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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.004 Å R factor = 0.042 wR factor = 0.120 Data-to-parameter ratio = 13.0

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The one-dimensional zigzag coordination polymer *catena*-poly[[[triaquazinc(II)]- μ -N,N'-(benzene-1,4-dicarboxamido)diacetato- $\kappa^2 O$:O'] dihydrate]

In the crystal structure of the title compound, $\{[Zn(C_{12}H_{10}N_2O_6)(H_2O)_3]\cdot 2H_2O\}_n$, the Zn^{II} atom is located on a twofold axis and is coordinated by two carboxylate O atoms and three water O atoms in a trigonal-bipyramidal geometry. The diacetate ligands connect Zn^{II} atoms to generate one-dimensional zigzag chains, which are linked together by intermolecular hydrogen bonding to form a three-dimensional supramolecular framework.

Comment

Considerable effort has been devoted to the design and synthesis of self-assembled coordination polymers owing to their potential applications as functional materials (Janiak, 2003; James, 2003). An emerging approach in the rational assembly of coordination polymers is through the organic ligand as a spacer for linking metal atoms (O'Keeffe et al., 2000). Accordingly, the structures of coordination polymers can be ascribed to the coordination interactions between organic ligands and metal atoms, as well as other non-covalent interactions such as hydrogen bonding and π - π stacking (Moulton & Zaworotko, 2001). Moreover, the conformations of organic ligands also have important influences on the structures. To avoid the structural complication arising from the conformations of the ligands, rigid ligands, such as 1,4benzenedicarboxylate, are widely employed as spacer linkers (Eddaoudi et al., 2001). Recently, flexible ligands have been used to construct coordination polymers because they may induce some unusual structures (Burrows et al., 2004: Li et al., 2004). However, ligands with both rigidity and flexibility are seldom used to assemble coordination polymers. For example, an N-substituted aromatic amide has rigidity owing to the amide group (-CO-N-) conjugating with the aromatic ring, while the σ -rotation about the N-C bond could induce flexibility in the molecule. Moreover, the imine group N-H could serve as a better hydrogen-bonding donor and the amide C=O could also act as a better hydrogen-bonding acceptor (Manoj & Vijayan, 2000). Therefore, we have designed and synthesized a semi-flexible amide ligand, 2.2'-[1,4-bis(benzamido)]diacetic acid (H_2L), and employed it with Zn^{II} ions to assemble the title coordination polymer, (I).



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Figure 1

The molecular structure of (I), with displacement ellipsoids at the 25% probability level (arbitrary spheres for H atoms). Dashed lines indicate hydrogen bonds. [Symmetry codes: (i) 2 - x, y, $\frac{3}{2} - z$; (vi) 1 - x, -y, 1 - z.]

As shown in Fig. 1, the asymmetric unit of (I) consists of one-half of a ligand dianion, one-half of a Zn^{II} ion, one and a half coordinated water molecules, and one uncoordinated water molecule. The Zn^{II} and O3 atoms are located on a twofold axis. The Zn^{II} atom is coordinated by five O atoms in a trigonal-bipyramidal geometry, with Zn–O distances ranging from 1.981 (3) to 2.183 (2) Å (Table 1). Two carboxyl O atoms, O2 and O2ⁱ [symmetry code: (i) 2 - x, y, $\frac{3}{2} - z$], and one water molecule (O4) occupy the equatorial positions, while two water O atoms, O1 and O1ⁱ, are located at the axial sites. The axial O1···O1ⁱ vector is almost perpendicular to the equatorial plane. The C3-amide group is almost coplanar with the benzene ring, while the C1-carboxyl group lies out of the benzamide plane, with a dihedral angle of 74.7 (4)°.

The ligand dianion L is located on an inversion center and bridges neighboring Zn^{II} ions to generate one-dimensional zigzag chains propagating along the [201] direction. The small folds near the Zn^{II} ions in the chain can be ascribed to the flexibility arising from rotation about the N–C bond. The C3–N1–C2–C1 torsion angle is –72.7 (4)°.

The dihedral angle between neighboring benzamide planes in the same chain is $68.93 (16)^{\circ}$, and the dihedral angle between the equatorial plane and the benzamide plane is $70.1 (5)^{\circ}$. The Zn···Zn intrachain distance is 16.524 (4) Å, while the Zn···Zn interchain distance is 5.1898 (8) Å.

All chains are arranged in a parallel fashion. The chains are connected by intermolecular hydrogen bonding to form the three-dimensional supramolecular structure (Fig. 2). The shortest centroid–centroid distance of 5.189 (3) Å between parallel phenyl rings from neighboring chains indicates no π - π stacking interaction (Janiak, 2000). Atom H1*A* is involved in a three-centered hydrogen bond (Table 2).

Experimental

The ligand *L* was synthesized from terephthaloyl chloride and glycine according to a similar method reported by Yuan *et al.* (2002). *L* (28.0 mg, 0.1 mmol) was dissolved in an aqueous solution (10 ml) of $Zn(CH_3COO)_2$ ·4H₂O (22.0 mg, 0.1 mmol). The resulting colorless solution was filtered and the filtrate was left at room temperature. Colorless platelet crystals of (I) were obtained (20.1 mg, yield *ca* 46%) after several weeks by slow evaporation of the solvent.





A packing diagram of (I), viewed down the a axis. All H atoms have been omitted for clarity. Dashed lines indicate hydrogen bonds.

Crystal data

 $\begin{bmatrix} Zn(C_{12}H_{10}N_2O_6)(H_2O)_3 \end{bmatrix} \cdot 2H_2O \qquad A \\ M_r = 433.69 \qquad M \\ Monoclinic, C2/c \qquad G \\ a = 6.8980 (14) Å \\ b = 7.7550 (16) Å \qquad G \\ c = 30.339 (6) Å \qquad G \\ \beta = 93.44 (3)^\circ \qquad G \\ V = 1620.0 (6) Å^3 \qquad G \\ Z = 4 \qquad G \\ \end{bmatrix}$

Data collection

Enraf-Nonius CAD-4 diffractometer ω scans Absorption correction: ψ scan (XCAD4; Harms & Wocadlo, 1995) $T_{\min} = 0.458, T_{\max} = 0.850$ 1723 measured reflections

1587 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.042$ $wR(F^2) = 0.120$ S = 1.121587 reflections 122 parameters H atoms treated by a mixture of independent and constrained

independent and constrained refinement

 $D_x = 1.778 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 25 reflections $\theta = 2.5 - 15.0^{\circ}$ $\mu = 1.58 \text{ mm}^{-1}$ T = 298 (2) KPlate, colorless $0.50 \times 0.46 \times 0.11 \text{ mm}$

1372 reflections with $I > 2\sigma(I)$ $R_{int} = 0.051$ $\theta_{max} = 26.0^{\circ}$ $h = -8 \rightarrow 0$ $k = 0 \rightarrow 9$ $l = -37 \rightarrow 37$ 3 standard reflections every 200 reflections intensity decay: 0.9%

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0961P)^2 \\ &+ 4.1191P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{max} < 0.001 \\ \Delta\rho_{max} = 0.41 \ e \ \text{\AA}^{-3} \\ \Delta\rho_{min} = -0.62 \ e \ \text{\AA}^{-3} \end{split}$$

Table 1

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

Zn-O1	2.183 (2)	Zn-O4	1.981 (3)
Zn-O2	1.988 (2)		
O4-Zn-O2	127.17 (7)	$O2^i - Zn - O1$	88.65 (9)
$O2^{i}-Zn-O2$	105.65 (13)	O2-Zn-O1	92.41 (9)
O4-Zn-O1	89.12 (6)	$O1-Zn-O1^{i}$	178.24 (11)

Symmetry code: (i) $-x + 2, y, -z + \frac{3}{2}$.

Table 2	
Hydrogen-bond geometry (Å, °).	

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$	
$O1-H1A\cdots O2^{ii}$	0.85	1.93	2.772 (3)	170	
$O1 - H1B \cdot \cdot \cdot O6^{iii}$	0.85	1.93	2.744 (4)	161	
$N1 - H3 \cdots O6$	0.86	2.15	2.913 (4)	148	
$O4-H4\cdots O1^{ii}$	0.84(2)	1.95 (2)	2.771 (3)	165 (3)	
$O6-H6A\cdots O5^{iv}$	0.85	1.91	2.747 (3)	167	
$O6-H6B\cdots O3^{v}$	0.85	1.97	2.738 (3)	149	

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

H atoms of the O1 and O6 water molecules were located in a difference Fourier map, idealized, and refined as riding, with O-H = 0.85 Å. Atom H4 was located in a difference Fourier map and refined subject to an O-H restraint of 0.85 (3) Å and an H···H restraint of 1.385 (3) Å. Other H atoms were placed at calculated positions, with C-H = 0.93 Å (aromatic) and 0.97 Å (methylene), and N-H = 0.86 Å, and refined using a riding model. In all cases, the $U_{\rm iso}({\rm H})$ value is 1.2 times $U_{\rm eq}$ of the parent atom.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000); software used to prepare material for publication: *SHELXTL*.

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