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Key indicators

Single-crystal X-ray study
 T = 100 K
 Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$
 R factor = 0.033
 wR factor = 0.082
 Data-to-parameter ratio = 10.2

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

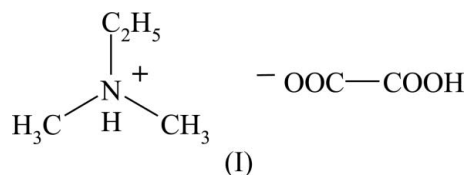
Ethylidimethylammonium oxalate

The title crystal structure, $\text{C}_4\text{H}_{12}\text{N}^+\cdot\text{C}_2\text{HO}_4^-$, contains discrete ethylidimethylammonium cations and oxalate monoanions. Linear chains of oxalate monoanions are formed by strong $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds. These chains are further connected by $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds through ethylidimethylammonium cations, forming channels along the *a* axis.

Received 14 November 2006
 Accepted 21 November 2006

Comment

Supramolecular hydrogen-bonded assemblies of oxalic acid and organic amines (alkylammonium, guanidine, 1,4-diazabicyclo[2.2.2]octane, *etc.*) have been examined by single-crystal X-ray diffraction and other techniques (MacDonald *et al.*, 2001; Vaidhyanathan *et al.*, 2001, 2002; Ejsmont & Zaleski, 2006*a,b*).



The asymmetric unit of the title compound, (I), consists of one ethylidimethylammonium cation and one oxalate monoanion (Fig. 1). The geometrical parameters of the ethylidimethylammonium cation (Table 1) compare well with those

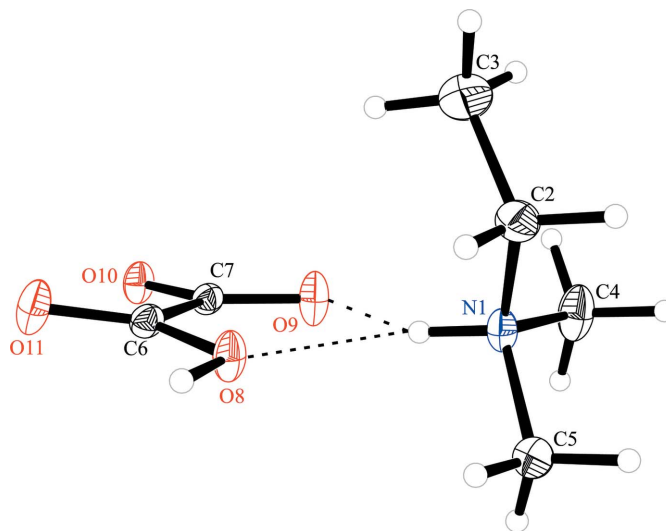


Figure 1
 The asymmetric unit of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. Hydrogen bonds are shown as dashed lines.

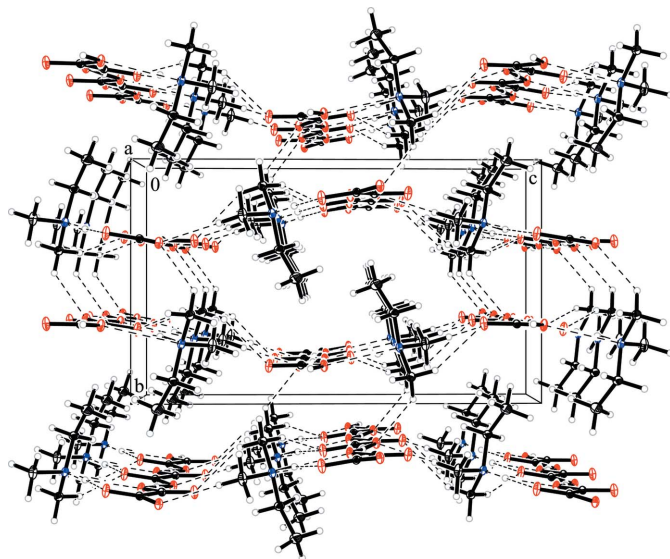


Figure 2

A packing plot of (I), showing the intermolecular hydrogen-bonding scheme (dashed lines).

found in other crystal structures which include this cation (Bujak & Zaleski, 1999; Scheibitz *et al.*, 2003; Molteni *et al.*, 2004). The oxalate monoanions are essentially planar, with dihedral angles between the carboxylate groups of less than 10° . A strong $O8-H8 \cdots O10^i$ (see Table 2 for symmetry code) hydrogen bond generates linear oxalate chains running parallel to the *a* axis. The geometrical parameters of this chain correlate well with the corresponding values found in related crystal structures (Vaidhyanathan *et al.*, 2002; Ejsmont & Zaleski, 2006b). In addition, the crystal structure of (I) is stabilized by ionic interactions between the ethyldimethylammonium cations and the oxalate monoanions chains, as well as by a network of $N-H \cdots O$ and $C-H \cdots O$ hydrogen bonds (Fig. 2 and Table 2). In the crystal structure of (I), there are channels running along the *a* axis. The dimensions of these channels, measured by the distance between methyl H atoms of the ethyldimethylammonium cation and between the mean planes of the oxalate monoanions chains, are 5.073 (5) and 3.267 (4) Å, respectively.

Experimental

Crystals of (I) were grown at room temperature by slow evaporation of an aqueous solution containing ethyldimethylamine and oxalic acid in a 2:1 stoichiometric ratio.

Crystal data

$C_4H_{12}N^+ \cdot C_2HO_4^-$	$Z = 4$
$M_r = 163.17$	$D_x = 1.305 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 5.5953 (7) \text{ \AA}$	$\mu = 0.11 \text{ mm}^{-1}$
$b = 9.3979 (9) \text{ \AA}$	$T = 100.0 (1) \text{ K}$
$c = 15.804 (2) \text{ \AA}$	Block, colourless
$\beta = 92.158 (9)^\circ$	$0.42 \times 0.40 \times 0.38 \text{ mm}$
$V = 830.45 (17) \text{ \AA}^3$	

Data collection

Oxford Diffraction Xcalibur diffractometer
 ω scans
 Absorption correction: none
 4871 measured reflections

1545 independent reflections
 1185 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$
 $\theta_{\text{max}} = 25.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.082$
 $S = 0.98$
 1545 reflections
 152 parameters
 All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.0778P)^2 + 0.1154P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.27 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

N1—C4	1.485 (2)	C6—O8	1.306 (1)
N1—C5	1.490 (2)	C6—C7	1.552 (2)
N1—C2	1.499 (2)	C7—O9	1.237 (2)
C6—O11	1.212 (2)	C7—O10	1.266 (2)
O11—C6—O8	125.9 (1)	O9—C7—O10	126.7 (1)
O11—C6—C7	121.7 (1)	O9—C7—C6	118.6 (1)
O8—C6—C7	112.4 (1)	O10—C7—C6	114.7 (1)
O11—C6—C7—O9	170.4 (1)	O11—C6—C7—O10	−8.4 (2)
O8—C6—C7—O9	−8.8 (2)	O8—C6—C7—O10	172.4 (1)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N1—H1 \cdots O8	0.92 (2)	2.19 (2)	2.931 (1)	136 (1)
N1—H1 \cdots O9	0.92 (2)	1.98 (2)	2.781 (1)	144 (1)
O8—H8 \cdots O10 ⁱ	1.12 (2)	1.35 (2)	2.473 (1)	177 (2)
C4—H4B \cdots O10 ⁱⁱ	1.00 (2)	2.30 (2)	3.264 (2)	161 (1)
C5—H5A \cdots O10 ⁱⁱⁱ	0.99 (2)	2.40 (2)	3.213 (2)	140 (1)
C5—H5B \cdots O9 ⁱ	0.98 (2)	2.49 (2)	3.421 (2)	160 (1)
C5—H5C \cdots O11 ⁱⁱ	1.00 (1)	2.54 (1)	3.449 (2)	151 (1)

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2002); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2002); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 1990); software used to prepare material for publication: *SHELXL97*.

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