

Diaquabis(thiocyanato- κN)bis[6-(4H-1,2,4-triazol-4-yl- κN^1)pyridin-2-amine]-cadmium

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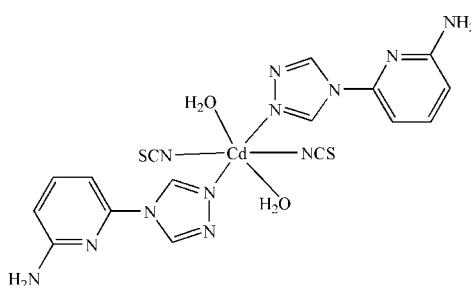
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Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.017; wR factor = 0.046; data-to-parameter ratio = 13.0.

In the title compound, $[\text{Cd}(\text{NCS})_2(\text{C}_7\text{H}_7\text{N}_5)_2(\text{H}_2\text{O})_2]$, the Cd^{II} cation lies on an inversion center and is coordinated by the N atoms of two thiocyanate anions, by N atoms of two 6-(4H-1,2,4-triazol-4-yl)pyridin-2-amine ligands and by the O atoms of two water molecules in a distorted N_4O_2 octahedral geometry. The dihedral angle between the triazole and pyridine rings is $23.15(12)^\circ$. In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{N}$ and $\text{O}-\text{H}\cdots\text{S}$ hydrogen bonds. Offset $\pi-\pi$ stacking between parallel pyridine rings of adjacent molecules is also observed, the centroid–centroid distance being $3.6319(14)\text{ \AA}$.

Related literature

For the preparation of the organic ligand, see: Gioia *et al.* (1988). For complexes with 4-3-pyridyl-1,2,4-triazole ligands, see: Moulton & Zaworotko (2001); Pan *et al.* (2001); Prior & Rosseinsky (2001); Ma *et al.* (2001); Ding *et al.* (2006); Liu *et al.* (2007).



Experimental

Crystal data

$[\text{Cd}(\text{NCS})_2(\text{C}_7\text{H}_7\text{N}_5)_2(\text{H}_2\text{O})_2]$
 $M_r = 586.94$
Triclinic, $P\bar{1}$
 $a = 7.5586(15)\text{ \AA}$

$b = 7.5876(15)\text{ \AA}$
 $c = 11.311(2)\text{ \AA}$
 $\alpha = 106.859(2)^\circ$
 $\beta = 95.790(2)^\circ$

$\gamma = 110.883(2)^\circ$
 $V = 564.7(2)\text{ \AA}^3$
 $Z = 1$
Mo $K\alpha$ radiation

$\mu = 1.19\text{ mm}^{-1}$
 $T = 293\text{ K}$
 $0.20 \times 0.16 \times 0.12\text{ mm}$

Data collection

Bruker APEXII CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2001)
 $T_{\min} = 0.796$, $T_{\max} = 0.870$

3064 measured reflections
1957 independent reflections
1893 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.010$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.017$
 $wR(F^2) = 0.046$
 $S = 1.05$
1957 reflections

151 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.31\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.22\text{ e \AA}^{-3}$

Table 1
Selected bond lengths (\AA).

Cd1–N1	2.2785 (16)	Cd1–O1	2.3501 (15)
Cd1–N6	2.3146 (19)		

Table 2
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1–H1A \cdots S1 ⁱⁱ	0.85	2.51	3.3468 (17)	168
O1–H1B \cdots S1 ⁱⁱⁱ	0.85	2.51	3.3575 (17)	172
N5–H5A \cdots N2 ^{iv}	0.86	2.23	3.080 (2)	169
N5–H5B \cdots N6 ^v	0.86	2.57	3.422 (3)	170

Symmetry codes: (ii) $-x + 2, -y + 2, -z + 1$; (iii) $x, y + 1, z$; (iv) $-x, -y + 1, -z$; (v) $x - 1, y, z - 1$.

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg, 1999); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5596).

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supporting information

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S1. Comment

Recently, considerable efforts have been devoted to crystal engineering of supramolecular architecture sustained by coordination covalent bonding, hydrogen bonding or some molecular interaction and their combination owing to their fascinating structural diversity and potential application in design of porous materials with novel inclusion or reactivity properties and in supramolecular devices such as sensor and indicator (Moulton *et al.*, 2001; Pan *et al.*, 2001; Ma *et al.*, 2001; Prior *et al.*, 2001). Our interest is to exploit the coordination chemistry of 1,2,4-triazole and its derivatives together with their potential application in material science (Liu *et al.*, 2007; Ding *et al.*, 2006).

In the report, the mono-nuclear Cadmium(II) complex was obtained *via* the reaction of 2-amino-6-(4-triazoyl)pyridine, NH₄NCS and corresponding Cadmium(II) salts. A view of the coordination compound [Cd(II)L₂(NCS)₂(H₂O)₂] is shown in Figure 1. Single crystal X-ray diffraction analysis reveals that the Cadmium(II) atom is six-coordinated by two pyridine nitrogen atoms, two NCS nitrogen atoms and two aqua oxygen atoms forming N₄O₂ donor set. Bond distances of Cd—N and Cd—O(Cd(1)—N(1):2.2785 (16) Å; Cd(1)—N(6):2.3146 (19) Å; Cd(1)—O(1):2.3500 (15) Å) are listed. The coordination geometry around the Cadmium(II) center in the molecular lattice lie in the inversion center and can be described as the Octahedral geometry.

L is mono-dentate terminal ligand coordinated *via* its pyridine nitrogen atoms. The weak N···N interactions between L triazole rings (N—H···N, 3.080 (2) and 3.422 (3) Å) between L triazole rings can be observed. The offset π ··· π stacking interactions between two neighboring pyridine rings are also important for the assembly of the supra-molecular structure, the ring centroid-centroid distance being 3.632 (3) Å. As shown in Figure 2, A two-dimensional supra-molecular network can be observed stabilized *via* N···N interactions and π ··· π stacking interactions.

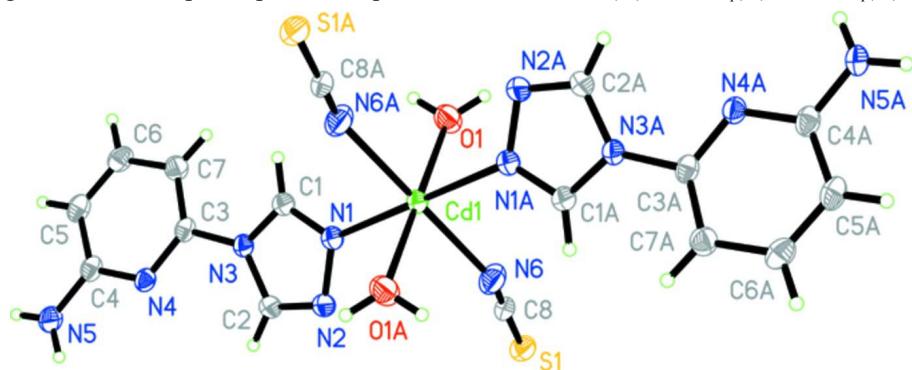
Further the non-classic O—H···S hydrogen bonds (O—H···S, 3.346 (8) and 3.357 (5) Å) also can be observed, which further assemble these two-dimensional supramolecular network to form a three-dimensional supra-molecular structure. The three-dimensional packing architecture in the unit cell of the complex is shown in Figure 3.

S2. Experimental

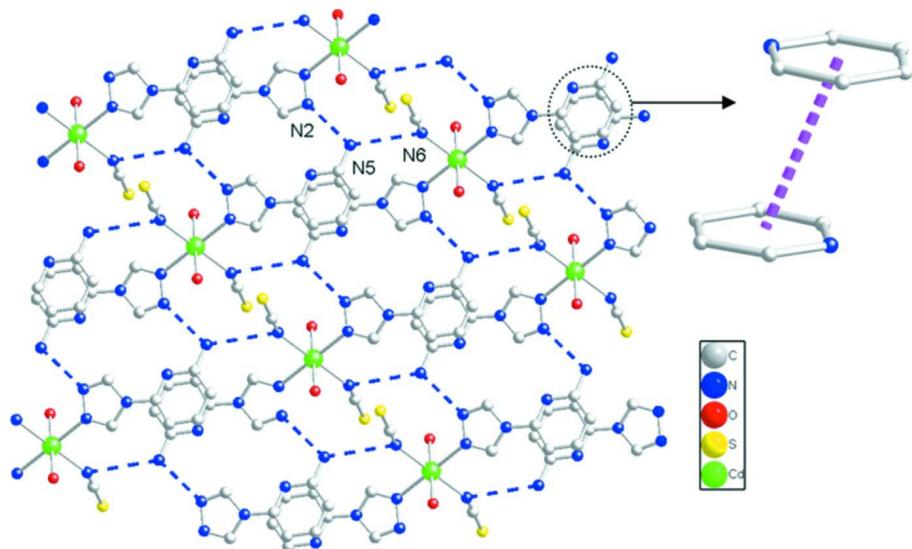
The organic ligand L was prepared according to the previously reported literature methods (Gioia *et al.*, 1988). A mixture of CdBr₂ (27.2 mg, 0.1 mmol), NH₄NCS (7.6 mg, 0.1 mmol), L (14.6 mg, 0.1 mmol) and water (10 ml) was stirred for 5 h and filtered. Suitable single crystals for X-ray diffraction study were obtained after a few days, yield 23% (based on Cd(II) salts). Anal. Calc. for C₁₆H₁₈CdN₁₂O₂S₂: C, 32.74%; H, 3.09%; N, 28.63%. Found: C, 32.86%; H, 3.18%; N, 28.74%. FT-IR (KBr): 3404w, 3281w, 3135w, 2969w, 2918w, 2069 s, 1625 s, 1524m, 1405 s, 1247m, 1096m, 1017m, 792w, 676w, 618w, 529w cm⁻¹.

S3. Refinement

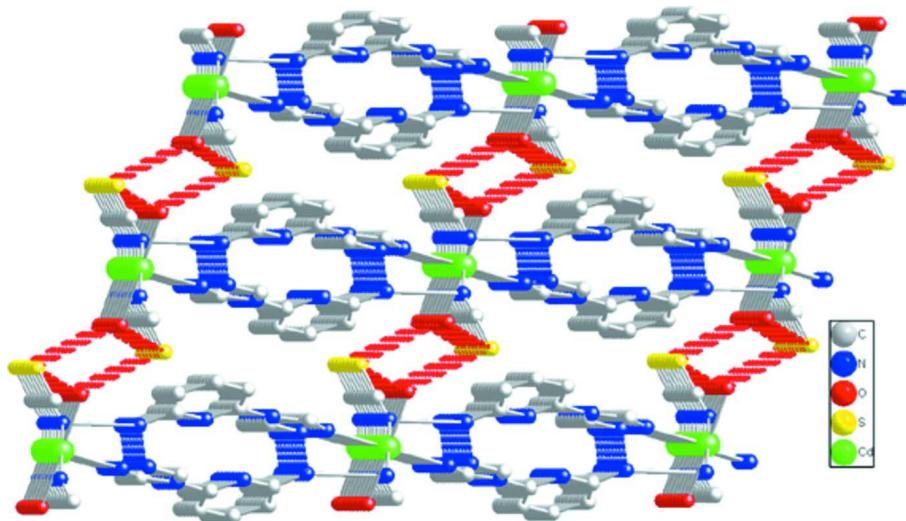
The H atoms of the aromatic rings were placed at calculated positions, with C—H = 0.93 \AA and O—H = 0.85 \AA. All H atoms were assigned fixed isotropic displacement parameters, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{O})$.

**Figure 1**

The molecular structure of (I) with atom labels and 30% probability displacement ellipsoids for non-H atoms..

**Figure 2**

The two-dimensional supra-molecular network stabilized *via* N—H···N hydrogen bonds, and offset π ··· π stacking interactions, Blue lines represent N—H···N hydrogen bonds.

**Figure 3**

The three-dimensional supramolecular packing architecture of (I). Red lines represent O—H···S hydrogen bonds and Blue lines represent N—H···N hydrogen bonds.

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Crystal data



$M_r = 586.94$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 7.5586 (15)$ Å

$b = 7.5876 (15)$ Å

$c = 11.311 (2)$ Å

$\alpha = 106.859 (2)^\circ$

$\beta = 95.790 (2)^\circ$

$\gamma = 110.883 (2)^\circ$

$V = 564.7 (2)$ Å³

$Z = 1$

$F(000) = 294$

$D_x = 1.726 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2419 reflections

$\theta = 3.0\text{--}27.8^\circ$

$\mu = 1.19 \text{ mm}^{-1}$

$T = 293$ K

Block, colorless

$0.20 \times 0.16 \times 0.12$ mm

Data collection

Bruker APEXII CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

phi and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2001)

$T_{\min} = 0.796$, $T_{\max} = 0.870$

3064 measured reflections

1957 independent reflections

1893 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.010$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 1.9^\circ$

$h = -8 \rightarrow 8$

$k = -9 \rightarrow 8$

$l = -11 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.017$

$wR(F^2) = 0.046$

$S = 1.05$

1957 reflections

151 parameters

0 restraints

H-atom parameters constrained

$$w = 1/[a^2(F_o^2) + (0.0263P)^2 + 0.1431P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.31 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.21 \text{ e \AA}^{-3}$$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cd1	0.5000	1.0000	0.5000	0.03514 (8)
S1	0.76959 (8)	0.51242 (8)	0.30994 (5)	0.04950 (14)
O1	0.7956 (2)	1.2313 (2)	0.48659 (14)	0.0558 (4)
H1A	0.8961	1.2929	0.5473	0.084*
H1B	0.7767	1.2993	0.4425	0.084*
N1	0.3864 (2)	0.8993 (2)	0.28661 (14)	0.0389 (4)
N2	0.3014 (2)	0.6986 (2)	0.20991 (15)	0.0444 (4)
N3	0.2754 (2)	0.8835 (2)	0.09699 (14)	0.0333 (3)
N4	0.0703 (2)	0.7895 (2)	-0.09637 (14)	0.0362 (3)
N5	-0.1479 (3)	0.6777 (3)	-0.28550 (16)	0.0536 (5)
H5A	-0.2021	0.5648	-0.2745	0.064*
H5B	-0.1931	0.6948	-0.3522	0.064*
N6	0.6234 (3)	0.7555 (3)	0.46977 (19)	0.0591 (5)
C1	0.3697 (3)	1.0051 (3)	0.21725 (17)	0.0381 (4)
H1	0.4162	1.1448	0.2465	0.046*
C2	0.2365 (3)	0.6948 (3)	0.09812 (18)	0.0419 (4)
H2	0.1716	0.5777	0.0277	0.050*
C3	0.2201 (3)	0.9376 (3)	-0.00828 (16)	0.0342 (4)
C4	0.0087 (3)	0.8284 (3)	-0.19785 (17)	0.0392 (4)
C5	0.0995 (3)	1.0167 (3)	-0.21024 (19)	0.0451 (5)
H5	0.0560	1.0407	-0.2815	0.054*
C6	0.2529 (3)	1.1644 (3)	-0.1160 (2)	0.0469 (5)
H6	0.3138	1.2902	-0.1227	0.056*
C7	0.3184 (3)	1.1270 (3)	-0.00964 (19)	0.0428 (4)
H7	0.4223	1.2246	0.0560	0.051*
C8	0.6818 (3)	0.6545 (3)	0.40295 (19)	0.0419 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cd1	0.03953 (12)	0.03434 (12)	0.02935 (11)	0.01460 (8)	0.00083 (8)	0.01088 (8)
S1	0.0553 (3)	0.0447 (3)	0.0527 (3)	0.0247 (2)	0.0111 (2)	0.0178 (2)
O1	0.0496 (8)	0.0550 (9)	0.0509 (9)	0.0073 (7)	0.0026 (7)	0.0226 (7)
N1	0.0424 (9)	0.0367 (8)	0.0337 (8)	0.0146 (7)	0.0009 (7)	0.0113 (7)
N2	0.0504 (9)	0.0345 (8)	0.0394 (9)	0.0105 (7)	-0.0045 (7)	0.0137 (7)

N3	0.0345 (8)	0.0358 (8)	0.0298 (7)	0.0149 (6)	0.0040 (6)	0.0121 (6)
N4	0.0438 (8)	0.0380 (8)	0.0302 (8)	0.0205 (7)	0.0055 (7)	0.0128 (6)
N5	0.0694 (12)	0.0486 (10)	0.0373 (9)	0.0225 (9)	-0.0090 (8)	0.0163 (8)
N6	0.0773 (13)	0.0569 (11)	0.0539 (11)	0.0424 (11)	0.0085 (10)	0.0178 (9)
C1	0.0420 (10)	0.0348 (9)	0.0351 (10)	0.0157 (8)	0.0034 (8)	0.0108 (8)
C2	0.0456 (10)	0.0352 (10)	0.0369 (10)	0.0126 (8)	-0.0027 (8)	0.0100 (8)
C3	0.0387 (9)	0.0424 (10)	0.0309 (9)	0.0233 (8)	0.0118 (7)	0.0162 (8)
C4	0.0492 (11)	0.0472 (11)	0.0312 (9)	0.0287 (9)	0.0115 (8)	0.0156 (8)
C5	0.0564 (12)	0.0572 (12)	0.0383 (10)	0.0315 (10)	0.0171 (9)	0.0276 (9)
C6	0.0522 (12)	0.0485 (12)	0.0539 (12)	0.0232 (10)	0.0211 (10)	0.0310 (10)
C7	0.0414 (10)	0.0451 (11)	0.0431 (11)	0.0155 (9)	0.0098 (9)	0.0197 (9)
C8	0.0450 (10)	0.0379 (10)	0.0432 (11)	0.0172 (9)	-0.0017 (9)	0.0181 (9)

Geometric parameters (\AA , $^\circ$)

Cd1—N1	2.2785 (16)	N4—C3	1.324 (2)
Cd1—N1 ⁱ	2.2786 (16)	N4—C4	1.345 (2)
Cd1—N6	2.3146 (19)	N5—C4	1.354 (3)
Cd1—N6 ⁱ	2.3146 (19)	N5—H5A	0.8600
Cd1—O1 ⁱ	2.3500 (15)	N5—H5B	0.8600
Cd1—O1	2.3501 (15)	N6—C8	1.152 (3)
S1—C8	1.644 (2)	C1—H1	0.9300
O1—H1A	0.8501	C2—H2	0.9300
O1—H1B	0.8501	C3—C7	1.368 (3)
N1—C1	1.301 (2)	C4—C5	1.403 (3)
N1—N2	1.381 (2)	C5—C6	1.366 (3)
N2—C2	1.297 (3)	C5—H5	0.9300
N3—C1	1.348 (2)	C6—C7	1.396 (3)
N3—C2	1.359 (2)	C6—H6	0.9300
N3—C3	1.438 (2)	C7—H7	0.9300
N1—Cd1—N1 ⁱ	180.00 (8)	C4—N5—H5A	120.0
N1—Cd1—N6	90.53 (6)	C4—N5—H5B	120.0
N1 ⁱ —Cd1—N6	89.47 (6)	H5A—N5—H5B	120.0
N1—Cd1—N6 ⁱ	89.47 (6)	C8—N6—Cd1	147.28 (17)
N1 ⁱ —Cd1—N6 ⁱ	90.53 (6)	N1—C1—N3	110.42 (17)
N6—Cd1—N6 ⁱ	179.999 (1)	N1—C1—H1	124.8
N1—Cd1—O1 ⁱ	89.76 (6)	N3—C1—H1	124.8
N1 ⁱ —Cd1—O1 ⁱ	90.24 (6)	N2—C2—N3	111.46 (17)
N6—Cd1—O1 ⁱ	89.56 (7)	N2—C2—H2	124.3
N6 ⁱ —Cd1—O1 ⁱ	90.45 (7)	N3—C2—H2	124.3
N1—Cd1—O1	90.24 (5)	N4—C3—C7	126.62 (17)
N1 ⁱ —Cd1—O1	89.76 (6)	N4—C3—N3	113.17 (15)
N6—Cd1—O1	90.45 (7)	C7—C3—N3	120.21 (17)
N6 ⁱ —Cd1—O1	89.55 (7)	N4—C4—N5	116.29 (17)
O1 ⁱ —Cd1—O1	180.0	N4—C4—C5	121.60 (18)
Cd1—O1—H1A	123.2	N5—C4—C5	122.09 (17)
Cd1—O1—H1B	111.3	C6—C5—C4	118.96 (18)

H1A—O1—H1B	115.7	C6—C5—H5	120.5
C1—N1—N2	107.86 (15)	C4—C5—H5	120.5
C1—N1—Cd1	129.58 (13)	C5—C6—C7	120.27 (19)
N2—N1—Cd1	122.24 (11)	C5—C6—H6	119.9
C2—N2—N1	105.99 (16)	C7—C6—H6	119.9
C1—N3—C2	104.27 (15)	C3—C7—C6	115.68 (18)
C1—N3—C3	128.48 (15)	C3—C7—H7	122.2
C2—N3—C3	127.20 (15)	C6—C7—H7	122.2
C3—N4—C4	116.86 (16)	N6—C8—S1	178.77 (19)
N1 ⁱ —Cd1—N1—C1	168 (6)	C3—N3—C1—N1	177.31 (16)
N6—Cd1—N1—C1	−146.87 (17)	N1—N2—C2—N3	0.3 (2)
N6 ⁱ —Cd1—N1—C1	33.13 (17)	C1—N3—C2—N2	0.0 (2)
O1 ⁱ —Cd1—N1—C1	123.58 (17)	C3—N3—C2—N2	−177.66 (17)
O1—Cd1—N1—C1	−56.42 (17)	C4—N4—C3—C7	0.3 (3)
N1 ⁱ —Cd1—N1—N2	−5 (6)	C4—N4—C3—N3	179.64 (15)
N6—Cd1—N1—N2	40.36 (15)	C1—N3—C3—N4	−155.32 (17)
N6 ⁱ —Cd1—N1—N2	−139.64 (15)	C2—N3—C3—N4	21.7 (2)
O1 ⁱ —Cd1—N1—N2	−49.19 (14)	C1—N3—C3—C7	24.1 (3)
O1—Cd1—N1—N2	130.81 (14)	C2—N3—C3—C7	−158.83 (19)
C1—N1—N2—C2	−0.5 (2)	C3—N4—C4—N5	−178.20 (17)
Cd1—N1—N2—C2	173.68 (13)	C3—N4—C4—C5	0.4 (3)
N1—Cd1—N6—C8	26.0 (3)	N4—C4—C5—C6	−0.7 (3)
N1 ⁱ —Cd1—N6—C8	−154.0 (3)	N5—C4—C5—C6	177.76 (19)
N6 ⁱ —Cd1—N6—C8	−19 (5)	C4—C5—C6—C7	0.5 (3)
O1 ⁱ —Cd1—N6—C8	115.8 (3)	N4—C3—C7—C6	−0.5 (3)
O1—Cd1—N6—C8	−64.2 (3)	N3—C3—C7—C6	−179.85 (16)
N2—N1—C1—N3	0.5 (2)	C5—C6—C7—C3	0.1 (3)
Cd1—N1—C1—N3	−173.11 (11)	Cd1—N6—C8—S1	123 (9)
C2—N3—C1—N1	−0.3 (2)		

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

Hydrogen-bond geometry (\AA , $^{\circ}$)

$D—\text{H}\cdots A$	$D—\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D—\text{H}\cdots A$
O1—H1A ⁱⁱ —S1 ⁱⁱ	0.85	2.51	3.3468 (17)	168
O1—H1B ⁱⁱⁱ —S1 ⁱⁱⁱ	0.85	2.51	3.3575 (17)	172
N5—H5A ^{iv} —N2 ^{iv}	0.86	2.23	3.080 (2)	169
N5—H5B ^v —N6 ^v	0.86	2.57	3.422 (3)	170

Symmetry codes: (ii) $-x+2, -y+2, -z+1$; (iii) $x, y+1, z$; (iv) $-x, -y+1, -z$; (v) $x-1, y, z-1$.