# 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 = 293 K Mean  $\sigma$ (C–C) = 0.007 Å R factor = 0.057 wR factor = 0.138 Data-to-parameter ratio = 17.5

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

# Dichlorobis[2-(o-tolyliminomethyl)phenolato]copper(II)

The title compound,  $[Cu(C_{14}H_{13}NO)_2Cl_2]$ , is a mononuclear copper(II) complex of a Schiff base. The Cu<sup>II</sup> atom is coordinated by two O atoms from two 2-(*o*-tolylimino-methyl)phenolate ligands, and by two chloride anions, forming a tetrahedral geometry.

Received 14 July 2005 Accepted 21 July 2005 Online 23 July 2005

## Comment

Copper(II) complexes are of great interest in coordination chemistry (Wagner & Walker, 1983; Margeat *et al.*, 2004). As an extension of work on the structural characterization of copper compounds, the title mononuclear copper(II) complex, (I) (Fig. 1), is reported.



The Cu atom is in a tetrahedral geometry (Table 1) and is coordinated by two O atoms from two 2-(o-tolyliminomethyl)phenolate ligands, and by two chloride anions. The Cu—O and Cu—Cl bond lengths are comparable to the values observed in other copper(II) complexes (MacLachlan *et al.*, 1996; Countryman *et al.*, 1974). There are no short intermolecular contacts (Fig. 2). Intramolecular hydrogen bonds are found in the ligands (Fig. 1 and Table 2).

## **Experimental**

*o*-Tolylamine (0.1 mmol, 10.7 mg), salicylaldehyde (0.1 mmol, 12.2 mg) and  $Cu(CH_3COO)_2 \cdot H_2O$  (0.1 mmol, 20.0 mg) were dissolved in methanol (15 ml). The mixture was stirred for 1 h and filtered. After keeping the filtrate in air for 7 d, blue block-shaped crystals were formed.

### Crystal data

$C_{\rm P}(C, \mathbf{H}, \mathbf{NO}) \subset \mathbf{I}$	$D = 1.442 \text{ M}_{\odot} \text{ m}^{-3}$
$[Cu(C_{14}H_{13}NO)_2Cl_2]$	$D_x = 1.443$ Mg m
$M_r = 556.95$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 726
a = 16.043 (3)  Å	reflections
b = 10.536 (2) Å	$\theta = 2.3-22.2^{\circ}$
c = 15.647 (3) Å	$\mu = 1.09 \text{ mm}^{-1}$
$\beta = 104.31 \ (3)^{\circ}$	T = 293 (2) K
V = 2562.8 (9) Å <sup>3</sup>	Block, blue
Z = 4	$0.32 \times 0.28 \times 0.27$ mm

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## Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. Intramolecular hydrogen bonds are shown as dashed lines.

#### Data collection

Bruker SMART CCD area-detector	2847 independent reflections
diffractometer	1462 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\rm int} = 0.073$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -16 \rightarrow 20$
$T_{\min} = 0.722, T_{\max} = 0.758$	$k = -13 \rightarrow 13$
8015 measured reflections	$l = -20 \rightarrow 18$

#### Refinement

Refinement on $F^2$	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.057$	independent and constrained
$wR(F^2) = 0.138$	refinement
S = 0.95	$w = 1/[\sigma^2(F_o^2) + (0.0522P)^2]$
2847 reflections	where $P = (F_0^2 + 2F_c^2)/3$
163 parameters	$(\Delta/\sigma)_{\rm max} < 0.001$
	$\Delta \rho_{\rm max} = 0.43 \ {\rm e} \ {\rm \AA}^{-3}$
	$\Delta \rho_{\rm min} = -0.28 \text{ e } \text{\AA}^{-3}$

#### Table 1

Selected	geometric	parameters	(Å,	°).	
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Cu1-O1	1.980 (3)	Cu1-Cl1	2.238 (2)
$O1^i$ -Cu1-O1	102.89 (17)	O1-Cu1-Cl1	106.04 (9)
O1 <sup>1</sup> -Cu1-Cl1	115.89 (9)	Cl1-Cu1-Cl1 <sup>1</sup>	110.23 (8)

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

## Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
N1-H1···O1	0.90 (1)	1.91 (4)	2.593 (4)	132 (4)

Atom H1 was located in a difference Fourier map and refined isotropically, with the N-H distance restrained to 0.90 (1) Å. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H distances in the range 0.93–0.96 Å and with  $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$  or  $1.5U_{eq}(\rm C)$ . The data are relatively weak, with almost half the reflections classed as 'unobserved'.



#### Figure 2

The crystal packing of (I), viewed along the a axis. Intramolecular hydrogen bonds are shown as dashed lines.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1998); 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 financially supported by the Jiangsu Province Education Commission Natural Science Foundation, People's Republic of China (grant No. 03KJB3101110), and by Natural Science Foundation of Jiangsu Province, China (grant No. BK2005042).

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