# organic compounds

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# Formation of ladders from $R_4^4(8)$ and $R_6^6(12)$ rings in 8-hydroxyquinolinium chloride monohydrate: comparisons with the supramolecular arrangements in related salts

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Molecules of the title compound,  $C_9H_8NO^+\cdot Cl^-\cdot H_2O$ , are linked into two rings by strong hydrogen bonding *via* the free water molecules and the Cl<sup>-</sup> anions. The two hydrogenbonded rings are joined to give a corrugated chain along [001]. Comparisons with other 8-hydroxyquinoline-based salts are also presented, highlighting similar ring structures in a 1:1 salt with Kemp's triacid (*r*-1,*c*-3,*c*-5-trimethylcyclohexane-1,3,5tricarboxylic acid) and in 8-hydroxy-1-methylquinolinium chloride monohydrate.

### Comment

In a continuation of our interest in the supermolecular arrangements of organic molecules, we now report the crystal structure of hydrated 8-hydroxyquinolinium chloride, 8-HO-QH<sup>+</sup>·Cl<sup>-</sup>. The title compound, (I), was isolated from a reaction mixture consisting of 2-chloronicotinoyl chloride and 8-hydroxyquinoline, 8-HOQ, in acetone, followed by recrystallization of the reaction residues from ethanol. Clearly, the water present in the solvent(s) had led to hydrolysis of the acyl chloride with formation of hydrogen chloride, which then formed the salt with 8-hydroxyquinoline.



Compound (I) crystallizes in the space group  $P\overline{1}$  and, for convenience, the reference positions of the free Cl atoms and water molecules were chosen to give the most direct



hydrogen-bonding scheme (Fig. 1 and Table 1). The hydrogen bonding was analysed with the aid of *PLATON* (Spek, 2003).

The 8-HOQ molecule is protonated at the N atom (8-HOQH<sup>+</sup>) and is very nearly planar; the angle between the fused rings is 0.32 (6)° and the Cremer & Pople (1975) total puckering amplitude, Q, is 0.0247 (17) Å.

The intramolecular  $N^+-H\cdots O$  hydrogen bond in 8-HOQH<sup>+</sup> has been observed in other 8-hydroxyquinolinium and related cations. In 7-iodo-8-hydroxyquinoline-5-sulfonic acid (Balasubramanian & Muthiah, 1996) and in an 8-hydroxyquinolinium salicylate–salicyclic acid complex, 8-HOQH<sup>+</sup>·C<sub>7</sub>H<sub>5</sub>O<sub>3</sub><sup>-</sup>·C<sub>7</sub>H<sub>6</sub>O<sub>3</sub> (Jebamony & Thomas Muthiah, 1998), this hydrogen bond was observed to result in an enhancement of the internal angle at the nitrogen centre. This interaction has also been highlighted in the crystal structures of 8-hydroxyquinoline-5-sulfonic acid dihydrate (Banerjee *et al.*, 1984), 8-hydroxy-7-nitroquinoline-5-sulfonic acid monohydrate (Balasubramanian & Thomas Muthiah, 1996) and 8-hydroxyquinolinium 3-carboxy-4-hydroxybenzenesulfonate monohydrate, 8-HOQH<sup>+</sup>·C<sub>7</sub>H<sub>5</sub>O<sub>6</sub>S<sup>-</sup>·H<sub>2</sub>O (Smith *et al.*, 2004).

The solvent water molecules and free chloride ions lead to a number of intra- and intermolecular hydrogen bonds (Table 1).











Part of the crystal structure of the title compound, showing the formation of an  $R_6^6(12)$  ring. Displacement ellipsoids are shown at the 50% probability level, H atoms are shown as circles of arbitrary radii and dashed lines indicate hydrogen bonds. [Symmetry code: (i) -x + 1, -y, -z.]

Taking these entities alone first, the water molecule at (x, y, z) participates in O1W-H1WA···Cl1 and O1W-H1WB···Cl1<sup>ii</sup> hydrogen bonds [symmetry code: (ii) -x + 1, -y, -z - 1], forming small  $R_4^4(8)$  rings (Bernstein *et al.*, 1995). This type of interaction is identical to that observed in 8-hydroxy-1-methylquinolinium chloride monohydrate (Rømming & Uggerud, 1983) and the isostructural iodide (Barczyński *et al.*, 2006), both of which also crystallize in  $P\overline{1}$ . In these latter structures, however, the methyl substitution at the N atom hinders further strong hydrogen-bonding interactions, and thus this centrosymmetric  $R_4^4(8)$  dimer is the only motif observed.

In (I), another ring forms from a combination of hydrogen bonds  $[O1-H1A\cdotsO1W, O1W-H1WA\cdotsCl1, N1-H1\cdots$  $Cl1^{i}$  and  $N1-H1\cdotsO1$ ; symmetry code: (i) -x + 1, -y, -z].



### Figure 3

Part of the crystal structure of the title compound, showing (a) the formation of 'ladders' and (b) the relative conformation of the molecules in the 'ladder' formed from the two hydrogen-bonded rings. Displacement ellipsoids are shown at the 50% probability level, H atoms are shown as circles of arbitrary radii and dashed lines indicate hydrogen bonds. [Symmetry code: (ii) -x + 1, -y, -z - 1.]

Together, these form an  $R_6^6(12)$  dimeric ring centred at  $(\frac{1}{2}, 0, 0)$  (Fig. 2).

The two rings join to form a polymeric ladder along [001] (Fig. 3a); viewed in the orientation shown in Fig. 3(a), the ladder appears flat, but viewed along the direction of the ladder, it can be seen that the molecules hydrogen bond in a chair-like configuration (Fig. 3b). The interactions forming this ladder are similar to those observed in the structure of 8-HOQH<sup>+</sup>·KTA<sup>-</sup> (KTA is Kemp's triacid, r-1,c-3,c-5-trimethylcyclohexane-1,3,5-tricarboxylic acid), in which a polymeric chain forms via the carboxylate groups of KTA and the hydroxy and N<sup>+</sup>H groups of 8-HOQH<sup>+</sup>, leading to similar dimeric  $R_6^6(12)$  rings which then form chains *via* the symmetric KTA molecule (Smith et al., 2000). The presence of the intramolecular N<sup>+</sup>-H···O hydrogen bond in 8-HOQH<sup>+</sup> was not shown in the cited work, but the effect of this intramolecular bond is to provide a 'short-cut' in the ring by providing an extra hydrogen bond, shortening the ring from  $R_4^4(18)$  to  $R_6^6(12)$ .

There is also similarity with 8-HOQH<sup>+</sup>·C<sub>7</sub>H<sub>5</sub>O<sub>6</sub>S<sup>-</sup>·H<sub>2</sub>O (Smith *et al.*, 2004), in that the hydroxy group of the 8-HOQH<sup>+</sup> ion acts as a donor to the free water molecule. In this case, the water molecule then goes on to act as a donor to O atoms from the sulfonate group, as does the carboxylate group, forming a three-dimensional network.

The hydrogen-bonding scheme in 8-HOQH<sup>+</sup> salicylate– salicyclic acid, 8-HOQH<sup>+</sup>·C<sub>7</sub>H<sub>5</sub>O<sub>3</sub><sup>-</sup>·C<sub>7</sub>H<sub>6</sub>O<sub>3</sub>, was described in terms of the interactions present but not in terms of the symmetry of these interactions (Jebamony & Thomas Muthiah, 1998). Two of the hydrogen bonds described are between molecules in the same asymmetric unit and do not lead to any continuity of the structure. The other three intermolecular interactions described occur between molecules at (x, y, z) and (-x + 1, -y + 1, -z + 1), forming an isolated dimer. Thus, no chain or network structure is formed in this compound.

The crystal structure of 8-HOQ 2,4,5-trichlorophenol (Singh *et al.*, 2001) shows an interesting phenomenon. The asymmetric unit consists of two molecules of 2,4,5-trichlorophenol and two of 8-HOQ. In one of these latter molecules, the H atom is located at the hetero-N atom, whereas in the other it is at the hydroxy group; thus, neither molecule is protonated as in the present study. The intramolecular N– $H \cdots O$  bond in the 8-HOQ molecule is still observed; hydrogen-bonded chains are supported by a number of intramolecular hydrogen bonds involving both forms of the 8-HOQ molecule, with the unprotonated N and O atoms acting as acceptors.

### **Experimental**

A solution of equimolar (2.0 mmol) 2-chloronicotinoyl chloride and 8-hydroxyquinoline in acetone (30 ml) was refluxed for 20 min; all volatiles were removed under vacuum and the residue was taken up in EtOH. Crystals of the title compound were deposited slowly. IR: 3500–2200 (KBr, with spike at 3327), 1631, 1602, 1552, 1500, 1472, 1398, 1300, 1268, 1220, 1205, 1098, 1059, 999, 661, 821, 756, 711, 523, 578, 540, 488, 412 cm<sup>-1</sup>.

Crystal data	
$\begin{array}{l} C_{9}H_{8}NO^{+}\cdot Cl^{-}\cdot H_{2}O\\ M_{r}=199.63\\ \text{Triclinic, }P\overline{1}\\ a=7.2865\ (3)\ \text{\AA}\\ b=8.2885\ (4)\ \text{\AA}\\ c=8.4596\ (3)\ \text{\AA}\\ \alpha=78.328\ (2)^{\circ}\\ \beta=86.105\ (2)^{\circ}\\ \gamma=66.388\ (2)^{\circ} \end{array}$	$V = 458.41 (3) Å^{3}$ Z = 2 $D_{x} = 1.446 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.38 \text{ mm}^{-1}$ T = 120 (2)  K Shard, light orange $0.36 \times 0.20 \times 0.14 \text{ mm}$
Data collection	
Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 2003) $T_{\min} = 0.802, T_{\max} = 0.928$	9570 measured reflections 2098 independent reflections 1835 reflections with $I > 2\sigma(I)$ $R_{int} = 0.031$ $\theta_{max} = 27.5^{\circ}$
Refinement	
Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.033$ $wR(F^2) = 0.085$ S = 1.14	$w = \frac{1}{[\sigma^2(F_o^2) + (0.0351P)^2} + 0.18P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$

 $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.32 \text{ e} \text{ Å}^{-3}$  $\Delta \rho_{\min} = -0.28 \text{ e} \text{ Å}^{-3}$ H atoms treated by a mixture of independent and constrained

Table 1

2098 reflections

131 parameters

refinement

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O1-H1A\cdots O1W$	0.84	1.80	2.6329 (15)	174
$N1-H1\cdots O1$	0.89 (2)	2.35 (2)	2.7050 (16)	103.7 (15)
$N1-H1\cdots Cl1^{i}$	0.89 (2)	2.37 (2)	3.1048 (13)	140.2 (17)
O1W−H1WA···Cl1 <sup>ii</sup>	0.82(2)	2.30 (3)	3.1245 (13)	176 (2)
$O1W-H1WB\cdots Cl1$	0.79 (2)	2.36 (3)	3.1550 (14)	175 (2)

Symmetry codes: (i) -x + 1, -y, -z; (ii) -x + 1, -y, -z - 1.

The space groups P1 and  $P\overline{1}$  were possible;  $P\overline{1}$  was selected and confirmed by the structure analysis. All H atoms were located from difference maps, and the parameters of water and NH H atoms were refined freely. All other H atoms were treated as riding, with C-H distances of 0.95 Å, O–H distances of 0.84 Å and  $U_{iso}(H)$  values of  $1.2U_{eq}(C)$  or  $1.5U_{eq}(O)$ .

Data collection: COLLECT (Hooft, 1998); cell refinement: DENZO (Otwinowski & Minor, 1997) and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure: OSCAIL (McArdle, 2003) and SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: OSCAIL and SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK3016). Services for accessing these data are described at the back of the journal.

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