Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.026 wR factor = 0.064 Data-to-parameter ratio = 17.9

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[2,6-Bis(2-pyridylamino)pyridine]dichlorocadmium(II)

In the title complex, $[CdCl_2(C_{15}H_{13}N_5)]$ or $[CdCl_2(tpdaH_2)]$ (tpdaH₂ is tripyridyldiamine), the geometry at the fivecoordinate Cd atom is distorted tetragonal pyramidal, with two Cl atoms and the three pyridine N atoms of the tripyridyldiamine molecule. The ligand (tpdaH₂) adopts an all-*anti* form. The observed Cd–N bond lengths are 2.2958 (19), 2.3254 (19) and 2.3067 (19) Å, slightly shorter than those in the Cd²⁺ complex with di-2-pyridylamine (dpa).

Comment

Transition metal complexes containing a polypyridylamine ligand are of interest to chemists because of their novel structures and special properties. The polypyridyl ligand, with its good σ electron-donating and π electron-withdrawing properties, is a popular chelating ligand used in modern coordination chemistry (Jing et al., 2000). Metal chain complexes are mostly used in the fundamental study of metalmetal interactions (Yang et al., 1997; Cotton et al., 1998) and investigated for their potential application as molecular metal wires. Previously, a series of polynuclear metal chain complexes have been successfully synthesized and characterized (Sheu et al., 1996; Shieh et al., 1997; Chang et al., 1999; Peng et al., 2000). Recently, we extended these studies by synthesizing some new transition metal complexes having metal atoms with larger atomic radii. We designed the reaction of the cadmium(II) ion with tripyridyldiamine (tpdaH₂) using earlier methods (Sheu et al., 1996; Shieh et al., 1997) and attempted to synthesize Cd-Cd metal chain complexes. The result shows that the synthesis of polynuclear metal chain complexes of cadmium is very difficult under these conditions, and only the mononuclear complex was obtained.



As shown in Fig. 1, the title complex, (I), is a neutral molecule and the geometry of the Cd atom is distorted tetragonal pyramidal, with two Cl atoms and three N atoms from pyridine rings; this structure is similar to others found in the literature (Zhou *et al.*, 2000; Bian *et al.*, 2003). Atom Cl1 is located at the apex of the pyramid, and atoms N1, N3, N5 and Cl2 lie at the four corners of the tetragonal pyramidal base, with a

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Figure 1

Molecular structure of the title compound. Displacement ellipsoids are shown at the 35% probability level and H atoms have been omitted.



Figure 2

The packing diagram of the title compound. Displacement ellipsoids are shown at the 15% probability level. Dashed lines indicate hydrogen bonds.

maximum deviation from the mean plane of these basal atoms (N1, N3, N5 and Cl2) of 0.2380 (3) Å for N3. The lengths of the two Cd-Cl bonds are 2.5298 (6) and 2.4691 (6) Å, both within normal ranges (Tao et al., 2002). The Cd-N bond lengths are 2.2958 (19), 2.3254 (19) and 2.3067 (19) Å, and are slightly shorter than those in the Cd^{II}–di-2-pyridylamine (dpa) complex (Zhu et al., 2001). Selected bond distances and angles are listed in Table 1. To the best of our knowledge, fivecoordinate Cd^{II} is found in only about 5% of cadmium complexes (Wang et al., 2003; Sigel & Martin, 1994). The tripyridyldiamine is a tridentate chelating ligand in this structure, and it exhibits an all-anti conformation. The three pyridine rings are not coplanar, and the dihedral angles between the planes of the central pyridine ring and the others are 15.6 (5) and 34.1 (3)°.

The molecular stacking is evident from the packing diagram (Fig. 2). The complex molecules overlap each other, forming layers linked by van der Waals forces. A closer inspection of the crystal structure reveals the presence of intermolecular hydrogen bonds between adjacent Cl atoms and amine N atoms of the tpdaH₂ ligand in the same layer, which play an important role in the assembly of the crystal structure. The main hydrogen bonds are listed in Table 2.

Experimental

Tripyridyldiamine (0.415 g), CdCl₂·2H₂O (0.74 g) and naphthalene (9.72 g) were placed in an Erlemeyer flask. The system was put under vacuum for 10 min and always kept under nitrogen. The mixture was heated for 10 min until boiling, and then anhydrous tert-butyl alcohol (8 ml) was added with an injector and the mixture was kept boiling for 1 h. A mixture of potassium tert-butoxide (0.5 g) and anhydrous tert-butyl alcohol (5 ml) was then injected, with continued heating for 4 h. The mixture was washed with n-hexane, and the solid was recrystallized from acetonitrile/diethyl ether. Light-yellow single crystals were obtained after 3 d.

Crystal data

$[CdCl_2(C_{15}H_{13}N_5)]$	$D_x = 1.823 \text{ Mg m}^{-3}$
$M_r = 446.60$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 6805
a = 7.7545 (2) Å	reflections
b = 23.8945 (6) Å	$\theta = 2.5 - 27.5^{\circ}$
c = 9.4462 (2) Å	$\mu = 1.68 \text{ mm}^{-1}$
$\beta = 111.644 \ (1)^{\circ}$	T = 295 (2) K
$V = 1626.88 (7) \text{ Å}^3$	Plate, light yellow
Z = 4	$0.40 \times 0.40 \times 0.05 \text{ mm}$

3733 independent reflections 3041 reflections with $I > 2\sigma(I)$

 $R_{\rm int}=0.023$ $\theta_{\rm max} = 27.5^{\circ}$ $h = -10 \rightarrow 10$ $k=-28\rightarrow 31$ $l = -12 \rightarrow 10$

Data collection

Bruker SMART CCD area detector
diffractometer
φ and ω scans
Absorption correction: multi-scan
(North et al., 1968)
$T_{\min} = 0.523, \ T_{\max} = 0.920$
14 214 measured reflections

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.026$	$w = 1/[\sigma^2(F_o^2) + (0.0393P)^2]$
$wR(F^2) = 0.064$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.01	$(\Delta/\sigma)_{\rm max} = 0.001$
3733 reflections	$\Delta \rho_{\rm max} = 0.46 \text{ e} \text{ Å}^{-3}$
208 parameters	$\Delta \rho_{\rm min} = -0.46 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cd-N1	2.2958 (19)	Cd-Cl1	2.4691 (6)
Cd-N5	2.3067 (19)	Cd-Cl2	2.5298 (6)
Cd-N3	2.3254 (19)		
N1-Cd-N5	158.84 (7)	N5-Cd-Cl1	100.01 (5)
N1-Cd-N3	82.90 (7)	N3-Cd-Cl1	105.69 (5)
N5-Cd-N3	83.57 (7)	N1-Cd-Cl2	88.43 (5)
N1-Cd-Cl1	99.27 (5)	Cl1-Cd-Cl2	114.43 (2)

Table	2	

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N2-H2A\cdots Cl2^{i}$ $N4-H4A\cdots Cl1^{ii}$ $C15-H15A\cdots Cl2$	0.86	2.44	3.303 (3)	170
	0.86	2.59	3.294 (6)	139
	0.93	2.73	3.408 (2)	131

Symmetry codes: (i) 2 - x, -y, 1 - z; (ii) $x - \frac{1}{2}, -\frac{1}{2} - y, z - \frac{1}{2}$.

H atoms were positioned geometrically, with N-H = 0.86 Å and C-H = 0.93 Å, and refined in the riding-model approximation, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C,N}).$

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL-Plus* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*93 (Sheldrick, 1993); molecular graphics: *SHELXTL-Plus*; software used to prepare material for publication: *SHELXTL-Plus*.

The authors thank the National Natural Science Foundation (grant No. 20261004) of China for aid.

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