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Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$
 R factor = 0.036
 wR factor = 0.077
Data-to-parameter ratio = 11.0For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis[*N*-(2-hydroxybenzoyl)-*N'*-[1-(2-pyridyl)-
ethylidene]hydrazinato]copper(II)

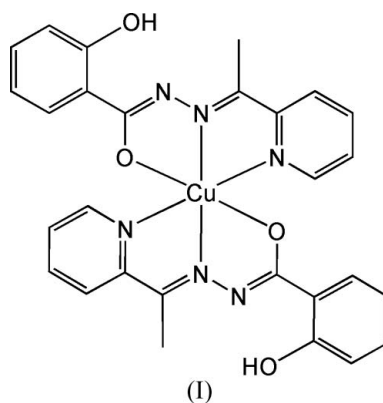
The title complex, $[\text{Cu}(\text{C}_{14}\text{H}_{12}\text{N}_3\text{O}_2)_2]$, consists of two deprotonated 2-acetylpyridine salicylhydrazone ligands coordinated to the copper as *N,N,O*-tridentate ligands, forming a neutral complex. The Cu atom lies on a twofold rotation axis. The structure is stabilized by intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bonding, and $\pi-\pi$ stacking interactions between the pyridyl and benzene rings lead to a two-dimensional network in the crystal structure.

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Comment

Diazine Schiff base ligands with two coordination compartments linked by a single N–N bridge have received considerable attention due to the diversity of their structural and functional properties (Grove *et al.*, 2004; Bai *et al.*, 2006). This type of polydentate ligand usually generates a variety of different structural forms (Bai *et al.*, 2005; Song *et al.*, 2004). Transition-metal-directed self-assembly has also become a powerful methodology for the construction of different coordination complexes with different geometries and spectroscopic characteristics and can give rise to extended solids with a variety of structures and physical properties (Leininger *et al.*, 2000).



The title copper(II) complex, (I), was obtained by the reaction of $\text{CuCl}_2\cdot 2\text{H}_2\text{O}$ with the pentadentate Schiff base ligand 2-acetylpyridine salicylhydrazone (H_2L). The X-ray structure determination shows that the molecule consists of two HL^- ligands coordinated to the copper as *N,N,O*-tridentate ligands, forming a neutral complex (Fig. 1). A feature of the structure is that the copper atom lies on a twofold rotation axis. The ligands bond to the copper in the *mer* configuration (with pairs of carbonyl O atoms and pyridyl N atoms each bearing a *cis* relationship, whereas the imino N atoms are *trans* to each other). The intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond

(Table 2) is an important factor in the stabilization of the molecule. In the crystal structure, molecules interact with each other, forming a two-dimensional network through π - π stacking interactions between the benzene and pyridine rings.

Experimental

2-Acetylpyridine salicylhydrazone (H_2L) was obtained in high yield by the direct combination of 1:1 molar equivalents of salicylaldehyde and 2-acetylpyridine in methanol. A solution of the H_2L ligand (0.025 g, 0.1 mmol) and Et_3N (0.011 g, 0.1 mmol) in methanol (6 ml) was layered on to a solution of DMF (6 ml) containing $CuCl_2 \cdot 2H_2O$ (0.017 g, 0.1 mmol) at the bottom. The solutions were left for few weeks at room temperature in darkness to give X-ray quality crystals (84% yield).

Crystal data

$[Cu(C_{14}H_{12}N_3O_2)_2]$	$Z = 4$
$M_r = 572.07$	$D_x = 1.516 \text{ Mg m}^{-3}$
Orthorhombic, $Aba2$	Mo $K\alpha$ radiation
$a = 12.570 (2) \text{ \AA}$	$\mu = 0.92 \text{ mm}^{-1}$
$b = 18.731 (4) \text{ \AA}$	$T = 293 (2) \text{ K}$
$c = 10.6482 (19) \text{ \AA}$	Block, green
$V = 2507.1 (8) \text{ \AA}^3$	$0.30 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer	1968 independent reflections
φ and ω scans	1628 reflections with $I > 2\sigma(I)$
Absorption correction: none	$R_{int} = 0.041$
6226 measured reflections	$\theta_{max} = 25.5^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.035P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.036$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.077$	$(\Delta/\sigma)_{max} < 0.001$
$S = 1.02$	$\Delta\rho_{max} = 0.52 \text{ e \AA}^{-3}$
1968 reflections	$\Delta\rho_{min} = -0.16 \text{ e \AA}^{-3}$
179 parameters	Absolute structure: Flack (1983),
H-atom parameters constrained	738 Friedel pairs
	Flack parameter: $-0.018 (19)$

Table 1

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

Cu1—N2	1.966 (3)	Cu1—O1	2.158 (3)
Cu1—N1	2.156 (3)		
N2—Cu1—N2 ⁱ	172.1 (2)	N1—Cu1—O1 ⁱ	93.46 (10)
N2—Cu1—N1 ⁱ	108.75 (13)	N2—Cu1—O1	76.59 (12)
N2—Cu1—N1	77.11 (13)	N1—Cu1—O1	152.95 (10)
N1 ⁱ —Cu1—N1	89.07 (16)	O1 ⁱ —Cu1—O1	96.32 (15)
N2—Cu1—O1 ⁱ	98.04 (11)		

Symmetry code: (i) $-x, -y, z$.

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O2-H2 \cdots N3$	0.82	1.82	2.550 (4)	148

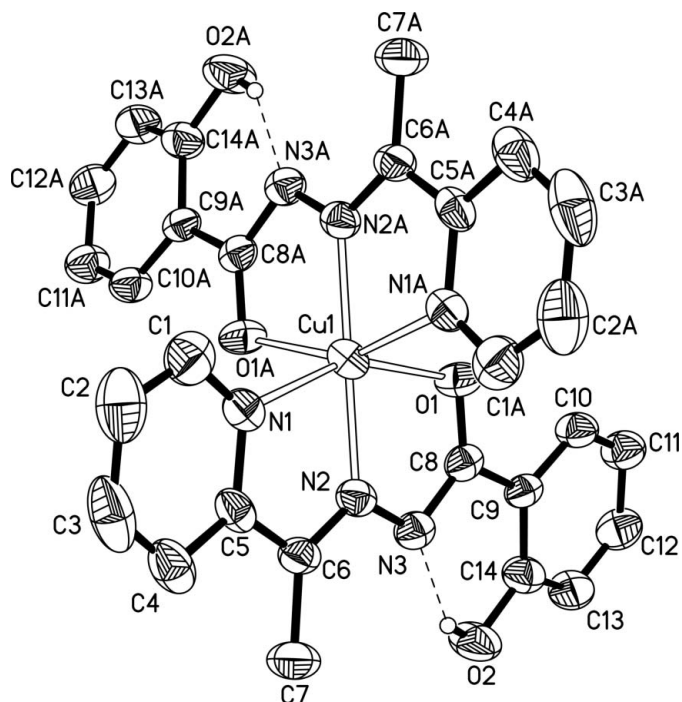


Figure 1

The molecular structure of the title complex, with displacement ellipsoids drawn at the 30% probability level. H atoms other than those of hydroxyl have been omitted for clarity. Hydrogen bonds are indicated by dashed lines. [Symmetry code: (A) $-x, -y, z$.]

The H atoms bonded to C and O atoms were constrained to ride on their parent atoms, with distances of 0.97 and 0.89 \AA , respectively, and $U_{iso}(H) = 1.2U_{eq}(C)$ and $1.5U_{eq}(O)$.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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