

Supporting information

for

[Re(CO)₃]⁺ complexes of exo-functionalized tridentate “click”

macrocycles: Synthesis, stability, photophysical properties, bio-

conjugation and antibacterial activity

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1 Selected ^1H NMR and HR-ESI-MS spectra of synthesized compounds

Macrocycle, 7

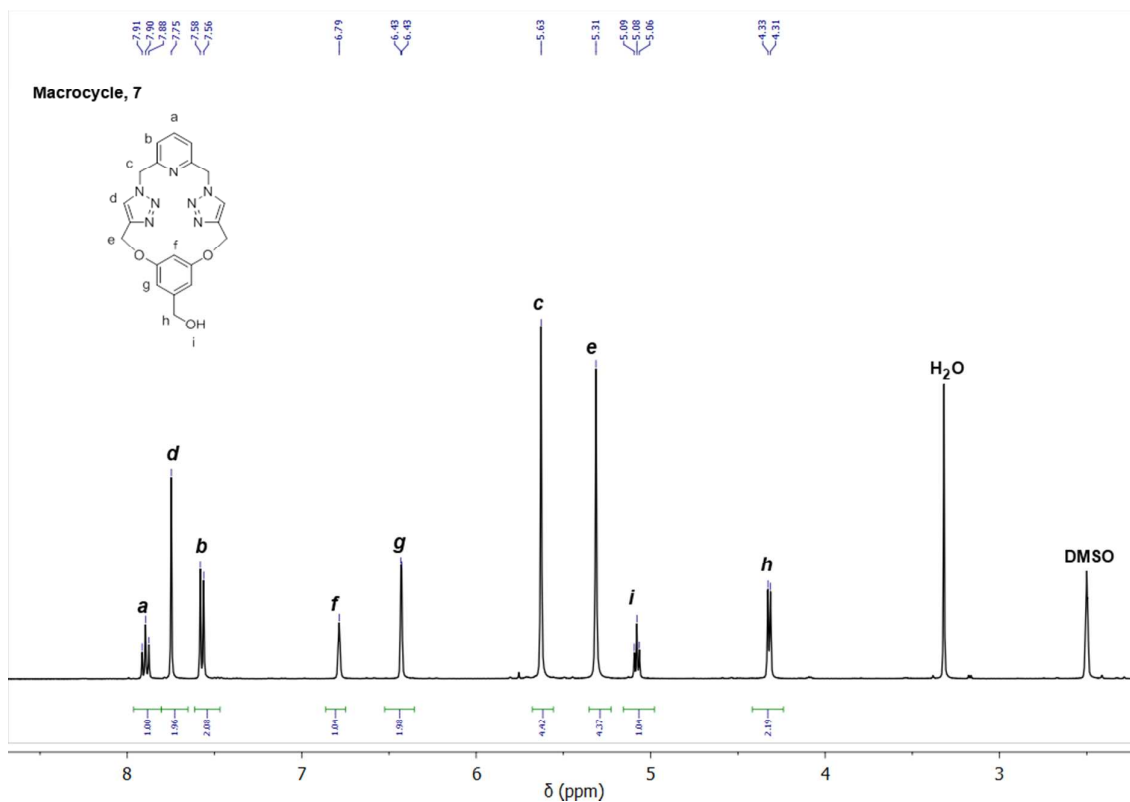


Figure S 1 ^1H NMR spectrum (400 MHz, $\text{DMSO}-d_6$, 298 K) of macrocycle 7

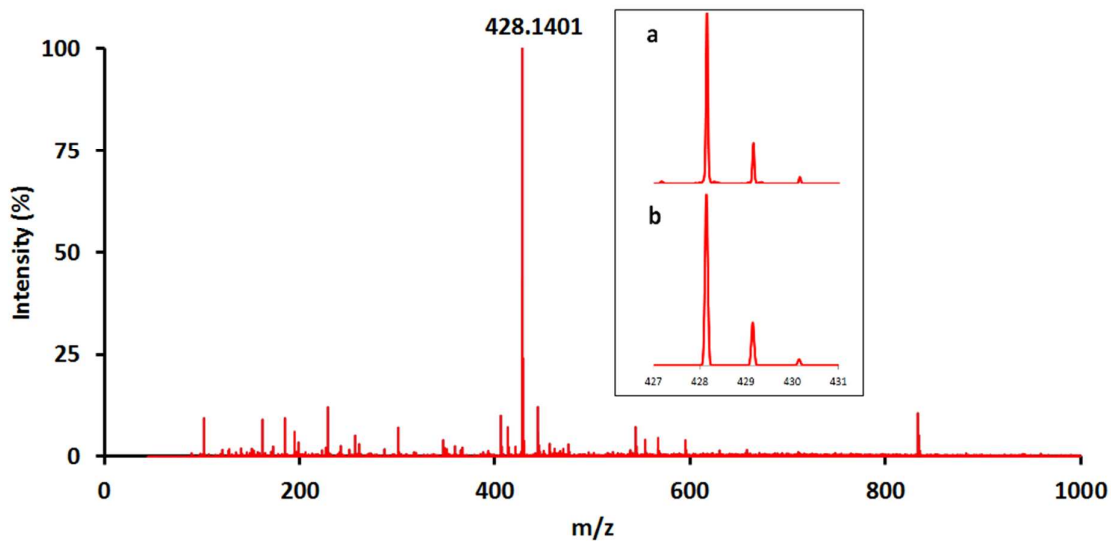


Figure S 2 HR-ESI-MS spectrum of macrocycle 7, inset a) observed and b) calculated isotopic patterns for the peaks at $m/z = 428.1404$ due to $[7+\text{Na}]^+$ ion.

Macrocycle, 8

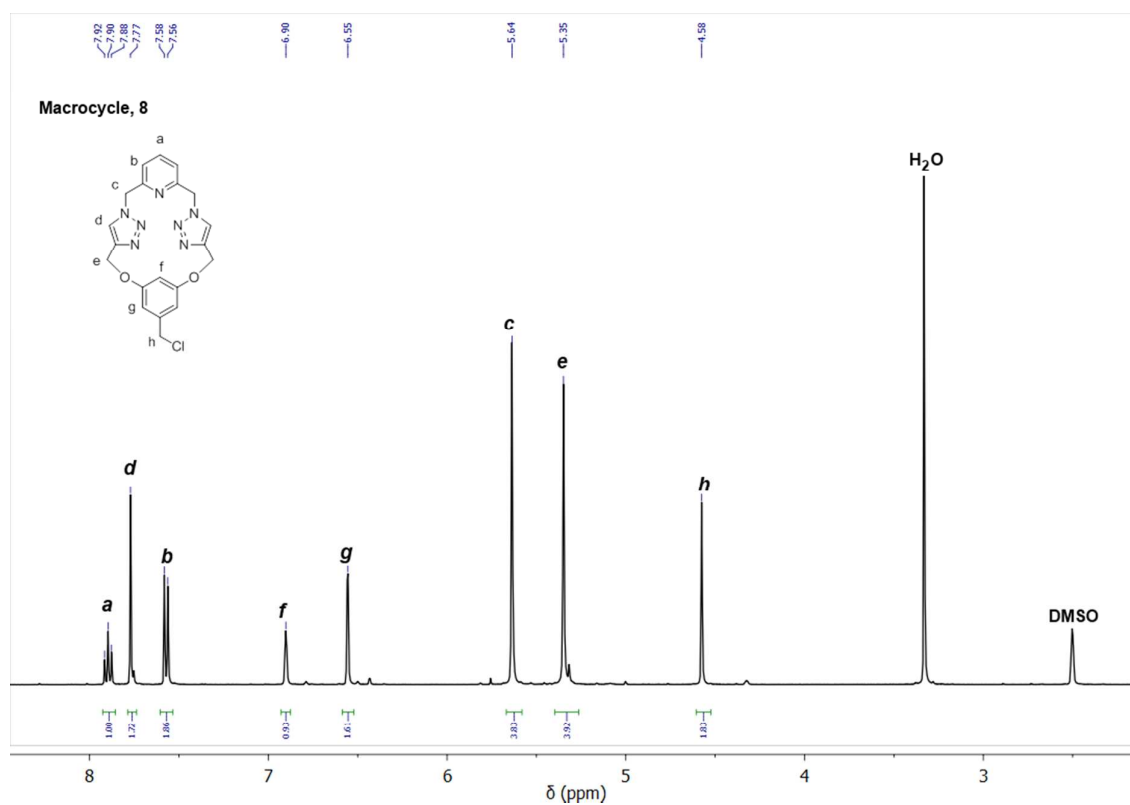


Figure S 3 ^1H NMR spectrum (400 MHz, $\text{DMSO}-d_6$, 298 K) of macrocycle **8**

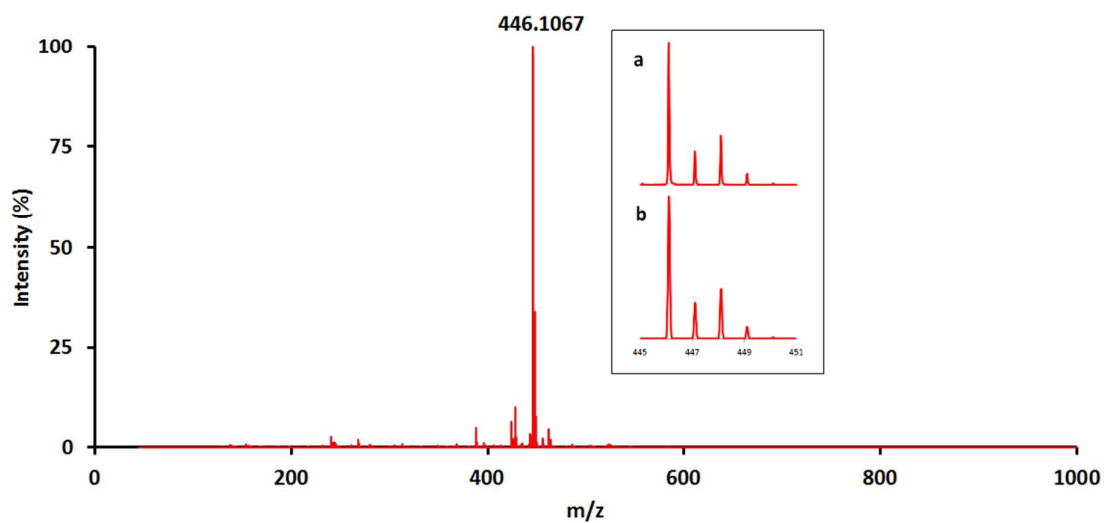


Figure S 4 HR-ESI-MS spectrum of macrocycle **8**, inset a) observed and b) calculated isotopic patterns for the peak at $m/z = 446.1067$ due to $[\mathbf{8}+\text{Na}]^+$ ion.

Macrocycle, 9

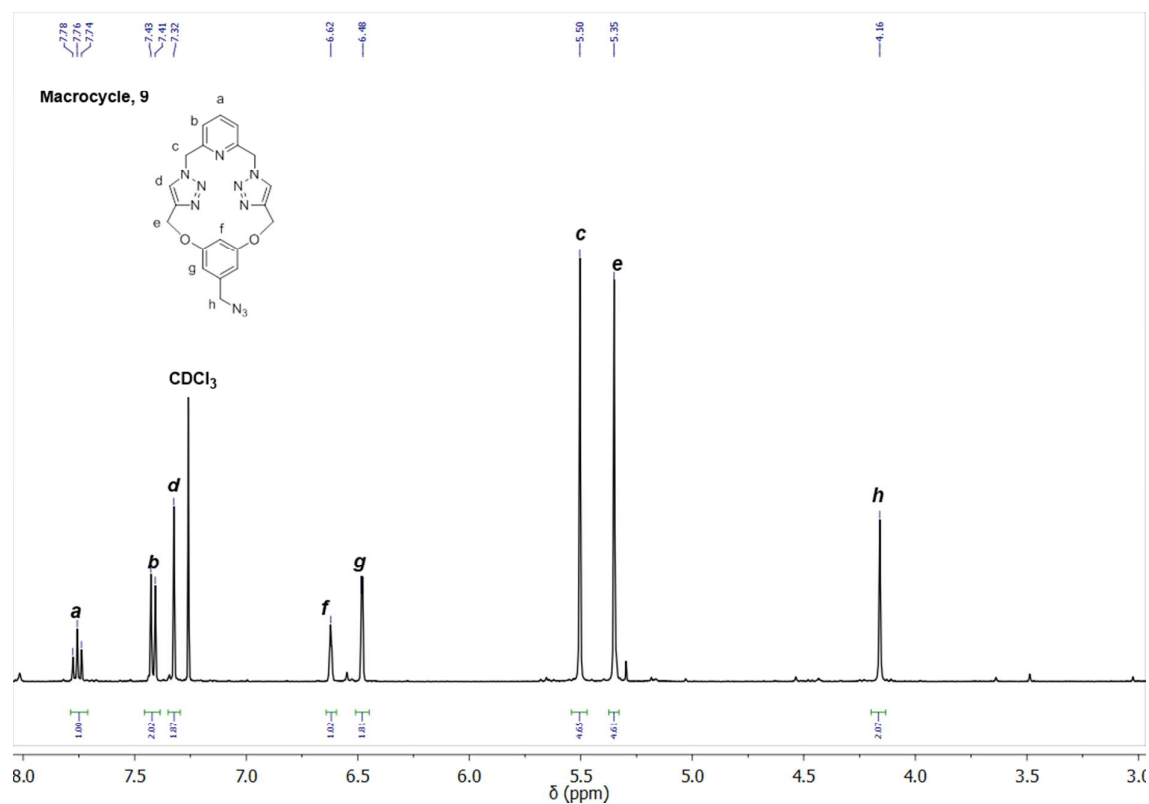


Figure S 5 ¹H NMR spectrum (400 MHz, CDCl₃, 298 K) of macrocycle 9

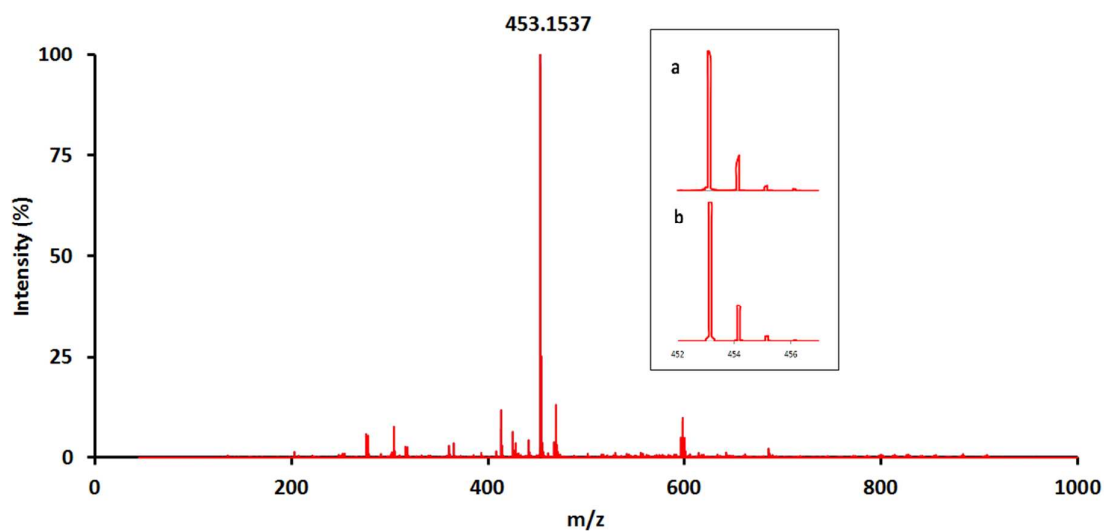


Figure S 6 HR-ESI-MS spectrum of macrocycle 9, inset a) observed and b) calculated isotopic patterns for the peak at $m/z = 453.1537$ due to $[9+Na]^+$ ion.

Acid functionalized macrocycle, 10a

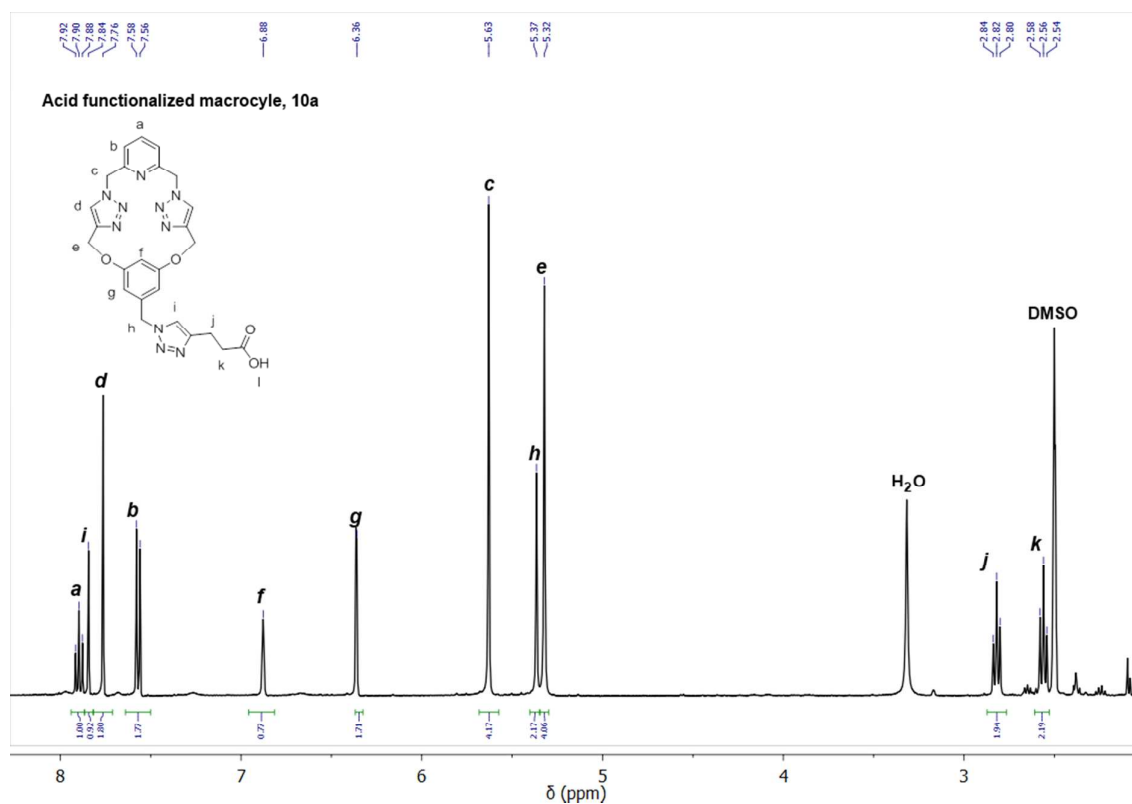


Figure S 7 ^1H NMR spectrum (400 MHz, $\text{DMSO}-d_6$, 298 K) of macrocycle **10a**

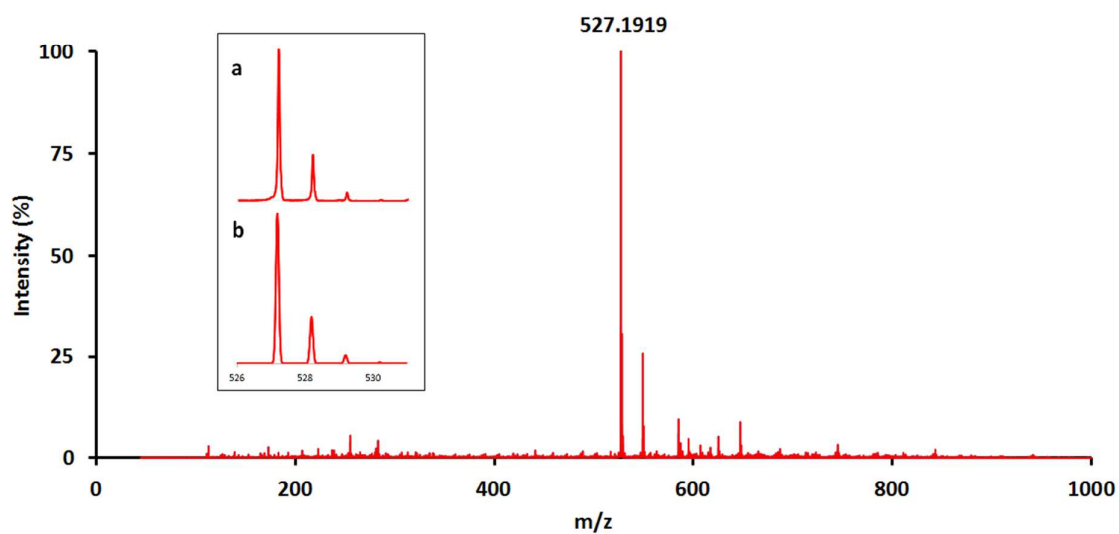


Figure S 8 HR-ESI-MS spectrum of macrocycle **10a**, inset a) observed and b) calculated isotopic patterns for the peak at $m/z = 527.1919$ due to $[\mathbf{10a-H}]^-$ ion.

NHS activated ester functionalized macrocycle, 10b

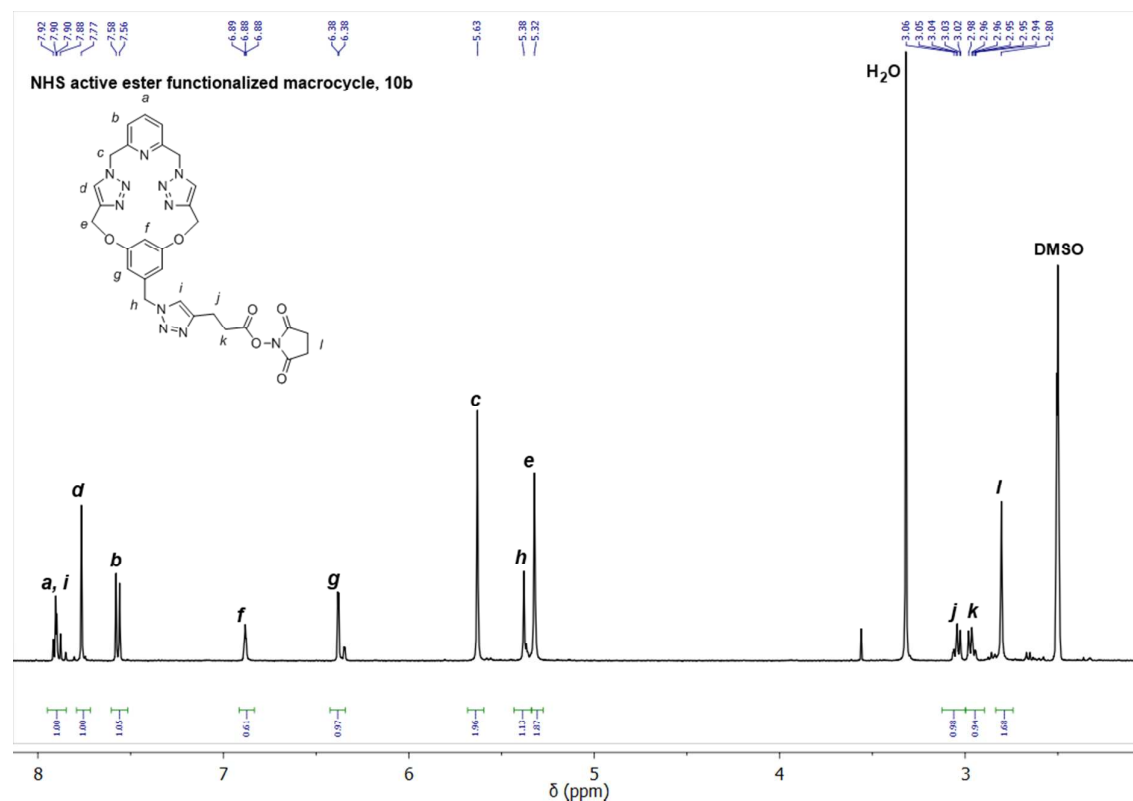


Figure S 9 ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of macrocycle **10b**.

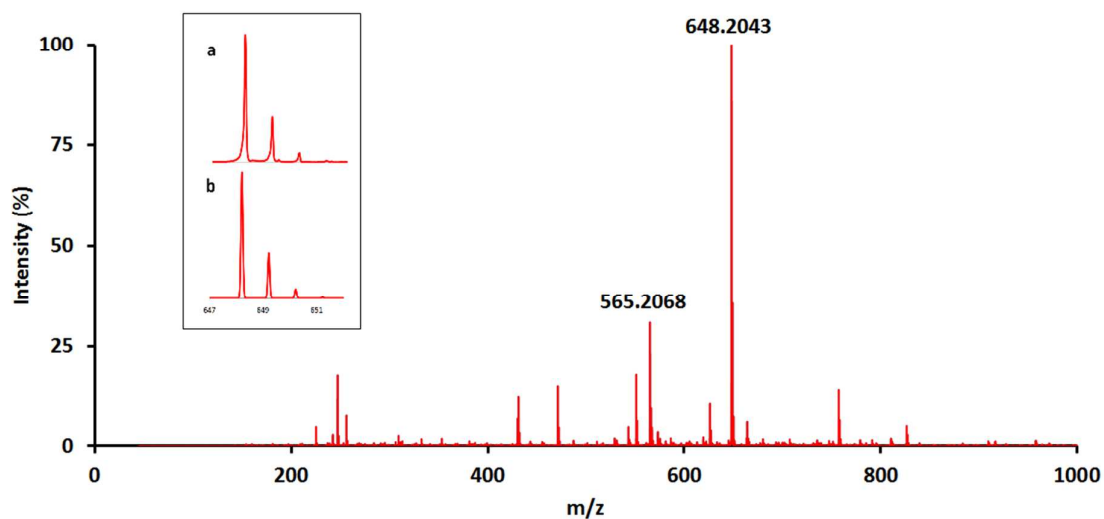


Figure S 10 HR-ESI-MS spectrum of macrocycle **10b**, inset a) observed and b) calculated isotopic patterns for the peak at m/z = 648.2043 due to [**10b**+Na]⁺ ion, also showing peak due to [**10b**-NHS+CH₃+Na]⁺ at m/z = 565.2068.

Gluco-conjugated macrocycle 10c

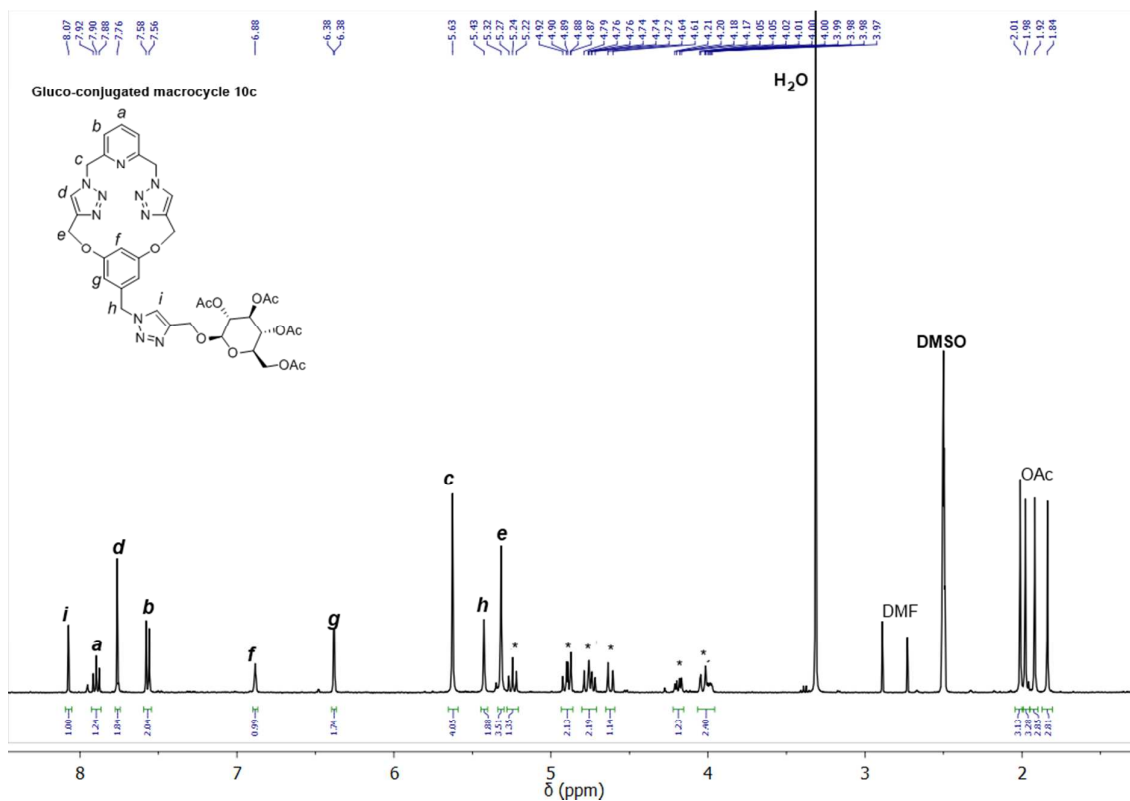


Figure S 11 ^1H NMR spectrum (400 MHz, $\text{DMSO-}d_6$, 298 K) of macrocycle **10c** (* peaks from sugar molecule).

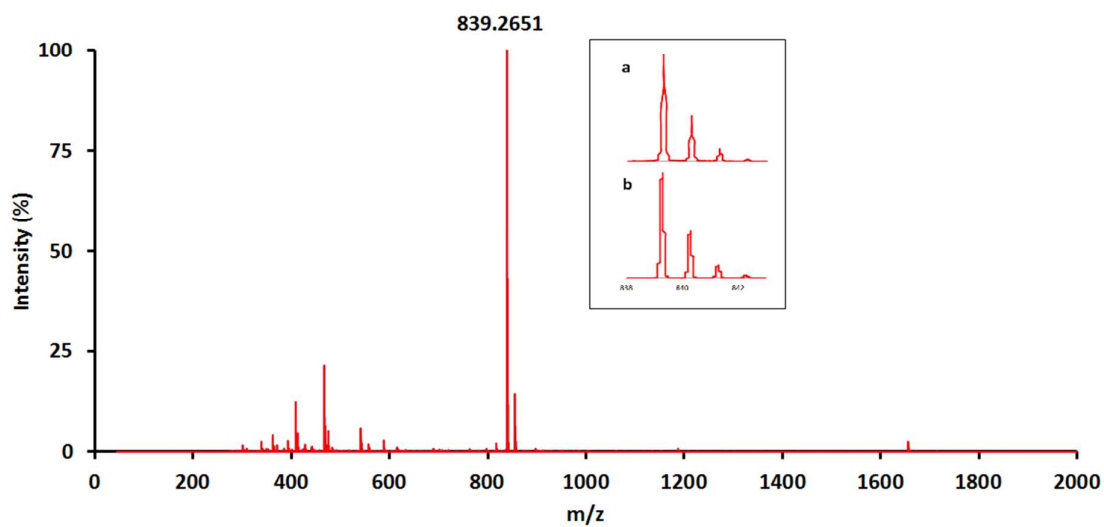


Figure S 12 HR-ESI-MS spectrum of macrocycle **10c**, inset a) observed and b) calculated isotopic patterns for the peak at $m/z = 839.2651$ due to $[\mathbf{10c}+\text{Na}]^+$ ion.

Gluco-conjugated macrocycle 10d

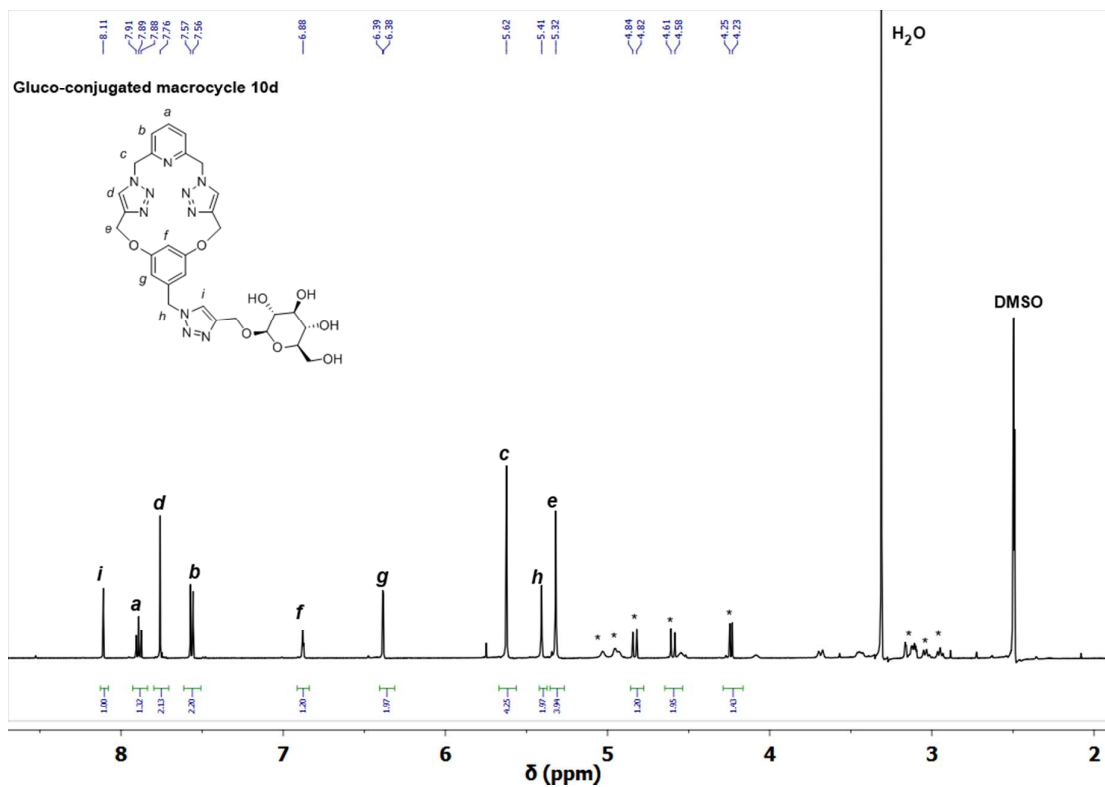


Figure S 13 ¹H NMR spectrum (500 MHz, DMSO-*d*₆, 298 K) of macrocycle **10d** (* peaks from sugar molecule).

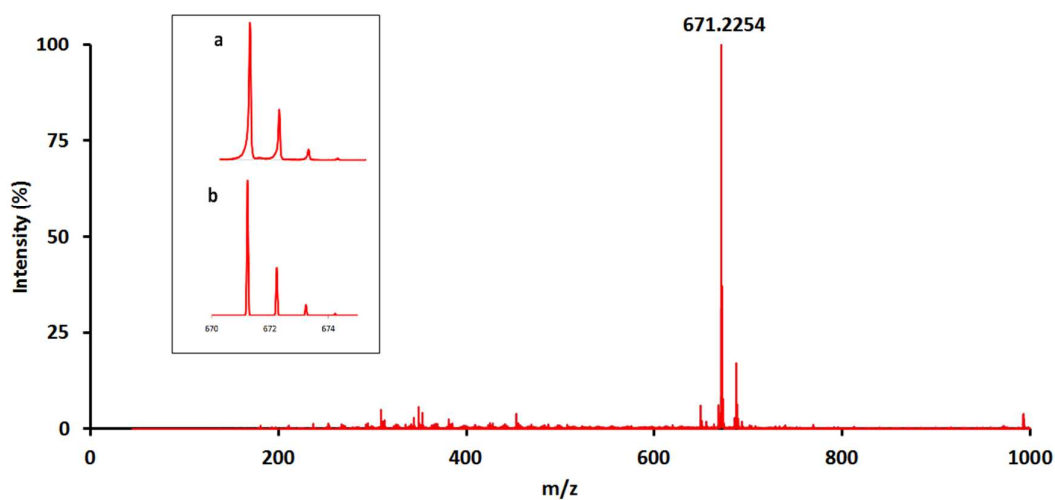


Figure S 14 HR-ESI-MS spectrum of macrocycle **10d**, inset a) observed and b) calculated isotopic patterns for the peak at m/z = 671.2254 due to $[\mathbf{10d}+\text{Na}]^+$ ion.

Steroid-conjugated macrocycle 10e

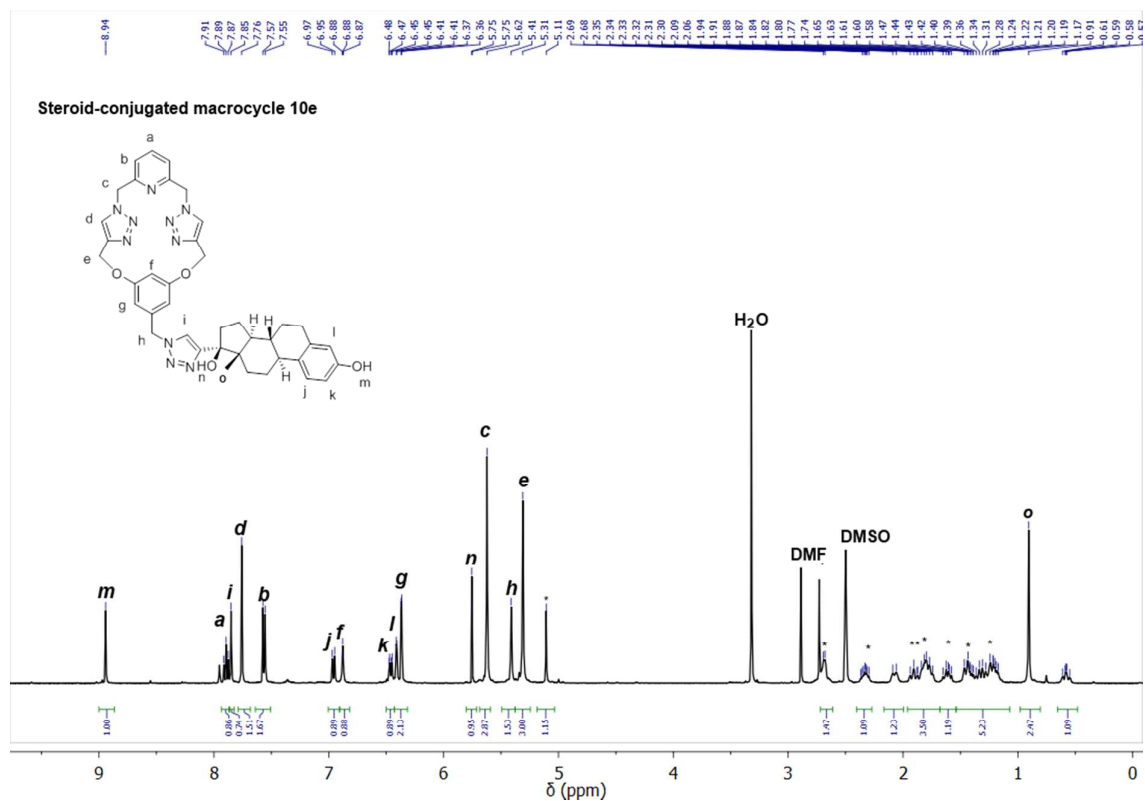


Figure S 15 ^1H NMR spectrum (400 MHz, $\text{DMSO}-d_6$, 298 K) of macrocycle **10e** (* peaks from steroid molecule).

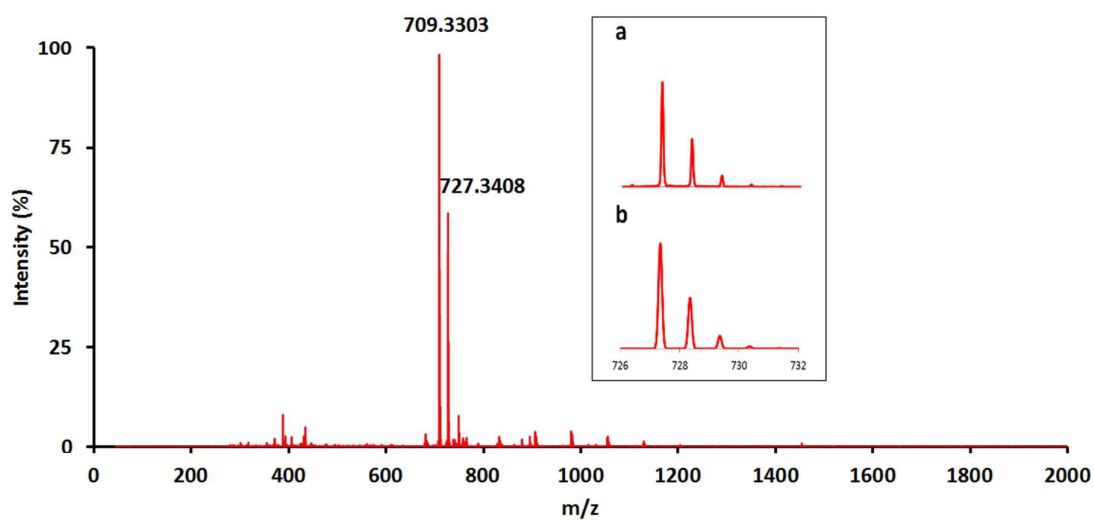


Figure S 16 HR-ESI-MS spectrum of macrocycle **10e**, inset a) observed and b) calculated isotopic patterns for the peak at $m/z = 727.3408$ due to $[\mathbf{10e}+\text{H}]^+$ ion.

[Re(CO)₃(7)]Br, (7-Re)

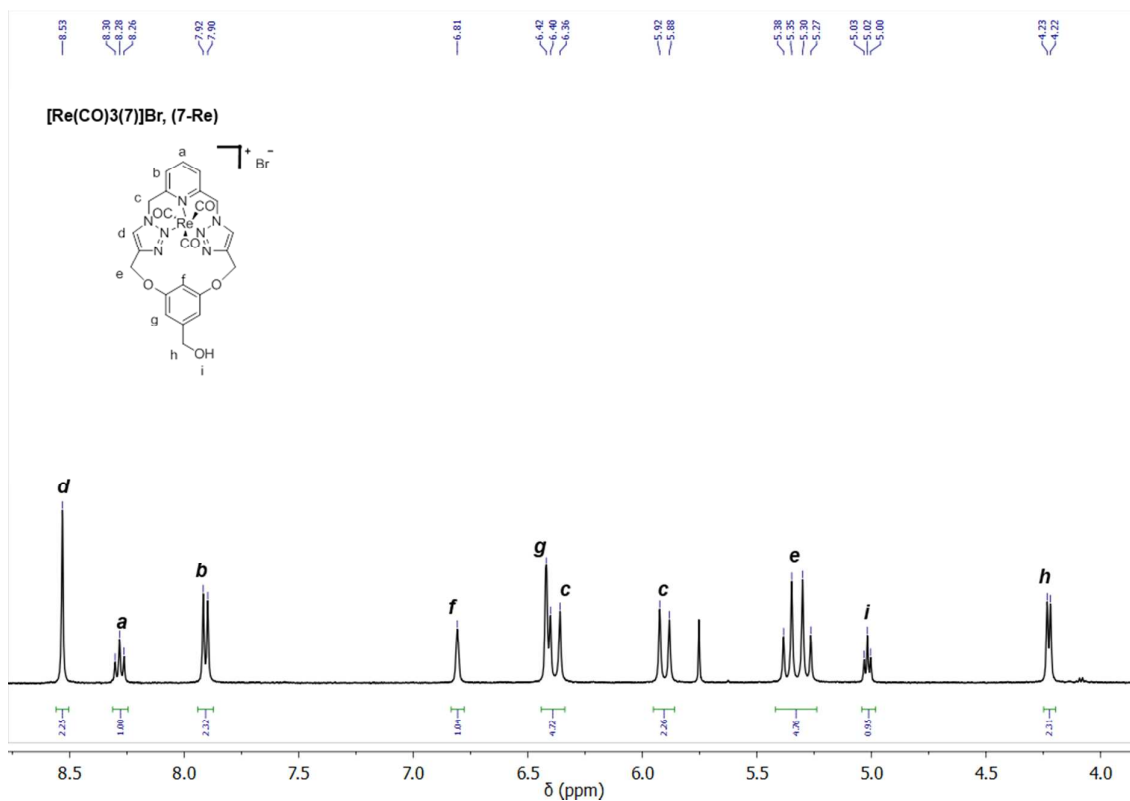


Figure S 17 ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of macrocycle **7-Re**.

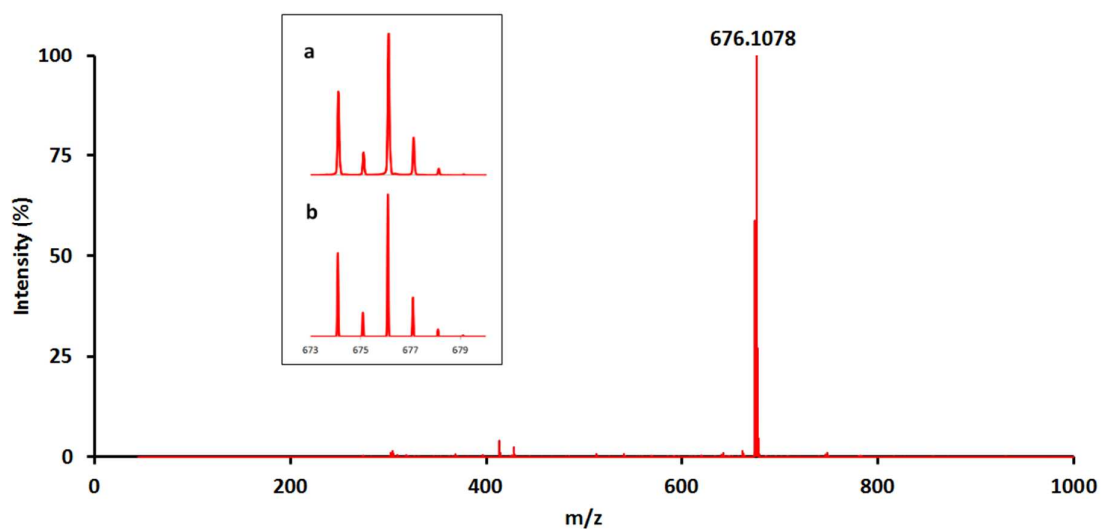


Figure S 18 HR-ESI-MS spectrum of macrocycle **7-Re**, inset a) observed and b) calculated isotopic patterns for the peak at m/z = 676.1078 due to $[\text{Re}(\text{CO})_3+7]^+$ ion.

[Re(CO)₃(10a)]Br, (Re-10a)

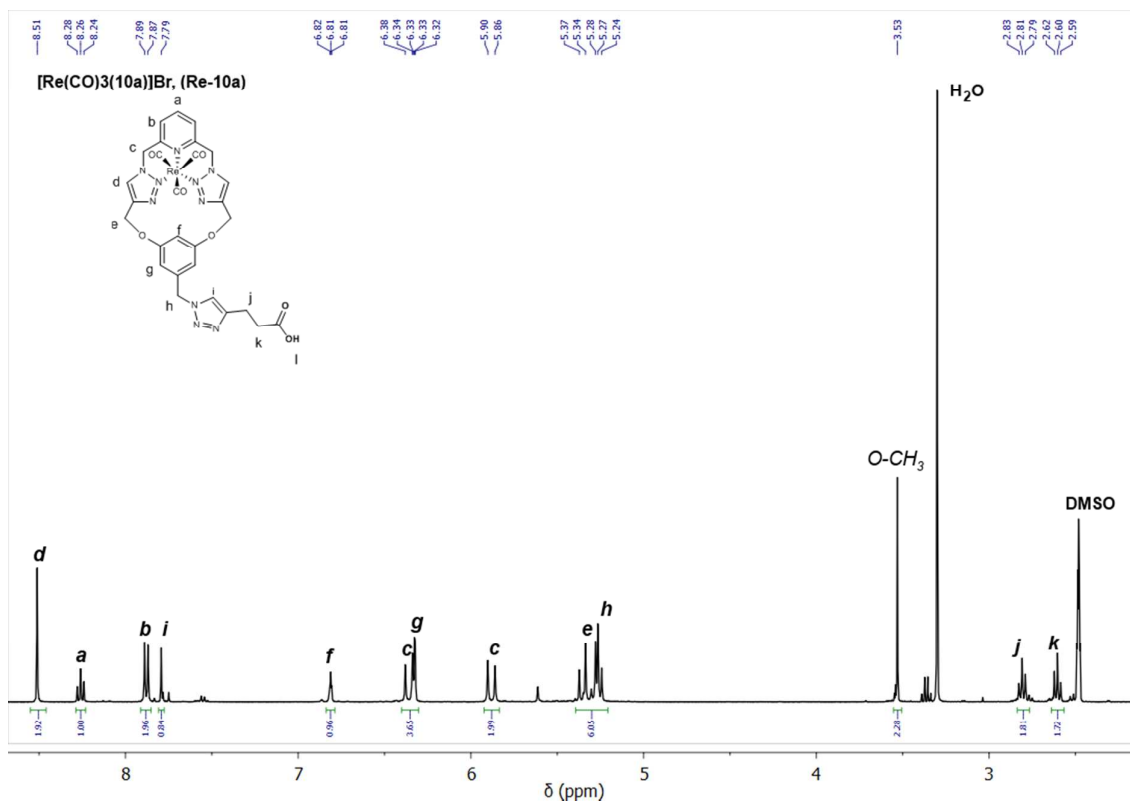


Figure S 19 ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of macrocycle **Re-10a**

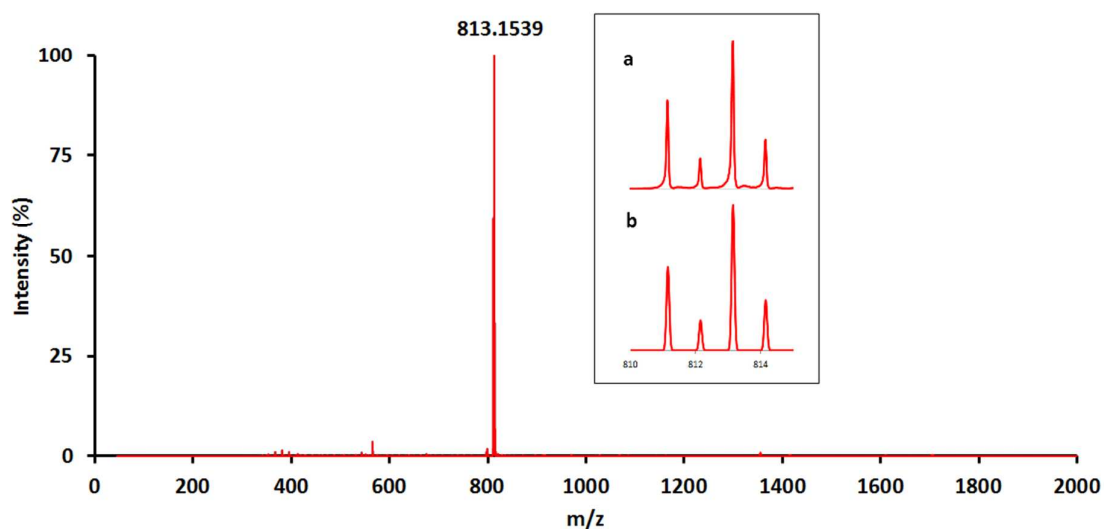


Figure S 20 HR-ESI-MS spectrum of macrocycle **Re-10a** showing peaks at $m/z = 813.1539$ due to esterified complex **Re-10a**, inset a) observed and b) calculated isotopic patterns for the peak at $m/z = 813.1539$ due to $[\text{Re}(\text{CO})_3 + 10\text{a} - (\text{H} + \text{CH}_3)]^+$ ion.

[Re(CO)₃(10b)]Br, (Re-10b)

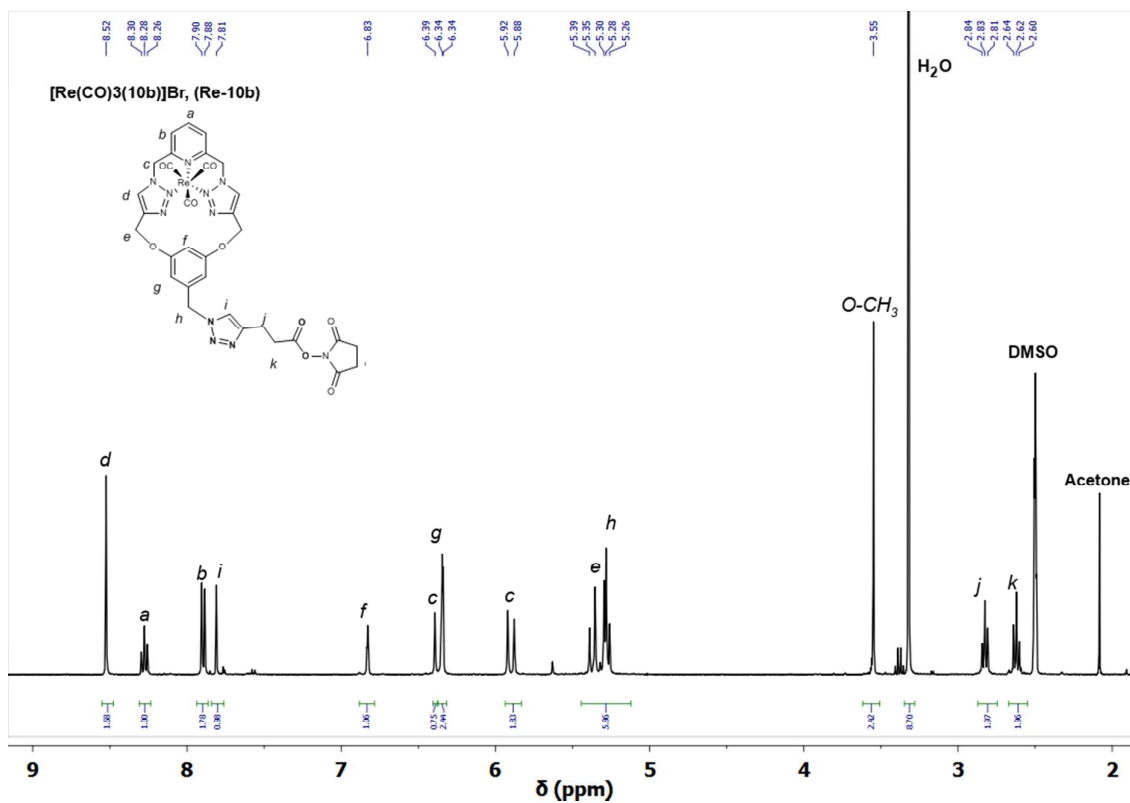


Figure S 21 ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of macrocycle **Re-10b**

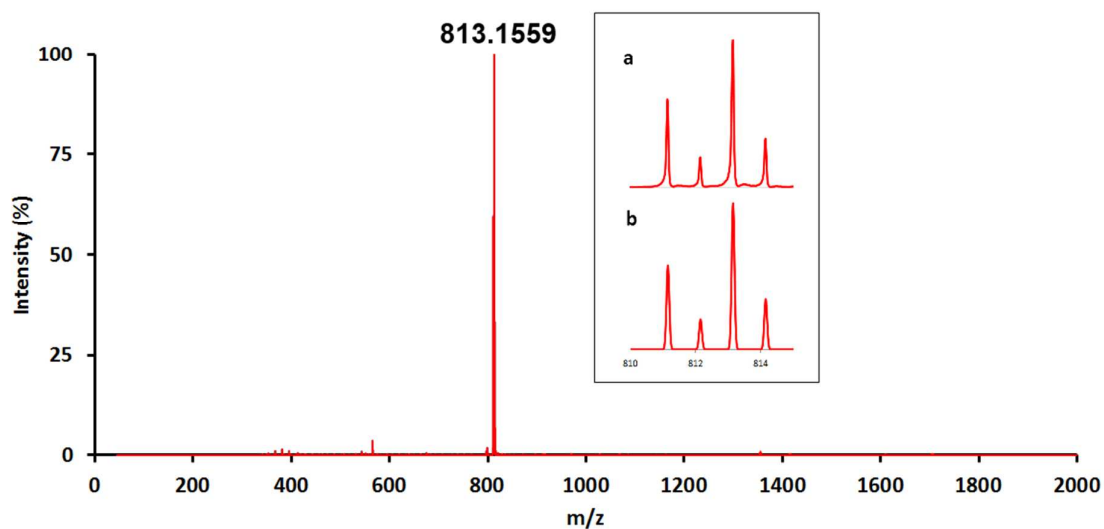


Figure S 22 HR-ESI-MS spectrum of macrocycle **Re-10b** showing peaks at *m/z* = 813.1559 due to esterified complex **Re-10b**, inset a) observed and b) calculated isotopic patterns for the peak at *m/z* = 813.1539 due to [Re(CO)₃+10a(-NHS+CH₃)]⁺ ion.

[Re(CO)₃(10c)]Br, (Re-10c)

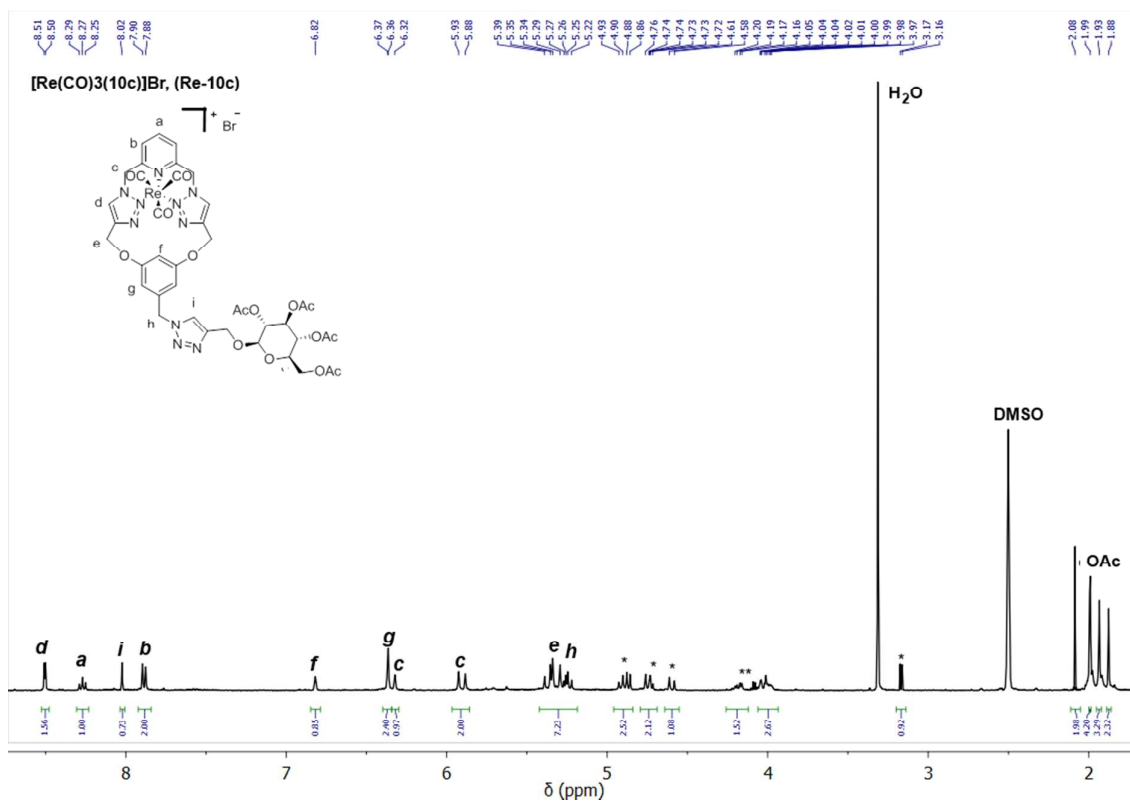


Figure S 23 ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of macrocycle **Re-10c** (* peaks from sugar molecule).

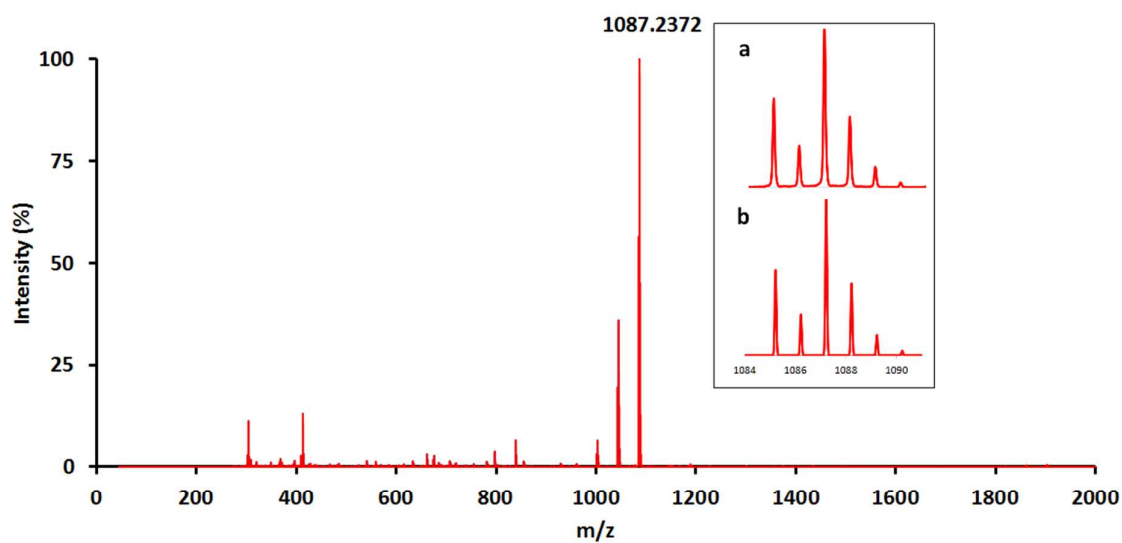


Figure S 24 HR-ESI-MS spectrum of macrocycle complex **Re-10c**, inset a) observed and b) calculated isotopic patterns for the peak at *m/z* = 1087.2372 due to [Re(CO)₃+**10c**]⁺ ion.

[Re(CO)₃(10d)]Br, (Re-10d)

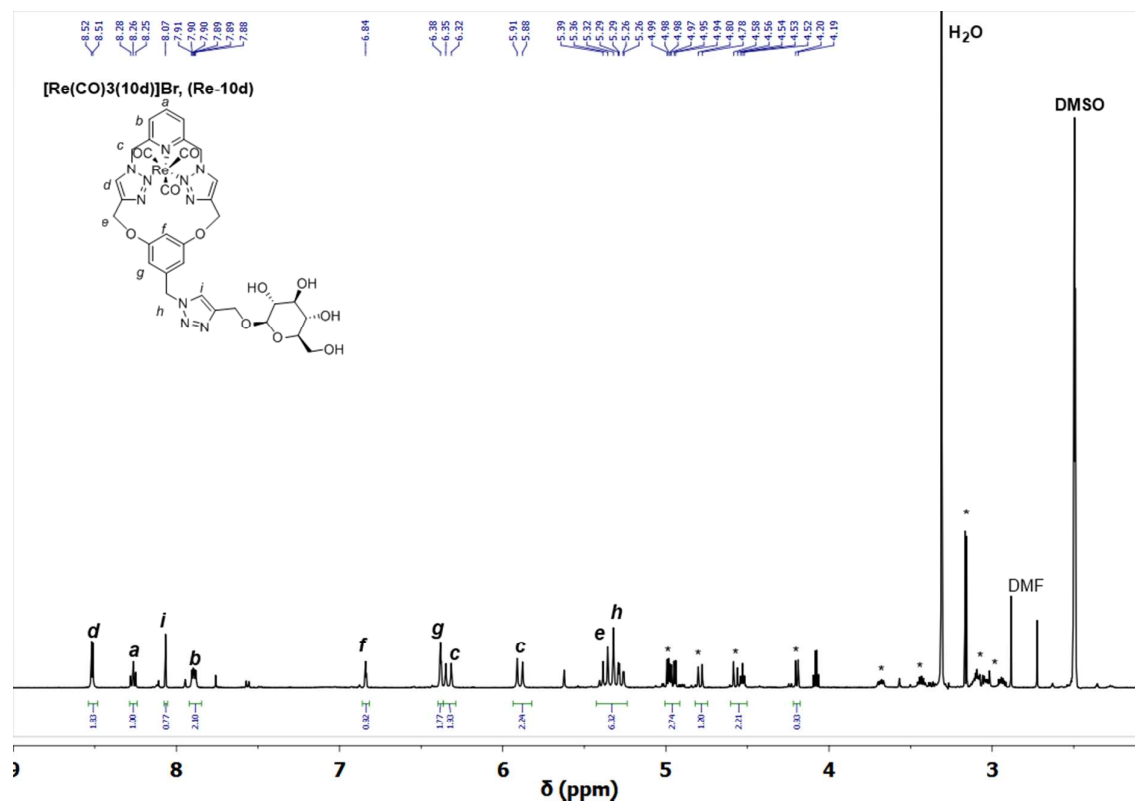


Figure S 25 ¹H NMR spectrum (500 MHz, DMSO-*d*₆, 298 K) of macrocycle **Re-10d** (* peaks from sugar molecule).

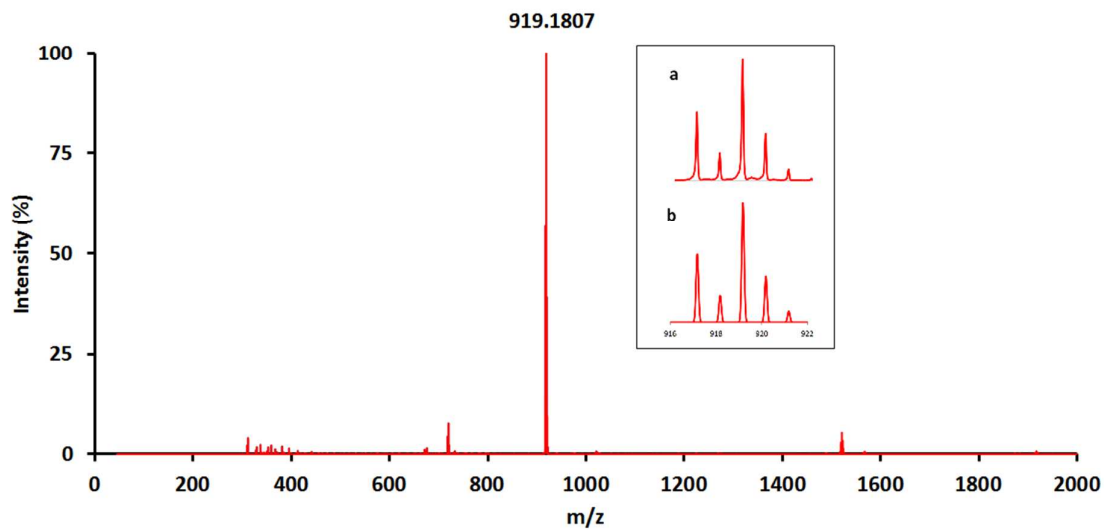


Figure S 26 HR-ESI-MS spectrum of macrocycle complex **Re-10d**, inset a) observed and b) calculated isotopic patterns for the peak at *m/z* = 919.1807 due to [Re(CO)₃+10d]⁺ ion.

[Re(CO)₃(10e)]Br, (Re-10e)

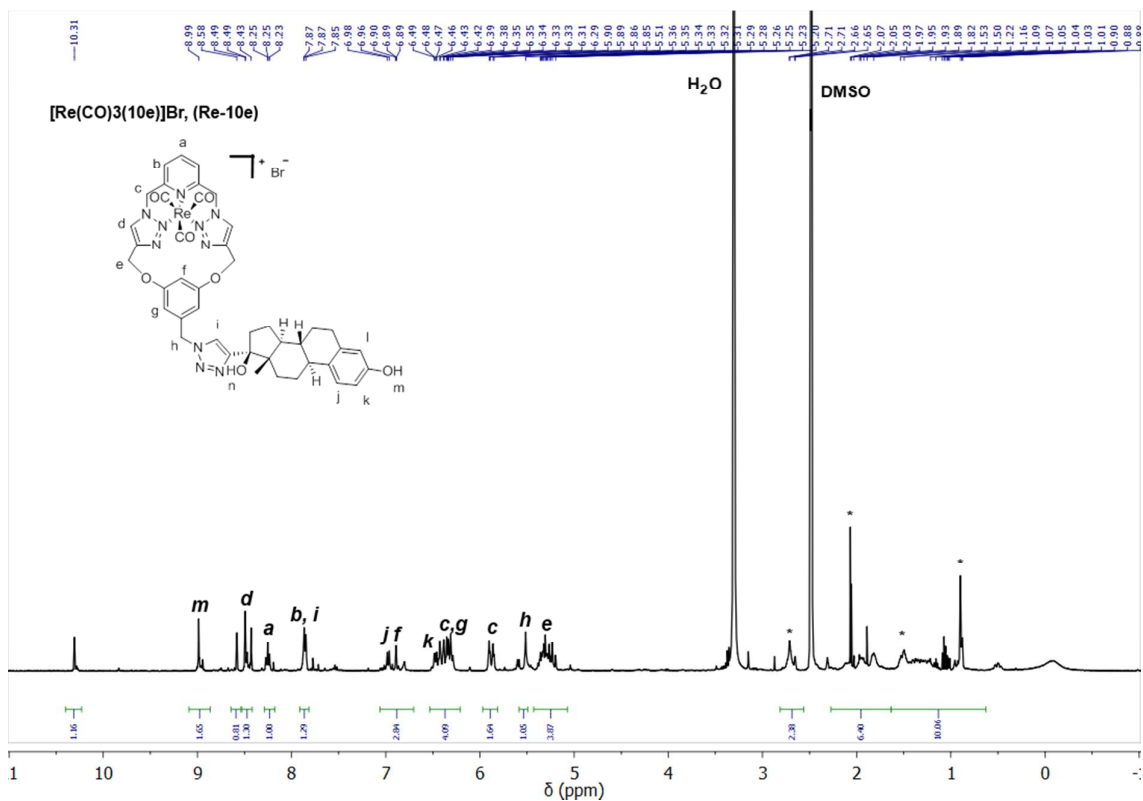


Figure S 27 ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of macrocycle **Re-10e** (* peaks from steroid molecule)

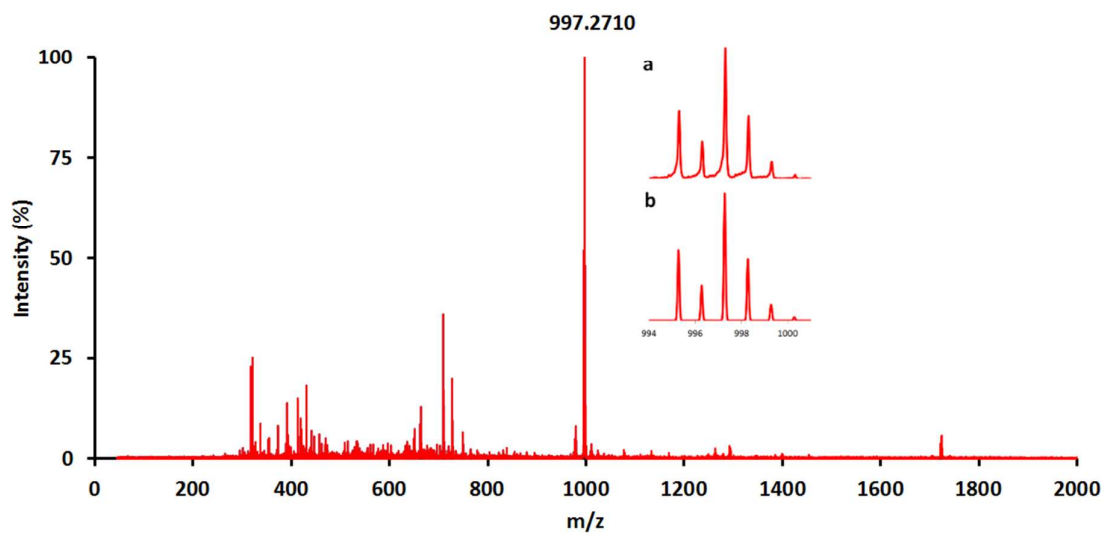


Figure S 28 HR-ESI-MS spectrum of macrocycle **Re-10e**, inset a) observed and b) calculated isotopic patterns for the peak at *m/z* = 997.2710 due to [Re(CO)₃+**10e**]⁺ ion.

[Re(CO)₃(10f)]Br, (Re-10f)

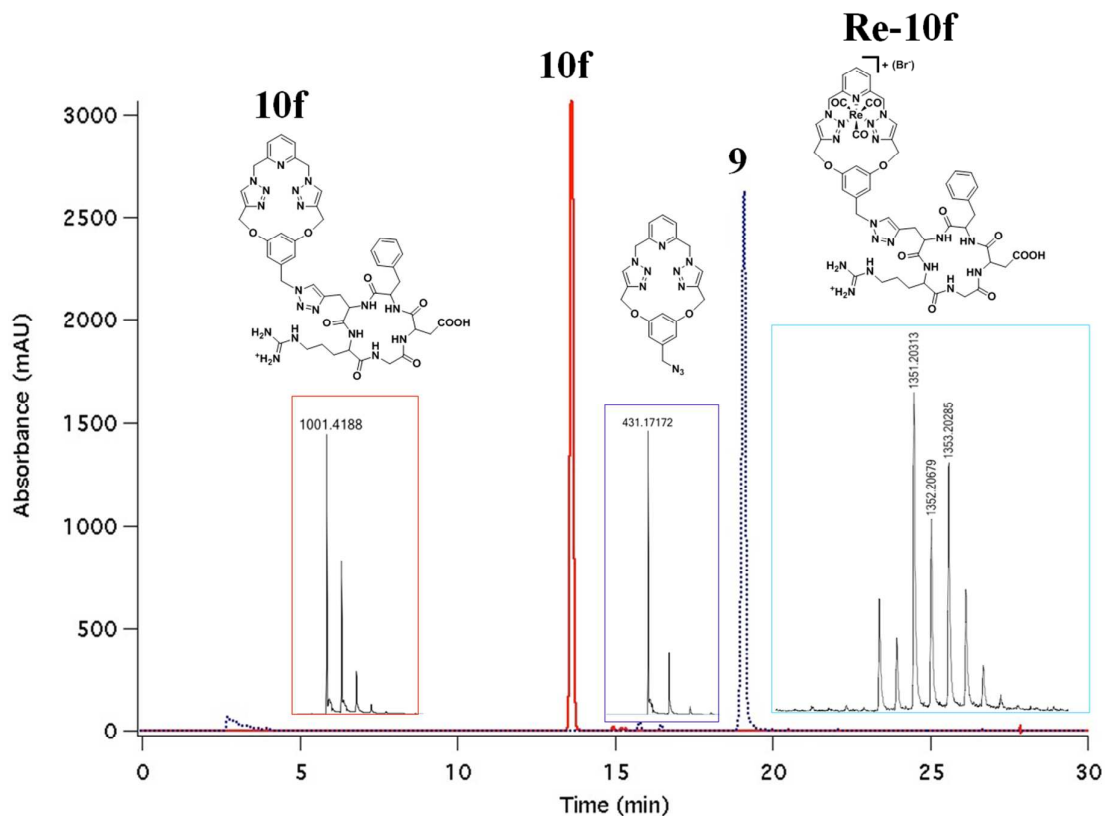


Figure S 29 RP-HPLC traces for **10f** (Method: 0-60% CH₃CN), **9** (Method: 0-100% CH₃CN) and observed ESI-MS isotopic patterns for **10f** ($m/z = 1001.4188$), **9** ($m/z = 431.1717$) and **Re-10f** ($m/z = 1351.2031$).

2 Selected stacked ^1H NMR spectra of the ligands and metal complexes

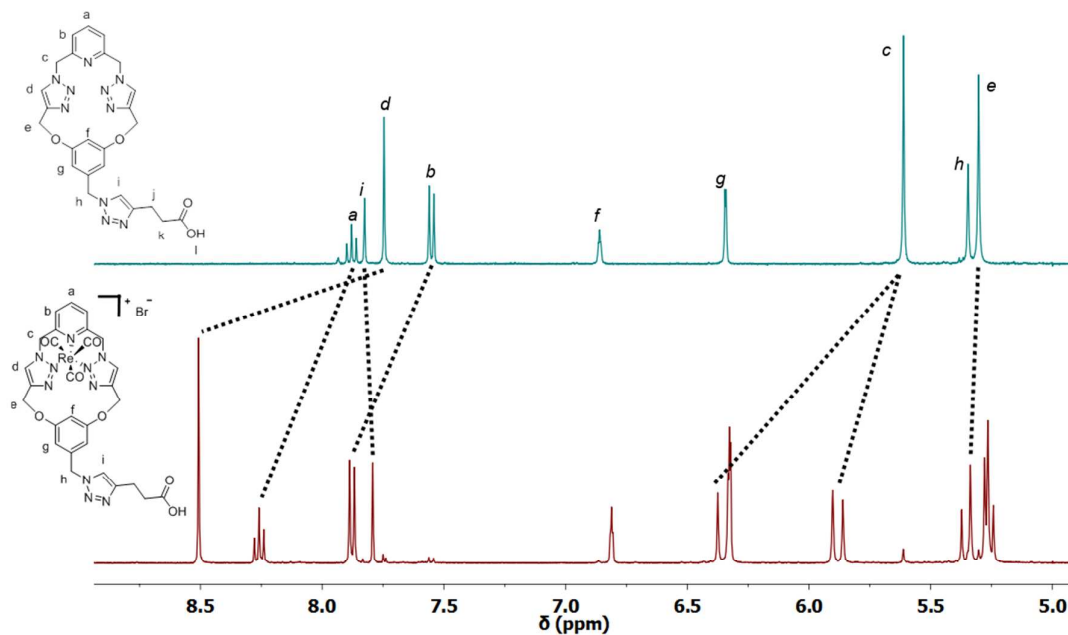


Figure S 30 Partial stacked ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$, 298 K) of acid-conjugated macrocycle **10a** and $[\text{Re}(\text{CO})_3]^+$ complex **Re-10a**.

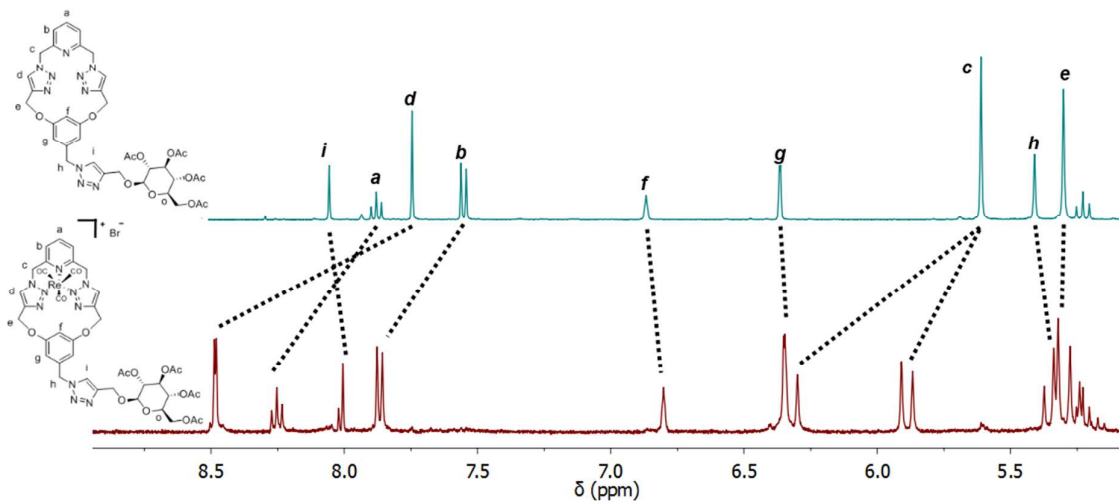


Figure S 31 Partial stacked ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$, 298 K) of sugar-conjugated macrocycle **10c** and $[\text{Re}(\text{CO})_3]^+$ complex **Re-10c**.

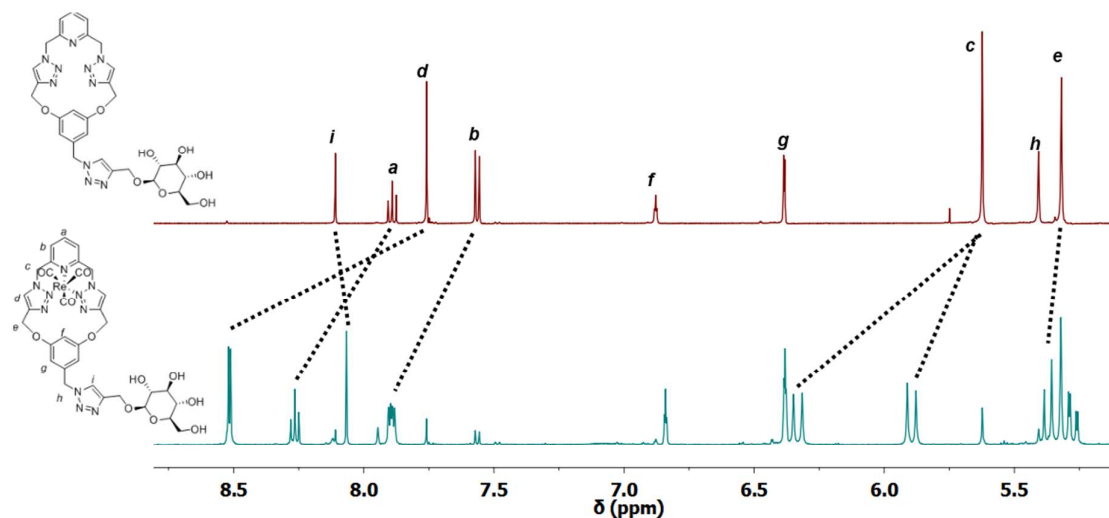


Figure S 32 Partial stacked ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$, 298 K) of sugar-conjugated macrocycle **10d** and $[\text{Re}(\text{CO})_3]^+$ complex **Re-10d**.

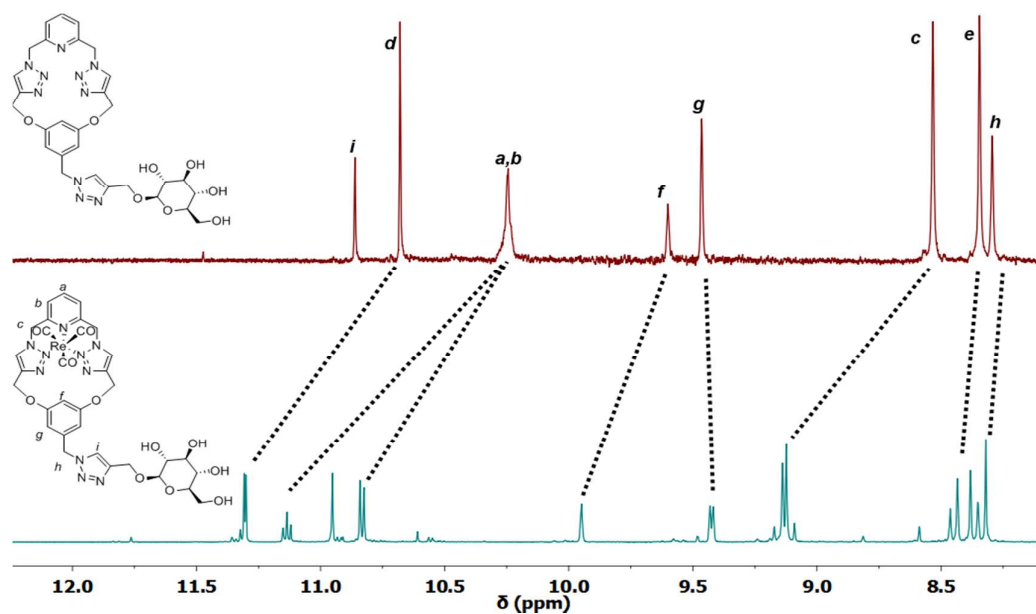


Figure S 33 Partial stacked ^1H NMR spectra (400 MHz, D_2O , 298 K) of sugar-conjugated macrocycle **10d** and $[\text{Re}(\text{CO})_3]^+$ complex **Re-10d** showing aqueous solubility of both ligand and complex at room temperature.

3 Stability and Re(I) labelling experiments

3.1 Temperature Stability

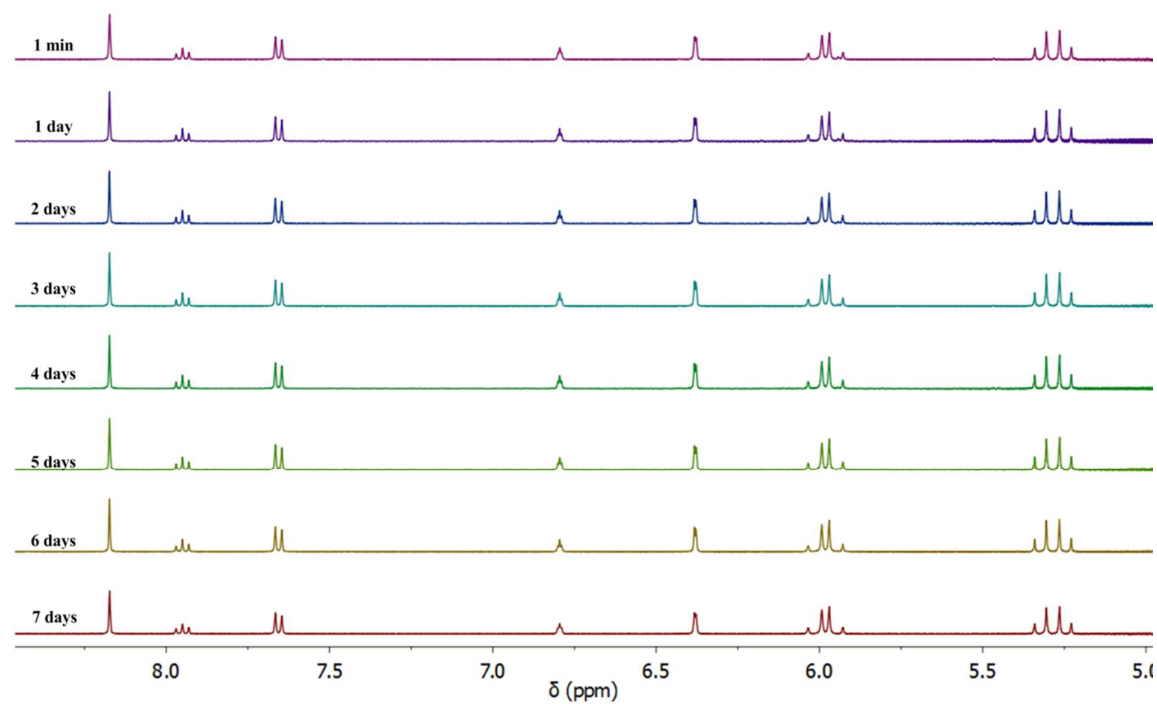


Figure S 34 Partial ^1H NMR spectra (400 MHz, D_2O , 298 K) of the Re(I) complex, **7-Re** over a period of one week.

3.2 Histidine competition experiments

7-Re vs Histidine

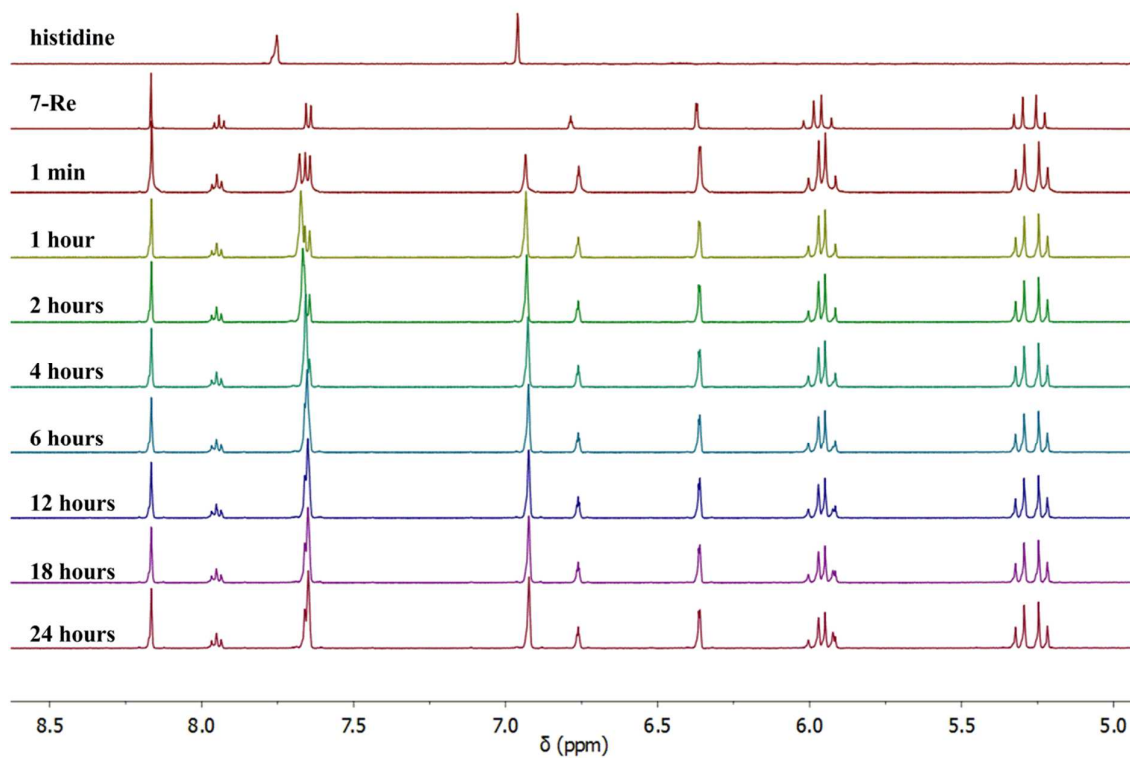


Figure S 35 Partial ^1H NMR spectra (500 MHz, D_2O , 313 K) of the histidine competition experiments showing top to bottom, histidine, the Re(I) complex **7-Re**, and the reaction mixtures of both after specified interval of time over 24 hours

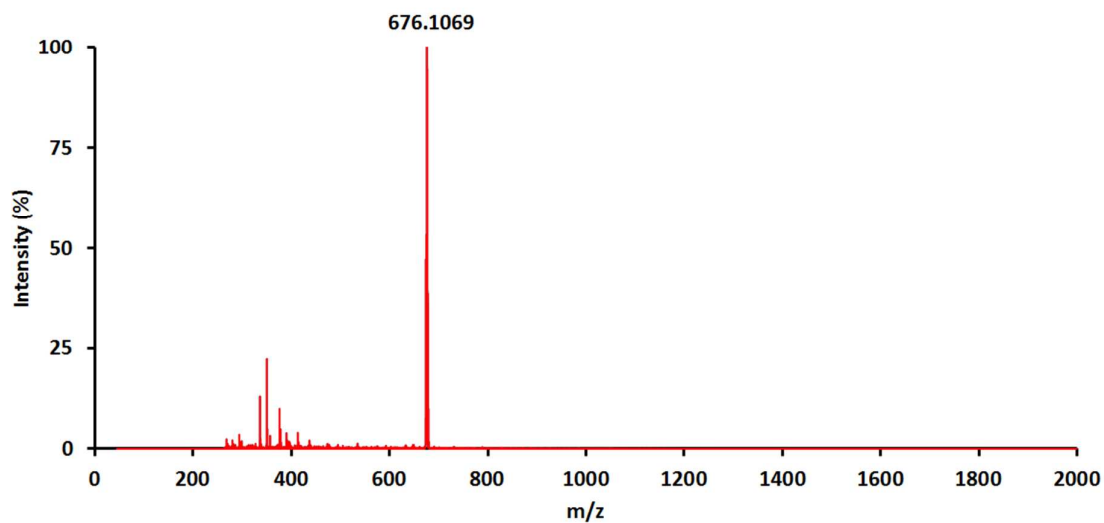


Figure S 36 HR-ESI-MS spectrum of the reaction mixture after completion of histidine competition experiment showing only major peak due to the $[\text{Re}(\text{CO})_3+7]^+$ ions at $m/z = 676.1069$

10d-Re vs Histidine

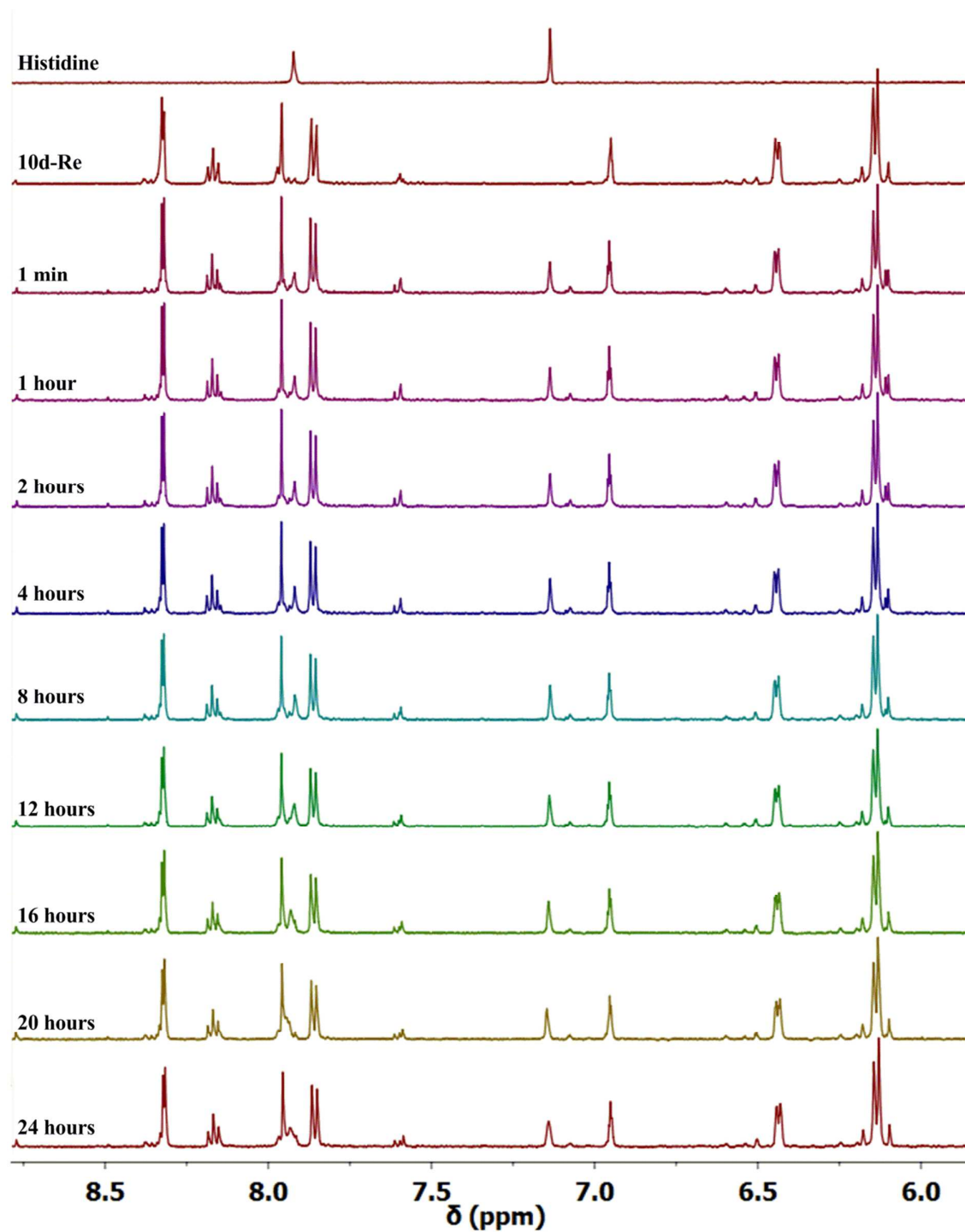


Figure S 37 Partial ¹H NMR spectra (500 MHz, D₂O, 313 K) of the histidine competition experiments showing top to bottom, histidine, the Re(I) complex **10d-Re**, and the reaction mixtures of both after specified interval of time over 24 hours

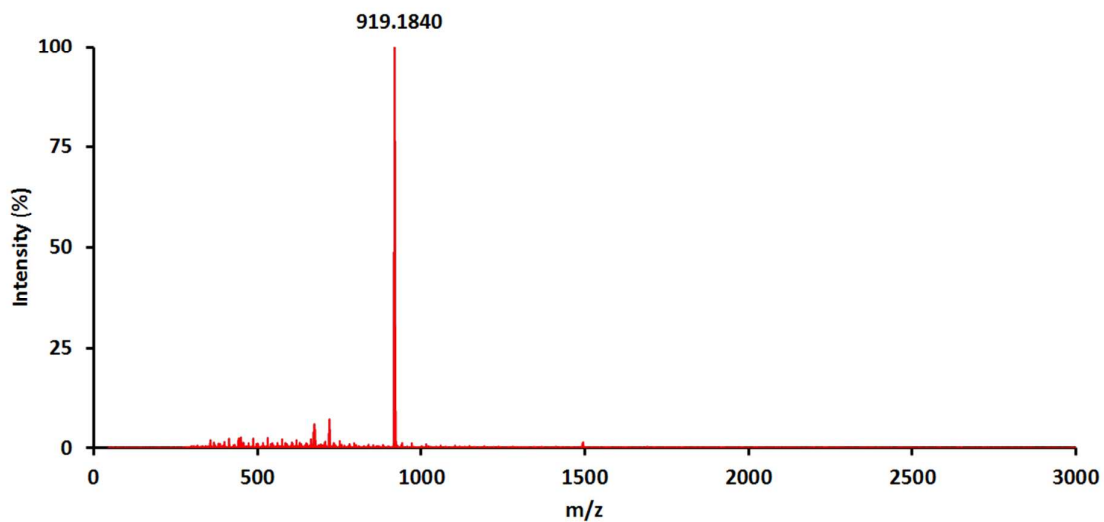


Figure S 38 HR-ESI-MS spectrum of reaction mixture after completion of histidine competition experiment showing only a major peak due to the $[\text{Re}(\text{CO})_3 + \mathbf{10d}]^+$ ions at $m/z = 919.1840$

3.3 Labelling experiment of 7-Re

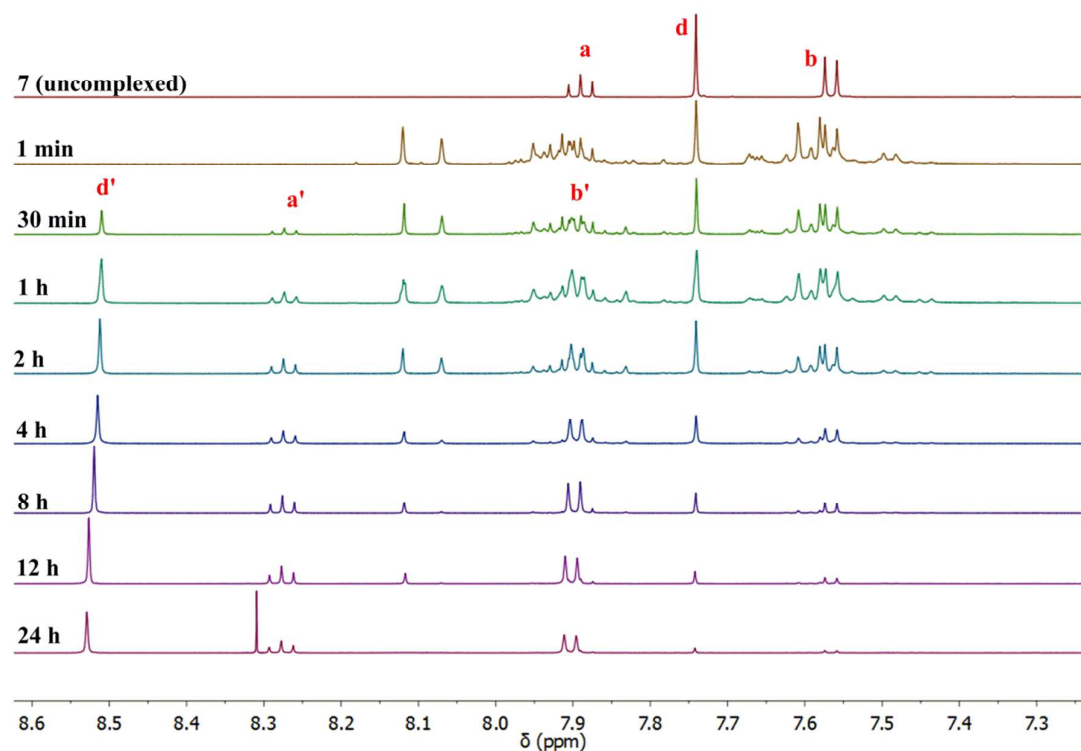


Figure S 39 Partial ^1H NMR spectra (500 MHz, $\text{DMSO-}d_6$, 298 K) of the labelling time experiments showing top to bottom, uncomplexed macrocycle **7**, and the reaction mixture containing **7** and **7-Re** after specified interval of time over 24 hours

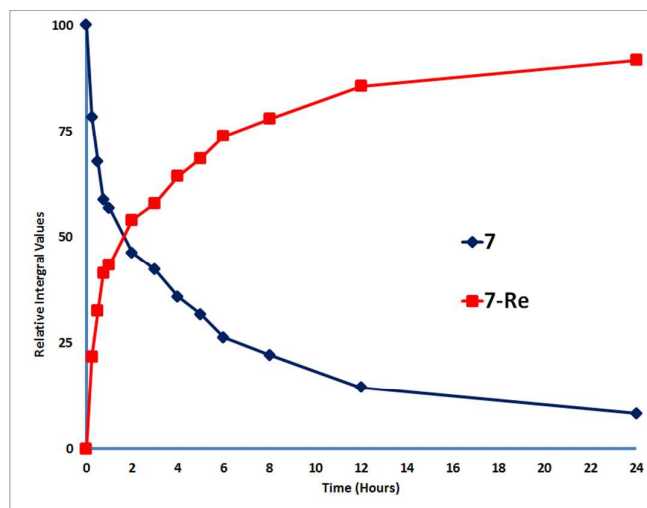


Figure S 40 Time vs integral values of the NMR labelling experiment of **7-Re**

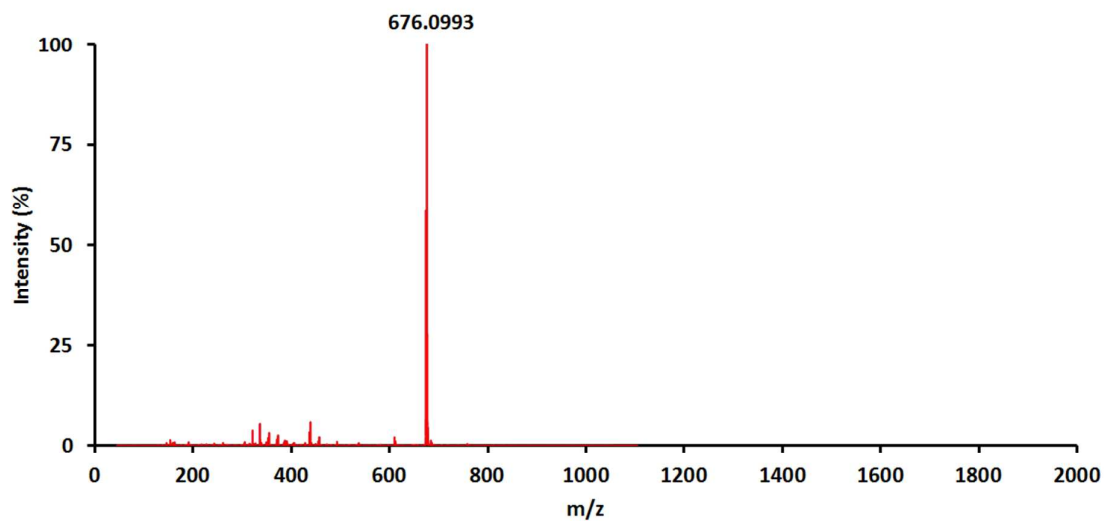


Figure S 41 HR-ESI-MS spectrum of reaction mixture after completion of labelling experiment showing only a major peak due to the $[\text{Re}(\text{CO})_3+7]^+$ ions at m/z 676.0993.

4 Antibacterial Activity

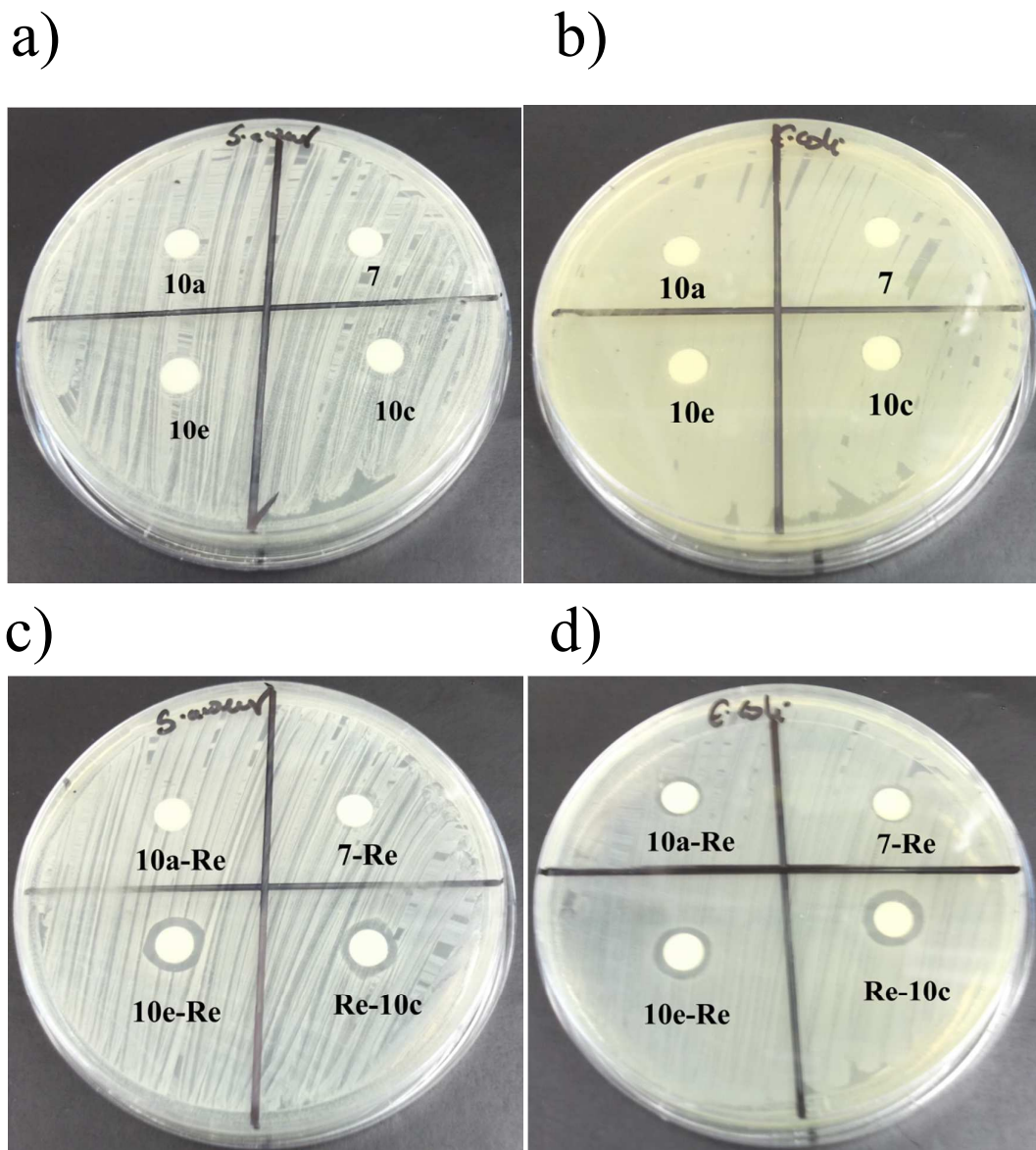


Figure S 42 a) & b) Disk diffusion assay of ligands on *S. aureus* and *E. coli*. c) & d) Disk diffusion assay of rhenium complexes on *S. aureus* and *E. coli*.

5 Photophysical Properties and Density Functional Theory Calculations

Table S 1 Calculated bond lengths and angles for **7-Re**

crystal structure	B3LYP		PBE0		CAM-B3LYP		M06		M062X	
Bond Length (Å)	Bond Length (Å)	Error	Bond Length (Å)	Error	Bond Length (Å)	Error	Bond Length (Å)	Error	Bond Length (Å)	Error
Re-N2	2.160		2.205	2.06%	2.175	0.67%	2.191	1.44%	2.200	1.86%
Re-N4	2.252		2.306	2.41%	2.259	0.31%	2.285	1.48%	2.295	1.93%
Re-N6	2.158		2.207	2.27%	2.177	0.87%	2.193	1.64%	2.205	2.16%
Angle (°)	Angle (°)	Error	Angle (°)	Error	Angle (°)	Error	Angle (°)	Error	Angle (°)	Error
N4-Re-N2	79.14		79.95	1.03%	80.46	1.66%	79.98	1.06%	79.76	0.79%
N4-Re-N6	80.44		79.90	0.68%	80.38	0.08%	79.93	0.63%	79.51	1.16%
N2-Re-N6	86.14		89.72	4.15%	89.65	4.08%	89.45	3.84%	89.34	3.71%

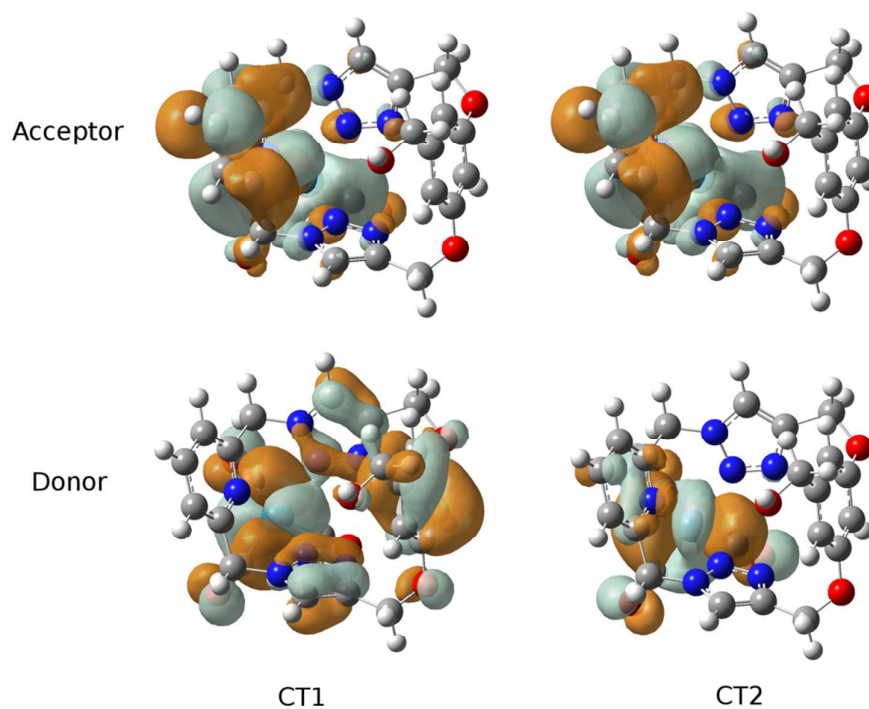


Figure S 43 Natural transitions orbitals for **7-Re** using M062X/6-31G(d)-LANL2DZ

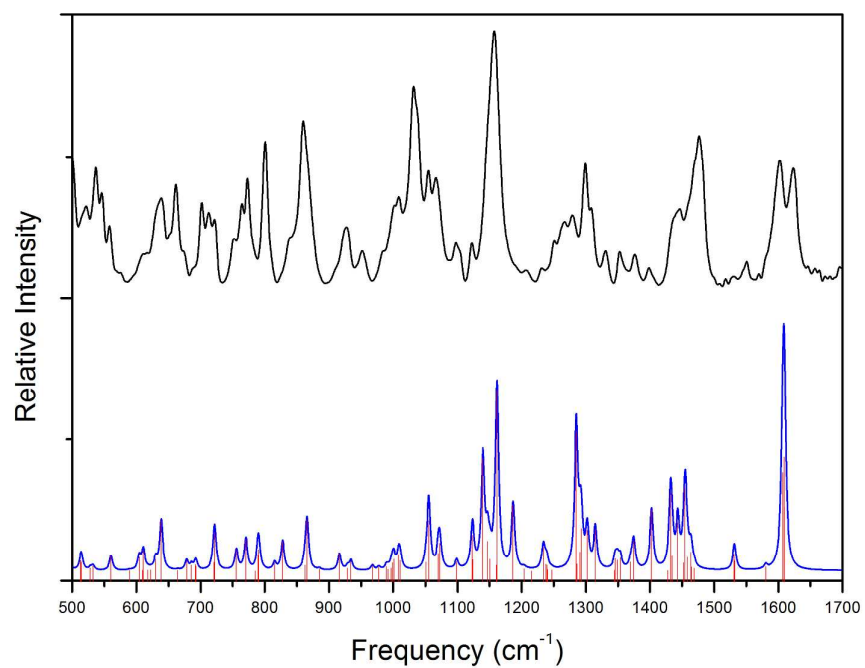


Figure S 44 ATR-IR spectrum of **7-Re** (top) and Calculated IR spectrum using CAM-B3LYP/6-31G(d)-LANL2DZ (bottom)

6 X-Ray data collection and refinement

6.1 Macrocycle (7)

X-Ray data for **7** was recorded using a Bruker APEX II CCD diffractometer using Mo-K α radiation ($\lambda = 0.71073$ Å). The structure was solved in the monoclinic space group $C2/c$ and refined to a R_1 value 7.2%. Hydrogen atoms were placed in calculated positions and refined using a riding model.

6.2 Re(I) complex of macrocycle (7-Re)

X-Ray data for **7-Re** was recorded using a Bruker APEX II CCD diffractometer using Mo-K α radiation ($\lambda = 0.71073$ Å). The structure was solved in triclinic space group $P\bar{1}$ using Fourier methods and refined using SHELXL¹ to a R_1 value 6.3%. Hydrogen atoms were placed in calculated positions and refined using a riding model.

6.3 Re(I) complex of macrocycle (Re-10a)

X-ray data for macrocycle **Re-10a** was collected at 100 K on a CrysAlisPro², Agilent Technologies diffractometer using Mo-K α radiation ($\lambda = 0.71073$ Å). The structure was solved by direct methods and refined against F^2 using anisotropic thermal displacement parameters for all non-hydrogen atoms using the xSeed³ and SHELXS-97⁴ program to a R_1 value 5.7%.. Hydrogen atoms were placed in calculated positions and refined using a riding model. The crystal lattice contained a small amount of diffuse electron density that could not be appropriately modelled. The SQUEEZE routine within PLATON was employed to resolve this problem. Void electron count of 58 electrons can be accounted for by four molecules of water (10 electrons each) and a molecule of methanol (18 electrons) to give a total electron count of 58.

6.4 Structure Refinement Data

Table S2 Crystal data and structure refinement for macrocycle **7**

CCDC	991260	
Empirical formula	C ₂₀ H ₁₉ N ₇ O ₃	
Formula weight	405.42	
Temperature	93(2) K	
Wavelength	0.71073 Å	
Crystal system	Monoclinic	
Space group	C ₂ /c	
Unit cell dimensions	a = 8.918(3) Å	α = 90°.
	b = 16.352(4) Å	β = 96.008(11)°.
	c = 25.482(7) Å	γ = 90°.
Volume	3695.7(17) Å ³	
Z	8	
Density (calculated)	1.457 Mg/m ³	
Absorption coefficient	0.103 mm ⁻¹	
F(000)	1696	
Crystal size	0.22 x 0.21 x 0.05 mm ³	
Theta range for data collection	1.61 to 20.21°.	
Index ranges	-8 ≤ h ≤ 8, -15 ≤ k ≤ 15, -24 ≤ l ≤ 24	
Reflections collected	11510	
Independent reflections	1761 [R(int) = 0.0826]	
Completeness to theta = 20.21°	99.5 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.9949 and 0.9777	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	1761 / 0 / 273	
Goodness-of-fit on F ²	1.144	
Final R indices [I > 2σ(I)]	R1 = 0.0725, wR2 = 0.1684	
R indices (all data)	R1 = 0.0873, wR2 = 0.1800	
Extinction coefficient	0.0024(5)	
Largest diff. peak and hole	0.332 and -0.250 e.Å ⁻³	

Table S3 Crystal data and structure refinement for Re(I) complex **7-Re**

CCDC	991258	
Empirical formula	C ₂₃ H ₁₉ BrN ₇ O ₆ Re	
Formula weight	755.56	
Temperature	89(2) K	
Wavelength	0.71073 Å	
Crystal system	Triclinic	
Space group	<i>P</i> $\bar{1}$	
Unit cell dimensions	a = 9.5826(17) Å	α = 66.484(11)°.
	b = 11.031(2) Å	β = 76.119(11)°.
	c = 13.594(3) Å	γ = 70.788(11)°.
Volume	1234.5(4) Å ³	
Z	2	
Density (calculated)	2.033 Mg/m ³	
Absorption coefficient	6.595 mm ⁻¹	
F(000)	728	
Crystal size	0.15 x 0.12 x 0.05 mm ³	
Theta range for data collection	1.65 to 26.54°.	
Index ranges	-11 ≤ h ≤ 11, -13 ≤ k ≤ 13, -17 ≤ l ≤ 17	
Reflections collected	32686	
Independent reflections	5000 [R(int) = 0.0988]	
Completeness to theta = 26.54°	97.6 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.7339 and 0.4379	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	5000 / 0 / 344	
Goodness-of-fit on F ²	1.070	
Final R indices [I > 2sigma(I)]	R1 = 0.0633, wR2 = 0.1573	
R indices (all data)	R1 = 0.0996, wR2 = 0.1866	
Largest diff. peak and hole	3.019 and -2.161 e.Å ⁻³	

Table S4 Crystal data and structure refinement for **Re-10a**

CCDC	991259	
Empirical formula	C ₂₉ H ₂₆ BrN ₁₀ O ₇ Re	
Formula weight	892.71	
Temperature	100.0(1) K	
Wavelength	0.71073 Å	
Crystal system	Triclinic	
Space group	<i>P</i> $\bar{1}$	
Unit cell dimensions	a = 10.4722(4) Å	α = 93.303(3)°.
	b = 16.0868(8) Å	β = 90.356(3)°.
	c = 23.1434(7) Å	γ = 108.644(4)°.
Volume	3686.8(3) Å ³	
Z	4	
Density (calculated)	1.608 Mg/m ³	
Absorption coefficient	4.434 mm ⁻¹	
F(000)	1744	
Crystal size	0.20 x 0.17 x 0.06 mm ³	
Theta range for data collection	2.88 to 26.37°.	
Index ranges	-13 ≤ h ≤ 12, -20 ≤ k ≤ 20, -28 ≤ l ≤ 28	
Reflections collected	31782	
Independent reflections	14778 [R(int) = 0.0483]	
Completeness to theta = 26.37°	98.6 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.7768 and 0.4708	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	14778 / 35 / 869	
Goodness-of-fit on F ²	1.043	
Final R indices [I > 2sigma(I)]	R1 = 0.0572, wR2 = 0.1491	
R indices (all data)	R1 = 0.0772, wR2 = 0.1598	
Largest diff. peak and hole	2.784 and -2.246 e.Å ⁻³	

Table S5 Selected bond lengths (Å) and angles (°) for **7-Re**, **Re-10a** and similar acyclic complex $[\text{Re}(\text{CO})_3(4 \text{ R} = \text{C}_3\text{H}_7)]\text{PF}_6$ (atom labels in bold correspond to the labelling of structure **Re-10a** in **Figure S19**)

	7-Re	Re-10a	$[\text{Re}(\text{CO})_3(4 \text{ R} = \text{C}_3\text{H}_7)]\text{PF}_6^5$
Bond Lengths (Å)			
Re(1)-C(30)/ C(29)	1.907(11)	1.906(9)	1.928(9)
Re(1)-C(32)/ C(27)	1.917(14)	1.926(10)	1.924(5)
Re(1)-C(31)/ C(28)	1.921(16)	1.934(10)	1.934(8)
Re(1)-N(6)	2.158(9)	2.179(6)	2.173(3)
Re(1)-N(2)/ N(3)	2.159(9)	2.171(6)	2.155(4)
Re(1)-N(4)/ N(1)	2.252(11)	2.241(6)	2.252(3)
Bond Angles (°)			
C(29) /C(30)-Re(1)-N(6)	92.9(5)	90.8(3)	91.47(16)
C(27) /C(32)-Re(1)-N(6)	178.4(4)	175.8(3)	176.92(15)
C(28) /C(31)-Re(1)-N(6)	91.7(4)	94.5(3)	94.64(16)
C(29) /C(30)-Re(1)-N(2)/ N(3)	178.8(4)	177.5(3)	177.96(15)
C(27) /C(32)-Re(1)-N(2)/ N(3)	95.5(4)	96.3(3)	95.55(16)
C(28) /C(31)-Re(1)-N(2)/ N(3)	91.7(4)	92.5(3)	91.77(17)
N(6)-Re(1)-N(2)/ N(3)	86.1(3)	86.9(2)	79.9(2)
C(29) /C(30)-Re(1)-N(4)/ N(1)	101.3(5)	101.8(3)	101.35(16)
C(27) /C(32)-Re(1)-N(4)/ N(1)	100.3(5)	98.5(3)	97.88(15)
C(28) /C(31)-Re(1)-N(4)/ N(1)	168.2(4)	169.6(3)	169.64(15)
N(6) -Re(1)-N(4)/ N(1)	80.3(4)	79.6(2)	80.9(2)
N(3) /N(2)-Re(1)-N(4)/ N(1)	79.1(3)	78.7(2)	78.95(13)

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