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Key indicators

Single-crystal X-ray study T = 291 K Mean σ (C–C) = 0.003 Å R factor = 0.038 wR factor = 0.108 Data-to-parameter ratio = 13.0

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

Quinolinium perchlorate

The crystal structure of the title compound, $C_9H_8N^+ \cdot ClO_4^-$, consists of cations and anions linked by $N-H \cdot \cdot \cdot O$ and $C-H \cdot \cdot \cdot O$ hydrogen bonds to form layers parallel to the (102) plane.

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Comment

The title compound, (I), has previously been analysed using single-crystal X-ray photographic methods, but only the unitcell dimensions and space group have been reported (Herbich & Lipkowski, 1975). The full structure determination of this compound is reported here.



In (I), the asymmetric unit is composed of one quinolinium cation and one perchlorate anion (Fig. 1). The cations and anions form hydrogen-bonded layers, parallel to the (102) plane and are linked together by a network of $N-H\cdots O$ and weak $C-H\cdots O$ hydrogen bonds (Fig. 2). The range of the observed hydrogen-bond parameters (Table 2) agrees with those accepted for $N-H\cdots O$ (Jeffrey, 1997) and $C-H\cdots O$ bonds (Desiraju & Steiner, 1999). Examination of the struc-



View of the title compound with the atom-numbering scheme.

Displacement ellipsoids for non-H atoms are drawn at the 50%

Figure 1

probability level.

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organic papers

ture with PLATON (Spek, 2003) reveals that parallel adjacent benzene rings are close together, with a perpendicular distance of 3.359 Å. However, a slippage of 2.426 Å and a distance between the ring centroids of 4.1434 (11) Å indicates no significant π - π stacking interactions between them.

The O atoms of the perchlorate anion are involved in different ways in the intermolecular hydrogen-bond system. An elongation of the Cl-O1 and Cl-O2 bond lengths [1.4400 (15) and 1.4377 (15) Å, respectively] caused by N - $H \cdots O$ bonds, in comparison with the remaining Cl-O3 and Cl-O4 bonds [1.4134 (18) and 1.4179 (15) Å, respectively], which are involved only in $C-H \cdots O$ interactions, is observed.

Experimental

The title compound was prepared by reaction of stoichiometric amounts of quinoline and perchloric acid (65%). The resulting solid was recrystallized from water.

 $\theta_{\rm max} = 27.5^{\circ}$

 $h = -5 \rightarrow 7$

 $k = -28 \rightarrow 28$

 $l = -10 \rightarrow 10$

Crystal data

| $C_9H_8N^+ \cdot ClO_4^-$ | $D_x = 1.576 \text{ Mg m}^{-3}$ |
|--------------------------------|--|
| $M_r = 229.61$ | Mo $K\alpha$ radiation |
| Monoclinic, $P2_1/c$ | Cell parameters from 7554 |
| a = 5.4935 (5) Å | reflections |
| b = 22.3315 (10) Å | $\theta = 1.8 - 30.3^{\circ}$ |
| c = 7.9970 (4) Å | $\mu = 0.39 \text{ mm}^{-1}$ |
| $\beta = 99.401 \ (6)^{\circ}$ | $T = 291 { m K}$ |
| $V = 967.88 (11) \text{ Å}^3$ | Prism, colourless |
| Z = 4 | $0.6 \times 0.3 \times 0.2 \text{ mm}$ |
| Data collection | |
| Kuma KM-4 CCD diffractometer | $R_{\rm int} = 0.023$ |

 ω scans Absorption correction: none 8865 measured reflections 2190 independent reflections 1921 reflections with $I > 2\sigma(I)$

Refinement

| Refinement on F^2 | $w = 1/[\sigma^2(F_0^2) + (0.0527P)^2]$ |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.038$ | + 0.3838P] |
| $wR(F^2) = 0.108$ | where $P = (F_0^2 + 2F_c^2)/3$ |
| S = 1.03 | $(\Delta/\sigma)_{\rm max} < 0.001$ |
| 2190 reflections | $\Delta \rho_{\rm max} = 0.26 \text{ e } \text{\AA}^{-3}$ |
| 169 parameters | $\Delta \rho_{\rm min} = -0.36 \text{ e } \text{\AA}^{-3}$ |
| All H-atom parameters refined | Extinction correction: SHELXL97 |
| | Extinction coefficient: $0.030(3)$ |

| Table | 1 |
|-------|---|
|-------|---|

Selected geometric parameters (Å, °).

| 1.4400 (15) | C4-C10 | 1.414 (3) |
|-------------|--|--|
| 1.4377 (15) | C5-C6 | 1.360 (3) |
| 1.4134 (18) | C5-C10 | 1.413 (3) |
| 1.4179 (15) | C6-C7 | 1.402 (3) |
| 1.324 (3) | C7-C8 | 1.361 (3) |
| 1.371 (2) | C8-C9 | 1.407 (3) |
| 1.378 (3) | C9-C10 | 1.415 (2) |
| 1.366 (3) | | |
| 107.55 (9) | O2-Cl-O4 | 109.75 (10) |
| 110.68 (12) | O3-Cl-O4 | 109.74 (11) |
| 109.77 (10) | C2-N1-C9 | 123.12 (17) |
| 109.31 (12) | | |
| | $\begin{array}{c} 1.4400 \ (15) \\ 1.4377 \ (15) \\ 1.4134 \ (18) \\ 1.4179 \ (15) \\ 1.324 \ (3) \\ 1.371 \ (2) \\ 1.378 \ (3) \\ 1.366 \ (3) \\ 107.55 \ (9) \\ 110.68 \ (12) \\ 109.77 \ (10) \\ 109.31 \ (12) \end{array}$ | $\begin{array}{ccccc} 1.4400 \ (15) & C4-C10 \\ 1.4377 \ (15) & C5-C6 \\ 1.4134 \ (18) & C5-C10 \\ 1.4179 \ (15) & C6-C7 \\ 1.324 \ (3) & C7-C8 \\ 1.371 \ (2) & C8-C9 \\ 1.378 \ (3) & C9-C10 \\ 1.366 \ (3) \\ \end{array}$ $\begin{array}{cccccccccccccccccccccccccccccccccccc$ |



Figure 2

Packing diagram showing one layer of hydrogen-bonded (dashed lines) cations and anions parallel to the (102) plane, projected along the c axis.

| Table 2 | |
|--------------------------------|--|
| Hydrogen-bond geometry (Å, °). | |

| $D - H \cdot \cdot \cdot A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - \mathbf{H} \cdot \cdot \cdot A$ |
|---|-----------------------------|---------------------------------------|----------------------------|--------------------------------------|
| $N1-H1\cdots O1^{i}$ | 0.83 (2) | 2.13 (2) | 2.958 (2) | 172 (2) |
| $N1 - H1 \cdots O2^i$ | 0.83(2) | 2.59 (2) | 3.156 (2) | 127 (2) |
| $C2-H2 \cdot \cdot \cdot O2^{i}$ | 0.89(2) | 2.58 (2) | 3.200 (3) | 128 (2) |
| $C5-H5\cdots O3^{ii}$ | 0.98(2) | 2.62 (2) | 3.550 (3) | 158 (2) |
| $C7-H7\cdots O4^{iii}$ | 0.89 (2) | 2.57 (2) | 3.395 (3) | 154 (2) |
| Symmetry codes: $-x + 1 + y + \frac{1}{2} - z$ | (i) $x + 1$, $\frac{1}{2}$ | $-y + \frac{3}{2}, +z - \frac{1}{2};$ | (ii) $x, -y + \frac{3}{2}$ | $,+z+\frac{1}{2};$ (iii) |

All H atoms were located in a difference map, and their coordinates and atomic displacement parameters were refined freely [C-H = 0.88 (3) - 0.98 (2) Å].

Data collection: CrysAlis CCD (Oxford Diffraction, 2004); cell refinement: CrysAlis RED (Oxford Diffraction, 2004); data reduction: CrysAlis RED; program(s) used to solve structure: SHELXTL (Sheldrick, 2003); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL and MERCURY (Version 1.3; Bruno et al., 2002); software used to prepare material for publication: PLATON (Spek, 2003).

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