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The C— $H \cdots O$ hydrogen bond in (dicyanomethyl)ammonium *p*-toluenesulfonate

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In the title compound, $C_3H_4N_3^+ \cdot C_7H_7O_3S^-$, the activated C— H group of the cation forms a short but bent C—H···O hydrogen bond with a sulfonate O atom of the anion; C···O = 3.075 (5) Å and C—H···O = 130°.

Comment

 $C-H \cdots O$ hydrogen bonds are known to have a very wide range of geometries and strengths. With highly polar C-H groups, such as in CHCl₃, C≡C−H, CH(NO₂)₃ etc., C− $H \cdots O$ interactions may have similar geometries to conventional $O/N - H \cdots O$ hydrogen bonds. On the other hand, with weakly polar methyl groups, C-H···O interactions have long contact distances and are only slightly directional. All intermediate situations exist between these extremes (see, e.g., Steiner, 1997; Steiner & Desiraju, 1998; Desiraju & Steiner, 1999). When studying the stronger kinds of $C-H \cdots O$ interactions, methyl groups which carry two or even three strongly electron-withdrawing substituents are of interest (Pedireddi & Desiraju, 1992). In this context, the (dicyanomethyl)ammonium cation is of obvious interest. In the p-toluenesulfonate salt, (I), formation of a short $C-H \cdots O$ hydrogen bond may be expected.



In the crystal structure of (I), the three ammonium H atoms form N-H···O hydrogen bonds with the sulfonate group (Table 2). As expected, the activated C-H group of the cation also interacts with a sulfonate O atom (Fig. 1). The C···O distance is very short at 3.075 (5) Å, but the geometry is strongly bent. Based on a normalized H-atom position (C-H = 1.08 Å), the H···O distance is 2.27 Å and the C- $H \cdots O$ angle is 130°. When compared to the shortest C- $H \cdots O$ interactions known, which have $H \cdots O$ distances around and below 2.0 Å (e.g. Bock et al., 1993; Kariuki et al., 1997), this may not appear to be so short. However, upon closer examination, it is seen that the C-H donor is involved in two additional intermolecular interactions with hydrogenbond geometry, but both have substantially longer distances (Table 2). The crystal packing as a whole is shown in Fig. 2 to illustrate the typical segregation of polar and apolar groups in the lattice. An important factor responsible for the poor C- $H \cdots O$ angle might be the competition between C-H and the neighbouring ammonium group for acceptors. The ammonium group certainly plays the more dominant role in determining intermolecular geometries, and optimizing N-H···O geometries might prevent the $C-H \cdots O$ angle from being linear.

To see how the C-H···O hydrogen bond in (I) compares with related interactions of dicyanomethyl C-H donors, a database search was performed (Cambridge Structural Database, version 5.20; Allen & Kennard, 1993). In the 15 relevant crystal structures found, the shortest occurring (C-)H···O distance is 2.12 Å (in a crown-ether complex of dicyanomethane; Grootenhuis *et al.*, 1986), with more typical values being greater than 2.2 Å. This indicates that (CN)₂C-H is a good hydrogen-bond donor in general, but clearly falls behind the classical strong C-H donors like CHCl₃ and C=C-H.



Figure 1

The molecular structure of (I) with displacement ellipsoids drawn at the 30% probability level.



Figure 2 The crystal packing shown in a projection on the *yz* plane.

Experimental

Compound (I) was obtained from Lancaster Synthesis and was recrystallized from MeOH by slow evaporation of the solvent.

> $D_x = 1.384 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 25 reflections $\theta=8.2{-}14.9^\circ$ $\mu = 0.27~\mathrm{mm}^{-1}$

T = 293 (2) KPlate, colourless $0.80 \times 0.30 \times 0.05 \text{ mm}$

 $h = -7 \rightarrow 4$

 $k = -25 \rightarrow 33$

 $l = -10 \rightarrow 10$

3 standard reflections

+ 0.4531P]

where $P = (F_o^2 + 2F_c^2)/3$

frequency: 90 min

intensity decay: 3%

Crystal data

$C_{3}H_{4}N_{3}^{+} \cdot C_{7}H_{7}O_{3}S$
$M_r = 253.28$
Monoclinic, $P2_1/n$
a = 5.604 (9) Å
b = 25.90(5) Å
c = 8.427 (12) Å
$\beta = 96.40 \ (14)^{\circ}$
$V = 1216 (4) \text{ Å}^3$
Z = 4

Data collection

Stoe four-circle diffractometer ω scans 3001 measured reflections 2660 independent reflections 2162 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.019$ $\theta_{\rm max} = 27.0^\circ$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0656P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.126$ S = 1.03 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.45 \ {\rm e} \ {\rm \AA}^{-3}$ 2660 reflections $\Delta \rho_{\rm min} = -0.39 \ {\rm e} \ {\rm \AA}^{-3}$ 171 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected geometric parameters (Å, °).

N1-C8	1.473 (4)	C8-C10	1.469 (4)
N2-C9 N3-C10	1.127 (4) 1.120 (4)	C8-C9	1.473 (4)
C10-C8-C9 C10-C8-N1	110.3 (2) 111.1 (2)	C9-C8-N1	109.7 (2)

H atoms bonded to C atoms of the anion were treated in the default riding model with $U_{\rm eq}$ values allowed to vary. The H atoms of the cation were located in difference Fourier calculations and refined isotropically [U_{eq} values: H(N1) = 0.043–0.057 and H(C8) = 0.040 Å²].

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N1-H1A\cdots O2^{i}$	0.91 (3)	1.84 (3)	2.741 (4)	175 (2)
$N1 - H1B \cdot \cdot \cdot O3^{ii}$	0.93 (3)	1.78 (3)	2.701 (4)	168 (2)
$N1 - H1C \cdot \cdot \cdot O1^{iii}$	0.92 (3)	1.87 (3)	2.778 (5)	167 (3)
C8−H8···O1	0.95 (3)	2.35 (2)	3.075 (5)	132.3 (19)
C8−H8···O2 ⁱⁱⁱ	0.95 (3)	2.73 (2)	3.373 (5)	125.8 (17)
$C8-H8\cdots N2^{iv}$	0.95 (3)	2.78 (3)	3.409 (6)	124.2 (17)
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Symmetry codes: (i) x, y, 1 + z; (ii) $x - \frac{1}{2}, \frac{1}{2} - y, \frac{1}{2} + z$; (iii) $\frac{1}{2} + x, \frac{1}{2} - y, \frac{1}{2} + z$; (iv) 1 + x, y, z.

Data collection: Stoe diffractometer software; cell refinement: Stoe diffractometer software: data reduction: Stoe diffractometer software; program(s) used to solve structure: SHELXS86 (Sheldrick, 1985); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 1990); software used to prepare material for publication: SHELXL97.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: DE1169). Services for accessing these data are described at the back of the journal.

References

Allen, F. H. & Kennard, O. (1993). Chem. Des. Autom. News, 8, 1, 31-37. Bock, H., Dienelt, R., Schödel, H. & Havlas, Z. (1993). J. Chem. Soc. Chem.

Commun. pp. 1792-1793. Desiraju, G. R. & Steiner, T. (1999). The Weak Hydrogen Bond in Structural Chemistry and Biology. Oxford University Press.

Grootenhuis, P. D. J., van Eerden, J., Dijkstra, P. J., Harkema, S. & Reinhoudt, D. N. (1986). J. Am. Chem. Soc. 109, 8044-8051.

Kariuki, B. M., Harris, K. D. M., Philip, D. & Robinson, J. M. A. (1997). J. Am. Chem. Soc. pp. 12679-12680.

Pedireddi, V. R. & Desiraju, G. R. (1992). J. Chem. Soc. Chem. Commun. pp. 988_990

Sheldrick, G. M. (1985). SHELXS86. University of Göttingen, Germany.

Sheldrick, G. M. (1997). SHELXL97. University of Göttingen, Germany.

Spek, A. L. (1990). Acta Cryst. A46, C-34.

Steiner, T. (1997). Chem. Commun. pp. 727-734.

Steiner, T. & Desiraju, G. R. (1998). Chem. Commun. pp. 891-892.