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Key indicators

Single-crystal X-ray study T = 153 KMean $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$ R factor = 0.029 wR factor = 0.060 Data-to-parameter ratio = 11.7

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. A macrocyclic binuclear copper(II) complex of the ligand *N*,*N*'-bis(pyridin-2-ylmethyl)pyrazine-2,3-dicarboxamide

In the complex, $bis[\mu-N,N'-bis(pyridin-2-ylmethyl)pyrazine-2,3-dicarboxamide]bis[sulfatocopper(II)] hexahydrate,$ [Cu₂(C₁₈H₁₆N₆O₂)₂(SO₄)₂]·6H₂O, the pyrazine ligand coordinates in a tridentate-monodentate manner with the Cu atomshaving N₃O₂ chromophores. A neutral centrosymmetricbinuclear macrocyclic structure is formed, with the Cu atomshaving almost perfect square-pyramidal geometry. Received 5 January 2004 Accepted 7 January 2004 Online 17 January 2004

Comment

Structures of both copper(II) (Hausmann et al., 2003; Cati et al., 2004) and nickel(II) (Cati et al., 2004) complexes of the *N*,*N*'-bis(pyridin-2-ylmethyl)pyrazine-2,3-dicarboxligand. amide, (H₂L), including $[2 \times 2]$ grids, have been described recently. Cati et al. (2004) also described the synthesis of H₂L and the structure of its triclinic polymorph. The structure of the orthorhombic polymorph of H_2L has also been described recently (Cati & Stoeckli-Evans, 2004a). Hausmann et al. (2003) also reported the structure of a dimeric $Cu(BF_4)_2$ complex with the ligand in a zwitterion form and coordinating in an unusual tridentate(N₃)-monodentate(O) fashion. A similar complex with $Cu(NO_3)_2$ has been observed by Cati & Stoeckli-Evans (2004b). The structure of a mononuclear cobalt(III) complex with the ligand in a zwitterion form, coordinating in a tridentate fashion, has also been reported (Cati & Stoeckli-Evans, 2004c). Binuclear copper(II) complexes with the ligand coordinating in the bis-tridentate coordination mode have also been observed (Cati et al., 2004; Cati & Stoeckli-Evans, 2004d).



In the complex, (I), formed by the reaction of CuSO₄ with H_2L , it can be seen that the ligand coordinates in a tridentatemonodentate mode. This gives rise to the formation of a centrosymmetric neutral binuclear macrocyclic structure (Fig. 1). Selected geometric parameters are given in Table 1. The Cu atoms are coordinated by three N atoms and two O atoms; one O atom from the non-coordinated (pyridin-2-ylmethyl)amide moiety and the other from the sulfate anion. The Cu atoms have almost perfect square–pyramidal geometry, with a τ value (Addison *et al.*, 1984) of 0.05 ($\tau = 0$ for SP, and = 1 for TBP). This is similar to the geometry observed in the Cu(BF₄)₂ and Cu(NO₃)₂ complexes

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Figure 1

View of the complex, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code: (a) 1-x, 1-y, 1-z.]



The molecular packing of the complex, viewed down the a axis. Hydrogen bonds are drawn as dashed lines.

mentioned above. The bond distances and angles are also comparable to those observed in these complexes.

The macrocyclic complex is formed with the ligand having an extended conformation, giving rise to a box-like structure (Fig. 1). This arrangement is similar to that observed in the $Cu(BF_4)_2$ or $Cu(NO_3)_2$ binuclear complexes. In the former the binuclear macrocycle is charged 4+, while in the latter it is charged 2+. This is in contrast to (I), which is neutral. The (pyridin-2-ylmethyl)amide moiety involving pyridine atom N4 is relatively planar, as shown by the dihedral angles about bonds C5-N3 and N3-C6 (Table 1). The pyridine ring involving atom N6, however, is twisted by almost 90° to the plane of the amide group, as shown by the dihedral angles involving bonds C13-N5 and N5-C12 (Table 1). The N4pyridine ring and the N6-pyridine ring are inclined to the pyrazine ring by 0.9(1) and $21.0(1)^{\circ}$, respectively. The dihedral angle between the two pyridine rings is $20.2 (2)^{\circ}$. Within the macrocycle, the pyrazine rings are parallel to one another and separated by 3.61 (1) Å.

In contrast to the $Cu(BF_4)_2$ or $Cu(NO_3)_2$ complexes mentioned above, there is no intramolecular hydrogen bond involving the carbonyl O atom, O1, of the coordinated (pyridin-2-ylmethyl)amide moiety, and the NH H atom of the protonated N6-pyridine ring [(pyridinium-2-ylmethyl)amide moiety], where the carbonyl O atom, O2, coordinates to the symmetry related Cu atom. In (I), the protonated pyridine ring is twisted in the opposite direction and hydrogen bonds to a water molecule of crystallization (O2Wⁱⁱ). The non-coordinated amide proton (N5-H) is hydrogen bonded to a sulfate

O atom (O12ⁱ) coordinated to a symmetry-related molecule; details are given in Table 2.

The complex crystallizes with six water molecules of crystallization. In the crystal structure, symmetry-related complex molecules are linked by an extensive hydrogen-bonding network involving the N-H protons, the sulfate O atoms and the water molecules, as shown in Fig. 2 and Table 2.

Experimental

The synthetic and analytical details concerning the ligand, H_2L , have been described elsewhere (Cati et al., 2004). The title complex was synthesized by adding 39 mg of CuSO₄.5H₂O (0.156 mmol) to a solution of H₂L (54 mg, 0.156 mmol) in 12 ml of EtOH and 2 ml of H₂O. The resulting green solution was refluxed for *ca* 4 h then filtered directly. The filtrate was left to cool and the solvent to slowly evaporate. After several days, blue-green block-like crystals appeared (yield 41 mg, 47%).

Crystal data

 $[Cu_2(C_{18}H_{16}N_6O_2)_2(SO_4)_2]\cdot 6H_2O$ Z = 1 $M_r = 1124.03$ $D_x = 1.700 \text{ Mg m}^{-3}$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation a = 8.9507 (8) Å Cell parameters from 5864 b = 10.0094 (9) Å reflections c = 13.9732 (12) Å $\theta=2.3{-}25.9^\circ$ $\mu = 1.16~\mathrm{mm}^{-1}$ $\alpha = 90.953 (10)^{\circ}$ $\beta = 100.442 (10)^{\circ}$ T = 153 (2) K $\gamma = 116.190 \ (9)^{\circ}$ Block, blue-green V = 1098.10 (17) Å³ $0.20 \times 0.20 \times 0.08 \text{ mm}$

Data collection

Stoe Image Plate Diffraction	3966 independent reflections
System diffractometer	2687 reflections with $I > 2\sigma(I)$
φ oscillation scans	$R_{\rm int} = 0.036$
Absorption correction: refined from	$\theta_{\rm max} = 25.9^{\circ}$
ΔF (DELABS in PLATON;	$h = -10 \rightarrow 10$
Spek, 2003)	$k = -12 \rightarrow 12$
$T_{\min} = 0.718, \ T_{\max} = 0.920$	$l = -17 \rightarrow 16$
8639 measured reflections	

Refinement

Refinement on F^2 $w = \left[\exp\{2(\sin\theta/\lambda)^2\}\right] / \left[\sigma^2(F_o^2) + \right]$ $R[F^2 > 2\sigma(F^2)] = 0.029$ $(0.0241P)^2$] where $P = (F_o^2 + 2F_c^2)/3$ $wR(F^2) = 0.060$ S = 0.99 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.35$ e Å 3966 reflections $\Delta \rho_{\rm min} = -0.35 \ {\rm e} \ {\rm \AA}^{-3}$ 340 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected geometric parameters (Å, °).

Cu1-N3	1.912 (2)	Cu1-N2	2.045 (2)
Cu1-O11	1.9426 (18)	Cu1-O2 ⁱⁱⁱ	2.3292 (19)
Cu1-N4	2.002 (2)		. ,
N3-Cu1-O11	165.57 (9)	N3-Cu1-O2 ⁱⁱⁱ	107.81 (9)
N3-Cu1-N4	81.69 (9)	O11-Cu1-O2 ⁱⁱⁱ	86.39 (7)
O11-Cu1-N4	100.81 (8)	N4-Cu1-O2 ⁱⁱⁱ	92.05 (8)
N3-Cu1-N2	81.08 (8)	N2-Cu1-O2 ⁱⁱⁱ	94.45 (8)
O11-Cu1-N2	95.58 (8)	C12-O2-Cu1 ⁱⁱⁱ	132.92 (15)
N4-Cu1-N2	162.72 (8)	S1-O11-Cu1	121.36 (10)
C6-N3-C5-C2	-177.0 (2)	C13-N5-C12-C1	-173.8(2)
C5-N3-C6-C7	174.9 (2)	C12-N5-C13-C14	-93.1 (3)
C5-N5-C0-C7	174.9 (2)	012-143-015-014	-95.1 (.

Symmetry code: (iii) 1 - x, 1 - y, 1 - z.

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

$D - H \cdots A$	$D-{\rm H}$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdots A$
$N5-H5N\cdotsO12^{i}$	0.88	2.05	2.864 (3)	153
N6-H6N···O2 W^{ii}	0.88	1.77	2.644 (5)	173
$O1W-H19\cdots O1^{iii}$	0.85(5)	1.93 (5)	2.770 (4)	172 (5)
$O1W-H20\cdots O13^{iv}$	0.82(5)	2.00(5)	2.821 (3)	176 (5)
$O2W - H21 \cdots O1W$	0.90(5)	1.88 (5)	2.772 (4)	171 (4)
$O2W - H22 \cdot \cdot \cdot O3W^{v}$	0.88(4)	1.87 (4)	2.715 (5)	160 (5)
O3W−H23···O13 ^{vi}	0.84(4)	2.00(4)	2.829 (3)	171 (4)
$O3W-H24\cdots O14^{vii}$	0.82 (5)	2.00 (5)	2.817 (4)	172 (4)

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

Both the N–H and C–H H atoms could be located in Fourier difference maps but were finally included in calculated positions and treated as riding atoms: N–H distances = 0.88 Å, C–H distances = 0.95–0.99 Å, and $U_{\rm iso}$ (H) = 1.2 times $U_{\rm eq}$ (parent N or C atom). The water H atoms were located in Fourier difference maps and refined isotropically; the distance O2W–H22 was restrained to be 0.86 (2) Å.

Data collection: *EXPOSE* in *IPDS Software* (Stoe & Cie, 2000); cell refinement: *CELL* in *IPDS Software*; data reduction: *INTE-GRATE* in *IPDS Software*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure:

*SHELXL*97 (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL*97.

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