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Key indicators

Single-crystal X-ray study
 $T = 295$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.036
 wR factor = 0.117
Data-to-parameter ratio = 14.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis(2-acetyl-5-hydroxyphenolato- $\kappa^2\text{O},\text{O}'$)copper(II)

The Cu atom in the title compound, $[\text{Cu}(\text{C}_8\text{H}_7\text{O}_3)_2]$, lies on a center of inversion and exists in a square-planar environment that is defined by the four chelating O atoms. Intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds [$\text{O}\cdots\text{O} = 2.711(3)$ Å] link adjacent molecules into a layer structure.

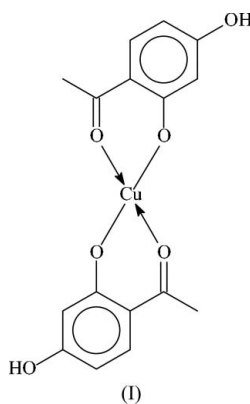
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Comment

The deprotonated 2-hydroxyacetophenone anion has been used to chelate to a number of divalent transition metal ions; the copper(II) derivatives include, for example, (2-acetylphenolato)(2,2'-bipyridyl)perchloratocopper (Elmali *et al.*, 2002) and bis(2-acetylphenolato)(4-methylpyridine)copper (Duckworth & Stephenson, 1969). In the second example, the metal atom is coordinated by two 4-picoline molecules; the title compound, (I) (Fig. 1), crystallizes from pyridine without any solvent. The anion chelates to the metal atom which lies on an inversion center, with different Cu—O distances (Table 1). The Cu atom exists in a square-planar geometry, and adjacent molecules are linked by intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 2) into a layer structure.



Experimental

2,4-Dihydroxyacetophenone (0.50 g, 3.6 mmol) and 4-nitroaniline (0.50 g, 3.6 mmol) were reacted in boiling ethanol, and to the mixture was added copper acetate monohydrate (0.36 g, 1.8 mmol). The heating was continued for an hour. The solid that was isolated upon removal of the solvent was recrystallized from pyridine.

Crystal data

$[\text{Cu}(\text{C}_8\text{H}_7\text{O}_3)_2]$
 $M_r = 365.81$
Orthorhombic, $Pbca$
 $a = 7.022(1)$ Å
 $b = 13.667(2)$ Å
 $c = 15.192(2)$ Å
 $V = 1458.1(3)$ Å³
 $Z = 4$

$D_x = 1.666$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 766
reflections
 $\theta = 2.7-27.0^\circ$
 $\mu = 1.53$ mm⁻¹
 $T = 295(2)$ K
Prism, dark brown

0.24 × 0.14 × 0.11 mm

Data collection

Bruker SMART area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 1999)
 $T_{\min} = 0.724$, $T_{\max} = 0.850$
 8105 measured reflections

1591 independent reflections
 1057 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.061$
 $\theta_{\max} = 27.0^\circ$
 $h = -6 \rightarrow 8$
 $k = -17 \rightarrow 17$
 $l = -18 \rightarrow 19$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.117$
 $S = 1.04$
 1591 reflections
 111 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0694P)^2 + 0.0274P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.35 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.34 \text{ e \AA}^{-3}$

Table 1

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

Cu1—O1	1.912 (2)	Cu1—O2	1.883 (2)
O1—Cu1—O1 ⁱ	180	O1—Cu1—O2 ⁱ	87.47 (8)
O1—Cu1—O2	92.53 (8)		

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

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O3—H3O ⁱⁱ ···O2 ⁱⁱ	0.83 (1)	1.90 (1)	2.710 (3)	166 (4)

Symmetry code: (ii) $x - \frac{1}{2}, -y + \frac{3}{2}, -z + 1$.

The C-bound H atoms were positioned geometrically (C—H = 0.93 \AA for the aromatic H atoms and 0.98 \AA for the methyl H atoms) and were included in the refinement in the riding-model approximation, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{parent atom})$ for the aromatic C atoms and 1.5 U_{eq} for the methyl H atoms. The methyl group was rotated to fit the electron density. The hydroxy H atom was located in a difference Fourier map, and was refined with a distance restraint of O—H = 0.85 (1) \AA .

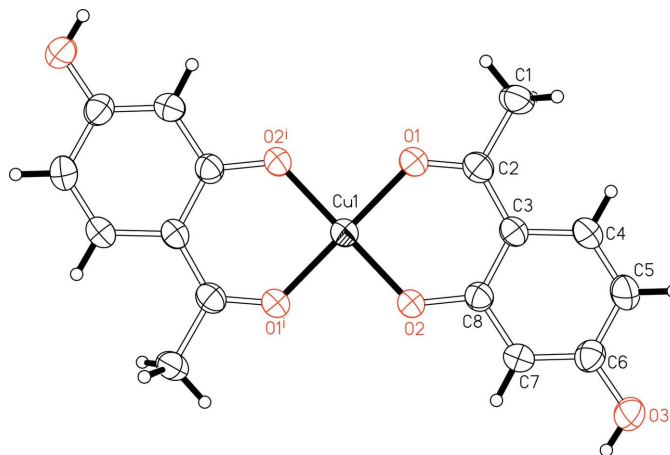


Figure 1

ORTEP plot (Johnson, 1976) of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as spheres of arbitrary radii. The Cu atom lies at the center of inversion ($\frac{1}{2}, \frac{1}{2}, \frac{1}{2}$) [symmetry code: (i) $1 - x, 1 - y, 1 - z$].

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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