

Bis(μ -8-hydroxy-7-iodoquinoline-5-sulfonato- $\kappa^3 N, O: O'$)bis[triaquazinc(II)] tetrahydrate

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Key indicators

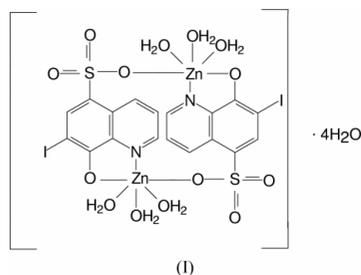
Single-crystal X-ray study
T = 293 K
Mean $\sigma(C-C)$ = 0.004 Å
R factor = 0.028
wR factor = 0.061
Data-to-parameter ratio = 17.0

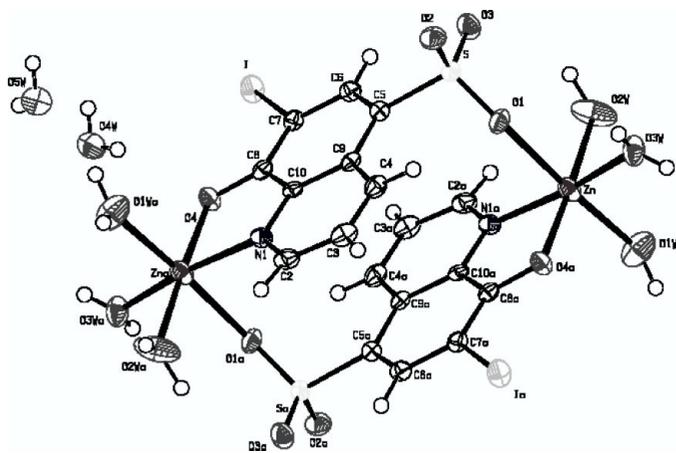
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

In the title compound, $[Zn_2(C_9H_4INO_4S)_2(H_2O)_6] \cdot 4H_2O$, the coordination geometry around the Zn^{II} atom is a distorted octahedron, formed by three water molecules, the N and deprotonated O atoms of the quinolinol ring and one of the O atoms of the sulfonate group. In the centrosymmetric binuclear complex, there is a π - π -stacking interaction between the quinolinol rings. The water molecules of crystallization play an important role in the hydrogen-bonding patterns of the three-dimensional network.

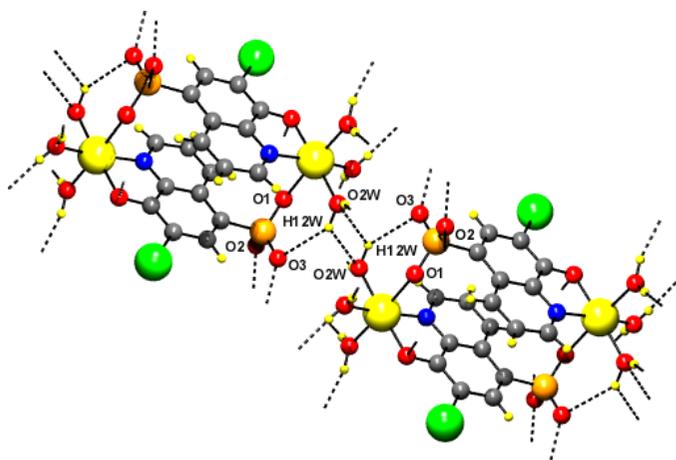
Comment

Oxine [8-hydroxyquinoline (HQ)] derivatives are well known analytical reagents and are also known for their anti-amoebic, antibacterial and antifungal activities, which could be correlated to their ability for bidentate chelation to metals. Many metal complexes of HQ have been reported (Palenik, 1964). The crystal structures of 8-hydroxy-7-nitroquinoline-5-sulfonic acid monohydrate (Balasubramanian & Muthiah, 1996) and 2-methyl-8-hydroxyquinoline-5-sulfonic acid monohydrate (Merrit & Duffin, 1970) have also been reported. As groups with coordination abilities are present on both sides, the sulfoxine ligand is capable of forming dimers. The iodo substitution at the 7-position in the sulfoxine ligand [7-iodo-8-hydroxyquinoline-5-sulfonic acid (ferron)] tends to result predominantly in isomorphous crystal structures with metal ions. Owing to its stability, a cage-like dimeric structure is formed rather than a polymeric network. Both types of crystal structure have already been reported (Petit, Coquerel & Perez, 1993; Petit, Ammor *et al.*, 1993) for the copper complex of sulfoxine [8-hydroxyquinoline-5-sulfonic acid (HQS)]. The structures of the cobalt complex of ferron (Balasubramanian, 1995), the nickel complex of ferron (Baskar Raj, Muthiah *et al.*, 2003), the lithium complex of HQS (Murugesan & Muthiah, 1997), the sodium complex of HQS (Baskar Raj *et al.*, 2002) and the nickel complex of HQS (Baskar Raj *et al.*, 2001) have been reported by our laboratory. The present study concerns the structure of the title zinc compound, (I).




Figure 1

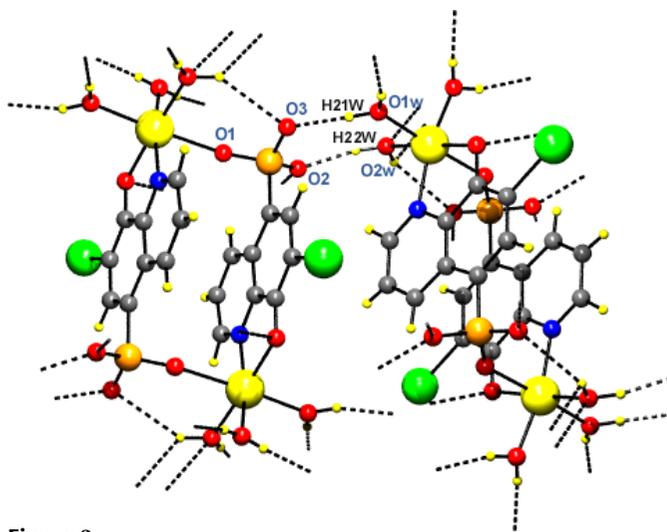
A view of the dimeric complex of (I), showing 50% probability displacement ellipsoids for non-H atoms.


Figure 2

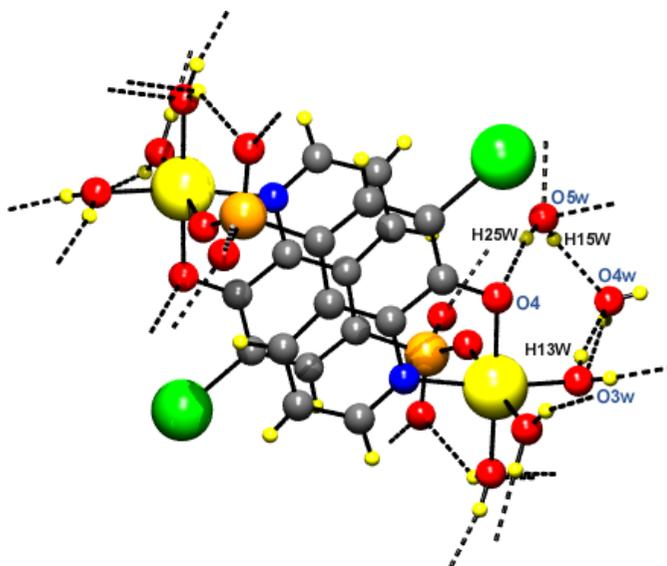
The tandem-type of hydrogen-bonding pattern in (I).

The asymmetric unit of (I) contains one monomeric unit of $[\text{Zn}(\text{ferron})(\text{H}_2\text{O})_3]$ and two uncoordinated water molecules. Two inversion-related ligands and two Zn atoms form a cage-like dimer (Fig. 1 and Table 1). The coordination of Zn is a distorted octahedron, the metal atom being surrounded by three water molecules, one of the O atoms of the sulfonate group (O1) and the N and O atoms of the oxine ring. In the octahedral geometry, two O atoms of coordinated water molecules (O2W and O3W) and N1, O4 of the oxine moiety occupy equatorial positions, and the sulfonate O atom and one O atom of a water molecule (O1W) are in axial positions. The Zn–N1, Zn–O and S–O bond distances and the N1–Zn–O4 bite angle $[80.56(9)^\circ]$ are very close to those found in the nickel complex of ferron (Baskar Raj, Muthiah *et al.*, 2003). In (I), π – π -stacking is observed between quinolinol rings within the dimer. The interplanar and centroid-to-centroid distances are 3.464 and 3.520 Å, respectively, and the slip angle (angle between the centroid vector and normal to the plane) is 9.45° .

The water molecules and the hetero O atoms play an important role in the formation of hydrogen-bonding patterns


Figure 3

A view of the fork-like hydrogen-bonding interaction in (I).


Figure 4

Hydrogen-bonding patterns involving water molecules of crystallization in (I).

(Table 2) and in the formation of supramolecular motifs. The non-coordinated O atom (O3) of the sulfonate group acts as an acceptor to the coordinated water molecule (O2W), forming a six-membered ring within the dimer. The tandem hydrogen bonding between each of the two coordinated water molecules of the adjacent dimeric units forms an $R_2^2(4)$ ring motif (Fig. 2). Atom O3 also acts as an acceptor to the coordinated water molecule (O1W) of the neighbouring unit. Another O atom of the SO_3 group (O2) is also involved in hydrogen bonds, one with an uncoordinated water molecule (O4W) and another with a neighbouring coordinated water molecule (O2W). Two coordinated water molecules (O1W and O2W) of one dimer donate H atoms to two of the O atoms (O2 and O3) of the sulfonate group of a neighbouring unit, reminiscent of the fork-like interactions (Fig. 3) observed in trimethoprim–sulfonate complexes (Baskar Raj, Sethuraman

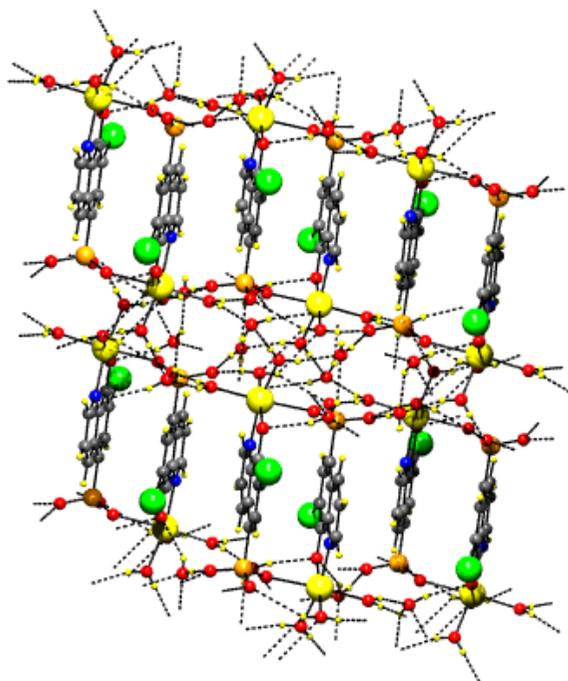


Figure 5
The supramolecular architecture in (I).

et al., 2003). The quinolinol O atom (O4) acts as a hydrogen-bonding acceptor to two uncoordinated water molecules, O4W and O5W, resulting in two types of hydrogen-bonding pattern. One (O4...H25W—O5W—H15W...O4W...H13W—O3W—Zn) makes an eight-membered ring (Fig. 4) and the other (O4...H24W—O4W—H14W...O2) is intermolecular in nature, bridges neighbouring moieties through the hydrogen-bonded chain. The overall supramolecular architecture consists of parallel supramolecular chains, made up of a metal atom and O atoms (of sulfonate and water molecules), linked by the aromatic part of the oxine moieties acting as ladders (Fig. 5).

Experimental

Hot aqueous solutions of trimethoprim (obtained as a gift from Shilpa Antibiotics Ltd, 0.072 g) and ferron (Riedel-de-Haen, 0.085 g) were mixed in a 1:1 molar ratio. Then a hot aqueous solution of ZnSO₄·7H₂O (S. D. Fine Chemicals Ltd, 0.072 g) was added to the above solution and warmed over a water bath for 8 h. After a few days of slow evaporation, needle-shaped yellow crystals of (I) separated out from the mother liquor.

Crystal data

[Zn₂(C₉H₄INO₄S)₂(H₂O)₆]₂·4H₂O
M_r = 1009.14
 Monoclinic, *C*2/*c*
a = 22.278 (4) Å
b = 10.076 (3) Å
c = 13.496 (2) Å
 β = 102.29 (3)°
V = 2960.1 (12) Å³
Z = 4

D_x = 2.264 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 25 reflections
 θ = 3.1–30.1°
 μ = 3.93 mm⁻¹
T = 293 (2) K
 Thin plate, yellow
 0.31 × 0.26 × 0.23 mm

Data collection

Philips PW1100 diffractometer
 ω scans
 Absorption correction: ψ scan
 (North *et al.*, 1968)
 T_{\min} = 0.303, T_{\max} = 0.405
 4484 measured reflections
 4326 independent reflections
 2975 reflections with $I > 2\sigma(I)$
 R_{int} = 0.021

θ_{\max} = 30.0°
 h = -31 → 30
 k = 0 → 14
 l = 0 → 18
 1 standard reflection
 every 100 reflections
 frequency: 60 min
 intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)]$ = 0.028
 $wR(F^2)$ = 0.061
 S = 0.90
 4326 reflections
 255 parameters

All H-atom parameters refined
 $w = 1/[\sigma^2(F_o^2) + (0.0239P)^2]$
 $(\Delta/\sigma)_{\max}$ = 0.001
 $\Delta\rho_{\max}$ = 0.91 e Å⁻³
 $\Delta\rho_{\min}$ = -1.00 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

I—C7	2.093 (3)	S—O1	1.461 (2)
Zn—O1	2.311 (2)	S—O2	1.466 (2)
Zn—O1W	2.132 (3)	S—O3	1.457 (2)
Zn—O2W	2.101 (3)	S—C5	1.771 (3)
Zn—O3W	2.017 (3)	O4—C8	1.311 (4)
Zn—O4 ⁱ	2.085 (2)	N1—C10	1.370 (4)
Zn—N1 ⁱ	2.073 (3)	N1—C2	1.325 (4)
O1—Zn—O1W	176.67 (11)	O2—S—O3	112.54 (14)
O1—Zn—O2W	89.46 (12)	O2—S—C5	105.36 (13)
O1—Zn—O3W	82.17 (10)	O3—S—C5	106.37 (14)
O1—Zn—O4 ⁱ	88.14 (9)	Zn—O1—S	140.92 (13)
O1—Zn—N1 ⁱ	87.41 (10)	Zn ⁱ —O4—C8	111.74 (18)
O1W—Zn—O2W	89.19 (14)	Zn ⁱ —N1—C2	128.6 (2)
O1W—Zn—O3W	94.70 (12)	C2—N1—C10	118.9 (3)
O1W—Zn—O4 ⁱ	93.25 (11)	Zn ⁱ —N1—C10	112.23 (18)
O1W—Zn—N1 ⁱ	95.80 (12)	N1—C2—C3	122.8 (3)
O2W—Zn—O3W	85.97 (13)	S—C5—C6	119.4 (2)
O2W—Zn—O4 ⁱ	177.50 (13)	S—C5—C9	121.2 (2)
O2W—Zn—N1 ⁱ	98.63 (11)	I—C7—C6	119.6 (2)
O3W—Zn—O4 ⁱ	94.40 (11)	I—C7—C8	117.6 (2)
O3W—Zn—N1 ⁱ	168.59 (11)	O4—C8—C7	124.5 (3)
O4 ⁱ —Zn—N1 ⁱ	80.56 (9)	O4—C8—C10	119.9 (3)
O1—S—O2	110.85 (13)	N1—C10—C9	122.0 (3)
O1—S—O3	113.43 (14)	N1—C10—C8	115.4 (3)
O1—S—C5	107.74 (14)		

Symmetry code: (i) $\frac{1}{2} - x, \frac{1}{2} - y, -z$.

Table 2

Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O1W—H11W...O5W ⁱ	0.91 (3)	1.97 (3)	2.855 (4)	163 (4)
O2W—H12W...O3	0.94 (5)	2.56 (5)	3.333 (5)	140 (5)
O2W—H12W...O2W ⁱⁱ	0.94 (5)	2.58 (5)	3.156 (5)	120 (5)
O3W—H13W...O4W ⁱⁱⁱ	0.83 (5)	1.97 (5)	2.777 (4)	165 (4)
O4W—H14W...O2 ^{iv}	0.79 (4)	2.04 (4)	2.813 (4)	168 (4)
O5W—H15W...O4W ^v	0.79 (5)	2.05 (5)	2.771 (4)	151 (5)
O1W—H21W...O3 ^{vi}	0.92 (3)	1.91 (3)	2.834 (4)	177 (4)
O2W—H22W...O2 ^{vi}	0.93 (3)	1.83 (3)	2.753 (4)	170 (4)
O3W—H23W...O5W ^{vii}	0.80 (4)	1.96 (4)	2.761 (4)	176 (4)
O4W—H24W...O4	0.72 (5)	2.29 (5)	2.947 (4)	152 (5)
O5W—H25W...O4 ^{viii}	0.84 (5)	2.16 (5)	2.988 (4)	168 (5)
C6—H6...O3	0.96 (4)	2.42 (4)	2.867 (4)	108 (3)

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

H atoms were located from difference Fourier maps and their coordinates and isotropic displacement parameters were refined. The

C—H and O—H bond lengths are 0.90 (3)–0.96 (4) and 0.72 (5)–0.94 Å, respectively.

Data collection: *FEBO* (Belletti, 1996); cell refinement: *MolEN* (Fair, 1990); data reduction: *MolEN*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON97* (Spek, 1997); software used to prepare material for publication: *PLATON97*.

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