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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.003 Å R factor = 0.028 wR factor = 0.072 Data-to-parameter ratio = 12.5

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

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The tetranuclear dianion of the title compound, Na₂[Zn₄(C₁₀H₂O₈)₂(OH)₂]·C₄H₄N₂, forms a network framework in which the Zn atom exists in a tetrahedral environment, as it is coordinated by the carboxyl O atoms belonging to three different $(C_{10}H_2O_8)^{4-}$ anions as well as by the bridging OH group. Adjacent OH groups are linked together through the uncoordinated heterocycle, which lies on a site of 2/*m* symmetry. The octahedrally coordinated Na ion, which lies on a site of 2 symmetry, interacts with one O atom of the carboxyl group of two anions and with two O atoms of the *ortho*-carboxyl groups of two other anions.

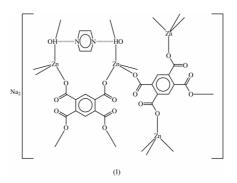
dihydroxytetrazincate(II) pyrazine

Disodium bis(1,2,4,5-benzenetetracarboxylato)-

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Comment

The 4,4'-bipyridine spacer molecule has been used in the formation of a plethora of adducts with zinc carboxylates; however, the spacer was not incorporated in hydrated disodium dihydrogen 1,2,4,5-benzenetetracarboxylatozinc (Wu *et al.*, 2001) when the synthesis was performed under acidic conditions. The use of pyrazine in the metathetical reaction of the sodium salt of this acid and a zinc salt did not lead to the formation of an N-coordinated Zn complex either. The title complex, (I), is formally a hydroxide (Fig. 1), and the heterocycle merely serves as a connector that links adjacent groups together.



The compound adopts a network framework in which the Zn atom exists in a tetrahedral geometry, being covalently bonded to the carboxyl O atoms belonging to three different $(C_{10}H_2O_8)^{4-}$ anions and to the bridging OH group. Adjacent OH groups are linked together through the uncoordinated lattice heterocycle (Fig. 2). The octahedrally coordinated Na ion interacts with one O atom of the carboxyl group of two anions and with two O atoms of the *ortho*-carboxyl groups of two other anions (Fig. 3).

In the assembly of coordination polymers the use of sodium hydroxide to deprotonate carboxylic acids sometimes

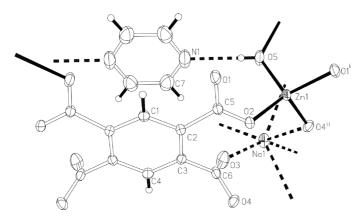


Figure 1

ORTEPII (Johnson, 1976) plot of a fragment of the structure of the title compound, with displacement ellipsoids drawn at the 75% probability level. H atoms are drawn as spheres of arbitrary radii. [Symmetry codes: (i) $\frac{1}{2} - x$, $\frac{1}{2} - y$, -z; (ii) $\frac{1}{2} - x$, $\frac{1}{2} + y$, 1 - z.]

furnishes compounds having sodium ions incorporated into the crystal structure (Robl, 1992; Wu *et al.*, 2001; Yang *et al.*, 2002) because the Na⁺ cation is able to form strong bonds with carboxylate groups or with water. Furthermore, it serves to compensate for the negative charge of the frameworks.

Experimental

1,2,4,5-Benzenetetracarboxylic acid anhydride (pyromellitic anhydride, 0.22 g, 1 mmol) and sodium hydroxide (0.16 g, 4 mmol) were dissolved in water (15 ml), and to the solution was added zinc dinitrate hexahydrate (0.59 g, 2 mmol) and pyrazine (0.16 g, 2 mmol) dissolved in water (5 ml). The mixture was placed in a 20 ml Teflon-lined stainless-steel bomb, which was then heated to 453 K for 100 h. The bomb was cooled to room temperature at 5 K h⁻¹. Block-shaped colorless crystals separated from the solution.

Crystal data

130 parameters

$M_{2}[Zn_{4}(C_{10}H_{2}O_{8})_{2}(OH)_{2}] \cdot C_{4}H_{4}N_{2}$ $M_{r} = 921.80$ Monoclinic, $C2/m$ a = 10.4073 (5) Å b = 17.8174 (8) Å c = 7.5278 (4) Å c = 0.474 (1) %	$D_x = 2.312 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 5344 reflections $\theta = 2.3-28.3^{\circ}$ $\mu = 3.71 \text{ mm}^{-1}$ T = 298 (2) K
$\beta = 108.474 \ (1)^{\circ}$ V = 1324.0 (1) Å ³	I = 298 (2) K Block, colorless
<i>Z</i> = 2	$0.29 \times 0.17 \times 0.12 \text{ mm}$
Data collection	
Bruker SMART APEX area- detector diffractometer φ and ω scans Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 1996) $T_{\min} = 0.463$, $T_{\max} = 0.641$ 7636 measured reflections	1629 independent reflections 1470 reflections with $I > 2\sigma(I)$ $R_{int} = 0.035$ $\theta_{max} = 28.3^{\circ}$ $h = -13 \rightarrow 13$ $k = -23 \rightarrow 23$ $l = -9 \rightarrow 9$
Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.072$ S = 1.03 1629 reflections	All H-atom parameters refined $w = 1/[\sigma^2(F_o^2) + (0.0429P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.51 \text{ e} \text{ Å}^{-3}$

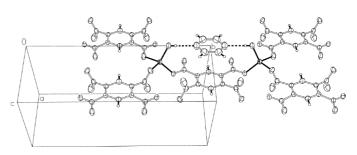


Figure 2

ORTEPII (Johnson, 1976) plot illustrating the hydroxyl-pyrazine-hydroxyl hydrogen bonds.

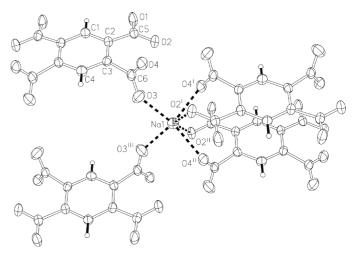


Figure 3

ORTEPII (Johnson, 1976) plot illustrating the coordination of the Na⁺ ion. [Symmetry codes: (i) $\frac{1}{2} - x$, $\frac{3}{2} - y$, 1 - z; (ii) $x - \frac{1}{2}$, $\frac{3}{2} - y$, z; (iii) -x, y, 1 - z.]

Table 1

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Selected geometric parameters (Å, °).
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-O1 ⁱ 1.973	$3(2)$ Na1 $-O2^{ii}$	2.542 (2)
-02 1.975	5 (2) Na1–O3	2.319 (2)
-O4 ⁱⁱ 1.960	$D(2)$ Na1 $-O4^{ii}$	2.406 (2)
-05 1.890	0(1)	
-Zn1-O2 120.3	$O2^{ii} - Na1 - O3^{iv}$	136.5 (1)
-Zn1-O4 ⁱⁱ 98.5	$5(1)$ $O2^{ii}-Na1-O4^{ii}$	73.7 (1)
-Zn1-O5 107.3	$O2^{ii} - Na1 - O4^{iii}$	79.6 (1)
-Zn1–O4 ⁱⁱ 113.8	3(1) O3-Na1-O3 ^{iv}	88.1 (1)
-Zn1-O5 108.1	1(1) O3-Na1-O4 ⁱⁱ	91.9 (1)
-Zn1-O5 108.2	2(1) O3-Na1-O4 ⁱⁱⁱ	148.8 (1)
-Na1-O2 ⁱⁱⁱ 136.2	$2(1)$ $O4^{ii}-Na1-O4^{iii}$	103.6 (1)
-Na1-O3 79.1	1 (1)	
-Na1-O3 79.1 metry codes: (i) $\frac{1}{2} - x, \frac{3}{2} - y, -$	()	-

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

Table	2	
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$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O5−H5···N1	0.84 (1)	1.88 (1)	2.720 (4)	180

(°))

The diffraction measurements were of sufficiently high quality to allow for the refinement of the four H atoms, three of which lie on

 $\Delta \rho_{\rm min} = -0.40 \text{ e } \text{\AA}^{-3}$

special positions, when restraints [O-H = 0.85 (1) Å and C-H 0.93 (1) Å] were applied.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2001); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP*II (Johnson, 1976); software used to prepare material for publication: *SHELXL*97.

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