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## Key indicators

Single-crystal X-ray study  
 $T = 293\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$   
 $R$  factor = 0.042  
 $wR$  factor = 0.104  
Data-to-parameter ratio = 15.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Bis[aquachlorobis(1,10-phenanthroline)-  
zinc(II)] benzene-1,4-dioxyacetate dihydrate

The title complex,  $[\text{ZnCl}(\text{C}_{12}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})]_2(\text{C}_{10}\text{H}_8\text{O}_6) \cdot 2\text{H}_2\text{O}$  ( $\text{C}_{10}\text{H}_8\text{O}_6 = \text{benzene-1,4-dioxyacetate}$  and  $\text{phen} = 1,10\text{-phenanthroline}$ ) consists of two  $[\text{ZnCl}(\text{phen})_2(\text{H}_2\text{O})]^{2+}$  cations, a benzene-1,4-dioxyacetate dianion and two water molecules. In the cation, the  $\text{Zn}^{\text{II}}$  atom is coordinated by four N atoms from two phen ligands, one  $\text{Cl}^-$  anion and one water molecule, forming a distorted octahedral coordination environment. The benzene-1,4-dioxyacetate dianion lies on an inversion center, with anions and cations forming one-dimensional chains through  $\text{O}-\text{H} \cdots \text{O}$  hydrogen bonds.

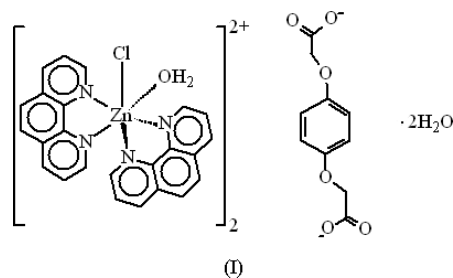
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## Comment

Recently, we have reported the crystal structure of the complex bis[aquachlorobis(1,10-phenanthroline)manganese(II)] benzene-1,4-dioxyacetate dihydrate (Gao *et al.*, 2004). The title zinc(II) analog, (I), was synthesized under similar reaction conditions and is reported here. A similar structural description of the Mn complex applies to the present isomorphous complex. As shown in Fig. 1, the  $\text{Zn}^{\text{II}}$  coordination is slightly distorted octahedral, defined by four N atoms from two phen ligands, one  $\text{Cl}^-$  anion and one water molecule. The Zn–N bond lengths are somewhat different from the Mn–N analogous ones, compared with 2.242 (2)–2.338 (2) Å in the Mn complex (Gao *et al.*, 2004), ranging from 2.263 (2) to 2.130 (2) Å. The water molecules form hydrogen bonds with carboxylate O atoms of adjacent anions and chloro ligands of adjacent cations, resulting in a one-dimensional chain (Table 2 and Fig. 2). In the crystal structure, the shortest ring-centroid to ring-centroid distances between symmetry-related phen ligands are 3.77 (4) and 3.78 (4) Å, and it is unlikely that these distances give rise to significant  $\pi$ – $\pi$  stacking interactions.



## Experimental

Benzene-1,4-dioxyacetic acid was prepared according to the method described for the synthesis of benzene-1,2-dioxyacetic acid by Mirci (1990). The title complex was prepared by the addition of phen (20 mmol) and zinc(II) acetate dihydrate (20 mmol) to an aqueous solution of benzene-1,4-dioxyacetic acid (40 mmol), and the pH was adjusted to 6 with 0.1 M sodium hydroxide. Colorless single crystals

were isolated from the filtered solution after several days. Analysis calculated for  $C_{58}H_{48}Cl_2N_8O_{10}Zn_2$ : C 62.95, H 4.37%; found: C 63.19, H 4.49%.

#### Crystal data

$[ZnCl(C_{12}H_8N_2)_2(H_2O)]_2 \cdot$   
 $(C_{10}H_8O_6) \cdot 2H_2O$   
 $M_r = 1218.72$   
 Triclinic,  $P\bar{1}$   
 $a = 9.950$  (6) Å  
 $b = 12.06$  (1) Å  
 $c = 12.81$  (1) Å  
 $\alpha = 61.95$  (3)°  
 $\beta = 80.65$  (4)°  
 $\gamma = 73.92$  (3)°  
 $V = 1302.6$  (17) Å<sup>3</sup>

$Z = 1$   
 $D_x = 1.554$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 6638 reflections  
 $\theta = 3.3$ – $27.5$ °  
 $\mu = 1.09$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 Prism, colorless  
 $0.37 \times 0.25 \times 0.18$  mm

#### Data collection

Rigaku R-Axis RAPID diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  
 $T_{min} = 0.687$ ,  $T_{max} = 0.827$   
 12 345 measured reflections

5884 independent reflections  
 4682 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.034$   
 $\theta_{max} = 27.5$ °  
 $h = -12 \rightarrow 11$   
 $k = -15 \rightarrow 15$   
 $l = -16 \rightarrow 16$

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.104$   
 $S = 1.04$   
 5884 reflections  
 373 parameters  
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0583P)^2 + 0.3253P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.001$   
 $\Delta\rho_{max} = 0.59$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.39$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

Zn1—Cl1	2.355 (2)	Zn1—N3	2.130 (2)
Zn1—N1	2.263 (2)	Zn1—N4	2.264 (2)
Zn1—N2	2.156 (2)	Zn1—O1W	2.138 (2)
N1—Zn1—Cl1	171.26 (5)	N3—Zn1—N4	76.07 (8)
N1—Zn1—N4	87.94 (8)	N3—Zn1—O1W	96.63 (8)
N2—Zn1—Cl1	97.48 (8)	N4—Zn1—Cl1	97.05 (7)
N2—Zn1—N1	75.33 (9)	O1W—Zn1—Cl1	92.24 (7)
N2—Zn1—N4	89.26 (8)	O1W—Zn1—N1	83.66 (8)
N3—Zn1—Cl1	97.56 (8)	O1W—Zn1—N2	95.68 (8)
N3—Zn1—N1	90.59 (9)	O1W—Zn1—N4	168.84 (7)
N3—Zn1—N2	160.13 (7)		

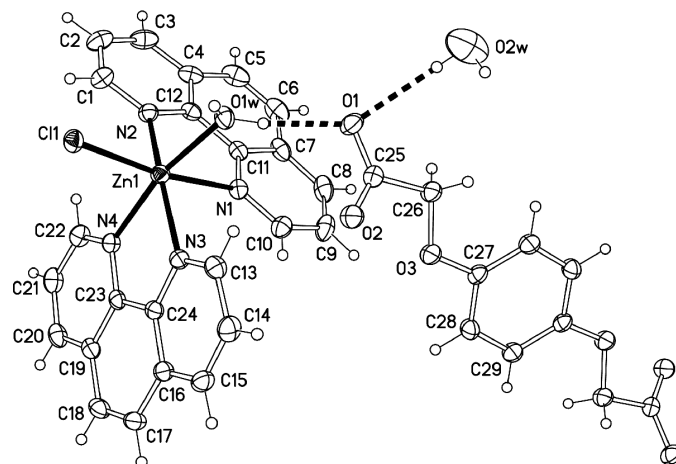
**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1W—H1W1 $\cdots$ O2 <sup>i</sup>	0.85 (4)	1.93 (3)	2.776 (3)	171 (3)
O1W—H1W2 $\cdots$ O1	0.84 (1)	1.94 (1)	2.778 (3)	172 (3)
O2W—H2W1 $\cdots$ Cl1 <sup>i</sup>	0.87 (7)	2.52 (7)	3.333 (5)	156 (7)
O2W—H2W2 $\cdots$ O1	0.87 (7)	2.22 (4)	3.013 (5)	152 (6)

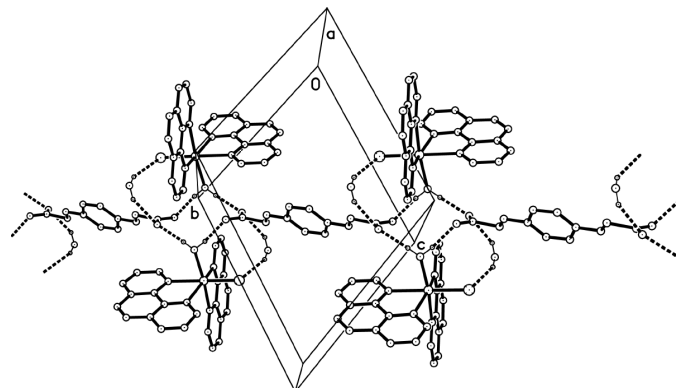
Symmetry code: (i)  $1 - x, -y, -z$ .

C-bound H atoms were placed in calculated positions [ $C-H = 0.93$  (aromatic) or  $0.97$  Å (aliphatic) and  $U_{iso}(H) = 1.2U_{eq}(C)$ ] in the riding-model approximation. The H atoms of water molecules were located from difference Fourier synthesis maps and refined with O—H and H $\cdots$ H distance restraints [ $0.85$  (1) and  $1.39$  (1) Å];  $U_{iso}(H) = 1.5U_{eq}(O)$ .



**Figure 1**

ORTEPII (Johnson, 1976) drawing of the title complex, showing 30% probability displacement ellipsoids. Hydrogen bonds are shown as dashed lines and the unlabeled atoms are related by the symmetry transformation  $(-x, 1 - y, 1 - z)$ .



**Figure 2**

The hydrogen-bonded chain structure of the title complex.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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