

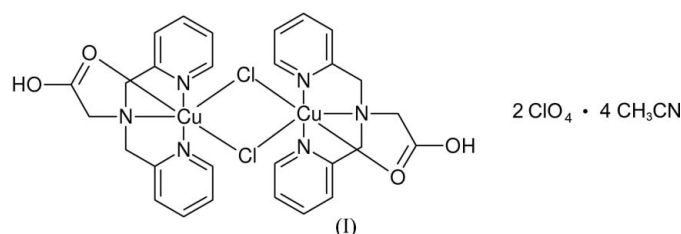
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Key indicators

Single-crystal X-ray study
 $T = 180$ K
Mean $\sigma(\text{C}-\text{C}) = 0.010$ Å
 R factor = 0.073
 wR factor = 0.191
Data-to-parameter ratio = 28.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Di- μ -chloro-bis{[*N,N*-bis(2-pyridylmethyl)-
glycine- κ^4N,N',N'',O]copper(II)} diperchlorate
acetonitrile tetrasolvateThe title compound, $[\text{Cu}_2\text{Cl}_2(\text{C}_{14}\text{H}_{15}\text{N}_3\text{O}_2)_2](\text{ClO}_4)_2 \cdot 4\text{C}_2\text{H}_3\text{N}$, contains dichloro-bridged dicopper(II) complexes lying on centres of inversion. The crystal examined was twinned by a 180° rotation about c^* .Received 23 August 2006
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Comment

Reaction of bis(2-pyridylmethyl)glycine (bpgH) and $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ in acidic methanol results in immediate precipitation of the title compound, (I). Compound (I) contains dichloro-bridged dicopper(II) complexes (Fig. 1) lying on centres of inversion, with the carboxylate groups protonated. The coordination geometry around the Cu^{II} atom (Table 1) is approximately octahedral, but exhibits significant elongation of the $\text{Cu1}-\text{O1}$ and $\text{Cu1}-\text{Cl1}^i$ bonds [symmetry code: (i) $1-x, 1-y, 1-z$] on account of the Jahn–Teller distortion associated with Cu^{II} . The protonated carboxyl groups form hydrogen bonds to perchlorate anions (Table 2).

Experimental

Bis(2-pyridylmethyl)glycine (0.1573 g, 0.611 mmol) and $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ (0.1047 g, 0.614 mmol) were dissolved in methanol (5 ml) and HClO_4 (0.25 ml of a 70% aqueous solution) was added. A turquoise powder (yield 0.2662 g, 93%) precipitated immediately. Recrystallization from hot acetonitrile afforded crystals of (I).

Crystal data

$[\text{Cu}_2\text{Cl}_2(\text{C}_{14}\text{H}_{15}\text{N}_3\text{O}_2)_2](\text{ClO}_4)_2 \cdot 4\text{C}_2\text{H}_3\text{N}$	$\gamma = 97.159$ (4) $^\circ$
$M_r = 1075.68$	$V = 1127.58$ (17) Å ³
Triclinic, $P\bar{1}$	$Z = 1$
$a = 7.1238$ (6) Å	$D_x = 1.584$ Mg m ⁻³
$b = 10.6609$ (10) Å	Mo $K\alpha$ radiation
$c = 14.9894$ (13) Å	$\mu = 1.25$ mm ⁻¹
$\alpha = 92.919$ (4) $^\circ$	$T = 180$ (2) K
$\beta = 91.266$ (4) $^\circ$	Plate, turquoise
	$0.24 \times 0.20 \times 0.05$ mm

Data collection

Bruker–Nonius X8APEX-II CCD diffractometer	26597 measured reflections
thin-slice ω and φ scans	8342 independent reflections
Absorption correction: multi-scan (<i>TWINABS</i> ; Sheldrick, 2004)	4142 reflections with $I > 2\sigma(I)$
$T_{\text{min}} = 0.768$, $T_{\text{max}} = 0.940$	$R_{\text{int}} = 0.076$
	$\theta_{\text{max}} = 26.6^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.073$
 $wR(F^2) = 0.191$
 $S = 1.01$
 8342 reflections
 289 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0828P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 1.11 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.65 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cu1—N1	1.970 (5)	Cu1—O1	2.316 (4)
Cu1—N2	2.076 (5)	Cu1—Cl1	2.2397 (17)
Cu1—N3	1.976 (5)	Cu1—Cl1 ⁱ	3.1116 (18)
N1—Cu1—N2	83.0 (2)	N3—Cu1—O1	91.58 (18)
N1—Cu1—N3	164.4 (2)	Cl1—Cu1—O1	100.18 (12)
N2—Cu1—N3	82.3 (2)	N1—Cu1—Cl1 ⁱ	84.87 (14)
N1—Cu1—Cl1	97.10 (15)	N2—Cu1—Cl1 ⁱ	84.60 (15)
N2—Cu1—Cl1	179.18 (17)	N3—Cu1—Cl1 ⁱ	88.40 (14)
N3—Cu1—Cl1	97.50 (15)	Cl1—Cu1—Cl1 ⁱ	94.61 (5)
N1—Cu1—O1	91.33 (17)	O1—Cu1—Cl1 ⁱ	165.08 (11)
N2—Cu1—O1	80.62 (18)		

Symmetry code: (i) $-x + 1, -y + 1, -z + 1$.

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O2-H2\cdots O2D^{\text{ii}}$	0.84	1.88	2.720 (7)	176

Symmetry code: (ii) $-x, -y, -z + 1$.

The crystal was twinned by 180° rotation about c^* . The diffraction pattern was indexed using *CELL_NOW* (Bruker–Nonius, 2004) and integrated as a two-component twin using *SAINT-Plus* (Bruker, 2003). 8445 data (2549 unique; $R_{\text{int}} = 0.077$) were associated with component 1 only, 8423 data (2539 unique; $R_{\text{int}} = 0.084$) were associated with component 2 only, and 9729 data (3277 unique; $R_{\text{int}} = 0.076$) were overlapped. Refinement was performed using the HKLF5 format in *SHELXTL* (Sheldrick, 2000). H atoms bound to C atoms were positioned geometrically and allowed to ride during subsequent refinement, with $C-H = 0.95 \text{ Å}$ for Csp^2 or 0.99 Å for the methylene groups, and with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$. The methyl groups of the acetonitrile molecules were positioned geometrically, with $C-H = 0.98 \text{ Å}$ and $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(C)$, and allowed to rotate about their local threefold axes. In the final cycles of refinement, rotation was not permitted to aid convergence. The H atom of the hydroxyl group was

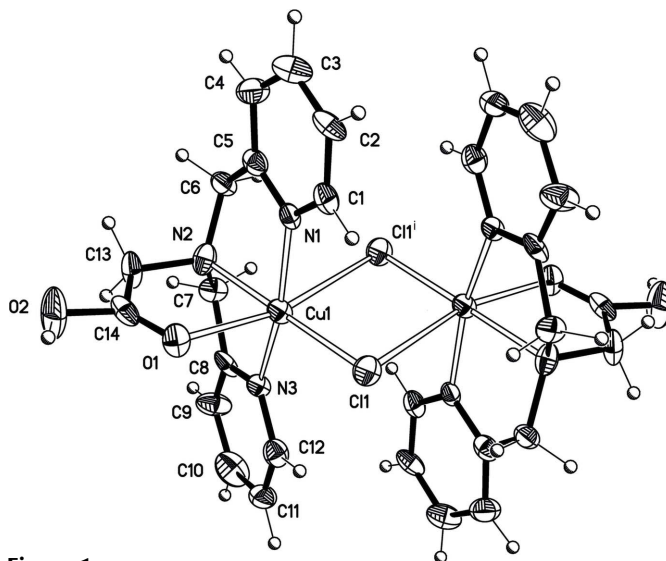


Figure 1

The dichloro-bridged dicopper(II) cationic complex in (I), showing displacement ellipsoids at the 50% probability level for non-H atoms. H atoms are shown as spheres of arbitrary radius. [Symmetry code (i) $1 - x, 1 - y, 1 - z$.]

placed in the plane of the carboxyl group, so as to form the best hydrogen bond (AFIX 83 in *SHELXTL*), with $O-H = 0.84 \text{ Å}$ and $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(O)$. The largest peak in the difference density lies in the vicinity of the perchlorate anion.

Data collection: *APEX2* (Bruker–Nonius, 2004); cell refinement: *SAINT-Plus* (Bruker, 2003); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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