metal-organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Cai-Feng Ding,^a Xue-Mei Li,^a Mei Zhu,^a Hong Xu^b and Shu-Sheng Zhang^a*

^aCollege of Chemistry and Molecular Engineering, Qingdao University of Science and Technology, 266042 Qingdao, Shandong, People's Republic of China, and ^bCollege of Life Science and Pharmaceutical Engineering, Nanjing University of Technology, 210093 Nanjing, Jiangsu, People's Republic of China

Correspondence e-mail: shushzhang@126.com

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.040 wR factor = 0.107 Data-to-parameter ratio = 15.5

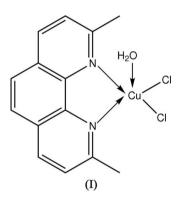
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Aquadichloro(2,9-dimethyl-1,10-phenanthroline- $\kappa^2 N, N$)copper(II)

In the title compound, $[CuCl_2(C_{14}H_{12}N_2)(H_2O)]$, there are two molecules in the asymmetric unit; both Cu atoms are fivecoordinate with a distorted trigonal-bipyramidal geometry. Molecules are linked into a three-dimensional network by O– $H \cdots Cl$ and $C-H \cdots Cl$ hydrogen bonds. The packing is stabilized by $\pi-\pi$ interactions between the phenanthroline ring systems.

Comment

In our ongoing studies of adipate–metal complexes, the title compound, (I), was obtained unexpectedly from the reaction of 2,9-dimethyl-1,10-phenanthroline and CuCl₂.



There are two molecules in the asymmetric unit; the chemically analogous bond lengths and angles of these two molecules are similar (Table 1). However, the molecules differ in the orientation of the phenanthroline ligands with respect to the Cu atoms (Fig. 1). Each Cu^{II} atom is five-coordinated by two N atoms from a 9,10-dimethylphenanthroline ligand, the O atom from a water molecule and two Cl atoms. The CuON₂Cl₂ unit forms a distorted trigonal-bipyramidal geometry, with one O atom and one N atom occupying the axial positions [O1-Cu1-N1 = 175.0 (1)° and O2-Cu2-N4 = 172.7 (2)°].

In the crystal structure, molecules are linked into a threedimensional network by O-H···Cl and C-H···Cl hydrogen bonds (Table 2 and Fig. 2). The short $Cg7 \cdots Cg7^{vi}$ distance of 3.551 Å [Cg7 is the centroid of the C5-C8/C12/C13 ring; symmetry code: (vi) 2 - x, 2 - y, -z] indicates π - π stacking interactions between the phenanthroline ring systems, which contribute to the crystal packing.

Experimental

© 2006 International Union of Crystallography All rights reserved To a solution of 2,9-dimethyl-1,10-phenanthroline (0.21 g, 1 mmol) and adipic acid (0.73 g, 0.5 mmol) in ethanol (10 ml) was added a

Received 5 January 2006 Accepted 20 February 2006 solution of CuCl₂ (0.14 g, 1 mmol) in distilled water (10 ml). The mixture was stirred and refluxed for 5 h. A hot solution was filtered into another flask containing a mixture of ethanol–methanol (1:2 ν/ν). Green crystals appeared over a period of a week by slow evaporation at room temperature.

Z = 4

 $D_{\rm r} = 1.578 {\rm Mg} {\rm m}^{-3}$

Cell parameters from 3160

 $0.22 \times 0.12 \times 0.07 \ \text{mm}$

5846 independent reflections 4664 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

reflections

 $\theta = 2.4 - 26.0^{\circ}$

 $\mu = 1.79~\mathrm{mm}^{-1}$

T = 293 (2) K

Plate, green

 $R_{\rm int} = 0.016$

 $\theta_{\rm max} = 26.0^{\circ}$

 $\begin{array}{l} h = -9 \rightarrow 9 \\ k = -9 \rightarrow 13 \end{array}$

 $l = -23 \rightarrow 19$

Crystal data

 $\begin{bmatrix} \text{CuCl}_2(\text{C}_{14}\text{H}_{12}\text{N}_2)(\text{H}_2\text{O}) \end{bmatrix} \\ M_r = 360.71 \\ \text{Triclinic, } P\overline{1} \\ a = 7.5047 \ (6) \text{ Å} \\ b = 11.3643 \ (9) \text{ Å} \\ c = 19.3670 \ (15) \text{ Å} \\ a = 100.981 \ (1)^{\circ} \\ \beta = 100.913 \ (1)^{\circ} \\ \gamma = 104.753 \ (1)^{\circ} \\ V = 1518.0 \ (2) \text{ Å}^{3} \end{bmatrix}$

Data collection

Siemens SMART 1000 CCD areadetector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.695, T_{\max} = 0.885$ 8719 measured reflections

Refinement

 Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0557P)^2]$
 $R[F^2 > 2\sigma(F^2)] = 0.040$ $w = 1/[\sigma^2(F_o^2) + (0.0557P)^2]$
 $wR(F^2) = 0.107$ where $P = (F_o^2 + 2F_c^2)/3$

 S = 1.02 $(\Delta/\sigma)_{max} = 0.001$

 5846 reflections
 $\Delta\rho_{max} = 0.48 \text{ e Å}^{-3}$

 377 parameters
 $\Delta\rho_{min} = -0.38 \text{ e Å}^{-3}$

 H atoms treated by a mixture of independent and constrained refinement
 $e^{A^{-3}}$

Table 1

Selected bond lengths (Å).

Cu1-O1	1.960 (3)	Cu2-O2	1.942 (3)
Cu1-N1	1.997 (3)	Cu2-N4	2.000(2)
Cu1-N2	2.227 (3)	Cu2-N3	2.241 (3)
Cu1-Cl2	2.3074 (10)	Cu2-Cl4	2.3109 (10)
Cu1-Cl1	2.3144 (10)	Cu2-Cl3	2.3194 (9)

Tab	le	2
-----	----	---

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
01-H101Cl3	0.84 (6)	2.31 (6)	3.121 (4)	164 (5)
$O2-H1O2 \cdot \cdot \cdot Cl2$	0.63 (5)	2.49 (5)	3.096 (5)	163 (5)
$O1-H2O1\cdots Cl4^{i}$	0.73 (6)	2.46 (6)	3.149 (4)	159 (6)
O2−H2O2···Cl1 ⁱⁱ	0.78 (6)	2.27 (6)	3.039 (5)	166 (5)
C4-H4A···Cl1 ⁱⁱⁱ	0.93	2.80	3.601 (5)	145
C10−H10A···Cl1 ^{iv}	0.93	2.80	3.624 (6)	149
$C23-H23A\cdots Cl3^{v}$	0.93	2.79	3.511 (4)	135

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

C-bound H atoms were refined using a riding model, with C–H = 0.93–0.96 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$ or $1.5U_{\rm eq}({\rm methyl}\ {\rm C})$. The H atoms of the water molecules were located in a difference Fourier

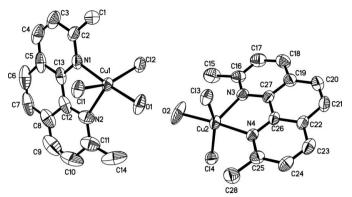


Figure 1

The asymmetric unit of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. H atoms have been omitted for clarity.

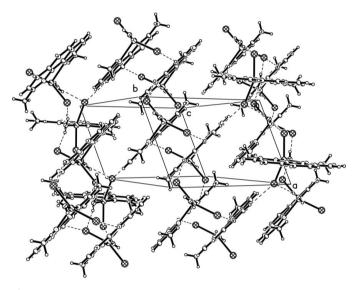


Figure 2

The molecular packing, showing the π - π interactions. Hydrogen bonds are indicated by dashed lines.

map and refined isotropically; the resulting distances are given in Table 2.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

This project was supported by the Project of Educational Administration of Shandong Province (No. J04B12) and the Outstanding Adult–Young Scientific Research Encouraging Foundation of Shandong Province (No. 2005BS04007).

References

- Nardelli, M. (1995). J. Appl. Cryst. 28, 659.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
- Sheldrick, G. M. (1997). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.
- Siemens (1996). *SMART* and *SAINT*. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.

Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.