

Trapping an ester hydrate intermediate in a π -stacked macrocycle by multiple hydrogen bonds

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Table S1. Crystallographic data of compound 1.

Compounds	1
Formula	C ₁₅₉ H _{203.33} Cl ₁₀ Co ₆ N ₄₂ S ₂ O _{30.66}
Formula weight	3805.60
Temp. (K)	200
Crystal System	Trigonal
Space group	$R\bar{3}$
<i>a</i> (Å)	22.8534(6)
<i>b</i> (Å)	22.8534(6)
<i>c</i> (Å)	28.8133(11)
<i>a</i> (°)	90
<i>β</i> (°)	90
<i>γ</i> (°)	120
<i>V</i> (Å ³)	13032.4(8)
<i>Z</i>	6
$\rho_{cal.}$ (g cm ⁻³)	1.423
μ	0.813
<i>F</i> (000)	5668.0
θ range (°)	2.176–27.494
Reflections (<i>I</i> > 2σ)	4323
<i>R</i> ₁ (<i>I</i> > 2σ)	0.0708
<i>wR</i> ₂ (<i>all</i>)	0.2171
<i>GOF</i> on <i>F</i> ²	1.027
CCDC#	2271453

^a $R = \sum ||F_0| - |F_c|| / \sum |F_0|$

^b $wR = [\sum w(F_0 - F_c)^2 / \sum w(F_0^2)]^{1/2}$

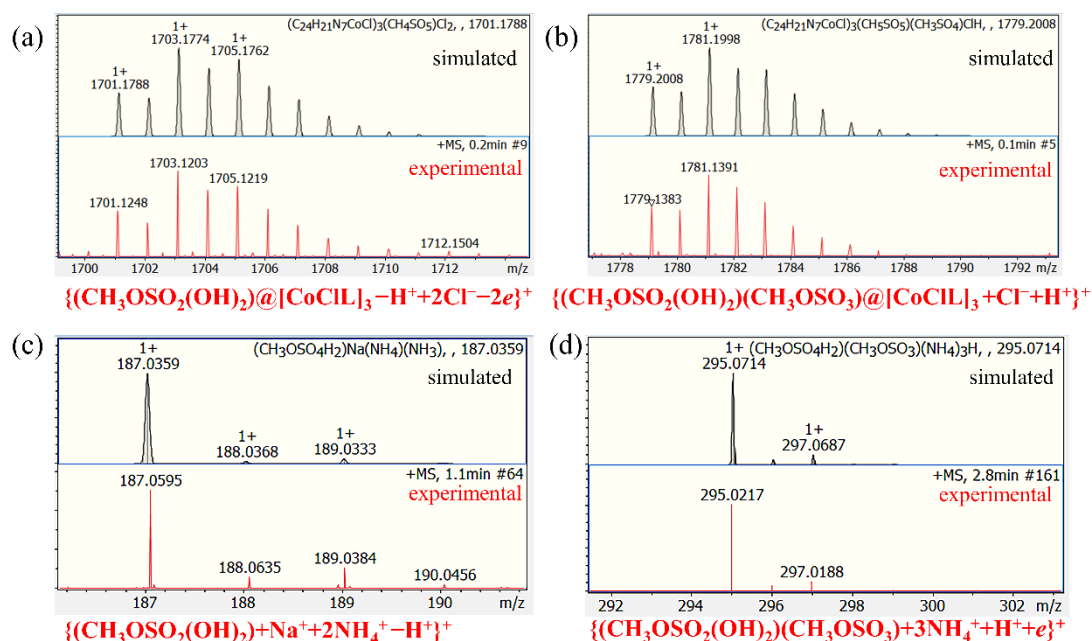


Figure S1. HR-MS of the reactant solution and the solution of compound **1** in methanol. (a) HR-MS of $\{(\text{CH}_3\text{OSO}_2(\text{OH})_2)@[\text{CILCo}^{\text{II}}]_3-\text{H}^++2\text{Cl}^--2e\}^+$ identified from the reactant solution. (b) HR-MS of $\{(\text{CH}_3\text{OSO}_2(\text{OH})_2)(\text{CH}_3\text{OSO}_3)@[\text{CILCo}^{\text{II}}]_3+\text{Cl}^--\text{H}^+\}^+$ identified from the reactant solution. (c) HR-MS of $\{(\text{CH}_3\text{OSO}_2(\text{OH})_2)+\text{Na}^++2\text{NH}_4^+-\text{H}^+\}^+$ identified from the solution of compound **1** in methanol. (d) HR-MS of $\{(\text{CH}_3\text{OSO}_2(\text{OH})_2)(\text{CH}_3\text{OSO}_3)+3\text{NH}_4^++\text{H}^++e\}^+$ identified from the solution of compound **1** in methanol.

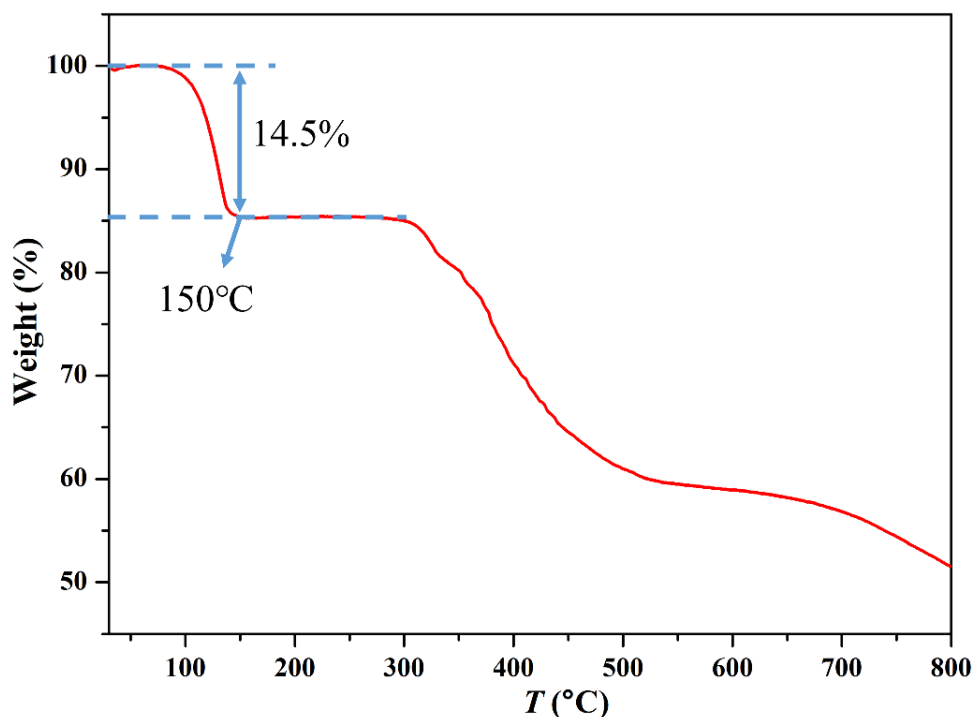


Figure S2. TGA for compound **1**.

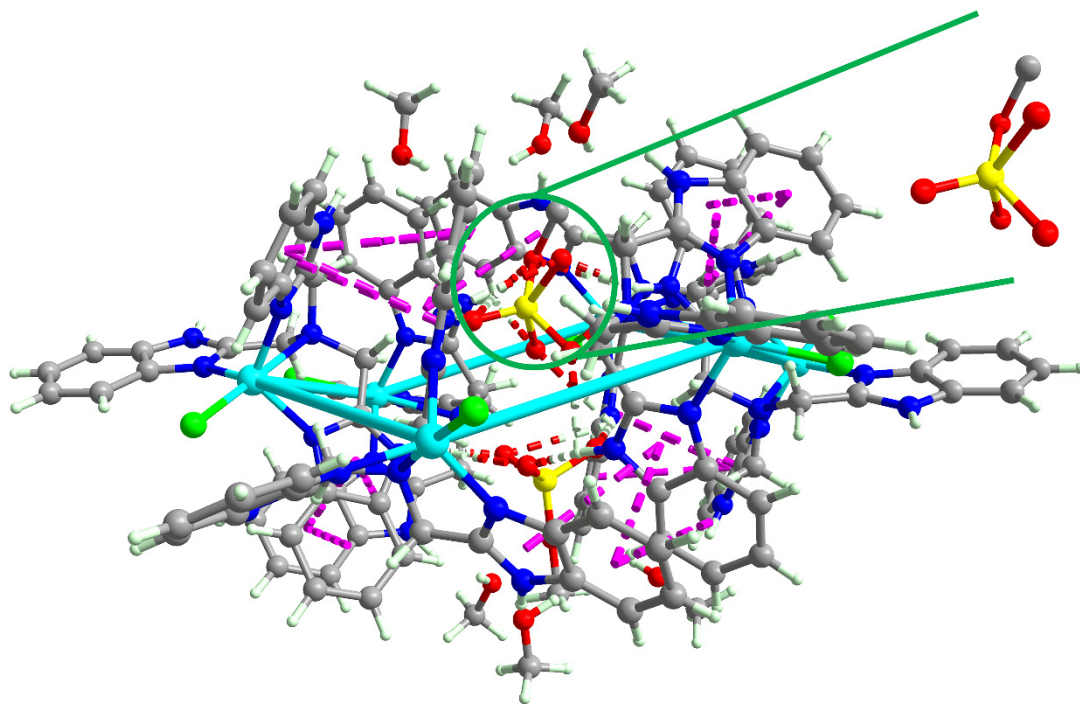


Figure S3. Side view of the structure of the boat-shaped macrocycle templating by $[\text{CH}_3\text{OSO}_2(\text{OH})_2]^-$ and $[\text{CH}_3\text{OSO}_3]^-$ in compound **1**. (inset: structure of the ester hydrate $[\text{CH}_3\text{OSO}_2(\text{OH})_2]^-$ in compound **1**.)

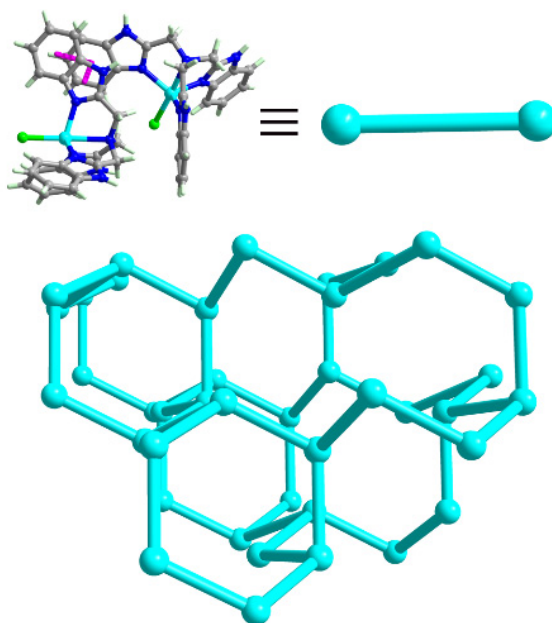


Figure S4. The pcu-h network of compound **1** when treating $[\text{CILCo}^{\text{II}}]^+$ as node and the π -stacked pair of benzimidazolymethyl arms as linker.

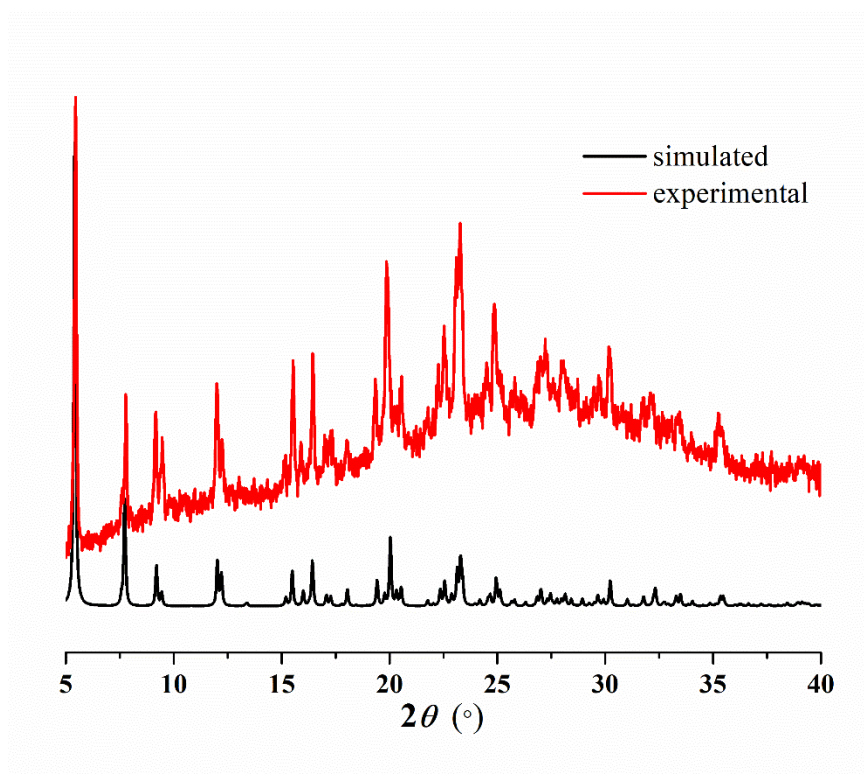


Figure S5. PXRD patterns for compound **1**.