Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.004 \text{ Å}$ Disorder in main residue R factor = 0.042 wR factor = 0.111 Data-to-parameter ratio = 15.1

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

$(3,5-Dinitrobenzoato-\kappa O)bis[(2-pyridyl)$ $methanol-<math>\kappa^2 N,O$]copper(II) 3,5-dinitrobenzoate

In the title compound, $[Cu(C_7H_3N_2O_6)(C_6H_7NO)_2](C_7H_3-N_2O_6)$, the Cu^{II} atom is coordinated by two chelating 2-pyridylmethanol ligands and a monodentate 3,5-dinitrobenzoate anion, forming a square-pyramidal coordination polyhedron. The cation and neighbouring anion are connected by two very strong $O-H \cdots O$ hydrogen bonds to form an ion pair.

Received 3 November 2005 Accepted 9 December 2005 Online 14 December 2005

Comment

As a part of our systematic study of copper(II) carboxylate complexes with biologically interesting molecular ligands, we report here the crystal structure of the title compound, (I).



The asymmetric unit of (I) is made up of a $[Cu(C_6H_7NO)_2(C_7H_3N_2O_6)]^+$ cation and a $C_7H_3N_2O_6^-$ anion. The (2-pyridyl)methanol ligand (also known as 2-pyridylcarbinol and abbreviated as 2-pycarb) prefers a chelating mode, in contrast with the bridging mode found for (3-pyridyl)methanol (known as ronicol) in copper(II) complexes (Stachová *et al.*, 2005). The 3,5-dinitrobenzoate ligand is monodentate, coordinating through a carboxylate O atom to give a complex cation with distorted square-pyramidal geometry and a CuO_2N_2O' coordination polyhedron. The basal plane is formed by carboxylate atom O1 from the coordinating 3,5-dinitrobenzoate ligand, atom O31 from one



© 2006 International Union of Crystallography Printed in Great Britain – all rights reserved The asymmetric unit of (I). Displacement ellipsoids are drawn at the 40% probability level. Dotted lines represent $O-H \cdots O$ hydrogen bonds.

2-pycarb ligand and two mutually trans pyridine atoms, N5 and N6, from two 2-pycarb ligands. The apical position of the coordination polyhedron is occupied by atom O41 of a 2pycarb ligand, with a Cu–O distance longer than those in the basal plane (Table 1). The Cu1–O2 distance of 2.735 (2) Å is longer than the Cu1-O1 distance. Also, the C11-O2 distance is significantly shorter than the C11-O1 distance. These confirm that atom O2 is not involved in coordination and the ligand is monodentate. Atom Cu1 is displaced out of the basal plane by 0.120(1) Å in the direction of the apical O41 atom. The 3.5-dinitrobenzoate anion is uncoordinated. With the exception of the C-O distances in the carboxylate groups, there are no significant differences in bond lengths or angles between the coordinated and uncoordinated anions and the values are comparable with those observed for bis(1,3diaminopropane)-3,5-dinitrobenzoatocopper(II) 3,5-dinitrobenzoate (Sundberg et al., 1994).

Two very strong hydrogen bonds link atoms O7 and O8 of the uncoordinated 3,5-dinitrobenzoate anion to hydroxyl atoms H41 and H31 of the coordinated 2-pycarb ligands (Table 2), giving rise to ion-pair formation. Similar hydrogen bonding is reported for bis(triethanolamine)copper(II) diacetate (Krabbes *et al.*, 1999). In the closely related benzoatobis(dimethylaminoetanol)copper(II) benzoate (Turpeinen *et al.*, 1985), the carboxylate anions form hydrogen bonds with hydroxyl H atoms of the coordinated ligands. Similar but weaker hydrogen bonds are found in copper(II) complexes containing coordinated NH groups (Sen *et al.*, 2000, Sundberg & Klinga, 1994).

Experimental

(2-Pyridyl)methanol (2 mmol) was added to copper(II) acetate (1 mmol) in aqueous solution (20 ml). 3,5-Dinitrobenzoic acid (2 mmol) was then added. The powdery blue product was filtered off and dried at room temperature. Blue prismatic crystals of (I) suitable for X-ray analysis were obtained from the mother liquor after slow room-temperature crystallization over a period of a few weeks.

Crystal data

$[Cu(C_7H_3N_2O_6)(C_6H_7NO)_2]$ -	Z
$(C_7H_3N_2O_6)$	Ľ
$M_r = 704.02$	N
Triclinic, P1	C
a = 9.870 (1) Å	
b = 12.181(1) Å	θ
c = 12.732 (1) Å	μ
$\alpha = 95.29 \ (1)^{\circ}$	T
$\beta = 108.68 \ (1)^{\circ}$	Р
$\gamma = 91.80 \ (1)^{\circ}$	0.
V = 1440.9 (2) Å ³	
Data collection	
Siemens P4 diffractometer	R
φ scans	θ_1

Absorption correction: ψ scan (*XEMP*; Siemens, 1990) $T_{\min} = 0.641$, $T_{\max} = 0.777$ 7697 measured reflections 6556 independent reflections 5266 reflections with $I > 2\sigma(I)$ Z = 2 $D_x = 1.623 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 25 reflections $\theta = 4.2-12.0^{\circ}$ $\mu = 0.84 \text{ mm}^{-1}$ T = 293 (2) K Prism, blue $0.55 \times 0.45 \times 0.30 \text{ mm}$

 $R_{int} = 0.029$ $\theta_{max} = 27.5^{\circ}$ $h = -1 \rightarrow 12$ $k = -15 \rightarrow 15$ $l = -16 \rightarrow 16$ 2 standard reflections every 100 reflections intensity decay: none Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0601P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 0.292P]
$wR(F^2) = 0.111$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.001$
6556 reflections	$\Delta \rho_{\rm max} = 0.52 \text{ e} \text{ \AA}^{-3}$
433 parameters	$\Delta \rho_{\rm min} = -0.63 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

Cu1-O1	1.967 (2)	Cu1-O41	2.301 (2)
Cu1-N5	1.972 (2)	O1-C11	1.274 (3)
Cu1-N6	1.993 (2)	O2-C11	1.224 (3)
Cu1-O31	2.000 (2)		
O1-Cu1-N5	92.43 (7)	N6-Cu1-O31	93.16 (7)
O1-Cu1-N6	93.46 (7)	O1-Cu1-O41	96.50 (7)
N5-Cu1-N6	173.81 (7)	N5-Cu1-O41	99.99 (7)
O1-Cu1-O31	162.24 (7)	N6-Cu1-O41	77.45 (8)
N5-Cu1-O31	81.75 (7)	O31-Cu1-O41	101.02 (6)

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

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
O31−H31···O8	0.87	1.65	2.512 (2)	173
O41−H41···O7	0.93	1.71	2.631 (3)	174

Atoms H31 and H41 were located in a difference Fourier map and their positions were not refined. The remaining H atoms were placed in geometrically calculated positions and allowed to ride on their parent atoms, with C–H distances of 0.93 (aromatic) or 0.97 Å (methylene). The $U_{\rm iso}$ value for all H atoms was fixed at 0.05 Å². Atom O11 of an NO₂ group is disordered over two positions, O11A and O11B. The occupancy factors were initially refined to 0.502 (2) and 0.498 (2), but later fixed at 0.5 each.

Data collection, cell refinement and data reduction: *XSCANS* (Siemens, 1994); program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); publication material: *WinGX* (Farrugia, 1999).

This work was supported by VEGA grant 1/2452/05.

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