

Supporting Information

A 3D Chiral Nanoporous Coordination Framework Consisting of Homochiral Nanotubes Assembled from Octuple Helices

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X-ray structure analysis: $C_{31}H_{38}N_3O_8Cd$: $M_r = 693.04$, hexagonal, space group $P6_122$, $a = b = 26.712(4)$ Å, $c = 15.919(3)$ Å, $\alpha = \beta = 90^\circ$, $\gamma = 120^\circ$, $V = 9837(3)$ Å³, $Z = 12$, $\rho_{\text{calcd}} = 1.404$ mg m⁻³, Flack parameter = $-0.01(3)$, final $R1 = 0.0384$ for 9677 independent reflections [$I > 2\sigma(I)$]. The data were collected on a Bruker Apex CCD diffractometer at 133(2) K with graphite-monochromated $Mo_{K\alpha}$ radiation ($\lambda = 0.71073$ Å).

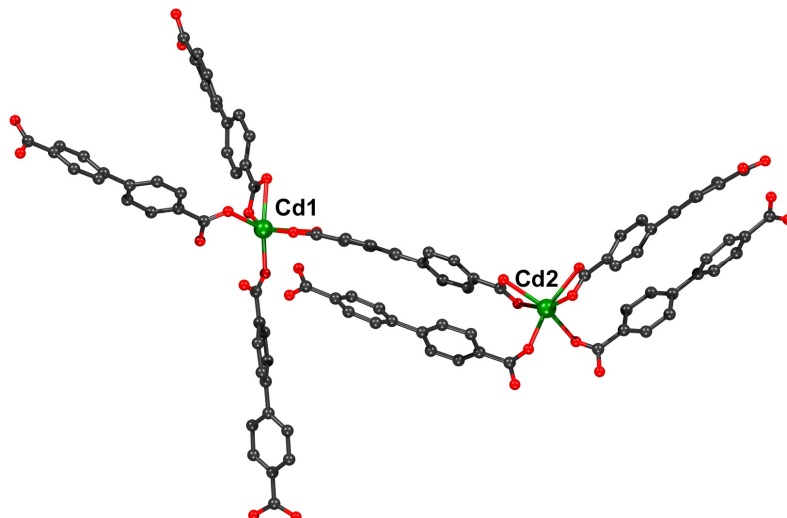
The data were collected on a Bruker Apex CCD diffractometer at 133(2) K with graphite-monochromated $Mo_{K\alpha}$ radiation ($\lambda = 0.71073$ Å). The structures were solved by direct method and refined by full-matrix least-squares methods with SHELXL.^[1] The anion and DMA molecules were highly disordered and could not be modeled properly, thus the SQUEEZE routine, a part of the PLATON package of crystallographic software, was applied to calculate the solvent disorder area and remove its contribution to the overall intensity data. The refinements of the guest-free structure on the squeezed data gave the final $R1 = 0.0384$ and $wR2 = 0.0798$ ($R1 = 0.0463$ and $wR2 = 0.1322$ without employing PLATON/SQUEEZE). The final formula was calculated from the SQUEEZE results combined with the following characterization techniques. Despite the disorder, identification of the guest molecules is readily accomplished by ¹H NMR spectroscopy. The ¹H NMR spectrum of **1** recorded in D₂O exhibits a set of well-resolved proton signals. The signals at 2.88, 2.73 and 1.91 ppm attributed to three CH₃ groups of DMA,^[2] while the signal at 2.54 ppm ascribes to the CH₃ group of dimethylamine. Furthermore, NMR analysis reveals that the relative molar ratio of dimethylamine and DMA is approximately 1:2. The IR

spectrum of **1**, as expected, exhibits a sharp band at 1619 cm^{-1} corresponding to $\tilde{\nu}_{\text{C=O}}$ stretching frequency, which is indicative of DMA molecule. According to the previous literature, the peaks of 1540 and 1398 cm^{-1} are attributed to the asymmetric and symmetric stretching vibrations of the carboxylate group of bpdc ligand.^[3] The peak at ca. 3419 cm^{-1} is attributed to the N-H absorption vibration of dimethylamine. Further information supporting the formula of **1** is obtained by thermogravimetric analysis, elemental analysis and the consideration of charge balance.

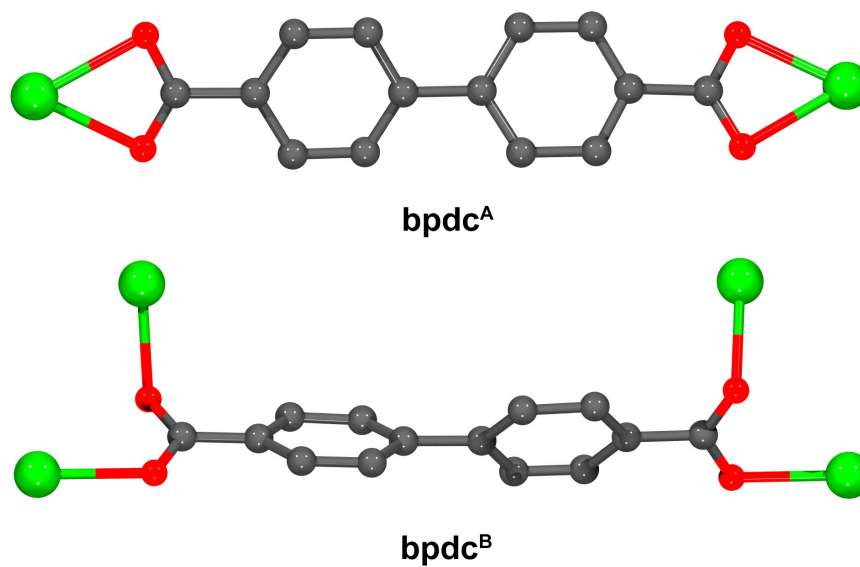
[1] G. M. Sheldrick, SHELXL, Version 5.1. Siemens Industrial Automation, Inc., Madison, WI, 1997.

[2] L. Carlucci, G. Ciani, D. M. Proserpio, F. Porta, *Angew. Chem. Int. Ed.* 2003, **42**, 317.

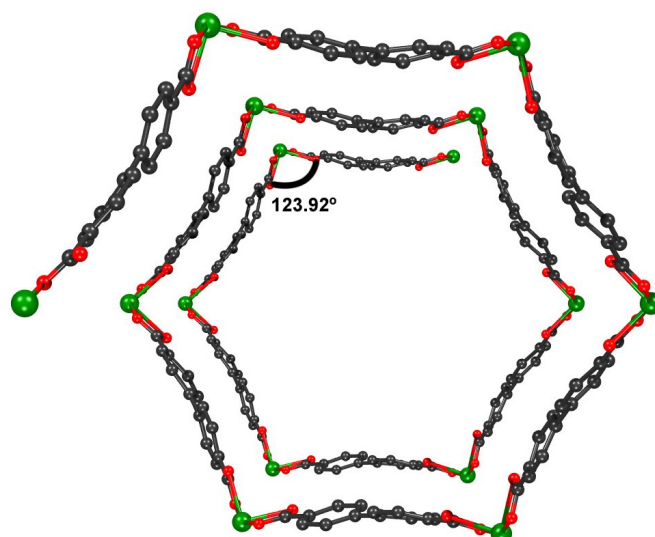
[3] a) F. A. Almeida Paz, Y. Z. Khimyak, A. D. Bond, J. Rocha, J. Klinowsk, *Eur. J. Inorg. Chem.* 2002, 2823; b) Q. R Fang, X. Shi, G. Wu, G. Tian, G. S. Zhu, R. W. Wang, S. L. Qiu, *J. Solid. State.* 2003, **176**, 1; c) Y. H. Wang, B. Breidenkötter, B. Riegera, D. Volkmer, *Dalton Trans.* 2007, 689.



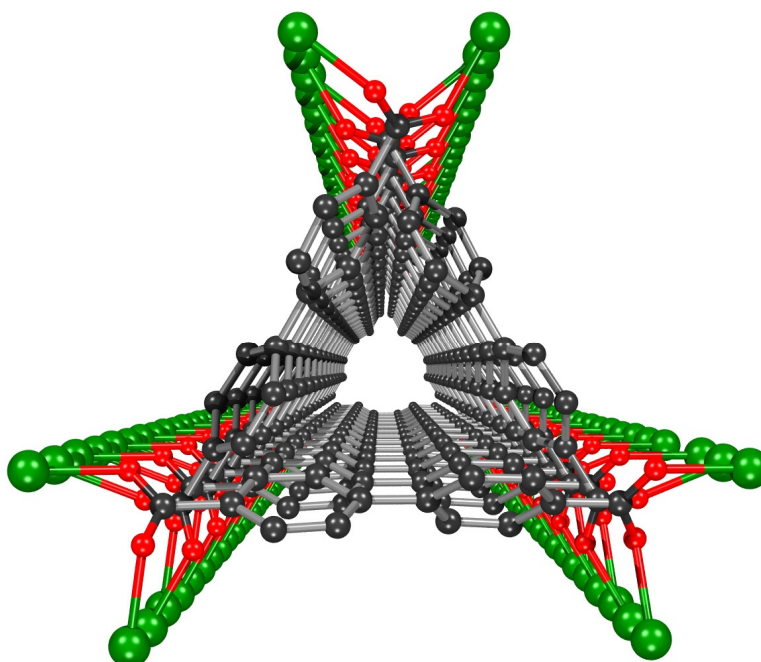
SI-1. Section of the X-ray crystal structure of **1**, showing the coordination environments of Cd^{2+} ions and their connectivity with bpdc anions. Color code: C black; O red; Cd green.



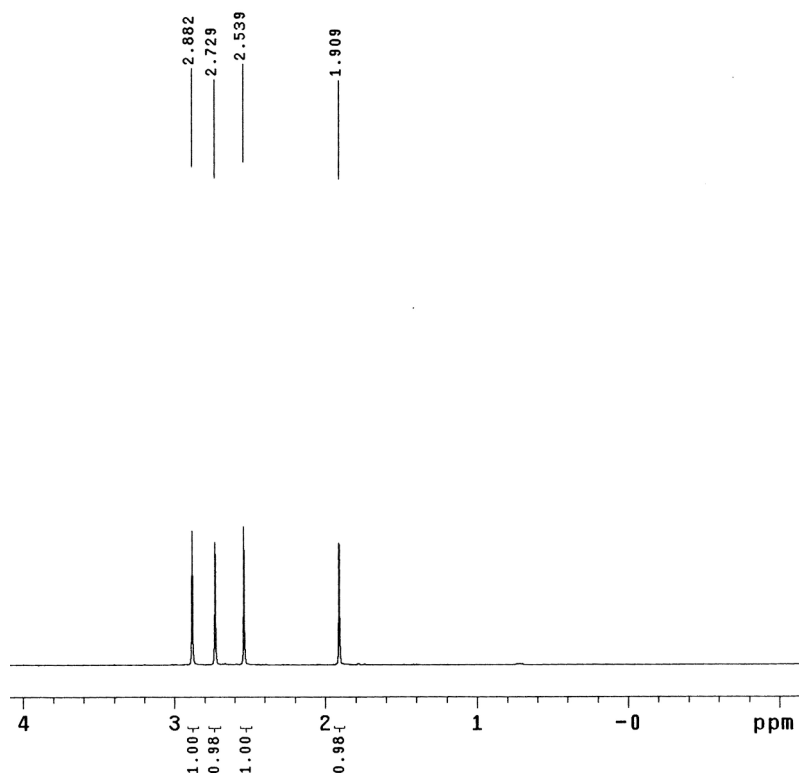
SI-2. Coordination modes of the bpdc ligands in the structure of **1**. bpdc^A: chelating bis(bidentate), bpdc^B: bridging bis(bidentate). Color code: C black; O red; Cd green.



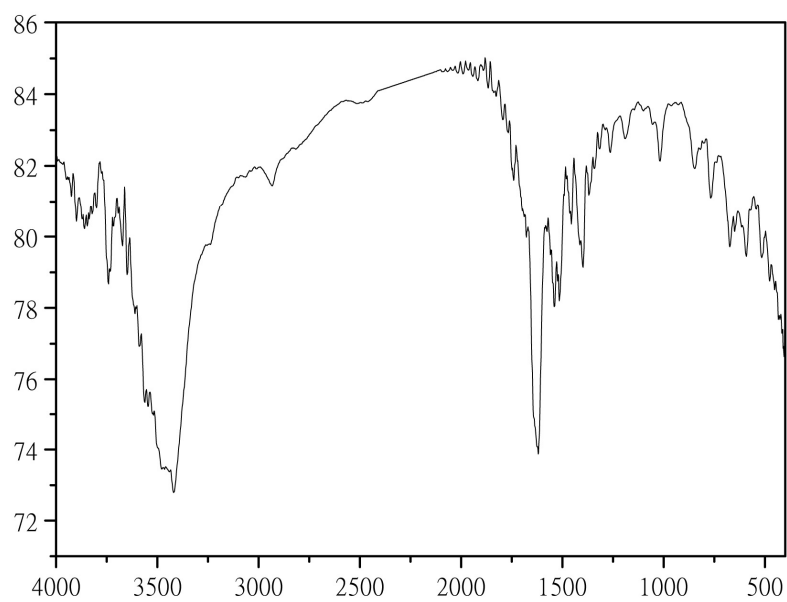
SI-3. An angle of 123.92° formed between the planes defined by the two C-O-Cd-O-C chelate rings at one metal center. Color code: C black; O red; Cd green.



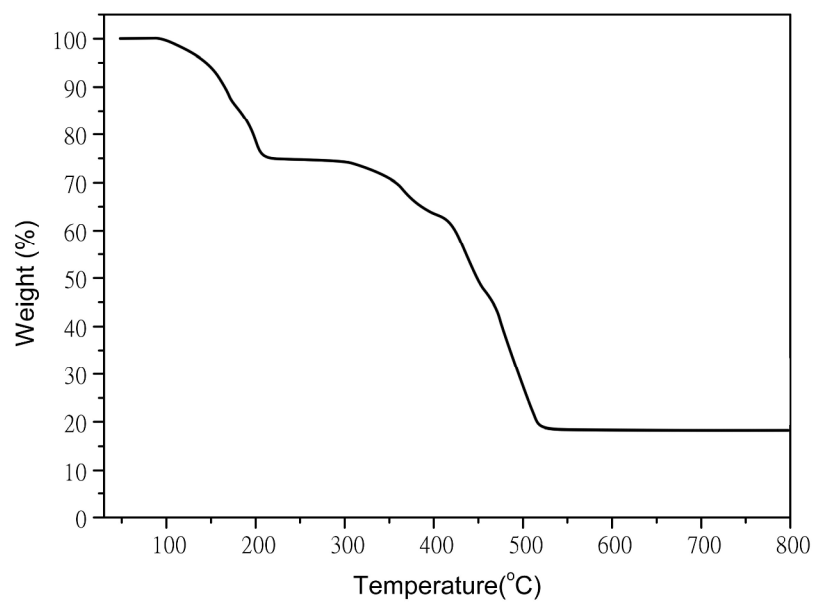
SI-4. Perspective view of the trigonal channel that is surrounded by bpdcb^B ligands. Color code: C black; O red; Cd green.



SI-5. The ^1H NMR spectrum of **1** recorded in D_2O .



SI-6. FT-IR spectrum of as-synthesized **1**.



SI-7. TG curve of 1.