

Jun Tao,^a Rong-Bin Huang,^a
Lan-Sun Zheng^a and Seik Weng
Ng^{b*}^aDepartment of Chemistry, Xiamen University,
Xiamen 361005, People's Republic of China,
and ^bDepartment of Chemistry, University of
Malaya, 50603 Kuala Lumpur, Malaysia

Correspondence e-mail: seikweng@um.edu.my

Key indicators

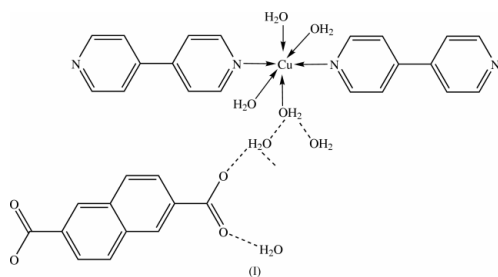
Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.042
 wR factor = 0.120
Data-to-parameter ratio = 15.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Tetraaquabis(4,4'-bipyridine)copper(II)
2,6-naphthalenedicarboxylate trihydrate

The Cu atom in $[\text{Cu}(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_4](\text{C}_{12}\text{H}_6\text{O}_4)\cdot 3\text{H}_2\text{O}$ exists in a *trans*- N_2O_4 octahedral coordination polyhedron that is formed by the N atoms of two 4,4'-bipyridine heterocycles and the O atoms of four water molecules. The carboxylate groups of the centrosymmetric anions interact indirectly with the Cu atoms through the coordinated water molecules. The cations and anions are linked by hydrogen bonds into a network motif. The Cu atom lies at a site of 2 symmetry.

Received 25 June 2003
Accepted 10 July 2003
Online 24 July 2003

Comment

The 4,4'-bipyridine spacer ligand has been used to react with a range of copper(II) salts; the products adopt a number of topologies such as squares, rectangles or rhombuses (Yuan *et al.*, 2003). At other times, the ligand links adjacent Cu atoms into a linear chain, and in one instance, the ligand merely interacts with the Cu atom through only one N site, the other being involved with hydrogen-bonding interactions (Abu-Shandi *et al.*, 2001). The structure of the 1/1 adduct with copper dibenzoate has not been authenticated, but a substituted derivative furnishes a heterocycle-coordinated compound in which the Cu atom is covalently linked to the carboxylate entity (He & Zhu, 2003). However, the extension of this study to copper terephthalate, a dicarboxylate compound, gave instead a tetraaquacopper complex in which the carboxylate anion interacts with the six-coordinate Cu atom through the coordinated water molecules (Long, 2003).



In the title compound, (I), the Cu atom, which lies on a twofold axis, is coordinated by four water molecules. The geometry is tetragonally distorted as the two $\text{Cu}-\text{O}_{\text{water}}$ distances [2.385 (3) and 2.457 (3) Å] are significantly longer than the pairs of $\text{Cu}-\text{O}$ [2.005 (2) Å] and $\text{Cu}-\text{N}$ [2.079 (2) Å] bonds. The coordinated water molecules interact with the uncoordinated water molecules, which, in turn, interact with the dicarboxylate dianion. One of them also interacts with the coordinated heterocycle. The cations and anions are linked by hydrogen bonds into a network motif (Table 2).

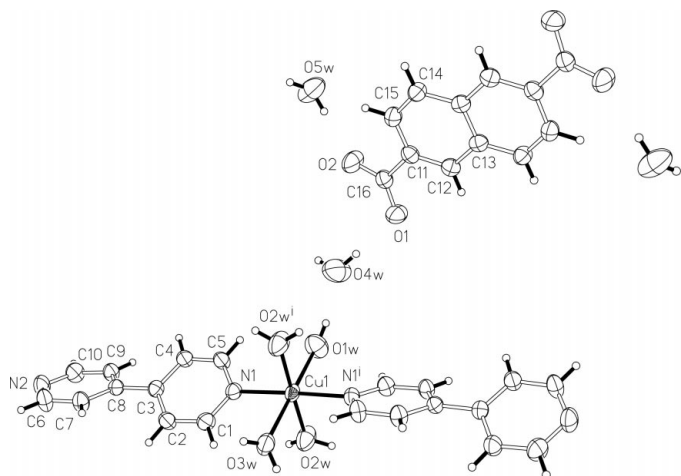


Figure 1
ORTEPII (Johnson, 1976) plot of the title compound, with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as circles of arbitrary radii. [Symmetry code: (i) $1 - x, y, \frac{1}{2} - z$.]

Experimental

Copper nitrate trihydrate (0.12 g, 0.5 mmol) and 2,6-naphthalenedicarboxylic acid (0.11 g, 0.5 mmol) were placed in water (5 ml) and ammonium hydroxide was added until the reagents dissolved completely. 4,4'-Bipyridine (0.08 g, 0.5 mmol) was added, and water was added to make up a total volume of 10 ml. The mixture was heated in a Teflon-lined stainless-steel bomb at 413 K for 16 h. The title compound deposited from the solution as well formed crystals when the bomb was cooled to room temperature.

Crystal data

$[\text{Cu}(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_4](\text{C}_{12}\text{H}_6\text{O}_4) \cdot 3\text{H}_2\text{O}$	$D_x = 1.49 \text{ Mg m}^{-3}$
$M_r = 716.19$	Mo $K\alpha$ radiation
Orthorhombic, $Pbcn$	Cell parameters from 2701 reflections
$a = 7.4063 (5) \text{ \AA}$	$\theta = 2.8\text{--}24.7^\circ$
$b = 18.157 (1) \text{ \AA}$	$\mu = 0.75 \text{ mm}^{-1}$
$c = 23.801 (2) \text{ \AA}$	$T = 298 (2) \text{ K}$
$V = 3200.7 (4) \text{ \AA}^3$	Plate, blue
$Z = 4$	$0.36 \times 0.10 \times 0.06 \text{ mm}$

Data collection

Bruker SMART area-detector diffractometer	3768 independent reflections
φ and ω scans	2292 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.045$
$T_{\text{min}} = 0.677, T_{\text{max}} = 0.914$	$\theta_{\text{max}} = 28.3^\circ$
17 912 measured reflections	$h = -9 \rightarrow 8$
	$k = -24 \rightarrow 17$
	$l = -30 \rightarrow 31$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.042$	$w = 1/[\sigma^2(F_o^2) + (0.0734P)^2]$
$wR(F^2) = 0.120$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.91$	$(\Delta/\sigma)_{\text{max}} = 0.001$
3768 reflections	$\Delta\rho_{\text{max}} = 0.67 \text{ e \AA}^{-3}$
240 parameters	$\Delta\rho_{\text{min}} = -0.52 \text{ e \AA}^{-3}$

Table 1

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

Cu1—O1w	2.385 (3)	Cu1—O3w	2.457 (3)
Cu1—O2w	2.005 (2)	Cu1—N1	2.079 (2)
O1w—Cu1—O3w	180	O2w—Cu1—N1	89.0 (1)
O1w—Cu1—O2w	88.8 (1)	O2w—Cu1—N1 ⁱ	91.1 (1)
O1w—Cu1—N1	91.6 (1)	O3w—Cu1—N1	88.4 (1)
O2w—Cu1—O2w ⁱ	177.6 (1)	N1—Cu1—N1 ⁱ	176.9 (1)
O2w—Cu1—O3w	91.2 (1)		

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

Table 2

Hydrogen-bonding geometry ($\text{\AA}, ^\circ$).

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
O1w—H1w1 \cdots O4w ⁱ	0.84	1.92	2.738 (3)	165
O2w—H2w1 \cdots N2 ⁱⁱ	0.87	2.34	3.202 (3)	169
O2w—H2w2 \cdots N2 ⁱⁱⁱ	0.86	2.40	3.257 (3)	173
O3w—H3w1 \cdots O2 ^{iv}	0.85	1.96	2.799 (2)	174
O4w—H4w1 \cdots O1	0.85	1.95	2.792 (3)	168
O4w—H4w2 \cdots O2 ^v	0.85	2.17	2.955 (3)	153
O5w—H5w1 \cdots O2	0.85	1.94	2.790 (2)	177

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

The water H-atoms were located and refined subject to O—H = 0.85 (1) \AA and H \cdots H = 1.39 (1) \AA . The C-bound H atoms were positioned geometrically (C—H = 0.93 \AA) and refined in the riding-model approximation; the displacement parameters of all H atoms were set to 1.2 times U_{eq} of the equivalent isotropic displacement parameters of the parent atoms.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

The authors thank the National Science Foundation of China (grant No. 20023001), the Innovation Foundation for Young Scientific Talent of Fujian Province, China (grant No. 2002 J004), Xiamen University (grant No. Y07015) and the University of Malaya for supporting this work.

References

- Abu-Shandi, K., Janiak, C. & Kersting, B. (2001). *Acta Cryst.* **C57**, 1261–1264.
- Bruker (2001). SAINT and SMART. Bruker AXS Inc., Madison, Wisconsin, USA.
- He, H.-Y. & Zhu, L.-G. (2003). *Acta Cryst.* **E59**, o174–o176.
- Johnson, C. K. (1976). ORTEPII. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
- Long, L. S. (2003). Unpublished results.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.
- Yuan, J. X., Hu, M. L. & Song, X. Y. (2003). *Z. Kristallogr. New Cryst. Struct.* **218**, 1–2.