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Dibromido[1,1'-dibenzyl-2,2'-(sulfaneyldimethylene)di-1*H*-benzimidazole]-cadmium(II) dimethylformamide solvate

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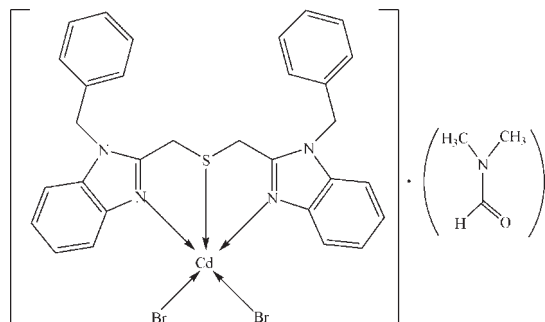
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 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; disorder in solvent or counterion; R factor = 0.029; wR factor = 0.077; data-to-parameter ratio = 15.7.

In the title compound, $[\text{CdBr}_2(\text{C}_{30}\text{H}_{26}\text{N}_4\text{S})] \cdot \text{C}_3\text{H}_7\text{NO}$, both the complex and solvent molecule lie on a crystallographic mirror plane. The Cd^{II} ion is coordinated in a distorted square-pyramidal $\text{CdBr}_2\text{N}_2\text{S}$ environment with one of the Br atoms in the apical site. In the crystal structure, the benzimidazole ring systems are involved in weak intermolecular $\pi-\pi$ stacking interactions [centroid-centroid distances = 3.606 (2) and 3.753 (2) Å]. Further stabilization is provided by weak intermolecular $\text{C}-\text{H} \cdots \text{O}$ hydrogen bonds. The methyl H atoms of the dimethylformamide solvent molecule are disordered about a mirror plane.

Related literature

For background to the synthesis and for related structures of 1,3-bis(benzimidazol-2-yl)-2-thiopropane and its derivatives, see: Dagdigian *et al.* (1979); Agh-Atabay *et al.* (2004); Wu *et al.* (2009).



Experimental

Crystal data

 $[\text{CdBr}_2(\text{C}_{30}\text{H}_{26}\text{N}_4\text{S})] \cdot \text{C}_3\text{H}_7\text{NO}$
 $M_r = 819.92$

 Monoclinic, $P2_1/m$
 $a = 9.7437$ (8) Å

 $b = 16.7792$ (14) Å

 $c = 10.5931$ (9) Å

 $\beta = 110.029$ (1)°

 $V = 1627.1$ (2) Å³
 $Z = 2$

 Mo $K\alpha$ radiation

 $\mu = 3.23$ mm⁻¹
 $T = 296$ K

 $0.36 \times 0.32 \times 0.28$ mm

Data collection

Bruker APEXII area-detector diffractometer

Absorption correction: multi-scan (SADABS; Bruker, 2006)

 $T_{\text{min}} = 0.390$, $T_{\text{max}} = 0.465$

9062 measured reflections

3305 independent reflections

 2742 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.077$
 $S = 1.05$

3305 reflections

211 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.78$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.64$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{C1}-\text{H1A} \cdots \text{O1}^i$	0.97	2.38	3.004 (5)	122

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

Data collection: APEX2 (Bruker, 2006); cell refinement: SAINT (Bruker, 2006); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5084).

References

- Agh-Atabay, N. M., Baykal, A. & Somer, M. (2004). *Transition Met. Chem.* **29**, 159–163.
- Bruker (2006). APEX2, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
- Dagdigian, J. V. & Reed, C. A. (1979). *Inorg. Chem.* **18**, 2624–2626.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Wu, H. L., Wang, K. T., Yun, R. R. & Huang, X. C. (2009). *Synth. React. Inorg. Met.-Org. Chem.* **39**, 629–632.

supporting information

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Dibromido[1,1'-dibenzyl-2,2'-(sulfanediyl dimethylene)di-1*H*-benzimidazole]-cadmium(II) dimethylformamide solvate

Kaitong Wang, Jingkun Yuan, Guisheng Chen, Qian Chen and Huilu Wu

S1. Comment

The asymmetric unit of the title complex is shown in Fig. 1. The Cd^{II} ion is coordinated by one tridentate 1,3-bis(1-benzylbenzimidazol-2-yl)-2-thiopropane ligand and two bromide ions in a distorted square-pyramidal geometry. In the crystal structure, the benzimidazole ring systems are involved in weak intermolecular π - π stacking interactions [centroid-centroid distances = 3.606 (2) and 3.753 (2) Å].

S2. Experimental

To a stirred solution of 1,3-bis(1-benzylbenzimidazol-2-yl)-2-thiopropane (0.237 g, 0.50 mmol) in hot MeOH (10 ml) was added Cd(C₆H₂N₃O₇)₂ (0.154 g, 0.25 mmol) and KBr (0.059 g, 0.50 mmol) in MeOH (5 ml). A yellow crystalline product formed rapidly. The precipitate was filtered off, washed with MeOH and absolute Et₂O, and dried *in vacuo*. The dried precipitate was dissolved in DMF resulting in a yellow solution. The deep yellow crystals suitable for X-ray diffraction studies were obtained by ether diffusion into a solution of the title compound in DMF after several days at room temperature. Yield, 0.29 g (73%). (found: C, 54.16; H, 4.49; N, 9.63. Calcd.: C, 54.22; H, 4.55; N, 9.58)

S3. Refinement

All H atoms were visible in difference Fourier maps and were subsequently refined in a riding-model approximation with C—H distances ranging from 0.93 to 0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ or $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{C}_{\text{methyl}})$.

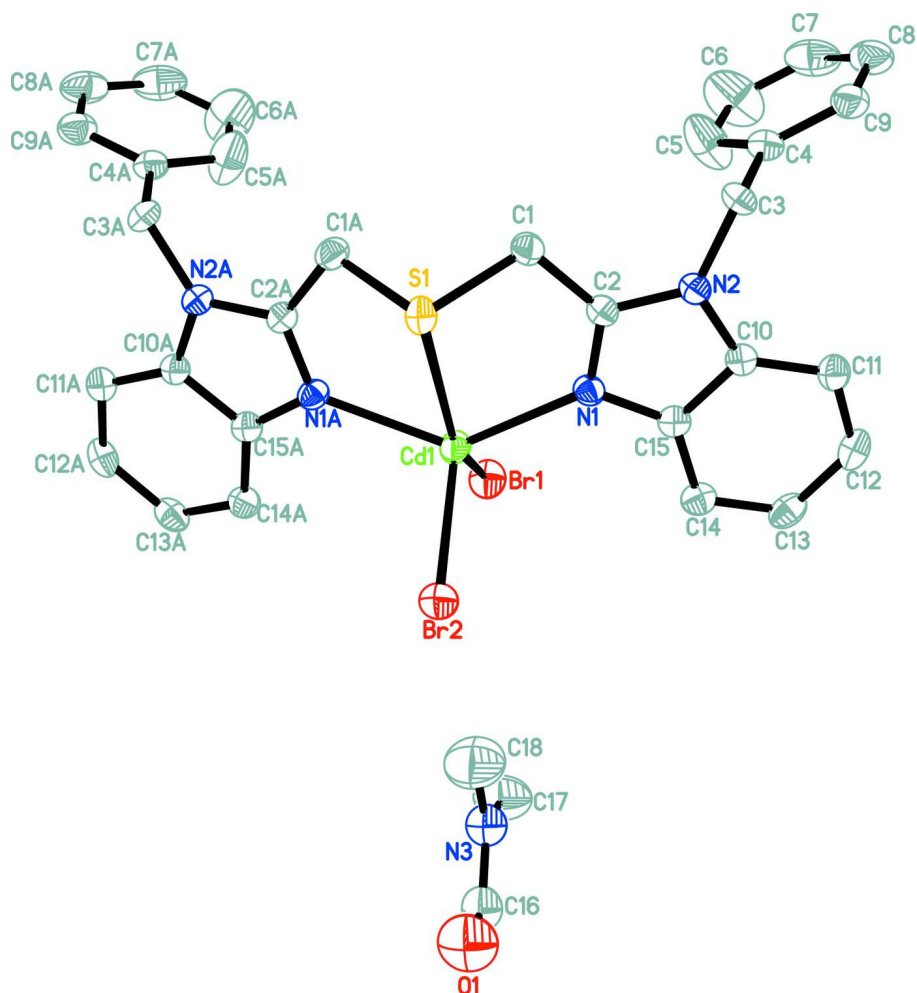


Figure 1

The asymmetric unit of the title compound. Displacement ellipsoids are drawn at the 30% probability level. H atoms have been omitted for clarity [symmetry code (A): $x, -y+1/2, z$].

Dibromido[1,1'-dibenzyl-2,2'-(sulfanediyldimethylene)di-1*H*-benzimidazole]cadmium(II) dimethylformamide solvate

Crystal data

$[\text{CdBr}_2(\text{C}_{30}\text{H}_{26}\text{N}_4\text{S})] \cdot \text{C}_3\text{H}_7\text{NO}$

$M_r = 819.92$

Monoclinic, $P2_1/m$

Hall symbol: $-P\ 2y\ b$

$a = 9.7437\ (8)\ \text{\AA}$

$b = 16.7792\ (14)\ \text{\AA}$

$c = 10.5931\ (9)\ \text{\AA}$

$\beta = 110.029\ (1)^\circ$

$V = 1627.1\ (2)\ \text{\AA}^3$

$Z = 2$

$F(000) = 816$

$D_x = 1.674\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 3975 reflections

$\theta = 2.2\text{--}27.3^\circ$

$\mu = 3.23\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Block, yellow

$0.36 \times 0.32 \times 0.28\ \text{mm}$

Data collection

Bruker APEXII area-detector diffractometer	9062 measured reflections
Radiation source: fine-focus sealed tube	3305 independent reflections
Graphite monochromator	2742 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.027$
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2006)	$\theta_{\text{max}} = 26.0^\circ$, $\theta_{\text{min}} = 2.1^\circ$
$T_{\text{min}} = 0.390$, $T_{\text{max}} = 0.465$	$h = -11 \rightarrow 12$
	$k = -20 \rightarrow 20$
	$l = -13 \rightarrow 6$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.029$	$w = 1/[\sigma^2(F_o^2) + (0.0374P)^2 + 0.7961P]$
$wR(F^2) = 0.077$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.05$	$(\Delta/\sigma)_{\text{max}} < 0.001$
3305 reflections	$\Delta\rho_{\text{max}} = 0.78 \text{ e } \text{\AA}^{-3}$
211 parameters	$\Delta\rho_{\text{min}} = -0.64 \text{ e } \text{\AA}^{-3}$
0 restraints	Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001x Fc^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.0025 (4)
Secondary atom site location: difference Fourier map	

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Br1	0.53761 (5)	0.2500	0.42042 (5)	0.05529 (15)	
Br2	0.95622 (6)	0.2500	0.72708 (5)	0.05828 (15)	
C1	1.0172 (3)	0.16482 (17)	0.2726 (3)	0.0434 (7)	
H1A	0.9562	0.1832	0.1844	0.052*	
H1B	1.1010	0.1377	0.2630	0.052*	
C2	0.9329 (3)	0.10786 (16)	0.3253 (3)	0.0343 (6)	
C3	0.9688 (3)	-0.01196 (18)	0.1929 (3)	0.0410 (7)	
H3A	1.0153	-0.0620	0.2295	0.049*	
H3B	1.0400	0.0207	0.1714	0.049*	
C4	0.8399 (3)	-0.02804 (19)	0.0656 (3)	0.0444 (7)	
C5	0.7349 (5)	0.0278 (3)	0.0121 (4)	0.0949 (16)	
H5	0.7429	0.0777	0.0523	0.114*	
C6	0.6156 (6)	0.0109 (4)	-0.1025 (5)	0.117 (2)	
H6	0.5439	0.0493	-0.1376	0.140*	
C7	0.6033 (5)	-0.0607 (3)	-0.1629 (4)	0.0825 (13)	

H7	0.5234	-0.0716	-0.2395	0.099*	
C8	0.7068 (5)	-0.1163 (3)	-0.1119 (3)	0.0691 (11)	
H8	0.6985	-0.1656	-0.1539	0.083*	
C9	0.8260 (4)	-0.1008 (2)	0.0029 (3)	0.0545 (8)	
H9	0.8967	-0.1398	0.0376	0.065*	
C10	0.8378 (3)	-0.00630 (16)	0.3597 (3)	0.0343 (6)	
C11	0.7924 (3)	-0.08440 (17)	0.3631 (3)	0.0432 (7)	
H11	0.8186	-0.1248	0.3157	0.052*	
C12	0.7067 (3)	-0.09879 (19)	0.4403 (3)	0.0491 (7)	
H12	0.6721	-0.1501	0.4436	0.059*	
C13	0.6703 (3)	-0.03818 (19)	0.5139 (3)	0.0468 (7)	
H13	0.6136	-0.0506	0.5662	0.056*	
C14	0.7157 (3)	0.03915 (18)	0.5110 (3)	0.0421 (7)	
H14	0.6912	0.0791	0.5603	0.051*	
C15	0.8005 (3)	0.05523 (17)	0.4308 (3)	0.0351 (6)	
C16	0.6054 (6)	0.2500	1.0757 (6)	0.0683 (14)	
H16	0.5334	0.2500	1.1148	0.082*	
C17	0.4073 (8)	0.2500	0.8662 (7)	0.111 (3)	
H17A	0.3912	0.2763	0.7819	0.166*	0.50
H17B	0.3553	0.2776	0.9152	0.166*	0.50
H17C	0.3728	0.1961	0.8502	0.166*	0.50
C18	0.6637 (11)	0.2500	0.8761 (10)	0.133 (3)	
H18A	0.6547	0.2014	0.8260	0.199*	0.50
H18B	0.7607	0.2540	0.9405	0.199*	0.50
H18C	0.6452	0.2946	0.8158	0.199*	0.50
Cd1	0.81788 (3)	0.2500	0.46873 (3)	0.04091 (11)	
N1	0.8612 (2)	0.12589 (13)	0.4073 (2)	0.0359 (5)	
N2	0.9223 (2)	0.02893 (13)	0.2940 (2)	0.0355 (5)	
N3	0.5618 (5)	0.2500	0.9435 (5)	0.0682 (12)	
O1	0.7309 (5)	0.2500	1.1506 (5)	0.1167 (18)	
S1	1.07918 (11)	0.2500	0.38353 (10)	0.0414 (2)	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0422 (3)	0.0582 (3)	0.0688 (3)	0.000	0.0232 (2)	0.000
Br2	0.0716 (3)	0.0552 (3)	0.0480 (3)	0.000	0.0204 (2)	0.000
C1	0.0503 (17)	0.0394 (16)	0.0495 (16)	-0.0028 (13)	0.0285 (14)	-0.0058 (13)
C2	0.0313 (13)	0.0363 (15)	0.0343 (13)	0.0002 (11)	0.0099 (11)	-0.0025 (11)
C3	0.0445 (16)	0.0412 (16)	0.0406 (15)	0.0036 (13)	0.0189 (13)	-0.0089 (12)
C4	0.0488 (17)	0.0510 (18)	0.0343 (15)	0.0001 (14)	0.0156 (13)	-0.0017 (12)
C5	0.096 (3)	0.082 (3)	0.070 (3)	0.035 (3)	-0.018 (2)	-0.023 (2)
C6	0.099 (4)	0.139 (5)	0.071 (3)	0.049 (4)	-0.023 (3)	-0.013 (3)
C7	0.067 (3)	0.132 (4)	0.041 (2)	-0.013 (3)	0.0088 (18)	-0.010 (2)
C8	0.083 (3)	0.081 (3)	0.0469 (19)	-0.034 (2)	0.027 (2)	-0.0184 (19)
C9	0.066 (2)	0.055 (2)	0.0441 (17)	-0.0117 (16)	0.0214 (16)	-0.0056 (14)
C10	0.0282 (13)	0.0381 (14)	0.0342 (14)	0.0013 (11)	0.0075 (11)	0.0008 (11)
C11	0.0428 (16)	0.0383 (16)	0.0468 (16)	0.0023 (13)	0.0131 (13)	0.0017 (13)

C12	0.0426 (16)	0.0401 (17)	0.0612 (19)	-0.0004 (13)	0.0134 (15)	0.0135 (14)
C13	0.0357 (15)	0.0560 (19)	0.0510 (18)	0.0053 (14)	0.0179 (14)	0.0176 (14)
C14	0.0362 (14)	0.0488 (17)	0.0434 (16)	0.0070 (13)	0.0164 (13)	0.0046 (13)
C15	0.0268 (13)	0.0409 (15)	0.0353 (14)	0.0014 (11)	0.0077 (11)	0.0006 (11)
C16	0.049 (3)	0.071 (4)	0.077 (4)	0.000	0.011 (3)	0.000
C17	0.075 (5)	0.161 (8)	0.077 (4)	0.000	0.001 (4)	0.000
C18	0.134 (8)	0.157 (9)	0.136 (7)	0.000	0.083 (6)	0.000
Cd1	0.04412 (19)	0.03567 (18)	0.0520 (2)	0.000	0.02820 (15)	0.000
N1	0.0345 (12)	0.0377 (12)	0.0394 (12)	-0.0018 (10)	0.0175 (10)	-0.0041 (10)
N2	0.0356 (12)	0.0363 (12)	0.0355 (12)	-0.0006 (10)	0.0132 (10)	-0.0036 (9)
N3	0.057 (3)	0.073 (3)	0.073 (3)	0.000	0.020 (2)	0.000
O1	0.062 (3)	0.149 (5)	0.116 (4)	0.000	0.002 (3)	0.000
S1	0.0379 (5)	0.0342 (5)	0.0486 (6)	0.000	0.0104 (4)	0.000

Geometric parameters (Å, °)

Br1—Cd1	2.6004 (6)	C10—C15	1.397 (4)
Br2—Cd1	2.6038 (6)	C11—C12	1.376 (4)
C1—C2	1.488 (4)	C11—H11	0.9300
C1—S1	1.817 (3)	C12—C13	1.399 (5)
C1—H1A	0.9700	C12—H12	0.9300
C1—H1B	0.9700	C13—C14	1.375 (4)
C2—N1	1.322 (3)	C13—H13	0.9300
C2—N2	1.360 (3)	C14—C15	1.399 (4)
C3—N2	1.468 (3)	C14—H14	0.9300
C3—C4	1.519 (4)	C15—N1	1.385 (3)
C3—H3A	0.9700	C16—O1	1.208 (7)
C3—H3B	0.9700	C16—N3	1.316 (7)
C4—C5	1.360 (5)	C16—H16	0.9300
C4—C9	1.374 (4)	C17—N3	1.446 (8)
C5—C6	1.393 (6)	C17—H17A	0.9600
C5—H5	0.9300	C17—H17B	0.9600
C6—C7	1.347 (7)	C17—H17C	0.9600
C6—H6	0.9300	C18—N3	1.408 (9)
C7—C8	1.344 (6)	C18—H18A	0.9600
C7—H7	0.9300	C18—H18B	0.9600
C8—C9	1.389 (5)	C18—H18C	0.9600
C8—H8	0.9300	Cd1—N1	2.264 (2)
C9—H9	0.9300	Cd1—N1 ⁱ	2.264 (2)
C10—N2	1.380 (3)	Cd1—S1	2.9784 (11)
C10—C11	1.387 (4)	S1—C1 ⁱ	1.817 (3)
C2—C1—S1	111.48 (19)	C12—C13—H13	119.1
C2—C1—H1A	109.3	C13—C14—C15	117.2 (3)
S1—C1—H1A	109.3	C13—C14—H14	121.4
C2—C1—H1B	109.3	C15—C14—H14	121.4
S1—C1—H1B	109.3	N1—C15—C10	109.2 (2)
H1A—C1—H1B	108.0	N1—C15—C14	130.7 (3)

N1—C2—N2	111.7 (2)	C10—C15—C14	120.1 (3)
N1—C2—C1	125.7 (2)	O1—C16—N3	125.8 (6)
N2—C2—C1	122.6 (2)	O1—C16—H16	117.1
N2—C3—C4	111.3 (2)	N3—C16—H16	117.1
N2—C3—H3A	109.4	N3—C17—H17A	109.5
C4—C3—H3A	109.4	N3—C17—H17B	109.5
N2—C3—H3B	109.4	H17A—C17—H17B	109.5
C4—C3—H3B	109.4	N3—C17—H17C	109.5
H3A—C3—H3B	108.0	H17A—C17—H17C	109.5
C5—C4—C9	118.4 (3)	H17B—C17—H17C	109.5
C5—C4—C3	121.5 (3)	N3—C18—H18A	109.5
C9—C4—C3	120.1 (3)	N3—C18—H18B	109.5
C4—C5—C6	120.4 (4)	H18A—C18—H18B	109.5
C4—C5—H5	119.8	N3—C18—H18C	109.5
C6—C5—H5	119.8	H18A—C18—H18C	109.5
C7—C6—C5	120.6 (5)	H18B—C18—H18C	109.5
C7—C6—H6	119.7	N1—Cd1—N1 ⁱ	133.74 (11)
C5—C6—H6	119.7	N1—Cd1—Br1	103.31 (6)
C8—C7—C6	119.7 (4)	N1 ⁱ —Cd1—Br1	103.31 (6)
C8—C7—H7	120.2	N1—Cd1—Br2	102.82 (6)
C6—C7—H7	120.2	N1 ⁱ —Cd1—Br2	102.82 (6)
C7—C8—C9	120.6 (4)	Br1—Cd1—Br2	109.732 (19)
C7—C8—H8	119.7	N1—Cd1—S1	69.50 (5)
C9—C8—H8	119.7	N1 ⁱ —Cd1—S1	69.50 (5)
C4—C9—C8	120.3 (4)	Br1—Cd1—S1	152.80 (3)
C4—C9—H9	119.9	Br2—Cd1—S1	97.46 (3)
C8—C9—H9	119.9	C2—N1—C15	106.0 (2)
N2—C10—C11	131.9 (3)	C2—N1—Cd1	126.29 (18)
N2—C10—C15	105.4 (2)	C15—N1—Cd1	127.02 (17)
C11—C10—C15	122.7 (3)	C2—N2—C10	107.6 (2)
C12—C11—C10	116.4 (3)	C2—N2—C3	128.0 (2)
C12—C11—H11	121.8	C10—N2—C3	123.7 (2)
C10—C11—H11	121.8	C16—N3—C18	120.8 (6)
C11—C12—C13	121.7 (3)	C16—N3—C17	119.8 (5)
C11—C12—H12	119.2	C18—N3—C17	119.4 (6)
C13—C12—H12	119.2	C1 ⁱ —S1—C1	103.7 (2)
C14—C13—C12	121.9 (3)	C1 ⁱ —S1—Cd1	93.83 (10)
C14—C13—H13	119.1	C1—S1—Cd1	93.83 (10)
S1—C1—C2—N1	25.0 (4)	N1 ⁱ —Cd1—N1—C2	5.0 (3)
S1—C1—C2—N2	-154.3 (2)	Br1—Cd1—N1—C2	128.6 (2)
N2—C3—C4—C5	42.8 (5)	Br2—Cd1—N1—C2	-117.2 (2)
N2—C3—C4—C9	-136.2 (3)	S1—Cd1—N1—C2	-23.9 (2)
C9—C4—C5—C6	0.8 (7)	N1 ⁱ —Cd1—N1—C15	-164.51 (14)
C3—C4—C5—C6	-178.2 (5)	Br1—Cd1—N1—C15	-40.9 (2)
C4—C5—C6—C7	-0.8 (9)	Br2—Cd1—N1—C15	73.3 (2)
C5—C6—C7—C8	0.2 (9)	S1—Cd1—N1—C15	166.6 (2)
C6—C7—C8—C9	0.4 (7)	N1—C2—N2—C10	0.2 (3)

C5—C4—C9—C8	-0.3 (5)	C1—C2—N2—C10	179.7 (2)
C3—C4—C9—C8	178.7 (3)	N1—C2—N2—C3	170.8 (2)
C7—C8—C9—C4	-0.3 (5)	C1—C2—N2—C3	-9.7 (4)
N2—C10—C11—C12	-179.7 (3)	C11—C10—N2—C2	179.8 (3)
C15—C10—C11—C12	0.4 (4)	C15—C10—N2—C2	-0.3 (3)
C10—C11—C12—C13	-1.4 (4)	C11—C10—N2—C3	8.7 (4)
C11—C12—C13—C14	1.2 (5)	C15—C10—N2—C3	-171.4 (2)
C12—C13—C14—C15	0.1 (4)	C4—C3—N2—C2	-100.7 (3)
N2—C10—C15—N1	0.3 (3)	C4—C3—N2—C10	68.5 (3)
C11—C10—C15—N1	-179.8 (2)	O1—C16—N3—C18	0.000 (6)
N2—C10—C15—C14	-179.0 (2)	O1—C16—N3—C17	180.000 (4)
C11—C10—C15—C14	0.9 (4)	C2—C1—S1—C1 ⁱ	-128.05 (17)
C13—C14—C15—N1	179.8 (3)	C2—C1—S1—Cd1	-33.2 (2)
C13—C14—C15—C10	-1.1 (4)	N1—Cd1—S1—C1 ⁱ	131.08 (12)
N2—C2—N1—C15	0.0 (3)	N1 ⁱ —Cd1—S1—C1 ⁱ	-27.04 (12)
C1—C2—N1—C15	-179.5 (3)	Br1—Cd1—S1—C1 ⁱ	52.02 (10)
N2—C2—N1—Cd1	-171.33 (16)	Br2—Cd1—S1—C1 ⁱ	-127.98 (10)
C1—C2—N1—Cd1	9.2 (4)	N1—Cd1—S1—C1	27.03 (12)
C10—C15—N1—C2	-0.2 (3)	N1 ⁱ —Cd1—S1—C1	-131.08 (12)
C14—C15—N1—C2	179.0 (3)	Br1—Cd1—S1—C1	-52.02 (10)
C10—C15—N1—Cd1	171.06 (17)	Br2—Cd1—S1—C1	127.98 (10)
C14—C15—N1—Cd1	-9.8 (4)		

Symmetry code: (i) $x, -y+1/2, z$.

Hydrogen-bond geometry (Å, °)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C1—H1A ⁱⁱ —O1 ⁱⁱ	0.97	2.38	3.004 (5)	122

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