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Key indicators

Single-crystal X-ray study
 $T = 150$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.029
 wR factor = 0.071
Data-to-parameter ratio = 19.6

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

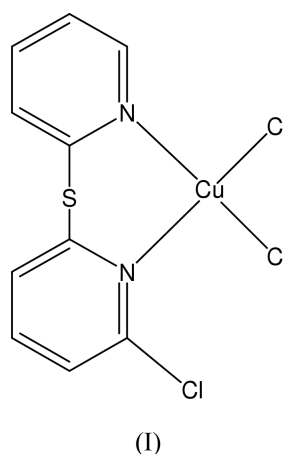
Dichloro[2-chloro-6-(mercaptopyridin-2-yl)pyridine]copper(II)

The title compound, $[\text{CuCl}_2(\text{C}_{10}\text{H}_7\text{ClN}_2\text{S})]$, crystallizes as neutral CuCl_2L [$L = 2\text{-chloro-6-(mercaptopyridin-2-yl)pyridine}$] molecular units. The copper(II) centre adopts a distorted four-coordinate geometry comprising the two pyridinyl N atoms of the bidentate chelating ligand [$\text{Cu}-\text{N}$ 2.012 (2) and 2.029 (2) Å] and the two chloride anions [$\text{Cu}-\text{Cl}$ 2.2049 (6) and 2.2335 (6) Å]. The dihedral angle δ quantifying the extent of the distortion of the geometry from square planar ($\delta = 0^\circ$) towards tetrahedral ($\delta = 90^\circ$) is 36.81 (6)°. There is evidence for the formation of dimeric units *via* $\text{Cl}\cdots\text{S}$ interactions.

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Comment

The structure of CuCl_2L [$L = 2\text{-chloro-6-(mercaptopyridin-2-yl)pyridine}$], (I), is that of a neutral molecular complex (Fig. 1) in which the copper(II) centre has a distorted four-coordinate geometry comprising the two N atoms of the bidentate chelating ligand L [$\text{Cu}-\text{N}$ 2.012 (2) and 2.029 (2) Å] and the two chloride anions [$\text{Cu}-\text{Cl}$ 2.2049 (6) and 2.2335 (6) Å]. The coordination geometry lies between square planar and tetrahedral, with a dihedral angle (δ) between the CuN_2 and CuCl_2 planes of 36.81 (6)°. Distortion from planarity is unusual for copper(II) complexes. Here, it is attributed to steric repulsion between a coordinated chloride anion and the Cl atom at the 2-position of the pyridine ring [$\text{Cl}\cdots\text{Cl}$ 3.787 (1) Å].



There is considerable evidence for dimer formation through $\text{Cl}\cdots\text{S}$ interactions (Fig. 2). Not only is the $\text{Cu}\cdots\text{Cl}$ distance to the chloride anion involved in dimer formation [2.2335 (6) Å] significantly longer than that to the other chloride anion [2.2049 (6) Å], but also the U_{eq} value for the former [0.02691 (13) Å²] is considerably smaller than that for the

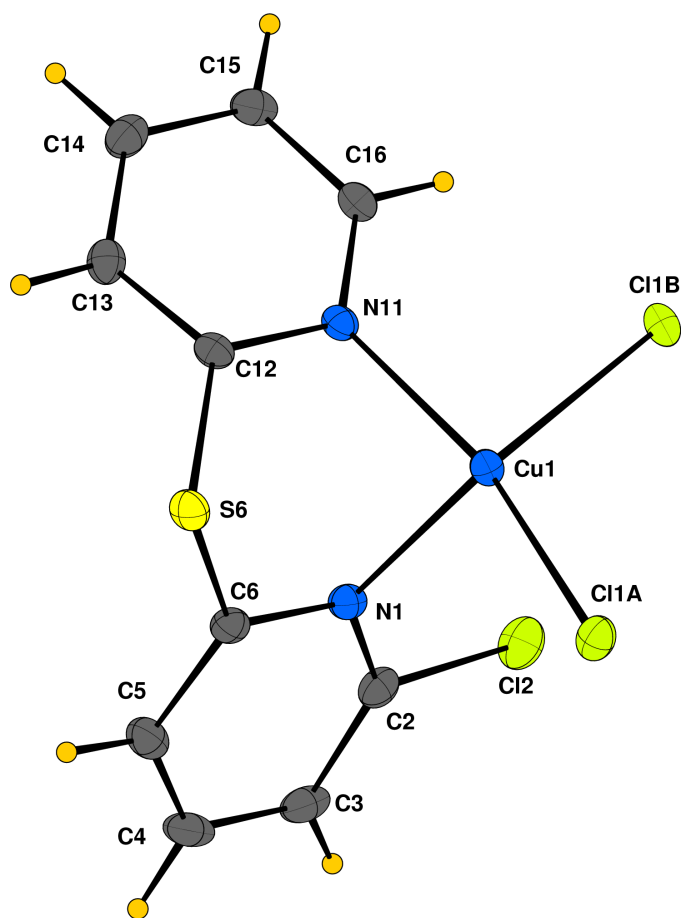


Figure 1
A view of the title compound showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

latter [0.03713 (16) Å²]. A search of the Cambridge Structural Database (Allen & Kennard, 1993) revealed that the intradimer Cl⋯S separation [3.4085 (9) Å] is typical for the distance separating the chloride anion of a CuCl₂ fragment and the S atom of either an aromatic or an aliphatic thioether.

Experimental

The title compound, (I), was obtained as a green lath-shaped crystal in the midst of a crop of green acicular crystals. Both had been produced by vapour phase diffusion of diethyl ether into a methanol solution of the product of the reaction between a methanol solution (10 ml) of copper(II) chloride dihydrate (0.199 g, 0.704 mmol) and a methanol solution (20 ml) of 2-(3,5-dimethylpyrazol-1-yl)-6-(mercaptopyridin-2-yl)pyridine (0.198 g, 0.701 mmol). This last ligand had been previously prepared by treatment of 2-chloro-6-(mercaptopyridin-2-yl)pyridine with 3,5-dimethylpyrazole in the presence of sodium hydride. Mass spectrometric (electrospray), IR and analytical studies of the bulk product suggested that it comprised dichloro[2-(3,5-dimethylpyrazol-1-yl)-6-(mercaptopyridin-2-yl)pyridine]copper(II) (yield: 0.123 g, 0.290 mmol, 42%). It is assumed that the lath-shaped crystal was a rogue crystal resulting from the presence of a small amount of unreacted 2-chloro-6-(mercaptopyridin-2-yl)pyridine in the sample of 2-(3,5-dimethylpyrazol-1-yl)-6-(mercaptopyridin-2-yl)pyridine.

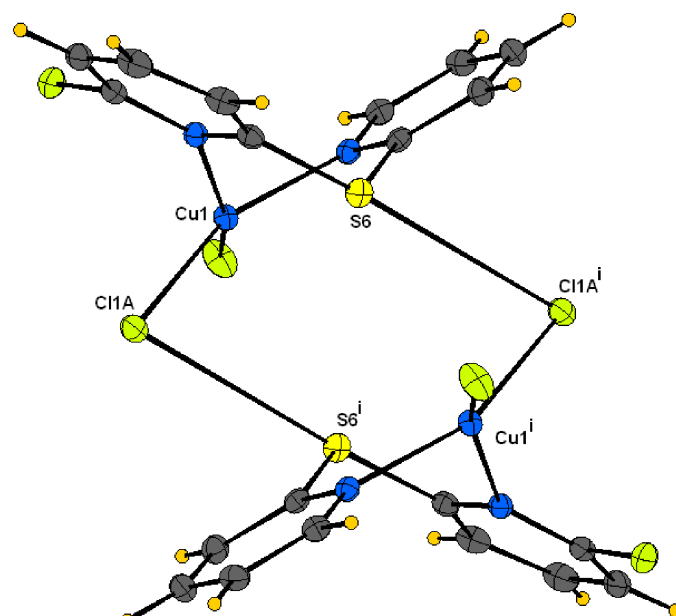


Figure 2
A view of the structure showing the pairwise centrosymmetric Cl⋯S interactions leading to dimer formation. Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code (i): 3/2 -x, 1/2 -y, 1 -z.]

Crystal data

C₁₀H₇Cl₃CuN₂S
M_r = 357.13
Monoclinic, C2/c
a = 13.1739 (12) Å
b = 9.6928 (9) Å
c = 21.137 (2) Å
β = 104.510 (2)°
V = 2612.9 (4) Å³
Z = 8

D_x = 1.816 Mg m⁻³
Mo Kα radiation
Cell parameters from 3638 reflections
θ = 2.7–26.8°
μ = 2.42 mm⁻¹
T = 150 (2) K
Lath, green
0.36 × 0.11 × 0.06 mm

Data collection

Bruker SMART APEX area-detector diffractometer
ω scans
Absorption correction: multi-scan (SADABS; Bruker, 2001)
T_{min} = 0.782, T_{max} = 0.870
12577 measured reflections

3191 independent reflections
2709 reflections with I > 2σ(I)
R_{int} = 0.026
θ_{max} = 28.2°
h = -16 → 17
k = -12 → 12
l = -25 → 26

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.029
wR(F²) = 0.071
S = 1.04
3019 reflections
154 parameters
H-atom parameters constrained

w = 1/[σ²(F_o²) + (0.035P)² + 3.807P]
where P = (F_o² + 2F_c²)/3
(Δ/σ)_{max} = 0.001
Δρ_{max} = 0.42 e Å⁻³
Δρ_{min} = -0.25 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Cu1—Cl1A	2.2335 (6)	Cu1—N1	2.029 (2)
Cu1—Cl1B	2.2049 (6)	Cu1—N11	2.012 (2)
Cl1A—Cu1—Cl1B	97.73 (2)	N1—Cu1—N11	87.58 (7)
Cl1A—Cu1—N1	90.65 (5)	C2—N1—Cu1	121.67 (15)
Cl1A—Cu1—N11	152.02 (5)	C6—N1—Cu1	119.37 (14)
Cl1B—Cu1—N11	95.96 (5)	C12—N11—Cu1	118.91 (14)
Cl1B—Cu1—N1	154.01 (5)	C16—N11—Cu1	123.11 (15)

After location from ΔF syntheses, the aromatic H atoms were placed geometrically and refined with a riding model in which the C—H distance was constrained to be 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT* and *SHELXTL/PC* (Bruker, 1997); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2001).

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