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Key indicators

Single-crystal X-ray study T = 297 KMean σ (C–C) = 0.007 Å R factor = 0.052 wR factor = 0.158 Data-to-parameter ratio = 16.9

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

trans-Bis(4-aminopyridine-*κN*)bis(succinimidato*κN*)copper(II) dihydrate

The title compound, $[Cu(C_4H_4NO_2)_2(C_5H_6N_2)_2]$ -2H₂O, has its metal atom in a square-planar *trans*- $[CuN_4]$ coordination environment; the Cu^{II} atom lies on a center of symmetry. Both succinimidate and 4-aminopyridine ligands behave as monodentate ligands, which coordinate through deprotonated imidate and pyridine (not amino) N atoms, respectively. The 4-aminopyridine is the first ligand in such Cu^{II} complexes containing two different N atoms to take potentially different coordination modes.

Comment

Recently, supramolecular self-assembling metal-organic frameworks of transition-metal complexes have been studied widely (Eddaoudi et al., 2001; Evans et al., 2002; Yaghi et al., 1998). In particular, rational design of crystals as well as molecular structures with anisotropy may be important fundamentals in the production of optical and electronic functional materials. It has been found that Cu^{II} complexes show flexible coordination geometries and electronic states (Hathaway & Billing, 1970). We have reported that the tetragonal Jahn-Teller distortion can be affected by the external temperature or light irradiation for Cu^{II} complexes with amine ligands (Akitsu & Einaga, 2003). Succinimidate ligands give rise to square-planar CuN₄ complexes, not only with four identical ligands (Akitsu, Komorita & Urushiyama, 2001) but also with mixed ligands such as trans-forms with two succinimidate and two monodentate amine ligands (Akitsu,



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The molecular structure of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code: (i) -x, -y, -z.]

Received 17 January 2006 Accepted 20 February 2006 Komorita & Kushi, 2001). Furthermore, five-coordinate CuN_4O or six-coordinate CuN_4O_2 square-bipvramidal chromophores have been reported for cis-forms with bidentate diimine (Akitsu et al., 1999) or bidentate amine (Akitsu & Komorita, 2002). However, the related Cu^{II} complexes with amine ligands containing two different coordinating N atoms have not been reported so far. Here, we report the crystal structure of the title compound, (I), incorporating 4-aminopyridine ligands with competitive amino and pyridine N-atom sites.

> NH₂ $2H_2O$ NH₂ (I)

In (I) (Fig. 1), the Cu atom exists in a square-planar CuN₄ coordination geometry; it lies on an inversion center. Both succinimidate and 4-aminopyridine ligands behave as monodentate. The Cu-N bond distances, the N-Cu-N bond angles, and other geometric parameters of the ligands (Table 1) are comparable to those of the analogous Cu^{II} complexes with pyridine derivative ligands (Latavalya & Taylor, 1975). The C1-N1-C4 angle of 110.2 (4)° indicates the sp^2 character of the deprotonated N1 atom; the O1-C1-N1 and O2-C4-N1 angles of 123.6 (4) and 124.6 (4) $^{\circ}$, respectively, also reflect electron delocalization in the π -conjugated system, similar to the Cu^{II} complexes with 5,5-diphenvlhydantoinate ligands (Akitsu & Einaga, 2004, 2005). The dihedral angle between the coordination plane [Cu1/N1/N2/N1ⁱ/N2ⁱ, symmetry code: (i) -x, -y, -z] and the succinimidate five-membered ring (N1/C1-C4) is 80.2 (2)°, and that between the coordination plane and the pyridine ring (N2/C5-C9) is $81.9(2)^{\circ}$. The dihedral angle between the succinimidate and pyridine rings is 85.0 (2)°.

As seen from Fig. 1, there is a water molecule of crystallization (together with an inversion-related one) near the Cu complex to form the $O1 \cdot \cdot \cdot H3E - O3 - H3F \cdot \cdot \cdot O2^{i}$ hydrogenbonding linkage (Table 2). There are also N-H···O intermolecular hydrogen bonds involving the amino groups.

Experimental

Treatment of copper(II) acetate (0.226 g, 1.25 mmol), succinimide (0.495 g, 5.00 mmol), and 4-aminopyridine (0.235 g, 2.50 mmol) in ethanol (50 ml) at 318 K for 2 h gave rise to a deep blue solution. Brown microcrystals (anhydrous) were obtained from the resulting solution. Yield 0.0752 g (13.2%). IR (KBr): 1618 cm⁻¹ (amide I). UV (reflectance spectra): 16700 cm⁻¹ [sh, $F(R_d) = 1.44$], 21000 cm⁻¹ $[F(R_d) = 2.36]$. Anal. Found: C 44.42, H 5.12, N 17.21%; calcd. for C18H24N6CuO6: C 44.67, H 5.00, N 17.36%. m.p. 559 K (decomposition). Brown crystals of (I) suitable for X-ray analysis were obtained from a methanol solution over a period of several days. The water solvent molecules came from undried methanol.

 $R_{\rm int} = 0.058$

 $\theta_{\rm max} = 27.5^{\circ}$ $h = -11 \rightarrow 10$

 $k = -11 \rightarrow 11$

3 standard reflections

every 150 reflections

intensity decay: 0.4%

 $l = -4 \rightarrow 10$

Crystal data

 $[Cu(C_4H_4NO_2)_2(C_5H_6N_2)_2].2H_2O$ Z = 1 $D_x = 1.521 \text{ Mg m}^{-3}$ $M_r = 483.98$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation a = 7.945 (2) Å Cell parameters from 20 b = 8.729 (3) Å reflections c = 9161(3) Å $\theta = 10.1 - 13.7^{\circ}$ $\mu = 1.08~\mathrm{mm}^{-1}$ $\alpha = 65.04 \ (2)^{\circ}$ T = 297 (2) K $\beta = 86.86(3)^{\circ}$ $\gamma = 67.59 (2)^{\circ}$ Prism, brown V = 528.2 (3) Å³ $0.20 \times 0.20 \times 0.20$ mm

Data collection

Rigaku AFC-7R diffractometer ω –2 θ scans Absorption correction: ψ scan (North et al., 1968) $T_{\min} = 0.696, T_{\max} = 0.807$ 2736 measured reflections 2430 independent reflections 1771 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_0^2) + (0.0697P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.052$ wR(F²) = 0.158 + 0.5544P] where $P = (F_0^2 + 2F_c^2)/3$ S = 1.06 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.58 \text{ e } \text{\AA}^{-3}$ 2430 reflections $\Delta \rho_{\rm min} = -0.54 \text{ e } \text{\AA}^{-3}$ 144 parameters H atoms treated by a mixture of independent and constrained

Cu1-N1	1.971 (3)	N1-C1	1.349 (6)
Cu1-N2	2.011 (4)	N1-C4	1.371 (6)
O1-C1	1.235 (6)	N2-C5	1.339 (6)
O2-C4	1.214 (6)	N2-C9	1.342 (6)
N1-Cu1-N2	89.93 (14)	C5-N2-C9	116.2 (4)
C1-N1-C4	110.2 (4)	C5-N2-Cu1	121.5 (3)
C1-N1-Cu1	127.0 (3)	C9-N2-Cu1	122.0 (3)
C4-N1-Cu1	122.8 (3)		. ,

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$\begin{array}{c} O3-H3E\cdots O1\\ O3-H3F\cdots O2^{i}\\ \end{array}$	0.73 (3)	2.13 (2)	2.827 (5)	161 (2)
	0.79 (3)	1.99 (2)	2.770 (7)	168 (2)
$N3 - H3C \cdots O1^{n}$	0.86	2.20	3.016 (5)	158
$N3 - H3D \cdots O3^{n}$	0.86	2.08	2.917 (6)	164

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

refinement Table 1 Selected geometric parameters (Å, °). The water H atoms were located in difference Fourier maps and their positions were refined with $U_{\rm iso} = 0.064 \text{ Å}^2$. H atoms bonded to C and N atoms were placed in calculated positions, with C–H and N–H = 0.93–0.97 and 0.86 Å, respectively, and with $U_{\rm iso}(H) = 1.2U_{\rm eq}(C,N)$.

Data collection: WinAFC Diffractometer Control Software (Rigaku, 1999); cell refinement: WinAFC Diffractometer Control Software ; data reduction: TEXSAN (Molecular Structure Corporation, 1989); program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: TEXSAN.

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