

## Hydrogen bonding in proton-transfer compounds of 8-quinolinol (oxine) with aromatic sulfonic acids

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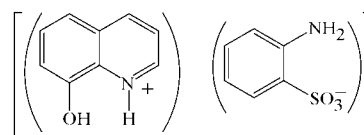
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The crystal structures of the proton-transfer compounds of 8-quinolinol (oxine) with the aromatic sulfonic acids 2-aminobenzenesulfonic acid (orphanic acid) and 8-hydroxy-7-iodoquinoline-5-sulfonic acid (ferron) have been determined. In both 8-hydroxyquinolinium 2-aminobenzenesulfonate,  $C_9H_8NO^+ \cdot C_6H_6NO_3S^-$ , (I), and 8-hydroxyquinolinium 8-hydroxy-7-iodoquinoline-5-sulfonate sesquihydrate,  $C_9H_8NO^+ \cdot C_9H_5INO_4S^- \cdot 1.5H_2O$ , (II), extensive hydrogen-bonding interactions, together with significant cation–cation [in (I)] and cation–anion [in (II)]  $\pi$ – $\pi$  stacking associations, give rise to layered polymer structures.

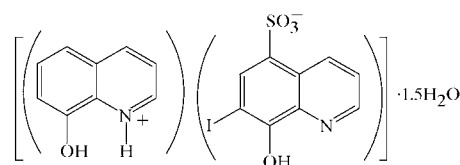
### Comment

8-Quinolinol (8-hydroxyquinoline, 8-HQ) is a particularly useful reagent (oxine) for metal complex formation, achieved through *N,O*-bidentate interaction, giving bis- and tris-chelate compounds, for example,  $[Pd(8-HQ)_2]$  (Prout & Wheeler, 1966). Oxine will react with most metal types, but the values of the dissociation constants for the phenol O and the heterocyclic N-donor groups ( $pK_{a1,2} = 4.9$  and 10.8) provide the textbook example of a reagent that has wide interactive selectivity, particularly through pH control, giving numerous analytical chemical applications (Skoog *et al.*, 1988). The molecule also readily forms adducts, both with metal complex species, for example,  $K^+ \cdot (8-HQ)^- \cdot (8-HQ)$  and  $K^+ \cdot (8-HQ)^- \cdot 2(8-HQ)$  (Hughes & Truter, 1979), and with neutral compounds, for example, chloranil (1:1; Prout & Wheeler, 1967) and 1,3,5-trinitrobenzene (1:1; Castellano & Prout, 1971). In the presence of relatively strong organic acids, protonation of the heterocyclic N atom of oxine occurs and the resulting 8-HQ<sup>+</sup> cation provides structure extension through hydrogen-bonding interactions. Known structures of this type are those with salicylic acid (SA, a discrete 2:2 dimer; Singh *et al.*, 2000; Smith, Wermuth & White, 2003), 2-nitrobenzoic acid (1:1 monohydrate), 3,5-dinitrobenzoic acid (1:1 trihydrate) and 3,5-dinitrosalicylic acid (1:1; Smith *et al.*, 2001),

5-sulfosalicylic acid (1:1 monohydrate; Smith, Wermuth & White, 2004), and Kemp's triacid (*cis-cis*-1,3,5-trimethylhexane-1,3,5-tricarboxylic acid, a 1:1 anhydrate; Smith *et al.*, 2000). Adduct salts with SA  $[(8-HQ)^+ \cdot (SA)^- \cdot (8-HQ)]$ ; Jebamony & Muthiah, 1998] and 1,2,3-trihydroxybenzene  $[(8-HQ)^+ \cdot (THB)^- \cdot (8-HQ)]$ ; Singh *et al.*, 1994] are also known. It is also of interest that, with a number of these compounds, the formation reactions can proceed in the solid state (Rastogi *et al.*, 1977; Singh *et al.*, 1994, 1999, 2000).



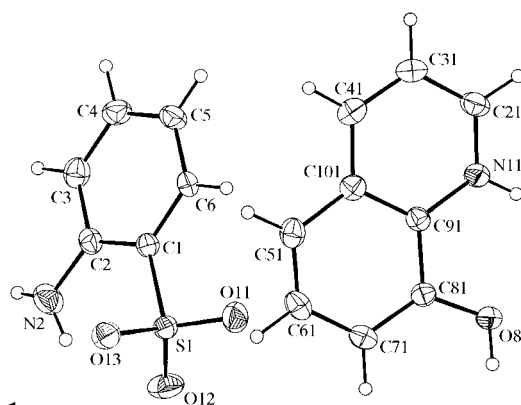
(I)



(II)

8-Hydroxy-7-iodoquinoline-5-sulfonic acid (ferron) is chemically analogous to oxine, with potential for complex formation but with steric constraints because of the presence of the bulky 7-iodo substituent. Therefore, ferron gives a colour reaction with iron(III) but not with iron(II) (Vogel, 1964). The crystal structure determination of ferron (Merritt & Duffin, 1970; Balasubramanian & Muthiah, 1996) has confirmed the presence of a sulfonate–quinolinium zwitterion. We have reported the structure of the bis(guanidinium) monohydrate salt of ferron, in which deprotonation of both the sulfonic acid and the phenol groups has occurred (Smith, Wermuth & Healy, 2003), while in the structure of the 1:1 salt with 4,4'-bipyridine (Hemamalini *et al.*, 2004), only the sulfonate group is deprotonated. We have also determined the 1:1:1 ferron/urea/water adduct structure (Smith, Wermuth & Healy, 2004).

The crystal structures reported here are those of the products of the reaction of 8-hydroxyquinoline with

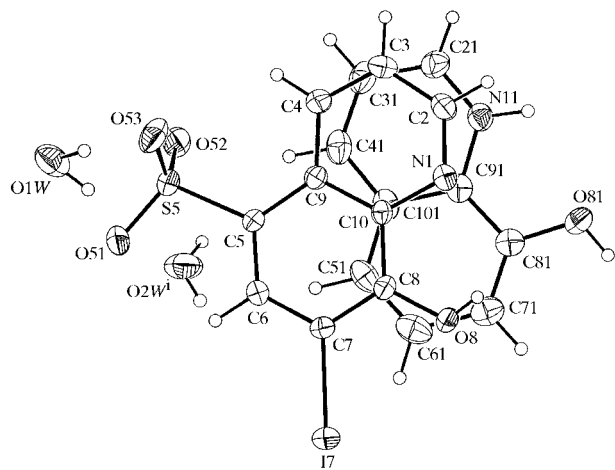


**Figure 1**  
The conformations and atom-numbering schemes for the 8-HQ<sup>+</sup> cation and 2-ABS<sup>-</sup> anion in (I). Non-H atoms are shown as 30% probability displacement ellipsoids.

2-aminobenzenesulfonic acid (orphanic acid), namely 8-hydroxyquinolinium 2-aminobenzenesulfonate [(8-HQ)<sup>+</sup>·(2-ABS)<sup>-</sup>], (I), and with ferron, 8-hydroxyquinolinium 8-hydroxy-7-iodoquinoline-5-sulfonate sesquihydrate [(8-HQ)<sup>+</sup>·C<sub>9</sub>H<sub>6</sub>INO<sub>4</sub>S·1.5H<sub>2</sub>O], (II). As expected, in both compounds, proton transfer from the sulfonic acid group to the quinoline N atom of the oxine molecule occurs. Fig. 1 shows the atom-numbering scheme used for the 8-HQ<sup>+</sup> cation and the 2-aminobenzenesulfonate anion (2-ABS<sup>-</sup>) in (I). A similar scheme is employed for the 8-HQ<sup>+</sup> cation in (II) (Fig. 2). Tables 1 and 2 list the geometric parameters for the hydrogen-bonding interactions in (I) and (II) and the symmetry codes used below.

Compound (I) exhibits a direct N<sup>+</sup>—H···O<sub>sulfonate</sub><sup>-</sup> interaction [N11—H11···O11 = 2.745 (4) Å; Fig. 3]. The 8-hydroxy substituent group of the 8-HQ<sup>+</sup> cation is strongly associated intermolecularly with a 2-ABS<sup>-</sup> sulfonate O atom [O81···O13<sup>ii</sup> = 2.623 (3) Å]. As is usual for 8-HQ<sup>+</sup> cations, there is an intramolecular association between the quinolinium H atom and the phenol O atom [N11···O81 = 2.675 (4) Å], while in the 2-ABS<sup>-</sup> anion, the conformation of the sulfonate group is stabilized by N<sub>amine</sub>—H···O and C<sub>ring</sub>—H···O interactions [2.937 (5) and 2.875 (4) Å, respectively].

The inversion-related 8-HQ<sup>+</sup> cation rings form stacks that extend approximately along the *c* direction, with approximate superimposition of the C51/C61/C71/C81/C91/C101 six-membered rings. These have perpendicular separations of 3.40 (1) (intradimer) and 3.39 (1) Å (interdimer), with a ring-centroid separation of 3.78 (1) Å, which is strongly indicative of π–π interaction. These ‘sandwich’ dimers are similar to those found in the discrete dimeric 2:2 compound of oxine with salicylic acid (Smith, Wermuth & White, 2003), in which the intra- and interdimer separations are 3.34 and 3.40 Å, respectively (the ring-centroid separation is 3.58 Å). In this salicylate compound, the oxine rings are edge-linked by bisalicylate cleats. In (I), the molecular pairs are peripherally linked to the sulfonate anions, forming sheets across the *ac*

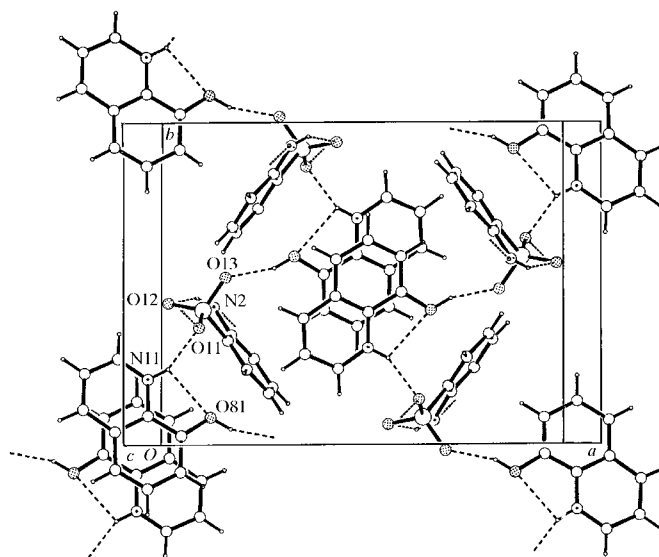


**Figure 2**

The conformation and atom-numbering scheme for (II), showing the 8-HQ<sup>+</sup> cation and ferron anion superimposition. Non-H atoms are shown as 30% probability displacement ellipsoids. [Symmetry code: (i) *x*, *y*, *z* − 1.]

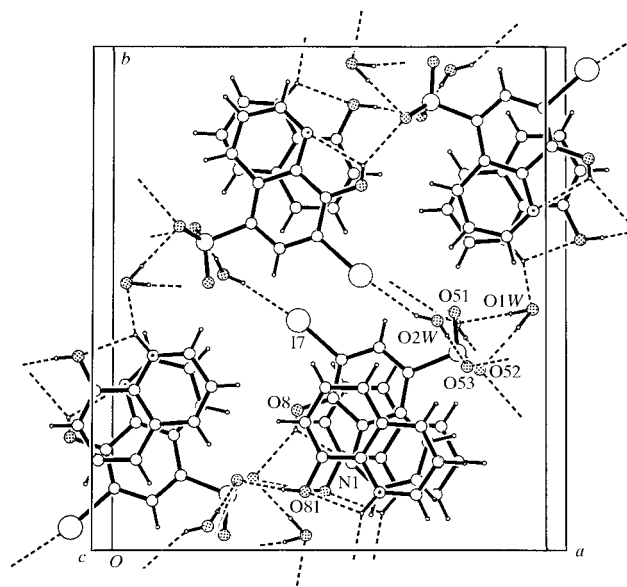
plane with only weak intersheet interactions [N2—H2B···O11<sup>i</sup> = 3.402 (4) Å].

In (II), the protonated heterocyclic N atom of the 8-HQ<sup>+</sup> cation forms an intramolecular hydrogen bond with the phenol O atom similar to that in (I) [N11···O81 = 2.694 (6) Å; Fig. 4]. However, in contrast to the reaction in (I), the propagating interaction in (II) is with a water molecule [N11—H11···O1W<sup>iv</sup> = 2.733 (5) Å] rather than with a sulfonate O atom. Although a direct N<sup>+</sup>—H···O<sub>sulfonate</sub><sup>-</sup> interaction is also found in the monohydrated compound of oxine with 5-sulfosalicylic acid (5-SSA; Smith, Wermuth & White, 2004), it is



**Figure 3**

A perspective view of the packing of (I) in the unit cell, viewed down *c*, showing homomolecular stacks of oxine cations and hydrogen-bonding associations (broken lines).



**Figure 4**

A perspective view of the packing of (II) in the unit cell, viewed down *c*, showing the 8-HQ<sup>+</sup>-cation–ferron-anion stacks linked through hydrogen-bonding interactions (broken lines) also involving the water molecules.

more usual that, in the hydrated 5-SSA compounds with the quinoline-type Lewis bases, this primary interaction is with a water molecule. The ferron anion molecules have an intramolecular aromatic C6—H6···O51<sub>sulfonate</sub> contact [2.823 (6) Å], while a short intermolecular OW2···I7<sup>vi</sup> contact [3.17 (1) Å] is also present.

The molecules of (II) form stacks of alternating 8-HQ<sup>+</sup> cations and ferron anions, having approximate superimposition of the six-membered rings N11/C21/C31/C41/C101/C91 and N1/C2/C3/C4/C9/C10, with perpendicular separations of 3.51 (1) (intradimer) and 3.40 (1) Å (interdimer), also indicative of significant π–π interaction. These stacks form along the *c* axis and are linked by the previously described hydrogen bonds, as well as by a number of other associations involving the water molecules. This stacking phenomenon is consistent with perfect crystal cleavage perpendicular to the short [6.977 (4) Å] *c* axis. The presence of a hemihydrate molecule (O2W) is consistent with some lability in this molecule, although it is strongly associated in the crystal and only minor crystal decomposition (1.1%) was observed during data collection.

Experimental

The title compounds were synthesized by heating under reflux 1 mmol quantities of 8-quinolinol and either 2-aminobenzenesulfonic acid [for (I)] or 8-hydroxy-7-iodoquinoline-5-sulfonic acid [for (II)] in 50% ethanol/water (50 ml) for 10 min. After concentration to ca 30 ml, partial room-temperature evaporation of the hot-filtered solutions gave large pale-brown crystal blocks of (I) (m.p. 454.6–454.7 K) and large yellow prisms of (II) (m.p. 475.7–478.7 K).

Compound (I)

Crystal data

C<sub>9</sub>H<sub>8</sub>NO<sup>+</sup>·C<sub>6</sub>H<sub>6</sub>NO<sub>3</sub>S<sup>-</sup>  
*M<sub>r</sub>* = 318.34  
 Monoclinic, *P*<sub>2</sub><sub>1</sub>/*a*  
*a* = 16.030 (4) Å  
*b* = 11.726 (3) Å  
*c* = 7.7024 (15) Å  
 β = 100.298 (17)°  
*V* = 1424.5 (6) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.484 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 25 reflections  
 θ = 12.7–17.1°  
 μ = 0.25 mm<sup>-1</sup>  
*T* = 298 (2) K  
 Fragment, pale brown  
 0.40 × 0.20 × 0.10 mm

Data collection

Rigaku AFC-7R diffractometer  
 ω–2θ scans  
 Absorption correction: ψ scan  
 (TEXSAN for Windows;  
 Molecular Structure Corporation,  
 1999)  
*T<sub>min</sub>* = 0.942, *T<sub>max</sub>* = 0.976  
 3780 measured reflections  
 3265 independent reflections

2047 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.053  
 θ<sub>max</sub> = 27.5°  
*h* = –8 → 20  
*k* = –15 → 6  
*l* = –10 → 9  
 3 standard reflections  
 every 150 reflections  
 intensity decay: 0.0%

Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.046  
*wR*(*F*<sup>2</sup>) = 0.163  
*S* = 0.90  
 3265 reflections  
 212 parameters  
 H atoms treated by a mixture of independent and constrained refinement

*w* = 1/[σ<sup>2</sup>(*F*<sub>o</sub><sup>2</sup>) + (0.1*P*)<sup>2</sup> + 3.934*P*]  
 where *P* = (*F*<sub>o</sub><sup>2</sup> + 2*F*<sub>c</sub><sup>2</sup>)/3  
 (Δ/σ)<sub>max</sub> < 0.001  
 Δρ<sub>max</sub> = 0.34 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = –0.32 e Å<sup>-3</sup>  
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.011 (2)

Table 1

Hydrogen-bonding geometry (Å, °) for (I).

<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>
N2–H2A···O12	0.83 (3)	2.26 (4)	2.937 (5)	139 (3)
N2–H2B···O11 <sup>i</sup>	0.86 (4)	2.60 (4)	3.402 (4)	157 (3)
N11–H11···O81	0.85 (3)	2.35 (4)	2.675 (4)	103 (3)
N11–H11···O11	0.85 (3)	1.95 (4)	2.745 (4)	155 (3)
O81–H81···O13 <sup>iii</sup>	0.84 (3)	1.81 (3)	2.623 (3)	161 (3)
C6–H6···O11	0.96	2.46	2.875 (4)	106

Symmetry codes: (i) *x*, *y*, *z* – 1; (ii) ½ – *x*, *y* – ½, 1 – *z*.

Compound (II)

Crystal data

C<sub>9</sub>H<sub>8</sub>NO<sup>+</sup>·C<sub>9</sub>H<sub>5</sub>INO<sub>4</sub>S<sup>-</sup>·1.5H<sub>2</sub>O  
*M<sub>r</sub>* = 523.30  
 Monoclinic, *P*<sub>2</sub><sub>1</sub>/*a*  
*a* = 15.857 (4) Å  
*b* = 17.667 (7) Å  
*c* = 6.977 (4) Å  
 β = 95.78 (4)°  
*V* = 1944.5 (14) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.791 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 25 reflections  
 θ = 12.6–16.6°  
 μ = 1.80 mm<sup>-1</sup>  
*T* = 298 (2) K  
 Block, yellow  
 0.50 × 0.50 × 0.35 mm

Data collection

Rigaku AFC-7R diffractometer  
 ω–2θ scans  
 Absorption correction: ψ scan  
 (TEXSAN for Windows;  
 Molecular Structure Corporation,  
 1999)  
*T<sub>min</sub>* = 0.434, *T<sub>max</sub>* = 0.533  
 5029 measured reflections  
 4473 independent reflections

3424 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.068  
 θ<sub>max</sub> = 27.5°  
*h* = –9 → 22  
*k* = 0 → 20  
*l* = –9 → 9  
 3 standard reflections  
 every 150 reflections  
 intensity decay: 1.1%

Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.049  
*wR*(*F*<sup>2</sup>) = 0.158  
*S* = 0.89  
 4473 reflections  
 291 parameters  
 H atoms treated by a mixture of independent and constrained refinement

*w* = 1/[σ<sup>2</sup>(*F*<sub>o</sub><sup>2</sup>) + (0.1*P*)<sup>2</sup> + 7.5983*P*]  
 where *P* = (*F*<sub>o</sub><sup>2</sup> + 2*F*<sub>c</sub><sup>2</sup>)/3  
 (Δ/σ)<sub>max</sub> = 0.002  
 Δρ<sub>max</sub> = 1.55 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = –1.30 e Å<sup>-3</sup>  
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.0024 (6)

Table 2

Hydrogen-bonding geometry (Å, °) for (II).

<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–H1A···O52	0.97 (6)	1.96 (6)	2.882 (6)	156 (5)
O1W–H1B···O2W <sup>i</sup>	0.97 (6)	2.22 (6)	3.175 (11)	169 (5)
O2W–H2A···I7 <sup>vi</sup>	0.79 (6)	2.44 (7)	3.169 (10)	155 (7)
O2W–H2B···O53	0.90 (6)	1.94 (6)	2.848 (11)	180 (6)
O8–H8···N1	0.86 (9)	2.26 (7)	2.687 (5)	111 (5)
O8–H8···O52 <sup>iii</sup>	0.86 (9)	2.13 (8)	2.836 (5)	139 (6)
N11–H11···O81	0.96 (7)	2.26 (7)	2.694 (6)	107 (5)
N11–H11···O1W <sup>iv</sup>	0.96 (7)	1.86 (7)	2.733 (5)	150 (6)
O81–H81···O53 <sup>v</sup>	0.86 (6)	1.80 (6)	2.656 (5)	175 (5)
C6–H6···O51	0.95	2.38	2.823 (6)	108

Symmetry codes: (i) *x*, *y*, *z* – 1; (iii) *x* – ½, ½ – *y*, *z*; (iv) ½ – *x*, *y* – ½, –*z*; (v) *x* – ½, ½ – *y*, *z* – 1; (vi) 1 – *x*, 1 – *y*, 1 – *z*.

H atoms potentially involved in hydrogen-bonding interactions were located by difference methods and their positional and isotropic displacement parameters were refined. Other H atoms were included in the respective refinements at calculated positions (C–H = 0.95 Å) as riding atoms, with *U*<sub>iso</sub>(H) values fixed at 1.2*U*<sub>eq</sub>(C). The large

difference electron-density maximum and minimum for (II) are adjacent to the I atom.

For both compounds, data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1999); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN for Windows* (Molecular Structure Corporation, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON for Windows* (Spek, 1999); software used to prepare material for publication: *PLATON for Windows*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: TA1460). Services for accessing these data are described at the back of the journal.

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