

Bis(3-methylbenzoato- κ^2O,O')(phenanthroline- κ^2N,N')copper(II)Tu-Gen Xu[‡] and Duan-Jun Xu*Department of Chemistry, Zhejiang University,
Hangzhou 310027, People's Republic of China[‡] Current address: Department of Chemistry,
Hangzhou Teachers' College, People's Republic
of China.

Correspondence e-mail: xudj@mail.hz.zj.cn

Key indicators

Single-crystal X-ray study

 $T = 295$ KMean $\sigma(C-C) = 0.003$ Å R factor = 0.037 wR factor = 0.091

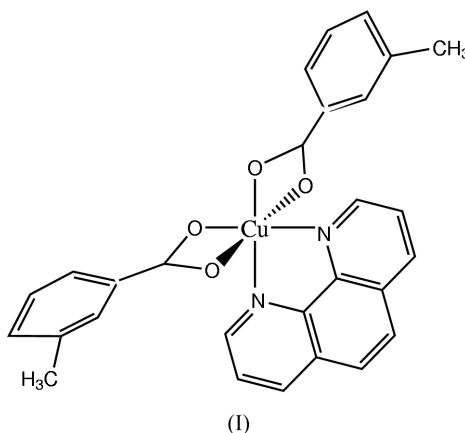
Data-to-parameter ratio = 16.7

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

In the title complex, $[Cu(C_8H_7O_2)_2(C_{12}H_8N_2)]$, the Cu^{II} atom is located on a twofold axis and assumes a distorted octahedral coordination geometry, formed by a phenanthroline molecule and two methylbenzoate anions, all in chelating mode. The carboxylate group chelates to the Cu^{II} atom with greatly differing $Cu-O$ distances, *viz.* 2.5923 (19) Å in the axial direction and 1.9435 (14) Å in the equatorial plane.

Comment

The Jahn–Teller distortion of copper(II) complexes is well known. Most copper(II) complexes display an elongated distortion, and the coordinate bonds in the axial direction are usually longer than those in the equatorial coordination plane by 0.2–0.6 Å. In the title copper(II) complex, (I), a pair of long $Cu-O$ bonds is observed.



The molecular structure of (I) is shown in Fig. 1. The Cu^{II} atom is located on a twofold axis and is surrounded by a phenanthroline (phen) molecule and two methylbenzoate anions in a distorted octahedral coordination geometry. Two O atoms from carboxyl groups and two phen N atoms form a tetrahedrally distorted equatorial coordination plane, with a dihedral angle of 9.7 (2)° between the $Cu/O1/O1^i$ and $Cu/N1/N1^i$ planes [symmetry code: (i) $1 - x, y, \frac{1}{2} - z$]. The bond lengths in the equatorial plane are normal (Table 1). In the axial direction, the $Cu-O2$ distance is longer than the $Cu-O1$ distance in the equatorial plane by 0.649 (2) Å.

The $Cu-O1-C7$ angle of 104.54 (13)° is similar to values found in copper(II) complexes with a chelating benzoate ligand, for example, 104.93 (16)° (Bakalbassis & Terzis, 1994), 106.1 (4)° (Cano *et al.*, 1997) and 100.50 (17)° (Parkanyi & Speier, 1995), but is much smaller than those in copper(II) complexes with a monodentate benzoate ligand, for example,

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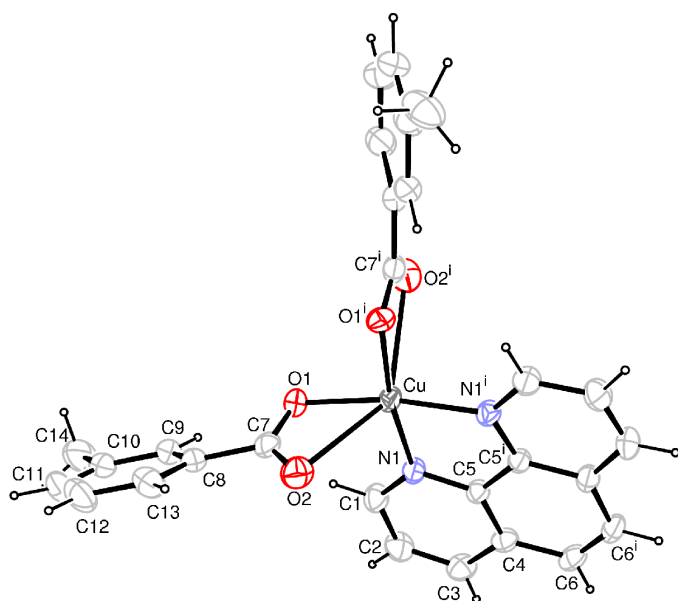


Figure 1
The molecular structure of (I), shown with 30% probability displacement ellipsoids. [Symmetry code: (i) $1 - x, y, \frac{1}{2} - z$.]

$127.3(3)^\circ$ (Hoang *et al.*, 1992) and $131.80(10)^\circ$ (Rodrigues *et al.*, 1999). This suggests the existence of a bonding interaction between atoms Cu and O2 in (I).

Besides the elongated Jahn–Teller distortion, the smaller O2–Cu–O2ⁱ angle of $137.53(8)^\circ$ is also the possible reason for the larger difference in Cu–O bond distances within the same carboxylate group.

A C–H··· π interaction occurs between the methyl group and the ring containing atom C6 (*Cg* denotes the centroid of this ring) of a phen ligand of a neighbouring molecule (Fig. 2), with an H···*Cg* distance of 3.01 Å and a C–H···*Cg* angle of 126° . The π – π stacking between phen ligands which is usually observed in metal complexes is not observed in (I).

Experimental

An acetonitrile–water solution (20 ml) containing CuSO₄·5H₂O (0.25 g, 1 mmol), 3-methylbenzoic acid (1.36 g, 10 mmol) and NaOH (0.40 g, 10 mmol) was refluxed for 1 h, yielding a blue precipitate. After adding phen (0.2 g, 1.1 mmol), this blue precipitate gradually dissolved. The resulting dark-blue solution was refluxed for a further 3 h and then filtered. Blue single crystals of (I) were obtained from the filtrate after one week.

Crystal data

[Cu(C₈H₇O₂)₂(C₁₂H₈N₂)]
M_r = 514.02
 Monoclinic, *C*2/*c*
a = 21.1482 (13) Å
b = 9.9476 (12) Å
c = 11.9730 (8) Å
 β = 112.690 (2)°
V = 2323.9 (4) Å³
Z = 4

D_x = 1.469 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 9404 reflections
 θ = 2.5–26.0°
 μ = 0.98 mm⁻¹
T = 295 (2) K
 Platelet, blue
 0.31 × 0.27 × 0.08 mm

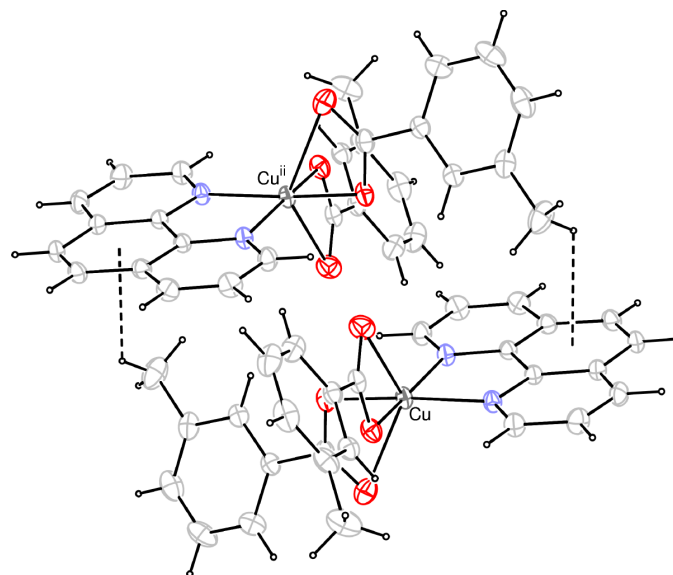


Figure 2
The C–H··· π interaction between neighbouring complex molecules [symmetry code: (ii) $1 - x, 1 - y, 1 - z$.]

Data collection

Rigaku R-AXIS RAPID diffractometer
 ω scans
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.740, T_{\max} = 0.920$
 10 454 measured reflections

2666 independent reflections
 2238 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$
 $\theta_{\max} = 27.5^\circ$
 $h = -27 \rightarrow 27$
 $k = -12 \rightarrow 12$
 $l = -15 \rightarrow 15$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.091$
 $S = 1.04$
 2666 reflections
 160 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0446P)^2 + 1.5001P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.30 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.24 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cu–O1	1.9435 (14)	Cu–N1	2.0113 (16)
Cu–O2	2.5923 (19)		
O2–Cu–N1	95.30 (6)	O2–Cu–O1 ⁱ	94.13 (6)
O2–Cu–N1 ⁱ	117.09 (6)	O2–Cu–O2 ⁱ	137.53 (8)
O2–Cu–O1	56.13 (6)	C7–O1–Cu	104.54 (13)

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

H atoms on aromatic rings were placed in calculated positions, with C–H = 0.93 Å, and were included in the final cycles of refinement in the riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. Methyl H atoms were placed in calculated positions, with C–H = 0.96 Å, and refined in the riding model, with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Shel-

drick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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