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Key indicators

Single-crystal X-ray study T = 150 KMean $\sigma(\text{C-C}) = 0.002 \text{ Å}$ R factor = 0.038 wR factor = 0.104Data-to-parameter ratio = 15.0

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

8-Hydrazinoquinoline dihydrochloride hydrate

The structure of the title compound, $C_9H_{11}N_3\cdot 2Cl\cdot H_2O$, comprises a planar cation containing two quaternary N atoms, packed in the solid-state with two Cl^- anions and one water molecule. Each component is involved in the hydrogen-bonding network, with one Cl^- anion and the O atom both being involved in three-centre associations, while the second Cl^- anion is involved in a four-centre association.

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$$H_3N_+$$
 H (I)

Experimental

The title compound was obtained from Key Organics Ltd and crystals were grown from ethanol solution.

Crystal data

 $C_9H_{11}N_3^-{\cdot}2Cl^-{\cdot}H_2O$ Z = 2 $D_x = 1.518 \text{ Mg m}^{-3}$ $M_r = 250.12$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation a = 6.9716 (14) ÅCell parameters from 3835 reflections b = 8.9866 (18) Åc = 9.986 (2) Å $\theta = 2.9-27.5^{\circ}$ $\mu = 0.57 \ \mathrm{mm}^{-1}$ $\alpha = 70.04 (3)^{\circ}$ T = 150 (2) K $\beta = 89.20 (3)^{\circ}$ $\gamma = 69.60 (3)^{\circ}$ Plate, colourless $V = 547.2 (3) \text{ Å}^3$ $0.40 \times 0.30 \times 0.10 \text{ mm}$

Data collection

Bruker–Nonius KappaCCD areadetector diffractometer φ and ω scans
Absorption correction: multi-scan (SORTAV; Blessing, 1995) $T_{\min} = 0.804, T_{\max} = 0.945$ 7519 measured reflections

2465 independent reflections 2168 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.075$ $\theta_{\rm max} = 27.5^{\circ}$ $h = -9 \rightarrow 9$ $k = -11 \rightarrow 11$ $l = -12 \rightarrow 12$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0551P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.104$ S = 1.04 $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} < 0.001$ $\Delta\rho_{\min} = -0.60$ e Å $^{-3}$ H atoms treated by a mixture of independent and constrained refinement

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Table 1 Hydrogen-bonding geometry (\mathring{A}, \circ) .

D $ H$ $\cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
N1—H1···O1W ⁱ	0.78 (2)	2.05 (2)	2.817 (2)	165 (2)
N81—H81···Cl1 ⁱⁱ	0.91 (2)	2.32 (2)	3.1983 (16)	164.5 (18)
N82—H82···Cl2 ⁱⁱ	0.91 (2)	2.25 (2)	3.1476 (18)	169.8 (18)
N82—H83···O1W ⁱⁱⁱ	0.87 (2)	1.94 (2)	2.808 (2)	178.7 (19)
$N82-H84\cdots Cl2^{iv}$	0.89 (3)	2.39 (3)	3.1805 (17)	149.1 (19)
$O1W-H1W\cdots Cl1^{v}$	0.84 (3)	2.21 (3)	3.0348 (17)	164 (2)
$O1W-H2W\cdots Cl2^{iii}$	0.76 (3)	2.47 (3)	3.180 (2)	158 (2)

Symmetry codes: (i) x, y, z - 1; (ii) x - 1, y, z; (iii) -x, 1 - y, 1 - z; (iv) -x, 1 - y, -z; (v) 1 - x, -y, 1 - z.

All aromatic H atoms were included in the refinement, at calculated positions, as riding models with C—H set to 0.95 Å. All H atoms involved in the hydrogen-bonding network were located in difference syntheses and both positional and displacement parameters were refined.

Data collection: *DENZO* (Otwinowski & Minor, 1997) and *COLLECT* (Hooft, 1998); cell refinement: *DENZO* and *COLLECT*; data reduction: *DENZO* and *COLLECT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *PLATON*-97 (Spek, 1997); software used to prepare material for publication: *SHELXL*97.

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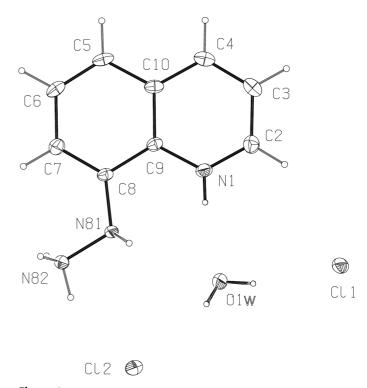


Figure 1Molecular configuration and atom-numbering scheme for the title compound, showing 50% probability ellipsoids.

Sheldrick, G. M. (1997). *SHELXS*97 and *SHELXL*97. University of Göttingen, Germany.

Spek, A. L. (1997). *PLATON-*97. Version of May 1997. University of Utrecht, The Netherlands.