

Supplementary Material (ESI) for Chemical Communications
This journal is © The Royal Society of Chemistry

An Effective Strategy to Boost the Robustness of Metal-Organic Framework via Introducing Size-Matching Ligand Braces

Xiuli Wang,^{a,b,*§} Wen-Yang Gao,^{b, §} Jian Luan,^a Lukasz Wojtas^b and Shengqian Ma^{b*}

^a *Department of Chemistry, Bohai University, Jinzhou, 121000, P.R. China;* ^b *Department of Chemistry, University of South Florida, 4202 E. Fowler Avenue, Tampa, Florida 33620, USA*

§ Equal contribution

ELECTRONIC SUPPLEMENTARY INFORMATION

General methods

Commercially available reagents were purchased as high purity from Fisher Scientific and used without further purification. 9,10-anthracenedicarboxylic acid (H₂adc) was synthesized using the similar procedures as reported.¹ Thermogravimetric analysis (TGA) was performed under nitrogen on a TA Instrument Q50 from 25°C to 800 °C at the speed of 10 °C/min. FT-IR data were recorded on a PerkinElmer Spectrum Two instrument in a range from 4000 to 500 cm⁻¹ and a resolution of 4 cm⁻¹. Powder X-ray diffraction patterns were collected on a Bruker D8 Advance X-ray diffractometer (CuK α = 1.54178 Å), at 40 kV, 40 mA with a scan speed of 0.5 s/step and a step size of 0.02 in 2 θ at room temperature. Gas adsorption isotherms were collected using a Micromeritics surface area analyzer ASAP-2020. Before the measurements, the freshly prepared samples of complex **1** and **2** were exchanged with HPLC-grade methanol for 3 days, and then activated with the “degas” port under the vacuum at 30 °C for 3 hours. CO₂ gas adsorption isotherms were collected at 195 K using an acetone-dry ice bath, 273K using a water-ice bath and at 298K with a water bath. The O₂, N₂, and H₂ adsorption isotherms were measured at 77K using a liquid nitrogen bath.

Single-crystal X-ray crystallography.

The single crystal X-ray data for complex **1** and **2** were collected using Bruker-AXS

SMART-APEXII diffractometer ($\text{CuK}\alpha = 1.54178 \text{ \AA}$). Indexing was performed using APEX2 (Difference Vectors method).² Data integration and reduction were performed using SaintPlus 6.01.³ Absorption correction was performed by multi-scan method implemented in SADABS.⁴ Space groups were determined using XPREP implemented in APEX2.² Structures were solved using SHELXS-97 (direct methods) and refined using SHELXL-97 (full-matrix least squares on F^2) contained in Win GX^{5,6,7,8} and Olex2 programs.⁹ All non-H atoms were found in the difference Fourier map. The atoms from adc ligands, metal ions, 4,4'-bpy ligands, $\mu_3\text{-O}$ and oxygen atoms of coordinated solvent molecules were refined anisotropically. Due to the disorder, the assignment of counterions was tentative. It has been assumed the dimethyl ammonium cation is a product of decomposition of DMA molecule and that those two share the position in the crystal structure. The geometry of disordered partially-occupied counter ions and coordinated DMA molecules were refined with restraints/constraints (DFIX, DELU, SIMU, DANG, FLAT, EXYZ, EADP, ISOR), in order to idealize the models of these ions and molecules. For complex **2** the solvent molecules were modeled as O atoms. Hydrogen atoms were placed in geometrically calculated positions and included in the refinement process using riding model with isotropic thermal parameters: $\text{Uiso(H)} = 1.2\text{Ueq(-CH)}$. Crystal data and refinement conditions of complex **1** and **2** are shown in Table S1 and S2, respectively.

Preparation of complexes 1 and 2.

Complex 1: *N,N*-dimethylacetamide (DMA, 0.5 mL) solution containing H_2adc (5 mg, 0.019 mmol) was mixed thoroughly with *N,N*-dimethylacetamide (DMA, 0.25 mL) solution containing $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (11 mg, 0.038 mmol), and then three drops of HBF_4 were added. The mixture was sealed in a Pyrex tube, heated at 120 °C for 3 days and then cooled to room temperature. The orange block crystals were obtained with a yield of 45% based on H_2adc .

Complex 2: A mixture of *N,N*-dimethylacetamide (DMA, 0.5 mL) solution containing H_2adc (5 mg, 0.019 mmol) and DMA (0.25 mL) solution containing 4,4'-bpy (3 mg, 0.019 mmol) was mixed thoroughly with DMA (0.25 mL) solution containing $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (11 mg, 0.038 mmol), and then four drops of HBF_4 were added. The mixture was sealed in a Pyrex tube, heated at 120 °C for 4 days and then cooled to room temperature. The red block crystals were obtained

with a yield of 51% based on H_2adc .

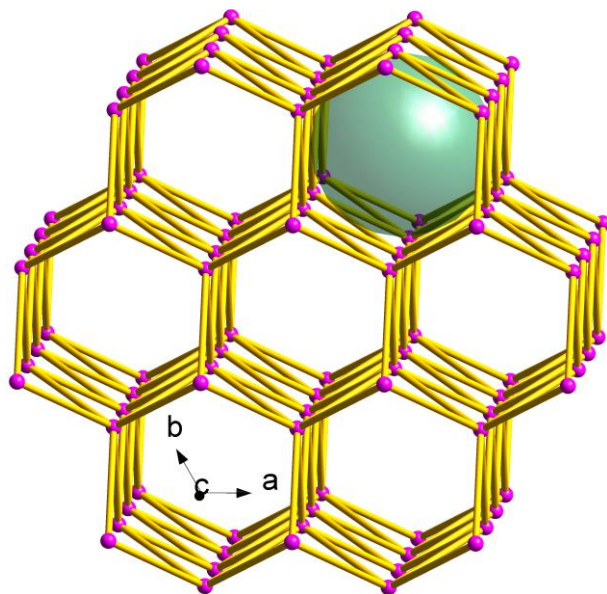


Fig. S1 View of the topological network of complex **1**.

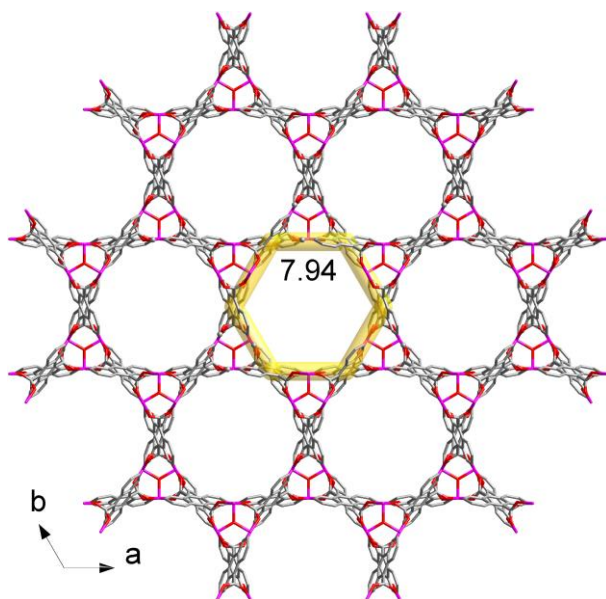


Fig. S2 View of 3D architecture with 1D channels by remove the coordinated DMA molecules in complex **1**.

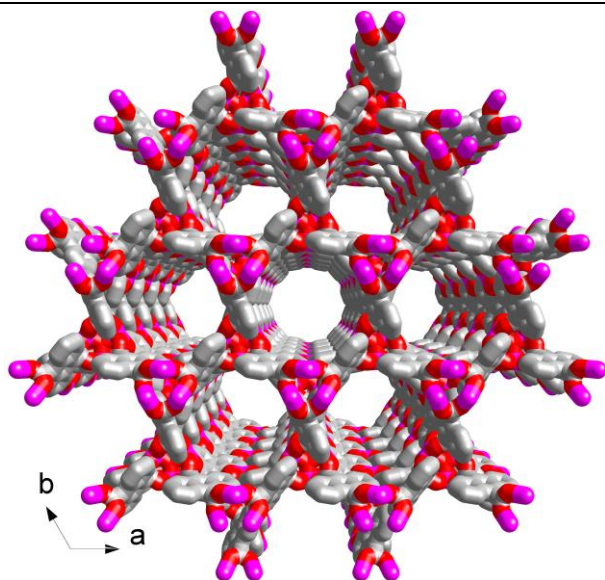


Fig. S3 The 1D nanochannels with terminal DMA removed from tricobalt SBUs in complex **1**.

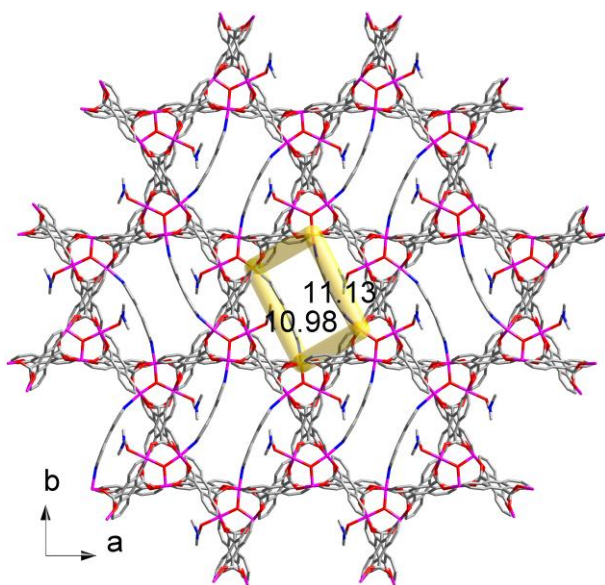


Fig. S4 View of 3D architecture with 1D channels decorated by bpy in complex **2**.

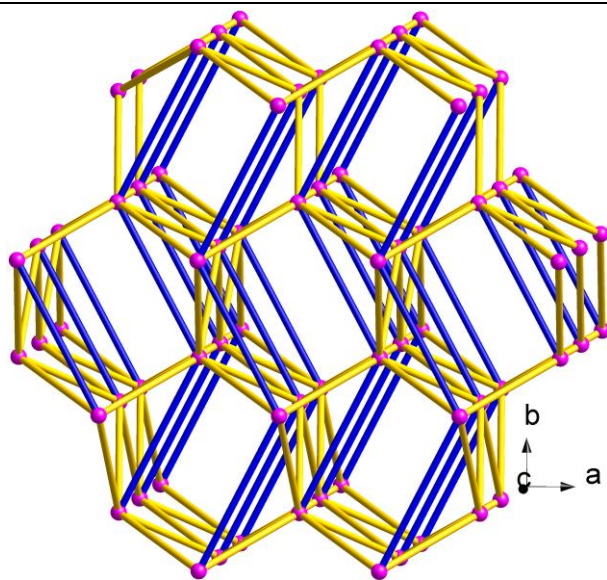


Fig. S5 View of the topological network of complex 2.

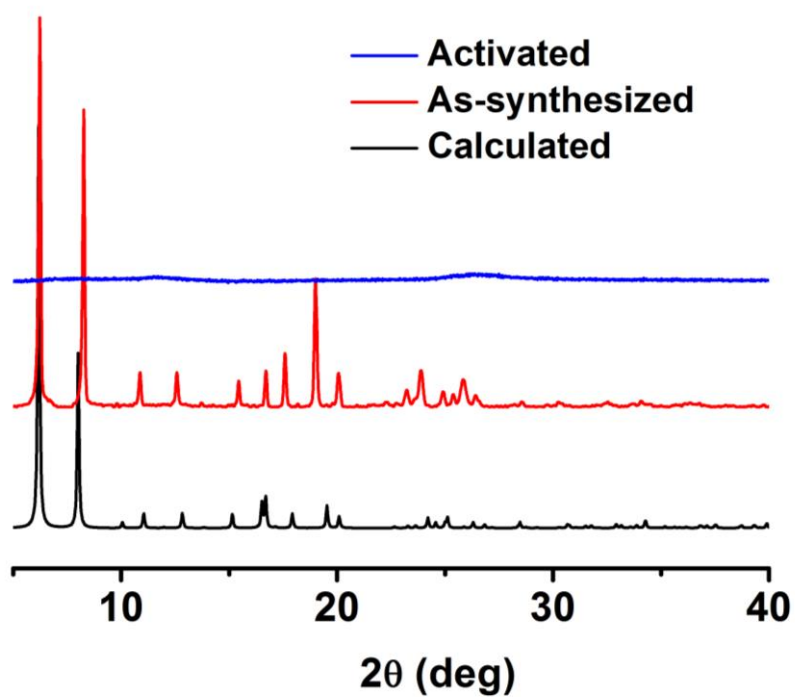


Fig. S6 The PXRD patterns of complexes 1

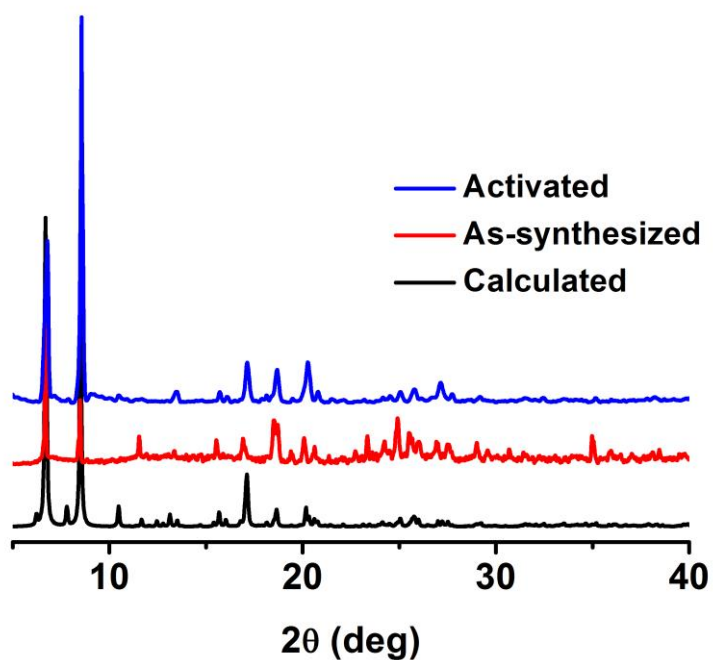


Fig. S7 The PXR D patterns for complex 2.

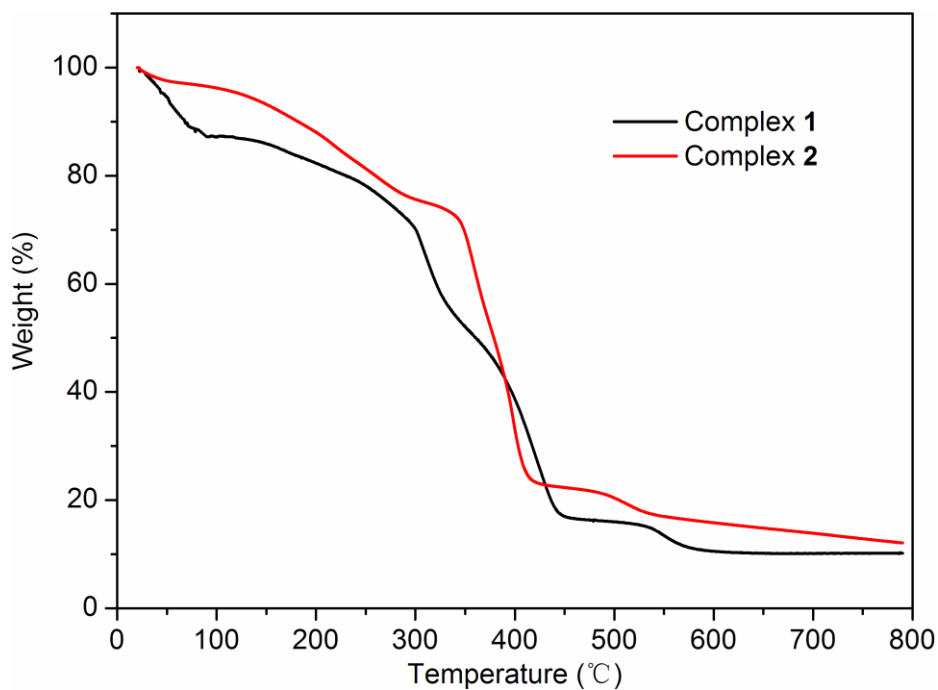


Fig. S8 The TGA curves of complexes 1 and 2.

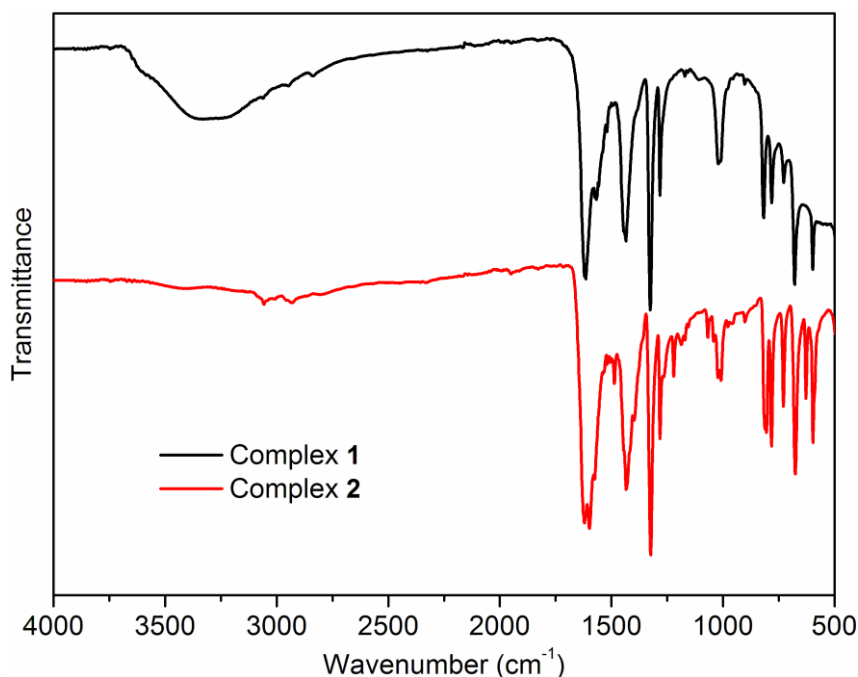


Fig. S9 The FT-IR spectra of complexes 1 and 2.

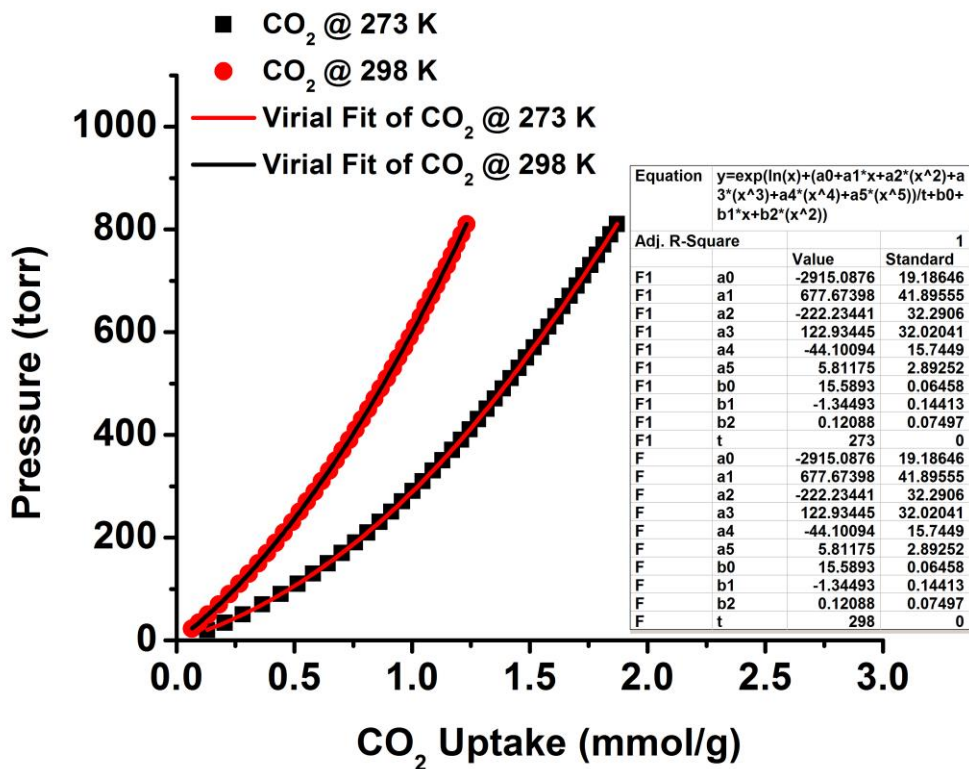


Fig. S10 The plots of virial equation of complex 2.

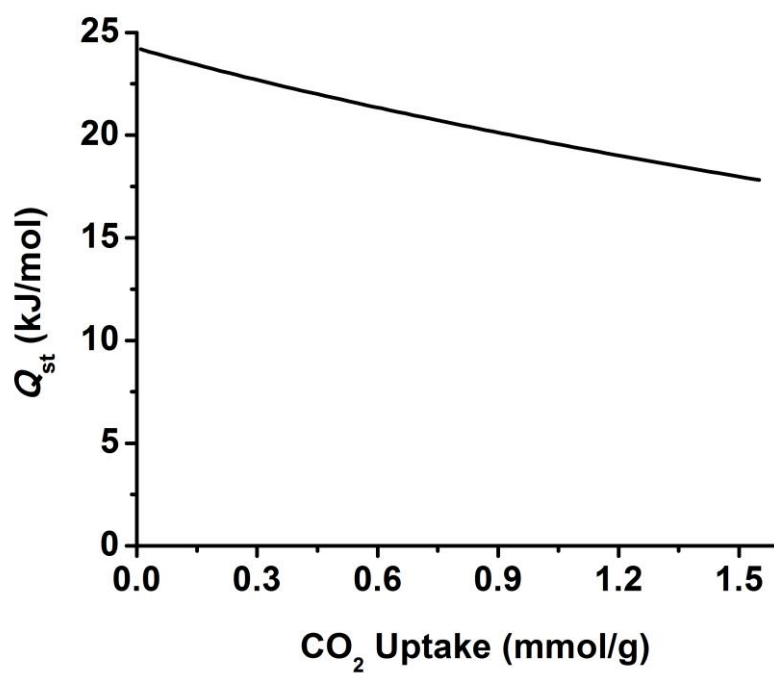


Fig. S11 Heats of adsorption of CO_2 of complex 2 using the virial method.

Identification code	Complex 1
Empirical formula	C ₆₄ H ₂₄ Co ₃ N ₅ O _{18.75}
Formula weight	1339.75
Temperature/K	233.15
Crystal system	trigonal
Space group	<i>P</i> -3 ₁ <i>c</i>
<i>a</i> /Å	15.3816(3)
<i>b</i> /Å	15.3816(3)
<i>c</i> /Å	16.8215(4)
α /°	90
β /°	90
γ /°	120
Volume/Å ³	3446.66(16)
Z	2
ρ_{calc} /g/cm ³	1.291
μ /mm ⁻¹	6.182
F(000)	1348.0
Crystal size/mm ³	0.14 × 0.09 × 0.08
Radiation	CuK α (λ = 1.54178)
2 θ range for data collection/°	6.636 to 133.01
Index ranges	-17 ≤ <i>h</i> ≤ 16, -18 ≤ <i>k</i> ≤ 16, -18 ≤ <i>l</i> ≤ 16
Reflections collected	15678
Independent reflections	1982 [<i>R</i> _{int} = 0.0693, <i>R</i> _{sigma} = 0.0411]
Data/restraints/parameters	1982/80/182
Goodness-of-fit on F ²	1.033
Final <i>R</i> indexes [<i>I</i> >= 2 σ (<i>I</i>)]	<i>R</i> ₁ = 0.0571, <i>wR</i> ₂ = 0.1561
Final <i>R</i> indexes [all data]	<i>R</i> ₁ = 0.0704, <i>wR</i> ₂ = 0.1671
Largest diff. peak/hole / e Å ⁻³	0.70/-0.36

Identification code	Complex 2
Empirical formula	C ₆₉ H ₃₂ C ₃ N _{5.75} O ₁₅
Formula weight	1358.29
Temperature/K	233.15
Crystal system	monoclinic
Space group	<i>P</i> 2 ₁ / <i>n</i>
<i>a</i> /Å	15.6069(4)
<i>b</i> /Å	26.4325(8)
<i>c</i> /Å	16.6221(5)
α /°	90
β /°	90.155(2)
γ /°	90
Volume/Å ³	6857.1(3)
Z	4
ρ_{calc} /g/cm ³	1.316
μ /mm ⁻¹	6.182
F(000)	2749.0
Crystal size/mm ³	0.24 × 0.15 × 0.13
Radiation	CuK α (λ = 1.54178)
2 θ range for data collection/°	8.45 to 137.242
Index ranges	-18 ≤ <i>h</i> ≤ 18, -31 ≤ <i>k</i> ≤ 30, -19 ≤ <i>l</i> ≤ 19
Reflections collected	62684
Independent reflections	12196 [<i>R</i> _{int} = 0.1311, <i>R</i> _{sigma} = 0.1057]
Data/restraints/parameters	12196/149/841
Goodness-of-fit on F ²	1.020
Final R indexes [<i>I</i> >= 2 σ (<i>I</i>)]	<i>R</i> ₁ = 0.0695, <i>wR</i> ₂ = 0.1786
Final R indexes [all data]	<i>R</i> ₁ = 0.1215, <i>wR</i> ₂ = 0.2069
Largest diff. peak/hole / e Å ⁻³	0.85/-0.49

Reference

1. S. Jones, J. C. C. Atherton, M. R. J. Elsegood, W. Clegg, *Acta Cryst.* 2000, **C56**, 881-883.
2. Bruker (2010). *APEX2*. Bruker AXS Inc., Madison, Wisconsin, USA.
3. Bruker (2009). *SAINT*. Data Reduction Software. Bruker AXS Inc., Madison, Wisconsin, USA.
4. G. M. Sheldrick, (2008). *SADABS*. *Program for Empirical Absorption Correction*.

University of Gottingen, Germany.

5. L. J. Farrugia, *J. Appl. Cryst.*, 1999, **32**, 837-838.
6. G. M. Sheldrick, (1997) SHELXL-97. *Program for the Refinement of Crystal*.
7. G. M. Sheldrick, *Acta Cryst.*, 1990, **A46**, 467-473.
8. G. M. Sheldrick, *Acta Cryst.*, 2008, **A64**, 112-122.
9. O. V. Dolomanov, L. J. Bourhis, R. J. Gildea, J. A. K. Howard and H. Puschmann, OLEX2: A complete structure solution, refinement and analysis program, *J. Appl. Cryst.*, 2009, **42**, 339-341.