

Redetermination of bis{[(2-hydroxyphenylmethyl)bis(2-pyridylmethyl)aminato]copper(II)} diperchlorate

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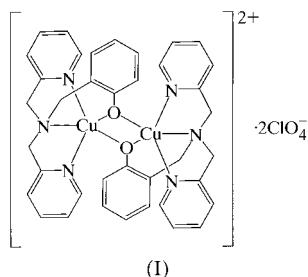
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The structure of the title compound, $[\text{Cu}_2(\text{C}_{19}\text{H}_{18}\text{N}_3\text{O})_2](\text{ClO}_4)_2$, was reported with insufficient accuracy because of a twinning problem by Adams, Bailey, Campbell, Fenton & He [J. Chem. Soc. Dalton Trans. (1996), pp. 2233–2237]. The dinuclear phenolate-bridged Cu^{II} complex has an inversion centre.

Comment

The title copper(II) complex, $[\text{Cu}_2(\text{phpy})_2](\text{ClO}_4)_2$ [phpy is (2-hydroxyphenylmethyl)bis(2-pyridylmethyl)aminato], (I), which has a long Cu–phenolic O atom bond, can be considered as a nobel model compound for galactose oxidase (Ito *et al.*, 1998).



Adams *et al.* (1996) noted that the structure was difficult to refine because of a twinning problem.

Experimental

In the present study, single crystals of (I) were grown from an acetonitrile solution as needles elongated along **a**.

Crystal data

$[\text{Cu}_2(\text{C}_{19}\text{H}_{18}\text{N}_3\text{O})_2](\text{ClO}_4)_2$	$D_x = 1.642 \text{ Mg m}^{-3}$
$M_r = 934.74$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 25 reflections
$a = 11.380 (1) \text{ \AA}$	$\theta = 10\text{--}15^\circ$
$b = 9.877 (2) \text{ \AA}$	$\mu = 1.334 \text{ mm}^{-1}$
$c = 17.400 (2) \text{ \AA}$	$T = 298 \text{ K}$
$\beta = 104.79 (1)^\circ$	Needle, green
$V = 1891.0 (5) \text{ \AA}^3$	$0.70 \times 0.15 \times 0.10 \text{ mm}$
$Z = 2$	

Data collection

Rigaku AFC-5S diffractometer	$R_{\text{int}} = 0.059$
$\theta\text{--}2\theta$ scans	$\theta_{\text{max}} = 27.5^\circ$
Absorption correction: by integration (Coppens <i>et al.</i> , 1965)	$h = 0 \rightarrow 15$
$T_{\text{min}} = 0.662$, $T_{\text{max}} = 0.897$	$k = 0 \rightarrow 13$
4819 measured reflections	$l = -23 \rightarrow 23$
4338 independent reflections	3 standard reflections
2393 reflections with $I > 2\sigma(I)$	every 100 reflections
	intensity decay: 5.5%

Refinement

Refinement on F^2	H-atom parameters not refined
$R(F) = 0.063$	$w = 1/[\sigma^2(F_o^2) + \{0.1(F_o^2 + 2F_c^2)/3\}^2]$
$wR(F^2) = 0.185$	$(\Delta/\sigma)_{\text{max}} = 0.009$
$S = 0.98$	$\Delta\rho_{\text{max}} = 0.59 \text{ e \AA}^{-3}$
4338 reflections	$\Delta\rho_{\text{min}} = -0.63 \text{ e \AA}^{-3}$
262 parameters	

Table 1
Selected geometric parameters (\AA , $^\circ$).

Cu1–O1	2.203 (4)	Cu1–N2	2.037 (4)
Cu1–O1 ⁱ	1.938 (3)	Cu1–N3	2.000 (5)
Cu1–N1	2.013 (5)		
O1–Cu1–O1 ⁱ	80.4 (2)	Cu1–O1–Cu1 ⁱ	99.6 (2)

Symmetry code: (i) $1 - x, -y, 1 - z$.

Positional parameters of all the H atoms were calculated geometrically and fixed with $U(\text{H}) = 1.2U_{\text{eq}}$ (parent atom).

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1993); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation, 1999); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *TEXSAN*; software used to prepare material for publication: *TEXSAN*.

References

- Adams, H., Bailey, N. A., Campbell, I. K., Fenton, D. E. & He, Q.-Y. (1996). J. Chem. Soc. Dalton Trans. pp. 2233–2237.
- Altomare, A., Cascarano, G., Giacovazzo, C., Guagliardi, A., Burla, M. C., Polidori, G. & Camalli, M. (1994). J. Appl. Cryst. **27**, 435.
- Coppens, P., Leiserowitz, L. & Rabinovich, D. (1965). Acta Cryst. **18**, 1035–1038.
- Ito, S., Nishino, S., Itoh, H., Ohba, S. & Nishida, Y. (1998). Polyhedron, **17**, 1637–1642.
- Molecular Structure Corporation (1993). *MSC/AFC Diffractometer Control Software*. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- Molecular Structure Corporation (1999). *TEXSAN*. Version 1.10. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.