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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.007 Å R factor = 0.043 wR factor = 0.103 Data-to-parameter ratio = 17.0

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µ-1,5-Dicyanamido-bis[chloro(1,4,7-triazacyclononane)copper(II)] perchlorate

In the title compound, $[Cu_2(C_2N_3)Cl_2(C_6H_{15}N_3)_2]ClO_4$, two $[CuCl_2(tacn)]$ units (tacn = 1,4,7-triazacyclononane) are linked by a single bridging dicyanamide ligand. The geometry about each Cu^{II} atom is deformed square-pyramidal or deformed trigonal-bipyramidal, formed by three N atoms of tacn, one of the terminal N atoms of the dicyanamide ligand and a Cl⁻ anion.

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Comment

Binuclear copper(II) complexes form an extremely active area of research in modern coordination chemistry (Sessler & Sibert, 1993; Edterurtas *et al.*, 1991) because of their crucial role in bioinorganic chemistry as model compounds for binuclear active sites in proteins (Collin *et al.*, 1988; Casellato *et al.*, 1979). Many model complexes containing various bridging ligands, such as oxalate (Zhang *et al.*, 2000), azide (Chaudhuri *et al.*, 1986), thiocyanate (Haasnoot *et al.*, 1984), terephthalate and tetracarboxylatobenzene (Chaudhuri *et al.*, 1988), and benzoate (Burger *et al.*, 1996), have been prepared and their structures characterized. During the course of our studies on binuclear copper(II) complexes, we obtained the title compound, (I). The synthesis and crystal structure of (I) are reported here.



The title compound consists of $[Cu_2\{N(CN)_2\}Cl_2(tacn)_2]^+$ cations and ClO_4^- anions (Fig. 1). Atom Cu1 is situated in a distorted square-pyramidal geometry, with atom N2 of tacn at the apical position. The Cu1–N2 bond distance of 2.272 (3) Å is longer than the other Cu1–N bonds of tacn [2.011 (3)–2.068 (3) Å; Table 1]. Atom Cu2 shows a distorted trigonal bipyramidal coordination geometry, with atoms N6 and N9 at the axial positions.

The N(CN)₂⁻ anion links two [Cu(tacn)Cl₂] units in a 1,5bridging mode. The C7-N5-C8 bond angle is 120.2 (4)°, which is typical for the dicyanamide ligand in a 1,5-bridging mode (Vangdal *et al.*, 2002; Carranza *et al.*, 2002).

Experimental

A solution of CuCl₂·2H₂O (0.5 mmol) in methanol (5 ml) was added to a solution of tacn (0.5 mmol) in methanol (15 ml), affording a clear deep-blue solution. The reaction mixture was stirred for 30 min at room temperature, then a solution of NaN(CN)₂ (0.045 g, 0.5 mmol) in water (5 ml) was slowly added dropwise. The deep-blue mixture was stirred for another 2 h, then filtered to remove any insoluble particles. Finally, NaClO₄ (*ca* 0.1 g) was added to the filtrate. After a month, crystals of (I) suitable for X-ray study were obtained by slow evaporation of the filtrate at room temperature (yield 65%).

Crystal data

[Cu₂(C₂N₃)Cl₂(C₆H₁₅N₃)₂]ClO₄ $D_x = 1.718 \text{ Mg m}^{-3}$ $M_r = 621.90$ Mo $K\alpha$ radiation Monoclinic, $P2_1/n$ Cell parameters from 864 a = 17.882 (6) Å reflections b = 7.494(3) Å $\theta = 2.3 - 26.0^{\circ}$ $\mu = 2.14 \text{ mm}^{-1}$ c = 18.596 (6) Å $\beta = 105.200(5)^{\circ}$ T = 293 (2) K $V = 2404.9 (14) \text{ Å}^3$ Block, blue $0.24 \times 0.18 \times 0.12 \text{ mm}$ Z = 4

4901 independent reflections

 $R_{\rm int}=0.047$

 $\theta_{\rm max} = 26.4^\circ$

 $h = -22 \rightarrow 22$

 $k = -6 \rightarrow 9$

 $l = -23 \rightarrow 23$

3372 reflections with $I > 2\sigma(I)$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{min} = 0.592, T_{max} = 0.773$ 13116 measured reflections

Refinement

 $\begin{array}{ll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.0444P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.043 & w \mbox{ere } P = (F_o^2 + 2F_c^2)/3 \\ w \mbox{Re}(F^2) = 0.103 & w \mbox{here } P = (F_o^2 + 2F_c^2)/3 \\ S = 1.05 & (\Delta/\sigma)_{\rm max} = 0.001 \\ 4901 \mbox{ reflections } & \Delta\rho_{\rm max} = 0.59 \mbox{ e } {\rm \AA}^{-3} \\ 289 \mbox{ parameters } & \Delta\rho_{\rm min} = -0.42 \mbox{ e } {\rm \AA}^{-3} \\ \mbox{H-atom parameters constrained } \end{array}$

Table 1

Selected geometric parameters (Å, °).

Cu1-N4	1.969 (4)	Cu2-N6	1.979 (4)
Cu1-N1	2.011 (3)	Cu2-N9	2.011 (3)
Cu1-N3	2.068 (3)	Cu2-N8	2.114 (3)
Cu1-N2	2.272 (3)	Cu2-N7	2.196 (4)
Cu1-Cl1	2.2909 (12)	Cu2-Cl2	2.2833 (12)
N4-Cu1-N1	168.44 (15)	N6-Cu2-N9	174.88 (14)
N4-Cu1-N3	90.63 (14)	N6-Cu2-N8	92.69 (14)
N1-Cu1-N3	82.20 (14)	N9-Cu2-N8	83.21 (12)
N4-Cu1-N2	106.10 (14)	N6-Cu2-N7	94.31 (16)
N1-Cu1-N2	81.89 (13)	N9-Cu2-N7	82.09 (13)
N3-Cu1-N2	81.48 (13)	N8-Cu2-N7	81.71 (13)
N4-Cu1-Cl1	93.88 (11)	N6-Cu2-Cl2	93.84 (11)
N1-Cu1-Cl1	91.37 (10)	N9-Cu2-Cl2	91.26 (9)
N3-Cu1-Cl1	167.41 (11)	N8-Cu2-Cl2	147.77 (10)
N2-Cu1-Cl1	108.43 (9)	N7-Cu2-Cl2	129.09 (10)



A perspective view of the complex cation in (I), shown with 30% probability displacement ellipsoids. H atoms have been omitted for clarity.

The C-bound H atoms were included in the riding-model approximation, with C-H distances of 0.97 Å and with $U_{\rm iso}$ (methylene H) = $1.5U_{\rm eq}$ (C).

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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