

Dichloridobis[2-(2-furyl)-1-(2-furylmethyl)-1H-benzimidazole- κN^3]cadmium(II)

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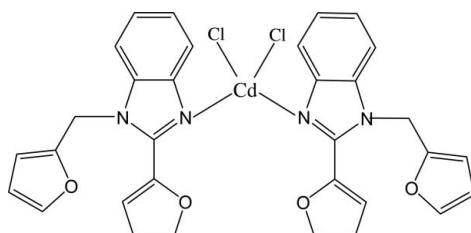
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Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$; R factor = 0.048; wR factor = 0.110; data-to-parameter ratio = 15.1.

In the title complex, $[\text{CdCl}_2(\text{C}_{16}\text{H}_{12}\text{N}_2\text{O}_2)_2]$, the Cd^{II} ion exhibits site symmetry 2. It shows a distorted tetrahedral coordination defined by two N atoms from symmetry-related 2-(2-furyl)-1-(2-furylmethyl)-1H-benzimidazole ligands and by two symmetry-related Cl atoms. Intramolecular C–H···O hydrogen bonds stabilize the molecular configuration. Adjacent molecules are linked through C–H···Cl hydrogen bonds into a network structure.

Related literature

For background to benzimidazoles, see: Shen & Yuan (2006); Yang *et al.* (2008). For background to Cd^{II} complexes, see: Meng *et al.* (2004); Yang *et al.* (2010).



Experimental

Crystal data

$[\text{CdCl}_2(\text{C}_{16}\text{H}_{12}\text{N}_2\text{O}_2)_2]$	$V = 3000.2\text{ (13) \AA}^3$
$M_r = 711.85$	$Z = 4$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 18.397\text{ (4) \AA}$	$\mu = 0.95\text{ mm}^{-1}$
$b = 10.451\text{ (2) \AA}$	$T = 293\text{ K}$
$c = 17.470\text{ (3) \AA}$	$0.21 \times 0.19 \times 0.16\text{ mm}$
$\beta = 116.72\text{ (3)}^\circ$	

Data collection

Rigaku Saturn diffractometer	10628 measured reflections
Absorption correction: multi-scan (<i>REQAB</i> ; Jacobson, 1998)	2953 independent reflections
$T_{\min} = 0.825$, $T_{\max} = 0.863$	2565 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$	195 parameters
$wR(F^2) = 0.110$	H-atom parameters constrained
$S = 1.10$	$\Delta\rho_{\max} = 0.34\text{ e \AA}^{-3}$
2953 reflections	$\Delta\rho_{\min} = -0.47\text{ e \AA}^{-3}$

Table 1
Selected geometric parameters (\AA , $^\circ$).

$\text{Cd1}-\text{N1}$	2.252 (3)	$\text{Cd1}-\text{Cl1}$	2.4513 (12)
$\text{N1}^i-\text{Cd1}-\text{N1}$	118.17 (15)	$\text{N1}-\text{Cd1}-\text{Cl1}$	106.27 (8)
$\text{N1}^i-\text{Cd1}-\text{Cl1}$	109.08 (8)	$\text{Cl1}^i-\text{Cd1}-\text{Cl1}$	107.57 (6)

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

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C5}-\text{H5A}\cdots\text{Cl1}^{ii}$	0.93	2.82	3.694 (5)	156
$\text{C9}-\text{H9A}\cdots\text{O2}$	0.93	2.49	3.256 (6)	140

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

Data collection: *CrystalClear* (Rigaku/MSC, 2006); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *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: WM2393).

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supporting information

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S1. Comment

Benzimidazole and its derivatives have attracted interest because of their biological activities as well as their abilities to bind to different metal ions (Shen & Yuan, 2006; Yang *et al.*, 2008). The Cd^{II} ion is a good model atom to construct complexes owing to its property to form bonds with different donors simultaneously, and to its various coordination modes (Meng *et al.*, 2004; Yang *et al.*, 2010). In this work, we describe the synthesis and structure of the title complex, $[\text{CdCl}_2(\text{C}_{16}\text{H}_{12}\text{N}_2\text{O}_2)_2]$, (I).

In the structure of (I), two 2-(furan-2-yl)-1-(furan-2-yl-methyl)-1*H*-1,3-benzimidazole ligands and two Cl atoms coordinate to the Cd^{II} ion which is located on a twofold rotation axis. As expected, the Cd—Cl bond length is slightly longer than the Cd—N bond length. The environment around the Cd^{II} ion can be best described as distorted tetrahedral (Fig. 1).

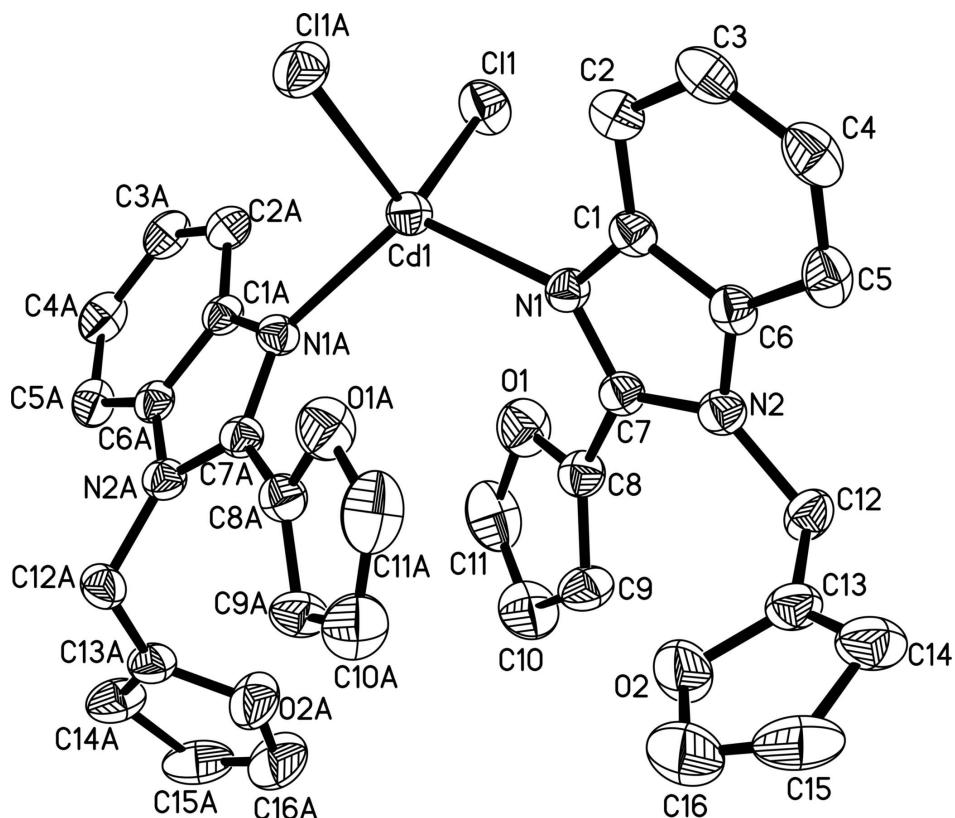
Intramolecular C—H···O hydrogen bonds stabilize the molecular configuration and C—H···Cl hydrogen bonds between adjacent molecules consolidate the crystal packing.

S2. Experimental

The ligand 2-(furan-2-yl)-1-(furan-2-yl-methyl)-1*H*-1,3-benzimidazole (0.04 mmol) in methanol (7 ml) was added dropwise to a methanol solution (5 ml) of CdCl₂ (0.02 mmol). The resulting solution was allowed to stand at room temperature. After one week colorless crystals with good quality were obtained from the filtrate and dried in air.

S3. Refinement

H atoms were positioned geometrically and refined as riding atoms, with C—H = 0.93 (aromatic) and 0.97 (CH₂) Å and with U_{iso}(H) = 1.2 U_{eq}(C).

**Figure 1**

View of the title complex, showing the labelled atoms and displacement ellipsoids at the 30% probability level. H atoms were omitted for clarity. [Symmetry code A: $-x+1, y, -z+0.5$.]

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Crystal data



$M_r = 711.85$

Monoclinic, $C2/c$

Hall symbol: $-C\bar{2}yc$

$a = 18.397 (4)$ Å

$b = 10.451 (2)$ Å

$c = 17.470 (3)$ Å

$\beta = 116.72 (3)^\circ$

$V = 3000.2 (13)$ Å³

$Z = 4$

$F(000) = 1432$

$D_x = 1.576 \text{ Mg m}^{-3}$

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

Cell parameters from 4133 reflections

$\theta = 2.3\text{--}27.9^\circ$

$\mu = 0.95 \text{ mm}^{-1}$

$T = 293$ K

Prism, colorless

$0.21 \times 0.19 \times 0.16$ mm

Data collection

Rigaku Saturn
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 28.5714 pixels mm⁻¹

ω scans

Absorption correction: multi-scan
(REQAB; Jacobson, 1998)

$T_{\min} = 0.825$, $T_{\max} = 0.863$

10628 measured reflections

2953 independent reflections

2565 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.039$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 2.3^\circ$

$h = -22 \rightarrow 22$

$k = -12 \rightarrow 10$

$l = -21 \rightarrow 21$

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.048$$

$$wR(F^2) = 0.110$$

$$S = 1.10$$

2953 reflections

195 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0536P)^2]$$
$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.34 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.47 \text{ e \AA}^{-3}$$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\text{sigma}(F^2)$ 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	-0.08924 (3)	0.2500	0.04422 (17)
C11	0.61483 (7)	-0.22781 (11)	0.27047 (8)	0.0670 (3)
O1	0.6047 (2)	0.1489 (3)	0.2577 (2)	0.0742 (9)
O2	0.3968 (2)	0.4328 (3)	0.0331 (2)	0.0822 (11)
N1	0.46860 (17)	0.0215 (3)	0.12856 (19)	0.0401 (7)
N2	0.46152 (18)	0.1695 (3)	0.0334 (2)	0.0439 (7)
C1	0.4133 (2)	-0.0189 (3)	0.0476 (2)	0.0400 (9)
C2	0.3666 (2)	-0.1304 (4)	0.0213 (3)	0.0487 (10)
H2A	0.3693	-0.1927	0.0605	0.058*
C3	0.3166 (2)	-0.1443 (4)	-0.0649 (3)	0.0552 (11)
H3A	0.2846	-0.2173	-0.0842	0.066*
C4	0.3126 (3)	-0.0521 (5)	-0.1240 (3)	0.0613 (12)
H4A	0.2782	-0.0653	-0.1818	0.074*
C5	0.3582 (3)	0.0584 (4)	-0.0994 (3)	0.0541 (11)
H5A	0.3556	0.1202	-0.1388	0.065*
C6	0.4083 (2)	0.0723 (3)	-0.0121 (3)	0.0426 (9)
C7	0.4950 (2)	0.1349 (3)	0.1171 (2)	0.0407 (8)
C8	0.5529 (2)	0.2124 (4)	0.1863 (3)	0.0476 (9)
C9	0.5623 (3)	0.3387 (4)	0.1971 (3)	0.0574 (11)
H9A	0.5330	0.4016	0.1576	0.069*
C10	0.6258 (3)	0.3579 (6)	0.2805 (4)	0.0835 (17)
H10A	0.6469	0.4363	0.3060	0.100*
C11	0.6496 (3)	0.2440 (7)	0.3154 (3)	0.0882 (17)
H11A	0.6905	0.2294	0.3705	0.106*
C12	0.4755 (2)	0.2850 (4)	-0.0060 (3)	0.0520 (10)

H12A	0.4817	0.2609	-0.0564	0.062*
H12B	0.5257	0.3254	0.0342	0.062*
C13	0.4077 (3)	0.3782 (4)	-0.0311 (3)	0.0532 (11)
C14	0.3522 (3)	0.4196 (4)	-0.1061 (3)	0.0719 (14)
H14A	0.3473	0.3972	-0.1597	0.086*
C15	0.3012 (3)	0.5054 (5)	-0.0888 (4)	0.0888 (19)
H15A	0.2564	0.5498	-0.1290	0.107*
C16	0.3296 (4)	0.5097 (5)	-0.0057 (5)	0.097 (2)
H16A	0.3073	0.5581	0.0232	0.117*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cd1	0.0472 (3)	0.0419 (3)	0.0376 (2)	0.000	0.01378 (19)	0.000
C11	0.0685 (7)	0.0712 (8)	0.0582 (7)	0.0256 (6)	0.0256 (6)	0.0060 (6)
O1	0.076 (2)	0.082 (2)	0.057 (2)	0.0048 (19)	0.0234 (18)	0.0035 (18)
O2	0.103 (3)	0.079 (2)	0.077 (2)	0.046 (2)	0.052 (2)	0.0202 (18)
N1	0.0410 (17)	0.0379 (18)	0.0373 (17)	0.0028 (14)	0.0141 (14)	0.0035 (13)
N2	0.0451 (18)	0.0402 (18)	0.0467 (19)	0.0071 (15)	0.0209 (15)	0.0041 (15)
C1	0.037 (2)	0.042 (2)	0.038 (2)	0.0097 (16)	0.0150 (17)	0.0002 (16)
C2	0.044 (2)	0.047 (2)	0.053 (2)	0.0025 (18)	0.0194 (19)	-0.0037 (19)
C3	0.046 (2)	0.061 (3)	0.050 (3)	0.001 (2)	0.013 (2)	-0.017 (2)
C4	0.053 (3)	0.076 (3)	0.038 (2)	0.012 (2)	0.006 (2)	-0.010 (2)
C5	0.053 (3)	0.065 (3)	0.042 (2)	0.018 (2)	0.019 (2)	0.008 (2)
C6	0.040 (2)	0.045 (2)	0.042 (2)	0.0131 (17)	0.0170 (17)	0.0022 (17)
C7	0.040 (2)	0.039 (2)	0.041 (2)	0.0060 (17)	0.0168 (17)	0.0031 (16)
C8	0.043 (2)	0.052 (2)	0.049 (2)	-0.0006 (19)	0.0220 (19)	0.0009 (19)
C9	0.062 (3)	0.040 (2)	0.063 (3)	-0.012 (2)	0.021 (2)	-0.003 (2)
C10	0.080 (4)	0.085 (4)	0.084 (4)	-0.045 (3)	0.035 (3)	-0.037 (3)
C11	0.060 (3)	0.134 (5)	0.056 (3)	-0.006 (4)	0.013 (3)	-0.018 (4)
C12	0.057 (3)	0.049 (2)	0.057 (3)	0.0098 (19)	0.033 (2)	0.0160 (19)
C13	0.059 (3)	0.040 (2)	0.065 (3)	0.0088 (19)	0.031 (2)	0.009 (2)
C14	0.079 (4)	0.052 (3)	0.066 (3)	0.014 (2)	0.016 (3)	0.002 (2)
C15	0.074 (4)	0.051 (3)	0.111 (5)	0.021 (3)	0.015 (4)	0.008 (3)
C16	0.099 (5)	0.077 (4)	0.125 (6)	0.045 (3)	0.059 (4)	0.012 (4)

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

Cd1—N1 ⁱ	2.252 (3)	C4—H4A	0.9300
Cd1—N1	2.252 (3)	C5—C6	1.389 (5)
Cd1—Cl1 ⁱ	2.4513 (12)	C5—H5A	0.9300
Cd1—Cl1	2.4513 (12)	C7—C8	1.447 (5)
O1—C8	1.354 (5)	C8—C9	1.334 (5)
O1—C11	1.392 (6)	C9—C10	1.414 (7)
O2—C13	1.352 (5)	C9—H9A	0.9300
O2—C16	1.372 (6)	C10—C11	1.319 (7)
N1—C7	1.330 (5)	C10—H10A	0.9300
N1—C1	1.386 (4)	C11—H11A	0.9300

N2—C7	1.356 (5)	C12—C13	1.484 (5)
N2—C6	1.387 (5)	C12—H12A	0.9700
N2—C12	1.469 (5)	C12—H12B	0.9700
C1—C6	1.386 (5)	C13—C14	1.320 (6)
C1—C2	1.396 (5)	C14—C15	1.424 (7)
C2—C3	1.374 (6)	C14—H14A	0.9300
C2—H2A	0.9300	C15—C16	1.305 (8)
C3—C4	1.390 (6)	C15—H15A	0.9300
C3—H3A	0.9300	C16—H16A	0.9300
C4—C5	1.378 (6)		
N1 ⁱ —Cd1—N1	118.17 (15)	N1—C7—C8	123.6 (3)
N1 ⁱ —Cd1—Cl1 ⁱ	106.27 (8)	N2—C7—C8	124.0 (3)
N1—Cd1—Cl1 ⁱ	109.08 (8)	C9—C8—O1	111.2 (4)
N1 ⁱ —Cd1—Cl1	109.08 (8)	C9—C8—C7	132.2 (4)
N1—Cd1—Cl1	106.27 (8)	O1—C8—C7	116.4 (4)
Cl1 ⁱ —Cd1—Cl1	107.57 (6)	C8—C9—C10	106.3 (4)
C8—O1—C11	105.1 (4)	C8—C9—H9A	126.8
C13—O2—C16	105.9 (4)	C10—C9—H9A	126.8
C7—N1—C1	105.5 (3)	C11—C10—C9	107.3 (4)
C7—N1—Cd1	130.1 (2)	C11—C10—H10A	126.4
C1—N1—Cd1	124.4 (2)	C9—C10—H10A	126.4
C7—N2—C6	106.5 (3)	C10—C11—O1	110.1 (5)
C7—N2—C12	129.4 (3)	C10—C11—H11A	125.0
C6—N2—C12	124.1 (3)	O1—C11—H11A	125.0
N1—C1—C6	109.2 (3)	N2—C12—C13	112.1 (3)
N1—C1—C2	130.6 (4)	N2—C12—H12A	109.2
C6—C1—C2	120.2 (4)	C13—C12—H12A	109.2
C3—C2—C1	117.3 (4)	N2—C12—H12B	109.2
C3—C2—H2A	121.4	C13—C12—H12B	109.2
C1—C2—H2A	121.4	H12A—C12—H12B	107.9
C2—C3—C4	121.8 (4)	C14—C13—O2	110.3 (4)
C2—C3—H3A	119.1	C14—C13—C12	133.0 (5)
C4—C3—H3A	119.1	O2—C13—C12	116.7 (4)
C3—C4—C5	121.8 (4)	C13—C14—C15	106.7 (5)
C3—C4—H4A	119.1	C13—C14—H14A	126.7
C5—C4—H4A	119.1	C15—C14—H14A	126.7
C4—C5—C6	116.2 (4)	C16—C15—C14	106.5 (5)
C4—C5—H5A	121.9	C16—C15—H15A	126.7
C6—C5—H5A	121.9	C14—C15—H15A	126.7
N2—C6—C1	106.3 (3)	C15—C16—O2	110.5 (5)
N2—C6—C5	130.9 (4)	C15—C16—H16A	124.7
C1—C6—C5	122.7 (4)	O2—C16—H16A	124.7
N1—C7—N2	112.4 (3)		
N1 ⁱ —Cd1—N1—C7	31.3 (3)	Cd1—N1—C7—C8	-0.1 (5)
Cl1 ⁱ —Cd1—N1—C7	152.8 (3)	C6—N2—C7—N1	1.3 (4)
Cl1—Cd1—N1—C7	-91.5 (3)	C12—N2—C7—N1	-178.8 (3)

N1 ⁱ —Cd1—N1—C1	−147.1 (3)	C6—N2—C7—C8	−178.6 (4)
C11 ⁱ —Cd1—N1—C1	−25.7 (3)	C12—N2—C7—C8	1.3 (6)
C11—Cd1—N1—C1	90.0 (3)	C11—O1—C8—C9	−0.9 (5)
C7—N1—C1—C6	0.8 (4)	C11—O1—C8—C7	−176.4 (4)
Cd1—N1—C1—C6	179.6 (2)	N1—C7—C8—C9	−147.9 (5)
C7—N1—C1—C2	−179.6 (4)	N2—C7—C8—C9	32.1 (7)
Cd1—N1—C1—C2	−0.8 (5)	N1—C7—C8—O1	26.4 (6)
N1—C1—C2—C3	−179.7 (4)	N2—C7—C8—O1	−153.6 (4)
C6—C1—C2—C3	−0.2 (5)	O1—C8—C9—C10	1.3 (5)
C1—C2—C3—C4	0.4 (6)	C7—C8—C9—C10	175.8 (5)
C2—C3—C4—C5	−0.3 (7)	C8—C9—C10—C11	−1.2 (6)
C3—C4—C5—C6	0.0 (6)	C9—C10—C11—O1	0.6 (7)
C7—N2—C6—C1	−0.7 (4)	C8—O1—C11—C10	0.1 (6)
C12—N2—C6—C1	179.4 (3)	C7—N2—C12—C13	−103.6 (5)
C7—N2—C6—C5	179.8 (4)	C6—N2—C12—C13	76.2 (5)
C12—N2—C6—C5	−0.1 (6)	C16—O2—C13—C14	1.3 (6)
N1—C1—C6—N2	0.0 (4)	C16—O2—C13—C12	−177.5 (4)
C2—C1—C6—N2	−179.7 (3)	N2—C12—C13—C14	−112.3 (6)
N1—C1—C6—C5	179.5 (3)	N2—C12—C13—O2	66.1 (5)
C2—C1—C6—C5	−0.1 (6)	O2—C13—C14—C15	−0.9 (6)
C4—C5—C6—N2	179.6 (4)	C12—C13—C14—C15	177.6 (5)
C4—C5—C6—C1	0.2 (6)	C13—C14—C15—C16	0.2 (7)
C1—N1—C7—N2	−1.3 (4)	C14—C15—C16—O2	0.6 (7)
Cd1—N1—C7—N2	180.0 (2)	C13—O2—C16—C15	−1.2 (7)
C1—N1—C7—C8	178.6 (4)		

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

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

$D—H\cdots A$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
C5—H5A ⁱⁱ —Cl1 ⁱⁱ	0.93	2.82	3.694 (5)	156
C9—H9A ⁱⁱ —O2	0.93	2.49	3.256 (6)	140

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