

Yong-Heng Xing,\* Jing Han,  
Bao-Li Zhang, Xing-Jing Zhang,  
Yuan-Hong Zhang and  
Guang-Hua ZhouCollege of Chemistry and Chemical Engineering,  
Liaoning Normal University, Dalian 116029,  
People's Republic of ChinaCorrespondence e-mail:  
yhxing2000@yahoo.com

## Key indicators

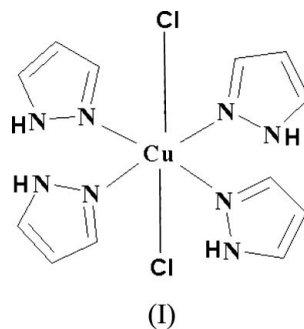
Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.007$  Å  
 $R$  factor = 0.049  
 $wR$  factor = 0.159  
Data-to-parameter ratio = 16.1For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Dichlorotetrakis(1*H*-pyrazole- $\kappa\text{N}^2$ )copper(II)

The reaction of  $\text{CuCl}_2$  and pyrazole (Hpz) in methanol solution at room temperature yields the title mononuclear complex,  $[\text{CuCl}_2(\text{C}_3\text{H}_4\text{N}_2)_4]$ . The Cu atom is six-coordinated by four N atoms from four pyrazole ligands and by two  $\text{Cl}^-$  ions. The overall symmetry of the complex is  $\bar{1}$ . The structure features  $\text{N}-\text{H}\cdots\text{Cl}$  and  $\text{C}-\text{H}\cdots\text{Cl}$  hydrogen bonds, yielding a three-dimensional network.

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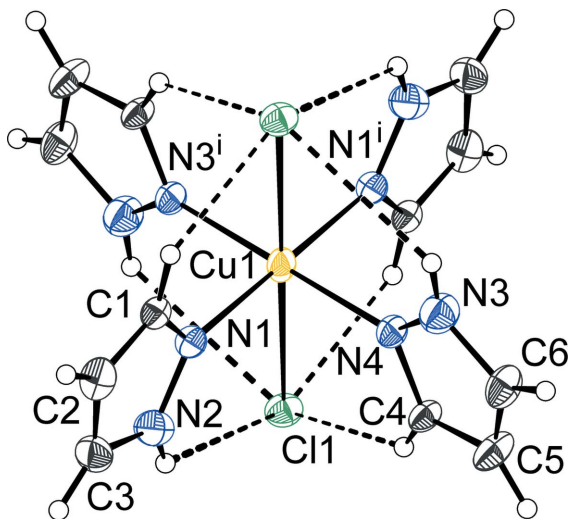
## Comment

Since Trofimenko discovered poly(pyrazolyl)borate in 1966 (Trofimenko, 1972), the coordination chemistry of pyrazole ligands has attracted much attention (Li *et al.*, 2004; Trofimenko, 1993; Mukherjee, 2000). The donor N atom of pyrazole can coordinate to metal atoms and the other N atom often forms hydrogen bonds with other atoms in the complexes, acting as either a hydrogen-bond donor or acceptor. Based on these findings, we attempted to determine the structure of the title complex,  $\text{Cu}(\text{pyrazole})_4\text{Cl}_2$ . (I).



The structure of (I) features a distorted octahedral complex of  $\text{Cu}^{2+}$ , which is coordinated in a square-planar fashion by four N atoms from four pyrazole ligands and two axial  $\text{Cl}^-$  ligands (Fig. 1). The Cu atom is located on a special position (site symmetry  $\bar{1}$ ). Selected bond lengths and angles are listed in Table 1.

The Cu—N distances of 2.001 (3) and 2.025 (3) Å are significantly longer than those in  $[\text{Cu}(\text{Hpz})_4\text{SO}_4](\text{H}_2\text{O})$  [mean 1.944 (6) Å; Shen *et al.*, 2004] or  $[\text{Cu}_3(\mu_3\text{-O})(\text{pz})_3(\text{Hpz})_3(\mu_1, \mu_2\text{-ClO}_4)](\text{CH}_3\text{OH})$  [mean 1.982 (5) Å; Shen *et al.*, 2004], but are close to the bond lengths found in  $[\text{Cu}(\text{Hpz})_4\text{SO}_4]_n(\text{CH}_3\text{OH})_n$  [mean 2.016 (3) Å; Shen *et al.*, 2004]. The  $\text{Cu}\cdots\text{Cl}$  distance of 2.839 (10) Å corresponds to a rather weak Cu—Cl bond. The *cis*-N—Cu—N angles deviate only slightly from  $90^\circ$  (Table 1). The  $\text{Cu1}-\text{N1}-\text{N2}-\text{C3}$  and  $\text{Cu1}-\text{N3}-\text{N4}-\text{C6}$  torsion angles are  $-179.2$  (3) and  $173.0$  (3) $^\circ$ , respectively.

**Figure 1**

The molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level. Unlabelled atoms and those labelled with a superscript *i* are related to labelled atoms by the symmetry code ( $\frac{1}{2} - x, \frac{1}{2} - y, 1 - z$ ). Hydrogen bonds are drawn as dashed lines.

Two kinds of hydrogen bond are present in the structure of (I) (Fig. 2 and Table 2). Firstly, there are hydrogen bonds between uncoordinated N atoms and the Cl atoms, and secondly there are hydrogen bonds between C atoms of the pyrazole ring and Cl atoms. Together, these hydrogen bonds yield a chain along the *ac* plane and the chains are further extended into a three-dimensional network in space.

## Experimental

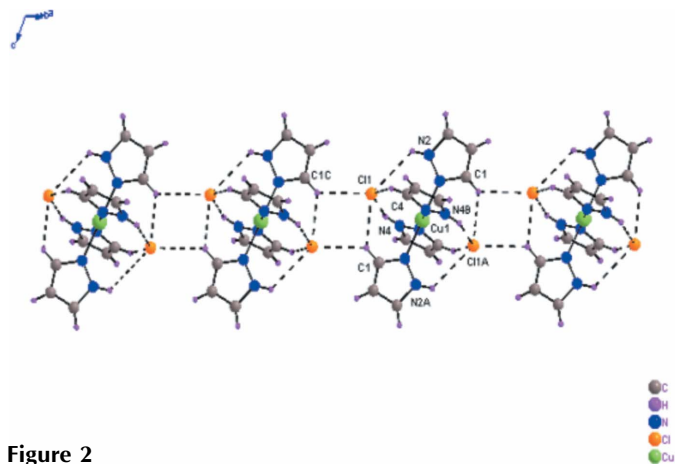
The starting materials were purchased from a commercial source (J & K CHEMICAL) and used without further purification. A methanol solution (10 ml) of  $\text{CuCl}_2$  (0.190 g) and pyrazole ( $\text{C}_3\text{H}_4\text{N}_2$ ) (0.536 g) was stirred for 4 h at room temperature, yielding a blue solution. This was set aside to crystallize, yielding analytically pure (I) as single crystals suitable for X-ray structure determination. Analysis, found: C 35.32, H 3.87, N 27.56%; calculated for  $\text{C}_{12}\text{H}_{16}\text{Cl}_2\text{Cu}_1\text{N}_8$ : C 35.40, H 3.93, N 27.53%.

### Crystal data

$[\text{CuCl}_2(\text{C}_3\text{H}_4\text{N}_2)_4]$	$Z = 4$
$M_r = 406.77$	$D_x = 1.637 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 13.674 (8) \text{ \AA}$	$\mu = 1.66 \text{ mm}^{-1}$
$b = 9.172 (5) \text{ \AA}$	$T = 293 (2) \text{ K}$
$c = 14.750 (8) \text{ \AA}$	Block, blue
$\beta = 116.852 (9)^\circ$	$0.23 \times 0.20 \times 0.15 \text{ mm}$
$V = 1650.5 (16) \text{ \AA}^3$	

### Data collection

Bruker APEXII diffractometer	4803 measured reflections
$\omega$ scans	1710 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	1603 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.690, T_{\max} = 0.847$	$R_{\text{int}} = 0.041$
(expected range = 0.635–0.780)	$\theta_{\max} = 26.5^\circ$

**Figure 2**

The two-dimensional hydrogen-bond structure along the *ac* plane in (I). [Symmetry codes: (A)  $-x + \frac{1}{2}, -y + \frac{1}{2}, -z + 1$ ; (B)  $x, y, z$ ; (C)  $x - \frac{1}{2}, y - \frac{1}{2}, z$ .]

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.049$   
 $wR(F^2) = 0.159$   
 $S = 1.17$   
 1710 reflections  
 106 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0886P)^2 + 4.2919P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.72 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.71 \text{ e \AA}^{-3}$

**Table 1**

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

Cu1—N3	2.001 (3)	Cu1—Cl1	2.8386 (15)
Cu1—N1	2.025 (3)		
N3 <sup>i</sup> —Cu1—N1	91.40 (12)	N3—Cu1—N1	88.60 (12)

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

**Table 2**

Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
C1—H1A $\cdots$ Cl1 <sup>i</sup>	0.93	2.53	3.211 (4)	130
C1—H1A $\cdots$ Cl1 <sup>ii</sup>	0.93	2.77	3.479 (5)	133
N2—H2A $\cdots$ Cl1	0.86	2.89	3.467 (5)	126
N2—H2A $\cdots$ Cl1 <sup>iii</sup>	0.86	2.95	3.646 (5)	139

Symmetry codes: (i)  $-x + \frac{1}{2}, -y + \frac{1}{2}, -z + 1$ ; (ii)  $x + \frac{1}{2}, y + \frac{1}{2}, z$ ; (iii)  $-x, y, -z + \frac{1}{2}$ .

H atoms were placed in calculated positions and constrained to ride on their parent C atoms, with  $C-H = 0.93 \text{ \AA}$  and  $N-H = 0.86 \text{ \AA}$ , and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{iso}}(\text{C}, \text{N})$ .

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: SHELXTL (Bruker, 1997); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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