

Dichloro[3,3'-bis(ethoxycarbonyl)-  
2,2'-bipyridyl- $\kappa^2N,N'$ ]copper(II)Ben-Lai Wu, You-Fu Zhou, Lei  
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## Key indicators

Single-crystal X-ray study

T = 293 K

Mean  $\sigma(C-C)$  = 0.009 Å

R factor = 0.068

wR factor = 0.138

Data-to-parameter ratio = 13.2

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

In the title compound,  $[CuCl_2(C_{16}H_{16}N_2O_4)]$ , which crystallizes in the monoclinic space group  $C2/c$ , molecules are located on twofold rotation axes passing through the Cu centers and bisecting the bipyridyl ligands. They are interconnected by weak  $C-H \cdots O$  and  $C-H \cdots Cl$  interactions to form a two-dimensional network which stacks along the  $c$  axis.

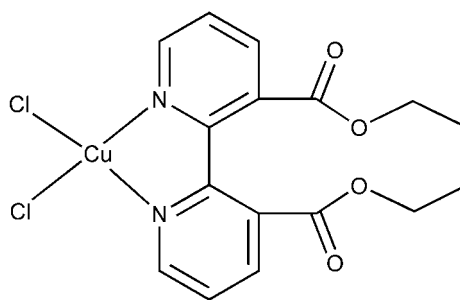
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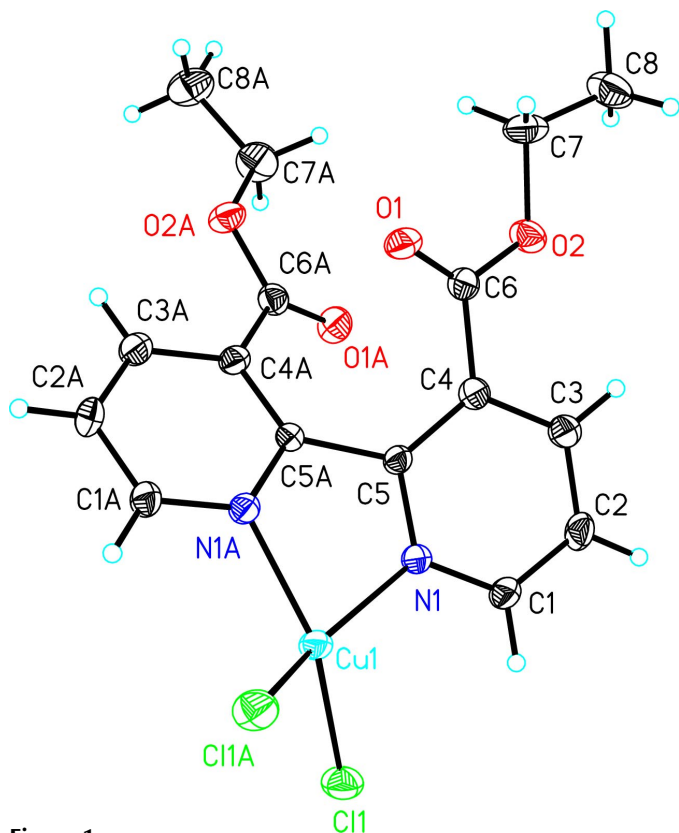
## Comment

Binicotinic acid, being a photoactive and multidentate ligand, has attracted much attention recently (Perkovic, 2000; Wu *et al.*, 2004). Moreover, the interesting emission properties of a platinum(II) complex with 3,3'-bis(methoxycarbonyl)-2,2'-bipyridyl has been reported by Miskowski *et al.* (1993). However, coordination compounds involving 3,3'-bis(ethoxycarbonyl)-2,2'-bipyridyl (DCEB) have not been exploited yet. We report here the first metal-DCEB coordination complex,  $[CuCl_2(DCEB)]$ , (I), prepared from the reaction of  $CuCl_2 \cdot 2H_2O$  and DCEB in  $CH_3CN$ .



(I)

X-ray diffraction analysis shows that (I) crystallizes in monoclinic space group  $C2/c$ . The local coordination geometry around each  $Cu^{II}$  center can be described as a severely distorted square (Fig. 1) in which each  $Cu^{II}$  center is coordinated by two N atoms from the 2,2'-pyridyl moiety of DCEB and two terminal chloride anions. The main deviation from the plane formed by atoms N1, Cl1, Cl1<sup>i</sup>, N1<sup>i</sup> (symmetry code as in Table 1) and Cu1 originates from its *trans* angles [146.67 (15)° for N1–Cu1–Cl1<sup>i</sup> and N1<sup>i</sup>–Cu1–Cl1], notably less than 180°, which leads to a coordination geometry close to that of a compressed tetrahedron. The dihedral angle between the two pyridine rings in (I) is 31.2 (5)°. The molecules are linked by weak  $C-H \cdots O$  and  $C-H \cdots Cl$  interactions to form a two-dimensional network which stacks along the  $c$  axis in the crystal (Fig. 2).



**Figure 1**  
Perspective view of (I) with the atom-numbering scheme. Displacement ellipsoids are shown at the 30% probability level. The suffix A corresponds to symmetry code (i) in Table 1.

## Experimental

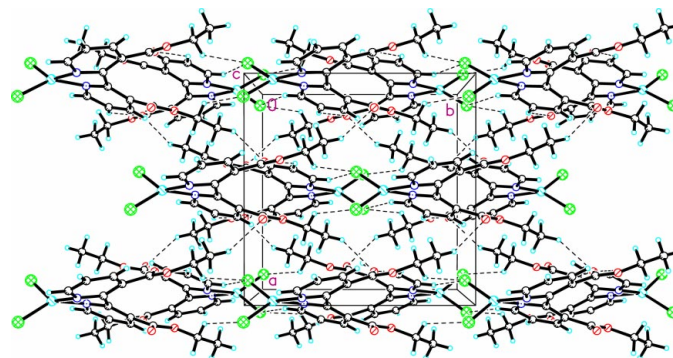
The organic ligand DCEB was prepared by substitution of methanol with ethanol in the literature method of Dholakia *et al.* (1985). A mixture of DCEB (30 mg, 0.1 mmol),  $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$  (17 mg, 0.1 mmol) and  $\text{CH}_3\text{CN}$  (5 ml) was vigorously stirred for about 1 h to give a green solution. Slow evaporation of the resultant solution for two weeks produced green plate-like crystals suitable for X-ray diffraction.

### Crystal data

|   |   |
|---|---|
| $[\text{CuCl}_2(\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_4)]$ | $D_x = 1.670 \text{ Mg m}^{-3}$           |
| $M_r = 434.75$  | Mo $K\alpha$ radiation                    |
| Monoclinic, $C2/c$  | Cell parameters from 2620 reflections     |
| $a = 11.3605 (16) \text{ \AA}$                                    | $\theta = 2.6\text{--}25.1^\circ$         |
| $b = 11.2772 (15) \text{ \AA}$                                    | $\mu = 1.60 \text{ mm}^{-1}$              |
| $c = 13.5039 (19) \text{ \AA}$                                    | $T = 293 (2) \text{ K}$                   |
| $\beta = 91.269 (3)^\circ$  | Plate, green                              |
| $V = 1729.6 (4) \text{ \AA}^3$                                    | $0.24 \times 0.18 \times 0.06 \text{ mm}$ |
| $Z = 4$   |   |

### Data collection

|   |  |
|---|--|
| Bruker SMART CCD diffractometer                             | 1523 independent reflections           |
| $\omega$ scans  | 1238 reflections with $I > 2\sigma(I)$ |
| Absorption correction: multi-scan (SADABS; Sheldrick, 1996) | $R_{\text{int}} = 0.035$               |
| $T_{\text{min}} = 0.673$ , $T_{\text{max}} = 0.909$         | $\theta_{\text{max}} = 25.1^\circ$     |
| 2620 measured reflections                                   | $h = -10 \rightarrow 13$               |
|   | $k = -13 \rightarrow 11$               |
|   | $l = -13 \rightarrow 16$               |



**Figure 2**  
Packing diagram of (I), viewed along the  $c$  axis. Hydrogen-bond interactions are indicated by dashed lines.

### Refinement

|                                 |  |
|---------------------------------|--|
| Refinement on $F^2$             | H-atom parameters constrained                        |
| $R[F^2 > 2\sigma(F^2)] = 0.068$ | $w = 1/[\sigma^2(F_o^2) + 18.8764P]$                 |
| $wR(F^2) = 0.138$               | where $P = (F_o^2 + 2F_c^2)/3$                       |
| $S = 1.23$                      | $(\Delta/\sigma)_{\text{max}} < 0.001$               |
| 1523 reflections                | $\Delta\rho_{\text{max}} = 0.51 \text{ e \AA}^{-3}$  |
| 115 parameters                  | $\Delta\rho_{\text{min}} = -0.48 \text{ e \AA}^{-3}$ |

**Table 1**

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

|                        |             |                          |             |
|------------------------|-------------|--------------------------|-------------|
| Cu1—N1                 | 2.009 (5)   | Cu1—Cl1                  | 2.2082 (18) |
| $\text{N1}^i$ —Cu1—N1  | 80.3 (3)    | N1—Cu1—Cl1               | 97.01 (15)  |
| $\text{N1}^i$ —Cu1—Cl1 | 146.67 (15) | Cl1—Cu1—Cl1 <sup>i</sup> | 102.45 (10) |

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

H-atoms were positioned geometrically ( $\text{C—H} = 0.95\text{--}0.99 \text{ \AA}$ ) and allowed to ride on their respective parent atoms, with  $U_{\text{iso}}(\text{H}) = 1.2$  or 1.5 times  $U_{\text{eq}}(\text{C})$ .

Data collection: SMART (Siemens, 1994); cell refinement: SAINT (Siemens, 1994); data reduction: SAINT and SHELXTL (Sheldrick, 1997); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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