metal-organic papers

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Tong-Tao Xu,^a Xing-You Xu,^b Lu-De Lu,^a* Jia Ni^c and Xu-Jie Yang^a

 ^aMaterials Chemistry Laboratory, Nanjing University of Science & Technology, Nanjing 210094, People's Republic of China,
^bDepartment of Chemical Engineering, Huaihai Institute of Technology, Lianyungang 222005, People's Republic of China, and ^cCentral Laboratory of Shantou University, Shantou 515062, People's Republic of China

Correspondence e-mail: xutongtao_1968@163.com

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.007 Å R factor = 0.053 wR factor = 0.144 Data-to-parameter ratio = 13.0

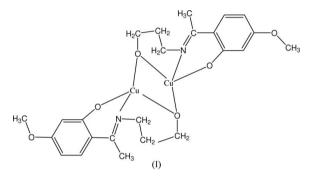
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Bis{µ-(*E*)-5-methoxy-2-[1-(3-oxidopropyl-ideneamino)ethyl]phenolato}copper(II)

The title dimeric Cu^{II} complex, $[Cu_2(C_{12}H_{15}NO_3)_2]$, is centrosymmetric. Two Cu^{II} atoms related by an inversion center are bridged by two O atoms of propanolamine. The Cu^{II} atom assumes a distorted square-planar geometry.

Comment

Paeonol, or 2-hydroxy-4-methoxyacetophenone, is an effective component of Chinese traditional medicines. As part of our ongoing investigation on Schiff base compounds of paeonol, we present here the structure of the title Cu^{II} complex, (I).



The molecular structure of (I) is shown in Fig. 1. While the 5-methoxy-2-[1-(3-oxidopropylideneamino)ethyl]phenolate dianion chelates to an individual Cu^{II} atom through the amino and two diprotonated hydroxy groups, the deprotonated hydroxy atom O3 bridges two Cu^{II} atoms, forming a centro-symmetric dimeric complex. The Cu^{II} atom assumes a distorted square-planar coordination (Table 1). Within the dimeric complex, the $Cu \cdot Cu$ separation is 3.0299 (11) Å.

Experimental

To a stirred solution of paeonol (1.0 mmol) and $\text{Cu}(\text{ClO}_4)_2 \cdot 6\text{H}_2\text{O}$ (1.0 mmol) in 25 ml absolute methanol was added dropwise a solution of propanolamine (1.0 mmol) in 10 ml absolute methanol at room temperature. After stirring for 3 h at 320 K, the resulting precipitate was filtered off, washed with methanol and dried *in vacuo*. Single crystals of (I) were obtained by slow evaporation of the filtrate after 5 d.

Crystal data $\begin{bmatrix} Cu_2(C_{12}H_{15}NO_3)_2 \end{bmatrix}$ $M_r = 569.6$ Monoclinic, $P2_1/c$ a = 6.774 (1) Å b = 6.703 (1) Å c = 25.775 (4) Å $\beta = 95.493$ (2)° V = 1165.1 (3) Å³

Z = 2 $D_x = 1.624 \text{ Mg m}^{-3}$ Mo K α radiation $\mu = 1.87 \text{ mm}^{-1}$ T = 293 (2) K Prism, blue $0.21 \times 0.08 \times 0.05 \text{ mm}$ Received 25 April 2006 Accepted 20 May 2006

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Bruker SMART APEX areadectector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002) $T_{\min} = 0.695, T_{\max} = 0.912$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.053$ $wR(F^2) = 0.144$ S = 1.052034 reflections 156 parameters H-atom parameters constrained 5982 measured reflections 2034 independent reflections 1788 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.032$ $\theta_{\text{max}} = 25.0^{\circ}$

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.0738P)^2 \\ &+ 3.3311P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} = 0.001 \\ \Delta\rho_{\rm max} = 0.74 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.54 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.873 (3)	N1-C8	1.302 (6)
Cu1-O3 ⁱ	1.935 (3)	N1-C10	1.475 (6)
Cu1-O3	1.912 (3)		
O1-Cu1-O3 ⁱ	92.16 (15)	O3-Cu1-N1	98.56 (16)
O1-Cu1-O3	167.99 (15)	O3 ⁱ -Cu1-N1	174.31 (16)
O1-Cu1-N1	93.28 (16)	Cu1 ⁱ -O3-Cu1	103.93 (16)
$O3^i$ -Cu1-O3	76.07 (16)		

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

Methyl H atoms were placed in calculated positions, with C–H = 0.96 Å, and torsion angles were refined to fit the electron density $[U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm C})]$. Other H atoms were positioned geometrically, with C–H = 0.93 (aromatic) and 0.97 Å (methylene), and refined as riding, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve

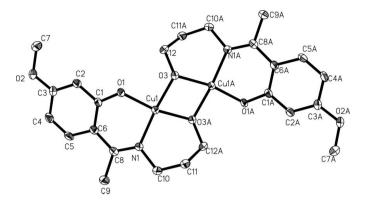


Figure 1

The molecular structure of (I), shown with 30% probability displacement ellipsoids [symmetry code: (i) -x, -y, -z]. H atoms have been omitted.

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*.

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References

Bruker (2003). SAINT (Version 6.45A) and SMART (Version 5.059). Bruker AXS Inc., Madison, Wisconsin, USA.

Sheldrick, G. M. (1997a). SHELXL97 and SHELXS97. University of Göttingen,Germany.

- Sheldrick, G. M. (1997b). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.
- Sheldrick, G. M. (2002). SADABS. Version 2.03. University of Göttingen,-Germany.