organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.051 wR factor = 0.153 Data-to-parameter ratio = 18.9

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

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N,N-Dimethyl-*N'*-phenylformamidinium perchlorate

The molecular structure of the title compound, $C_9H_{13}N_2^+ \cdot ClO_4^-$, consists of an *N*,*N*-dimethyl–*N'*-phenyl-formamidinium cation and a perchlorate anion. The latter forms five hydrogen bonds with three different formamidinium cations, using three O atoms. The cations and anions are linked through C–H···O and N–H···O hydrogen bonds, forming a ring; the centrosymmetrically related rings form a staircase *via* C–H···O hydrogen bonds.

Comment

There is considerable interest in the chemistry and structure of iminium salts, because of their importance in organic synthesis (Boehme & Viehe, 1976). The chromophores of visual pigments contain polyunsaturated iminium salts (Uhl & Abrahamson, 1981; Brige, 1981). The iminium salts exhibit optical properties and the position of the counterion plays a role in determining the optical properties of the iminium cation (Arnaboldi *et al.*, 1979).



The cation of the title compound, (I), is nearly planar and the dihedral angle between the phenyl ring and the Schiff base chain is 7.7 (1)°. Protonation occurs at N9, and the positive charge is delocalized between atoms N9 and N7, as evidenced by the bond distances N7–C8 [1.316(3) Å] and C8=N9 [1.294 (3) Å]. This is confirmed by the coplanarity of atoms C1, N7, C8, N9, C10 and C11 [maximum deviation is 0.031 (4) Å for C10]. The perchlorate anion accepts five hydrogen bonds from three different formamidinium cations, using three O atoms. One of the perchlorate O atoms (O4) acts as an acceptor in three hydrogen bonds and this O atom is also characterized by an extended bond of length 1.426 (2) Å. The cations and anions are linked through $N7-H7\cdots O4^{i}$ and C8-H8···O3ⁱⁱ hydrogen bonds and form a ring, which is further reinforced by two C2-H2···O4ⁱ and C11-H11D···O4ⁱ hydrogen bonds. The centrosymmetrically related rings are linked via C11-H11E···O1ⁱⁱⁱ interactions and form a staircase, which runs parallel to the b axis (see Table 2 for symmetry codes).

Experimental

Aniline (0.05 mol) was dissolved in 6 ml DMF and kept under icecold conditions. To this, 1.4 ml $POCl_3$ (0.035 ml) was added dropwise for 15 min with stirring at 273 K. The reaction mixture was stirred at room temperature for 30 minutes and then at 303 K for 15 h. The Received 4 March 2003 Accepted 4 April 2003 Online 23 April 2003

 $\Delta \rho_{\rm max} = 0.34 \ {\rm e}$

 $\Delta \rho_{\min}$

−0.22 e Å

Extinction correction: SHELXL97

Extinction coefficient: 0.007 (2)



Figure 1

The molecular structure of the title compound, (I), showing 35% probability displacement ellipsoids.



Figure 2

Part of the crystal packing of the title compound, showing hydrogenbonded networks of cations and anions. Atoms marked with a dollar (\$), hash (#) or asterisk (*) are at the symmetry positions (1 - x, -y, -z), (1 - x, -y, 1 - z) and $(1 - x, y - \frac{1}{2}, \frac{1}{2} - z)$, respectively.

reaction mixture was finally poured into crushed ice and treated with sodium perchlorate or dilute perchloric acid. The resulting creamy yellow precipitate was filtered, washed with water and dried. This crude product was recrystallized using CHCl₃ to give pure title compound.

Crystal data

$C_9H_{13}N_2ClO_4$	$D_x = 1.433 \text{ Mg m}^{-3}$
$M_r = 248.66$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 3903
a = 7.8382 (2) Å	reflections
b = 17.5688 (6) Å	$\theta = 2.3 - 28.3^{\circ}$
c = 9.1123 (2) Å	$\mu = 0.33 \text{ mm}^{-1}$
$\beta = 113.269 \ (1)^{\circ}$	T = 293 (2) K
V = 1152.77 (6) Å ³	Prism, pale yellow
Z = 4	$0.28 \times 0.16 \times 0.12 \text{ mm}$

Data collection

Siemens SMART CCD area- detector diffractometer	2854 independent reflections 1917 reflections with $I > 2\sigma(I)$
w scans	$R_{int} = 0.039$
Absorption correction: multi-scan	$\theta_{\rm max} = 28.3^{\circ}$
(SADABS; Siemens, 1996)	$h = -10 \rightarrow 8$
$T_{\rm min} = 0.913, T_{\rm max} = 0.961$	$k = -15 \rightarrow 23$
7777 measured reflections	$l = -11 \rightarrow 12$
Refinement	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0752P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.051$	+ 0.2933P]
$wR(F^2) = 0.153$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} = 0.001$

2854 reflections 151 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected geometric parameters (Å, °).

Cl1-O1	1.406 (3)	N7-C8	1.316 (3)
Cl1-O2	1.407 (2)	C8-N9	1.294 (3)
Cl1-O3	1.419 (2)	N9-C10	1.465 (3)
Cl1-O4	1.426 (2)	N9-C11	1.466 (3)
C1-N7	1.422 (3)		
C2-C1-N7	117.6 (2)	C8-N9-C10	120.0 (2)
C6-C1-N7	122.6 (2)	C8-N9-C11	123.2 (2)
C8-N7-C1	125.3 (2)	C10-N9-C11	116.7 (2)
N9-C8-N7	126.0 (2)		
C2-C1-N7-C8	173.9 (2)	N7-C8-N9-C10	177.0 (2)
C6-C1-N7-C8	-6.5(4)	N7-C8-N9-C11	-0.7(4)
C1-N7-C8-N9	-179.9 (2)		()

Table 2		
Hydrogen-bonding geometry	(Å,	°).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D{\cdots}A$	$D - \mathbf{H} \cdots A$
C11-H11DN7	0.96	2.47	2.875 (3)	105
$N7 - H7 \cdots O4^i$	0.82 (3)	2.09 (3)	2.877 (3)	160 (3)
$C2-H2\cdots O4^i$	0.93	2.56	3.295 (3)	134
$C11 - H11D \cdots O4^{i}$	0.96	2.41	3.329 (3)	160
C8-H8···O3 ⁱⁱ	0.93	2.40	3.174 (4)	140
$C11 - H11E \cdots O1^{iii}$	0.96	2.38	3.310 (4)	162
	(**) 1	1	() 1	1.1

Symmetry codes: (i) 1 - x, -y, -z; (ii) 1 - x, -y, 1 - z; (iii) $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$.

The H atom on N7 was located in a difference Fourier map and refined, while all other H atoms were positioned geometrically and were allowed to ride on their attached atoms. One of the methyl groups (C11) was found to be disordered; it was treated as an idealized disordered methyl group, with two positions rotated from each other by 60° , and the site-occupation factors were fixed at 0.5.

Data collection: SMART (Siemens, 1996): cell refinement: SAINT: data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ZORTEP (Zsolnai, 1997) and PLUTON (Spek, 1990); software used to prepare material for publication: SHELXL97 and PARST (Nardelli, 1995).

Financial support from the University Grants Commission (UGC) and the Department of Science and Technology (DST), Government of India, are gratefully acknowledged.

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