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Crystal structure and characterization of a new onedimensional copper(II) coordination polymer containing a 4-aminobenzoic acid ligand

Alisha Gogia,^a* Egor M. Novikov,^a Ilia A. Guzei,^b Marina S. Fonari^c and Tatiana V. Timofeeva^a

^aDepartment of Chemistry, New Mexico Highlands University, Las Vegas, New, Mexico, 87701, USA, ^bChemistry Department, University of Wisconsin-Madison, 1101 University Ave, Madison, WI 53706, USA, and ^cInstitute of Applied Physics, Moldova State University, Academy str., 5 MD2028, Chisinau, Moldova. *Correspondence e-mail: alishagogia@nmhu.edu

A Cu^{II} coordination polymer, *catena*-poly[[[aquacopper(II)]-bis(μ -4-aminobenzoato)- $\kappa^2 N:O;\kappa^2 O:N$] monohydrate], {[Cu(pABA)₂(H₂O)]·H₂O}_n (pABA = p-aminobenzoate, C₇H₄NO₂⁻), was synthesized and characterized. It exhibits a one-dimensional chain structure extended into a three-dimensional supramolecular assembly through hydrogen bonds and π - π interactions. While the twinned crystal shows a metrically orthorhombic lattice and an apparent space group *Pbcm*, the true symmetry is monoclinic (space group *P2/c*), with disordered Cu atoms and mixed roles of water molecules (aqua ligand/crystallization water). The luminescence spectrum of the complex shows an emission at 345 nm, *cf.* 349 nm for *pABAH*.

1. Chemical context

Coordination polymers (CPs), which can be categorized in the class of lower dimensional metal-organic frameworks (MOFs), have received great attention in the past few decades owing to the multitude of applications they offer, such as gas storage and separation (Férey, 2008), sensing (Horcajada et al., 2012), drug delivery (Liu et al., 2020), electrochemical applications (Morozan & Jaouen, 2012), adsorption and remediation (Baruah, 2022), magnetic properties (Maspoch et al., 2004), etc. Despite advancements, the anticipation of MOF structures remains an ongoing challenge. Even with reticular synthesis initiated by geometrically analogous ligands, the outcome of structures or ligand behaviors under elevated temperature and pressure conditions, prevalent during synthesis, remains complicated (Szczypiński et al., 2021). Occasionally, in the pursuit of creating porous architectures, our efforts yield coordination polymers with unexpected features. In the present work, we attempted to synthesize a porous metal-organic framework based on Cu^{II} and a flexible tricarboxylic acid ligand, 4,4',4''-{[(1E,1'E,1"E)-benzene-1,3,5trivltris(methaneylylidene)] tris(azaneylylidene)}tribenzoic acid (H₃bttta) (Fig. 1). Instead, we obtained a one-dimensional CP, {[Cu(pABA)₂(H₂O)]·H₂O}_n (I), with the anion of p-aminobenzoic acid (pABAH), the latter presumably formed by disintegration of H₃bttta in the course of hydrothermal synthesis. Subsequently we synthesized compound (I) from $Cu(NO_3)_2$:2.5H₂O and pABAH under the same synthetic conditions. Compound (I) was characterized by single-crystal X-ray diffraction, FTIR spectroscopy and thermogravimetric analysis (TGA).



Its anion, *p*ABA, is capable of versatile binding with metal ions *via* amino and carboxylic groups (Fig. 2), as well as strong hydrogen bonds and π - π stacking interactions, enhancing the overall stability of the CP. Moreover, *p*ABAH has a variety of applications, *viz*. as precursor in the synthesis of pharmaceutical compounds, UV absorbers, components in hair dyes, antioxidants, food additives, *etc*.

2. Structural commentary

Compound (I) crystallizes in a monoclinic space group P2/c, although the unit cell is metrically orthorhombic. The asymmetric unit comprises half of Cu atom, one *p*ABA ligand and one water molecule. The Cu atom is disordered between two alternative sites, Cu1 and Cu2, both located on crystallographic twofold axes, with crystallographic occupancies of 0.3098 (8) and 0.1902 (8), respectively. The carboxylic group is also disordered, the atomic sites C1A and O1A are occupied simultaneously with Cu1 and have occupancies of 0.6196 (16), whereas C1B and O1B are occupied simultaneously with Cu2 and have occupancies of 0.3804 (16). The H atoms of the amino group are also disordered between two sets of positions with the same occupancies, depending on whether the adja-



Figure 1

Tricarboxylic ligand (H_3 btta) used and its fragmentation to *p*ABAH under hydrothermal reaction conditions.



N: η^1 mode, O: η^2 mode

Figure 2

Binding modes of the *p*ABA ligand in coordination polymers with Co^{II} , Ni^{II} , Cu^{II} , Zn^{II} or Cd^{II} (shown as blue spheres).

cent Cu1 or Cu2 site is occupied and coordinated with N1. The disorder is illustrated in Fig. 3.



Figure 3

Disorder in the crystal of (I). (a) The asymmetric unit, showing atomic displacement ellipsoids at the 30% probability level. The major (solid) and minor (stippled) components have occupancies of 0.6196 (16) and 0.3804 (16), respectively. (b), (c) Crystal packing for these components. In the former, molecule O4H₂ acts as an aqua ligand, O3H₂ as crystallization water, and *vice versa* in the latter. Hydrogen bonds are shown as dotted lines, π - π stacking as dashed lines between the centroids of arene rings. Symmetry codes: (1) 1 - x, 1 - y, 1 - z; (2) 1 - x, y, $\frac{3}{2} - z$; (3) x, 1 - y, $\frac{1}{2} + z$.

research communications

0.91

0.91

 $N - H1B \cdot \cdot \cdot O2^{ii}$

 $N-H2B\cdots O3^{iv}$

Table 1Hydrogen-bond geometry (Å, °).						
$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot$		
$O3-H3\cdots O2^i$	0.96(1)	1.84 (1)	2.789 (3)	173 (2)		
$O4-H4\cdots O2^{i}$	0.96(1)	1.88 (2)	2.791 (3)	159 (3)		
$N-H1A\cdots O2^{ii}$	0.91	2.26	2.954 (3)	133		
$N-H2A\cdots O4^{iii}$	0.91	2.17	3.050 (4)	163		

 $-H \cdot \cdot \cdot A$

133

163

2.20 Symmetry codes: (i) x, y + 1, z; (ii) $x, -y + 1, z - \frac{1}{2}$; (iii) -x + 2, -y + 1, -z + 1; (iv) -x + 1, -y + 1, -z + 1.

2.26

2.954 (3)

3.086 (4)

It is noteworthy that the atomic positions (including those of the disordered atoms) approximately comply with the orthorhombic symmetry (apparent space group Pbcm), but their occupancies do not, therefore refinement of the structure in this symmetry gives a computationally unstable, as well as chemically and crystallographically unreasonable, model.

Both the Cu1 and Cu2 sites have an N₂O₃ square-pyramidal coordination environment, in which the apical position is occupied by an aqua ligand (i.e. the O3 or O4 atom, respectively), also located on a twofold axis. Note that the water sites, unlike the Cu ones, are fully occupied. Thus, if the Cu1 site is occupied and Cu2 is vacant, O3H₂ is an aqua ligand and O4H₂ is a water molecule of crystallization and vice versa if the Cu2 site is occupied.

The pABA ligand bridges two adjacent Cu atoms (related by the *c* glide plane) through amine nitrogen and carboxylate oxygen atoms in a μ_2 -O:N binding mode. Thus each Cu atom is linked with two symmetry-equivalent ones by pairs of antiparallel pABA ligands (whose two O and two N atoms comprise the basal plane of the pyramid), to form a polymeric chain parallel to the c axis.

3. Supramolecular features

The one-dimensional catena-Cu(pABA) chains of (I) are combined into a three-dimensional supramolecular structure by a network of hydrogen bonds (Table 1). Both water molecules (whether coordinated or not) donate hydrogen bonds to the non-coordinated carboxylic atom O2 (and its equivalents). forming an infinite zigzag chain $O2 \cdots H - O3 - H \cdots O2 \cdots H - O4 - H \cdots O2$ along the *a*-axis direction. The amino group, which is disordered over two orientations (see above), in either case donates one hydrogen bond to a trans-annular O2 and the other to the water molecule, which is not coordinated (the adjacent Cu site being vacant). Thus, while an aqua ligand donates two hydrogen bonds, the crystallization water at the same site donates two and accepts two, from different adjacent Cu(pABA) chains.

There is $\pi - \pi$ stacking of practically parallel arene rings of pABA (Fig. 3). Infinite stacks run parallel to the a axis, with alternating interplanar separations of 3.41(6) and 3.49(6) Å, lateral shifts between adjacent rings of 1.72 (8) and 1.42 (9) Å, and distances between ring centroids of 3.82 (4) and 3.77 (4) Å, respectively.



FTIR spectra of *p*ABAH (black) and compound (I) (red).

4. Spectroscopic and thermal properties

The FTIR spectra of pABAH and (I) (Fig. 4) demonstrated successful incorporation of the pABA ligand in (I). In comparison to the free ligand, pABAH, the peaks corresponding to the amine group suffer a decrease in the wavenumber and intensity upon binding to the Cu^{II} atom in (I), similar to what is observed in other cases in the literature (Crisan *et al.*, 2019). In addition, the peak at 1661 cm^{-1} , corresponding to the free carboxylic acid in pABAH is diminished upon metal coordination in (I), Fig. 4. The strong bands at 1606 cm⁻¹ and 1404 cm⁻¹ correspond to the asymmetric (v_{asym}) and symmetric (v_{sym}) stretching vibrations of the carboxylate group of pABA in (I). The difference in the asymmetric and symmetric vibrations ($\Delta v = 202 \text{ cm}^{-1}$) corresponds to monodentate binding of the carboxylate which corroborates well with the structure of (I).





The stability of (I) was studied by thermal gravimetric analysis in the range of 30-500°C, which shows that (I) is stable up to 300°C. The initial loss of 2 wt% corresponds to the loss of coordinated water molecules, and the complete decomposition (94 wt%) corresponds to the evolution of CO₂ upon the decomposition of the carboxylate group in the ligand, pABA, leaving behind metal oxide ash (Fig. 5). The percentage of ash left behind is surprisingly lower than expected and might be due to the heterogeneity of the material.

5. Luminescence properties

The emission spectra of (I) and the *p*ABA ligand were recorded at room temperature to assess the luminescence properties of the samples. For this, 1 mg of each sample was finely dispersed in 2 mL of water through ultrasonication. Their respective emission spectra were then recorded at an excitation wavelength of 280 nm, and excitation and emission slit widths of 1 and 1 nm, respectively, in the range 300 to 450 nm. It was found that the emission intensity of (I) is much more intense compared to the emission intensity of the pure *p*ABAH ligand in water. Compound (I) also undergoes a slight blue shift of $\Delta\lambda = 4$ nm, which is representative of the binding of ligand (*p*ABA) with the metal center (Cu^{II}) (Fig. 6).

6. Database survey

Although *p*ABA is widely used as a ligand in the synthesis of coordination polymers and metal–organic frameworks, a survey of the Cambridge Structural Database (version 5.45, updated on 01/01/2024; Groom *et al.*, 2016) revealed no Cu complexes containing only *p*ABA ligands and coordinated or crystallization water, while such complexes are known for Co^{II} , Ni^{II}, Zn^{II} and Cd^{II}. Most of these are one-dimensional coordination polymers, although $[Co(pABA)(H_2O)_4]$



Figure 6

Luminescence emission spectra of pure *p*ABAH and (I) measured at room temperature in water ($\lambda_{\text{excitation}} = 280 \text{ nm}$).

Tá	able 2	
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Experiment	tal c	letail	s.
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Crystal data	
Chemical formula	$[Cu(C_7H_4NO_2)_2(H_2O)]\cdot H_2O$
M _r	371.83
Crystal system, space group	Monoclinic, P2/c
Temperature (K)	100
a, b, c (Å)	6.9143 (14), 6.2111 (12), 17.169 (3)
β (°)	90.05 (3)
$V(A^3)$	737.3 (3)
Ζ	2
Radiation type	Μο Κα
$\mu \text{ (mm}^{-1})$	1.52
Crystal size (mm)	$0.3 \times 0.2 \times 0.2$
Data collection	
Diffractometer	Bruker SMART APEXII
Absorption correction	Multi-scan (SADABS; Krause et al., 2015)
T_{\min}, T_{\max}	0.618, 0.745
No. of measured, independent and	8617, 1477, 1311
observed $[I > 2\sigma(I)]$ reflections	
R _{int}	0.027
$(\sin \theta / \lambda)_{\rm max} ({\rm \AA}^{-1})$	0.642
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.033, 0.087, 1.09
No. of reflections	1477
No. of parameters	131
No. of restraints	4
H-atom treatment	H atoms treated by a mixture of independent and constrained
$\Delta \rho = \Delta \rho + (e \text{ Å}^{-3})$	0.37 - 0.29
$\rightarrow P \max$, $\rightarrow P \min (\circ I \bullet)$	0.07, 0.22

Computer programs: APEX2 (Bruker, 2019), SAINT-Plus (Bruker, 2020), SHELXT (Sheldrick, 2015a), SHELXL (Sheldrick, 2015b), and OLEX2 (Dolomanov et al., 2009).

(ABZACO10; Amiraslanov *et al.*, 1979*a*) crystallizes as discrete molecular units, $[Zn(pABA)_2(H_2O)] \cdot H_2O$ (IWORET; Ibragimov *et al.*, 2016) as a two-dimensional polymer, and $[Zn(pABA)_2] \cdot H_2O$ (RUPZIM; Li *et al.*, 2009) as a threedimensional MOF. The carboxylic group of *p*ABA is usually monodentate (Amiraslanov *et al.*, 1978; Prondzinski & Merz, 2008), except in Cd^{II} complexes ABZCUH (Amiraslanov *et al.*, 1979*b*) and BESRAS (Turner, *et al.*, 1982), where it is bidentate, and in RUPZIM where both mono- and bidentate coordination is present. Thus, compound (I) shows the most typical structural features, being a 1D coordination polymer with the *p*ABA bridge coordinated *via* the amino group and one carboxylic O atom (Fig. 2*b*).

It is noteworthy that an isomer of the two-dimensional polymer IWORET (IWORET01; Crisan *et al.*, 2019) is onedimensional and essentially isostructural with (I), with the same space group P2/c and similar unit-cell parameters, a = 7.0013 (4), b = 6.1301 (2), c = 17.1919 (7) Å, $\beta = 92.148$ (4)°, albeit without disorder. Another isomer of these, YIMDEO (Prondzinski & Merz, 2008) is 1D-polymeric, but with a tetrahedral (O₃N) metal coordination and different *p*ABA modes (Fig. 2*a*,*b*).

7. Synthesis and crystallization

Synthesis of (I). A mixture of $Cu(NO_3)_2 \cdot 2.5H_2O$ (117 mg, 0.5 mmol), pABAH (68.6 mg, 0.5 mmol) and 10 mL of H₂O was placed in a 15 mL stainless steel-jacketed Teflon reactor.

The reactor was carefully sealed, placed in the center of a programmable oven (Nabertherm 30–3000°C, S/N. 432847, 2022), and subjected to heating at a gradual rate of 0.1 K min⁻¹ to 358 K, kept at the same temperature for a duration of 24 h, followed by gradual cooling of K min⁻¹ to 298 K over 12 h. This afforded green block-shaped clear crystals. The obtained crystals were collected *via* filtration, washed with water (3 × 4 mL), then with ethanol (2 × 4 mL) and air-dried. Yield: 58 mg (65%), based on metal salt. Selected FTIR peaks (KBr, cm⁻¹): 3250 (*br*), 3139 (*br*), 1606 (*s*), 1576 (*s*), 1304 (*s*), 1092 (*m*), 854 (*w*), 775 (*m*). The reaction synthesis is similar to that synthesized with H₃bttta, except that 0.034 mmol (174 mg) of H₃bttta were used instead of 0.5 mmol (34.8 mg) of *p*ABAH.

8. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The crystal studied was a merohedral twin with the twin components of equal size related by a 180° rotation about the *c* axis. The water H atoms were refined in isotropic approximation, other H atoms as riding in idealized positions, with $U_{iso}(H) = 1.2 \times U_{eq}$ of the bearing C or N atom.

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Crystal structure and characterization of a new one-dimensional copper(II) coordination polymer containing a 4-aminobenzoic acid ligand

Alisha Gogia, Egor M. Novikov, Ilia A. Guzei, Marina S. Fonari and Tatiana V. Timofeeva

Computing details

catena-Poly[[[aquacopper(II)]-bis(μ-4-aminobenzoato)-κ²N:O;κ²O:N] monohydrate]

Crystal data

[Cu(C₇H₄NO₂)₂(H₂O)]·H₂O $M_r = 371.83$ Monoclinic, P2/c a = 6.9143 (14) Å b = 6.2111 (12) Å c = 17.169 (3) Å $\beta = 90.05$ (3)° V = 737.3 (3) Å³ Z = 2

Data collection

Bruker SMART APEXII diffractometer Radiation source: sealed X-ray tube, EIGENMANN GmbH Graphite monochromator Detector resolution: 7.9 pixels mm⁻¹ ω and φ scans Absorption correction: multi-scan (SADABS; Krause *et al.*, 2015)

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.033$ $wR(F^2) = 0.087$ S = 1.091477 reflections 131 parameters 4 restraints Primary atom site location: dual F(000) = 382 $D_x = 1.675 \text{ Mg m}^{-3}$ Mo K\alpha radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 2372 reflections $\theta = 3.0-26.0^{\circ}$ $\mu = 1.52 \text{ mm}^{-1}$ T = 100 KBlock, clear dark green $0.3 \times 0.2 \times 0.2 \text{ mm}$

 $T_{\min} = 0.618, T_{\max} = 0.745$ 8617 measured reflections 1477 independent reflections 1311 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.027$ $\theta_{\text{max}} = 27.1^{\circ}, \theta_{\text{min}} = 1.2^{\circ}$ $h = -8 \rightarrow 8$ $k = -7 \rightarrow 4$ $l = -20 \rightarrow 21$

Hydrogen site location: mixed H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0395P)^2 + 0.5003P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.37$ e Å⁻³ $\Delta\rho_{min} = -0.29$ e Å⁻³

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refined as a 2-component twin.

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	Occ. (<1)
Cu1	0.500000	0.70928 (13)	0.750000	0.0206 (2)	0.6196 (16)
Cu2	1.000000	0.7095 (2)	0.750000	0.0204 (4)	0.3804 (16)
O1A	0.6443 (6)	0.6835 (6)	0.6512 (2)	0.0262 (8)	0.6196 (16)
O1B	0.8560 (9)	0.6813 (9)	0.6519 (4)	0.0243 (13)	0.3804 (16)
O2	0.7512 (5)	0.3523 (3)	0.68132 (10)	0.0363 (5)	
O3	0.500000	1.0567 (5)	0.750000	0.0468 (9)	
Н3	0.586 (4)	1.1507 (11)	0.7225 (18)	0.070*	
O4	1.000000	1.0536 (5)	0.750000	0.0450 (9)	
H4	0.933 (5)	1.1481 (11)	0.7152 (15)	0.067*	
Ν	0.7528 (6)	0.3218 (3)	0.30662 (11)	0.0259 (5)	
H2A	0.820104	0.197454	0.299519	0.031*	0.6196 (16)
H1A	0.820685	0.428119	0.282487	0.031*	0.6196 (16)
H1B	0.684525	0.428327	0.282761	0.031*	0.3804 (16)
H2B	0.685060	0.197642	0.299792	0.031*	0.3804 (16)
C1A	0.7092 (8)	0.5003 (14)	0.6330 (5)	0.0224 (11)	0.6196 (16)
C1B	0.7949 (14)	0.492 (3)	0.6320 (9)	0.0224 (11)	0.3804 (16)
C2	0.7486 (7)	0.4541 (4)	0.54737 (13)	0.0284 (6)	
C3	0.7403 (8)	0.6201 (4)	0.49334 (15)	0.0424 (8)	
H3A	0.731206	0.764858	0.510783	0.051*	
C4	0.7451 (8)	0.5766 (4)	0.41429 (14)	0.0344 (7)	
H4A	0.742225	0.691677	0.377838	0.041*	
C5	0.7542 (8)	0.3680 (4)	0.38847 (13)	0.0268 (5)	
C6	0.7623 (8)	0.2011 (4)	0.44200 (17)	0.0494 (10)	
H6	0.768947	0.056284	0.424427	0.059*	
C7	0.7608 (8)	0.2452 (4)	0.52089 (16)	0.0420 (8)	
H7	0.768247	0.130184	0.557248	0.050*	

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (A^2)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cu1	0.0307 (4)	0.0221 (4)	0.0090 (3)	0.000	-0.0030 (12)	0.000
Cu2	0.0274 (7)	0.0220 (6)	0.0119 (6)	0.000	-0.005 (2)	0.000
O1A	0.038 (2)	0.031 (2)	0.0097 (17)	0.0077 (18)	0.0008 (19)	0.0004 (17)
O1B	0.037 (3)	0.016 (3)	0.020 (3)	-0.005 (3)	-0.002 (4)	0.000 (3)
02	0.0657 (13)	0.0277 (8)	0.0155 (8)	0.0000 (13)	0.001 (2)	0.0047 (7)
03	0.0460 (19)	0.0244 (16)	0.070 (2)	0.000	0.017 (4)	0.000
O4	0.0417 (18)	0.0358 (18)	0.057 (2)	0.000	-0.017 (4)	0.000
Ν	0.0398 (13)	0.0253 (9)	0.0127 (9)	0.0015 (17)	-0.001 (2)	-0.0010 (8)

supporting information

C1A	0.027 (3)	0.0214 (14)	0.0188 (14)	0.002 (4)	0.002 (4)	-0.0026 (11)
C1B C2	0.027 (3)	0.0214(14) 0.0221(12)	0.0188(14) 0.0121(11)	0.002 (4)	0.002 (4)	-0.0026(11) -0.0001(9)
C3	0.093 (3)	0.0178 (12)	0.0171 (12)	-0.006 (3)	-0.002(3)	-0.0025 (10)
C4	0.066 (2)	0.0218 (11)	0.0155 (11)	0.000 (2)	0.003 (3)	0.0031 (9)
C5	0.0412 (14)	0.0268 (12)	0.0123 (10)	0.002 (2)	0.000 (2)	-0.0026 (9)
C6	0.107 (3)	0.0222 (13)	0.0189 (13)	0.008 (3)	0.000 (3)	-0.0037 (10)
C7	0.090 (3)	0.0196 (11)	0.0167 (13)	0.004 (2)	-0.001 (2)	0.0046 (10)

Geometric parameters (Å, °)

Cu1—O1A ⁱ	1.975 (4)	N—H2A	0.9100
Cu1—O1A	1.975 (4)	N—H1A	0.9100
Cu1—O3	2.158 (3)	N—H1B	0.9100
Cu1—N ⁱⁱ	2.009 (4)	N—H2B	0.9100
Cu1—N ⁱⁱⁱ	2.009 (4)	N—C5	1.434 (3)
Cu2—O1B	1.964 (6)	C1A—C2	1.523 (9)
Cu2—O1B ^{iv}	1.964 (6)	C1B—C2	1.506 (16)
Cu2—O4	2.137 (3)	C2—C3	1.388 (3)
Cu2—N ⁱⁱ	1.976 (4)	C2—C7	1.377 (3)
Cu2—N ^v	1.976 (4)	С3—НЗА	0.9500
O1A—C1A	1.262 (9)	C3—C4	1.384 (3)
O1B—C1B	1.294 (17)	C4—H4A	0.9500
O2—C1A	1.272 (9)	C4—C5	1.371 (3)
O2—C1B	1.251 (16)	C5—C6	1.386 (4)
O3—H3 ⁱ	0.957 (3)	С6—Н6	0.9500
O3—H3	0.957 (3)	C6—C7	1.382 (4)
O4—H4	0.958 (3)	С7—Н7	0.9500
O4—H4 ^{iv}	0.958 (3)		
O1A ⁱ —Cu1—O1A	170.7 (2)	C5—N—Cu2 ^v	119.8 (3)
O1A—Cu1—O3	94.66 (12)	C5—N—H2A	107.4
O1A ⁱ —Cu1—O3	94.66 (12)	C5—N—H1A	107.4
O1A ⁱ —Cu1—N ⁱⁱ	90.96 (15)	C5—N—H1B	107.4
O1A ⁱ —Cu1—N ⁱⁱⁱ	88.14 (15)	C5—N—H2B	107.4
O1A—Cu1—N ⁱⁱⁱ	90.96 (15)	O1A—C1A—O2	124.8 (7)
O1A—Cu1—N ⁱⁱ	88.14 (15)	O1A—C1A—C2	118.3 (7)
N ⁱⁱⁱ —Cu1—O3	95.52 (6)	O2—C1A—C2	116.9 (6)
N ⁱⁱ —Cu1—O3	95.52 (6)	O1B—C1B—C2	117.8 (12)
N ⁱⁱⁱ —Cu1—N ⁱⁱ	168.96 (12)	O2—C1B—O1B	122.1 (13)
O1B ^{iv} —Cu2—O1B	169.8 (3)	O2—C1B—C2	119.5 (11)
O1B—Cu2—O4	95.11 (17)	C3—C2—C1A	119.8 (4)
O1B ^{iv} —Cu2—O4	95.11 (17)	C3—C2—C1B	122.4 (7)
O1B ^{iv} —Cu2—N ⁱⁱ	90.4 (2)	C7—C2—C1A	120.5 (4)
O1B—Cu2—N ⁱⁱ	88.6 (2)	C7—C2—C1B	117.0 (6)
N ⁱⁱ —Cu2—O4	95.64 (7)	C7—C2—C3	118.8 (2)
N ^v —Cu2—O4	95.64 (7)	С2—С3—НЗА	119.7
N^{ii} —Cu2—N ^v	168.72 (13)	C4—C3—C2	120.6 (2)

C1A—O1A—Cu1	117.8 (5)	С4—С3—Н3А	119.7
C1B—O1B—Cu2	118.2 (8)	C3—C4—H4A	119.9
Cu1—O3—H3 ⁱ	127.6 (4)	C5—C4—C3	120.2 (2)
Cu1—O3—H3	127.6 (4)	C5—C4—H4A	119.9
H3—O3—H3 ⁱ	104.8 (8)	C4—C5—N	120.4 (2)
Cu2—O4—H4	127.8 (4)	C4—C5—C6	119.6 (2)
$H4-O4-H4^{iv}$	104.4 (8)	C6—C5—N	120.0 (2)
Cu1 ⁱⁱⁱ —N—H2A	107.4	С5—С6—Н6	120.0
Cu1 ⁱⁱⁱ —N—H1A	107.4	C7—C6—C5	120.1 (2)
Cu2 ^v —N—H1B	107.4	С7—С6—Н6	120.0
Cu2 ^v —N—H2B	107.4	C2—C7—C6	120.7 (2)
H2A—N—H1A	106.9	С2—С7—Н7	119.6
H1B—N—H2B	106.9	С6—С7—Н7	119.6
C5—N—Cu1 ⁱⁱⁱ	119.9 (3)		
Cu1—O1A—C1A—O2	-26.2 (8)	O2—C1B—C2—C3	161.0 (7)
Cu1—O1A—C1A—C2	156.5 (4)	O2—C1B—C2—C7	-34.5 (11)
$Cu1^{iii}$ N—C5—C4	-86.6 (6)	NC5C6C7	-179.1 (5)
Cu1 ⁱⁱⁱ —N—C5—C6	92.6 (5)	C1A—C2—C3—C4	-169.7 (5)
Cu2—O1B—C1B—O2	30.0 (12)	C1A—C2—C7—C6	168.5 (5)
Cu2—O1B—C1B—C2	-159.1 (6)	C1B—C2—C3—C4	163.9 (6)
Cu2 ^v —N—C5—C4	93.1 (5)	C1B—C2—C7—C6	-165.9 (7)
Cu2 ^v —N—C5—C6	-87.7 (5)	C2—C3—C4—C5	1.4 (9)
O1A—C1A—C2—C3	9.4 (9)	C3—C2—C7—C6	-0.8 (8)
O1A—C1A—C2—C7	-159.8 (5)	C3—C4—C5—N	178.0 (5)
O1B—C1B—C2—C3	-10.2 (12)	C3—C4—C5—C6	-1.3 (9)
O1B—C1B—C2—C7	154.3 (7)	C4—C5—C6—C7	0.2 (9)
O2—C1A—C2—C3	-168.2 (5)	C5—C6—C7—C2	0.9 (9)
O2—C1A—C2—C7	22.7 (8)	C7—C2—C3—C4	-0.3 (8)

Symmetry codes: (i) -x+1, y, -z+3/2; (ii) x, -y+1, z+1/2; (iii) -x+1, -y+1, -z+1; (iv) -x+2, y, -z+3/2; (v) -x+2, -y+1, -z+1.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	$D \cdots A$	D—H···A	
O3—H3…O2 ^{vi}	0.96(1)	1.84 (1)	2.789 (3)	173 (2)	
$O4$ — $H4$ ··· $O2^{vi}$	0.96 (1)	1.88 (2)	2.791 (3)	159 (3)	
N—H1A····O2 ^{vii}	0.91	2.26	2.954 (3)	133	
$N - H2A - O4^{v}$	0.91	2.17	3.050 (4)	163	
N—H1 B ····O2 ^{vii}	0.91	2.26	2.954 (3)	133	
N—H2B····O3 ⁱⁱⁱ	0.91	2.20	3.086 (4)	163	

Symmetry codes: (iii) -*x*+1, -*y*+1, -*z*+1; (v) -*x*+2, -*y*+1, -*z*+1; (vi) *x*, *y*+1, *z*; (vii) *x*, -*y*+1, *z*-1/2.