metal-organic papers

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Key indicators

Single-crystal X-ray study $T = 293 K$ Mean σ (C-C) = 0.008 Å R factor = 0.066 W_R factor = 0.103 Data-to-parameter ratio = 14.7

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

Interplay of hydrogen bonding and coordination modes in diaquabis(8-hydroxy-7-iodoquinoline-5-sulfonato)calcium(II) monohydrate

In the crystal structure of the title compound, $[Ca(C_9H_5I NO_4S$ ₂(H₂O)₂.H₂O, the asymmetric unit consists of one calcium ion, two organic ligands, two coordinated water molecules and one uncoordinated water molecule. The calcium ion has a pentagonal-bipyramidal geometry, formed by O atoms of three different sulfonate groups, deprotonated O atoms of two different oxine rings and two water molecules. In the ligands of type I, two O atoms of the sulfonate group are involved in coordination, whereas in type II ligands, only one O atom is involved. Ligands of type I are stacked over one another, each pair of adjacent members of the stack being related by an inversion centre. Ligands of type II are also stacked in a similar manner. Both of these molecular stacks are cross-linked by $Ca-O(sulfonate)$ and $Ca-O(quinolinol)$ bonds, leading to a three-dimensional network. This network is further stabilized by a number of $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonds.

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Comment

Calcium is essential for physiological functions. Studies of metal complexes of drugs will contribute towards the understanding of metal-drug interactions at the molecular level (Guo & Sadler, 2000). The coordination chemistry of the metal-sulfonate system has not been extensively explored and rationalized compared to the well studied metal-phosphonates; this is due to the weak coordination strength of sulfonates with respect to metal ions. Various coordination modes of the sulfonate group in the crystal structures of metal-aryl sulfonates have been reported (Cai, Chen, Feng et al., 2001; Cote et al., 2002). The crystal structures of the calcium complex of 1,5-naphthalenedisulfonate dihydrate (Cai, Chen, Liao et al., 2001), two calcium salts of o -toluidine-6,6'-disulfonate (Gunderman & Squattrito, 1996) and heptaaquacalcium 1,2-benzenedisulfonimide (Moers et al., 1997) have been reported.

8-Hydroxyquinoline (oxine) and its derivatives are well known for their antifungal, antibacterial and antiamoebic activities (Bambury, 1979). The biological activities of oxine derivatives have been correlated with their capacity for metal chelation. Crystal structures have been reported for the

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Figure 1

A view of the asymmetric unit of (I), showing 30% probability displacement ellipsoids.

Figure 2

The coordination geometry of Ca in (I).

The polymeric network of (I) . Suffixes a, b and d correspond to the symmetry codes (i), (ii) and (iii), respectively, given in Table 1.

copper complex of 8-hydroxyquinoline-5-sulfonic acid (HQS) (Petit, Ammor et al., 1993; Petit, Coquerel et al., 1993), the nickel complex of HQS (Raj et al., 2001), the lithium complex of HQS (Murugesan & Muthiah, 1997), the sodium complex of HQS (Raj et al., 2002), the nickel complex of 8-hydroxy-7 iodoquinoline-5-sulfonic acid (ferron) (Raj et al., 2003), the cobalt complex of ferron (Balasubramanian, 1995) and the zinc complex of ferron (Francis et al., 2003).

In the present crystal structure, (I), the asymmetric unit contains a Ca^{2+} ion, two 8-hydroxy-7-iodoquinoline-5-sulfonate (ferron) anions and three water molecules. The metalligand ratio is 1:2. A view of the asymmetric unit is shown in Fig. 1. In this complex, the coordination geometry around calcium is that of a seven-coordinate pentagonal bipyramid, formed by O atoms (O4 and O24) of the oxine ring, O atoms (O1, O22 and O23) of three different sulfonate groups of the ferron anions and two water molecules (Fig. 2). The $Ca-O$

Figure 4 The supramolecular architecture of (I).

Figure 5 The hydrogen-bonded network in the complex (I).

distances (Table 1) agree with those in related calcium sulfonate crystal structures (Cai, Chen, Liao et al., 2001; Onoda et al., 2001).

In the ligands of type I, two O atoms of the sulfonate group are involved in coordination, whereas in type II ligands, only one O atom is involved. Ligands of type I (atom labels N2 and C22–C30) are stacked over one another, each pair of adjacent members of the stack being related by an inversion centre. Ligands of type II (atom labels $N1$ and $C2-C10$) are also stacked in a similar manner. Both of these molecular stacks are cross-linked by $Ca-O(sulfonate)$ and $Ca-O(quinolinol)$ bonds, leading to a three-dimensional network (Fig. 3). In the overall three-dimensional network, the planes of the oxine rings in adjacent layers are slightly twisted, by $16.4 \,(1)^{\circ}$, with respect to each other (Fig. 4).

The O atoms (O1W and O2W) of the coordinated water molecules donate H atoms to the sulfonate O atoms (O21 and O3) of two sulfonate groups to form six-membered hydrogenbonded rings on either side of the Ca atom. Another H atom of the coordinated water molecule (O2W) acts as a donor to one of the sulfonate O atoms, O2, generating a ten-membered hydrogen-bonded ring. The non-coordinated water molecule donates H atoms to the O atoms (O2 and O21) of the sulfonate groups to form a 16-membered ring. It also acts as an acceptor with one of the coordinated water molecules (O1W), to produce an eight-membered ring and bridges the two complex moieties, through hydrogen bonding, as shown in Fig. 5. In both type I and type II ligands, the protonated N atom of the oxine ring forms an unsymmetrical bifurcated hydrogen bond with the O atom of the same ring and the coordinated water molecule (Fig. 6).

In this crystal structure, $\pi-\pi$ stacking interactions are also observed between the oxine rings. For the type I ligand, the interplanar and centroid-to-centroid distances are 3.463 and 3.628 (3) Å, respectively, and the slip angle (defined as the angle between the plane normal and the line joining the centroids) is 17.69° . The corresponding values for the type II ligand are 3.318 and 3.509 (3) \AA and 21.09°, respectively. Similar stacking has also been observed in the crystal structures of the nickel complex of ferron (Raj et al., 2003), and the zinc complex of ferron (Francis et al., 2003). In the title complex, two I \cdots O interactions are observed, viz. I2 \cdots O2 [3.470 (4) Å; symmetry code: $\frac{1}{2} - x$, $y + \frac{1}{2}$, $1 - z$] and I2. \cdot O3W [3.592 (4) Å; symmetry code: $1 - x$, $1 - y$, $1 - z$]. This type of oxygen-halogen interaction has also been observed in the crystal structures of ferron (Balasubramanian & Muthiah, 1996) and its nickel complex (Raj et al., 2003), among others.

Experimental

Hot aqueous solutions of ferron (88 mg, Riedel de-Haen) and salicylic acid (35 mg, Merck) were mixed in a 1:1 molar ratio. The resultant solution was added to $CaCO₃$ (25 mg, LOBA) in a 1:1:1 molar ratio and warmed over a water bath for 90 min. The red solution was filtered and allowed to evaporate slowly. After a few days, red plate-shaped crystals separated from the solution.

 \overline{D} = 2.182 Mg mix = 2.

Crystal data

Figure 6 Bifurcated hydrogen-bonding in (I).

Refinement

Table 1

Selected geometric parameters (\mathring{A}, \degree) .

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

Table 2

Hydrogen-bonding geometry (A, \circ) .

Symmetry codes: (i) $x, y - 1, z$; (iii) $2 - x, 1 - y, 2 - z$; (iv) $1 - x, \frac{1}{2} + y, \frac{3}{2} - z$; (v) $x, 1 + y, z$; (vi) $x, \frac{3}{2} - y, z - \frac{1}{2}$; (vii) $x, \frac{5}{2} - y, \frac{1}{2} + z$.

H atoms were located in a difference map and were treated as riding atoms, with C $-H$, O $-H$ and N $-H$ distances of 0.96, 0.85 and 0.90 Å, respectively; U_{iso} values were set at 1.2 U_{eq} of the carrier atom.

The highest residual electron-density peak is located 0.941 Å from atom I2.

Data collection: KM-4 CCD Software (Kuma, 1999); cell refinement: KM-4 CCD Software; data reduction: KM-4 CCD Software; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 1997); software used to prepare material for publication: SHELXL97.

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