Received 18 February 2006 Accepted 23 February 2006

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.005 Å R factor = 0.045 wR factor = 0.117 Data-to-parameter ratio = 12.4

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

[2-Oxido-1-naphthaldehyde (2-hydroxybenzoyl)hydrazonato]pyridinecopper(II)

The approximately planar complex molecule, $[Cu(C_{18}H_{12}N_2O_3)(C_5H_5N)]$, contains one L^{2-} ligand $(L^{2-}$ is the dianion of 2-hydroxy-1-naphthaldehyde 2-hydroxybenzoylhydrazone), one Cu atom and one pyridine molecule. The Cu centre shows square-planar N_2O_2Cu coordination. The tridentate dianion has an intramolecular $N \cdots H - O$ hydrogen bond. Each pair of adjacent molecules is linked together by $\pi - \pi$ stacking and Cu $\cdots N$ interactions, which lead to the existence of a dimer. Owing to C $-H \cdots O$ hydrogen bonds, these dimers are further assembled into a two-dimensional framework.

Comment

Interest in the chemistry of aroylhydrazones arises from their ability to bind to metal ions (Singh et al., 1982; Salem, 1998) as well as their biological activity (Singh et al., 1982; Carcelli et al., 1995). In fact, with some hydrazones, their metal complexes are endowed with significantly improved biological and pharmaceutical activity (Ainscough et al., 1998; Anford et al., 1998). In particular, the copper complex of salicylaldehyde benzoylhydrazone possesses exceptional antibacterial activity (Ainscough et al., 1998). The coordination chemisty of copper hydrazonates, such as $[Cu(C_{15}H_{12}N_2O_3)(C_5H_5N)]$ (Chen et al., 2004), $[Cu(H_{2}sb)(CCl_{3}CO_{2})_{2}]_{2}$ (Ainscough *et al.*, 1998), and [Cu₄(POAPZ-H)₄(H₂O)₂](NO₃)₄·4H₂O and [Cu₄(POAP- H_{4} (ClO₄)₄ (POAPZ-H is *N*-imidopyrazine picolinic hydrazide and POAP-H is N-imidopicoline picolinic hydrazide; Matthews et al., 1999), have been reported. The present study describes the title compound, (I) (Fig. 1).



The complex, (I), contains of one L^{2-} ligand, one Cu atom and one pyridine molecule. The geometry around the Cu centre can be described as a square plane composed of the N₂O₂ set of atoms; the two O atoms and one N atom are supplied by the tridentate L^{2-} , while the other N atom comes

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Figure 1

The molecular structure of the title complex. Displacement ellipsoids are drawn at the 30% probability level and H atoms have been omitted for clarity.



Figure 2

The dimer structure formed by Cu···N and π - π interactions (dashed lines). [Symmetry code: (A) 1 - x, 1 - y, 2 - z]. H atoms have been omitted.



Figure 3

The crystal packing of the title complex. Dashed lines indicate hydrogen bonds.

from the pyridine molecule. The free hydroxy group forms an intramolecular hydrogen bond. The two chelate rings are nearly coplanar.

Inversion-related pairs of molecules are linked together by weak $Cu \cdots N$ interactions, with $Cu 1 \cdots N1^{i}$ and $Cu 1 \cdots N2^{i}$

separations of more than 3.2 Å [symmetry code: (i) 1 - x, 1 - y, -z]. As with [Ni(C₁₈H₁₂N₂O₃)(C₅H₅N)] (Liu *et al.*, 2005), there are $\pi - \pi$ interactions between each pair of naphthalene and phenyl ring planes, giving rise to a weakly held dimer (Fig. 2).

The packing of (I) (Fig. 3) shows that atom C20 interacts with atom $O3^{i}$ at a distance of 3.347 (5) Å [symmetry code: (i) $x, \frac{1}{2} - y, z + \frac{1}{2}$, and this weak interaction leads to the formation of a layer motif ing the *bc* plane.

Experimental

The ligand L (0.25 mmol, 0.077 g), KOH (0.5 mmol, 0.028 g), $Cu(ClO_4)_2$ (0.25 mmol, 0.086 g), dimethylformamide (8 ml) and methanol (8 ml) were mixed. After stirring the solution for about 4 h. the green precipitate which formed was dissolved in pyridine (10 ml). A week later, dark-green crystals of (I) were isolated from the solution [m.p. >573 K (decomposition)]. Spectroscopic analysis: FT-IR (KBr pellets, v, cm⁻¹): 3443 (s), 3047 (w), 1615 (vs), 1602 (s), 1527 (s), 1452 (s), 1484 (s), 1391 (s), 1334 (s), 1256 (s), 1302 (m), 1239 (w), 1138 (m), 1017 (s), 971 (m), 855 (w), 757 (vs), 696 (s), 676 (m), 526 (m), 452 (m).

Crystal data

$[Cu(C_{18}H_{12}N_2O_3)(C_5H_5N)]$	$D_{\rm x} = 1.575 {\rm Mg} {\rm m}^{-3}$
$M_r = 446.94$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2185
a = 11.745 (2) Å	reflections
b = 8.1666 (17) Å	$\theta = 2.4-22.8^{\circ}$
c = 20.380 (4) Å	$\mu = 1.19 \text{ mm}^{-1}$
$\beta = 105.397 \ (3)^{\circ}$	T = 298 (2) K
V = 1884.5 (7) Å ³	Block, dark green
Z = 4	$0.28 \times 0.21 \times 0.12 \text{ mm}$

Data collection

Siemens SMART CCD area-3369 independent reflections detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

 $T_{\min} = 0.732, T_{\max} = 0.870$ 9587 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.117$ S = 1.023369 reflections 271 parameters

2557 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.038$ $\theta_{\rm max} = 25.2^{\circ}$ $h = -10 \rightarrow 14$ $k = -9 \rightarrow 9$ $l = -23 \rightarrow 24$

H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0671P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.42 \text{ e} \text{ \AA}^{-3}$ $\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.880(2) 1 904(3)	N1 - N2 N2 - C12	1.395(3) 1 319(4)
Cu1-O2	1.904 (3)	01-C3	1.302 (4)
Cu1-N3 N1-C1	1.997(3) 1.284(4)	O2-C12 O3-C14	1.288(3) 1.353(4)
O1-Cu1-N1	92.75 (10)	O1-Cu1-N3	93.94 (10)
O1 - Cu1 - O2	173.84 (9)	N1-Cu1-N3	171.13 (11)
n = Cu = 02	81.00 (10)	02-Cu1-N3	91.89 (10)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
C20-H20···O3 ⁱⁱ	0.93	2.81	3.347 (5)	118
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Symmetry code: (ii) $x, -y + \frac{1}{2}, z + \frac{1}{2}$.

All H atoms were positioned geometrically and refined as riding on their parent atoms; C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$, and O-H = 0.82 Å and $U_{iso}(H) = 1.5U_{eq}(O)$.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

The authors acknowledge the support of the Natural Science Foundation of Shandong Province, China (grant No. Y2003B01).

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