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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.024 wR factor = 0.064 Data-to-parameter ratio = 41.8

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Poly[[diaquazinc(II)]- μ_2 -4,4'-bipyridine- $\kappa^2 N:N'$ - μ_2 -*p*-phenylenebis(oxyacetato)- $\kappa^2 O:O'$]

The title compound, $[Zn^{II}(C_8H_8O_6)(C_8H_8N_2)(H_2O)_2]_n$, contains a Zn atom coordinated by 4,4'-bipyridine, benzene-1,4-dioxydiacetate and two water molecules. The Zn atom lies on an inversion centre and the benzene-1,4-dioxydiacetate and 4,4-bipyridine ligands lie about other inversion centres.

Comment

The title compound, (I), contains a Zn atom coordinated by 4,4'-bipyridine, benzene-1,4-dioxydiacetate and two water molecules. The Zn atom lies on an inversion centre and the benzene-1,4-dioxydiacetate and 4,4-bipyridine ligands lie about other inversion centres. The benzene-1,4-dioxydiacetate ligands bridge the octahedral Zn^{II} coordination centres to form a one-dimensional zigzag chain. The chains are further bridged *via* 4,4-bipyridine, forming a two-dimensional architecture.



Hydrogen-bonding interactions between the coordinated water molecules and the carboxylate O atoms lead to the formation of a three-dimensional network structure (Table 1). Compound (I) is isostructural with its cobalt analogue (Dai *et al.*, 2005).

Experimental

Zn(OAc)₂·H₂O (0.5 mmol, 0.10 g), 4,4'-bpy (0.5 mmol, 0.078 g), benzene-1,4-dioxydiacetic acid (0.5 mmol, 0.113 g) and NaOH (0.04 g, 1 mmol) were mixed in H₂O (15 ml) and heated at 433 K for 3 d in a sealed 25 ml Teflon-lined stainless-steel vessel under autogenous pressure. After cooling to room temperature at 50 K h⁻¹, yellow block crystals of (I) were isolated, and were washed with ethanol and dried in air.

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metal-organic papers

Crystal data

$$\begin{split} & [Zn(C_8H_8O_6)(C_8H_8N_2)(H_2O)_2] \\ & M_r = 481.77 \\ & Triclinic, P\overline{1} \\ & a = 5.7279 \ (17) \ \text{\AA} \\ & b = 8.076 \ (3) \ \text{\AA} \\ & c = 10.668 \ (3) \ \text{\AA} \\ & \alpha = 105.924 \ (4)^{\circ} \\ & \beta = 96.726 \ (2)^{\circ} \\ & \gamma = 97.816 \ (4)^{\circ} \end{split}$$

Data collection

Bruker SMART 1K CCD areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.725, T_{\max} = 0.871$

Refinement

Refinement on F^2	H-atom parameters constrained		
$R[F^2 > 2\sigma(F^2)] = 0.024$	$w = 1/[\sigma^2(F_o^2) + (0.0424P)^2]$		
$wR(F^2) = 0.065$	where $P = (F_0^2 + 2F_c^2)/3$		
S = 1.02	$(\Delta/\sigma)_{\rm max} < 0.001$		
2131 reflections	$\Delta \rho_{\rm max} = 0.90 \ {\rm e} \ {\rm \AA}^{-3}$		
51 parameters	$\Delta \rho_{\rm min} = -0.90 \text{ e} \text{ Å}^{-3}$		

V = 463.9 (3) Å³

 $D_x = 1.724 \text{ Mg m}^{-3}$

 $0.30 \times 0.20 \times 0.10 \text{ mm}$

2644 measured reflections

2131 independent reflections

1953 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 1.38 \text{ mm}^{-1}$

T = 293 (2) K

Prism, yellow

 $\begin{array}{l} R_{\rm int}=0.020\\ \theta_{\rm max}=27.5^\circ\end{array}$

Z = 1

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$ \begin{array}{c} \hline O1W-H1WA\cdots O3^{i} \\ O1W-H1WB\cdots O2^{ii} \end{array} $	0.81 (2)	2.16 (2)	2.9090 (17)	153 (2)
	0.83 (2)	1.81 (2)	2.6084 (17)	163 (3)

Symmetry codes: (i) -x, -y, -z + 1; (ii) -x + 1, -y, -z + 1.

The organic H atoms were positioned geometrically and refined using a riding model, with C-H = 0.93 and 0.97 Å, and with $U_{iso}(H) =$ $1.2U_{eq}(C)$. The water H atoms were located in difference maps and refined isotropically, with the O-H distances restrained to 0.82 (2) Å and $U_{iso}(H) = 1.5U_{eq}(O)$.



Figure 1

A view of the title compound, showing the atomic labelling scheme and 50% probability displacement ellipsoids. [Symmetry codes: (i) 1 - x, -y, 1 - z; (ii) 1 - x, -1 - y, -z; (iii) -x, -y, -z; (iv) 1 + x, y, 1 + z; (v) x, 1 + y, 1 + z].

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1994); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Siemens, 1994); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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