

Electronic Supplementary Information

Synthesis and characterization of cyclodiphosphazanes embedded macrocycle tetrabromo-resorcin[4]arene-tetrakis(cyclodiphosphazane) and its tetra-rhodium(I) complex

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Table S1 Crystallographic information for compounds **1** and **2**.

	1	2
Empirical formula	C ₇₆ H ₁₁₅ Br ₄ N ₈ O ₈ P ₈ ·1(C ₆ H ₁₄)	C ₂₁₆ H ₃₀₄ Br ₈ Cl ₈ N ₁₆ O ₁₆ P ₁₆ Rh ₈
M _w	1922.32	5622.41
Cryst. System	Triclinic	Orthorhombic
Space group	P-1	Amm2
<i>a</i> , Å	12.7606(3)	32.9022(9)
<i>b</i> , Å	19.6694(5)	37.7479(10)
<i>c</i> , Å	19.8216(5)	12.5306(4)
α , deg	89.631(2)	90
β , deg	72.434(2)	90
γ , deg	83.770(2)	90
<i>V</i> , Å ³	4713.3(2)	15562.9(8)
<i>Z</i>	2	2
<i>D</i> _{calc} , g cm ⁻³	1.355	1.200
μ (Mo Ka), mm ⁻¹	1.898	1.637
<i>F</i> (000)	2002.0	5712.0
crystal size, mm	0.271 × 0.091 × 0.072	0.254 × 0.165 × 0.092
<i>T</i> (K)	100	100
2θ range, deg	4.168 to 49.998	3.642 to 50
total no. Reflns	49983	87929
no. of indep reflns	16572 [R _{int} = 0.0819]	14156 [R _{int} = 0.1771]
<i>S</i>	1.033	1.024
<i>R</i> _I	0.0582	0.0719
<i>wR</i> ₂	0.1445	0.1877

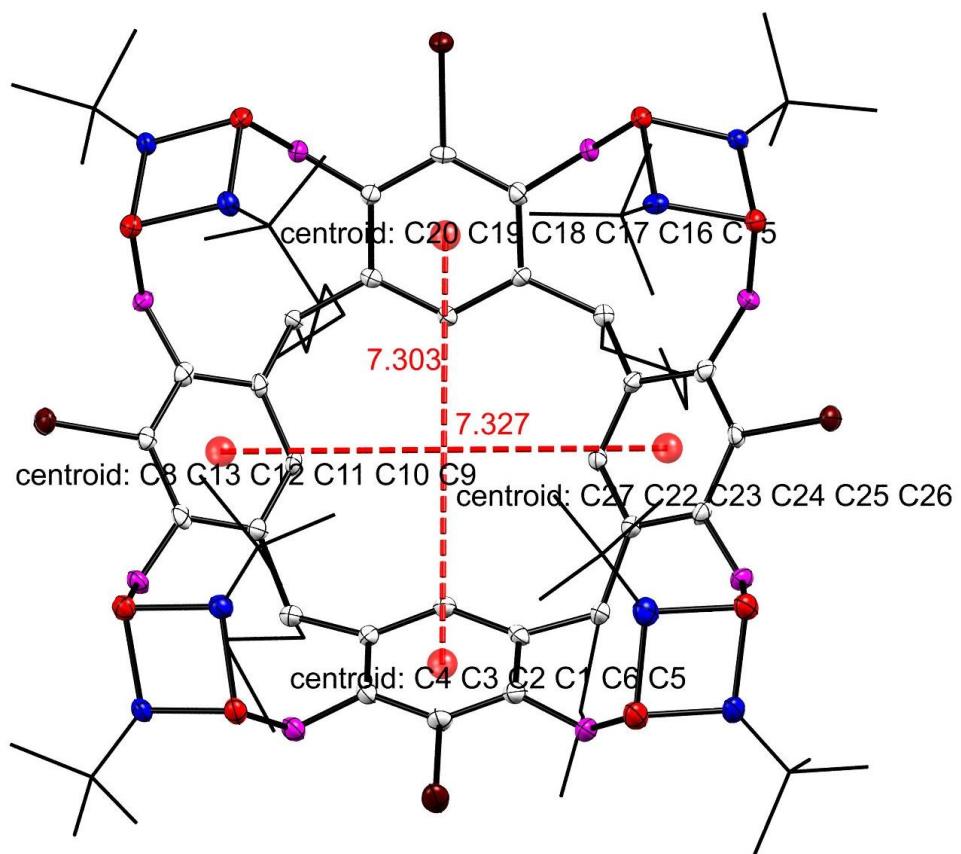


Fig. S1 Molecular structure of **1**(top view). Distances between the centroids of the phenyl groups of resorcin[4]arene core showing C_{4v} crown conformation of **1**.

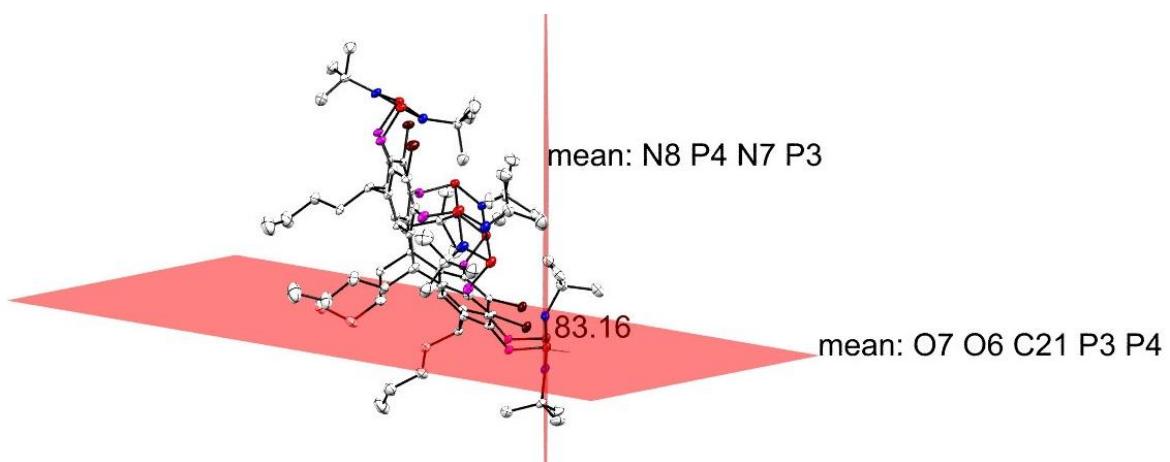


Fig. S2 Molecular structure of **1** showing the angle between planes ($O7O6C21P3P4$) and ($N8P4N7P3$).

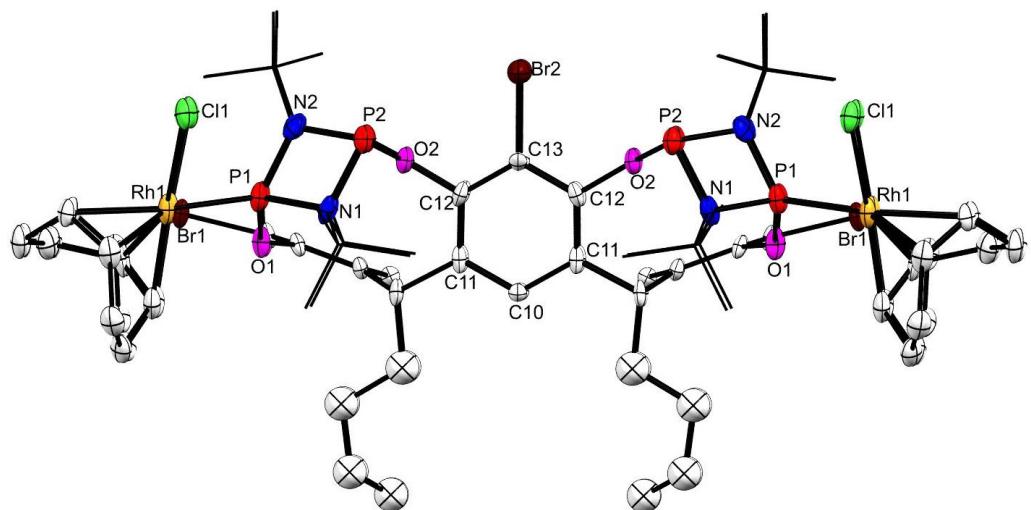


Fig. S3 Molecular structure of **2**. Depicting the structural arrangements (front view).

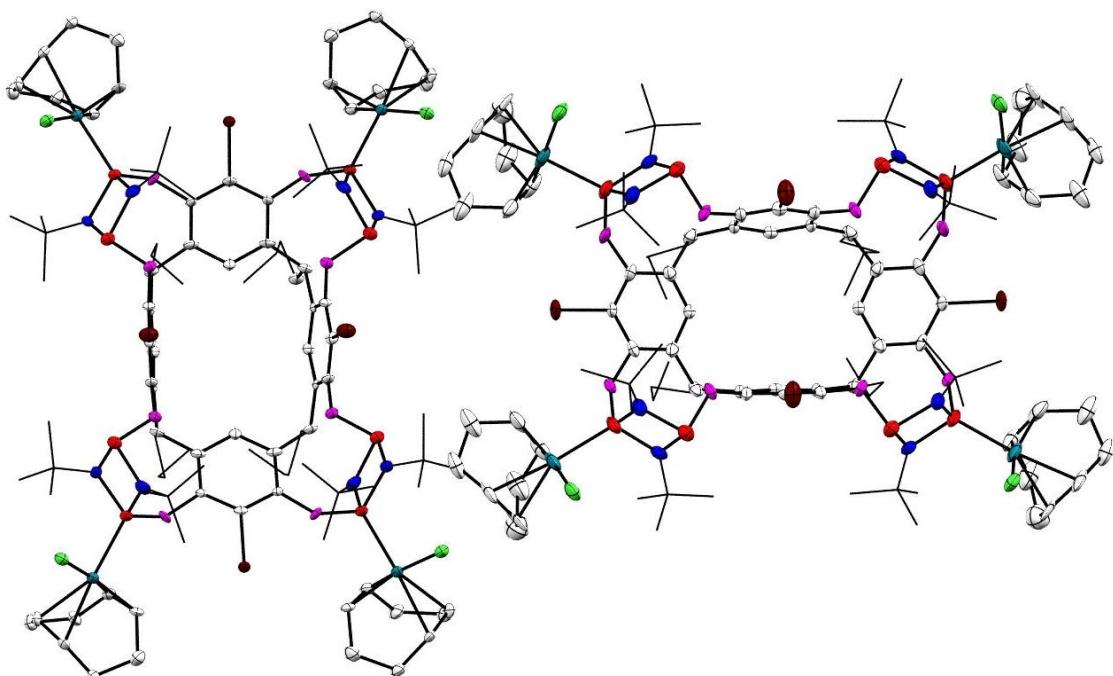


Fig. S4 Molecular structure of **2**(top view). Selected bond lengths (\AA) and bond angles ($^\circ$): Rh1–P1 2.237(3), Rh1–Cl1 2.370(4), P1–O1 1.646(9), P1–N1 1.707(9), O1–P1–N1 108.0(5), P1–Rh1–Cl1 90.74(13), O1–P1–Rh1 111.4(3).

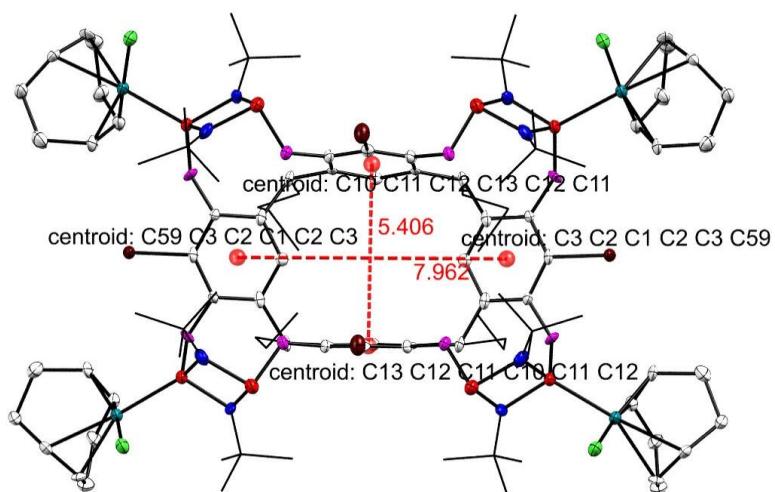


Fig. S5 Molecular structure of **2**(top view). Distances between the centroids of the phenyl groups of resorcin[4]arene showing C_{2v} boat conformation of **2**.

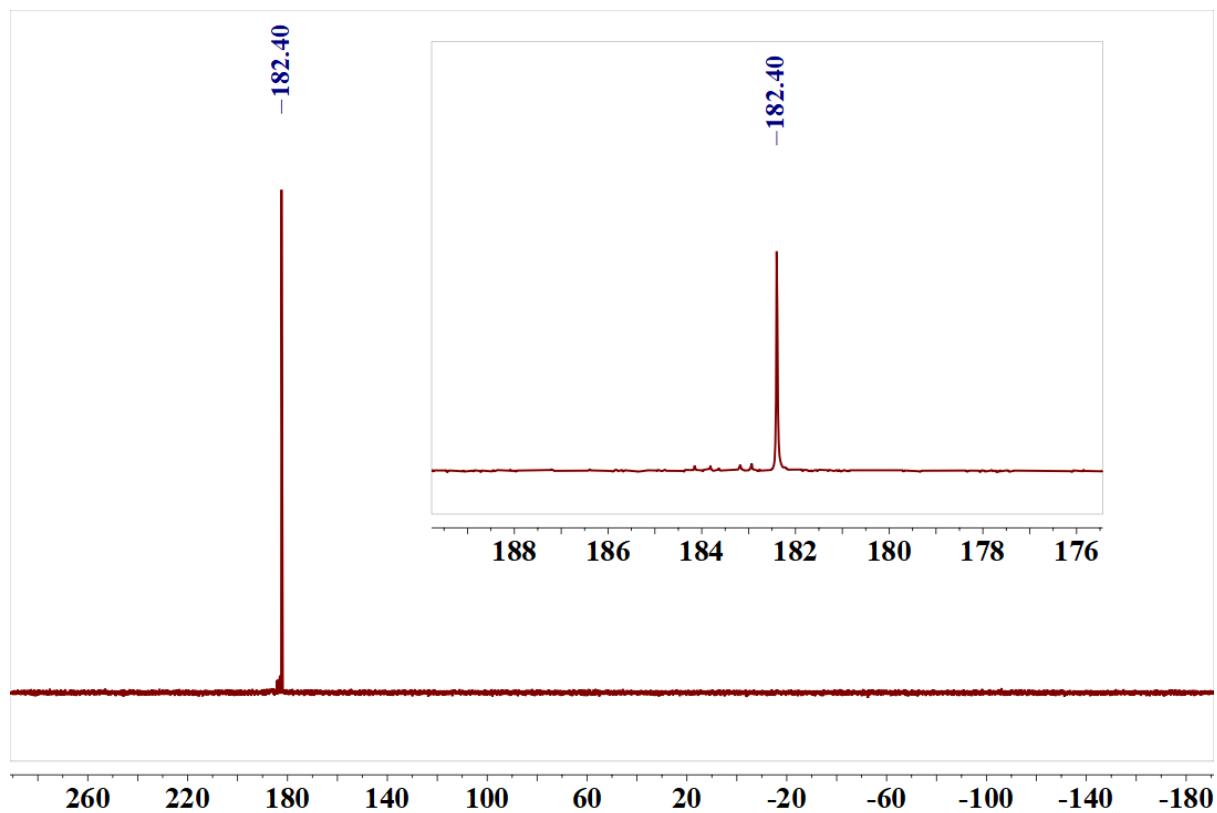


Fig. S6 $^{31}\text{P}\{\text{H}\}$ NMR spectrum of **1** in CDCl_3 (162 MHz).

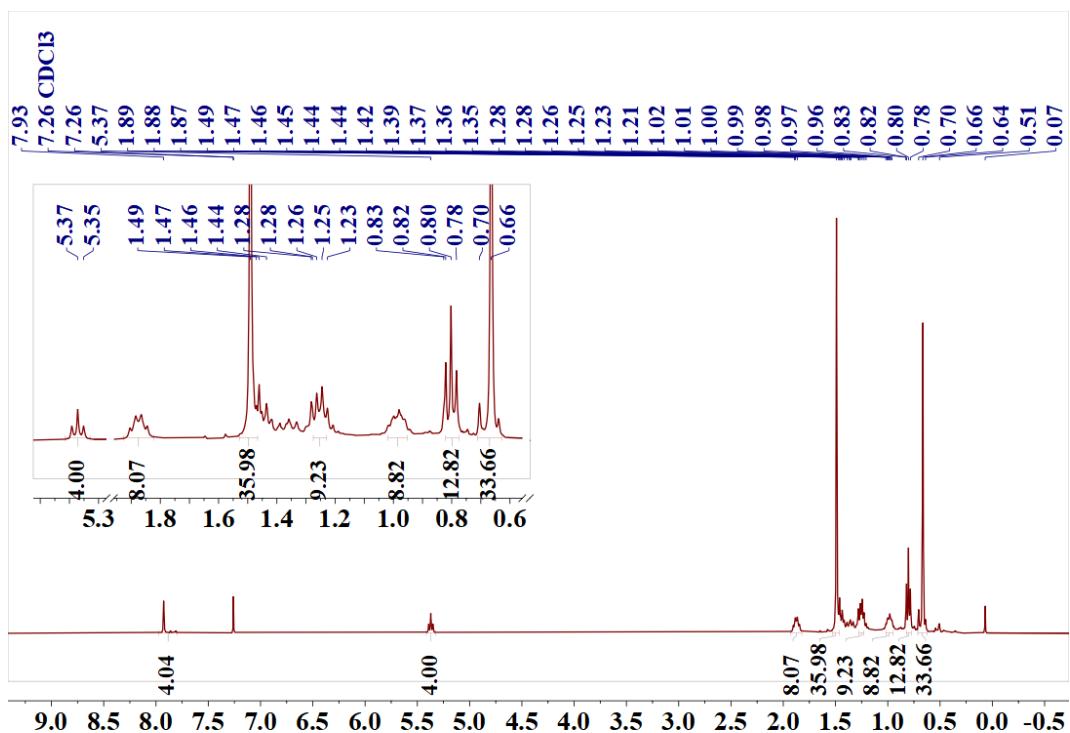


Fig. S7 ^1H NMR spectrum of **1** in CDCl_3 (400 MHz).

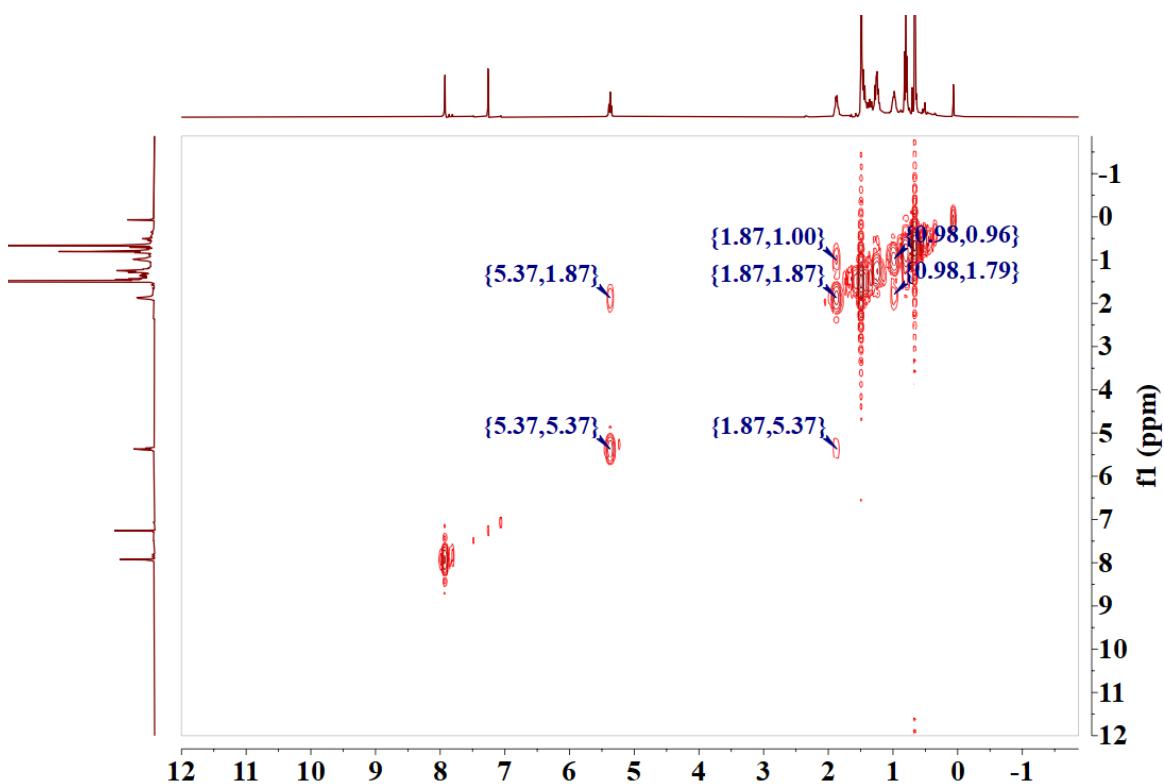


Fig. S8 ^1H - ^1H COSY NMR spectrum of **1** showing the correlation between 5.37, 1.87 ppm peaks and 1.87, 0.98 ppm peaks.

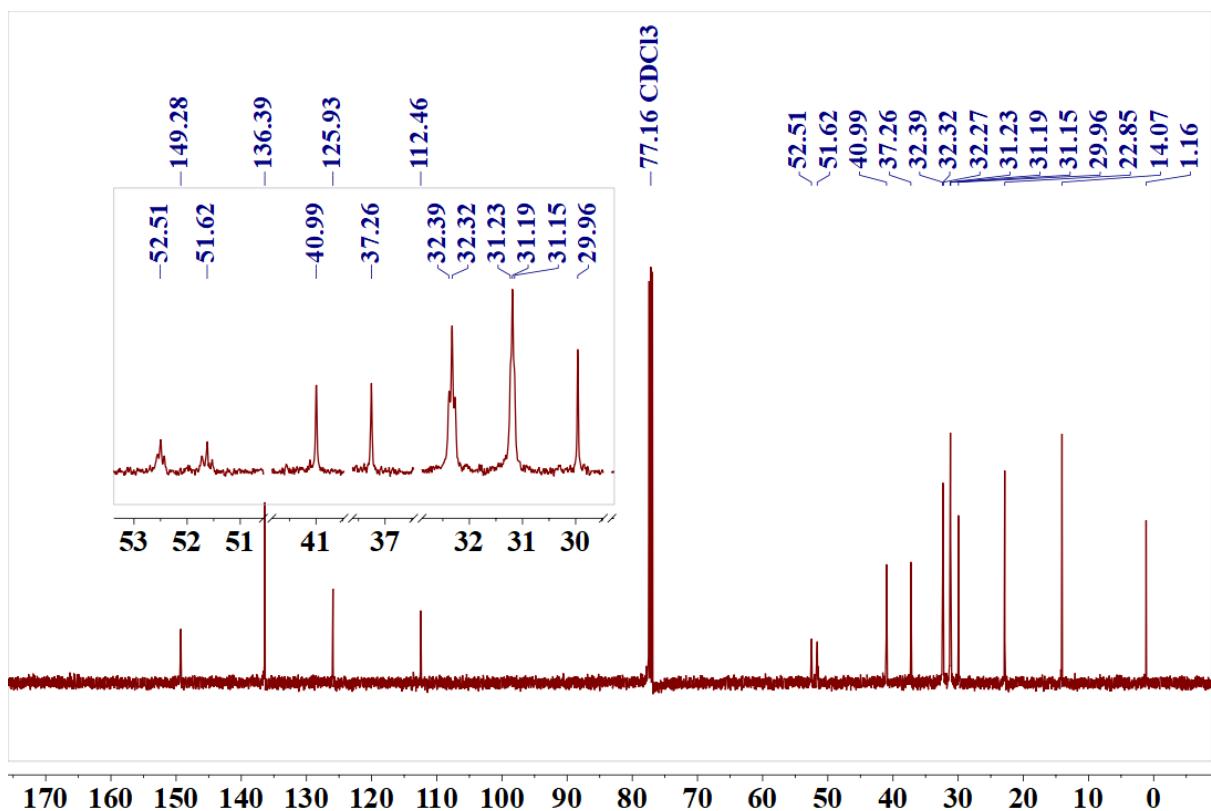


Fig. S9 $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of **1** in CDCl_3 (126 MHz).

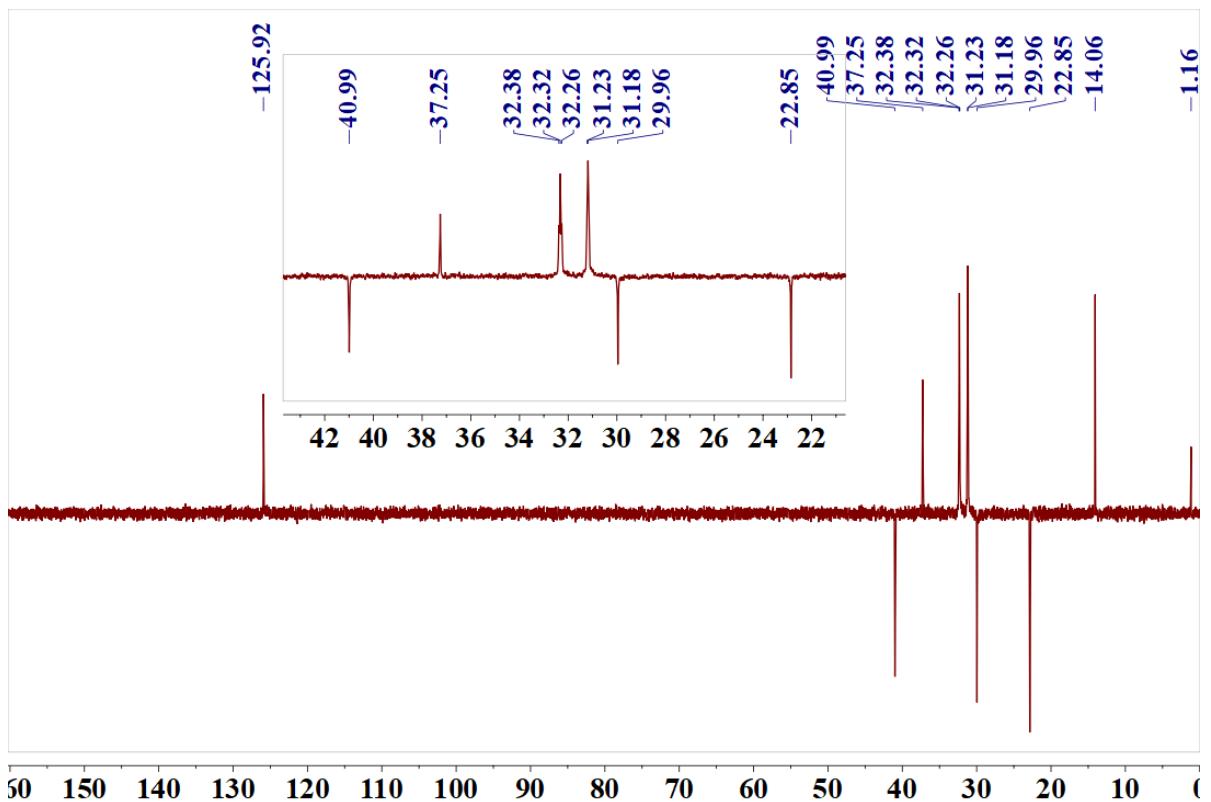


Fig. S10 ^{13}C DEPT-135 NMR spectrum of **1** in CDCl_3 .

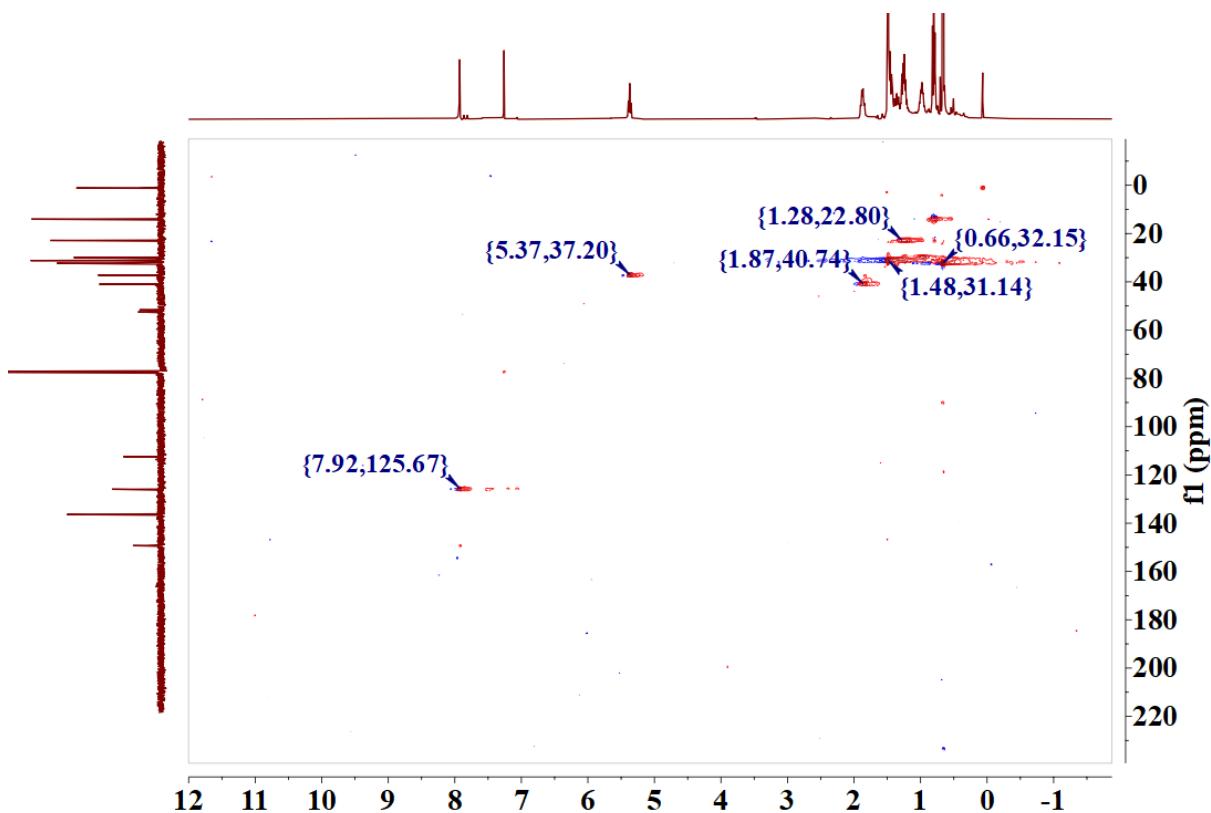


Fig. S11 $^1\text{H}-^{13}\text{C}$ HSQC NMR spectrum of **1** in CDCl_3 .

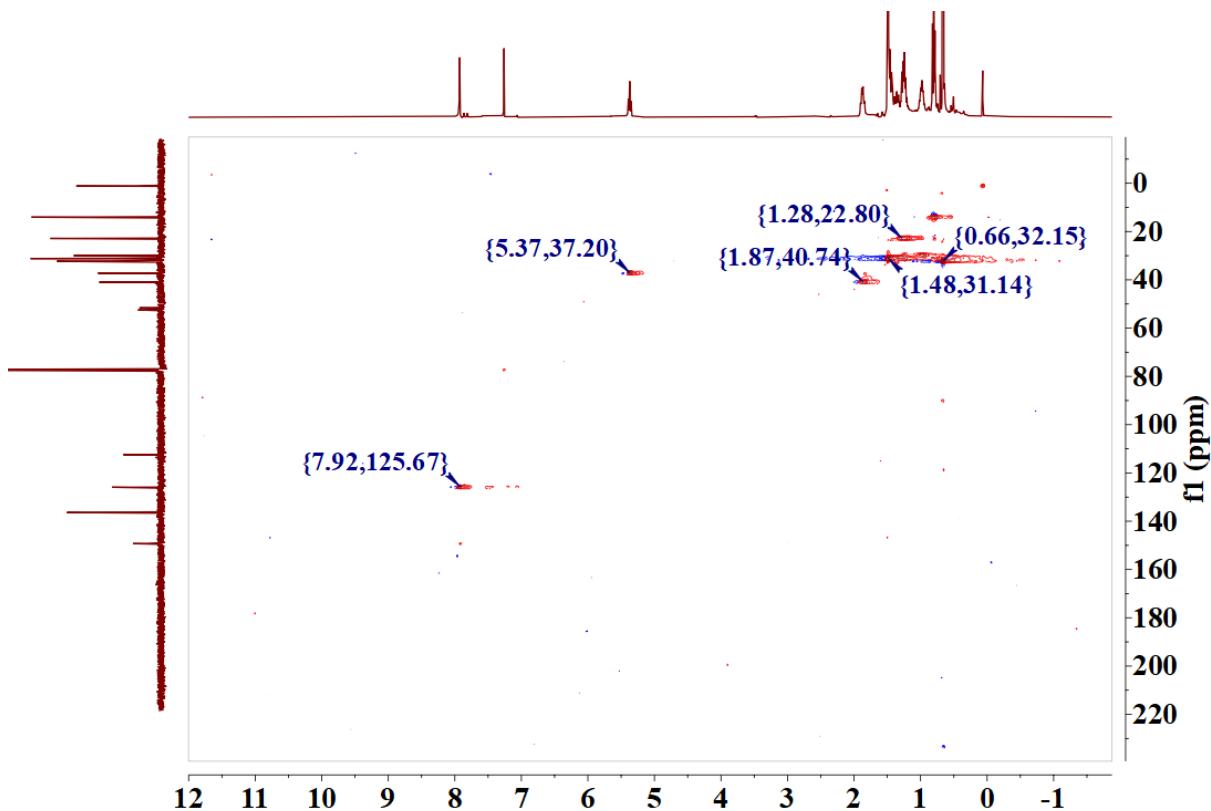


Fig. S12 $^1\text{H}-^{13}\text{C}$ HSQC NMR spectrum of **1** in CDCl_3 (Aliphatic region)

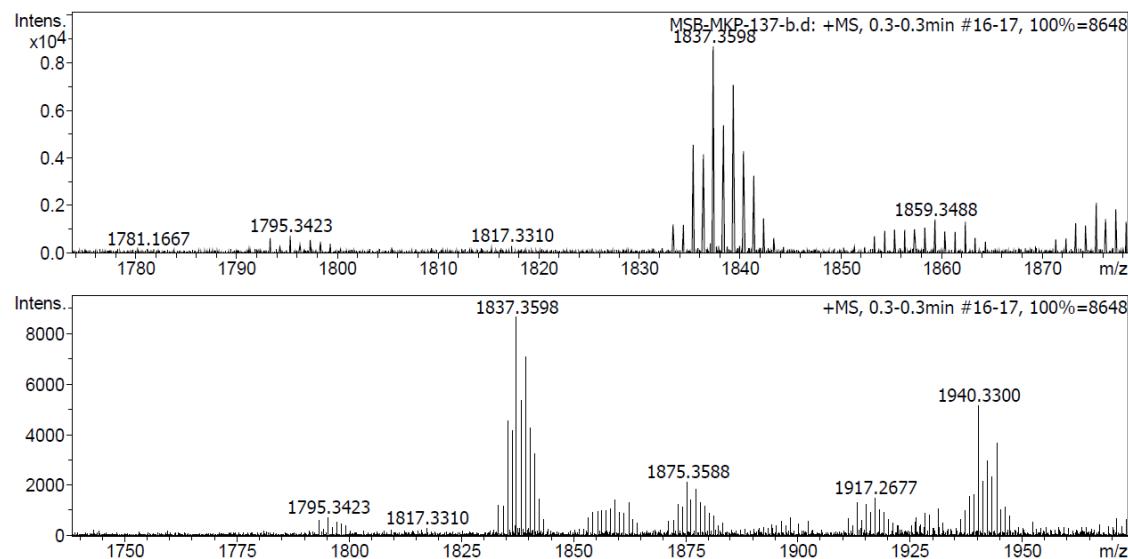
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Analysis Info

Analysis Name	D:\Data\SEPT-2016\MSB-MKP-137-b.d	Acquisition Date	9/3/2016 4:07:28 PM
Method	Tune_pos_NAICSI-3000a.m	Operator	MSB IN
Sample Name	MSB-MKP-137-b	Instrument	maXis impact 282001.00081
Comment	C76H116O8N8P8Br4		

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	0.3 Bar
Focus	Active	Set Capillary	3800 V	Set Dry Heater	180 °C
Scan Begin	50 m/z	Set End Plate Offset	-500 V	Set Dry Gas	4.0 l/min
Scan End	3000 m/z	Set Collision Cell RF	2100.0 Vpp	Set Divert Valve	Source



Meas. m/z	#	Ion Formula	m/z	err [ppm]	mSigma	# Sigma	Score	rdb	e⁻ Conf	N-Rule
1837.3598	1	C76H117Br4N8O8P8	1833.3623	-0.1	43.9	1	100.00	24.5	even	ok

Fig. S13 HRMS spectrum of **1**.

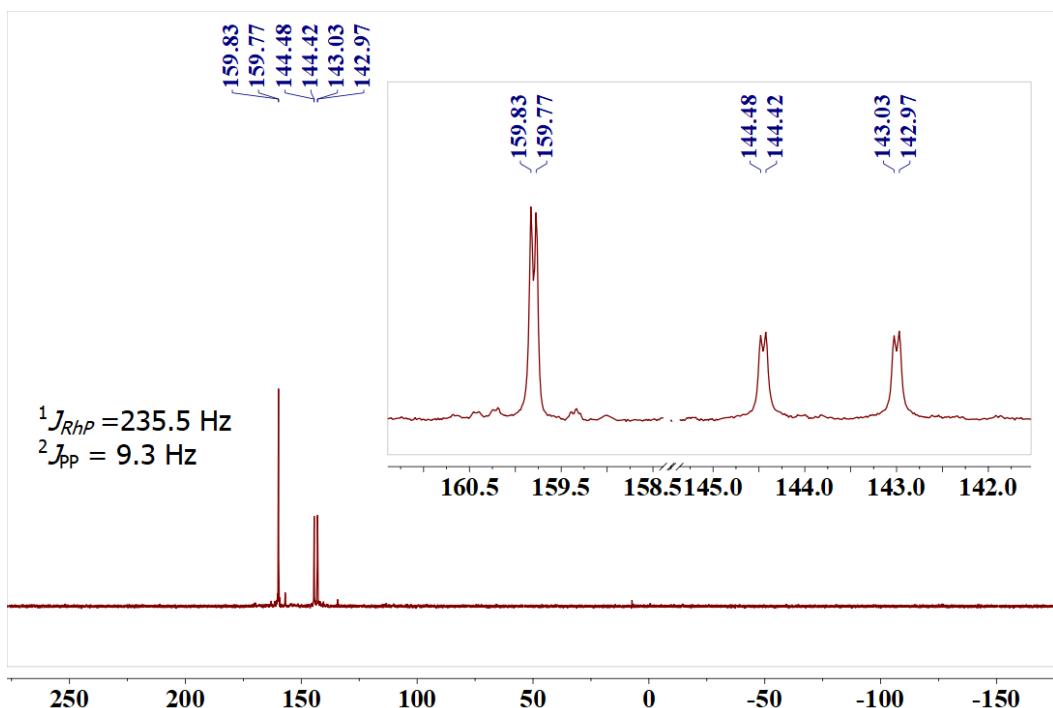


Fig. S14 $^{31}\text{P}\{\text{H}\}$ NMR spectrum of **2** in CDCl_3 (162 MHz).

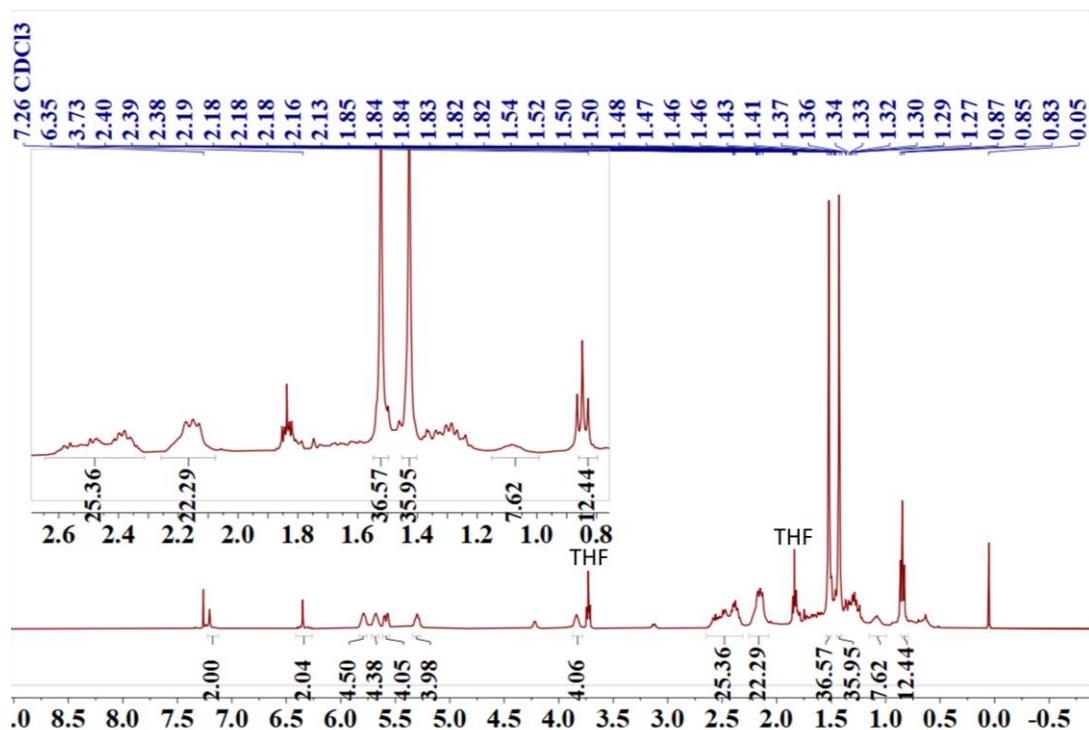


Fig. S15 ^1H NMR spectrum of **2** in CDCl_3 (400 MHz).

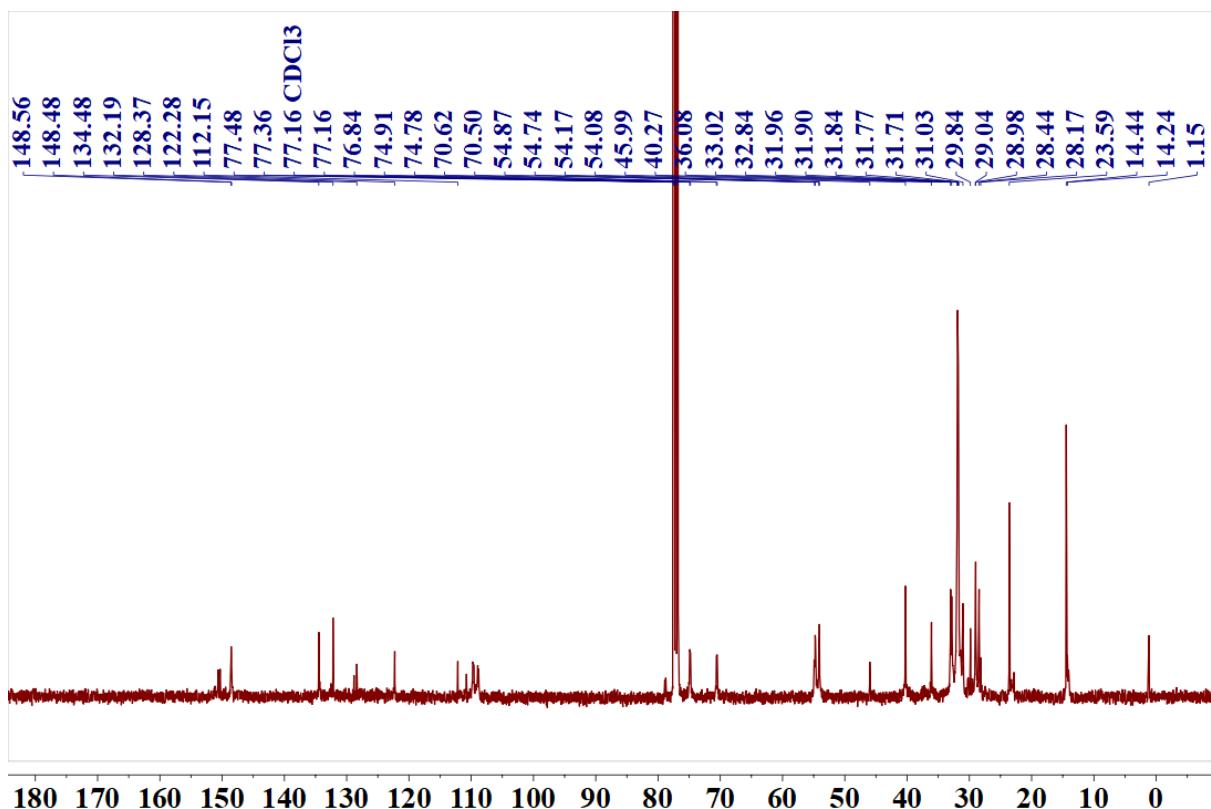


Fig. S16 $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of **2** in CDCl_3 (101 MHz).

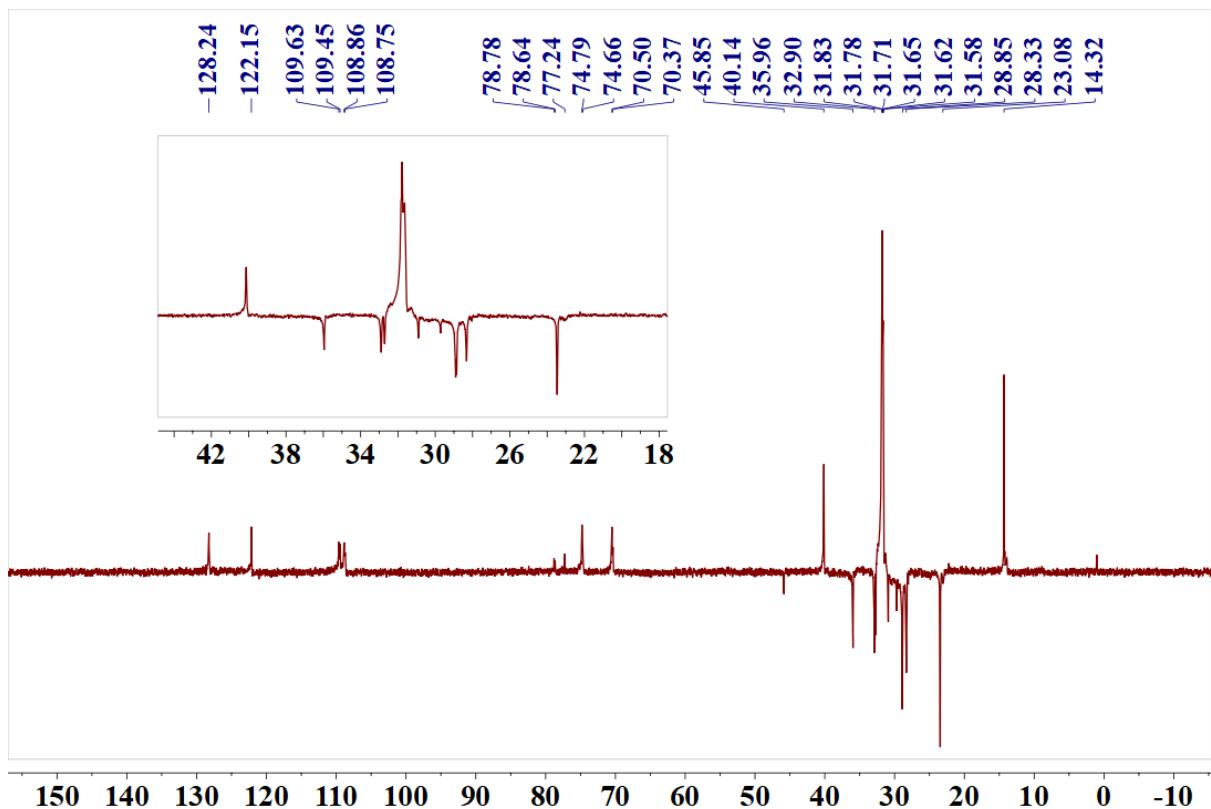


Fig. S17 ^{13}C DEPT-135 NMR spectrum of **2** in CDCl_3 .

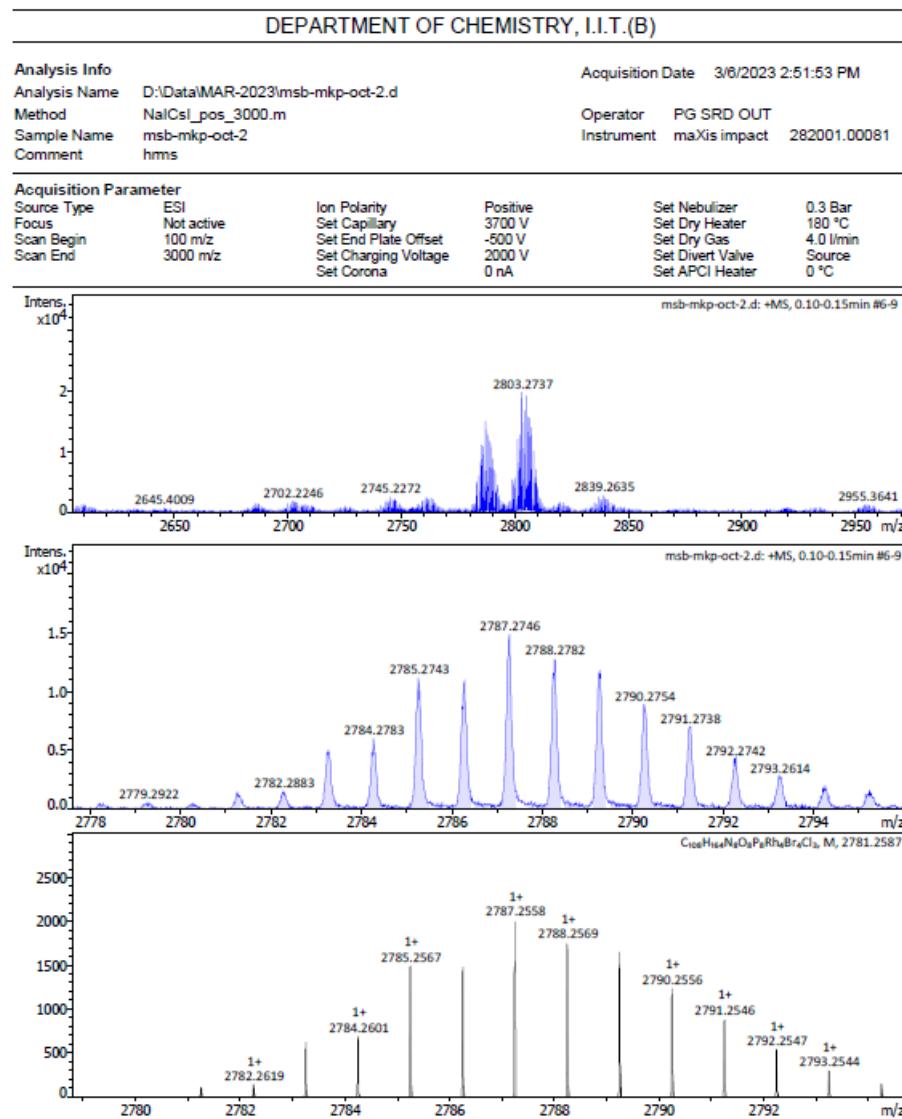


Fig. S18 HRMS spectrum of **2** showing $[M-Cl]^+$.molecular ion peak at 2787.2746.

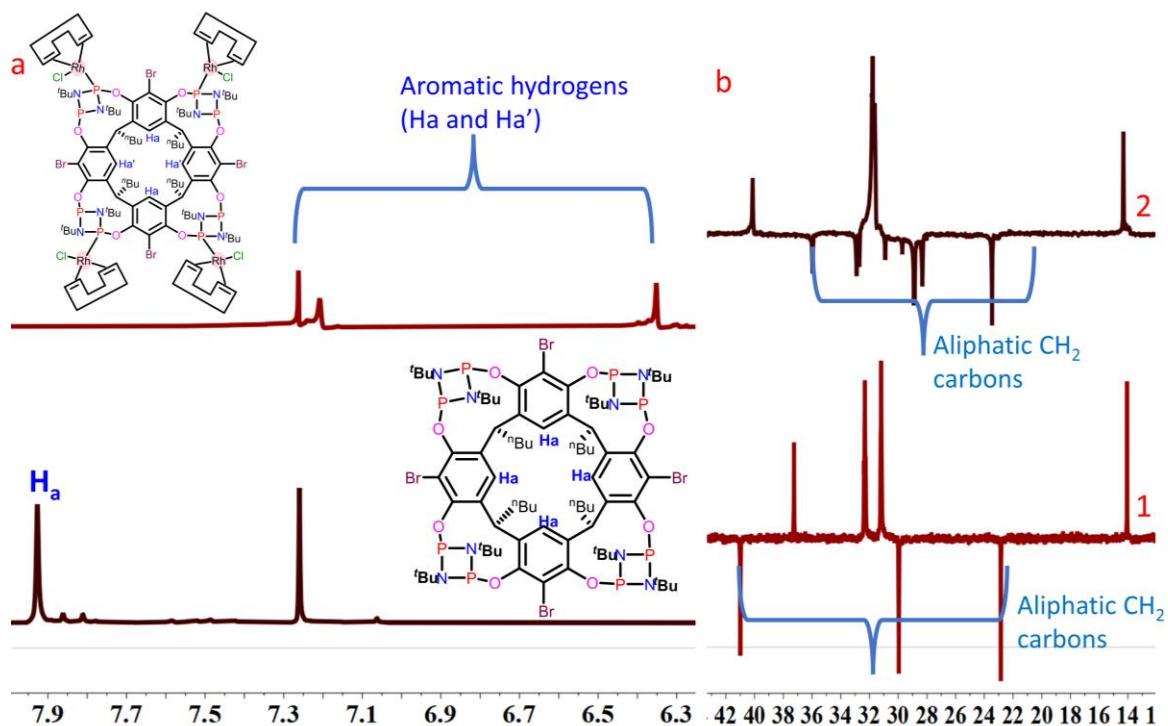


Fig. S19 Stack plots of ¹H (a: Left portion) and ¹³C-DEPT (b: Right portion) NMR spectra of **1** and **2** showing the changes after complex formation.