organic papers

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Key indicators

Single-crystal X-ray study T = 297 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.038 wR factor = 0.111 Data-to-parameter ratio = 11.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. Anhydrous quinolinium 2,4,6-trinitrobenzenesulfonate

The title compound, $C_9H_8N^+\cdot C_6H_2N_3O_9S_7^-$ forms a simple dimer structure through a single $N^+-H^{-1}\cdot O$ hydrogen bond involving a sulfonate O atom of the cation. Some cation-cation $\pi-\pi$ interactions are also present.

Comment

Quinoline $(pK_a = 4.8)$ can be readily protonated by sulfonic acids as well as by many carboxylic acids. The single N⁺-H donor group of the resultant quinolinium cation is not a particularly efficient one for molecular assembly via hydrogen bonding. However, cation-cation or cation-anion π - π associations together with aromatic C-H···O interactions are often present in the structures of quinolinium compounds with aromatic carboxylic acids that give stable crystalline materials, e.g. in the anhydrous salt with 3,5-dinitrosalicylic acid (Smith et al., 2006). With aromatic sulfonic acids, the presence of the additional O acceptor atom of the sulfonate group usually results in the incorporation of donor-rich solvent molecules such as water in the crystal structure, e.g. quinolinium 5sulfosalicylate trihydrate (Smith et al., 2004). Therefore, the formation of the title compound, (I), from the 1:1 stoichiometric reaction of quinoline with picrylsulfonic acid in 50% propan-2-ol-water was unusual and its structure is reported here.



In compound (I) (Fig. 1), the quinolinium H atom has a single interaction with a sulfonate-O acceptor of the anion $[N11-H11\cdotsO1B^{i} = 2.725 (3) \text{ Å} and N-H\cdotsO = 162 (2)^{\circ};$ symmetry code: (i) -x + 1, -y, -z + 1], giving a simple hydrogen-bonded heterodimer. The quinolinium cations form stacks through alternating inversion-related molecules in the unit cell (Fig. 2) [ring centroid separation = 3.666 (2) Å; interring dihedral angle = 0°]. The second and third sulfonate O atoms and the six nitro-O atoms are unassociated except for minor weak C-H···O interactions [C21-H···O1B = 3.342 (4) Å, C81-H···O1A^{i} = 3.466 (3) Å and C5-H···O2A^{ii} = 3.241 (4) Å; symmetry code: (ii) x + 1, y, z]. An

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Figure 1

The asymmetric unit of (I). Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

A perspective view of the packing in the unit cell of (I), viewed approximately down the *a*-axis direction, showing hydrogen-bonded associations as dashed lines. [Symmetry code (i): -x + 1, -y, -z + 1.]

unusually short non-bonding intermolecular sulfonate-O···nitro-N interaction is also present $[O1A \cdot \cdot \cdot N4^{iii} = 2.840 (3) \text{ Å}$; symmetry code: (iii) -x + 1, y, z].

The *ortho*-related nitro groups of the picrylsulfonate anions are significantly rotated out of the plane of the benzene ring [torsion angles C1-C2-N2-O2A = -61.6 (3)° and C1-C6-N6-O6B = 61.6 (3)°] compared with the essentially coplanar *para*-related group [C3-C4-N4-O4B = -176.8 (3)°].

Experimental

The title compound was synthesized by heating together 2,4,6-trinitrobenzenesulfonic acid (picrylsulfonic acid) (1 mmol) and quinoline (1 mmol) in 50% propan-2-ol-water (50 ml) under reflux for 10 min. After concentration to ca 30 ml, partial room-temperature evaporation of the hot-filtered solution gave yellow crystal plates (m.p. 532.7–533.9 K).

Crystal data

 $\begin{array}{l} C_9H_8N^+ \cdot C_6H_2N_3O_9S^- \\ M_r = 422.33 \\ \text{Triclinic, } P\overline{1} \\ a = 8.1070 \ (19) \text{ Å} \\ b = 9.486 \ (2) \text{ Å} \\ c = 12.581 \ (2) \text{ Å} \\ a \approx 79.41 \ (2)^\circ \\ \beta = 82.347 \ (16)^\circ \\ \gamma = 64.909 \ (16)^\circ \end{array}$

Data collection

Rigaku AFC 7*R* diffractometer $\omega/2\theta$ scans Absorption correction: ψ scan *TEXSAN* for Windows (Molecular Structure Corporation, 1999) $T_{min} = 0.90, T_{max} = 0.97$ 3552 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.111$ S = 0.95 3026 reflections 266 parameters H atoms treated by a mixture of independent and constrained refinement $V = 859.6 (3) \text{ Å}^{3}$ Z = 2 $D_x = 1.632 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.25 \text{ mm}^{-1}$ T = 297 (2) KPlate, yellow $0.36 \times 0.28 \times 0.11 \text{ mm}$

3026 independent reflections 1992 reflections with $I > 2\sigma(I)$ $R_{int} = 0.017$ $\theta_{max} = 25.0^{\circ}$ 3 standard reflections frequency: 150 min intensity decay: 0.5%

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\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.0556P)^2 \\ &+ 0.1982P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.18 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.30 \ {\rm e} \ {\rm \AA}^{-3} \end{split}
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The single quinolinium H atom was located by difference methods and its positional and isotropic displacement parameters were refined [refined N-H = 0.89 (3) Å]. The aromatic ring H atoms were included in the refinement in calculated positions, with C-H = 0.95 Å, and treated using a riding-model approximation, with $U_{iso}(H)$ = 1.2 $U_{eq}(C)$.

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: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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