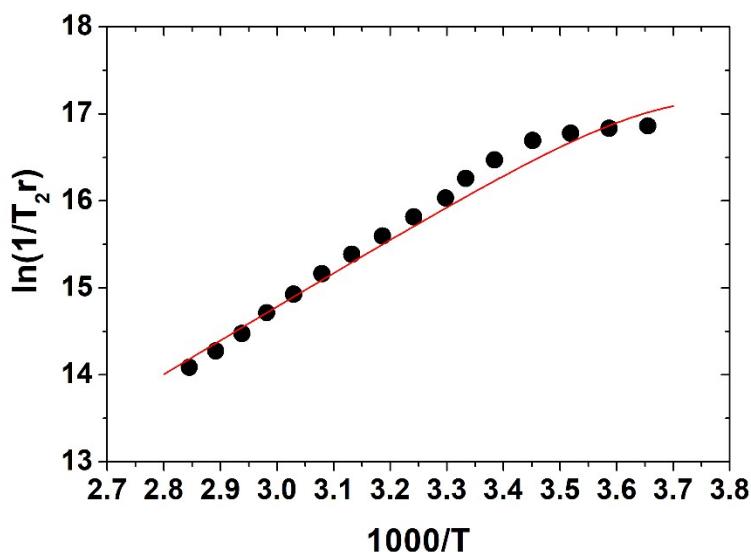


Figure S3. Reduced transverse ^{17}O relaxation rates for a 4.02 mM solution of MnTDDTA measured at 11.74 T and pH 6.5

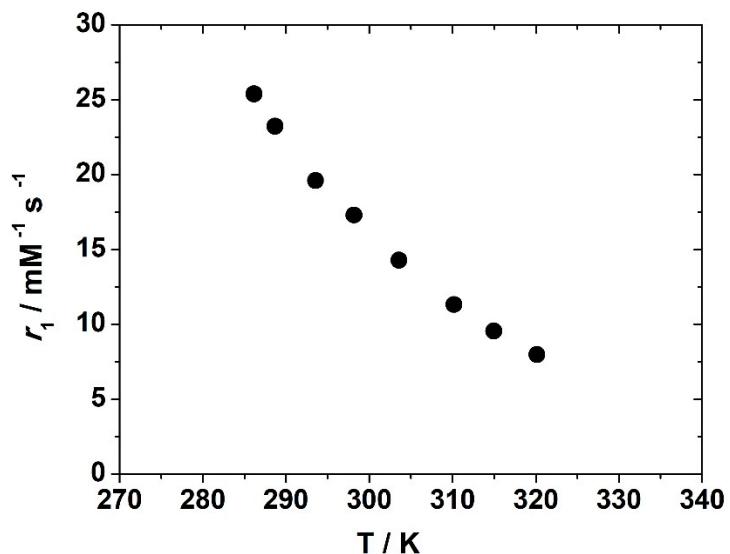


Figure S4. Temperature dependence of the longitudinal water proton relaxivity for Mn-HCDTA mixed micelle at 20 MHz and pH = 7.

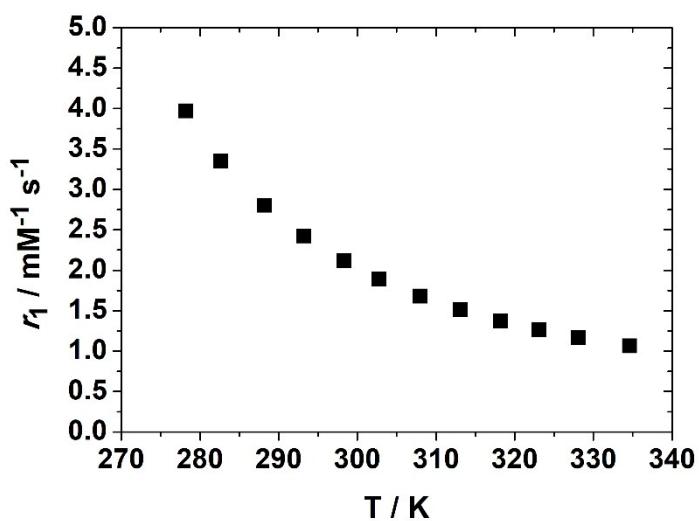


Figure S5. Temperature dependence of the longitudinal water proton relaxivity for Mn-DB-DO2A at 20 MHz and pH = 7.

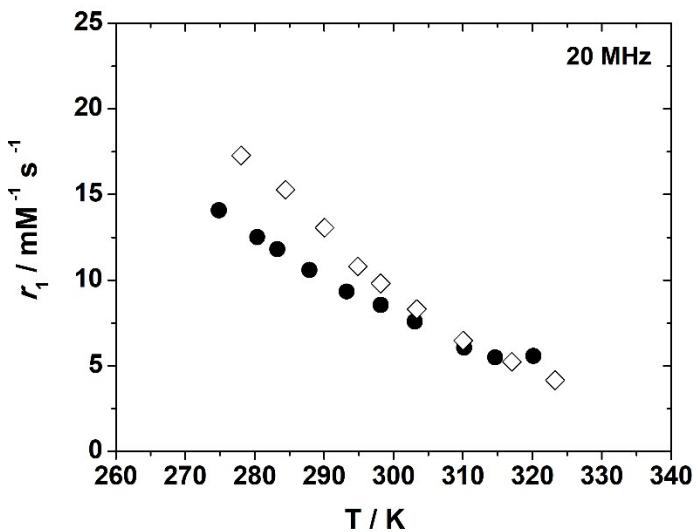


Figure S6. Temperature dependence of the longitudinal water proton relaxivity for Mn-**DD-DO2A** (empty diamonds) and Mn-**DH-DO2A** (black circles) mixed micelles at 20 MHz and pH = 7.

2.2 NMRD profiles

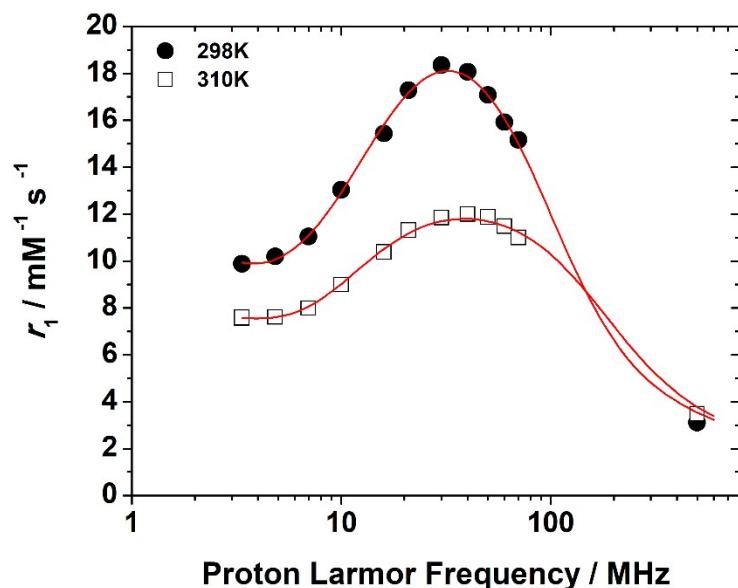


Figure S7. ^1H NMRD profiles of Mn-**HCDTA** mixed micelles at 298 (black circles) and 310 K (open squares).

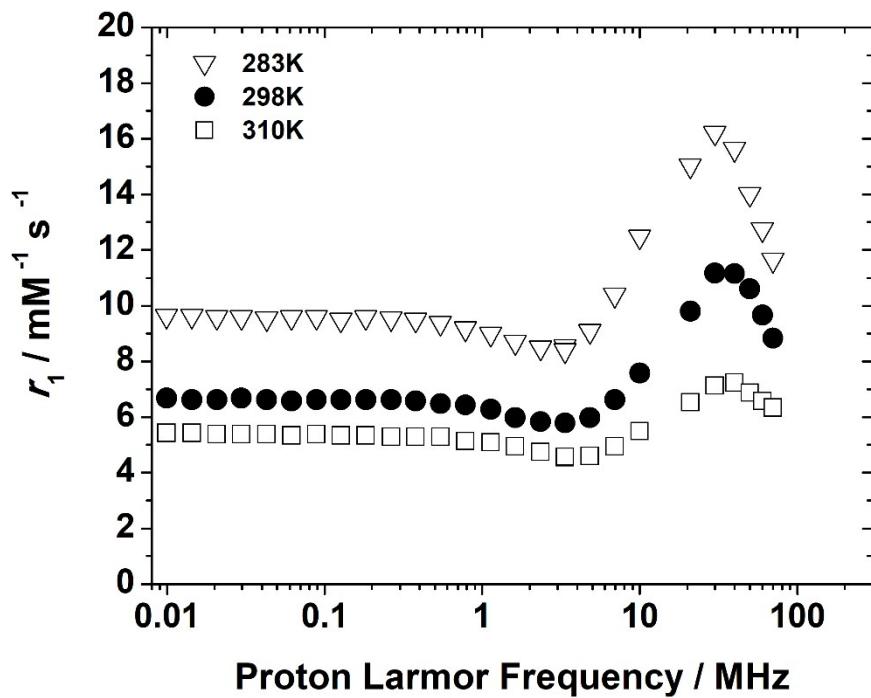


Figure S8. ^1H NMRD profiles of Mn-DD-DO2A at 283 (open triangles), 298 (black circles) and 310 K (open squares).

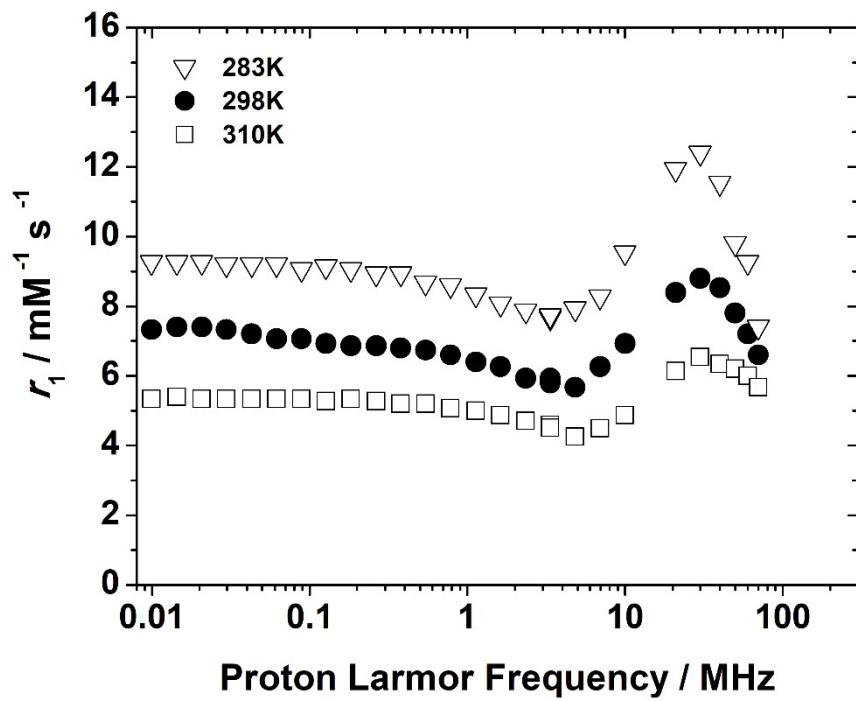


Figure S9. ^1H NMRD profiles of Mn-DH-DO2A at 283 (open triangles), 298 (black circles) and 310 K (open squares).

2.3 Human serum albumin (HSA) interaction

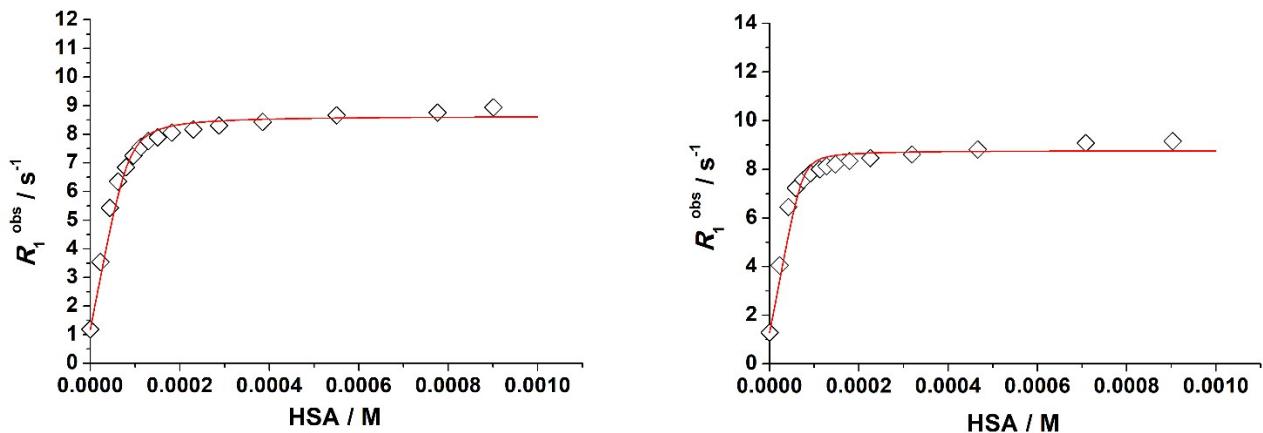


Figure S10. Water proton relaxation rate of an aqueous solution of 0.17 mM of Mn-TDDTA (left) and 0.15 mM of Mn-ODDTA (right) as a function of increasing amounts of HSA (20 MHz, 298 K, pH 7.0).

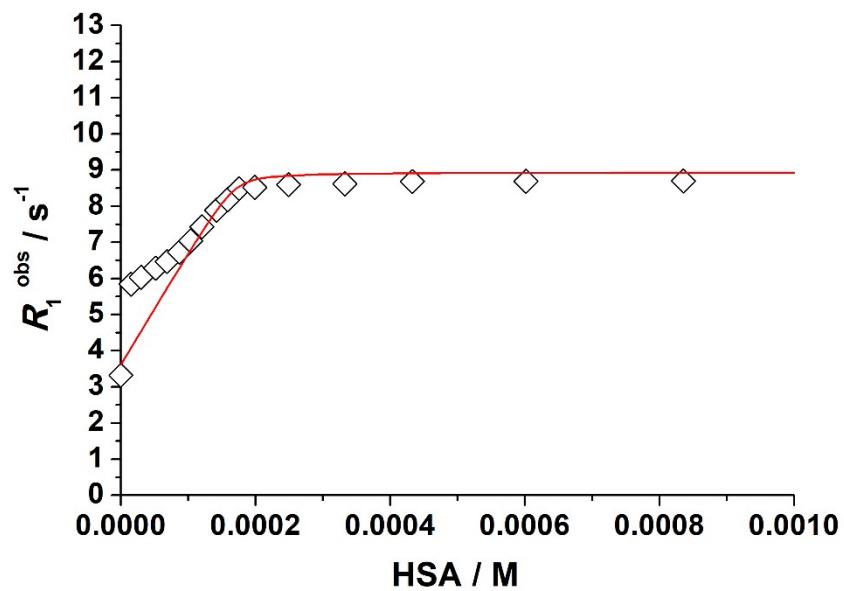


Figure S11. Water proton relaxation rate of an aqueous solution of 0.17 mM of Mn-HCDTA as a function of increasing amounts of HSA (20 MHz, 298 K, pH 7.0).

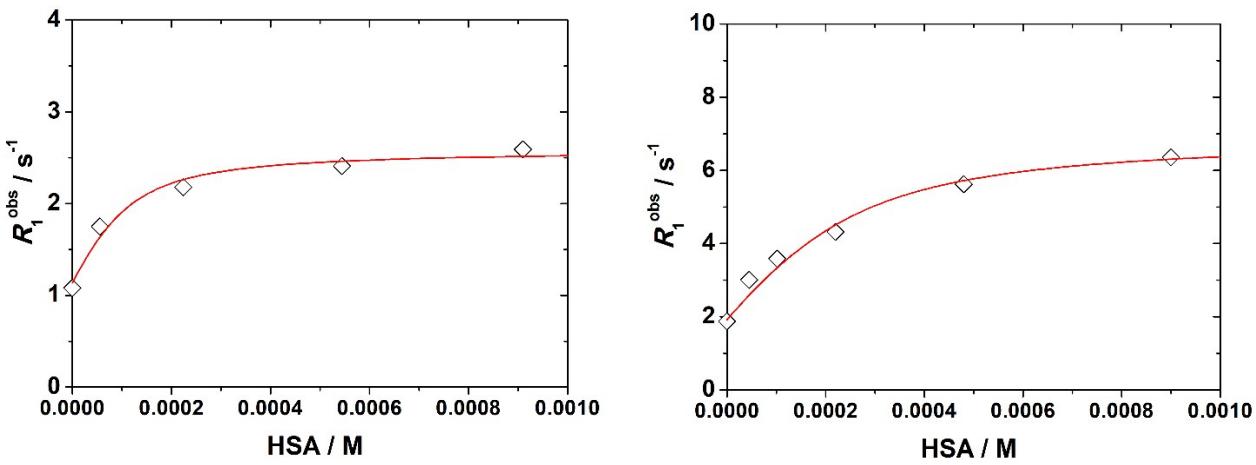


Figure S12. Water proton relaxation rate of an aqueous solution of 0.095 mM of **Mn-DD-DO2A** (left) and 0.22 mM of **Mn-DH-DO2A** (right) as a function of increasing amounts of HSA (20 MHz, 298 K, pH 7.2). Each titration point was left stirring 24 h to allow complete formation of the adducts.

2.4 NMRD profiles of Mn-complexes-HSA adducts

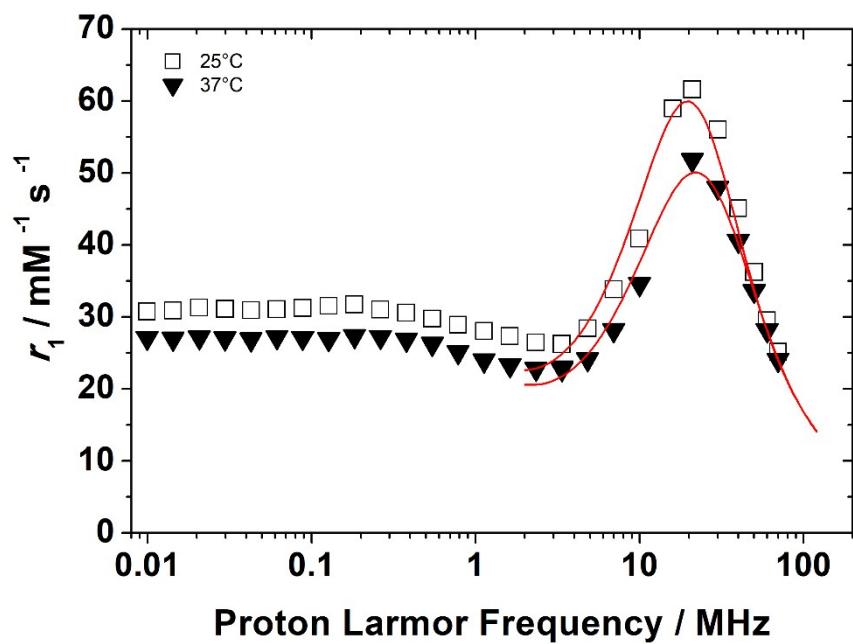


Figure S13. NMRD profiles of **Mn-TDDTA-HSA** adduct at 298 (empty squares) and 310 K (black triangles).

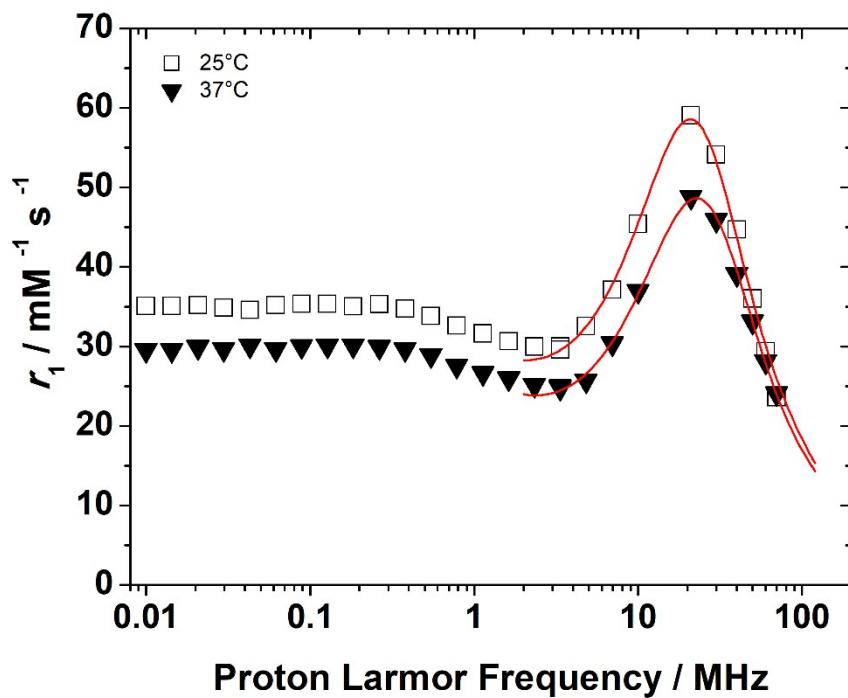


Figure S14. NMRD profiles of Mn-ODDTA-HSA adduct at 298 (empty squares) and 310 K (black triangles).

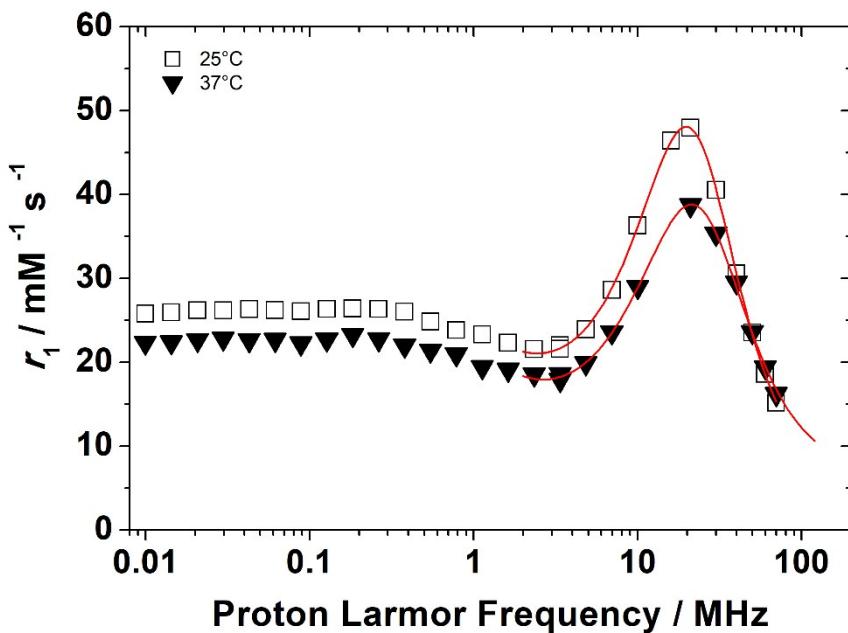


Figure S15. NMRD profiles of Mn-HCDTA-HSA adduct at 298 (empty squares) and 310 K (black triangles).

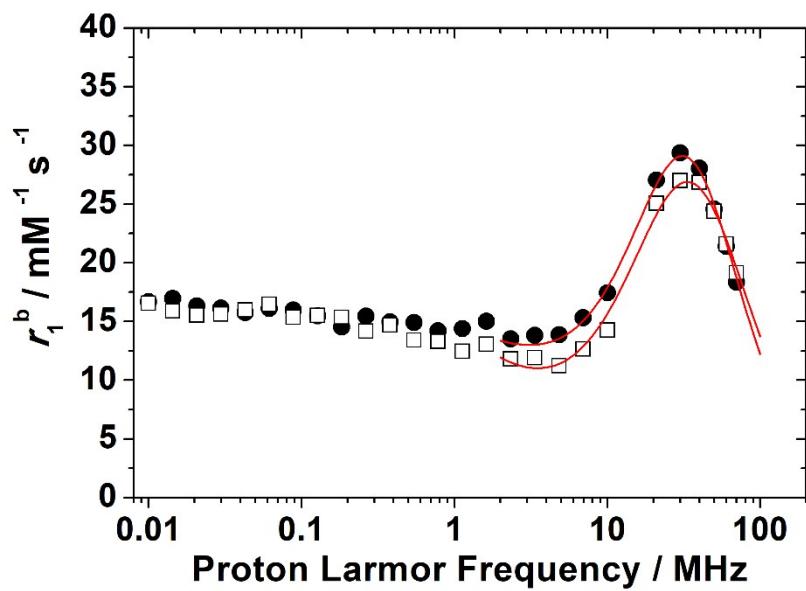


Figure S16. NMRD profiles of Mn-DD-DO2A–HSA adduct at 298 (black circles) and 310 K (empty squares).

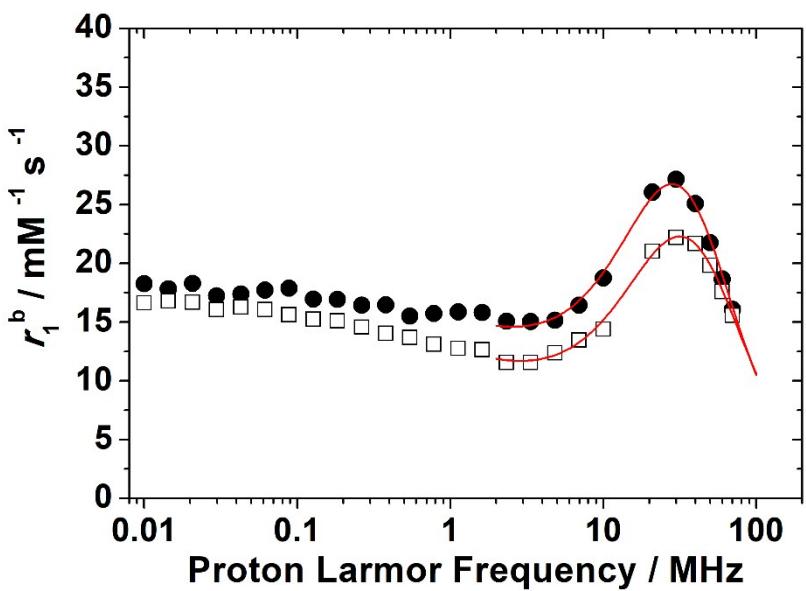


Figure S17. NMRD profiles of Mn-DH-DO2A–HSA adduct at 298 (black circles) and 310 K (empty squares).