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Jing Zhang, Ling Ye, Jing-Sheng Yu and Li-Xin Wu*

Key Laboratory for Supramolecular Structure and Materials, Jilin University, Changchun 130012, People's Republic of China

Correspondence e-mail: wulx@jlu.edu.cn

Key indicators

Single-crystal X-ray study T = 273 K Mean σ (C–C) = 0.006 Å R factor = 0.043 wR factor = 0.127 Data-to-parameter ratio = 17.4

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

4,4'-Ethylenebipyridinium tetrachlorocuprate

The crystal structure of the title compound, $(C_{12}H_{12}N_2)$ -[CuCl₄], consists of Cu^{II} complex anions and diprotonated bis(4-pyridyl)ethylene. The cations and anions lie across individual inversion centers. The square-planar Cu^{II} complex anions link with cations *via* N-H····Cl hydrogen bonding. Received 22 June 2005 Accepted 22 July 2005 Online 27 July 2005

Comment

As a part of our ongoing study on organic–inorganic hybrid crystal engineering involving transition metals with azoaromatic molecules, we report here the crystal structure of the title compound, (I).



The crystal structur of (I) (Fig. 1) consists of $[CuCl_4]^{2-}$ anions and diprotonated 1,2-bis(4-pyridyl)ethylene cations. The anion and cation lie across individual inversion centers. The Cu^{II} atom is coordinated by four Cl⁻ anions in a squareplanar geometry (Table 1). The bis(4-pyridyl)ethylene cations link with the Cu^{II} complex anions *via* N-H···Cl hydrogen bonding (Table 2). The centroid-to-centroid distance of 3.498 (2) Å suggests the existence of π - π stacking between neighboring parallel pyridyl rings (Fig. 2).



Figure 1

The molecular structure of (I), shown with 40% probability displacement ellipsoids (arbitrary spheres for H atoms). [Symmetry codes: (ii) 1 - x, 1 - y, 1 - z; (iii) -x, 1 - y, 2 - z.]

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Experimental

Equimolar amounts of bis(4-pyridyl)ethylene and copper acetate were mixed in ethanol, followed by addition of several drops of concentrated HCl solution. Blue single crystals of (I) were obtained from the solution after 5 d.

Z = 1

 $D_r = 1.801 \text{ Mg m}^{-3}$

Cell parameters from 1725

 $0.43 \times 0.24 \times 0.23~\text{mm}$

Mo $K\alpha$ radiation

reflections

 $\theta = 3.1 - 27.1^{\circ}$

 $\mu = 2.25~\mathrm{mm}^{-1}$

T = 273 (2) K

Block, blue

Crystal data

 $\begin{array}{l} ({\rm C}_{12}{\rm H}_{12}{\rm N}_2)[{\rm CuCl}_4] \\ M_r = 389.58 \\ {\rm Triclinic}, \ P\overline{1} \\ a = 7.0207 \ (14) \ {\rm \mathring{A}} \\ b = 7.0602 \ (11) \ {\rm \mathring{A}} \\ c = 8.1978 \ (19) \ {\rm \mathring{A}} \\ \alpha = 68.66 \ (1)^\circ \\ \beta = 73.748 \ (6)^\circ \\ \gamma = 76.204 \ (11)^\circ \\ V = 359.10 \ (12) \ {\rm \mathring{A}}^3 \end{array}$

Data collection

Rigaku R-AXIS RAPID	1547 independent reflections
diffractometer	1357 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.038$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.2^{\circ}$
(ABSCOR; Higashi, 1995)	$h = 0 \rightarrow 9$
$T_{\min} = 0.531, T_{\max} = 0.603$	$k = -8 \rightarrow 9$
3071 measured reflections	$l = -9 \rightarrow 10$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F^2) + (0.069P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.043$	+ 0.374P
$wR(F^2) = 0.127$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.14	$(\Delta/\sigma)_{\rm max} < 0.001$
1547 reflections	$\Delta \rho_{\rm max} = 0.56 \text{ e} \text{ Å}^{-3}$
89 parameters	$\Delta \rho_{\rm min} = -0.82 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXTL
-	Extinction coefficient: 0.086 (11)

Table 1

Selected geometric parameters (Å, °).

Cu1-Cl1	2.2489 (10)	Cu1-Cl2	2.2701 (10)
Cl1-Cu1-Cl2	89.90 (3)		

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$N1 - H1 \cdots Cl1^{iv}$ $N1 - H1 \cdots Cl2^{iv}$	0.86	2.48	3.174 (3) 3.187 (3)	138 142
	0.80	2.47	5.167 (5)	142

Symmetry code: (iv) x, y - 1, z.



Figure 2

A packing diagram showing π - π stacking and hydrogen bonding (dashed lines). [symmetry codes: (i) -x, -y, 2 - z; (ii) x, -2 + y, z; (iii) -x, 1 - y, 2 - z; (iv) x, -i + y, z; (v) 1 - x, -y, 1 - z; (vi) 1 + x, -2 + y, -1 + z; (vii) -1 + x, y, 1 + z; (viii) x, -2 + y, z; (ix) 1 - x, 1 - y, 1 - z; (x) 2 - x, -1 - y, -z; (xi) 1 + x, -1 + y, -1 + z.]

H atoms were placed in calculated positions, with C-H = 0.93 Å and N-H = 0.86 Å, and refined in riding mode, with $U_{iso}(H) = 1.2U_{eq}(carrier)$.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/ MSC, 2002); program(s) used to solve structure: *SHELXTL* (Bruker, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *DIAMOND* (Brandenburg, 1999) and *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXTL*.

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