metal-organic compounds

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Bis(µ-hexamethylenetetramine)bis-(aquadibromocadmium)diaquadibromocadmium dihydrate

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The title compound, poly[[diaquadibromocadmium- μ -(1,3,5,7-tetraazatricyclo[3.3.1.1^{3,7}]decane- $N^1:N^5$)-aquacadmium-di- μ -bromo-aquacadmium- μ -(1,3,5,7-tetraazatricyclo-[3.3.1.1^{3,7}]decane- $N^1:N^5$)-di- μ -bromo] dihydrate], [Cd₃Br₆(C₆-H₁₂N₄)₂(H₂O)₄]·2H₂O, is made up of two-dimensional neutral rectangular coordination layers. Each rectangular subunit is enclosed by a pair of Cd₃(μ_2 -Br)₆(H₂O)₃ fragments and a pair of (μ_2 -hmt)Cd(H₂O)₂Br₂(μ_2 -hmt) fragments as sides (hmt is hexamethylenetetramine). The unique Cd^{II} atom in the Cd₂Br₂ ring in the Cd₃(μ_2 -Br)₆(H₂O)₃ fragment is in a slightly distorted octahedral CdNOBr₄ geometry, surrounded by one

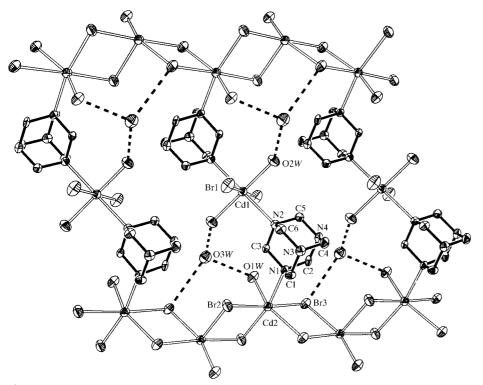


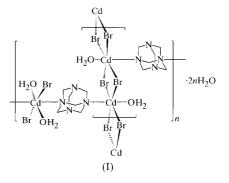
Figure 1

ORTEP (Sheldrick, 1998) plot (35% probability displacement ellipsoids) of the coordination layer with rectangular cavities in (I).

hmt ligand [2.433 (5) Å], one aqua ligand [2.273 (4) Å] and four Br atoms [2.6409 (11)–3.0270 (14) Å]. The Cd^{II} atom in the $(\mu_2$ -hmt)Cd(H₂O)₂Br₂(μ_2 -hmt) fragment lies on an inversion center and is in a highly distorted octahedral CdN₂O₂Br₂ geometry, surrounded by two *trans*-related N atoms of two hmt ligands [2.479 (5) Å], two *trans*-related aqua ligands [2.294 (4) Å] and two *trans*-related Br atoms [2.6755 (12) Å]. Adjacent two-dimensional coordination sheets are connected into a three-dimensional network by hydrogen bonds involving lattice water molecules, and the aqua, bromo and hmt ligands belonging to different layers.

Comment

Hexamethylenetetramine (hmt) as a potential polydentate ligand has been of interest in recent years (Miyamae *et al.*,



1988; Mak, 1984; Pickardt & Droas, 1985). These compounds may exhibit interesting topological structures. We and others

have recently reported some interesting two- and three-dimensional non-interpenetrating open networks constructed from Ag^I ions and the μ_4 -hmt ligand (Bertelli *et al.*, 1997; Carlucci *et al.*, 1997; Tong *et al.*, 1999). We describe herein the crystal structure of the title compound, (I), with a two-dimensional rectanglelike network, in which hmt acts in a μ_2 -bridging mode.

The title complex is made up of two-dimensional neutral layers of rectangular subunits and lattice water molecules. As shown in Fig. 1, each rectangular subunit is enclosed by a pair of $Cd_3(\mu_2-Br)_6(H_2O)_3$ fragments and pair а of $Cd(H_2O)_2Br_2(\mu_2-hmt)_2$ fragments as sides. The $Cd(H_2O)_2Br_2(\mu_2-hmt)_2$ fragment has inversion symmetry with octahedral coordination by two trans-related monodentate Br atoms [Cd1-Br1 2.6755 (12) Å], two transrelated aqua ligands [Cd1-O2W 2.294 (4) Å] and two trans-related N atoms [Cd1-N2 2.479 (5) Å] from

different hmt ligands. The Cd2 atom in the Cd₃(μ_2 -Br)₆(H₂O)₃ fragment is in a highly distorted octahedral geometry and coordinated by four Br atoms [2.6409 (11)–3.0270 (14) Å], one aqua [2.273 (4) Å] and one hmt ligand [2.433 (5) Å]. Along the *a* direction, the Cd^{II} atoms share common Br atoms at the corners to form Cd(μ_2 -Br)₂(H₂O) infinite chains, similar to related compounds (Rogers *et al.*, 1993; Rogers & Bond, 1996; Krishnan *et al.*, 1991). These chains are further interlinked by (μ_2 -hmt)–CdBr₂(H₂O)₂–(μ_2 -hmt) fragments, resulting in a two-dimensional coordination rectangular layer. The hmt ligand acts in a μ_2 -mode and bridges two Cd^{II} atoms belonging to two types of Cd^{II} atoms. The Cd–Br(μ_2 -mode) bonds are largely unequivalent compared with those [2.668 (3) and 2.780 (4) Å] found in a related compound, *catena*-[(μ_2 -nicotinato-*O*,*O'*,*N*)- μ_2 -aquacadmium(II)] (Zhang *et al.*, 1996).

The structure of (I) differs significantly from that of a closely related compound, $bis(\mu_2$ -hexamethylenetetramine)bis(diiodocadmium)diaquadiiodocadmium dihydrate, in which the Cd^{II} atoms have an octahedral or tetrahedral coordinate geometry, resulting in a three-dimensional coordination network (Mak, 1981, 1982; Pickardt, 1981).

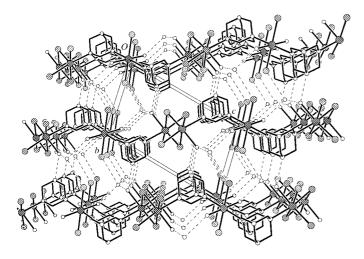


Figure 2 Perspective view of the molecular packing of (I).

It is also noteworthy that the extensive interlayer hydrogen bonds between the adjacent layers may play a role in consolidating the crystal architecture. Each O1W ligand forms one hydrogen bond with a lattice water molecule and one interlayer hydrogen bond with the hmt ligand belonging to the adjacent layer. Each O2W ligand donates only one hydrogen bond, which is the lattice water molecule. Interestingly, each lattice water molecule forms two acceptor hydrogen bonds with two aqua ligands and two donor hydrogen bonds with two bromo ligands $[O1W \cdots O3W \ 2.812 (7), O2W \cdots O3W^{i} 2.837 (7), O3W \cdots Br1^{ii} 3.332 (5), O3W \cdots Br3^{iii} 3.415 (5),$ $O1W \cdots N3^{ii} 2.771 (7) and O2W - Br1^{i} 3.522 (5) Å; symmetry$ codes: (i) <math>-x, -y, -z; (ii) x, y - 1, z; (iii) x - 1, y, z]. These hydrogen bonds extend the two-dimensional coordination layers into three-dimensional molecular networks (Fig. 2).

Experimental

To a solution (5 ml) of $CdBr_2 \cdot 4H_2O$ (1 mmol), a solution (5 ml) of hmt (1 mmol) was added slowly with stirring at 323 K for 15 min. The resulting solution was allowed to stand in air at room temperature and colorless crystals were deposited within several days (90% yield).

Crystal data

 $\begin{bmatrix} Cd_3Br_6(C_6H_{12}N_4)_2(H_2O)_4 \end{bmatrix} \cdot 2H_2O \\ M_r = 1205.15 \\ Triclinic, P\overline{I} \\ a = 7.577 (2) Å \\ b = 8.030 (2) Å \\ c = 13.464 (3) Å \\ \alpha = 76.60 (3)^{\circ} \\ \beta = 89.17 (3)^{\circ} \\ \gamma = 82.60 (3)^{\circ} \\ \gamma = 82.60 (3)^{\circ} \\ V = 790.2 (3) Å^{3} \end{bmatrix}$

Data collection

Siemens R3M diffractometer ω scans Absorption correction: semiempirical absorption based on ψ scans (North *et al.*, 1968) $T_{\min} = 0.112, T_{\max} = 0.214$ 3630 measured reflections 3630 independent reflections

Refinement

Refinement on F^2 w $R[F^2 > 2\sigma(F^2)] = 0.043$ w $wR(F^2) = 0.104$ SS = 1.025(23630 reflections Δ 161 parameters Δ H atoms treated by a mixture of
independent and constrained
refinementE

Z = 1 $D_x = 2.533 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 25 reflections $\theta = 7.5-15.0^{\circ}$ $\mu = 9.625 \text{ mm}^{-1}$ T = 293 (2) K Block, colorless $0.30 \times 0.20 \times 0.16 \text{ mm}$

2716 reflections with $I > 2\sigma(I)$ $\theta_{\text{max}} = 27.57^{\circ}$ $h = 0 \rightarrow 9$ $k = -10 \rightarrow 10$ $l = -17 \rightarrow 17$ 2 standard reflections every 150 reflections intensity decay: none

$$\begin{split} & w = 1/[\sigma^2(F_o^2) + (0.0557P)^2 \\ & + 0.0981P] \\ & \text{where } P = (F_o^2 + 2F_c^2)/3 \\ & (\Delta/\sigma)_{\text{max}} = 0.002 \\ & \Delta\rho_{\text{max}} = 1.00 \text{ e } \text{ Å}^{-3} \\ & \Delta\rho_{\text{min}} = -0.93 \text{ e } \text{ Å}^{-3} \\ & \text{Extinction correction: } SHELXL97 \\ & (\text{Sheldrick, 1997}) \\ & \text{Extinction coefficient: } 0.0046 (5) \end{split}$$

Table 1

Selected geometric parameters (Å, °).

Cd1 - O2W	2.294 (4)	Cd2-Br3 ⁱ	2.6409 (11)
Cd1-N2	2.479 (5)	Cd2-Br2 ⁱⁱ	2.7287 (13)
Cd1-Br1	2.6755 (12)	Cd2-Br2	2.7482 (13)
Cd2 - O1W	2.273 (4)	Cd2-Br3	3.0270 (14)
Cd2-N1	2.433 (5)		
O2W-Cd1-N2	88.47 (17)	Br3 ⁱ -Cd2-Br2 ⁱⁱ	99.99 (4)
O2W-Cd1-Br1	90.06 (13)	O1W-Cd2-Br2	96.00 (13)
N2-Cd1-Br1	91.75 (13)	N1-Cd2-Br2	85.88 (13)
O1W-Cd2-N1	83.48 (17)	Br3 ⁱ -Cd2-Br2	106.02 (4)
O1W-Cd2-Br3 ⁱ	157.16 (13)	Br2 ⁱⁱ -Cd2-Br2	87.17 (4)
N1-Cd2-Br3i	91.96 (12)	O1W-Cd2-Br3	76.72 (13)
O1W-Cd2-Br2 ⁱⁱ	86.88 (12)	N1-Cd2-Br3	95.03 (13)
N1-Cd2-Br2 ⁱⁱ	167.46 (11)		

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

Data collection: *R3M Software* (Siemens, 1990); cell refinement: *R3M Software*; data reduction: *R3M Software*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 1998); software used to prepare material for publication: *SHELXL*97.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: DE1139). Services for accessing these data are described at the back of the journal.

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