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Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.008 Å R factor = 0.052 wR factor = 0.123 Data-to-parameter ratio = 16.7

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

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In the crystal structure of the title complex, $[Cu(C_{10}H_9NO_3)-(H_2O)]\cdot H_2O$, the Cu atom is tetracoordinated by O and N atoms, and facilitates an intermolecular Cu···O short contact of 2.809 (3) Å. The two symmetrically-related coordinated Cu ions form distorted square pyramids of coordination CuO₄N. In the asymmetric unit, a free water molecule is hydrogen bonded to the coordinated water molecule. In the packing, the molecules are interconnected by two types of $O-H\cdots O$ hydrogen bonds.

acetato]copper(II) monohydrate

Aqua[{[2-(2-hydroxyphenyl)ethylidene]amino}-

Comment

Considerable attention has been paid to the structures of Cu complexes with Schiff bases formed from salicylaldehydeamino acid (Ueki *et al.*, 1967, 1969; Bkouche-Waksman *et al.*, 1988). These complexes are expected to be catalytic intermediates in non-enzymatic transamination reactions (Eichhorn & Marchand, 1956; Longenecker & Snell, 1957). Recently, X-ray structural studies on tin complexes with such ligands exhibited a variety of coordination and bonding for the tin^{TV} ion (Dakternieks *et al.*, 1998; Basu Baul *et al.*, 1999, 2001, 2002). The usefulness of such systems has, in turn, prompted research into the chemistry of Cu, since the bond distances involving the N atoms are of interest. In this study, we report the X-ray structural analysis of the title complex, (I).



The asymmetric unit comprises a ligand-coordinated Cu^{II} cation, a coordinated water molecule, and a free water molecule. The Cu atom is tetracoordinated by two O atoms and one N atom from the *O*,*N*,*O*-tridentate ligand and by one O atom of a water molecule. The Cu–O and Cu–N bond distances are in the range 1.881 (3)–1.959 (3) Å, in agreement with reported values [1.892 (2)–1.993 (2) Å; Warda, 1997*a*,*b*; Warda *et al.*, 1997; Warda, 1998*a*,*b*)]. Atom Cu1 also facilitates an intermolecular Cu···O1ⁱ short contact (Table 1), giving a distorted square pyramidal CuO₄N coordination with O1ⁱ occupying the apical position. The basal plane is formed by Cu1/O1/N1/O3/O1*W*. The bond angles around Cu1 are listed in Table 1.

Received 2 June 2003 Accepted 3 June 2003 Online 17 June 2003

2718 independent reflections

 $R_{\rm int} = 0.080$

 $\theta_{\rm max} = 28.3^{\circ}$

 $h = -26 \rightarrow 20$

 $k = -12 \rightarrow 14$ $l = -14 \rightarrow 18$

refinement

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.97 \ {\rm e} \ {\rm \AA}^{-3}$ $\Delta \rho_{\rm min} = -0.73 \ {\rm e} \ {\rm \AA}^{-3}$

1739 reflections with $I > 2\sigma(I)$

H atoms treated by a mixture of

 $w = 1/[\sigma^2(F_o^2) + (0.0475P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

independent and constrained



Figure 1

The asymmetric unit of the title compound (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.



Figure 2

Packing diagram of (I), showing the links via O-H···O hydrogen bonds.

The ligand and the atom Cu1 form two nearly planar rings, one five-membered [ring A: Cu1-O3-C9-C8-N1; Q₂ puckering amplitude (Cremer & Pople, 1975) 0.090 (5) Å] and the other six-membered [ring B: Cu1-O1-C1-C6-C7-N1; Q_T puckering amplitude 0.184 (4) Å]. These two rings adopt envelope conformations, with atoms C8 and Cu1 deviating by 0.134 (1) Å and 0.222 (1) Å from the N1/Cu1/O3/ C9 and O1/C1/C6/C7/N1 planes, respectively. The sixmembered, five-membered and aromatic (C) rings are nearly coplanar, with ring B making dihedral angles of 7.1 (2) $^{\circ}$ and 4.8 (2)° with A and C, respectively.

In the asymmetric unit, the Cu^{II} complex cation and the free water molecule are interconnected by $O1W-H1W1\cdots O2W$ hydrogen bonds. In the crystal packing, the two water molecules also play an important role in the intermolecular hydrogen-bonding scheme, in which there are two types of intermolecular O-H···O bonds, viz.O1W-H2W1···O3¹ and $O2W - H2W2 \cdots O2^{ii}$ (Table 2 and Fig. 2). Two molecules are interconnected by the two intermolecular hydrogen bonds into molecular chains.

Experimental

A methanolic solution of potassium ((2-(2-hydroxyphenyl)ethylidene)amino)acetate (Dakternieks et al., 1998) was added dropwise to a stirred aqueous solution of Cu(OAc)₂·H₂O containing a few

Crystal data

$[Cu(C_{10}H_{11}NO_4)]H_2O$	$D_x = 1.759 \text{ Mg m}^{-3}$
$M_r = 290.75$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 3297
a = 20.2250 (3) Å	reflections
b = 10.6741 (3) Å	$\theta = 2.8 - 28.3^{\circ}$
c = 14.2089 (3) Å	$\mu = 2.00 \text{ mm}^{-1}$
$\beta = 134.294 \ (1)^{\circ}$	T = 293 (2) K
V = 2195.59 (9) Å ³	Block, green
Z = 8	$0.20 \times 0.12 \times 0.10 \text{ mm}$

Data collection

Siemens SMART CCD areadetector diffractometer ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\min} = 0.691, T_{\max} = 0.825$ 7657 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.052$ $wR(F^2) = 0.123$ S = 0.952718 reflections 163 parameters

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.881 (3)	Cu1-O1W	1.959 (3)
Cu1-N1	1.936 (3)	Cu1-O1 ⁱ	2.809 (3)
Cu1-O3	1.948 (3)		
O1-Cu1-N1	93.9 (1)	O3-Cu1-O1W	88.6 (1)
O1-Cu1-O3	179.0 (1)	O1-Cu1-O1 ⁱ	78.3 (1)
N1-Cu1-O3	86.0(1)	N1-Cu1-O1 ⁱ	101.4 (2)
O1-Cu1-O1W	91.7 (1)	O3-Cu1-O1 ⁱ	100.8 (2)
N1-Cu1-O1W	170.9 (1)	$O1W-Cu1-O1^{i}$	86.7 (1)

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

Table 2		
Hydrogen-bonding	geometry (Å, °).

$D-\mathrm{H}\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1W - H1W1 \cdots O2W$	1.02 (9)	1.65 (9)	2.649 (9)	164 (7)
$O1W - H2W1 \cdots O3^{n}$	0.83 (7)	2.02(7)	2.850 (8)	172 (6)
$O2W - H2W2 \cdot \cdot \cdot O2^{iii}$	1.04	1.82	2.838 (5)	166
			1	

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

All H atoms attached to C were positioned geometrically, with C-H 0.93–0.97 Å. Though the H atoms attached to O1W were calculated and refined isotropically, with OW-H 0.84 (6)-1.02 (7) Å, those attached to O2W were located in a difference map, with OW-H 1.04 Å and treated as riding atoms.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine

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structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

The authors thank the Malaysian Government and Universiti Sains Malaysia for research grant R&D No. 304/ PFIZIK/670011, and AU thanks Universiti Sains Malaysia for a Visiting Post-Doctoral Fellowship. The financial support of the Department of Science & Technology, New Delhi, India (grant No. SP/S1/F26/99, TSBB) is gratefully acknowledged.

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