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Key indicators

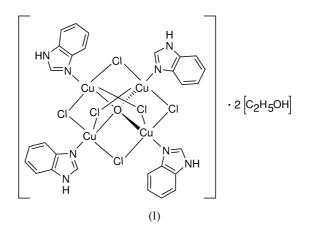
Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.012 Å H-atom completeness 84% Disorder in solvent or counterion R factor = 0.051 wR factor = 0.185 Data-to-parameter ratio = 20.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. μ_4 -Oxo-hexa- μ_2 -chloro-tetrakis[(benzimidazole- κN)copper(II)] ethanol disolvate: a reformulation

The title compound $[Cu_4Cl_6O(C_7H_6N_2)_4]\cdot 2C_2H_5OH$, consists of the tetranuclear copper(II) complex and uncoordinated ethanol molecules. The structure was reported previously by Tosik, Bukowska-Strzyzewska & Mrozinski [*J. Coord. Chem.* (1991), **24**, 113–120], but only 0.68 ethanol molecules were found in the asymmetric unit. The present redetermination indicates that two ethanol molecules occur in the asymmetric unit. One ethanol molecule is ordered while the other one is disordered, with the hydroxyl group in two alternative sites. Both disordered hydroxyl groups hydrogen bond to the ordered ethanol molecule with $O \cdots O$ separations of 2.845 (16) and 2.67 (2) Å, and $O-H \cdots O$ angles of 156 and 147°, respectively.

Comment

Crystals of the title compound, (I), were occasionally obtained as a by-product in the preparation of a complex of (4hydroxybenzoato)copper(II) with benzimidazole.



The molecular structure of (I) is illustrated in Fig. 1. There are one tetranuclear copper(II) complex and two ethanol molecules in the asymmetric unit. The structure of the copper(II) complex is the same as that reported previously (Tosik *et al.*, 1991). One ordered ethanol molecule, which was ignored in the previous determination, hydrogen bonds to a benzimidazole of the adjacent copper(II) complex (Fig. 1 and Table 2). The other ethanol molecule is disordered in the structure, the hydroxyl group located in different sites with site-occupancy factors of 0.68 (O3*a*) and 0.32 (O3*b*), respectively. Atom O3*b* was ignored in the previous determination. As shown in Fig. 1, both O3*a* and O3*b* hydrogen bond to the ordered ethanol, with reasonable O···O separations and O– $H \cdots O$ angles (Table 2). The H atoms on the disordered

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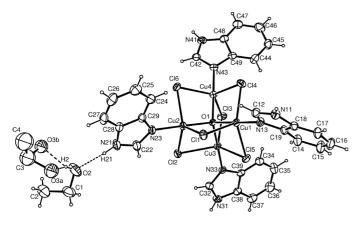


Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level. Dashed lines indicate hydrogen bonds. The suffixes a and b indicate the disordered hydroxyl groups.

ethanol molecule were not located in the structure, but the separation of 2.60 (3) Å suggests the existence of intermolecular hydrogen bonding between atoms O3b and O3b(2 - x, 2 - y, 2 - z).

Experimental

Sodium 3-hydroxybenzoate (0.16 g 1 mmol), benzimidazole (0.12 g, 1 mmol) and CuCl₂·2H₂O (0.17 g, 1 mmol) were dissolved in absolute ethanol (15 ml). The solution was refluxed for 5 h and a red precipitate appeared. After cooling to room temperature, the solution was filtered. Brown crystals were obtained from the filtrate after 5 d.

Crystal data

$[Cu_4Cl_6O(C_7H_6N_2)_4]\cdot 2C_2H_6O$	Z = 2
$M_r = 1047.55$	$D_x = 1.681 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 13.0005 (14) Å	Cell parameters from 10395
b = 13.2362 (15) Å	reflections
c = 13.8867 (9) Å	$\theta = 3.0-26.0^{\circ}$
$\alpha = 68.062 (2)^{\circ}$	$\mu = 2.46 \text{ mm}^{-1}$
$\beta = 74.315 \ (2)^{\circ}$	T = 298 (2) K
$\gamma = 71.376 \ (2)^{\circ}$	Prism, brown
V = 2069.2 (4) Å ³	$0.41\times0.21\times0.11~\rm{mm}$
Data collection	
Rigaku R-AXIS-RAPID	9365 independent reflections
diffractometer	6936 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.034$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(ABSCOR; Higashi, 1995)	$h = -16 \rightarrow 16$
$T_{\rm min} = 0.358, T_{\rm max} = 0.758$	$k = -17 \rightarrow 17$
19494 measured reflections	$l = -18 \rightarrow 16$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.051$ wR(F²) = 0.185 S = 0.869365 reflections 467 parameters H-atom parameters constrained I)

 $w = 1/[\sigma^2(F_o^2) + (0.1372P)^2]$ + 6.2288Pwhere $P = (F_o^2)^2$ $(\Delta/\sigma)_{\rm max} = 0.002$ $\Delta \rho_{\rm max} = 1.01 \text{ e} \text{ Å}$ $\Delta \rho_{\rm min} = -0.89 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1		
Selected	geometric parameters	(Å).

Cu1-O1	1.905 (3)	Cu3-O1	1.908 (3)
Cu1-N13	1.955 (4)	Cu3-N33	1.947 (4)
Cu1-Cl4	2.3538 (15)	Cu3-Cl5	2.3833 (15)
Cu1-Cl5	2.4068 (15)	Cu3-Cl3	2.4331 (15)
Cu1-Cl1	2.5095 (14)	Cu3-Cl2	2.4436 (15)
Cu2-O1	1.923 (3)	Cu4-O1	1.916 (3)
Cu2-N23	1.957 (4)	Cu4-N43	1.948 (4)
Cu2-Cl1	2.3958 (15)	Cu4-Cl3	2.3629 (15)
Cu2-Cl6	2.4435 (16)	Cu4-Cl6	2.4231 (14)
Cu2-Cl2	2.4571 (15)	Cu4-Cl4	2.4596 (15)

Table 2		
Hydrogen-bonding geometry	(Å,	°).

$D - H \cdot \cdot \cdot A$	$D-{\rm H}$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
O2−H2···O3a	0.87	2.03	2.845 (16)	156
$O2-H2 \cdot \cdot \cdot O3b$	0.87	1.88	2.67 (2)	147
$N11 - H11 \cdot \cdot \cdot Cl6^{i}$	0.86	2.70	3.445 (6)	146
N21-H21···O2	0.86	1.96	2.761 (10)	155
N31-H31···Cl1 ⁱⁱ	0.86	2.59	3.372 (5)	152
$N41 - H41 \cdots Cl2^{iii}$	0.86	2.69	3.408 (5)	141

Symmetry codes: (i) 2 - x, 1 - y, 1 - z; (ii) 1 - x, 1 - y, 2 - z; (iii) 1 - x, 2 - y, 1 - z.

The disordered ethanol molecule was refined isotropically, and its H atoms were not located. The hydroxyl H atom of the ordered ethanol was placed in a calculated position (Nardelli, 1999), and included in the final cycles of refinement with fixed coordinates and an isotropic displacement parameter of 0.10 Å². Other H atoms were placed in calculated positions, with C-H = 0.93-0.97 Å and N-H =0.86 Å, and included in the final cycles of refinement in a riding model, with $U_{iso}(H) = 1.2U_{eq}$ or $1.5U_{eq}$ of the carrier atoms. The peak with the maximum electron density in the difference Fourier map was close (0.87 Å) to atom C3.

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/ MSC, 2002); program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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References

Altomare, A., Cascarano, G., Giacovazzo, C. & Guagliardi, A. (1993). J. Appl. Cryst. 26, 343-350.

Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.

Farrugia, L. J. (1999). J. Appl. Cryst. 32, 837-838.

Higashi, T. (1995). ABSCOR. Rigaku Corporation, Tokyo, Japan.

- Nardelli, M. (1999). J. Appl. Cryst. 32, 563-571.
- Rigaku (1998). PROCESS-AUTO. Rigaku Corporation, Tokyo, Japan.
- Rigaku/MSC (2002). CrystalStructure. Version 3.00. Rigaku/MSC, 9009 New Trails Drive, The Woodlands, TX 77381-5209, USA.
- Sheldrick, G. M. (1997). SHELXL97. University of Göttingen, Germany.
- Tosik, A., Bukowska-Strzyzewska, M. & Mrozinski, J. (1991). J. Coord. Chem. 24, 113-120.