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Supporting information

A new dynamic framework with direct in situ visualisation of breathing under CO2 gas pressure

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1. Synthesis and characterisation

Materials

All chemicals and solvents were purchased from Aldrich and used without further purification. The ligand 4-bis(pyridin-4-ylmethyl)piperazine was prepared according to a literature procedure.¹

Preparation of COB1-DMF

{[Co₂(OBA)₂(BPMP)]·2.5(DMF)_n (COB1-DMF): CO₂(NO₃)₂6H₂O (1 mmol), 4,4'-oxybis(benzoic acid) (OBA, 1 mmol) and (4-bis(pyridin-4-ylmethyl)piperazine (BPMP, 1 mmol) were mixed in 2 ml methanol (MeOH) and dimethylformamide (DMF) and placed in a pre-heated oven at 80 °C. Blue block-shaped crystals of solvated (COB1-DMF) were obtained after 48 h.

+
$$\frac{\text{Co(NO_3)_2 6H_2O, 80 °C 48 h}}{2:1\text{eqv, DMF and MeOH}}$$
 {[Co₂(OBA)₂(BPMP)]·2.5(DMF)}_n (COB1-DMF)

4,4'-oxydibenzoic acid 1,4-bis(pyridin-4ylmethyl)piperazine
(OBA) (BPMP)

Scheme S1. Synthesis of COB1-DMF.

2. Methods

Thermogravimetric analysis (TGA)

Thermogravimetric analysis was carried out using a TA Instruments Q500 analyser. The sample was loaded onto an aluminium pan and heated at 10 °C/min from room temperature to 600 °C.

Activation method of COB1-DMF

Pure bulk material of **COB1-DMF** was activated at 120 °C under dynamic vacuum $(3.1 \times 10^{-2} \text{ mbar})$ for 12 hours to yield the fully desolvated form **COB1-act**. TGA shows that **COB1-act** is desolvated and PXRD analysis shows that it is still crystalline. Direct activation does not yield single crystals suitable for SCD analysis. Single crystals of suitable quality were obtained by activation via solvent exchange with diethyl ether for 2 days, followed by activation in a Schlenck tube at ca 195 K (slurry of dry ice and acetone) under a dynamic vacuum of 3.1×10^{-2} mbar for 48 hours. The desolvated form absorbs moisture and needed to be pre-activated under dynamic vacuum at room temperature for 2 hours before any further experiments were carried out.

Powder X-ray Diffraction (PXRD)

Experiments were carried out on a PANalytical X'Pert PRO instrument with Bragg-Brentano geometry. Intensity data were recorded using an X'Celerator detector and 2θ scans in the range of 5-40° were performed. During the experiment the powdered sample was exposed to Cu-K α radiation (λ = 1.5418 Å). The PXRD patterns of the as-synthesised (**COB1-DMF**) and activated (**COB-act**) samples were recorded at 298 K.

Single-crystal X-ray Diffraction

Single crystal X-ray diffraction data were collected on a Bruker APEX-II Quasar CCD area-detector diffractometer equipped with an Oxford Cryostream 700Plus cryostat, and on a Bruker D8 Venture diffractometer equipped with a PHOTON II CPAD detector and an Oxford Cryostream 800Plus cryostat. A multilayer monochromator with $Mo_{K\alpha}$ radiation ($\lambda=0.71073$ Å) from an Incoatec $I_{\mu S}$ microsource was used. Data reduction was carried out by means of standard procedures using the Bruker software package SAINT² and absorption corrections and the correction of other systematic errors were performed using SADABS.³ The structures were solved by direct methods using SHELXS-2016 and refined using SHELXL-2016⁴ and X-Seed⁵ or Olex2⁵ were used as the graphical interface for the SHELX program suite. Hydrogen atoms were placed in calculated positions using riding models.

Environmental gas cell experiments

An environmental gas cell (developed in-house) was used to determine the crystal structures of the various phases at 298 K under controlled pressures by means of single-crystal X-ray diffraction. First, the structure of COB1⁰ was determined under vacuum after activating a crystal of COB1-DMF. The structure of COB1-CO2¹⁰ was determined after pressurising COB1⁰ with 10 bar of CO₂ gas for 8 hours. The structure of COB1-CO2¹⁹ was determined after pressurising the crystal of COB1-CO2¹⁰ with 19 bar CO₂ and allowing it to equilibrate for 8 hours. The structure of COB1-CO2³⁰ was determined after pressurising COB1-CO2¹⁹ with 30 bar CO₂ and equilibrating for 8 hours. We note that R1 for the gas-loaded structures increases with pressure, and this is most likely due to increasing strain experienced by the crystal.

Volumetric Sorption Analysis

A Setaram PCTPro-E&E gas sorption analyser with a MicroDoser attachment was utilised to conduct high pressure gas sorption experiments with CO_2 and N_2 . The instrument utilises Sievert's volumetric method. The sample temperature was maintained to an accuracy of ± 1 °C using a Grant refrigerated recirculation bath filled with antifreeze and water. A sample at known pressure and volume was connected to a reservoir of known volume and pressure through an isolation valve. The valve was opened, and the system allowed to equilibrate. The difference between the measured and calculated pressures was used to determine the amount of gas adsorbed. National Institute of Standards and Technology (NIST) software was used to calculate the thermodynamic corrections to account for the non-ideal behaviour of the gases at relatively high pressures. The PCTPro-E&E with the MicroDoser attachment is used for small sample sizes and has a range of vacuum to 60 bar. Sample sizes of 80 mg were used and activated in situ using vacuum and heat, if necessary. Blank runs for each gas were recorded to further correct for any other residual systematic errors in the experiment. Figure preparation and data analyses were performed using Microsoft Excel and OriginPro.

Pressure-Gradient Differential Scanning Calorimetry for CO₂ (PG-DSC)

Experiments were carried out using a Setaram Micro-DSC7 Evo instrument. Heat flow was recorded at 298 K in the pressure range of 4-35 bar for CO_2 at 298 K.

Table S1. Crystallographic table of COB1 under controlled pressure of CO₂.

Complex	COB1-DMF	COB1 ⁰	COB1-CO2 ¹⁰	COB1-CO2 ¹⁹	COB1-CO2 ³⁰	
Temperature (K)	100	298	298	298	298	
Empirical formula	$C_{51.50}H_{53.50}Co_2N_{6.50}$ $O_{12.50}$	C ₄₄ H ₃₆ N ₄ O ₁₀ Co ₂	C _{44.9} H ₃₆ N ₄ O _{11.8} Co ₂	C ₄₅ H ₃₆ N ₄ O ₁₃ Co ₂	C ₄₇ H ₃₆ N ₄ O ₁₆ Co ₂	
Formula weight	1081.36	898.66	938.23	958.64	1030.66	
Wavelength (Å)	0.71073	0.71073	0.71073	0.71073	0.71073	
Crystal system	monoclinic	monoclinic	monoclinic	monoclinic	monoclinic	
Space group	$P2_1/n$	C2/c	C2/c	C2/c	C2/c	
a, (Å)	29.7813(12)	23.7504(14)	23.7632(11)	30.071(3)	30.075(5)	
b, (Å)	15.7987(7)	9.7971(6)	9.8043(5)	7.8976(6)	7.8801(13)	
c, (Å)	23.7492(9)	20.3230(12)	20.3897(10)	23.8471(17)	23.860(4)	
β, (°)	101.808(2)	104.156(2)	102.360(2)	101.951(4)	102.013(4)	
Volume (ų)	10937.7(8)	4585.3(5)	4640.3(4)	5540.6(8)	5531.0(16)	
Z	8	4	4	4	4	
Calculated density (g cm ⁻³)	1.313	1.302	1.343	1.149	1.238	
Absorption coefficient (mm ⁻¹)	0.671	0.781	0.772	0.646	0.647	
F ₀₀₀	4496	1848	1927	1968	2112	
Approx. Crystal size (mm³)	0.300 × 0.170 × 0.100	0.268 × 0.135 × 0.088	0.268 × 0.135 × 0.088	0.268 × 0.135 × 0.088	0.268 × 0.135 × 0.088	
θ range for data collection (°)	1.97 to 26.0	2.26 to 26.45	2.21 to 25.09	1.99 to 26.8	1.99 to 21.3	
Miller index ranges	$-36 \le h \le 36$ $-19 \le k \le 19$ $-36 \le l \le 36$	$ -29 \le h \le 29 -12 \le k \le 12, -25 \le l \le 2 $	$ -28 \le h \le 28 $ $ -11 \le k \le 11 $ $ -24 \le l \le 24 $	$-37 \le h \le 36,$ $0 \le k \le 9$ $0 \le 1 \le 30$	$-35 \le h \le 35$ $-9 \le k \le 9$ $-28 \le l \le 28$	
Reflections collected	135400	60406	55175	11597	144436	
Independent reflections	$21507 \\ R_{\text{int}} = 0.1097 \\ R_{\text{sigma}} = 0.0818$	4721 R _{int} =0.1258 R _{sigma} = 0.0611	$\begin{array}{c} 4116 \\ R_{int} = 0.1010 \\ R_{sigma} = 0.0448 \end{array}$	5776 $R_{\text{int}} = 0.1658$ $R_{\text{sigma}} = 0.1368$	$4952 \atop R_{\text{int}} = 0.2636 \atop R_{\text{sigma}} = 0.0658$	
Completeness to θ max (%)	0.995	99.9	99.5	98.0	99.9	
Refinement method		Fu	ll-matrix least-squares	on F ²		
Data / restraints / parameters	21507 / 0 / 1081	4721 /0 / 271	4116 / 0 / 271	5776 / 48 / 430	4952 / 0 / 462	
Goodness-of-fit on F	1.021	1.034	1.021	1.236	1.120	
Final R indices [I > 2 Θ(I)]	R1 = 0.0848 $wR2 = 0.2354$	R1 = 0.0494 $wR2 = 0.1000$	R1 = 0.0428 wR2 = 0.0843	R1 = 0.1368 $wR2 = 0.2858$	R1 = 0.1143 wR2 = 0.2554	
R indices	R1 = 0.1453 wR2 = 0.2855	R1 = 0.1217 w $R_2 = 0.1371$	$R1 = 0.0791$ $wR_2 = 0.0983$	R1 = 0.1658 $wR2 = 0.3045$	R1 = 0.1594 $wR2 = 0.2956$	

Table S2. Structural phase classification at 298 K.

Table S2. Structural phase classification at 298 K.				
COB1-DMF	As-synthesised version containing DMF guest molecules			
COB10	Host framework determined under vacuum			
COB1-CO210	OB1-CO2 ¹⁰ Form determined under 10 bar of CO ₂ gas pressure			
COB1-CO219	Form determined under 19 bar of CO ₂ gas pressure			

Table S3. Selected distances and angles of **COB1-DMF** and **COB1**^x under controlled CO₂ pressures.

Compound	OBA length (centroid _{SBU} ··· centroid _{SBU}) (Å)	OBA Bend angle (centroid _{SBU} ···O _{ether} ···centroid _{SBU}) (°)	BPMP length (centroid _{SBU} ··· centroid _{SBU}) (Å)	Acute angle of pyrazine relative to P_{ac}	Co···Co distance (Å)	Paddlewheel acute angle with P_{ac}
COB1-DMF	14.263	119.764	19.029	26.401	2.629	43.910
COB10	14.141	119.208	18.902	84.11	2.749	35.892
COB1-CO2 ¹⁰	14.144	119.288	18.906	83.903	2.749	35.885
COB1-CO219	14.302	120.444	19.142	25.662	2.727	43.991

1. TGA thermograms

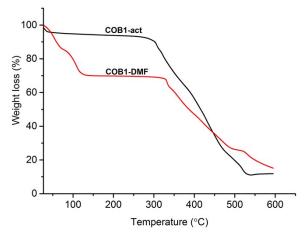


Figure S1. TGA of COB1-DMF and COB1-act.

2. Unit cell dimensions and volume as a function of pressure

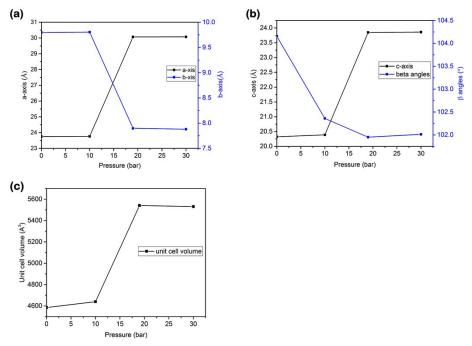


Figure S2. Changes in unit cell parameters for the VP-SCD structures $COB1^x$, where x = 0, 10, 19 and 30. In situ data were collected at 298 K.

3. Variable temperature PG-DSC under CO₂ pressure

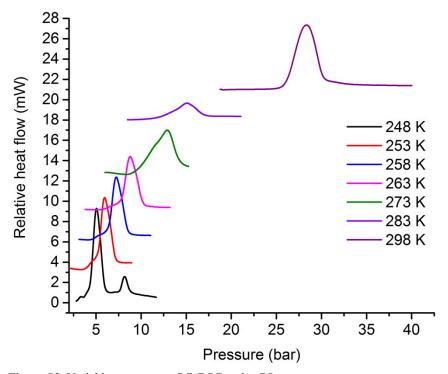


Figure S3. Variable temperature PG-DSC under CO₂ gas pressure.

4. Theoretical investigation

Owing to positional disorder, the apohost crystallographic models obtained under CO_2 pressure were geometry optimized using the CASTEP⁶ code implemented in Materials Studio.⁷ The density functional theory (DFT) generalised gradient approximation of Perdew-Burke-Ernzerhof (PBE)⁸ in combination with Grimme's DFT-D dispersion correction⁹ was employed with on the fly generated Vanderbilt-type ultrasoft pseudopotentials¹⁰ in combination with the Koelling-Harmon scalar-relativistic approach¹¹ and a planewave expansion to an energy cutoff of 489.8 eV. Integration in the reciprocal lattice was performed using a Monkhorst-Pack grid¹² with a 0.05 Å⁻¹ k-point separation and self-consistent field convergence was set to 2.0×10^{-6} eV. A 50% admixture of the charge density¹³ was applied in conjunction with a DIIS (direct inversion in an iterative subspace) ¹⁴ size of 20 to speed up convergence. Convergence tolerances for geometry optimization using the BFGS (Broyden-Fletcher-Goldfarb-Shanno) algorithm¹⁵ were set to 2.0×10^{-5} eV atom⁻¹, 0.05 eV Å⁻¹ and 0.002 Å on energy, maximum force and maximum displacement respectively. To reduce computational expense, calculations were carried out in the primitive cell representation with

$$a' = b' = \sqrt{a^2 + b^2}/2$$

$$c' = c$$

$$\cos(\gamma'/2) = a/\sqrt{a^2 + b^2}$$

$$\cos^2 \alpha' = \cos^2 \beta' = \frac{\sin^2 \gamma' - (2ab\sin \beta/(a^2 + b^2))^2}{2 - 2\cos \gamma'}$$

since V' = V/2, as shown in Figure S4.

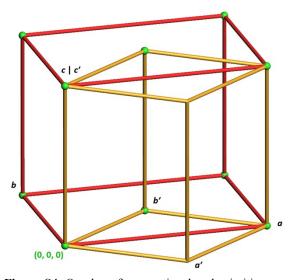


Figure S4. Overlay of conventional and primitive representations of a face-centred monoclinic unit cell in red and orange, respectively. Lattice points are indicated as green spheres.

Table S3. Guest-omitted unit cell energies (kcal mol-1) relative to that of COB10.

COB1-DMF	54.65
COB1 ⁰	0
COB1-CO2 ¹⁰	0.058
COB1-CO2 ¹⁹	45.26
COB1-CO2 ³⁰	56.36

Attempting to model the CO_2 -induced gate opening through an incremental perturbing of the unit cell parameters (10 bar \rightarrow 19 bar) rendered the ligands far removed from their crystallographic orientations. A linear synchronous transit approach was subsequently applied to geometrically follow the structural transformation. A molecular fragment representing a complete coordination sphere of the Co_2OBA_2 paddlewheel was extracted from the geometry optimised 10 bar and 19 bar crystal structures and each atom paired after translocation to the same origin. The resulting pathway shown in Video S1 reveals a major contortion of the BPMP ligand, while the Co_2OBA_2 paddlewheel undergoes a reorientation. Since OBA links paddlewheels along the (0±12) Miller planes, viewing down [0±21] sheds light on the BPMP contribution to the transformation. Similarly, BPMP bridges paddlewheels along (3±10) allowing the [1±30] view to highlight the contortion of the OBA ligand. The rotation of the piperazine moiety from nearly perpendicular (83.90°) to acute (25.66°) during the course of the transformation is discernible in the view down the c axis in **Figure S4** below.

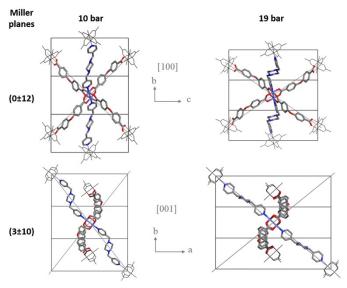


Figure S4. Comparison of CO₂-adsorption induced gate opening structural transformation. Miller planes along which the two ligands interconnect paddlewheel moieties are shown.

5. Description of video files

Video S1 shows the CO₂-induced gate opening transformation of the framework from COB1-CO2¹⁰ to COB1-CO2¹⁹. Expanding the mechanism derived for the fragment into a crystallographic representation, Video S2 shows the gradual increase in solvent-accessible space. These video files are best viewed with the video player set to loop.

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