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Hexaaquazinc(II) bis(6-hydroxypyridine-3-carboxylate)
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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{n}-\mathrm{O})=0.003 \AA$
Disorder in main residue
$R$ factor $=0.038$
$w R$ factor $=0.107$
Data-to-parameter ratio $=8.8$

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

The Zn atom in the title compound, $\left[\mathrm{Zn}\left(\mathrm{H}_{2} \mathrm{O}\right)_{6}\right]\left(\mathrm{C}_{6} \mathrm{H}_{4} \mathrm{NO}_{3}\right)_{2}$, lies on a special position of $2 / m$ site symmetry in an octahedron made up of water molecules. The anion is disordered across a mirror plane. The cation interacts with the anion indirectly through hydrogen bonds to form a threedimensional network.

## Comment

A number of syntheses of the zinc(II) derivatives of organic acids have yielded the hexaaquazinc(II) cation, the deprotonated acid remaining in the outer coordination sphere of the coordination octahedron as noted in, for example, the dihy-drogen-1,2,4,5-benzenetetracarboxylate (Sun et al., 2002), the bis[(4-oxo-4H-pyridin-1-yl)acetate] dihydrate (Gao et al., 2004), the 1,5 -naphthalenedisulfonate (Huo et al., 2005) and the 3-carboxy-4-hydroxybenzenesulfonate tetrahydrate (Ma et al., 2003). The reaction of the zinc cation and the 6-hydroxy-pyridine-3-carboxylate anion in a $1: 1$ molar stoichiometry yielded a similar compound, (I), with a hexaaquazinc(II) cation and an uncoordinated 6-hydroxpyridine-3-carboxylate anion (Fig. 1). As the O1w water molecule lies on a general position, whereas atom Zn 1 lies on a special position of $2 / m$ site symmetry, atom Zn 1 exists within a square made up of the four equivalent water molecules; the other two sites are occupied by the $\mathrm{O} 2 w$ (which lies on a mirror plane) and $\mathrm{O} 2 w^{\mathrm{ii}}$ water molecules [symmetry code: (ii) $1-x, 1-y, 1-z$ ]. The two ions interact through hydrogen bonds to lead to a tightly held three-dimensional framework (Table 2). The anion lies on a mirror plane bisecting the carboxyl group; its two $\mathrm{C}-\mathrm{O}$ distances are equal. The hydroxypyridine portion is disordered across this mirror plane. Pairs of adjacent anions are linked by hydrogen bonds to furnish a hydrogen-bonded dianion (Fig. 1).


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Figure 1
ORTEPII (Johnson, 1976) plot showing the numbering scheme for (I). Displacement ellipsoids are shown at the $50 \%$ probability level, and H atoms are drawn as spheres of arbitrary radii. [Symmetry codes: (i) $1-x, y, 1-z$; (ii) $1-x, 1-y, 1-z$; (iii) $x, 1-y, z$; (iv) $\frac{5}{2}-x, \frac{3}{2}-y, 2-z$.]

## Experimental

A mixture of zinc nitrate trihydrate $(0.298 \mathrm{~g}, 1 \mathrm{mmol})$, 6-hydroxy-pyridine-3-carboxylic acid ( $0.139 \mathrm{~g}, 1 \mathrm{mmol}$ ), sodium hydroxide $(0.040 \mathrm{~g}, 1 \mathrm{mmol})$ and water $(10 \mathrm{ml})$ were sealed in a 23 ml Teflonlined stainless steel Parr bomb. The bomb was heated to 433 K for 2 d . It was cooled to room temperature at $10 \mathrm{~K} \mathrm{~h}^{-1}$ to yield colorless crystals.

## Crystal data

$\left[\mathrm{Zn}\left(\mathrm{H}_{2} \mathrm{O}\right)_{6}\right]\left(\mathrm{C}_{6} \mathrm{H}_{4} \mathrm{NO}_{3}\right)_{2}$
$M_{r}=449.67$
Monoclinic, $C 2 / m$
$a=11.568$ (1) A
$b=9.780$ (1) $\AA$
$c=7.6255$ (7) $\AA$
$\beta=91.286(3)^{\circ}{ }^{\circ}$
$V=862.5$ (1) $\AA^{3}$
$Z=2$
$D_{x}=1.732 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 1640 reflections
$\theta=2.7-27.5^{\circ}$
$\mu=1.49 \mathrm{~mm}^{-1}$
$T=295$ (2) K
Block, colorless
$0.34 \times 0.31 \times 0.24 \mathrm{~mm}$

## Data collection

Bruker APEX area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\text {min }}=0.211, T_{\text {max }}=0.716$
2428 measured reflections

$$
\begin{aligned}
& 992 \text { independent reflections } \\
& 979 \text { reflections with } I>2 \sigma(I) \\
& R_{\text {int }}=0.033 \\
& \theta_{\max }=27.5^{\circ} \\
& h=-14 \rightarrow 14 \\
& k=-12 \rightarrow 12 \\
& l=-7 \rightarrow 9
\end{aligned}
$$

## Refinement

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Refinement on \(F^{2}\)
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$w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0522 P)^{2}\right.$
$+1.9207 P]$
where $P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}^{2}\right) / 3$
$(\Delta / \sigma)_{\text {max }}=0.001$
$\Delta \rho_{\max }=0.40 \mathrm{e}^{-3}$
$\Delta \rho_{\min }=-0.35 \mathrm{e}^{-3}$

Table 2
Hydrogen-bond geometry ( $\AA$, ${ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O} 1 w-\mathrm{H} 1 w 1 \cdots \mathrm{O} 1$ | $0.84(1)$ | $1.82(1)$ | $2.655(3)$ | $174(3)$ |
| $\mathrm{O} 1 w-\mathrm{H} 1 w 2 \cdots 1^{\text {viii }}$ | $0.84(1)$ | $1.89(1)$ | $2.706(3)$ | $166(3)$ |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 1 \cdots \mathrm{O}^{\mathrm{v}}$ | $0.85(1)$ | $2.01(2)$ | $2.788(5)$ | $153(3)$ |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 1 \cdots \mathrm{O}^{\mathrm{v}}$ | $0.85(1)$ | $2.01(2)$ | $2.788(5)$ | $153(3)$ |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 2 \cdots \mathrm{O}^{\text {vi }}$ | $0.85(1)$ | $1.96(1)$ | $2.758(6)$ | $157(1)$ |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 2 \cdots \mathrm{O}^{\text {vii }}$ | $0.85(1)$ | $1.96(1)$ | $2.758(6)$ | $157(1)$ |
| $\mathrm{O} 2-\mathrm{H} 2 \mathrm{o} \cdots \mathrm{N}^{\mathrm{iv}}$ | 0.85 | 2.03 | $2.86(1)$ | 165 |
| Symmetry codes: | (iv) | $-x+\frac{5}{2},-y+\frac{3}{2},-z+2 ;$ | (v) $x-1, y, z-1 ;$ | (vi) |
| $-x+2,-y+1,-z+1 ;$ (vii) $-x+2, y,-z+1 ;\left(\right.$ (viii) $-x+\frac{3}{2},-y+\frac{3}{2},-z+1$. |  |  |  |  |

The carboxylate anion is disordered over a mirror plane. The $\mathrm{C}-\mathrm{C}$ distances were restrained to 1.39 (1) $\AA$, and the two $\mathrm{N}-\mathrm{C}$ distances were restrained to within $0.01 \AA$ of each other. Additionally, the ring was restrained to near planarity. C-bound H atoms were placed at calculated positions $(\mathrm{C}-\mathrm{H}=0.93 \AA)$ and were included in the refinement in the riding-model approximation, with $U_{\text {iso }}(\mathrm{H})$ set to $1.2 U_{\text {eq }}(\mathrm{C})$. The water H atoms were located in difference Fourier maps, and were refined with a distance restraint of $\mathrm{O}-\mathrm{H}=$ 0.85 (1) Å. The hydroxy group was allowed to rotate about the $\mathrm{C}-\mathrm{O}$ bond, with $\mathrm{O}-\mathrm{H}=0.85 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{O})$.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997; program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976).

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