

## Electronic Supplementary Information

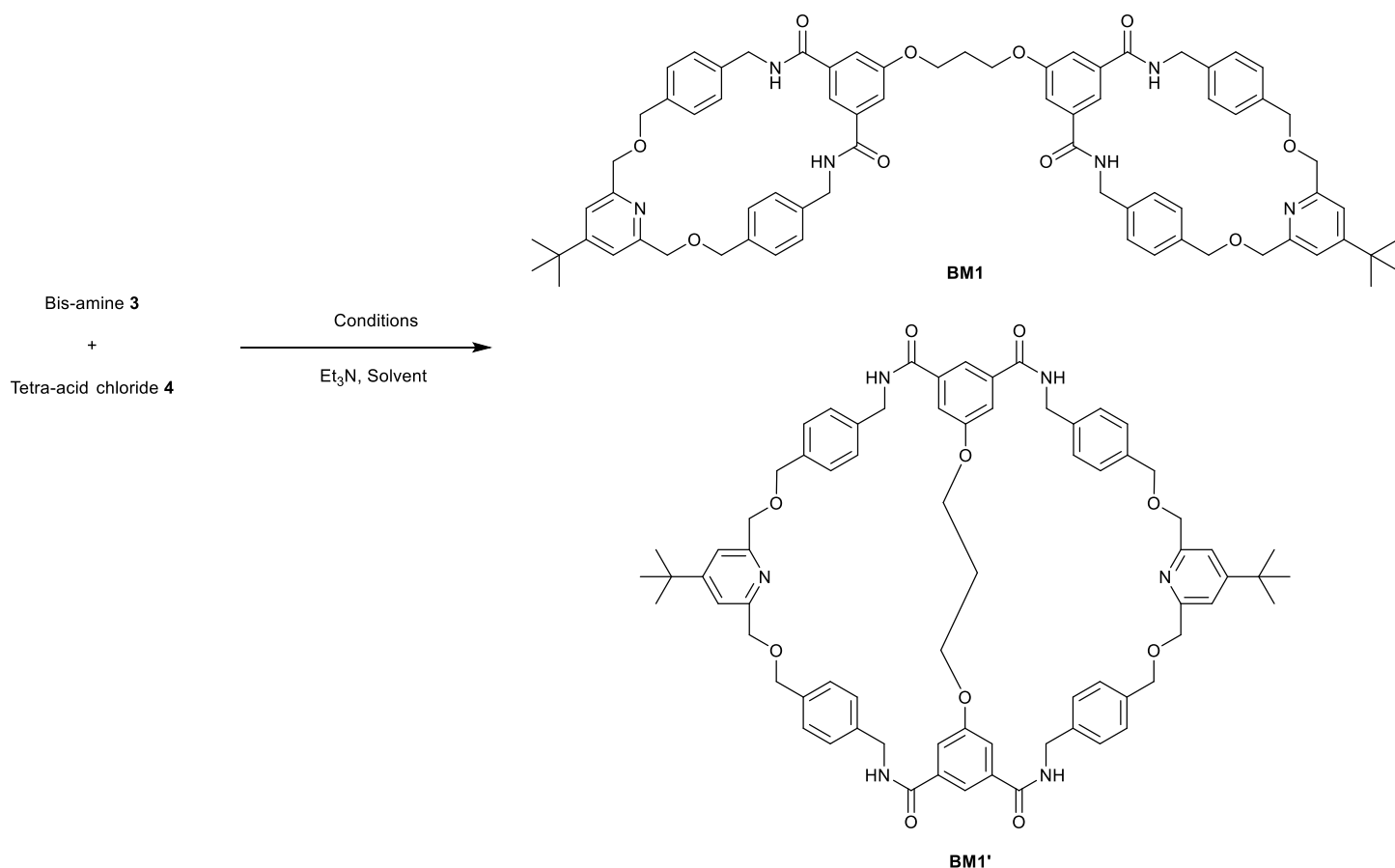
### Rapid Synthesis of Hydrogen Bond Templated Handcuff Rotaxanes

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## Part 1: Further details on attempted synthesis of bis-macrocycle **BM1**



Attempt	Solvent	Template	High <sup>a</sup> / Semi-high <sup>b</sup> dilution	(%) Yield <b>BM1</b>	(%) Yield <b>BM1'</b>
1	CH <sub>2</sub> Cl <sub>2</sub>	<b>5.CI</b>	Semi-high dilution	2	5
2	CH <sub>2</sub> Cl <sub>2</sub>	<b>5.CI</b>	High dilution	< 2	3
3	CH <sub>2</sub> Cl <sub>2</sub>	-	High dilution	-	-
4	CHCl <sub>3</sub>	<b>5.CI</b>	Semi-high dilution	< 2	6
5	CHCl <sub>3</sub>	-	High dilution	-	5

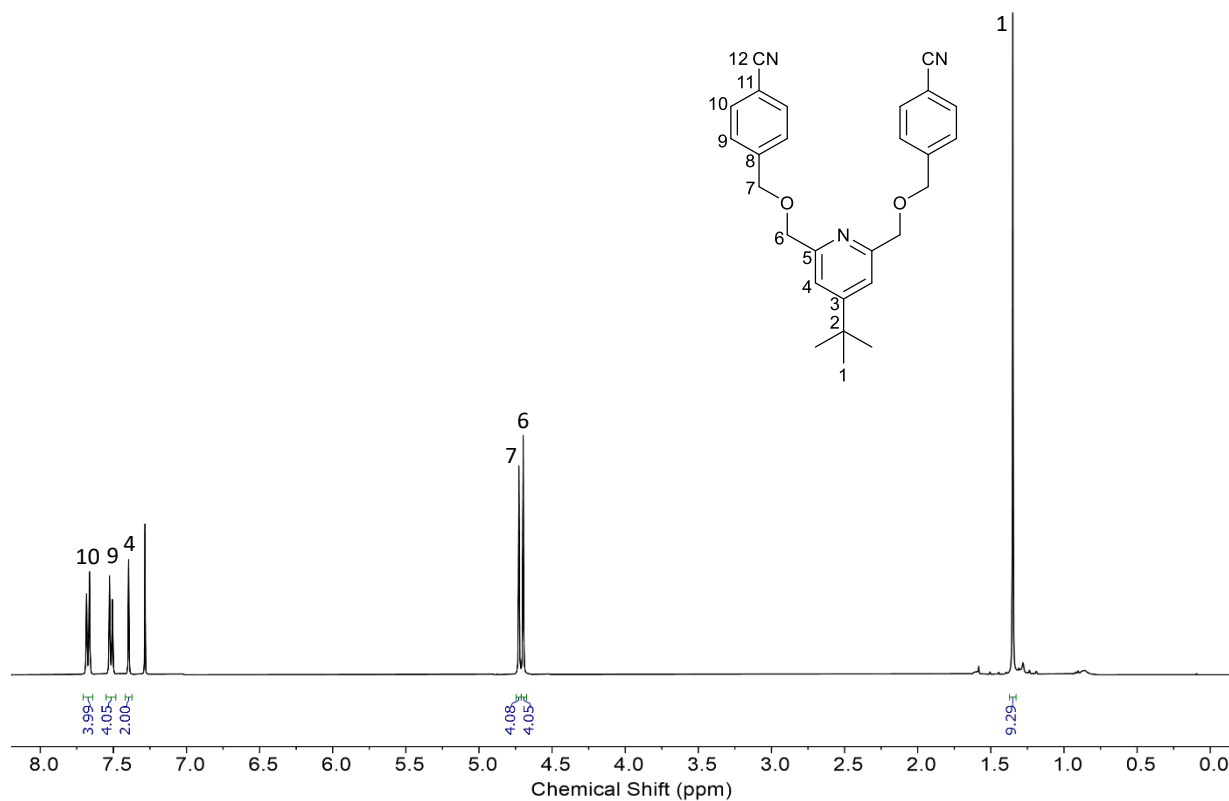
<sup>a</sup>High dilution attempts were conducted as follows: Et<sub>3</sub>N (5.0 eq) was dissolved in dry solvent (100 mL). Solutions of tetra-acid chloride **4** (1.0 eq) in dry solvent (100 mL) and bis-amine **3** (2.0 eq, with or without the presence of template **5.CI**, 2.0 eq) in dry solvent (100 mL) were added dropwise simultaneously to the above basic solution. The reaction was stirred for 16 hours, then washed with 10% citric acid (2 x 100 mL) and brine (1 x 100 mL). The organic layer was dried (MgSO<sub>4</sub>) and concentrated *in vacuo*. The crude material was then purified by silica gel column chromatography.

<sup>b</sup>Semi-high dilution attempts were conducted as follows: Bis-amine **3** (2.0 eq) and template **5.CI** (2.0 eq) were dissolved in dry solvent (60 mL). Et<sub>3</sub>N (5.0 eq) was added followed immediately by a dropwise solution of tetra-acid chloride **4** (1.0 eq) in dry solvent (30 mL). Following addition, the solution was allowed to stir for 4 hours. Then the reaction mixture was washed with 10% citric acid (2 x 100 mL) and brine (1 x 100 mL). The organic layer was dried (MgSO<sub>4</sub>) and concentrated *in vacuo*. The crude material was then purified by silica gel column chromatography.

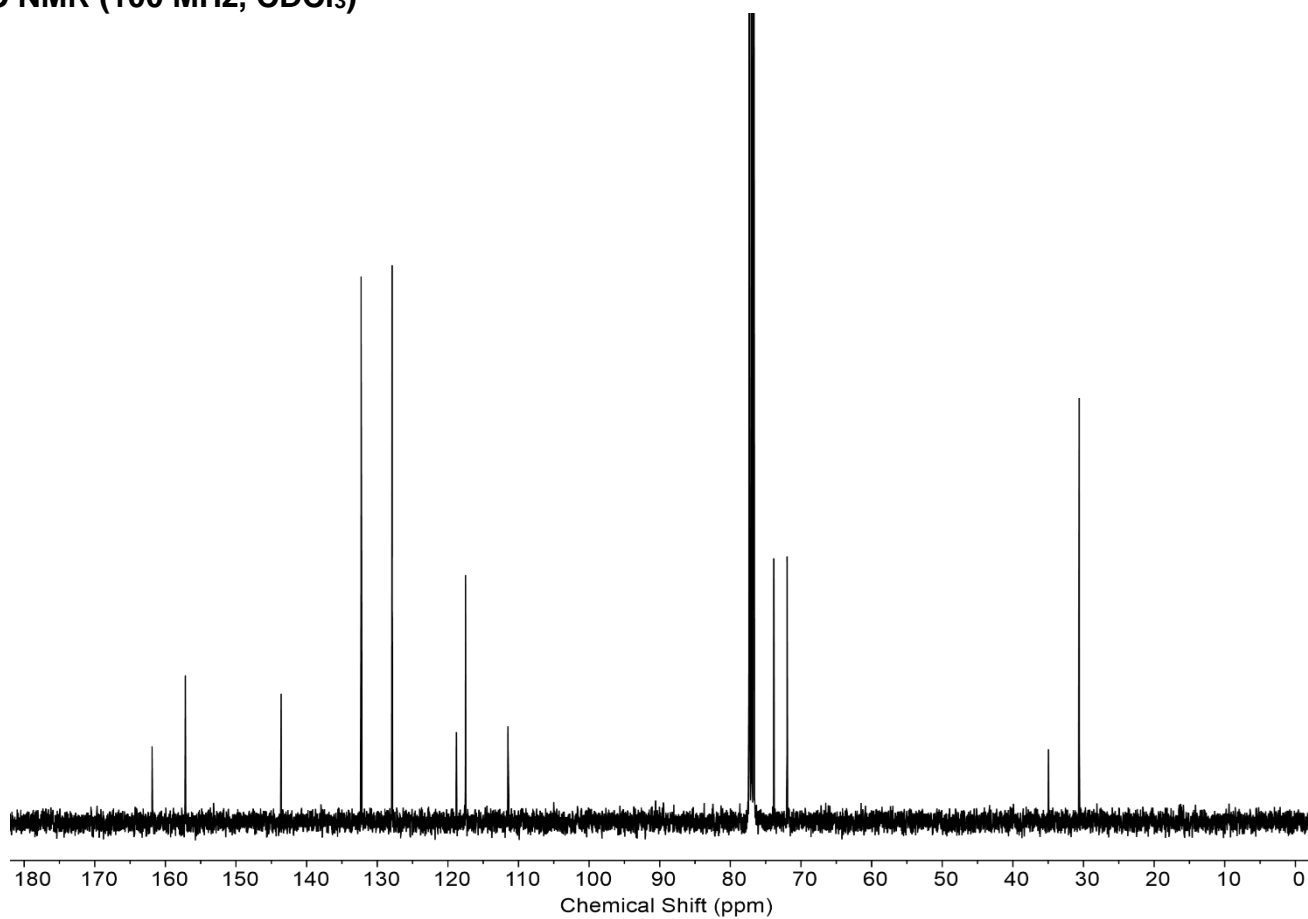
## Part 2: Characterization Spectra

### Bis-Nitrile 2

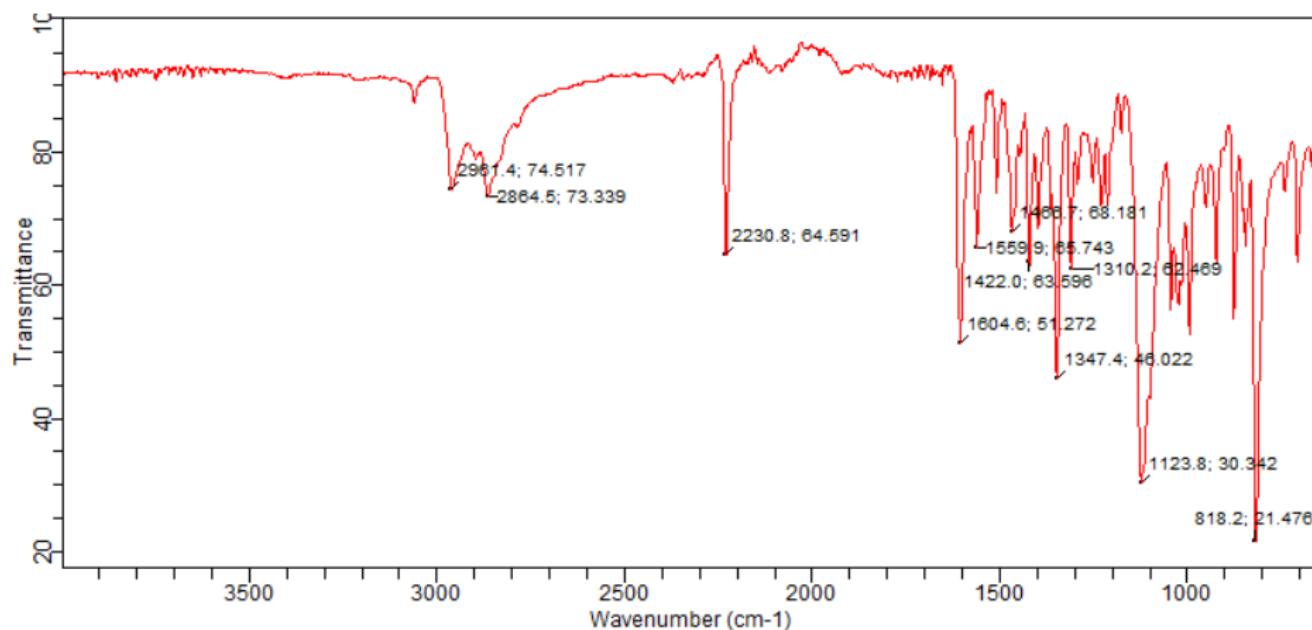
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ )



$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ )

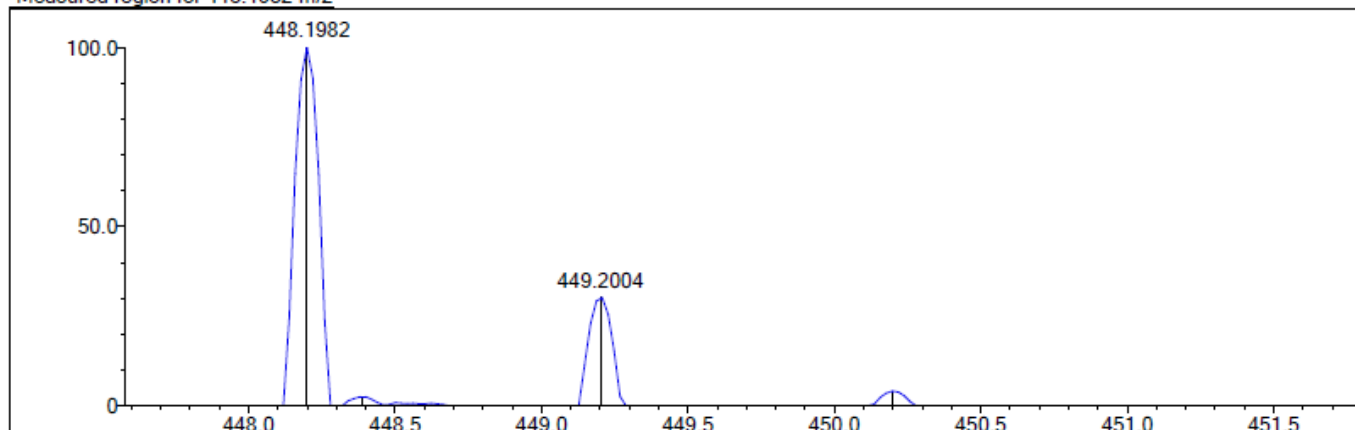


## IR Spectrum (neat)

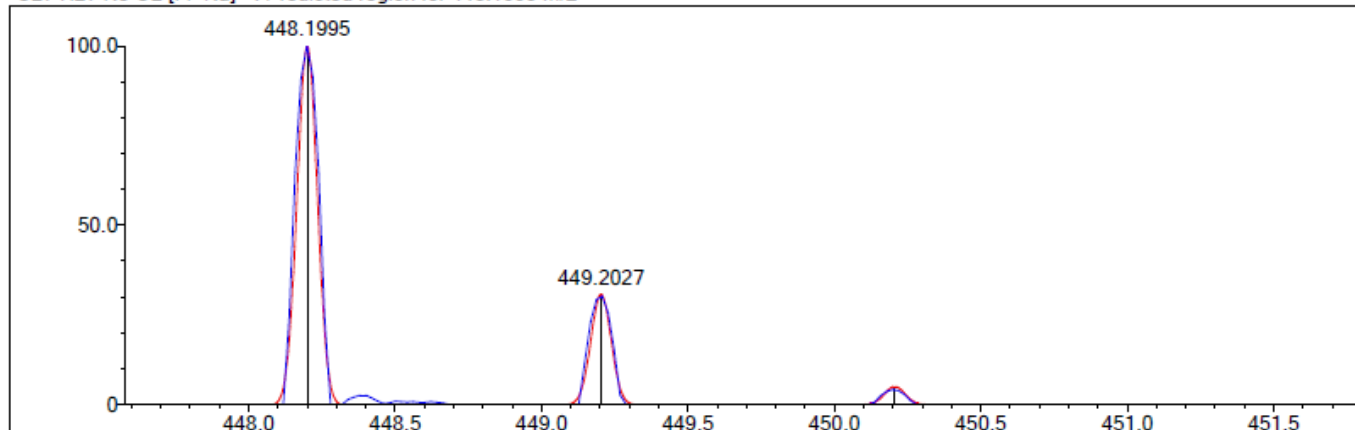


## HRMS (ES +ve)

Measured region for 448.1982 m/z



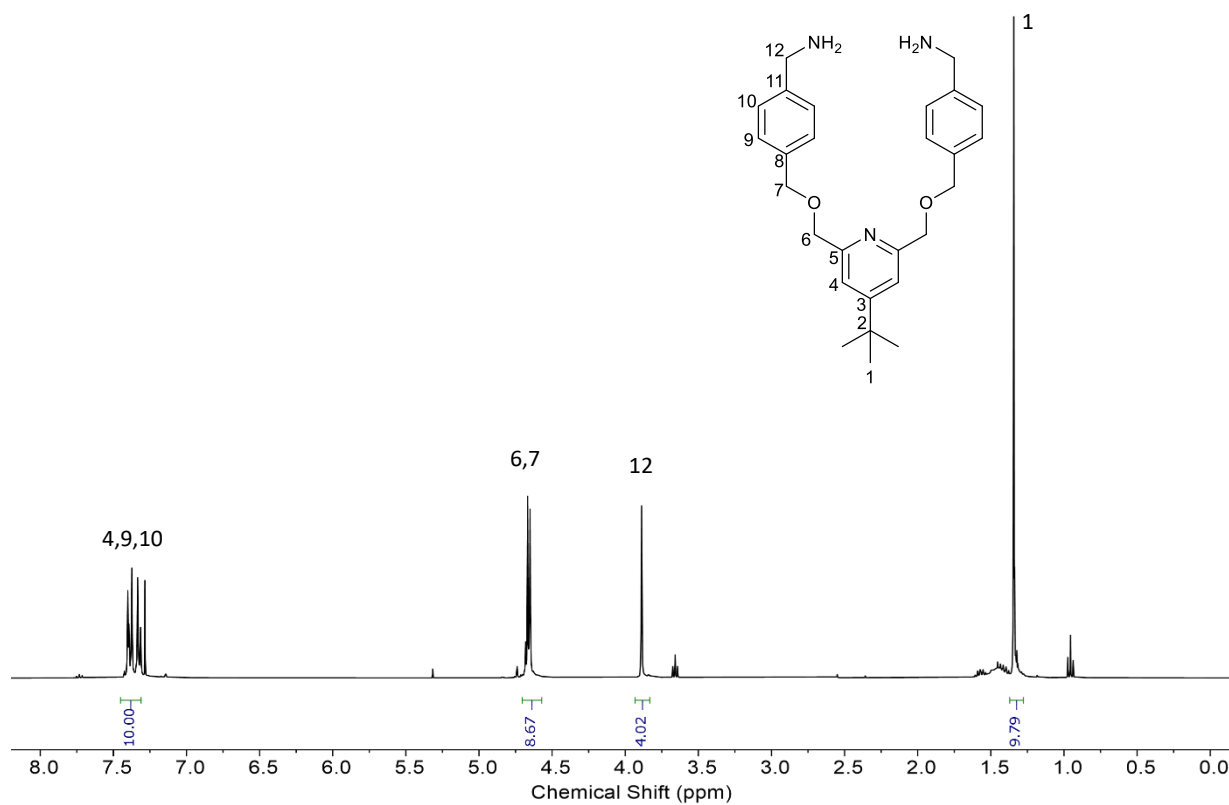
C<sub>27</sub>H<sub>27</sub>N<sub>3</sub>O<sub>2</sub> [M+Na]<sup>+</sup>: Predicted region for 448.1995 m/z



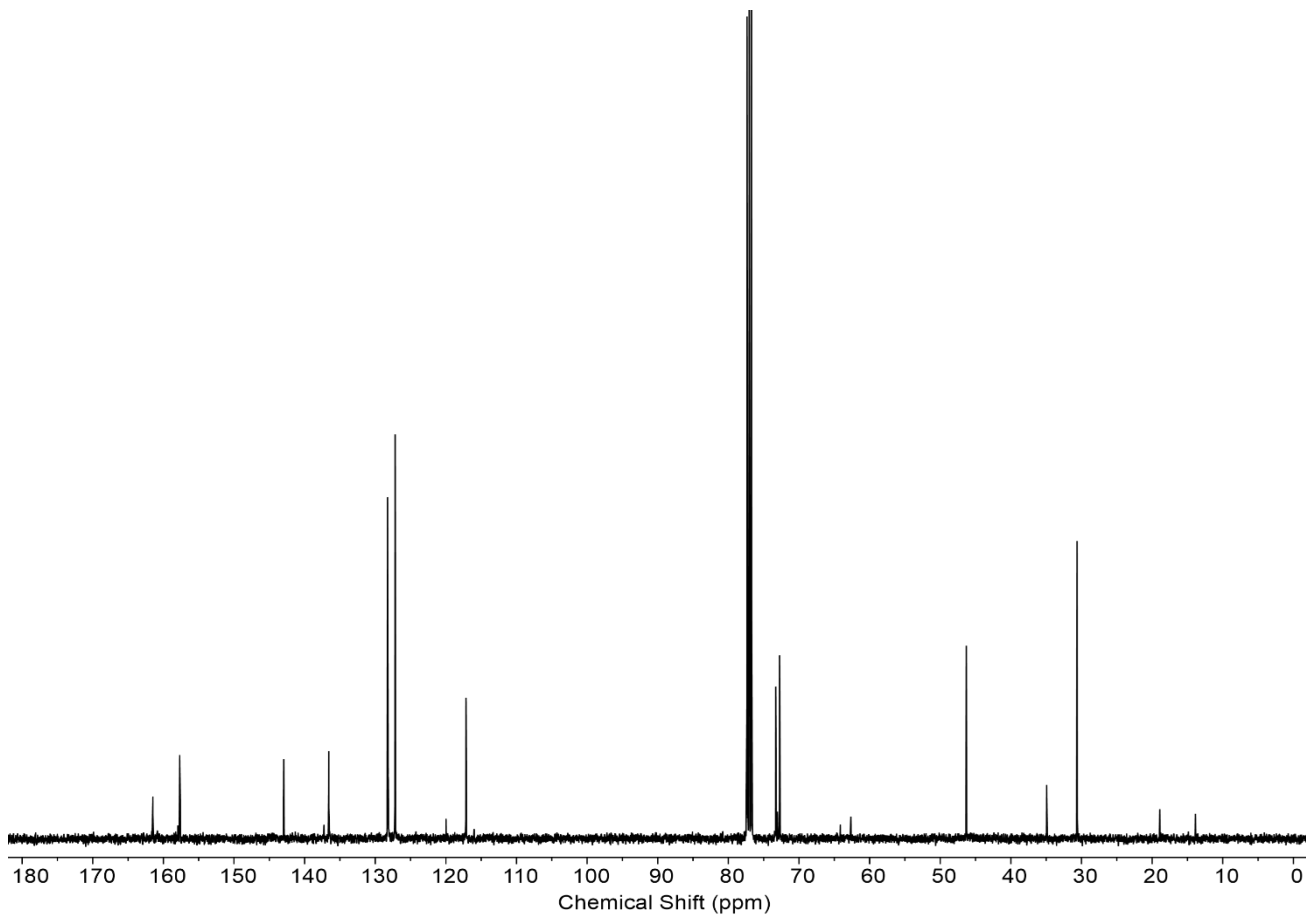
Rank	Score	Formula (M)	Ion	Meas. m/z	Pred. m/z	Df. (mDa)	Df. (ppm)	Iso	DBE
1	95.25	C <sub>27</sub> H <sub>27</sub> N <sub>3</sub> O <sub>2</sub>	[M+Na] <sup>+</sup>	448.1982	448.1995	-1.3	-2.90	100.00	16.0

## Bis-Amine 3

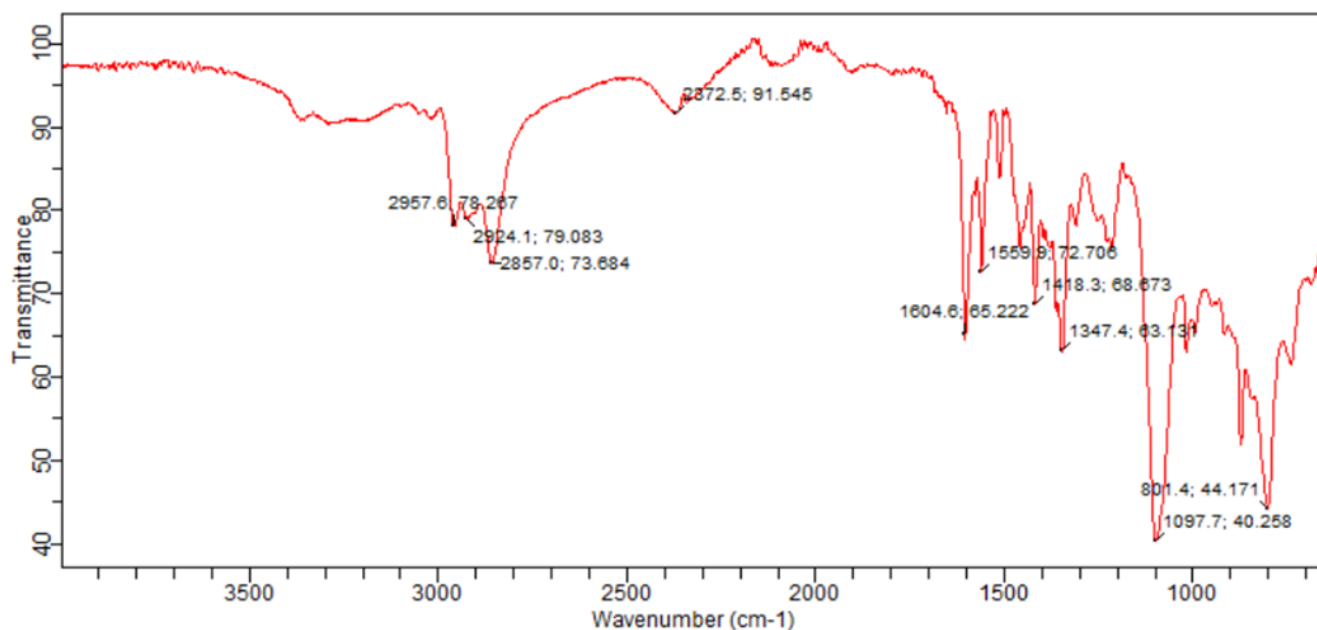
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ )



$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ )

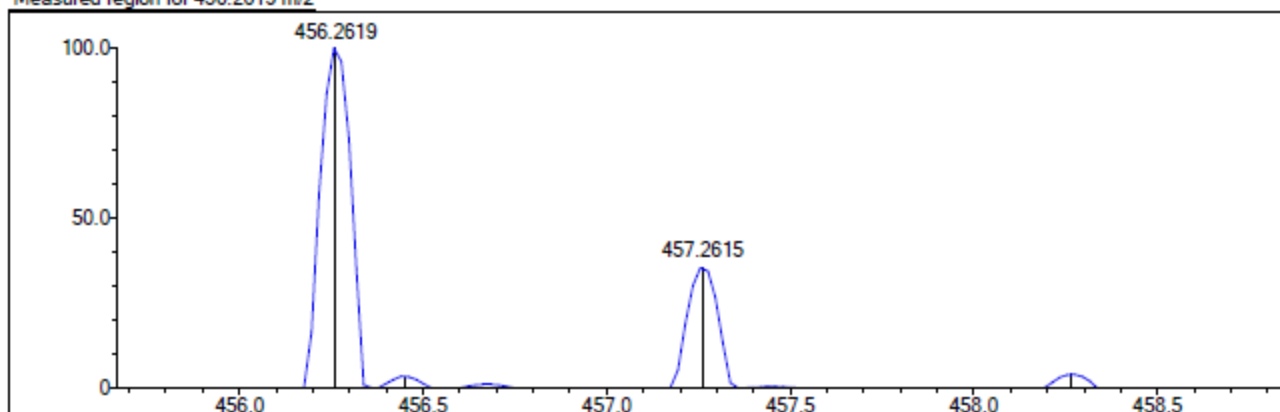


## IR Spectrum (neat)

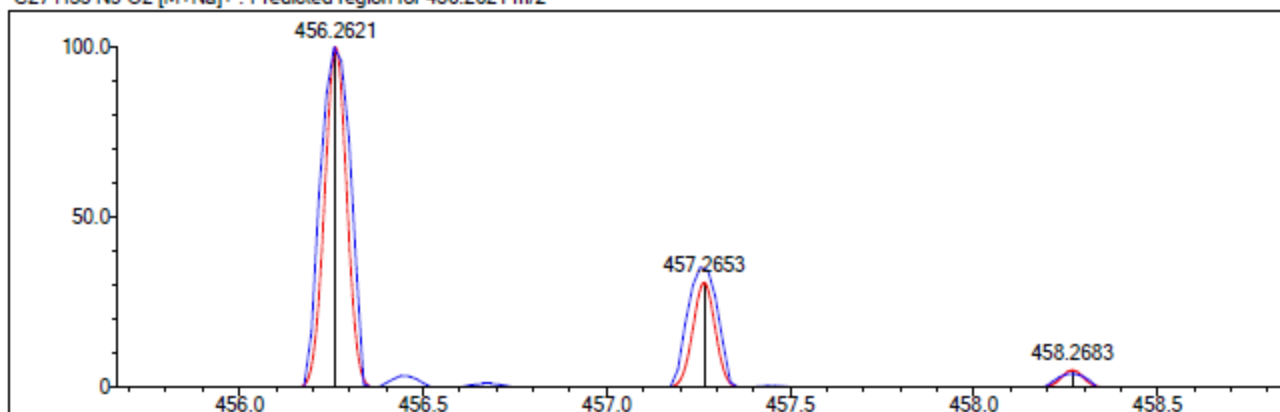


## HRMS (ES +ve)

Measured region for 456.2619 m/z



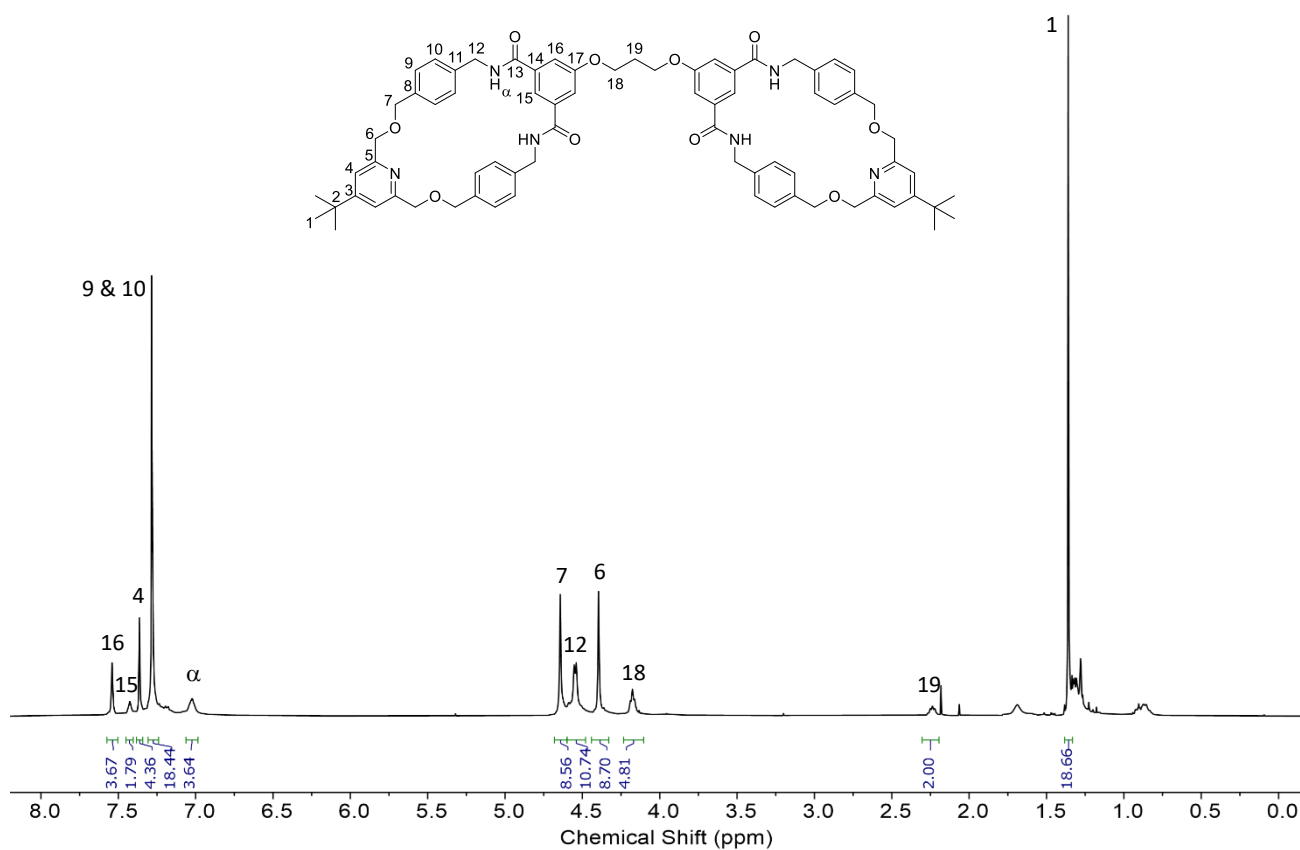
C<sub>27</sub>H<sub>35</sub>N<sub>3</sub>O<sub>2</sub> [M+Na]<sup>+</sup> : Predicted region for 456.2621 m/z



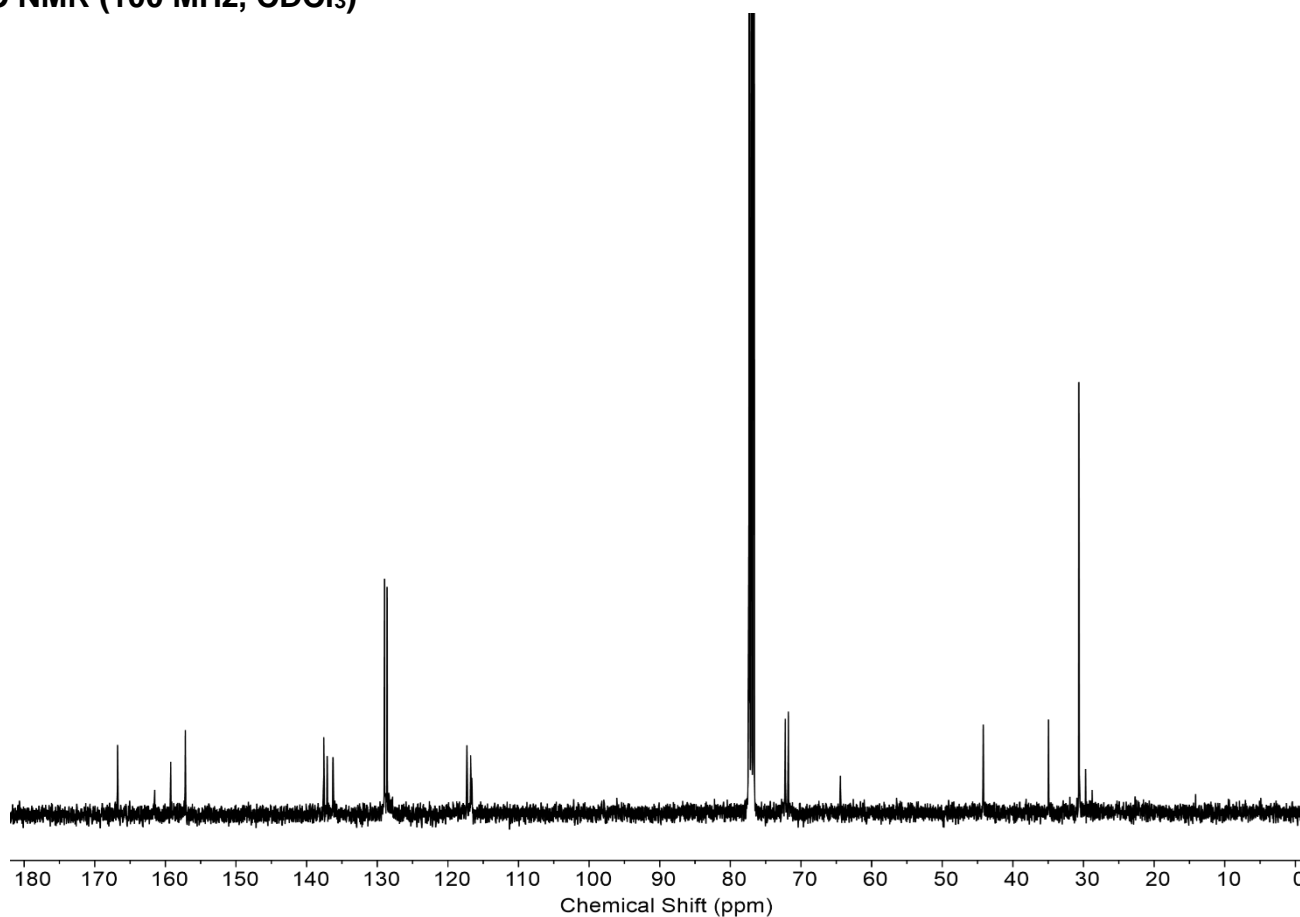
Rank	Score	Formula (M)	Ion	Meas. m/z	Pred. m/z	Df. (mDa)	Df. (ppm)	Iso	DBE
1	95.60	C <sub>27</sub> H <sub>35</sub> N <sub>3</sub> O <sub>2</sub>	[M+Na] <sup>+</sup>	456.2619	456.2621	-0.2	-0.44	95.60	12.0

# Bis-Macrocyclic BM1

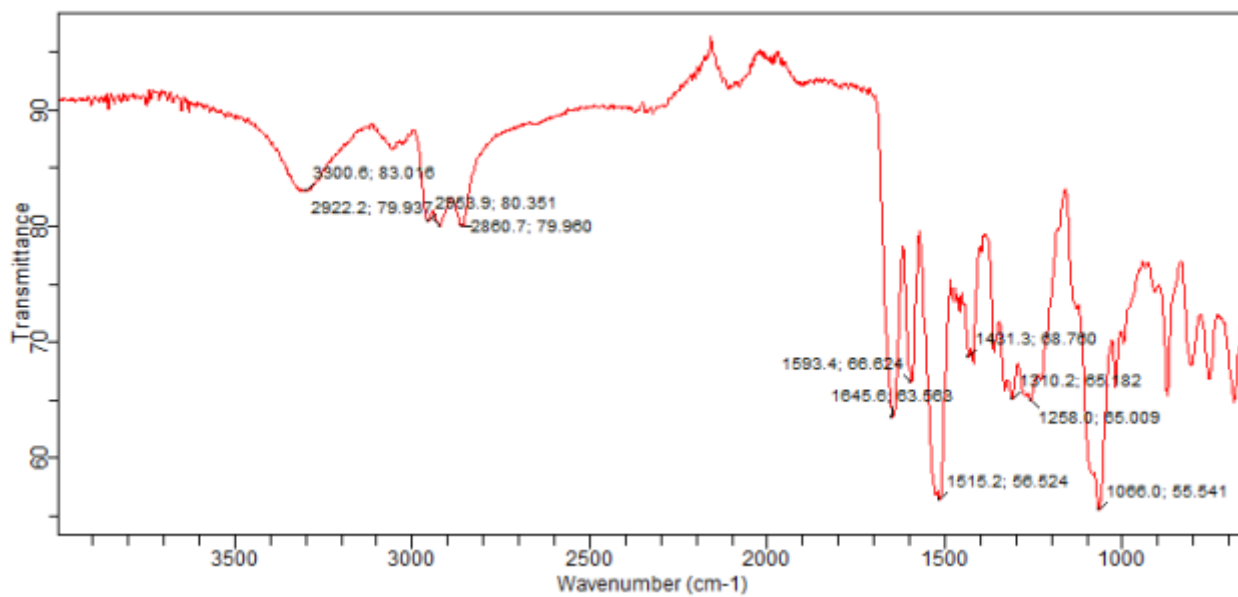
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ )



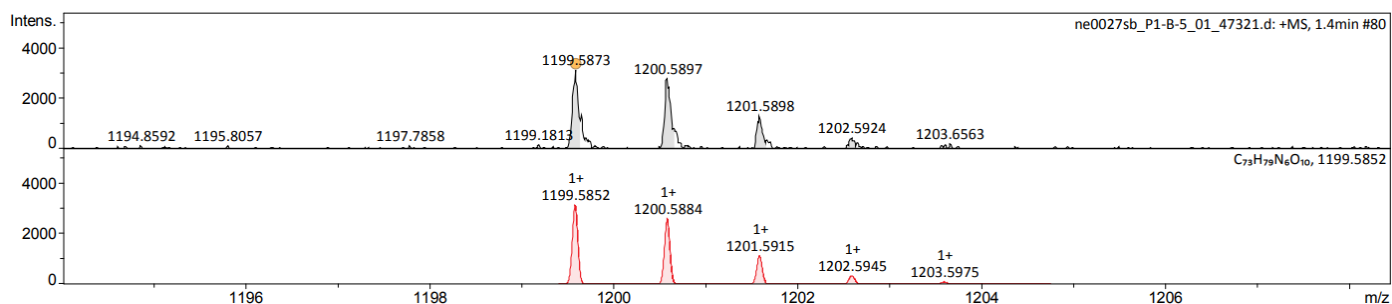
$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ )



## IR Spectrum (neat)



## HRMS (ES +ve)

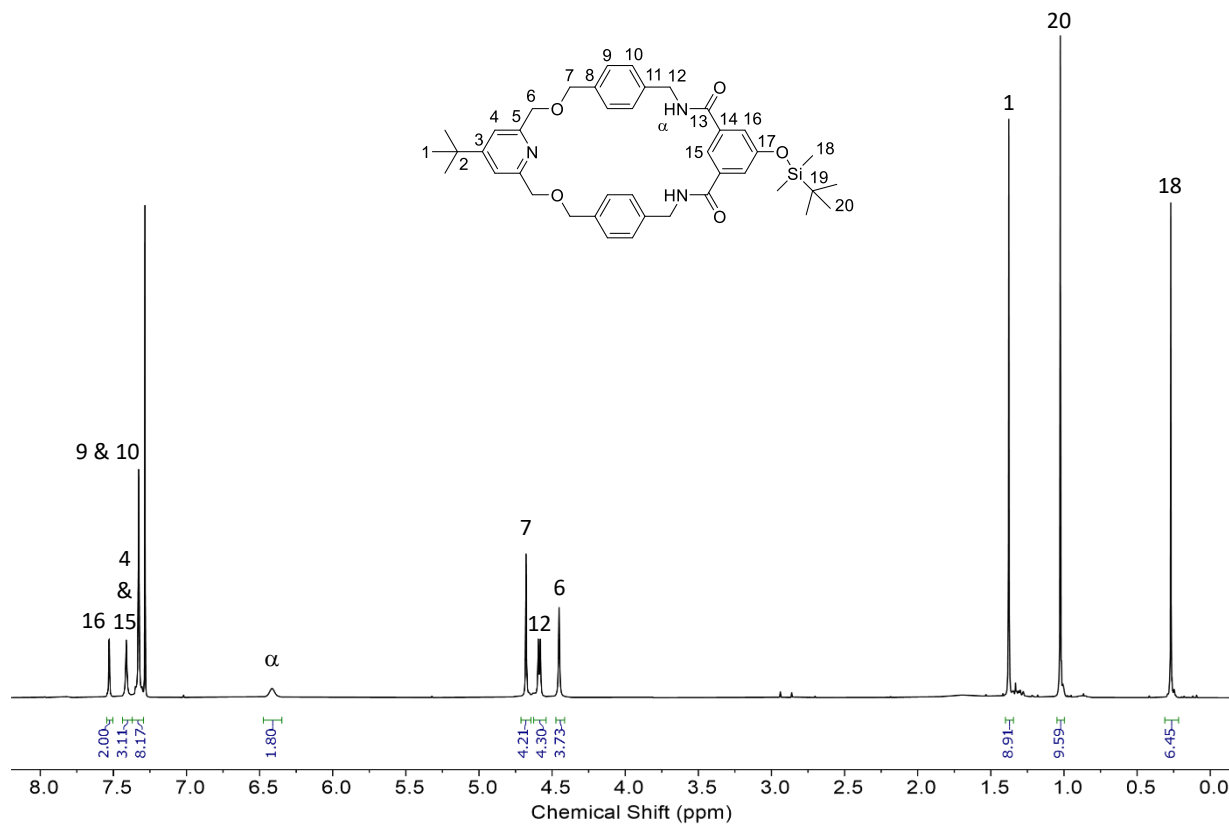


Meas. m/z	#	Ion Formula	m/z	err [ppm]	err [mDa]	mSigma	Mean err [ppm]
1199.5873	1	C <sub>73</sub> H <sub>79</sub> N <sub>6</sub> O <sub>10</sub>	1199.5852	-1.7	-2.0	45.1	-0.4

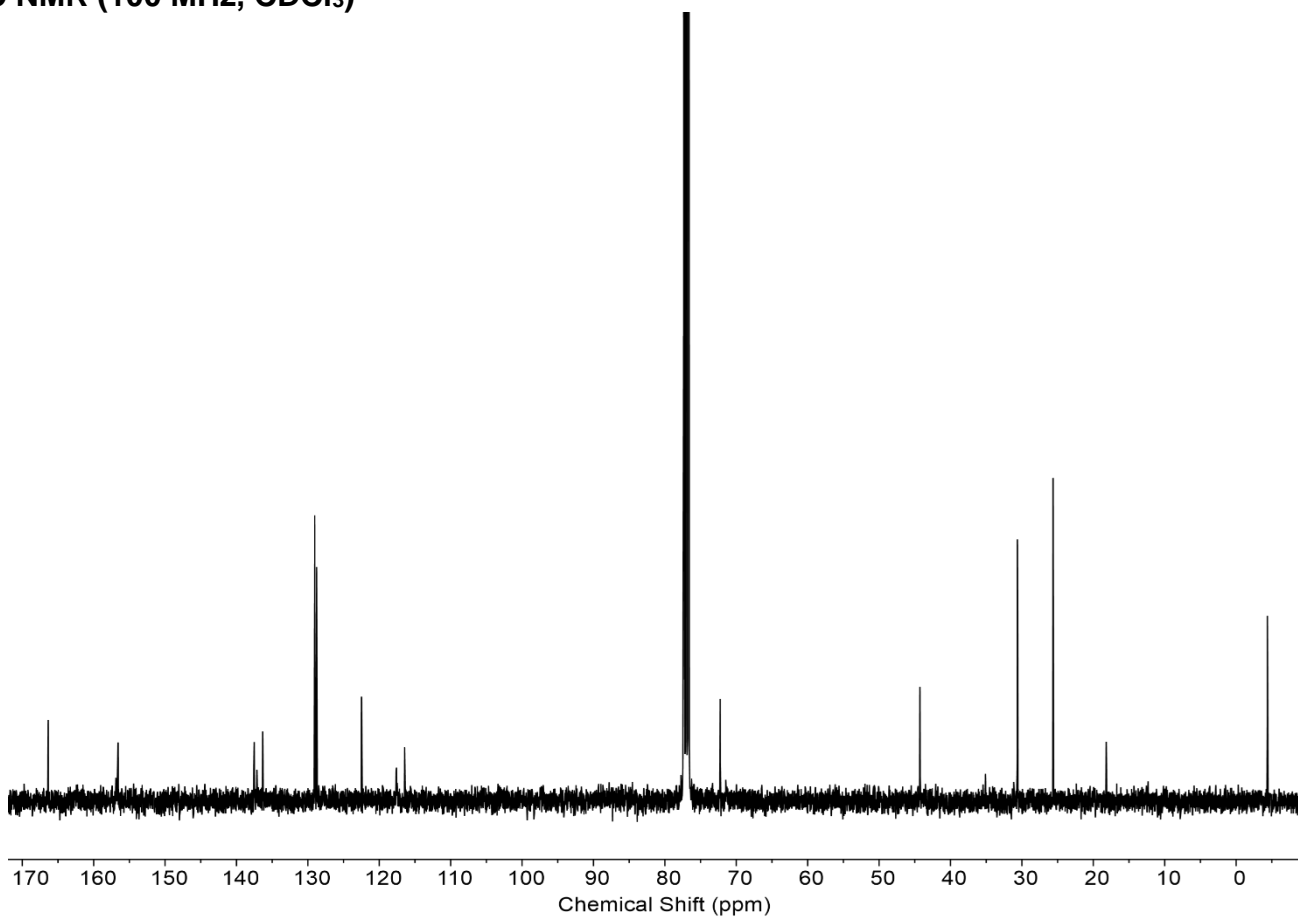


# TBDMS Protected Macrocycle 7

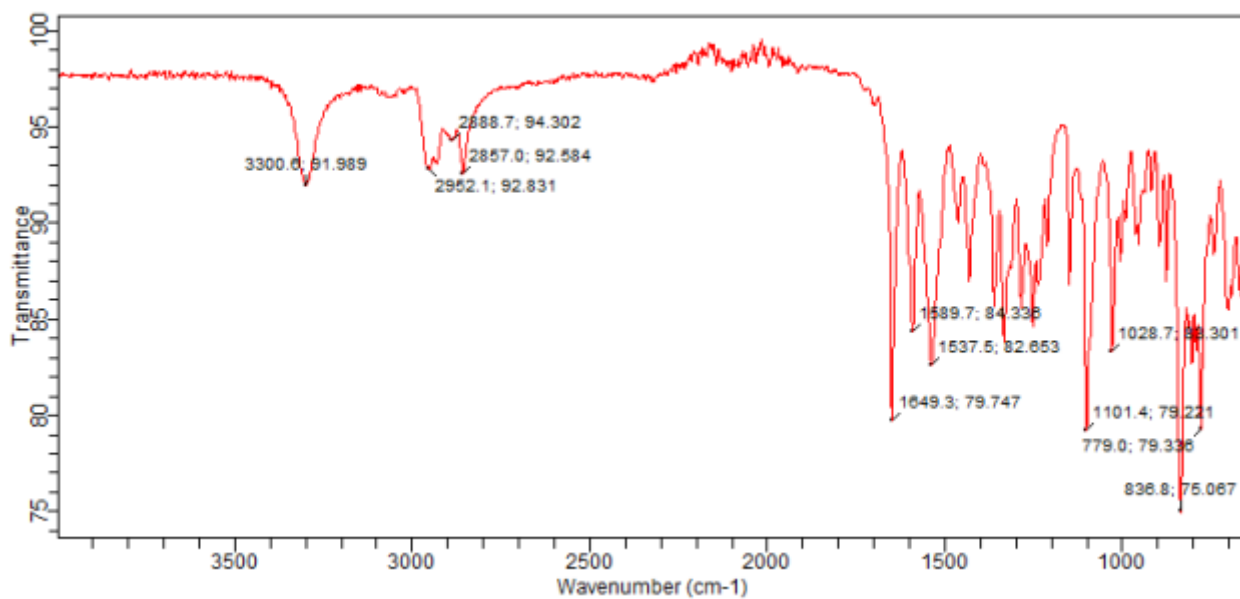
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ )



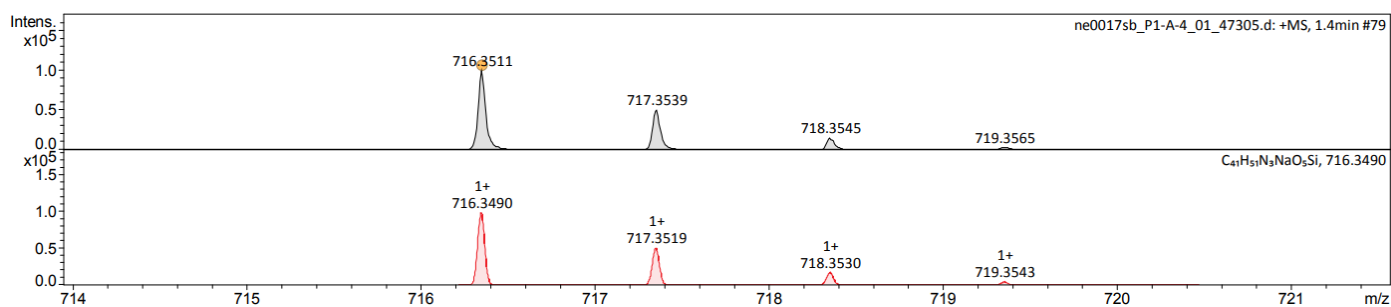
$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ )



## IR Spectrum (neat)



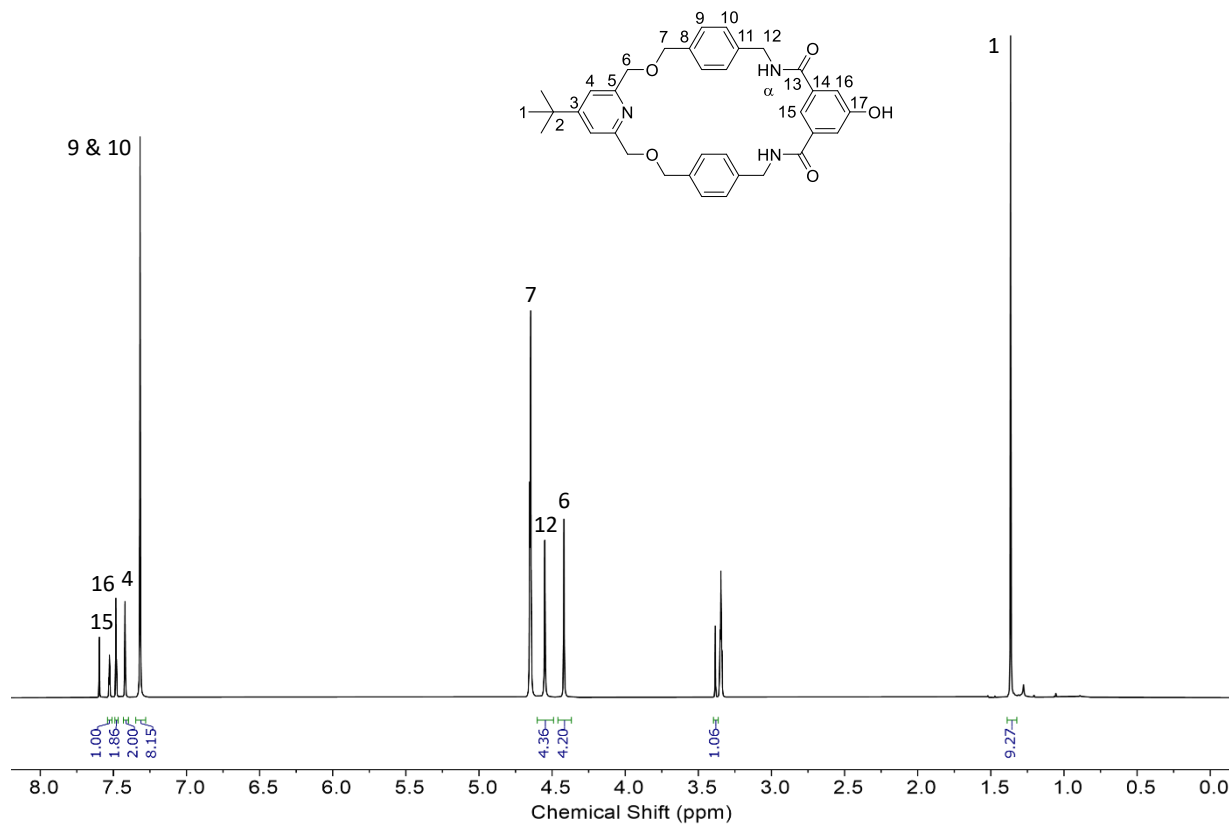
## HRMS (ES +ve)



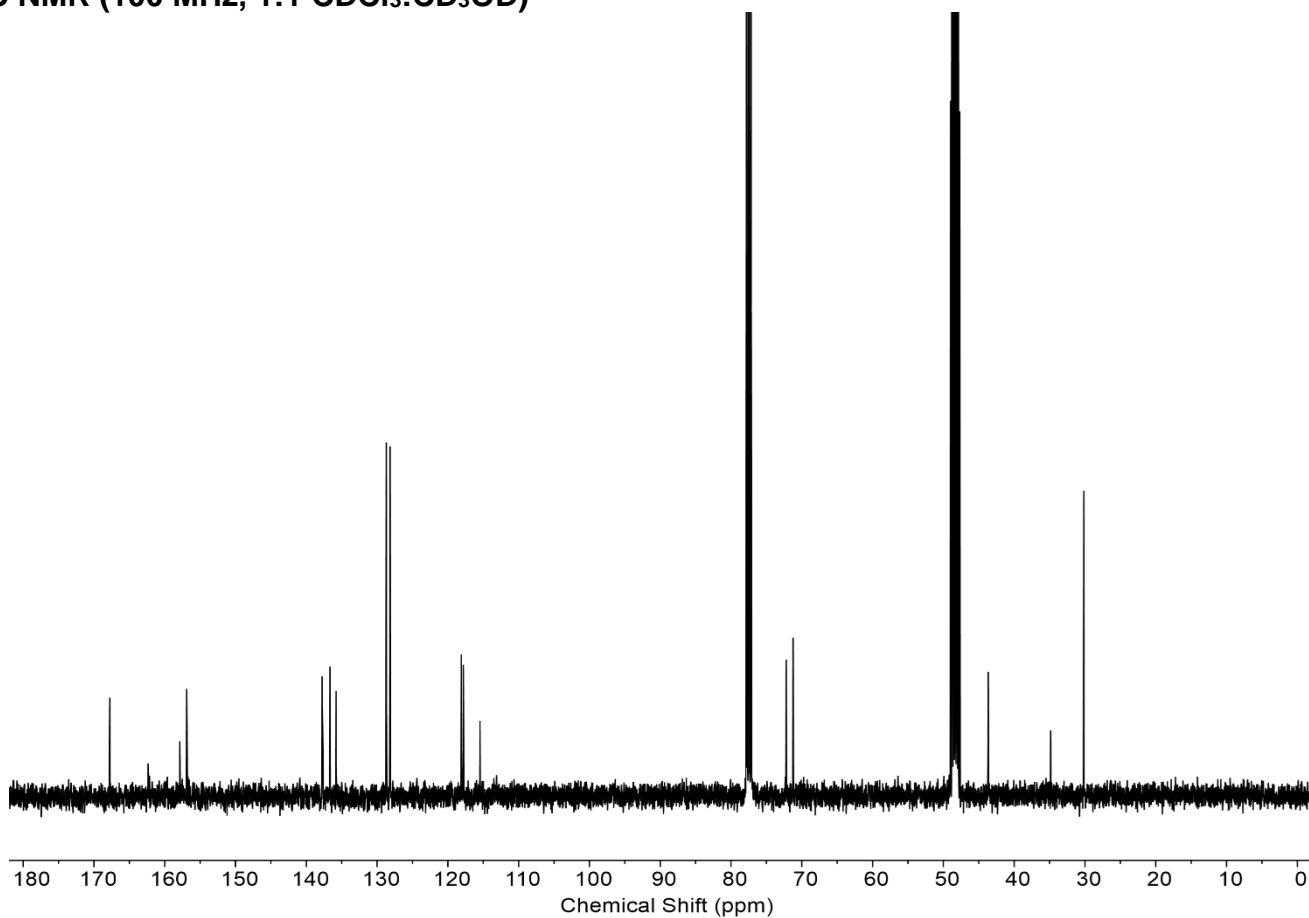
Meas. m/z	#	Ion Formula	m/z	err [ppm]	err [mDa]	mSigma	Mean err [ppm]
694.3684	1	C <sub>41</sub> H <sub>52</sub> N <sub>3</sub> O <sub>5</sub> Si	694.3671	-1.9	-1.3	6.0	-3.9
716.3511	1	C <sub>41</sub> H <sub>51</sub> N <sub>3</sub> NaO <sub>5</sub> Si	716.3490	-2.9	-2.1	10.2	-3.5

# Phenol Macrocycle 8

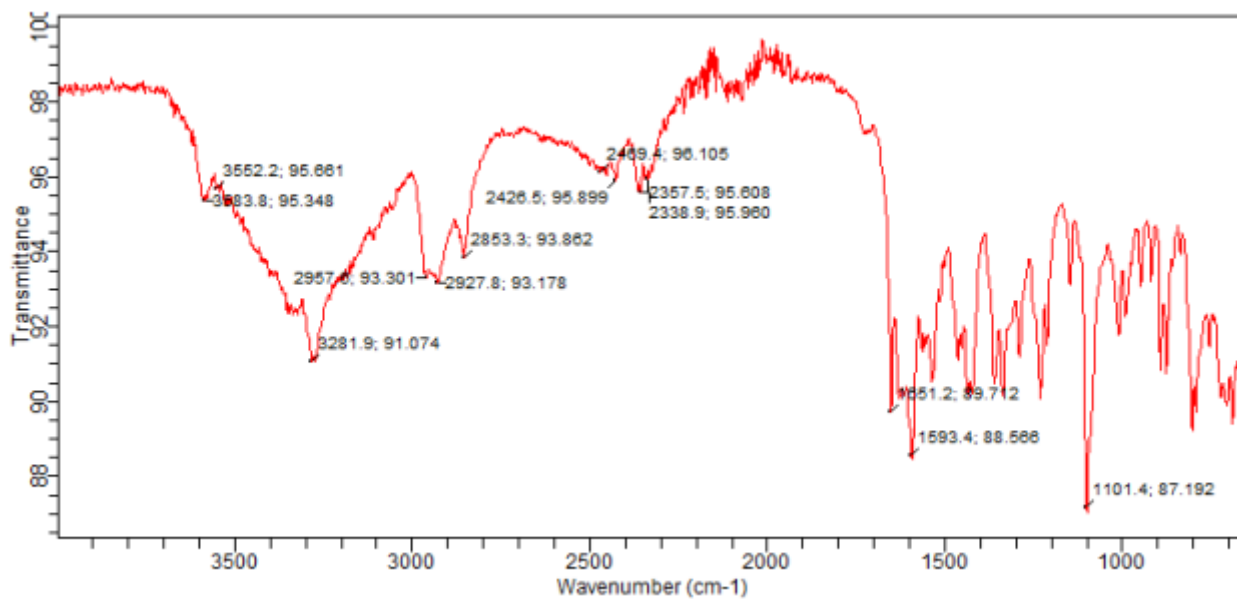
$^1\text{H}$  NMR (400 MHz, 1:1  $\text{CDCl}_3:\text{CD}_3\text{OD}$ )



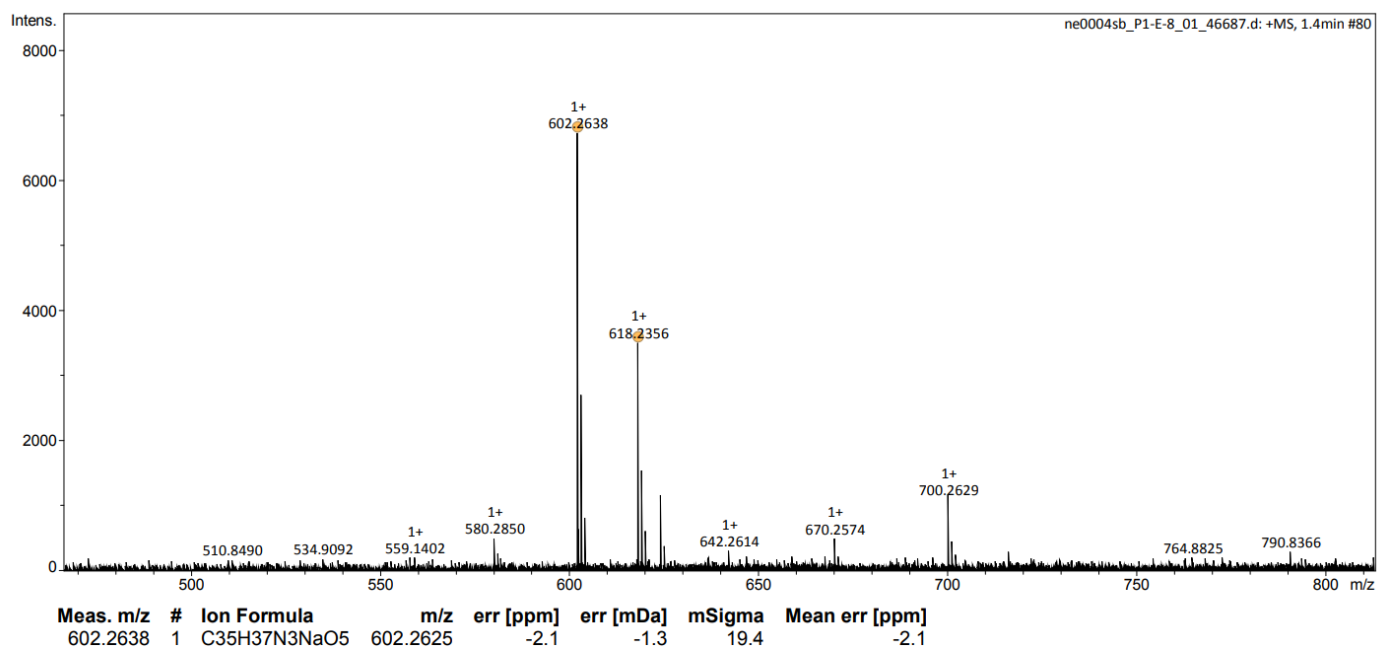
$^{13}\text{C}$  NMR (100 MHz, 1:1  $\text{CDCl}_3:\text{CD}_3\text{OD}$ )



## IR Spectrum (neat)

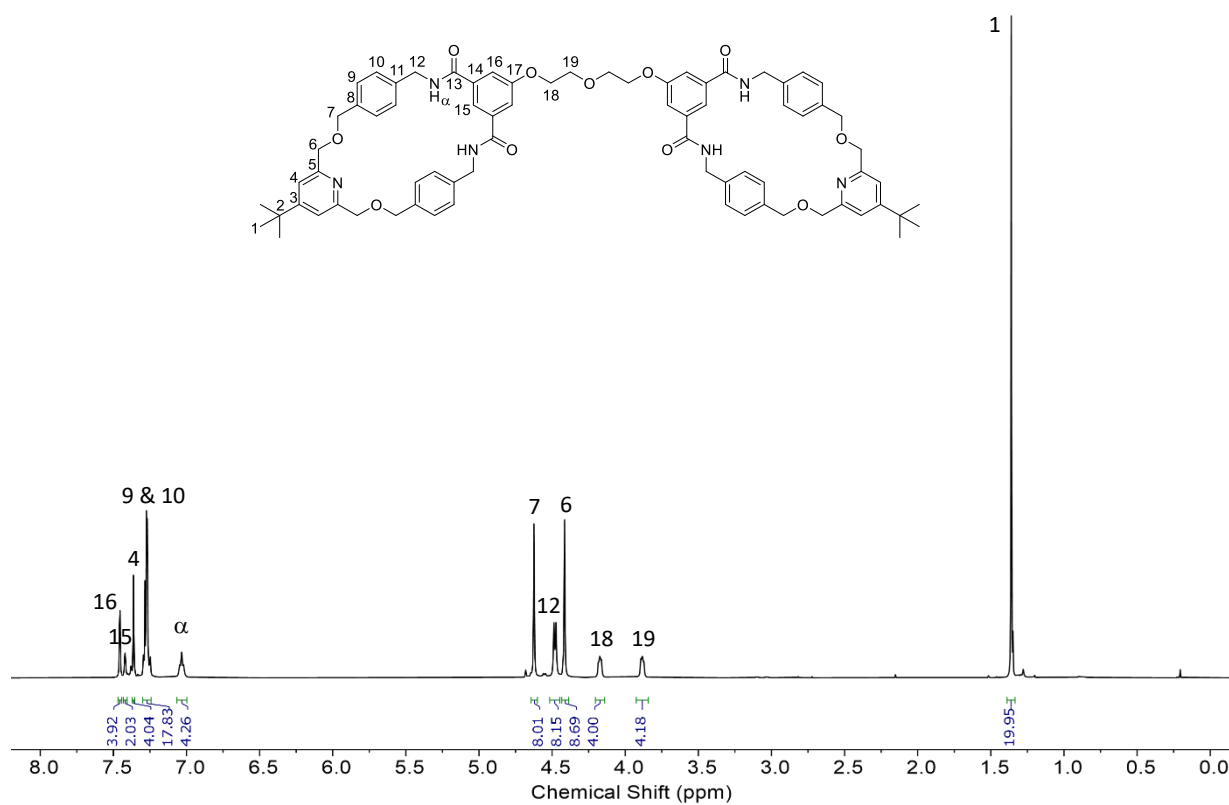


## HRMS (ES +ve)

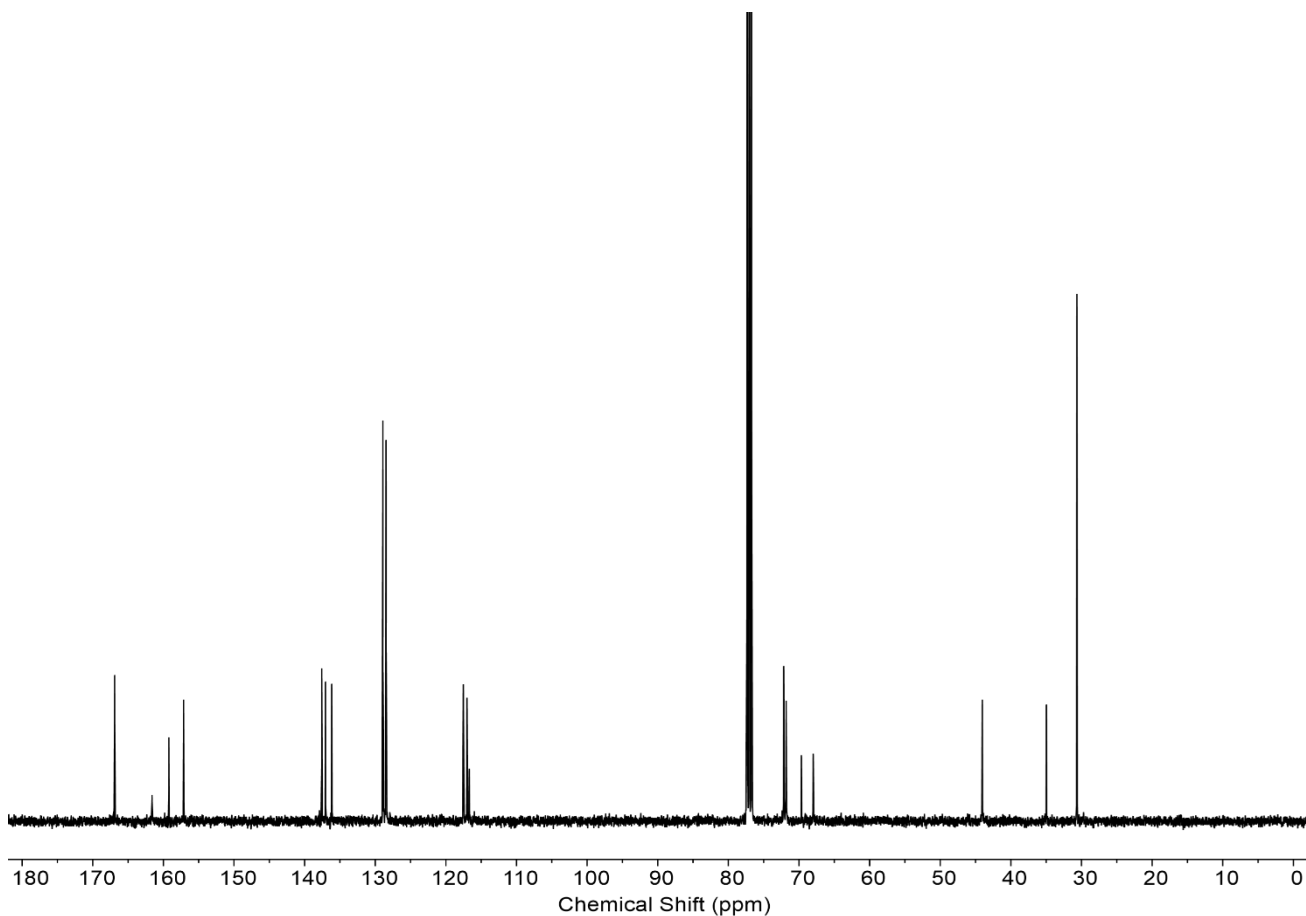


## Bis-Macrocycle BM2

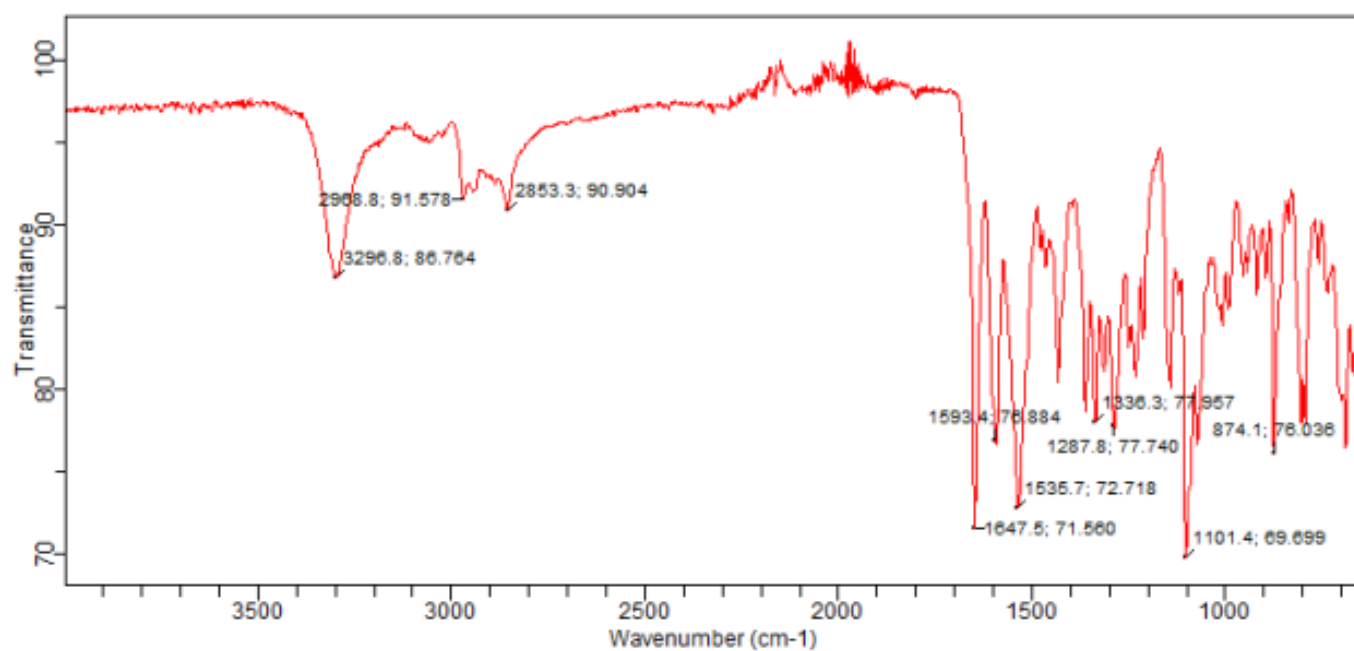
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ )



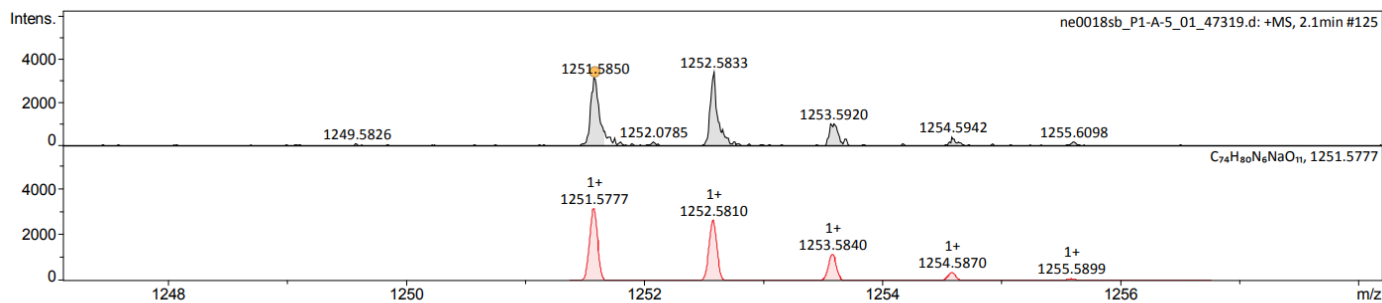
$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ )



## IR Spectrum (neat)



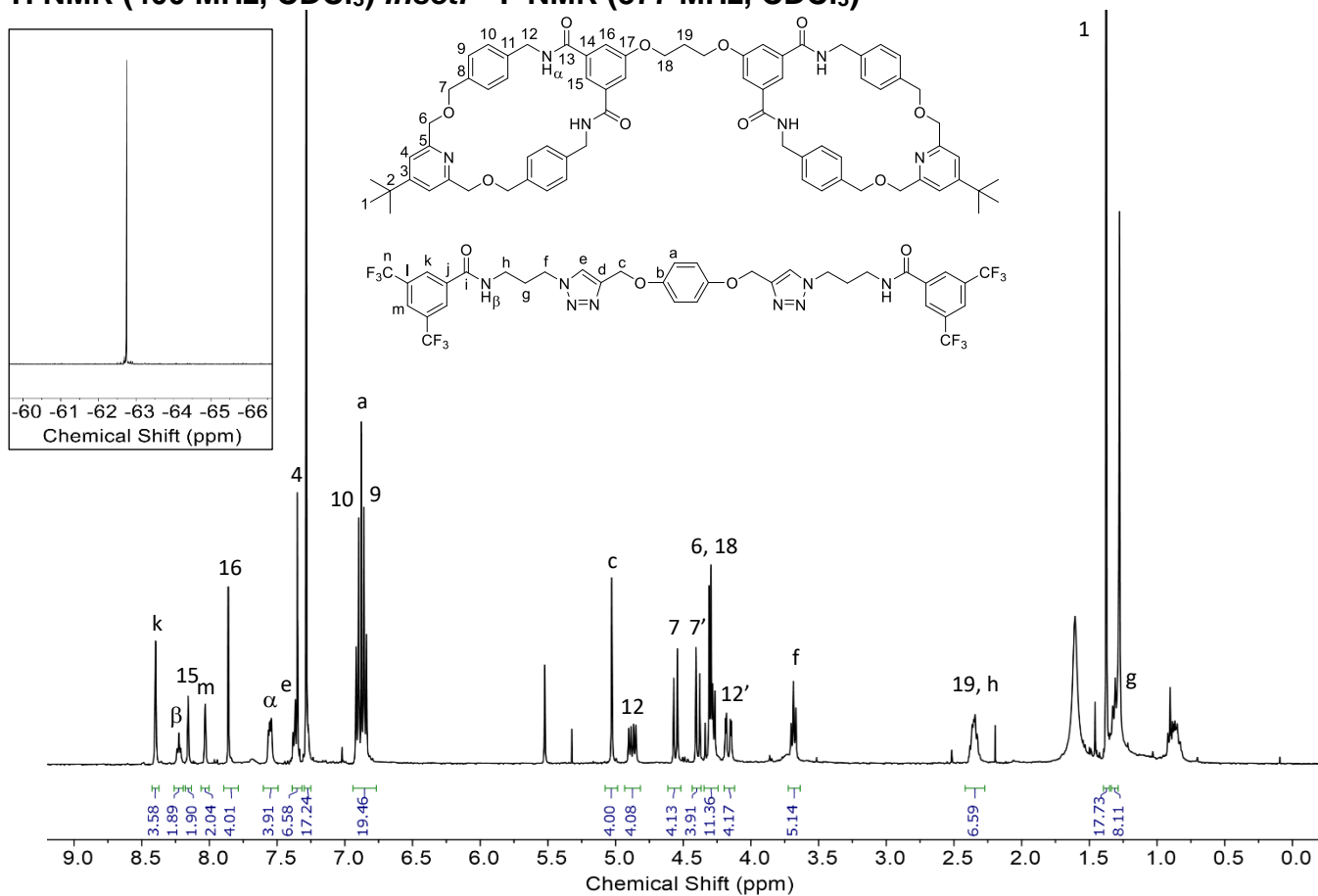
## HRMS (ES +ve)



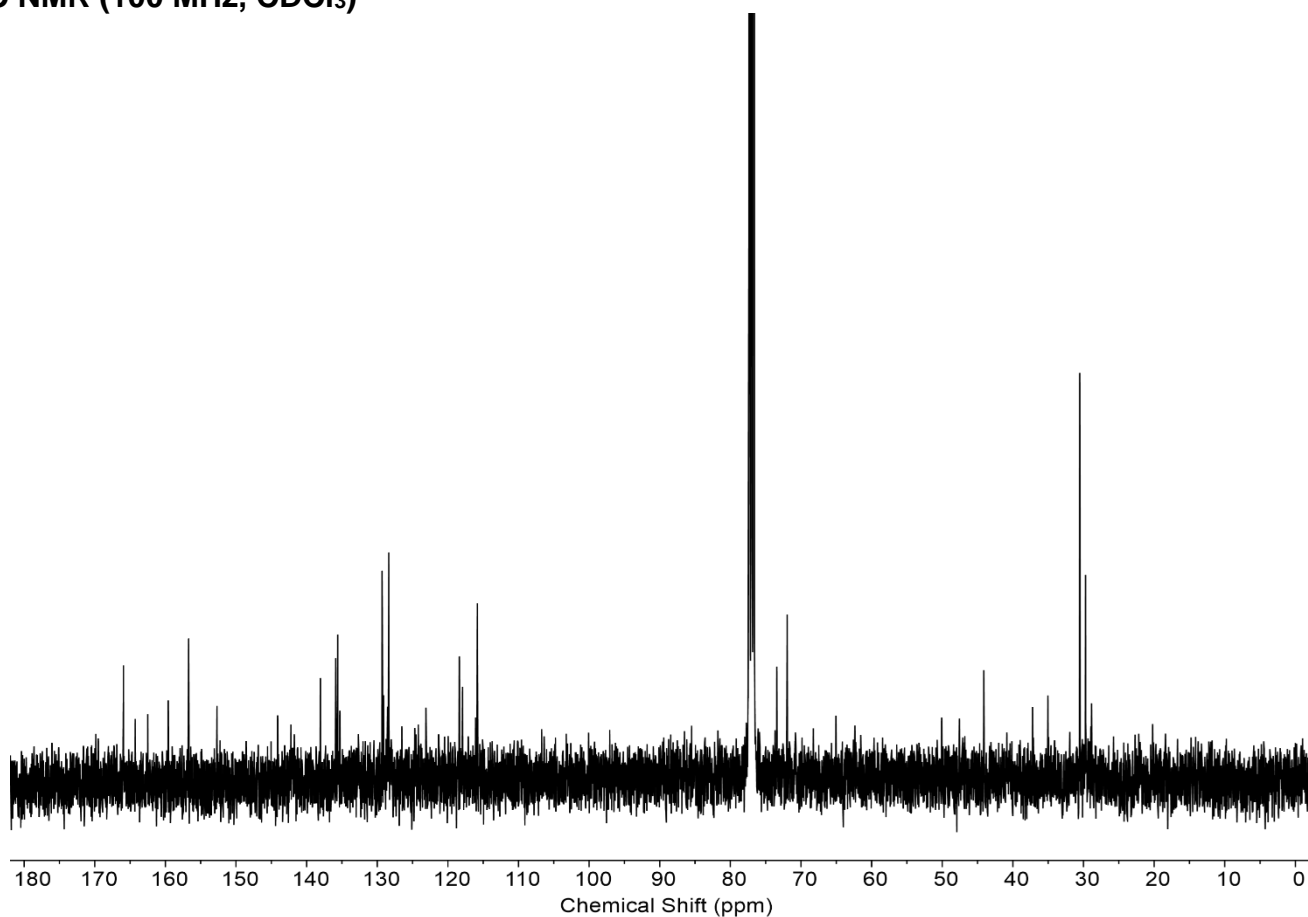
Meas. m/z	#	Ion Formula	m/z	err [ppm]	err [mDa]	mSigma	Mean err [ppm]
1229.5958	1	C <sub>74</sub> H <sub>81</sub> N <sub>6</sub> O <sub>11</sub>	1229.5958	-0.0	-0.0	93.3	-2.5
1251.5850	1	C <sub>74</sub> H <sub>80</sub> N <sub>6</sub> NaO <sub>11</sub>	1251.5777	-5.8	-7.3	88.1	-4.2

# Handcuff Rotaxane HR1

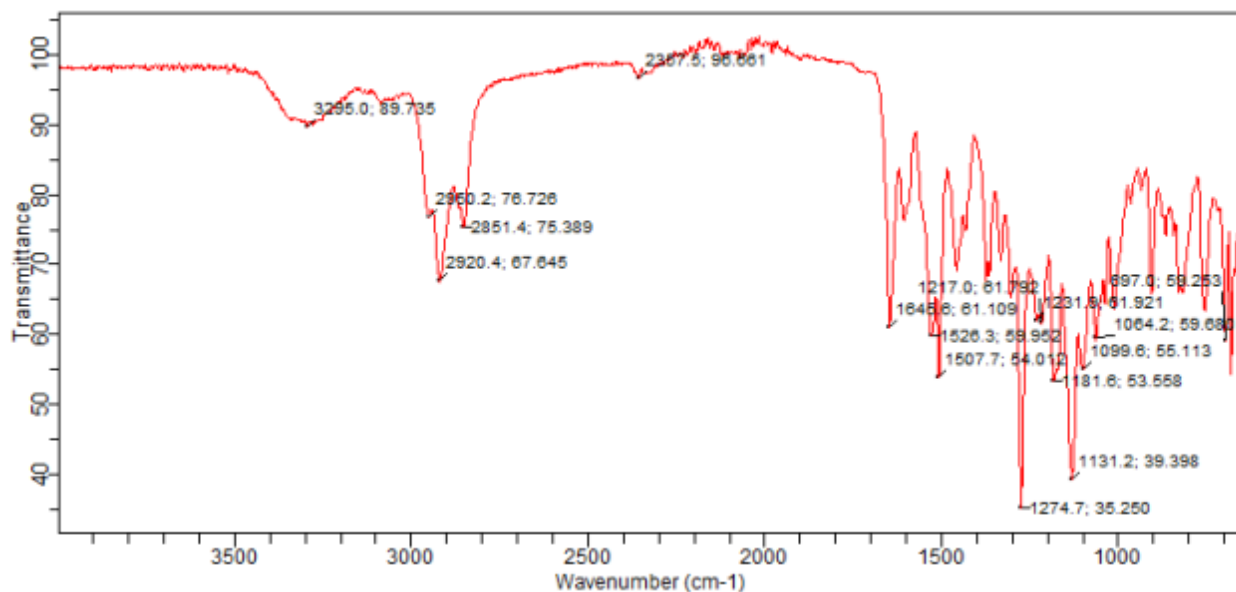
**<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) Inset: <sup>19</sup>F NMR (377 MHz, CDCl<sub>3</sub>)**



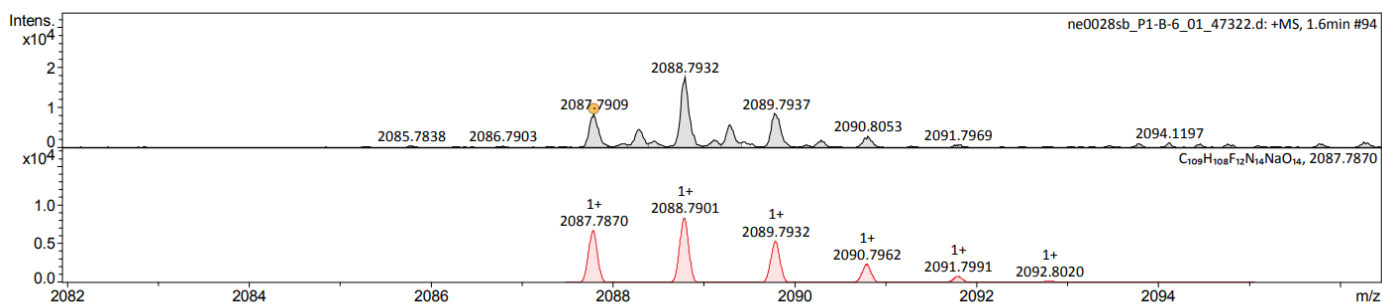
**<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)**



## IR Spectrum (neat)

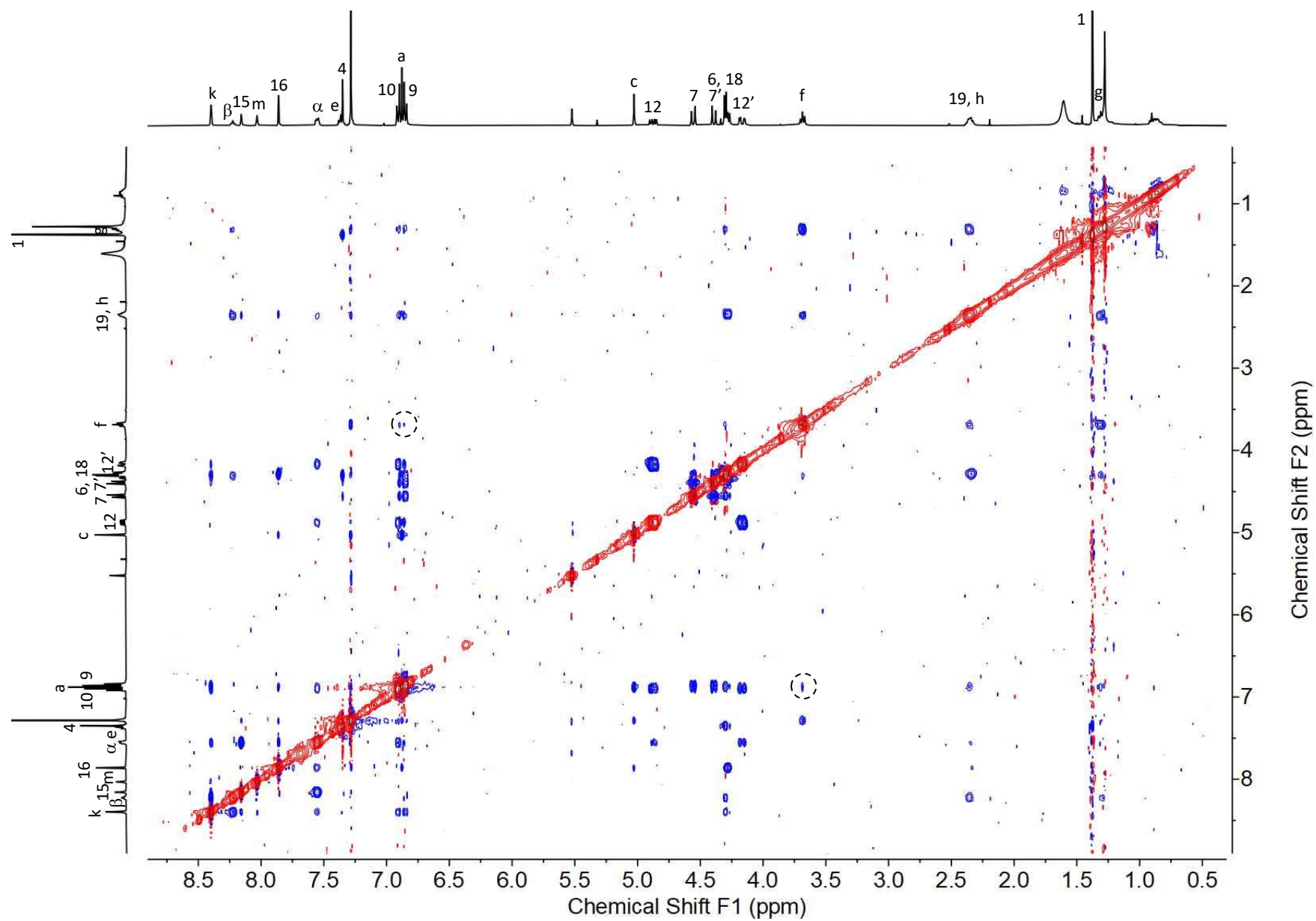


## HRMS (ES +ve)



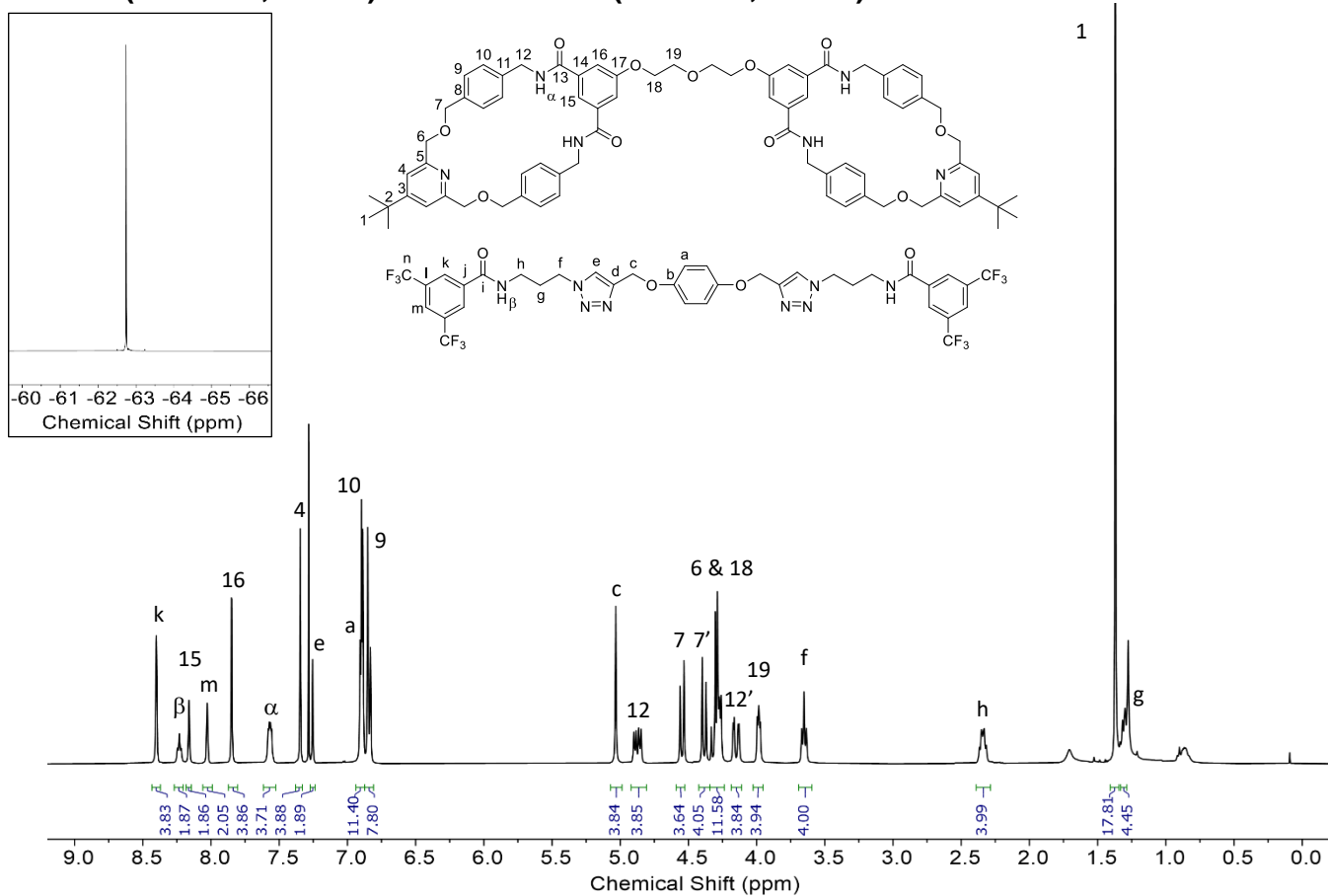


**$^1\text{H}$ - $^1\text{H}$  ROESY NMR (400 MHz,  $\text{CDCl}_3$ ) of HR1 (with intercomponent couplings highlighted)**

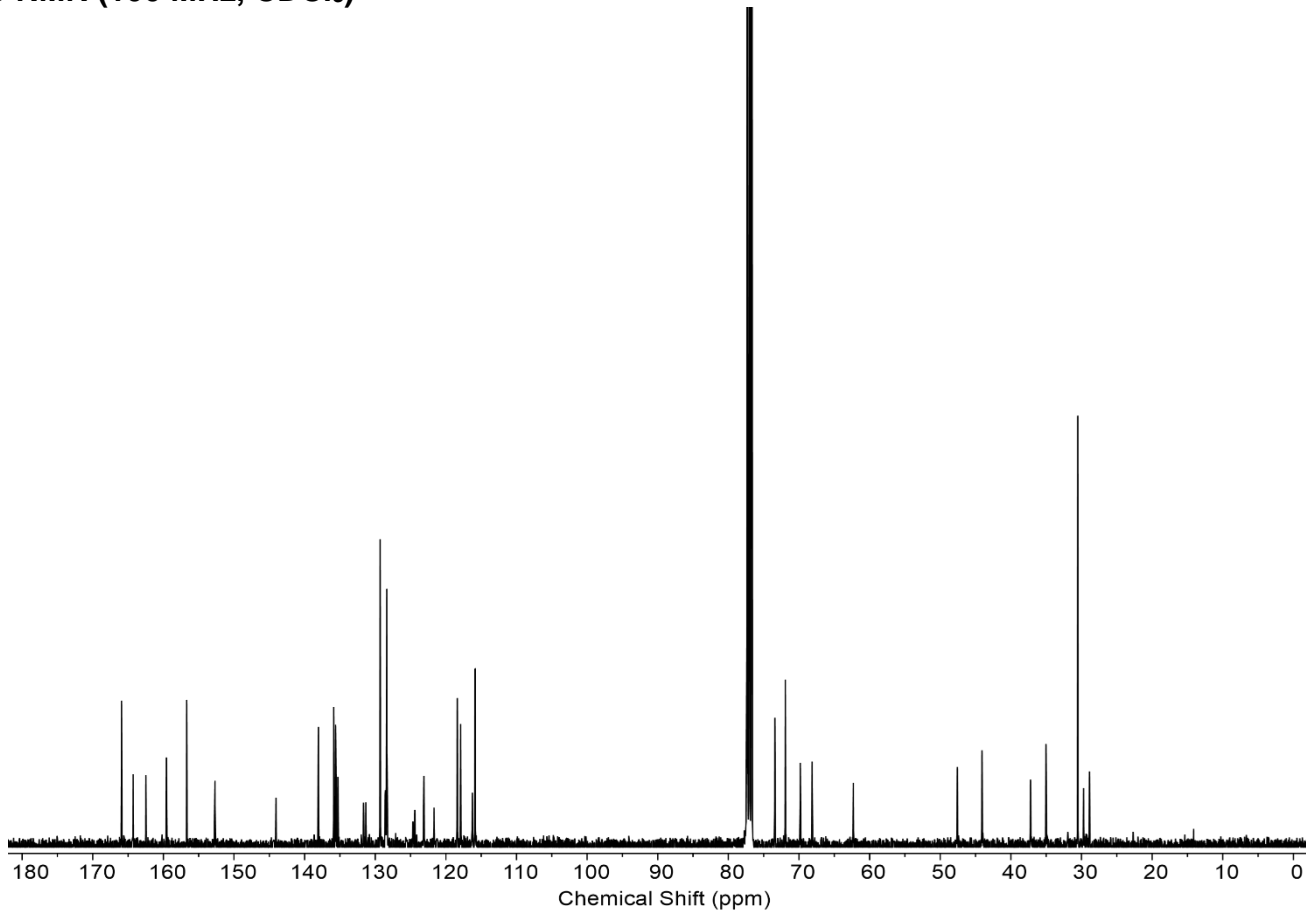


# Handcuff Rotaxane HR2

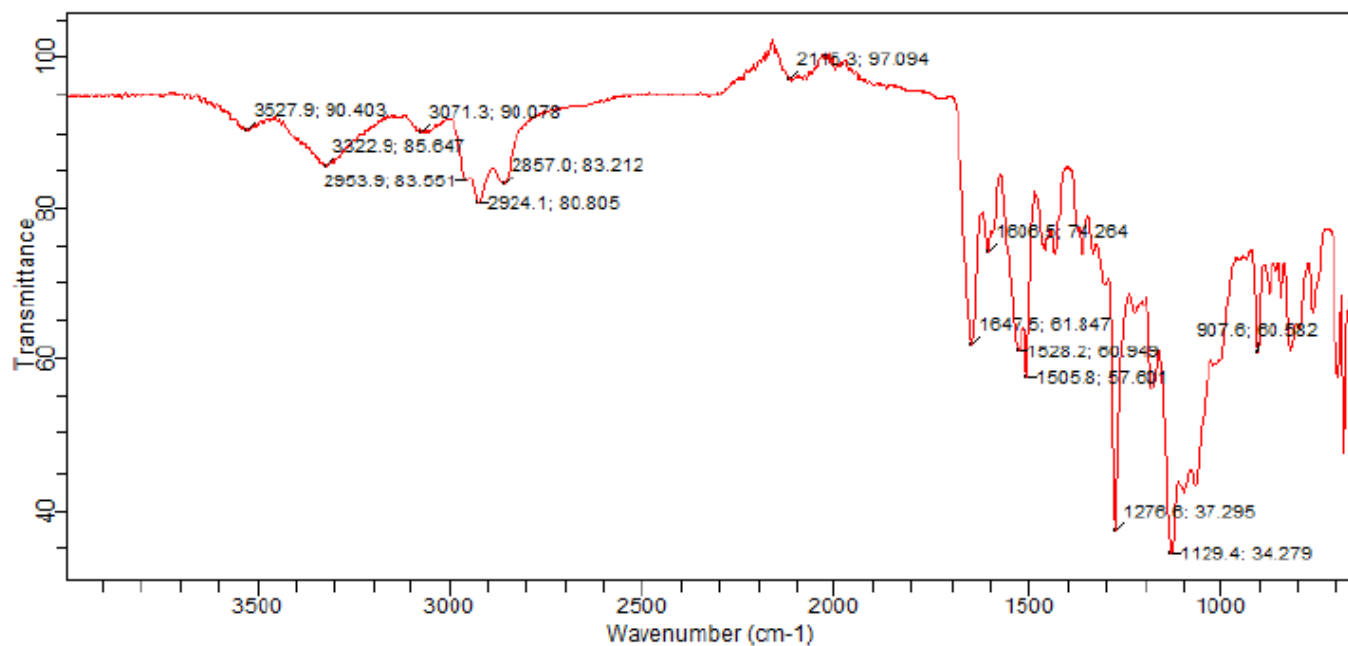
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ ) Inset:  $^{19}\text{F}$  NMR (377 MHz,  $\text{CDCl}_3$ )



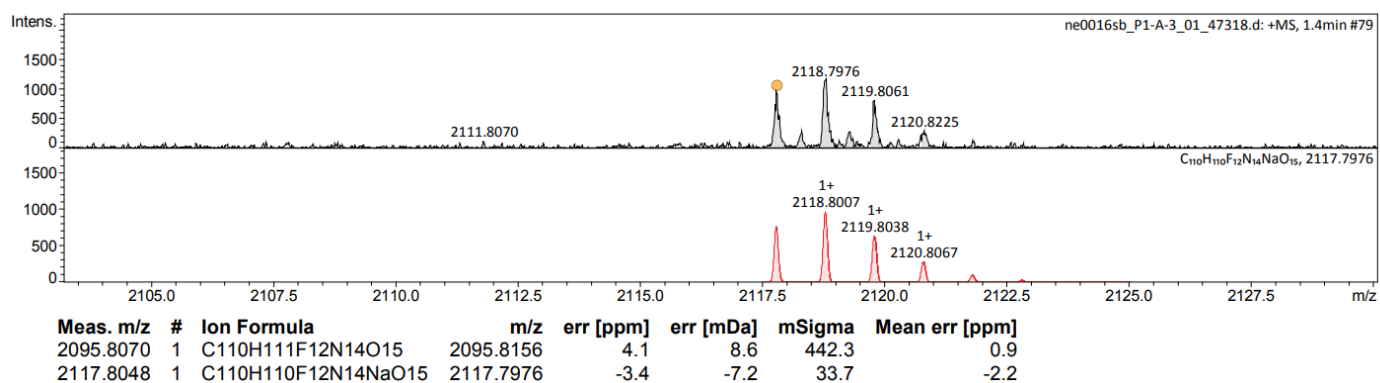
$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ )



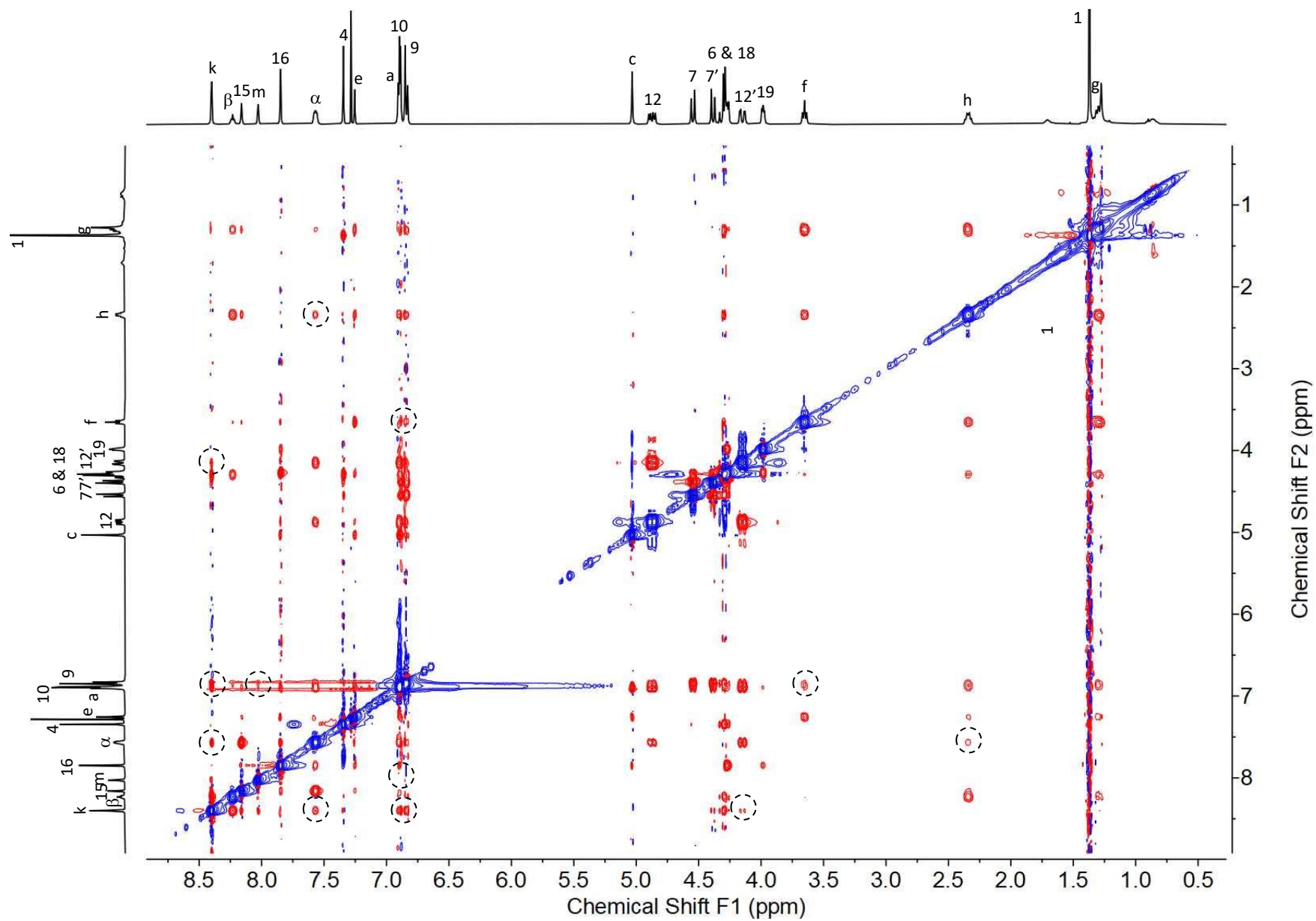
## IR Spectrum (neat)



## HRMS (ES +ve)

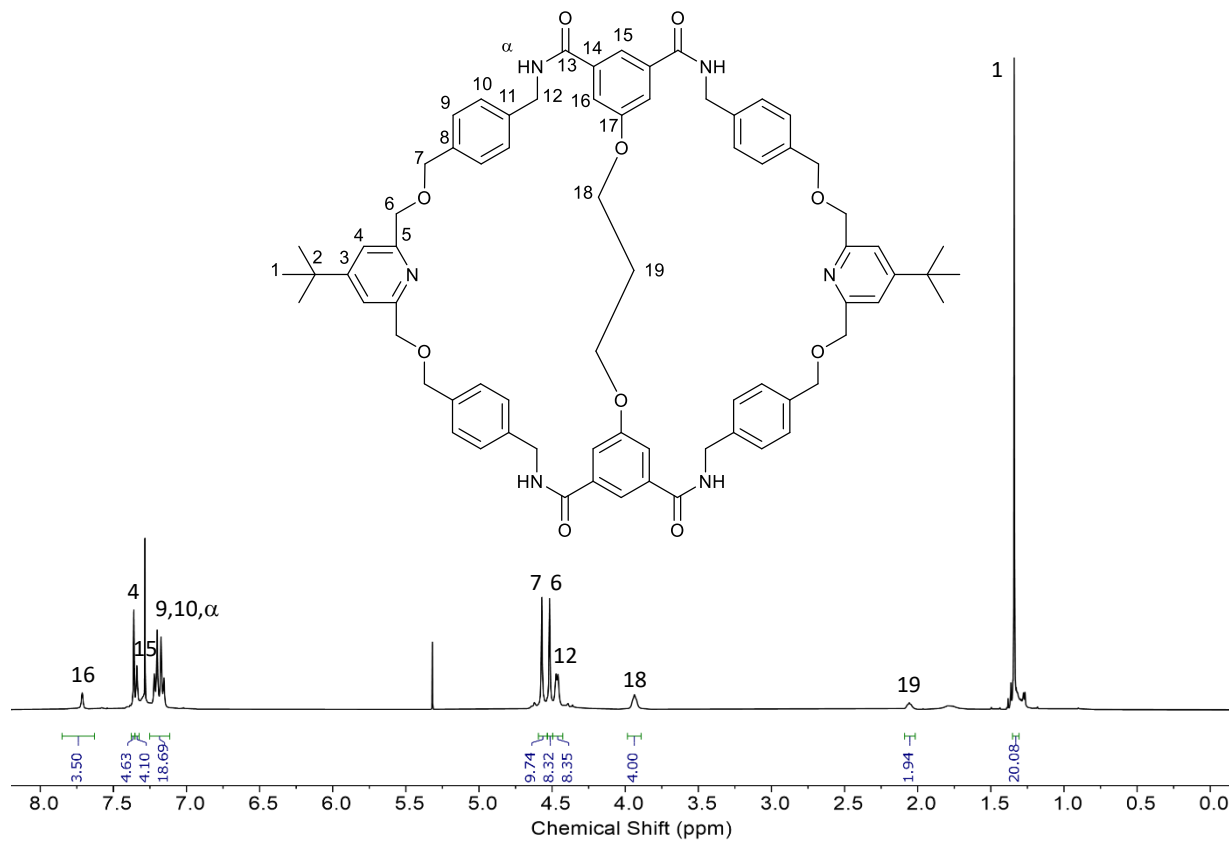


**$^1\text{H}$ - $^1\text{H}$  ROESY NMR (400 MHz,  $\text{CDCl}_3$ ) of HR2 (with intercomponent couplings highlighted)**

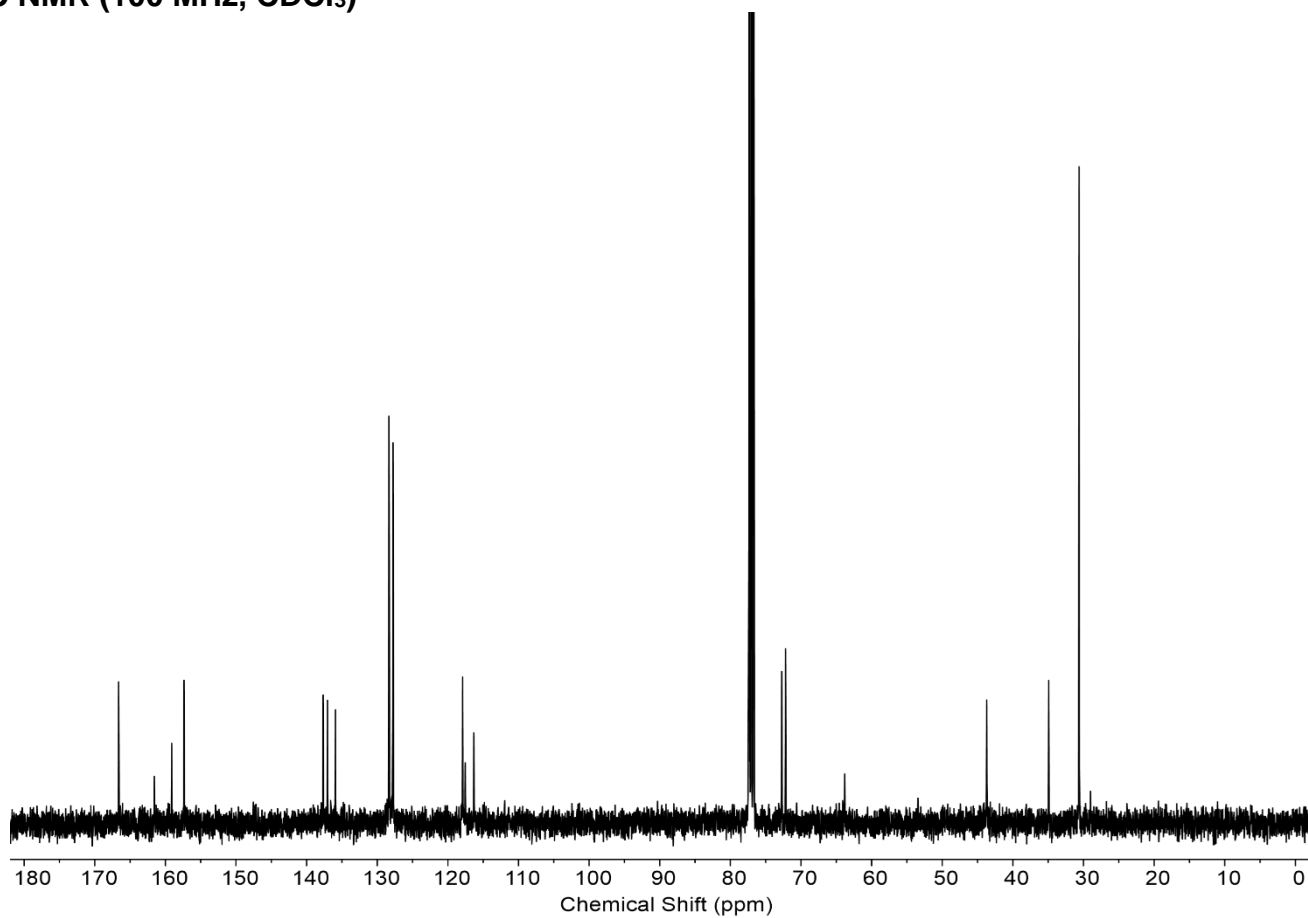


# Bis-Macrocycle BM1'

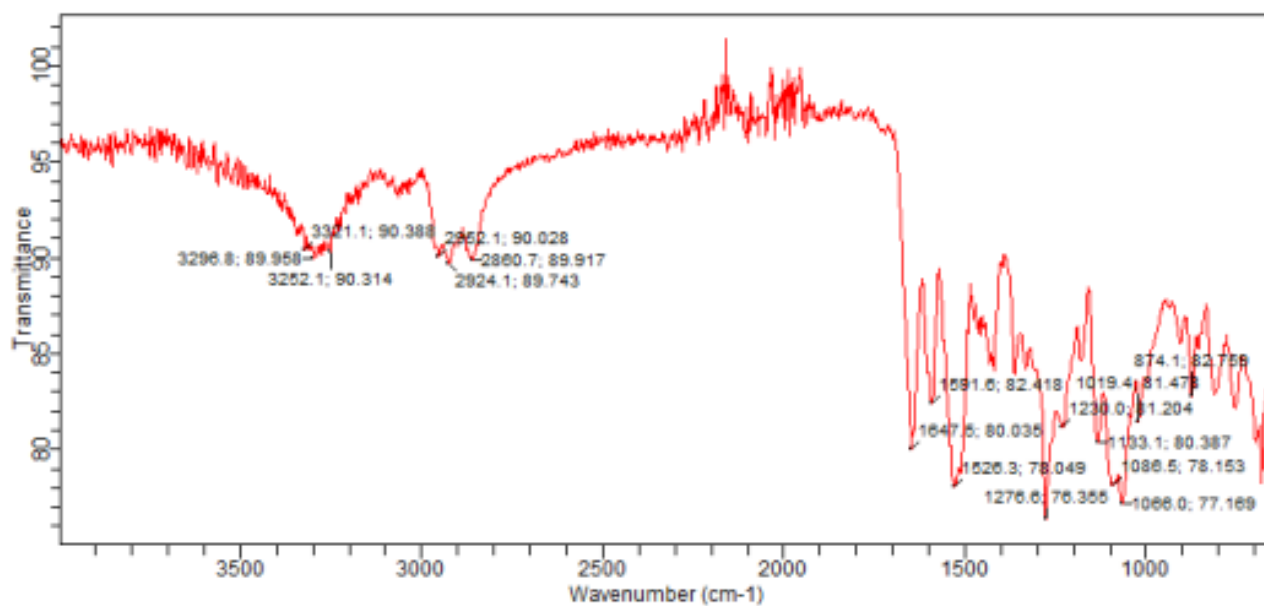
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>)



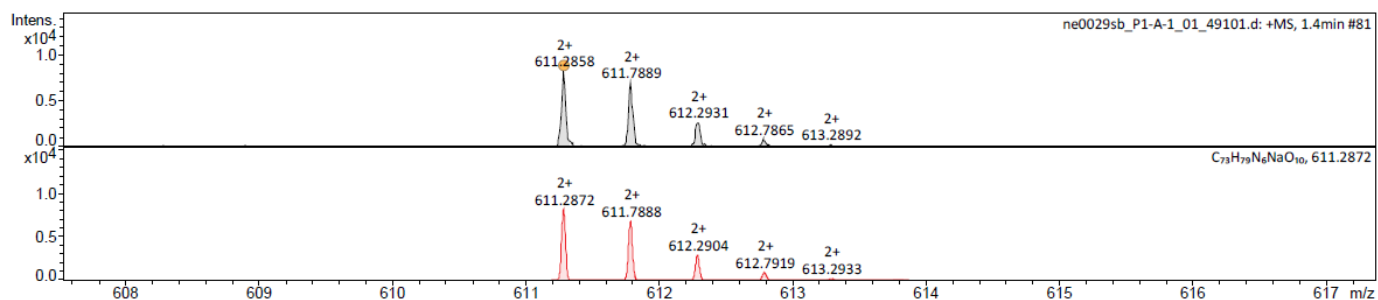
<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)



## IR Spectrum (neat)



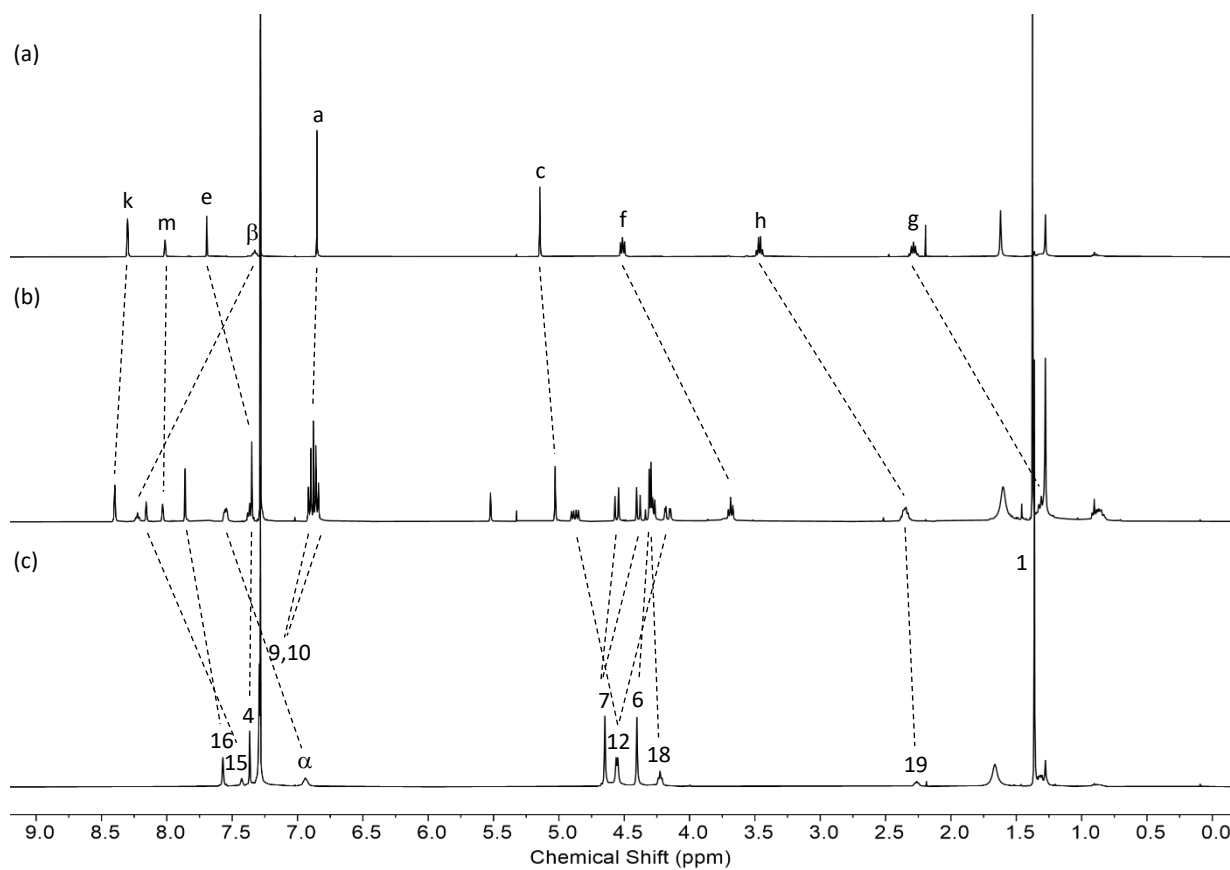
## HRMS (ES +ve)



Meas. m/z	#	Ion Formula	m/z	err [ppm]	err [mDa]	mSigma	Mean err [ppm]
611.2858	1	C <sub>73</sub> H <sub>79</sub> N <sub>6</sub> NaO <sub>10</sub>	611.2872	2.4	1.5	20.4	1.9
622.2761	1	C <sub>73</sub> H <sub>78</sub> N <sub>6</sub> Na <sub>2</sub> O <sub>10</sub>	622.2782	3.3	2.1	16.9	1.8
1199.5863	1	C <sub>73</sub> H <sub>79</sub> N <sub>6</sub> O <sub>10</sub>	1199.5852	-0.9	-1.1	51.9	-2.7
1221.5740	1	C <sub>73</sub> H <sub>78</sub> N <sub>6</sub> NaO <sub>10</sub>	1221.5672	-5.6	-6.8	181.0	-1.9

NB: Depicted molecular ion peak corresponds to  $[M+H+Na]^{2+}$

## Stacked $^1\text{H}$ NMR Spectra of Handcuff Rotaxane HR1

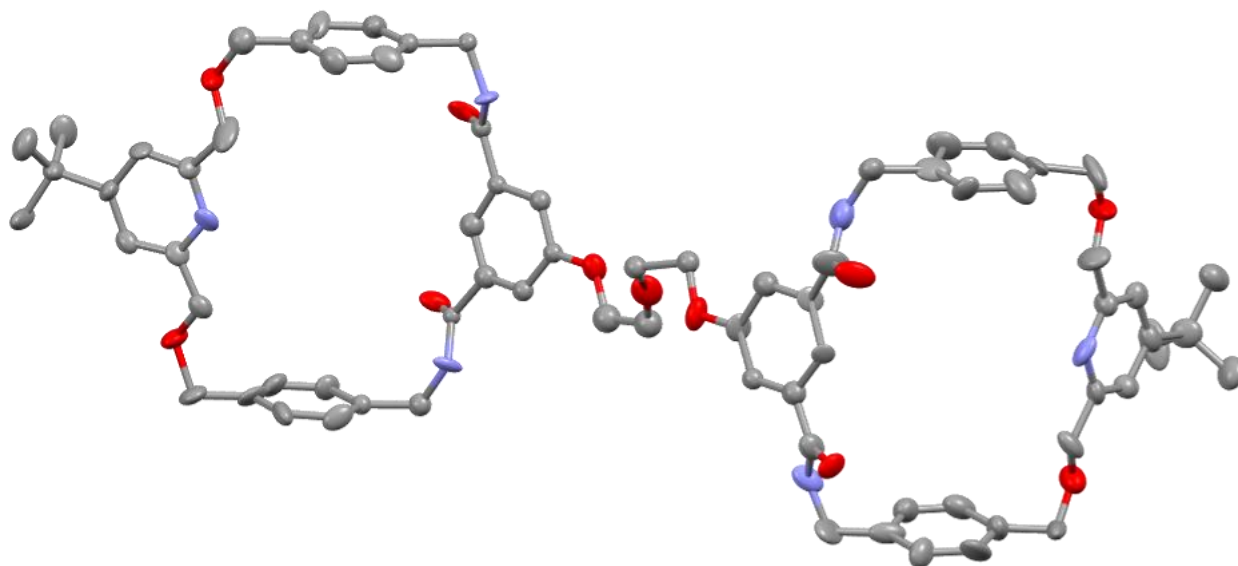


Stacked  $^1\text{H}$  NMR spectra of (a) axle **Ax**, (b) handcuff rotaxane **HR1** and (c) bis-macrocycle **BM1**.

### Part 3: Crystallographic Data

#### Bis-macrocycle **BM2**:

Single crystals of bis-macrocycle **BM2** were grown by slow evaporation of a chloroform solution. A suitable crystal was selected and studied using an Agilent SuperNova AtlasS2 diffractometer. Using Olex2<sup>1</sup> the structure was solved with the ShelXT<sup>2</sup> structure solution program using Intrinsic Phasing and refined with the ShelXL<sup>3</sup> refinement package using Least Squares minimisation.



X-ray crystal structure of bis-macrocycle **BM2**. Thermal ellipsoids are displayed at 50% probability.



Crystal data and structural refinement for bis-macrocycle **BM2**:

CCDC Number	2343052
Empirical formula	C <sub>74</sub> H <sub>80</sub> N <sub>6</sub> O <sub>11</sub>
Formula weight	1229.44
Temperature/K	100.01(10)
Crystal system	monoclinic
Space group	Pc
a/Å	27.7031(18)
b/Å	13.0253(10)
c/Å	9.5007(5)
α/°	90
β/°	93.854(6)
γ/°	90
Volume/Å <sup>3</sup>	3420.5(4)
Z	2
ρ <sub>calc</sub> /cm <sup>3</sup>	1.194
μ/mm <sup>-1</sup>	0.648
F(000)	1308.0
Crystal size/mm <sup>3</sup>	0.17 × 0.08 × 0.03
Radiation	Cu Kα (λ = 1.54184)
2θ range for data collection/°	7.502 to 145.604
Index ranges	-34 ≤ h ≤ 33, -16 ≤ k ≤ 16, -11 ≤ l ≤ 11
Reflections collected	8629
Independent reflections	8629 [R <sub>int</sub> = ?, R <sub>sigma</sub> = 0.0577]
Data/restraints/parameters	8629/187/828
Goodness-of-fit on F <sup>2</sup>	1.220
Final R indexes [ I  ≥ 2σ (I)]	R <sub>1</sub> = 0.1052, wR <sub>2</sub> = 0.3054
Final R indexes [all data]	R <sub>1</sub> = 0.1300, wR <sub>2</sub> = 0.3275
Largest diff. peak/hole / e Å <sup>-3</sup>	1.14/-0.44

#### Part 4: References

- 1 O. V. Dolomanov, L. J. Bourhis, R. J. Gildea, J. A. K. Howard and H. Puschmann, *J. Appl. Cryst.*, 2009, **42**, 339-341.
- 2 G. M. Sheldrick, *Acta Cryst. A*, 2015, **71**, 3-8.
- 3 G. M. Sheldrick, *Acta Cryst. C*, 2015, **71**, 3-8.