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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.009 Å R factor = 0.094 wR factor = 0.170 Data-to-parameter ratio = 13.2

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catena-Poly[[μ -1,2-di-4-pyridylethylene- $\kappa^2 N$,N'-[aqua(2-sulfonatobenzoato- $\kappa^2 O$,O')copper(II)]] N,N-dimethylformamide]

In the title compound, $[Cu(C_{12}H_8N_2)(C_7H_4SO_5)(H_2O)]$ - C_3H_7NO , the Cu^{II} atom has a square-pyramidal coordination geometry formed by three O atom and two N atoms. The 1,2-di-4-pyridylethylene ligands function as μ_2 -bridging ligands to form a zigzag chain. The 2-sulfonatobenzoate ligands protrude on alternate sides of the chain. Intermolecular O-H···O hydrogen bonds link the chains into a two-dimensional network structure.

Comment

The 2-sulfobenzoic acid $(o-H_2sb)$ ligand, containing one sulfonate group and one carboxyl group, exhibits diverse coordination behaviour in the preparation of metal–organic coordination polymers (Li & Yang, 2004; Su *et al.*, 2005; Xiao *et al.*, 2005). Numerous novel coordination polymers with the 1,2-di-4-pyridylethylene (bpe) ligand have been synthesized in recent years (Hong & You, 2004; Kondo *et al.*, 2004; Yin & Xiao, 2005). Here, we present the crystal structure of the title compound, (I), from the same family.



In compound (I), the Cu^{II} atom has a square-pyramidal environment defined by one water O atom, two O atoms belonging to the 2-sulfonatobenzoate ligand and two N atoms from the two bpe ligands (Fig. 1). In the basal plane, the $N2^{i}$ -Cu1-N1 [symmetry code: (i) $x, \frac{1}{2} - y, z - \frac{1}{2}$] and O1-Cu1-O6 bond angles are 163.59 (19) and 179.47 $(19)^\circ$, respectively, while the other angles around the Cu^{II} centre are in the range 86.00 (17)–104.44 (17)°. The apical position is occupied by sulfonate atom O3, with an axial Cu-O bond distance of 2.268 (4) Å (Table 1). The *o*-sb ligand chelates to the Cu^{II} centre to form a six-membered ring, leading to a coordination mode observed previously in [Ni(o-sb)(bpe)(H₂O)₂]·0.25H₂O (Xiao et al., 2005). The dihedral angle between the planes of the o-sb ring and its carboxylate group is 62.5 (2)°. The C13-O1 bond [1.307 (7) Å] is longer than C13–O2 [1.212 (7) Å], indicating more keto character in the latter. Bpe ligands

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metal-organic papers

function as μ_2 -bridging ligands, forming a zigzag polymeric chain. The *o*-sb ligands protrude on alternate sides of this chain (Fig. 2). Intermolecular O-H···O hydrogen bonds (Table 2) link the chains into a two-dimensional network structure (Fig. 3).

Experimental

An aqueous solution (10 ml) containing $\text{Cu}(\text{ClO}_4)_2 \cdot 6\text{H}_2\text{O}$ (0.25 mmol, 0.910 g) was added slowly to a solution (10 ml) of *N*,*N*dimethylformamide containing 1,2-di-4-pyridylethylene (0.25 mmol, 0.991 g) and 2-sulfobenzoic acid (0.25 mmol, 0.g). The mixture was left to stand at room temperature. Blue crystals of (I) were obtained after 5 d.

 $D_r = 1.579 \text{ Mg m}^{-3}$

Cell parameters from 2377

 $0.20 \times 0.17 \times 0.15~\mathrm{mm}$

 $w = 1/[\sigma^2(F_0^2) + (0.0339P)^2]$

+ 17.9772*P*] where $P = (F_0^2 + 2F_c^2)/3$

 $\Delta \rho_{\rm min} = -0.81 \text{ e } \text{\AA}^{-3}$

 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.64 \text{ e} \text{ Å}^{-3}$

Mo $K\alpha$ radiation

reflections

 $\theta = 2.5 - 21.8^{\circ}$ $\mu = 1.11 \text{ mm}^{-1}$

T = 298 (2) K

Prism, blue

Crystal data

$$\begin{split} & [\mathrm{Cu}(\mathrm{C}_{12}\mathrm{H}_8\mathrm{N}_2)(\mathrm{C}_7\mathrm{H}_4\mathrm{SO}_3)(\mathrm{H}_2\mathrm{O})] & \cdot \\ & \mathrm{C}_3\mathrm{H}_7\mathrm{NO} \\ & M_r = 537.03 \\ & \mathrm{Orthorhombic}, Pbca \\ & a = 17.3586 \ (19) \ \mathrm{\AA} \\ & b = 9.8854 \ (11) \ \mathrm{\AA} \\ & c = 26.324 \ (3) \ \mathrm{\AA} \\ & V = 4517.1 \ (9) \ \mathrm{\AA}^3 \\ & Z = 8 \end{split}$$

Data collection

Bruker SMART CCD area-detector	4102 independent reflections
diffractometer	3679 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.071$
Absorption correction: multi-scan	$\theta_{\rm max} = 25.3^{\circ}$
(SADABS; Bruker, 2002)	$h = -19 \rightarrow 20$
$T_{\min} = 0.809, T_{\max} = 0.851$	$k = -11 \rightarrow 11$
22 879 measured reflections	$l = -28 \rightarrow 31$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.094$ $wR(F^2) = 0.170$ S = 1.424102 reflections 310 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.958 (4)	Cu1-O3	2.268 (4)
Cu1-N2 ⁱ	1.992 (5)	O1-C13	1.307 (7)
Cu1-O6	1.993 (4)	O2-C13	1.212 (7)
Cu1-N1	2.018 (5)		
O1-Cu1-N2 ⁱ	92.38 (19)	O6-Cu1-N1	90.7 (2)
O1-Cu1-O6	179.47 (19)	O1-Cu1-O3	94.43 (18)
$N2^{i}-Cu1-O6$	87.82 (19)	N2 ⁱ -Cu1-O3	104.44 (17)
O1-Cu1-N1	88.98 (19)	O6-Cu1-O3	86.00 (17)
N2 ⁱ -Cu1-N1	163.59 (19)	N1-Cu1-O3	91.76 (18)

Symmetry code: (i) $x, -y + \frac{1}{2}, z - \frac{1}{2}$.

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O6-H6B\cdotsO1^{ii}$	0.82	1.88	2.693 (6)	170
$O6-H6A\cdots O7^{iii}$	0.82	1.84	2.642 (7)	164

Symmetry codes: (ii) $-x + \frac{3}{2}$, $y - \frac{1}{2}$, z; (iii) x, y - 1, z + 1.

Figure 1





Figure 2

An illustration of the zigzag chain formed in the structure of (I).



Figure 3

The two-dimensional network structure formed by $O-H\cdots O$ hydrogen bonds (dashed lines).

The H atoms of the water molecule were placed in calculated positions and refined using a riding model, with O–H distances of 0.82 Å and $U_{iso}(H) = 1.5U_{eq}(O)$. All H atoms of the dimethyl-formamide molecule were placed in calculated positions and refined using a riding-model approximation, with C–H distances of 0.93 Å for O—C–H and 0.96 Å for the methyl H atoms, and $U_{iso}(H) = 1.2$ (for CH) or 1.5 (for CH₃) times U_{eq} (parent atom). The remaining H atoms were positioned geometrically, with C–H = 0.93 Å, and allowed to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Bruker, 2002); software used to prepare material for publication: *SHELXL97*.

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