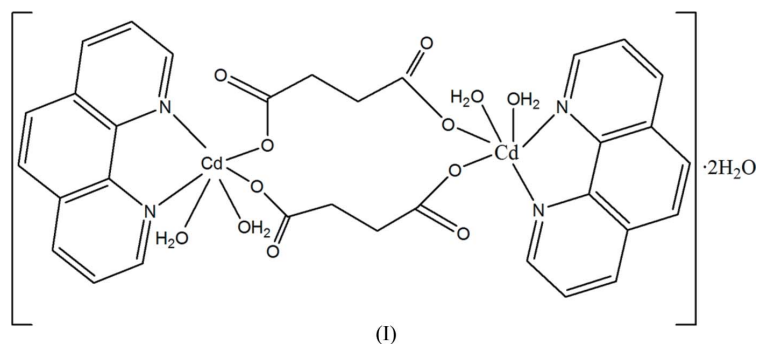


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Republic of ChinaCorrespondence e-mail:  
lixm20032006@yahoo.com.cn**Key indicators**Single-crystal X-ray study  
*T* = 292 K  
Mean  $\sigma(\text{C}-\text{C})$  = 0.003 Å  
*R* factor = 0.021  
*wR* factor = 0.056  
Data-to-parameter ratio = 14.2For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.**Di- $\mu$ -succinato- $\kappa^4\text{O}:\text{O}'$ -bis[*diaqua*(1,10-  
phenanthroline- $\kappa^2\text{N},\text{N}'$ )cadmium(II)]  
dihydrate**In the centrosymmetric title compound,  $[\text{Cd}_2(\text{C}_4\text{H}_4\text{O}_4)_2(\text{C}_{12}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_4]\cdot 2\text{H}_2\text{O}$ , the  $\text{Cd}^{\text{II}}$  atom is six-coordinated in an octahedral environment by two N atoms from one 1,10-phenanthroline, two O atoms from two different succinate and two water molecules. The crystal structure features  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds between carboxylate O atoms and water molecules as well as  $\pi-\pi$  stacking interactions between phenanthrolines.Received 16 November 2006  
Accepted 17 November 2006**Comment**Metal-organic complexes show a variety of supramolecular architectures (Eddaoudi *et al.*, 2001) and metal succinates are one such class. The succinate anion can connect metal ions to form one-, two- and three-dimensional supramolecular structures (Zheng *et al.*, 2000; Padmanabhan *et al.*, 2005; Ghoshal *et al.*, 2004). 1,10-Phenanthroline in its complexes gives rise to  $\pi-\pi$  interactions (Chen & Liu, 2002). These two features are combined in the dinuclear title compound, (I).The  $\text{Cd}^{\text{II}}$  atom is six-coordinated in an octahedral environment by two N atoms from a phenanthroline, two O atoms from two different succinates, and two water molecules (Fig. 1). The compound is dimeric, arising from bridging by the succinates across a center of inversion. A set of  $\pi-\pi$  interactions between the phenanthroline molecules, the shortest atom-to-atom distance being 3.68 (1) Å, leads to the formation of a linear chain structure. The supramolecular chains are linked into layers by  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds (Fig. 2 and Table 2).**Experimental**Compound (I) was prepared from a mixture of  $\text{Cd}(\text{CH}_3\text{COO})_2$  (0.133 g, 0.5 mmol),  $\text{H}_2\text{suc}$  (0.118 g, 1.0 mmol), phen (0.099 g, 0.5 mmol) and  $\text{H}_2\text{O}$  (18 ml) in a 30 ml Teflon-lined autoclave under autogenous pressure at 453 K for 3 d. After cooling to room

temperature, pale-yellow crystals suitable for X-ray structure analysis were obtained. Analysis calculated for  $C_{32}H_{36}Cd_2N_4O_{14}$ : C 72.7, H 5.1, N 14.1%; found: C 72.5, H 4.8, N 113.9%.

## Crystal data

$[Cd_2(C_4H_4O_4)_2(C_{12}H_8N_2)_2 \cdot (H_2O)_4] \cdot 2H_2O$   
 $M_r = 925.45$   
 Triclinic,  $P\bar{1}$   
 $a = 8.1775 (16) \text{ \AA}$   
 $b = 8.4059 (17) \text{ \AA}$   
 $c = 13.064 (3) \text{ \AA}$   
 $\alpha = 95.13 (3)^\circ$   
 $\beta = 104.18 (3)^\circ$

$\gamma = 104.86 (3)^\circ$   
 $V = 830.4 (4) \text{ \AA}^3$   
 $Z = 1$   
 $D_x = 1.851 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 $\mu = 1.36 \text{ mm}^{-1}$   
 $T = 292 (2) \text{ K}$   
 Block, pale yellow  
 $0.27 \times 0.23 \times 0.21 \text{ mm}$

## Data collection

Bruker APEX CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SAINT; Bruker, 1998)  
 $T_{\min} = 0.661$ ,  $T_{\max} = 0.755$

5179 measured reflections  
 3670 independent reflections  
 3585 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.011$   
 $\theta_{\text{max}} = 28.3^\circ$

## Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.021$   
 $wR(F^2) = 0.056$   
 $S = 1.13$   
 3670 reflections  
 259 parameters  
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0333P)^2 + 0.2504P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.003$   
 $\Delta\rho_{\text{max}} = 0.36 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.91 \text{ e \AA}^{-3}$

**Table 1**

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

Cd1—N1	2.3664 (17)	Cd1—O1W	2.3812 (15)
Cd1—N2	2.3661 (16)	Cd1—O2W	2.4204 (18)
Cd1—O1	2.2536 (15)	Cd1—O3 <sup>i</sup>	2.2567 (16)
O1—Cd1—O3 <sup>i</sup>	110.17 (6)	N2—Cd1—O1W	157.66 (6)
O1—Cd1—N2	95.59 (6)	N1—Cd1—O1W	86.82 (6)
O3 <sup>i</sup> —Cd1—N2	116.03 (6)	O1—Cd1—O2W	157.09 (7)
O1—Cd1—N1	87.23 (6)	O3 <sup>i</sup> —Cd1—O2W	79.63 (7)
O3 <sup>i</sup> —Cd1—N1	159.65 (6)	N2—Cd1—O2W	98.41 (7)
N2—Cd1—N1	70.84 (6)	N1—Cd1—O2W	80.39 (7)
O1—Cd1—O1W	83.12 (6)	O1W—Cd1—O2W	77.04 (7)
O3 <sup>i</sup> —Cd1—O1W	84.98 (6)		

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

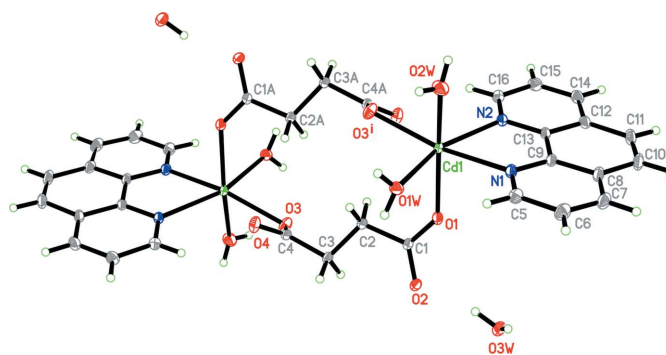
**Table 2**

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

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1W—HW12 $\cdots$ O2 <sup>ii</sup>	0.82 (3)	1.92 (3)	2.733 (2)	172 (3)
O1W—HW11 $\cdots$ O3 <sup>iii</sup>	0.76 (4)	2.16 (4)	2.855 (3)	153 (3)
O2W—HW21 $\cdots$ O3 <sup>iii</sup>	0.77 (3)	2.13 (3)	2.885 (3)	169 (3)
O2W—HW22 $\cdots$ O3W <sup>iv</sup>	0.78 (3)	1.94 (3)	2.721 (3)	173 (3)
O3W—HW31 $\cdots$ O2	0.84 (4)	1.94 (4)	2.759 (2)	163 (3)
O3W—HW32 $\cdots$ O4 <sup>ii</sup>	0.86 (3)	1.92 (3)	2.777 (2)	171 (3)

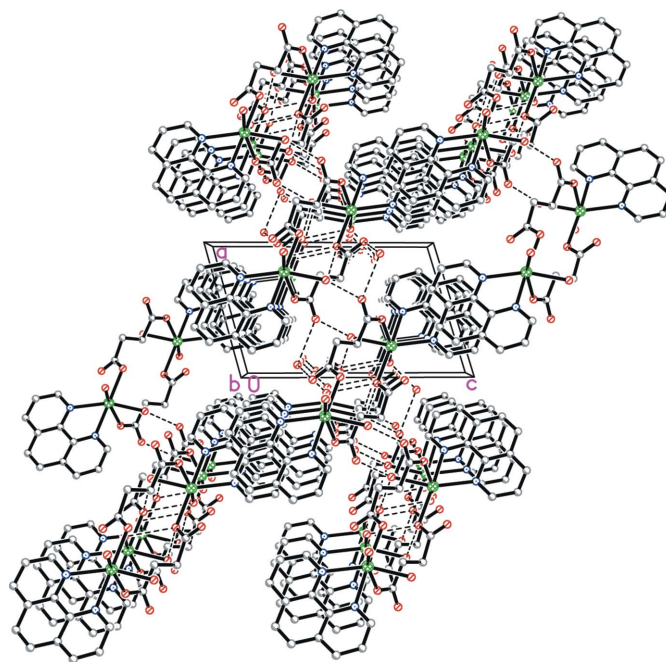
Symmetry codes: (ii)  $-x + 1, -y + 1, -z + 1$ ; (iii)  $x, y - 1, z$ ; (iv)  $x - 1, y - 1, z$ .

All H atoms on C atoms were positioned geometrically and refined as riding atoms, with  $C-H = 0.93 \text{ \AA}$  and  $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$ . The H



**Figure 1**

The molecular structure of (I), showing the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code: (i)  $-x, 1 - y, 1 - z$ .]



**Figure 2**

View of the two-dimensional layer, formed via  $\pi$ - $\pi$  stacking and hydrogen-bond interactions (dashed lines), along the  $b$  axis. H atoms have been omitted.

atoms of the water molecules were located in a difference Fourier map and were refined freely.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); 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, 1998); software used to prepare material for publication: SHELXTL.

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