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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.005 Å R factor = 0.035 wR factor = 0.077 Data-to-parameter ratio = 17.7

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

Chloro{2-[tris(hydroxymethyl)methyliminomethyl]phenolato}copper(II)

In the title compound, $[Cu(C_{11}H_{14}NO_4)Cl]$, the tridentate Schiff base ligand coordinates to the metal atom through the N and O atoms, forming a square-planar coordination geometry.

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Comment

The chemistry of transition metal ion complexes of hydroxy (aryl and alkyl OH) rich molecules containing imine/amine groups is important in biomimetic chemistry (Cornman *et al.*, 1992). Many complexes of this kind have been reported (Asgedom & Rao, 1996; Dey, Rao, Saarenketo & Rissanen, 2002; Dey, Rao, Saarenketo, Rissanen & Kolehmainen, 2002). We report here a new copper(II) complex, (I), with a tridentate Schiff base ligand.



In compound (I), the Cu^{II} center is four-coordinated in a square-planar configuration by one N and two O atoms of the Schiff base ligand and one Cl atom. The Cu–O bond lengths



scheme. Displacement ellipsoids are drawn at the 50% probability level.

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H atoms are drawn as spheres of arbitrary radii.

metal-organic papers

are 1.885 (2) and 1.979 (2) Å; the shorter distance between copper and the phenoxy O atom indicates that the electronegativity of atom O1 is stronger than that of the other O atoms of the ligand.

Experimental

The ligand 2-[tris(hydroxymethyl)methyliminomethyl]phenol was prepared according to the literature procedure of Asgedom et al. (1996). Cuprous chloride (0.105 g, 0.5 mmol) was added to a solution of the ligand (0.111 g, 0.5 mmol) in water (10 ml). After stirring for a short time, the solution turned dark green. The filtrate was left for 2 d at room temperature and green needle-shaped crystals were obtained in about 62% yield.

Crystal data

 $[Cu(C_{11}H_{14}NO_4)Cl]$ $M_r = 323.22$ Tetragonal, $P\overline{4}2_1c$ a = 16.7345 (6) Å c = 8.7634 (6) Å V = 2454.1 (2) Å³ Z = 8 $D_x = 1.750 \text{ Mg m}^{-3}$ Data collection

Bruker APEX area-dectector diffractometer φ and φ scans Absorption correction: multi-scan (SADABS; Bruker, 2002) $T_{\min} = 0.679, T_{\max} = 0.873$ 20984 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.035$ wR(F²) = 0.077 S = 1.012918 reflections 165 parameters H-atom parameters constrained Cell parameters from 1879 reflections $\theta = 2.4 - 23.3^{\circ}$ $\mu = 2.00 \text{ mm}^{-1}$ T = 293 (2) K Needle, green $0.21 \times 0.09 \times 0.07 \text{ mm}$ 2010 · 1

Mo $K\alpha$ radiation

2661 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.049$ $\theta_{\text{max}} = 27.9^{\circ}$ $h = -21 \rightarrow 21$
$\begin{split} R_{\rm int} &= 0.049 \\ \theta_{\rm max} &= 27.9^{\circ} \\ h &= -21 \rightarrow 21 \end{split}$
$\theta_{\max} = 27.9^{\circ}$ $h = -21 \rightarrow 21$
$h = -21 \rightarrow 21$
$k = -21 \rightarrow 21$
$l = -11 \rightarrow 11$

 $w = 1/[\sigma^2(F_0^2) + (0.0354P)^2]$ + 1.812P] where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.36 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), 1265 Friedel pairs Flack parameter: 0.007 (15)

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.885 (2)	Cu1-O2	1.979 (2)
Cu1-N1	1.951 (2)	Cu1-Cl1	2.2366 (8)
O1-Cu1-N1	94.92 (10)	O1-Cu1-Cl1	93.56 (7)
O1-Cu1-O2	169.50 (10)	N1-Cu1-Cl1	171.25 (8)
N1-Cu1-O2	81.81 (10)	O2-Cu1-Cl1	90.22 (7)

H atoms were placed in idealized positions [N-H = 0.82 Å, C-H = 0.93–0.97 Å and $U_{iso}(H) = 1.2U_{eq}(C,N)$ and were included in the refinement in the riding-model approximation.

Data collection: SMART (Bruker, 2002): cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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References

Asgedom, G. & Rao, C. P. (1996). Inorg. Chem. 35, 5674-5683.

- Bruker (2002). SADABS, SAINT and SMART. Bruker AXS Inc., Madison, Wisconsin, USA.
- Cornman, C. R., Colpas, G. J., Hoeschele, J. D., Kampf, J. & Pecoraro, V. L. (1992). J. Am. Chem. Soc. 114, 9925-9933.

Dey, M., Rao, C. P., Saarenketo, P. K. & Rissanen, K. (2002). Inorg. Chem. Comm. pp. 380-383.

Dev, M., Rao, C. P., Saarenketo, P., Rissanen, K. & Kolehmainen, E. (2002). Eur. J. Inorg. Chem. pp. 2207-2215.

Flack, H. D. (1983). Acta Cryst. A39, 876-881.

- Johnson, C. K. (1976). ORTEPII. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.