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Key indicators

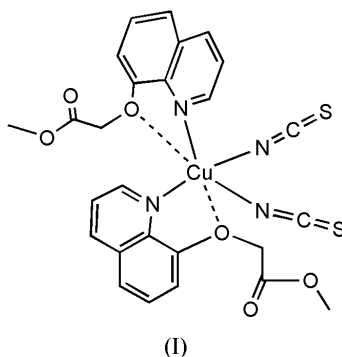
Single-crystal X-ray study
 $T = 193$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
 R factor = 0.055
 wR factor = 0.101
Data-to-parameter ratio = 17.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis(isothiocyanato- κN)bis[8-(methoxycarbonyl-
methoxy)quinoline- κN]copper(II)

In the title mononuclear complex, $[\text{Cu}(\text{NSC})_2(\text{C}_{13}\text{H}_{11}\text{NO}_3)_2]$, the isothiocyanate ions and 8-(methoxycarbonylmethoxy)-quinoline molecules act as monodentate ligands. The ligand geometry about the Cu centre, which lies on a twofold rotation axis, can be described as square-planar or grossly distorted octahedral, formed by two N atoms of the isothiocyanate ions, two quinoline N atoms and two methoxy O atoms of the 8-(methoxycarbonylmethoxy)quinoline ligands. The N—Cu—N *cis* angles around the Cu atom vary from $88.38(12)$ to $94.27(15)^\circ$ and the Cu—N bond distances are $1.950(2)$ and $2.024(2)$ Å.

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Comment

Derivatives of quinoline are well-known ligands in transition metal coordination compounds (Ghedini *et al.*, 2002; Inomata *et al.*, 1999; Jotterand *et al.*, 2001; Sivakolunthu *et al.*, 1998). Such compounds are being increasingly studied in modelling biological systems, one target being the metal binding site of copper metalloenzymes (Sorrel, 1989). In the light of this interest, we have prepared the biologically relevant ligand 8-(methoxycarbonylmethoxy)quinoline and its title copper complex, (I).



As shown in Fig. 1, (I) is mononuclear. Each Cu^{II} centre has a slightly distorted square geometry (Table 1) comprising of two isothiocyanate N atoms and two quinoline N atoms of the 8-(methoxycarbonylmethoxy)quinoline ligands. The coordination plane is defined by N1, N2, N2ⁱ and N1ⁱ [symmetry code: (i) $1 - x, y, \frac{1}{2} - z$]. Atoms N1, N2, N2ⁱ and N1ⁱ show deviations of $0.3295(1)$, $-0.3248(1)$, $0.3248(1)$ and $-0.3295(1)^\circ$, respectively, from the least-squares mean plane, and the Cu centre is constrained to lie in the plane by symmetry. The N—Cu—N angles vary from $88.38(12)$ to $94.27(15)^\circ$. The Cu—N_q ($q = \text{quinoline}$) bond length of

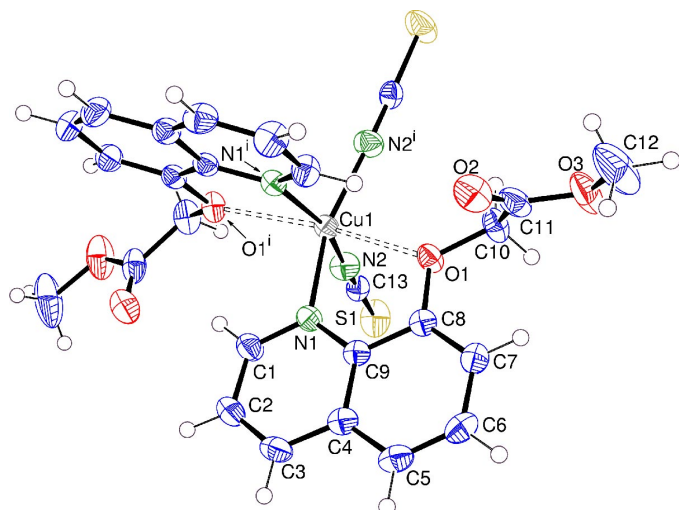


Figure 1
View of (I), shown with 30% probability displacement ellipsoids and small spheres for H atoms [symmetry code: (i) $1 - x, y, \frac{1}{2} - z$]. The long Cu1...O1 interactions are indicated by dashed lines.

2.024 (2) Å is somewhat longer than the Cu—N_{NSC} bond length of 1.950 (2) Å, although both distances fall within the expected range of distances for Cu—N bonds (Breneman & Parker, 1993; Kožišek & Kabešová, 1990). The NSC groups in (I) are almost linear [179.0 (3)°], and slight bending is observed in the Cu—N—C linkage [177.1 (2)°]. The mean planes of the two quinoline moieties form a dihedral angle of 86.5 (1)°. There is also a pair of O atoms, O1 and O1ⁱ, each some 2.4708 (19) Å from Cu1. If these are considered to be chemically significant interactions, a grossly distorted octahedral coordination about Cu1 results (Fig. 1).

The molecular packing in (I) is influenced by intermolecular π – π interactions which exist between the parallel quinoline rings of neighbouring complex molecules, with a separation of approximately 3.56 Å.

Experimental

Solid KSCN (19.4 mg, 2 mmol) was added slowly with continuous stirring to a solution of Cu(ClO₄)₄·6H₂O (57 mg, 1 mmol) in distilled water (5 ml). 8-(Methoxycarbonylmethoxy)quinoline (21.8 mg, 1 mmol) was dissolved in methanol (20 ml) and the two solutions were mixed and stirred for 5 h. The resulting solution was filtered. Pale green square-prismatic crystals suitable for X-ray diffraction analysis were obtained by slow evaporation of the resulting solution at room temperature. Analysis found: C 50.12, H 3.57, N 9.09%; calculated for C₂₆H₂₂CuN₄O₆S₂: C 50.85, H 3.61, N 9.12%

Crystal data

[Cu(NSC) ₂ (C ₁₃ H ₁₁ NO ₃) ₂]	$D_x = 1.509 \text{ Mg m}^{-3}$
$M_r = 614.14$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 5205 reflections
$a = 16.921 (4) \text{ \AA}$	$\theta = 3.0\text{--}27.5^\circ$
$b = 10.073 (2) \text{ \AA}$	$\mu = 1.01 \text{ mm}^{-1}$
$c = 17.590 (5) \text{ \AA}$	$T = 193 (2) \text{ K}$
$\beta = 115.647 (4)^\circ$	Block, green
$V = 2702.6 (11) \text{ \AA}^3$	$0.38 \times 0.24 \times 0.10 \text{ mm}$
$Z = 4$	

Data collection

Rigaku Mercury diffractometer	2849 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.042$
Absorption correction: multi-scan (Jacobson, 1998)	$\theta_{\text{max}} = 27.5^\circ$
$T_{\text{min}} = 0.700, T_{\text{max}} = 0.906$	$h = -21 \rightarrow 17$
14799 measured reflections	$k = -13 \rightarrow 13$
3086 independent reflections	$l = -20 \rightarrow 22$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0246P)^2 + 6.2754P]$
$R[F^2 > 2\sigma(F^2)] = 0.055$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.101$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.22$	$\Delta\rho_{\text{max}} = 0.42 \text{ e \AA}^{-3}$
3086 reflections	$\Delta\rho_{\text{min}} = -0.43 \text{ e \AA}^{-3}$
179 parameters	
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

Cu1—N2	1.950 (2)	Cu1—O1	2.4708 (19)
Cu1—N1	2.024 (2)		
N2 ⁱ —Cu1—N2	94.27 (15)	N2—Cu1—O1	101.17 (9)
N2—Cu1—N1	91.79 (10)	N1—Cu1—O1	72.38 (7)
N2—Cu1—N1 ⁱ	160.81 (9)	N2—Cu1—O1 ⁱ	88.58 (9)
N1—Cu1—N1 ⁱ	88.38 (12)	N1—Cu1—O1 ⁱ	97.17 (8)

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

H atoms were included in calculated positions and refined as riding, with C—H distances of 0.93 (aromatic), 0.99 (ethyl) and 0.98 Å (methyl), and with U_{iso} (aromatic and ethyl) = $1.2U_{\text{eq}}(\text{C})$ and U_{iso} (methyl) = $1.5U_{\text{eq}}(\text{C})$.

Data collection: *CrystalClear* (Molecular Structure Corporation, 2000; Rigaku Corporation, 1999); cell refinement: *CrystalClear*; data reduction: *CrystalStructure* (Rigaku/MS, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

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