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## Key indicators

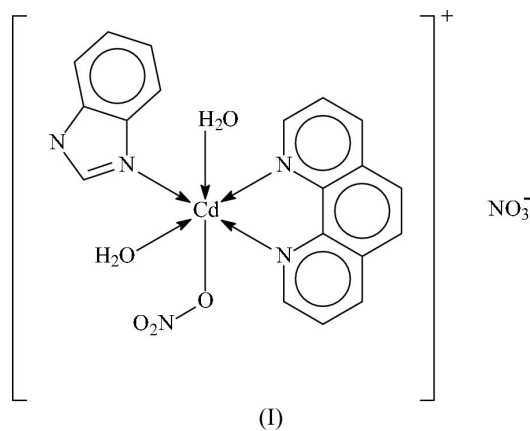
Single-crystal X-ray study  
 $T = 295$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.008$  Å  
 $R$  factor = 0.046  
 $wR$  factor = 0.137  
Data-to-parameter ratio = 15.1For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Diaqua(benzimidazole)(nitrate- $\kappa O$ )(1,10-phenanthroline)cadmium(II) nitrateIn the crystal structure of the title compound,  $[\text{Cd}(\text{NO}_3)(\text{C}_7\text{H}_6\text{N}_2)(\text{C}_{12}\text{H}_8\text{N}_2)(\text{H}_2\text{O})_2]\text{NO}_3$ , the phenanthroline-chelated Cd atom is bonded to the benzimidazole, two water molecules and a monodentate nitrate group in a six-coordinate octahedral geometry. The cation interacts with the nitrate anion to form a three-dimensional hydrogen-bonding network.

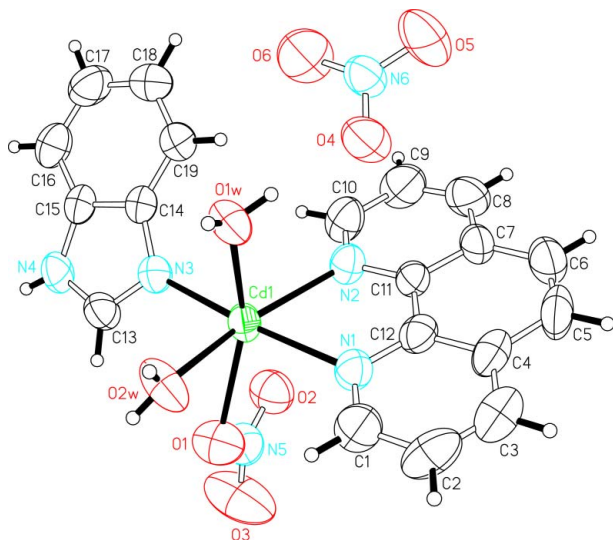
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## Comment

Cadmium salts, such as cadmium carboxylates, yield adducts with 1,10-phenanthroline in which the heterocycle binds in the typical chelating mode, and a number of such complexes can be found in the Cambridge Structural Database (Version 5.26; Allen, 2002). Cadmium nitrate affords only two complexes, (nitrate- $\kappa^2 O, O'$ )(nitrate- $\kappa O$ )bis(1,10-phenanthroline)-cadmium (Shi *et al.*, 2004) and di(nitrate- $\kappa^2 O, O'$ )bis(1,10-phenanthroline) (Tadjarodi *et al.*, 2001), probably because the two ligands present severe crowding around the metal atom.The present study used a hydrothermal method to synthesize a cadmium nitrate adduct with both 1,10-phenanthroline and benzimidazole as donor ligands. Monodentate benzimidazole has been used to bind to, for example, cadmium succinate (Liu & Xu, 2004*a*), cadmium phthalate (Liu & Xu, 2004*b*) and the cadmium derivative of pentaazapentacycloheptacosatriecaene (Sessler *et al.*, 1989). The synthesis yielded diaqua(benzimidazole)(nitrate- $\kappa O$ )(1,10-phenanthroline)cadmium(II) nitrate, (I), in which the nitrate anion interacts indirectly with the Cd atom through the coordinated water molecules. In this mixed-ligand compound, the Cd atom in the cation exists in an octahedral environment; the donor atoms of the ligands and a water molecule comprise a square around it. The second water molecule and the O atom of the monodentate nitrate group occupy the other two positions



**Figure 1**  
ORTEP plot (Johnson, 1976) of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms as spheres of arbitrary radii.

(Fig. 1). The cation and anion interact through H atoms to furnish a tightly held three-dimensional network (Table 2).

## Experimental

The title compound was synthesized hydrothermally from cadmium nitrate dihydrate (1 mmol, 0.27 g), 1,10-phenanthroline (2 mmol, 0.36 g), benzimidazole (2 mmol, 0.24 g) and water (20 ml). The reagents were heated in a 30 ml Teflon-lined stainless steel Parr bomb at 426 K for 5 d. The bomb was slowly cooled to room temperature to yield colourless crystals that were collected and washed with water.

### Crystal data

$[\text{Cd}(\text{NO}_3)(\text{C}_7\text{H}_6\text{N}_2)(\text{C}_{12}\text{H}_8\text{N}_2)(\text{H}_2\text{O})_2]\text{NO}_3$	$D_x = 1.740 \text{ Mg m}^{-3}$
$M_r = 570.79$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 5541 reflections
$a = 7.5119(4) \text{ \AA}$	$\theta = 2.4\text{--}26.8^\circ$
$b = 16.9551(9) \text{ \AA}$	$\mu = 1.06 \text{ mm}^{-1}$
$c = 17.1127(9) \text{ \AA}$	$T = 295(2) \text{ K}$
$\beta = 90.821(1)^\circ$	Block, colourless
$V = 2179.3(2) \text{ \AA}^3$	$0.32 \times 0.24 \times 0.21 \text{ mm}$
$Z = 4$	

### Data collection

Bruker SMART APEX area-detector diffractometer	4878 independent reflections
$\varphi$ and $\omega$ scans	4191 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 2002)	$R_{\text{int}} = 0.029$
$T_{\text{min}} = 0.336$ , $T_{\text{max}} = 0.808$	$\theta_{\text{max}} = 27.5^\circ$
13 046 measured reflections	$h = -8 \rightarrow 9$
	$k = -19 \rightarrow 21$
	$l = -17 \rightarrow 21$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0611P)^2 + 2.947P]$
$R[F^2 > 2\sigma(F^2)] = 0.046$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.137$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.13$	$\Delta\rho_{\text{max}} = 0.80 \text{ e \AA}^{-3}$
4878 reflections	$\Delta\rho_{\text{min}} = -0.85 \text{ e \AA}^{-3}$
322 parameters	
H atoms treated by a mixture of independent and constrained refinement	

**Table 1**

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

Cd1—O1	2.405 (4)	Cd1—N1	2.343 (4)
Cd1—O1w	2.345 (4)	Cd1—N2	2.348 (4)
Cd1—O2w	2.316 (3)	Cd1—N3	2.246 (4)
O1—Cd1—O1w	151.9 (1)	O1w—Cd1—N3	90.0 (1)
O1—Cd1—O2w	71.7 (1)	O2w—Cd1—N1	90.3 (2)
O1—Cd1—N1	89.9 (2)	O2w—Cd1—N2	158.7 (2)
O1—Cd1—N2	117.2 (1)	O2w—Cd1—N3	96.1 (2)
O1—Cd1—N3	92.0 (2)	N1—Cd1—N2	71.2 (1)
O1w—Cd1—O2w	80.2 (1)	N1—Cd1—N3	173.6 (1)
O1w—Cd1—N1	91.2 (1)	N2—Cd1—N3	102.6 (1)
O1w—Cd1—N2	89.7 (1)		

**Table 2**

Hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D\text{—}H\cdots A$	$D\text{—}H$	$H\cdots A$	$D\cdots A$	$D\text{—}H\cdots A$
O1w—H1w1 $\cdots$ O4	0.848 (10)	1.873 (13)	2.717 (5)	173 (5)
O1w—H1w2 $\cdots$ O2 <sup>i</sup>	0.849 (10)	1.975 (12)	2.824 (5)	179 (6)
O2w—H2w1 $\cdots$ O6 <sup>ii</sup>	0.843 (10)	1.96 (2)	2.775 (5)	163 (5)
O2w—H2w2 $\cdots$ O3 <sup>i</sup>	0.844 (10)	1.883 (15)	2.721 (6)	172 (7)
N4—H4n $\cdots$ O4 <sup>iii</sup>	0.848 (10)	2.03 (3)	2.789 (5)	149 (5)

Symmetry codes: (i)  $1 + x, y, z$ ; (ii)  $\frac{3}{2} - x, y - \frac{1}{2}, \frac{3}{2} - z$ ; (iii)  $x - \frac{1}{2}, \frac{3}{2} - y, \frac{1}{2} + z$ .

The aromatic H atoms were placed at calculated positions ( $C\text{—}H = 0.93 \text{ \AA}$ ) and were included in the refinement in the riding-model approximation, with  $U_{\text{iso}}(H)$  values tied to the  $U_{\text{eq}}$  of the parent atoms by a factor of 1.2. The water and amino H atoms were located in difference Fourier maps and were refined, with distance restraints of  $O\text{—}H = N\text{—}H = 0.85(1) \text{ \AA}$  and  $H\cdots H = 1.39(1) \text{ \AA}$ ; their displacement parameters could not be satisfactorily refined and were instead similarly tied.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2002); software used to prepare material for publication: SHELXL97.

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