# **Supplementary Information**

Co-crystals built from macrocycles with pyromellitic diimides and naphthalene derivatives: transformation of a binary co-crystal solvate to a solvate through solvent exchange

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# Single crystal X-ray diffraction experiment for crystal 1a

The yellowish-brown prismatic crystal ( $0.120 \times 0.050 \times 0.050 \text{ mm}^3$ ), obtained from slow evaporation of a mixture of chloroform and acetone containing **1** and **2**, was immersed in Paraton-N oil and placed in the N<sub>2</sub> cold stream at 100 K. Data were collected using diffractometer with a CMOS detector (Brukerd D8 VENTURE, CuKa:  $\lambda = 1.54178$  Å). Absorption correction was performed by an empirical method implemented in SADABS.<sup>1</sup> Structure solution and refinement were performed by using SHELXT-2018/2<sup>2</sup> and SHELXL-2019/2<sup>3</sup>.

 $C_{57.8}H_{52.8}Br_2Cl_{2.39}N_4O_{11}, Mr = 1223.90$ ; triclinic, space group *P*-1, *Z* = 2,  $D_{calc} = 1.555$  g·cm<sup>-3</sup>, a = 9.7752(10), b = 11.4458(12), c = 24.731(3) Å,  $a = 94.352(3), \beta = 93.154(3), \gamma = 108.044(3)^\circ$ , *V* = 2614.4(5) Å<sup>3</sup>, 36841 observed and 10049 [*I* > 2 $\sigma$ (*I*)], 10481 [all] independent reflections, 782 parameters, 48 restraints, final  $R_1 = 0.0343, wR_2 = 0.0885, S = 1.025$  [*I* > 2 $\sigma$ (*I*)] and  $R_1 = 0.0357, wR_2 = 0.0907, S = 1.052$  [all data]. CCDC 2389266.

All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were refined isotropically on the calculated positions using a riding model (AFIX 13, 137, 147, 23, and 43) with  $U_{iso}$  values constrained to 1.2/1.5  $U_{eq}$  of their parent atoms. The two disordered chloroform molecules (C58, C11-Cl3 and C59, Cl4-Cl6) overlapped in almost identical positions. In addition, the disordered another chloroform molecule (C60, Cl7-Cl9) was also overlapped in the same position. They were refined with PART –n. Each occupancy was as follows. C58, Cl1-Cl3: 50%, C59, Cl4-Cl6: 15%, C60, Cl7-Cl9: 15%.

The ratio of **1**:**2**:CHCl<sub>3</sub>:acetone included in the crystal was 1:1:0.8:1.



Fig. S1 ORTEP drawing of 1, 2, chloroform, and acetone in an asymmetric unit in crystal 1a (50% probability).



Fig. S2 ORTEP drawing of disordered chloroform molecules in crystal 1a (50% probability).

# Single crystal X-ray diffraction experiment for crystal 1b

The yellowish-brown prismatic crystal ( $0.120 \times 0.120 \times 0.080 \text{ mm}^3$ ), obtained from slow evaporation of an acetone solution of **1** and **2**, was immersed in Paraton-N oil and placed in the N<sub>2</sub> cold stream at 100 K. Data were collected using diffractometer with CMOS detector (Bruker D8 VENTURE, CuKa:  $\lambda = 1.54178$  Å). Absorption correction was performed by an empirical method implemented in SADABS.<sup>1</sup> Structure solution and refinement were performed by using SHELXT-2018/2<sup>2</sup> and SHELXL-2019/2<sup>3</sup>.

 $C_{60}H_{58}Br_2N_4O_{12}$ , Mr = 1186.92; triclinic, space group *P*-1, Z = 2,  $D_{calc} = 1.522$  g·cm<sup>-3</sup>, a = 9.7166(6), b = 11.3692(7), c = 24.7966(16) Å, a = 94.639(2),  $\beta = 92.734(2)$ ,  $\gamma = 107.934(2)^\circ$ , V = 2590.0(3) Å<sup>3</sup>, 38075 observed and 10351 [ $I > 2\sigma(I)$ ], 10549 [all] independent reflections, 709 parameters, final  $R_1 = 0.0256$ ,  $wR_2 = 0.0680$ , S = 1.065 [ $I > 2\sigma(I)$ ] and  $R_1 = 0.0261$ ,  $wR_2 = 0.0683$ , S = 1.065 [all data]. CCDC 2389267.

All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were refined isotropically on the calculated positions using a riding model (AFIX 13, 137, 147, 23, and 43) with  $U_{iso}$  values constrained to 1.2  $U_{eq}$  of their parent atoms.

The ratio of 1:2: acetone included in the crystal was 1:1:2.



Fig. S3 ORTEP drawing of 1, 2, and acetone in an asymmetric unit in crystal 1b (50% probability).

# Single crystal X-ray diffraction experiment for crystal 1c

The yellowish-brown plate crystal ( $0.120 \times 0.120 \times 0.080 \text{ mm}^3$ ), obtained from slow evaporation of a chloroform solution of **1** and **2**, was immersed in Paraton-N oil and placed in the N<sub>2</sub> cold stream at 100 K. Data were collected using diffractometer with CMOS detector (Bruker D8 VENTURE, CuKa:  $\lambda = 1.54178$  Å). Absorption correction was performed by an empirical method implemented in SCALE3 ABSPACK. Structure solution and refinement were performed by using SHELXT-2018/2<sup>2</sup> and SHELXL-2019/2<sup>3</sup>.

 $C_{59}H_{51}Br_2Cl_5N_4O_{10}, Mr = 1667.60$ ; monoclinic, space group  $P2_1/m, Z = 2, D_{calc} = 1.635 \text{ g}\cdot\text{cm}^{-3}$ , a = 9.6670(8), b = 22.9197(19), c = 15.4712(13) Å,  $\beta = 98.877(3)^\circ$ , V = 3386.8(5) Å<sup>3</sup>, 44054 observed and 6548 [ $I > 2\sigma(I)$ ], 7202 [all] independent reflections, 498 parameters, 7 restraints, final  $R_1 = 0.0573, wR_2 = 0.1451, S = 1.056$  [ $I > 2\sigma(I)$ ] and  $R_1 = 0.0622, wR_2 = 0.1494, S = 1.056$  [all data]. CCDC 2389268.

All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were refined isotropically on the calculated positions using a riding model (AFIX 13, 147, 23, and 43) with  $U_{iso}$  values constrained to 1.2  $U_{eq}$  of their parent atoms. The crystals contained 2.5 chloroform molecules in five positions in an asymmetric unit. One-half chloroform molecule (C32, C11, C12, C11b) encapsulated in the host molecule located on the symmetry axis was refined to 50% occupancy. Two pairs of disordered chloroform molecules (C33, C13, C14, C15; C34, C16, C17, C18 and C35, C19, C110, C111; C36, C112, C113, C114) filled in the void are refined with 61.9; 38.1% and 51.7; 48.3% occupancy (with PART -n), respectively.

The ratio of **1**:**2**:CHCl<sub>3</sub> included in the crystal was 1:1:5.



Fig. S4 ORTEP drawing of complete structure 1, 2, and chloroform in crystal 1c (50% probability).



Fig. S5 ORTEP drawing of 1, 2, and chloroform in an asymmetric unit in crystal 1c (50% probability).

#### Single crystal X-ray diffraction experiment for crystal 1d

The reddish-brown plate crystal ( $0.200 \times 0.200 \times 0.020 \text{ mm}^3$ ), obtained from slow evaporation of a mixture of chloroform and acetone containing **1** and **3**, was immersed in Paraton-N oil and placed in the N<sub>2</sub> cold stream at 100 K. Data were collected using diffractometer with CMOS detector (Bruker D8 VENTURE, CuK $\alpha$ :  $\lambda = 1.54178$  Å). Absorption correction was performed by an empirical method implemented in SADABS.<sup>1</sup> Structure solution and refinement were performed by using SHELXT-2018/2<sup>2</sup> and SHELXL-2019/2<sup>3</sup>.

 $C_{58}H_{52}Cl_{12}N_4O_{10}$ , Mr = 1390.43; monoclinic, space group  $P2_1/c$ , Z = 4,  $D_{calc} = 1.548$  g·cm<sup>-3</sup>, a = 9.3975(6), b = 23.0975(14), c = 27.7307(17) Å,  $\beta = 97.683(3)^\circ$ , V = 5965.2(6) Å<sup>3</sup>, 76857 observed and 9908 [ $I > 2\sigma(I)$ ], 12549 [all] independent reflections, 796 parameters, 6 restraints, final  $R_1 = 0.0909$ ,  $wR_2 = 0.2720$ , S = 1.037 [ $I > 2\sigma(I)$ ] and  $R_1 = 0.1061$ ,  $wR_2 = 0.2905$ , S = 1.037 [all data]. CCDC 2389269.

All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were refined isotropically on the calculated positions using a riding model (AFIX 13, 147, 23, and 43) with  $U_{iso}$  values constrained to 1.2/1.5  $U_{eq}$  of their parent atoms. The crystals contained four chloroform molecules in five positions in an asymmetric unit. Three chloroform molecules (C55, C11-C13 and C56, C14-C16 and C57, C17-C19) in the void were refined normally. Two disordered chloroform molecules (C60, C110-C112 and C61, C113-C115) in the void were refined with 58.1 and 41.9% occupancy (with PART -n), respectively.

The ratio of **1**:**3**:CHCl<sub>3</sub> included in the crystal was 1:1:4.



Fig. S6 ORTEP drawing of 1, 3, and chloroform in an asymmetric unit in crystal 1d (50% probability).

# Single crystal X-ray diffraction experiment for crystal 1e

The colorless prismatic crystal ( $0.200 \times 0.150 \times 0.080 \text{ mm}^3$ ), obtained from slow evaporation of a mixture of chloroform and acetone containing **1** and **4**, was immersed in Paraton-N oil and placed in the N<sub>2</sub> cold stream at 100 K. Data were collected using diffractometer with CMOS detector (Bruker D8 VENTURE, CuK $\alpha$ :  $\lambda = 1.54178$  Å). Absorption correction was performed by an empirical method implemented in SADABS.<sup>1</sup> Structure solution and refinement were performed by using SHELXT-2018/2<sup>2</sup> and SHELXL-2019/2<sup>3</sup>.

 $C_{103.50}H_{93.75}Br_2Cl_{9.75}N_8O_{16.75}$ , Mr = 2223.03; triclinic, space group *P*-1, Z = 1,  $D_{calc} = 1.485$ g·cm<sup>-3</sup>, a = 9.590(2), b = 15.202(4), c = 18.069(3) Å, a = 107.935(14),  $\beta = 95.420(14)$ ,  $\gamma = 92.865(17)^\circ$ , V = 2486.5(10) Å<sup>3</sup>, 35907 observed and 9339 [ $I > 2\sigma(I)$ ], 10087 [all] independent reflections, 708 parameters, 19 restraints, final  $R_1 = 0.0529$ ,  $wR_2 = 0.1358$ , S = 1.117 [ $I > 2\sigma(I)$ ] and  $R_1 = 0.0563$ ,  $wR_2 = 0.1394$ , S = 1.130 [all data]. CCDC 2389270.

All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were refined isotropically on the calculated positions using a riding model (AFIX 13, 137, 23, and 43) with  $U_{iso}$  values constrained to 1.2/1.5  $U_{eq}$  of their parent atoms. The disordered and overlapped chloroform (C50, C11-Cl3) and acetone (C54-C56, O9) molecules were located in almost identical positions. They were refined with PART –n. The occupancies of C50, C11-Cl3: 62.5% and C54-C56, O9: 37.5% have been refined. In addition, other disordered chloroform molecules C51, Cl4-Cl6, and C52, Cl7-Cl9 in the void were normally refined with 51.8% and 48.2% occupancy (with PART –n), respectively.

The ratio of 1:4:CHCl<sub>3</sub>:acetone included in the crystal was 1:1:3:1.



Fig. S7 ORTEP drawing of 1, 4, chloroform, and acetone in crystal 1e (50% probability).

# Single crystal X-ray diffraction experiment for crystal 1f

The colorless prismatic crystal ( $0.080 \times 0.080 \times 0.060 \text{ mm}^3$ ), obtained from crystal of **1d** soaked with diethyl ether at room temperature for 3 d, was immersed in Paraton-N oil and placed in the N<sub>2</sub> cold stream at 100 K. Data were collected using diffractometer with CMOS detector (Bruker D8 VENTURE, CuK $\alpha$ :  $\lambda = 1.54178$  Å). Absorption correction was performed by an empirical method implemented in SADABS.<sup>1</sup> Structure solution and refinement were performed by using SHELXT-2018/2<sup>2</sup> and SHELXL-2019/2<sup>3</sup>.

 $C_{48}H_{50}N_4O_9$ , Mr = 826.92; triclinic, space group *P*-1, Z = 4,  $D_{calc} = 1.319$  g·cm<sup>-3</sup>, a = 16.4443(5), b = 16.6197(5), c = 17.6078(6) Å,  $\alpha = 113.668(2)$ ,  $\beta = 105.994(2)$ ,  $\gamma = 92.943(2)^\circ$ , V = 4164.5(2) Å<sup>3</sup>, 45481 observed and 9797 [ $I > 2\sigma(I)$ ], 15905 [all] independent reflections, 1104 parameters, final  $R_1 = 0.0918$ ,  $wR_2 = 0.2143$ , S = 1.053 [ $I > 2\sigma(I)$ ] and  $R_1 = 0.1505$ ,  $wR_2 = 0.2668$ , S = 1.053 [all data]. CCDC 2444562.

All non-hydrogen atoms were refined anisotropically. Hydrogen atoms were refined isotropically on the calculated positions using a riding model (AFIX 13, 137, 23, and 43) with  $U_{iso}$  values constrained to 1.2/1.5  $U_{eq}$  of their parent atoms. The ratio of 1:Et<sub>2</sub>O included in the crystal was 1:1.



Fig. S8 ORTEP drawing of 1 and diethyl ether in crystal 1f (50% probability).



Fig. S9 Packing diagrams of 1 and naphthalene derivatives in crystals (a) 1a, (b) 1b, (c) 1c, (d) 1d, and (e) 1e.

# The stoichiometric ratio of 1 and diethyl ether in crystal 1f by <sup>1</sup>H NMR experiments

Crystal **1d** was immersed into diethyl ether at room temperature for 3 d. The resulting crystals **1f** were collected, and dried under air for 0.5 h on a filter paper. The crystals were dissolved in CDCl<sub>3</sub> followed by <sup>1</sup>H NMR analysis (Fig. S10). The composition of crystals was evaluated from the integral ratios of the signal of the aromatic proton at  $\delta$  8.22 ppm of **1** and the corresponding protons of diethyl ether. The molar ratio of **1** and diethyl ether was approximately 1:1.



**Fig. S10** <sup>1</sup>H NMR spectra (400 MHz, CDCl<sub>3</sub>); (a) diethyl ether, (b) crystal **1f**, and (c) macrocycle **1**.

# References

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