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#### **Key indicators**

Single-crystal X-ray study T = 130 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.039 wR factor = 0.090 Data-to-parameter ratio = 11.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. Received 14 March 2005

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# O-(2,3,4,5,6-Pentafluorobenzyl)hydroxylaminium chloride

The crystal structure of the title compound,  $C_7H_5F_5NO^+\cdot Cl^-$ , consists of two-dimensional networks of hydrogen-bonded primary  $NH_3^+$  cations and chloride anions with the pentafluorobenzyl moieties located on one side of the layer. To maximize the Coulombic interactions, the  $Cl^-$  anions approach the  $NH_3^+$  groups of the neighbouring network along the O-N bond direction  $[O-N\cdots Cl = 174.3 (2)^\circ]$ . The N $\cdots$ Cl distance between the ions belonging to two adjacent networks is 3.066 (2) Å and is shorter than the  $N\cdots Cl$ distances in the  $N^+-H\cdots Cl^-$  hydrogen bonds [3.099 (3)-3.133 (3) Å].

### Comment

Primary ammonium halides are known to assemble in crystal structurs *via*  $N^+-H\cdots Cl^-$  hydrogen bonds into ladders, cubanes and two-dimensional networks fully utilizing the hydrogen-bonding ability of the *R*NH<sub>3</sub><sup>+</sup> group (Gray & Jones, 2002*a*,*b*; Lopez-Dupla *et al.*, 2003; Bond, 2005). Besides hydrogen bonds, to maximize Coulombic interactions within the crystal structure, ammonium groups and halide anions often form additional short N<sup>+</sup>…·Cl<sup>-</sup> contacts (Bond, 2005).



Our interest in the title compound, (I), originates from recent reports on the interactions between anions and electron-deficient  $\pi$ -acidic rings (Quiñonero *et al.*, 2002; Garau, Frontera *et al.*, 2003; Garau, Quiñonero *et al.*, 2003; Domeshko *et al.*, 2004). Quiñonero *et al.* (2002) provided computational evidence for non-bonding interactions between hexafluorobenzene and several anions and found 27 structures in the Cambridge Structural Database (Allen, 2002) where the interaction of anions with pentafluorobenzene derivatives could be postulated. By solving the crystal structure of (I), we aimed to investigate whether the pentafluorophenyl moiety is able to compete for Coulombic interactions with the NH<sub>3</sub><sup>+</sup> group. We expected that in (I) the ammonium group attached *via* a flexible  $-CH_2-O-$  arm to a pentafluorophenyl ring should co-operate with the aromatic ring in the anion binding.

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## organic papers



View of the asymmetric unit of (I), shown with 50% displacement ellipsoids.







The crystal structure of (I), showing (a) the hydrogen-bonded layer of cations and anions parallel to the (100) plane and (b) the arrangement of two layers in the crystal structure. Hydrogen bonds are shown as brown dashed lines and short  $N^+ \cdots Cl^-$  interlayer contacts as violet dashed lines.

The structure of (I), with its atom numbering, is shown in Fig. 1. Analysis of the crystal packing revealed that the postulated anion- $\pi$  ring interactions do not exist in (I). Primary ammonium groups and chloride anions form three  $N^+$ -H···Cl<sup>-</sup> hydrogen bonds (Table 1), generating twodimensional networks composed of  $R_6^3$  motifs (Fig. 2a). The six-membered rings of the network have a very flattened chair conformation. The pentafluorobenzyl moieties are grouped on one side of the layer, resulting in its polar structure. A similar polar two-dimensional assembly was observed in anilinium chloride (Lopez-Dupla et al., 2003) and is rather exceptional among ammonium halides (Bond, 2005). However, there is a striking difference between the two structures. Crystals of anilinium chloride are polar. *i.e.* adjacent lavers are in contact via their hydrophilic and hydrophobic faces. In (I), the Coulombic attraction between oppositely charged units located on the hydrophilic side of the layer leads to a centrosymmetric structure (Fig. 2b). To maximize the Coulombic interactions, the Cl<sup>-</sup> anions approach the NH<sub>3</sub><sup>+</sup> groups of the neighbouring network along the cation O-N bond direction  $[O - N \cdots Cl = 174.3 (2)^{\circ}]$ . The N  $\cdots Cl$  distance between the two networks is 3.066 (2) Å and is shorter than the N···Cl distances of the N<sup>+</sup>-H···Cl<sup>-</sup> hydrogen bonds [3.099 (3)-3.133 (3) Å].

#### **Experimental**

The title compound was purchased from Aldrich and recrystallized from ethanol [m.p. 500 K (sublimation)].

Crystal data	
$C_7H_5F_5NO^+\cdot Cl^-$	$D_{\rm r} = 1.725 {\rm Mg} {\rm m}^{-3}$
$M_r = 249.57$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2766
a = 19.994 (3) Å	reflections
b = 5.4020 (8)  Å	$\theta = 4-25^{\circ}$
c = 9.0305 (15)  Å	$\mu = 0.45 \text{ mm}^{-1}$
$\beta = 99.939 \ (13)^{\circ}$	T = 130 (2)  K
$V = 960.7 (3) \text{ Å}^3$	Needle, colourless
Z = 4	$0.42 \times 0.12 \times 0.03 \text{ mm}$
Data collection	
Kuma KM4CCD κ-geometry	1042 reflections with $I > 2\sigma(I)$
diffractometer	$R_{\rm int} = 0.038$
$\omega$ scans	$\theta_{\rm max} = 25.0^{\circ}$
Absorption correction: none	$h = -23 \rightarrow 23$
3889 measured reflections	$k = -6 \rightarrow 6$
1702 independent reflections	$l = -10 \rightarrow 7$
Refinement	
Refinement on $F^2$	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.039$	independent and constrained
$wR(F^2) = 0.090$	refinement
S = 0.90	$w = 1/[\sigma^2(F_o^2) + (0.0436P)^2]$
1702 reflections	where $P = (F_0^2 + 2F_c^2)/3$

Table 1	
Hydrogen-bonding geometry	(Å, °).

148 parameters

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H1A \cdots Cl1^{i}$ $N1 - H1B \cdots Cl1^{ii}$ $N1 - H1C \cdots Cl1^{iii}$	0.98 (4) 1.01 (3) 1.02 (4)	2.15 (4) 2.11 (3) 2.11 (4)	3.107 (3) 3.099 (3) 3.133 (3)	165 (3) 166 (2) 177 (3)
	1.3	(11) A A		1.3

 $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.52 \ {\rm e} \ {\rm \AA}^{-3}$  $\Delta \rho_{\rm min} = -0.26 \ {\rm e} \ {\rm \AA}^{-3}$ 

Symmetry codes: (i)  $1 - x, \frac{1}{2} + y, \frac{3}{2} - z$ ; (ii) 1 - x, 1 - y, 1 - z; (iii)  $1 - x, y - \frac{1}{2}, \frac{3}{2} - z$ .

All H atoms were located in electron-density difference maps. The C-H bond distances were standardized to 0.97 Å and these H atoms were then refined as riding. Positional parameters of H atoms from the NH<sub>3</sub><sup>+</sup> group and isotropic displacement parameters of all H atoms were allowed to refine.

Data collection: CrysAlis CCD (Oxford Diffraction, 2004); cell refinement: CrysAlis CCD; data reduction: CrysAlis RED (Oxford Diffraction, 2004); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: *Stereochemical Workstation Operation Manual* (Siemens, 1989) and *MERCURY* (Bruno *et al.*, 2002); software used to prepare material for publication: *SHELXL97*.

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