metal-organic papers

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) = 0.003 Å R factor = 0.032 wR factor = 0.091 Data-to-parameter ratio = 17.4

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

Acetato{2-[2-(diethylamino)ethyliminomethyl]-4-nitrophenolato}copper(II)

In the title mononuclear copper(II) complex, $[Cu(C_{13}H_{18}-N_3O_3)(C_2H_3O_2)]$, the Cu^{II} ion is four-coordinated by one Schiff base ligand and one acetate anion in a slightly distorted square-planar geometry.

Received 22 March 2007 Accepted 22 March 2007

Comment

Recently, we have reported the crystal structures of a few Schiff base-metal complexes (You *et al.*, 2006*a,b*; You, Han *et al.*, 2006). As an extension of this work, the crystal structure of the title mononuclear copper(II) complex, (I) (Fig. 1), is reported here.



The Cu^{II} ion in (I) is four-coordinated by the *NNO* donor set of the Schiff base ligand, and by one O atom of the acetate anion in a slightly distorted square-planar geometry. The Cu– O and Cu–N bond lengths (Table 1) are comparable to those in other Schiff base–copper(II) complexes (You, 2006; You & Zhu, 2006; You *et al.*, 2006*a,b*; You, Han *et al.*, 2006; You, Jiao *et al.*, 2006). The two *trans* angles at the metal centre are 171.54 (6) and 166.21 (7)°; all other angles around Cu1 are close to 90°, ranging from 84.63 (7) to 93.55 (6)°, indicating a slightly distorted square-planar geometry for Cu1.

In the crystal structure, molecules are linked through intermolecular C-H···O hydrogen bonds (Table 2), forming a ribbon along the *b* axis.

Experimental

All reagents were of commercially available grade and were used without further purification. 5-Nitro-2-hydroxybenzaldehyde (0.1 mmol, 16.5 mg) and *N*,*N*-diethylethane-1,2-diamine (0.1 mmol, 11.6 mg) were dissolved in MeOH (10 ml). The mixture was stirred at room temperature for 10 min, giving a clear orange solution. To this solution was added an aqueous solution (1 ml) of Cu(CH₃COO)·H₂O (0.1 mmol, 20.0 mg) with stirring. The mixture was stirred for a furher 10 min at room temperature. After leaving the filtrate to stand in air for 6 d, blue block-shaped crystals of (I) were formed.

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Crystal data

 $\begin{bmatrix} Cu(C_{13}H_{18}N_3O_3)(C_2H_3O_2) \end{bmatrix} \\ M_r = 386.89 \\ Monoclinic, P2_1/c \\ a = 18.458 (2) Å \\ b = 6.5665 (7) Å \\ c = 13.8915 (15) Å \\ \beta = 91.111 (2)^{\circ} \\ \end{bmatrix}$

Data collection

Bruker SMART 1000 CCD areadetector diffractometer Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) *T*_{min} = 0.668, *T*_{max} = 0.806

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$	220 parameters
$wR(F^2) = 0.091$	H-atom parameters constrained
S = 1.05	$\Delta \rho_{\rm max} = 0.31 \text{ e } \text{\AA}^{-3}$
3837 reflections	$\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$

V = 1683.4 (3) Å³

Mo $K\alpha$ radiation

 $0.33 \times 0.22 \times 0.17 \text{ mm}$

13908 measured reflections

3837 independent reflections

3241 reflections with $I > 2\sigma(I)$

 $\mu = 1.33 \text{ mm}^{-1}$

T = 298 (2) K

 $R_{\rm int} = 0.024$

Z = 4

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.9296 (15)	Cu1-O4	1.9539 (13)
Cu1-N1	1.9320 (16)	Cu1-N2	2.0675 (16)
O1-Cu1-N1	93.09 (6)	O1-Cu1-N2	166.21 (7)
O1-Cu1-O4	90.60 (6)	N1-Cu1-N2	84.63 (7)
N1-Cu1-O4	171.54 (6)	O4-Cu1-N2	93.55 (6)

Table 2

		~	
Hydrogen-bond	1 geometry	(À.	°)
,	- <u>-</u> ,	7	

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\overline{C7-H7\cdots O3^{i}}$	0.93	2.46	3.284 (3)	148
$C8-H8A\cdots O3^{i}$	0.97	2.54	3.439 (3)	153
$C8-H8B\cdots O4^{ii}$	0.97	2.43	3.338 (3)	157

Symmetry codes: (i) -x, -y + 2, -z; (ii) x, y + 1, z.

H atoms were placed in idealized positions and constrained to ride on their parent atoms, with C-H distances in the range 0.93–0.97 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$ and $1.5U_{eq}(methyl C)$.



Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme. H atoms have been omitted for clarity.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); 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*.

This work was supported by the Scientific Research Foundation of the Education Office of Liaoning Province (Project No. 2005226).

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