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#### Key indicators

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.007 Å R factor = 0.054 wR factor = 0.150 Data-to-parameter ratio = 15.0

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# A two-dimensional hydrogen-bonding supramolecular architecture of bis(2,2'-bipyridine)[(p-phenylenedioxy)diacetato]copper(II) tetrahydrate

In the crystal structure of the title complex,  $[Cu(bda)-(bipy)_2]\cdot 4H_2O$   $[H_2bda = (p-phenylenedioxy)diacetic acid or benzene-1,4-dioxyacetic acid <math>(C_{10}H_{10}O_6)$  and bipy = 2,2'-bipyridine  $(C_{10}H_8N_2)]$ , the Cu<sup>II</sup> centre is pentacoordinated, with a square-pyramidal geometry comprising four N atoms from two chelating bipy ligands and one carboxylate O atom of the bda ligand. The other three uncoordinated carboxylate O atoms of the bda ligand act as the acceptors of four O-H···O hydrogen bonds from the water molecules, among which another four O-H···O hydrogen-bonding interactions are also observed, resulting in a two-dimensional supramolecular pattern.

## Comment

Self-assembly processes directed by either hydrogen-bonding interactions or metal coordination have been extensively utilized in crystal engineering to construct supramolecular systems with novel structures and properties due to their inherent strength and reliability (Braga et al., 2003; Zhang et al., 2004; Chen & Liu, 2002). Among synthetic strategies in the context of coordination supramolecules, one of the most fruitful approaches is to take advantage of suitable bridging ligands, such as aromatic carboxylic acids, which possess diverse coordination modes and favour the formation of hydrogen bonds (Yaghi et al., 2003). Phenylenedioxydiacetic acids, especially (p-phenylenedioxy)diacetic acid (or benzene-1,4-dioxyacetic acid), H<sub>2</sub>bda, could act as a family of flexible multidentate ligands with versatile binding abilities, the coordination chemistry of which has been documented very little to date. Curiously, according to results in recent reports on coordination supramolecular systems based on mixed ligands, H<sub>2</sub>bda usually plays an uncoordinated anionic role in these complexes (Gao et al., 2004) or is not involved in the final product (Liu et al., 2004). In the present paper, we report the crystal structure of the title compound, (I), a new Cu<sup>II</sup> complex based on the mixed ligands bipy and H<sub>2</sub>bda, in which very interesting hydrogen-bonding interactions are observed.



The crystal structure of (I) consists of a neutral mononuclear  $[Cu(bipy)_2(bda)]$  molecule and four uncoordinated water molecules. Selected bond lengths and angles are listed in Received 21 June 2004 Accepted 13 July 2004 Online 17 July 2004



Figure 1

The molecular strucure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms and the water molecules have been omitted for clarity.



#### Figure 2

A view of the two-dimensional supramolecular structure of (I), showing the hydrogen-bonding interactions (dashed lines) and the A and B ring motifs. The bipy molecules have been omitted for clarity.

Table 1. As shown in Fig. 1, the Cu<sup>II</sup> centre is coordinated by four N atoms from two bipy ligands, with Cu–N bond lengths in the range 1.977 (3)–2.172 (4) Å, and by one additional O donor from the carboxylate group of the bda anion, with a Cu–O bond distance of 1.975 (3) Å. Thus, the coordination polyhedron around the Cu<sup>II</sup> cation could be best described as a distorted square-based pyramid, with a  $\tau$  parameter of 0.21 (Addison *et al.*, 1984).

The Cu<sup>II</sup> centre deviates from the least-squares basal plane of atoms N1/N2/O1/N4 by approximately 0.12 Å toward the apical N3 site. The bipy ligands combine with the Cu<sup>II</sup> centre to give two five-membered chelate rings (N1-C5-C6-N2-Cu1 and N3-C15-C16-N4-Cu1), forming an N2-Cu1-N1 angle of 80.76 (15)° and an N3-Cu1-N4 angle of 79.28 (14)°.

It should be noted that only one (O1-C21-O2) of the two carboxylate groups is involved in coordination to the Cu<sup>II</sup> cation, exhibiting the monodentate mode *via* atom O1; this is also be reflected by the fact that the C21-O1 bond [1.259 (5) Å] is significantly longer than C21-O2 [1.237 (5) Å]. The two C-O bond distances of the other carboxylate group (O5-C30-O6) are almost equivalent [1.241 (6) and 1.243 (5) Å], agreeing with its uncoordinated structural feature and delocalized state.

Analysis of the crystal packing of (I) shows the existence of eight  $O-H \cdots O$  interactions between the water molecules and the complex molecules, resulting in a two-dimensional supramolecular network, as depicted in Fig. 2. All relevant hydrogen-bonding geometries are listed in Table 2. Water molecules O7 and O8 form two O7-H7A···O6 and O8-H8A...O5 hydrogen bonds with the two O atoms of the uncoordinated carboxylate group. The same two water molecules are also involved in the formation of  $O7-H7B\cdots O2^{i}$ and  $O8 - H8B \cdot \cdot \cdot O2^{1}$  hydrogen bonds with the free atom O2 of the monodentate carboxylate subunit [symmetry code: (i) 1 - x, 1 - y, 2 - z]. The resulting motif B (Fig. 2), in the formalism of graph-set analysis of hydrogen-bond patterns (Etter, 1990), is characterized as  $N_4 = R_4^3(10)$ . Furthermore, four other types of intermolecular  $O-H \cdots O$  interactions are observed among the water molecules (Table 2). Thus, a hydrogen-bonded ring A,  $N_7 = R_7^5(14)$ , is generated, which comprises six water molecules (two O7, two O9, one O8 and one O10) and one free O2 atom of the monodentate carboxylate group. In summary, eight types of O-H···O interactions connect the water and complex subunits of (I) to form an infinite two-dimensional layered architecture (Fig. 2).

Examination of the structure of (I) with *PLATON* (Spek, 2003) showed that there were no solvent-accessible voids or  $\pi$ - $\pi$  stacking interactions in the crystal structure.

## Experimental

A methanol solution (5 ml) of 2,2'-bipyridine (33 mg, 0.2 mmol) was added to a solution of Cu(OAc)<sub>2</sub>·H<sub>2</sub>O (42 mg, 0.2 mmol) in CH<sub>3</sub>OH-H<sub>2</sub>O (10 ml, 1:1) with vigorous stirring. A methanol solution (10 ml) of H<sub>2</sub>bda (23 mg, 0.1 mmol) was then added and the mixture stirred for *ca* 30 min. The reaction solution was filtered and left to stand at room temperature. Blue block-like single crystals of (I) suitable for X-ray diffraction were obtained after several days by slow evaporation of the solvent (yield 45%). Spectroscopic analysis: IR (KBr pellet, v, cm<sup>-1</sup>): 3384 (*b*), 3108 (*m*), 3070 (*m*), 3030 (*m*), 2953 (*m*), 2894 (*m*), 1619 (*vs*), 1597 (*vs*), 1509 (*s*), 1475 (*s*), 1445 (*s*), 1423 (*s*), 1342 (*s*), 1320 (*m*), 1287 (*m*), 1233 (*s*), 1162 (*m*), 1121 (*m*), 1055 (*s*), 1016 (*m*), 938 (*w*), 909 (*w*), 830 (*m*), 809 (*m*), 772 (*s*), 731 (*m*), 690 (*m*), 658 (*w*), 622 (*w*), 535 (*w*), 491 (*w*). Analysis calculated for (I): C 53.60, H 4.80, N 8.33%; found: C 53.97, H 4.55, N 8.28%.

Crystal data

$[Cu(C_{10}H_8O_6)(C_{10}H_8N_2)_2] \cdot 4H_2O$	$D_x = 1.498 \text{ Mg m}^{-3}$
$M_r = 672.14$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 800
a = 14.460 (4)  Å	reflections
b = 10.112 (3)  Å	$\theta = 2.6-21.6^{\circ}$
c = 20.394(5)  Å	$\mu = 0.80 \text{ mm}^{-1}$
$\beta = 91.459 \ (5)^{\circ}$	T = 293 (2)  K
$V = 2981.1 (13) \text{ Å}^3$	Block, blue
Z = 4	$0.16 \times 0.14 \times 0.10 \text{ mm}$
Data collection	
Bruker SMART CCD area-detector	6108 independent reflections
diffractometer	3301 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.089$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.4^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -18 \rightarrow 16$
$T_{\min} = 0.796, T_{\max} = 0.923$	$k = -12 \rightarrow 9$
16 900 measured reflections	$l = -25 \rightarrow 25$

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.054$	$w = 1/[\sigma^2(F_o^2) + (0.0677P)^2]$
$wR(F^2) = 0.150$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.98	$(\Delta/\sigma)_{\rm max} < 0.001$
6108 reflections	$\Delta \rho_{\rm max} = 0.35 \text{ e} \text{ Å}^{-3}$
406 parameters	$\Delta \rho_{\rm min} = -0.43 \text{ e} \text{ Å}^{-3}$

#### Table 1

Selected geometric parameters (Å,  $^{\circ}$ ).

Cu1-O1	1.975 (3)	O1-C21	1.259 (5)
Cu1-N1	1.977 (3)	O2-C21	1.237 (5)
Cu1-N4	1.990 (3)	O5-C30	1.241 (6)
Cu1-N2	2.032 (4)	O6-C30	1.243 (5)
Cu1-N3	2.172 (4)		
O1-Cu1-N1	93.94 (14)	N4-Cu1-N2	95.58 (14)
O1-Cu1-N4	91.00 (13)	O1-Cu1-N3	98.89 (13)
N1-Cu1-N4	173.98 (15)	N1-Cu1-N3	96.54 (14)
O1-Cu1-N2	161.33 (14)	N4-Cu1-N3	79.28 (14)
N1-Cu1-N2	80.76 (15)	N2-Cu1-N3	99.49 (14)

 Table 2

 Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
O7−H7A···O6	0.85	1.83	2.661 (5)	165
$O7 - H7B \cdot \cdot \cdot O2^{i}$	0.85	2.07	2.908 (5)	170
O8−H8A···O5	0.85	1.87	2.681 (5)	160
$O8-H8B\cdots O2^{i}$	0.85	2.08	2.834 (5)	148
$O9-H9A\cdots O8^{ii}$	0.85	1.87	2.713 (6)	173
$O9-H9B\cdots O7^{iii}$	0.85	2.16	2.930 (6)	151
$O10-H10A\cdots O7^{iv}$	0.85	2.13	2.905 (6)	152
$O10-H10B\cdots O9^{ii}$	0.85	2.04	2.839 (7)	156

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

Although all H atoms were visible in difference maps, they were finally placed in geometrically calculated positions, with C-H

distances in the range 0.93–0.97 Å and O–H distances of 0.85 Å, and included in the final refinement in the riding model approximation, with  $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$  for aromatic and methylene H atoms, and  $1.5U_{eq}(\rm O)$  for water H atoms.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1998); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg & Berndt, 1999); software used to prepare material for publication: *SHELXL*97.

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### References

- Addison, A. W., Rao, T. N., Reedijk, J., van Rijn, J. & Verschoor, G. C. (1984). J. Chem. Soc. Dalton Trans. pp. 1349–1356.
- Braga, D., Maini, L., Polito, M., Tagliavini, E. & Grepioni, F. (2003). Coord. Chem. Rev. 246, 53–71.
- Brandenburg, K. & Berndt, M. (1999). *DIAMOND*. Version 2.1c. Crystal Impact GbR, Bonn, Germany.

Bruker (1998). SMART (Version 5.0), SAINT (Version 4.0) and SHELXTL. Bruker AXS Inc., Madison, Wisconsin, USA.

- Chen, X. M. & Liu, G. F. (2002). Chem. Eur. J. 8, 4811-4817.
- Etter, M. C. (1990). Acc. Chem. Res. 23, 120-126.
- Gao, S., Liu, J.-W., Huo, L.-H., Zhao, H. & Zhao, J.-G. (2004). Acta Cryst. E60, m620–m621, and references cited therein.

Liu, J.-W., Gao, S., Huo, L.-H. & Ng, S. W. (2004). Acta Cryst. E60, m501-m503.

Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.

Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.

- Yaghi, O. M., O'Keeffe, M., Ockwig, N. W., Chae, H. K., Eddaoudi, M. & Kim, J. (2003). *Nature (London)*, **423**, 705–714.
- Zhang, J. P., Zheng, S. L., Huang, X. C. & Chen, X. M. (2004). Angew. Chem. Int. Ed. 43, 206–209.