



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.

Crystal data

(C<sub>12</sub>H<sub>12</sub>N<sub>2</sub>)[CuCl<sub>4</sub>]  
 M<sub>r</sub> = 389.58  
 Triclinic, P $\bar{1}$   
 a = 7.0207 (14) Å  
 b = 7.0602 (11) Å  
 c = 8.1978 (19) Å  
 α = 68.66 (1)°  
 β = 73.748 (6)°  
 γ = 76.204 (11)°  
 V = 359.10 (12) Å<sup>3</sup>

Z = 1  
 D<sub>x</sub> = 1.801 Mg m<sup>-3</sup>  
 Mo Kα radiation  
 Cell parameters from 1725 reflections  
 θ = 3.1–27.1°  
 μ = 2.25 mm<sup>-1</sup>  
 T = 273 (2) K  
 Block, blue  
 0.43 × 0.24 × 0.23 mm

Data collection

Rigaku R-AXIS RAPID diffractometer  
 ω scans  
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  
 T<sub>min</sub> = 0.531, T<sub>max</sub> = 0.603  
 3071 measured reflections

1547 independent reflections  
 1357 reflections with I > 2σ(I)  
 R<sub>int</sub> = 0.038  
 θ<sub>max</sub> = 27.2°  
 h = 0 → 9  
 k = -8 → 9  
 l = -9 → 10

Refinement

Refinement on F<sup>2</sup>  
 R[F<sup>2</sup> > 2σ(F<sup>2</sup>)] = 0.043  
 wR(F<sup>2</sup>) = 0.127  
 S = 1.14  
 1547 reflections  
 89 parameters  
 H-atom parameters constrained

w = 1/[σ<sup>2</sup>(F<sub>o</sub><sup>2</sup>) + (0.069P)<sup>2</sup> + 0.374P]  
 where P = (F<sub>o</sub><sup>2</sup> + 2F<sub>c</sub><sup>2</sup>)/3  
 (Δ/σ)<sub>max</sub> < 0.001  
 Δρ<sub>max</sub> = 0.56 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = -0.82 e Å<sup>-3</sup>  
 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...A	D—H	H...A	D...A	D—H...A
N1—H1...Cl1 <sup>iv</sup>	0.86	2.48	3.174 (3)	138
N1—H1...Cl2 <sup>iv</sup>	0.86	2.47	3.187 (3)	142

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

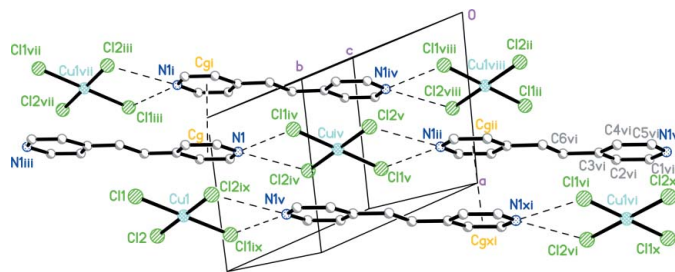


Figure 2 A packing diagram showing π-π stacking and hydrogen bonding (dashed lines). [symmetry codes: (i) -x, -y, 2 - z; (ii) x, -2 + y, z; (iii) -x, 1 - y, 2 - z; (iv) x, -i + y, z; (v) 1 - x, -y, 1 - z; (vi) 1 + x, -2 + y, -1 + z; (vii) -1 + x, y, 1 + z; (viii) x, -2 + y, z; (ix) 1 - x, 1 - y, 1 - 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<sub>iso</sub>(H) = 1.2U<sub>eq</sub>(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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References

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