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Bis(2-aminopyridinium) maleate

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Two cyclic eight-membered hydrogen-bonded rings exist in the title compound, $2C_5H_7N_2^+ \cdot C_4H_2O_4^{2-}$, involving the 2-aminopyridinium and maleate ions. The dihedral angle between the two pyridinium rings hydrogen bonded to the maleate ion is 74.80 (4) $^{\circ}$. The maleate anion lies on a twofold axis and is linked to the pyridinium cations by intermolecular N-H···O hydrogen bonds. The heterocycle is fully protonated, which enables amino-imino tautomerization.

Comment

The present work is part of a structural study of complexes of 2-aminopyridinium systems with hydrogen-bond donors, and we report here the structure of bis(2-aminopyridinium) maleate, (I). A similar series of complexes formed from 2-aminopyridine and carboxylate have been reported recently (Büyükgüngör & Odabaşoğlu, 2002; Odabaşoğlu et al., 2003).



A view of (I) is shown in Fig. 1. The complex owes its formation to two hydrogen-bond pairs, one between atoms O1 and O2 of the maleate ions and hydrogen-bond donors N1-H1 and N2-H2A, and the other between their symmetryrelated pairs. There are two eight-membered rings in the structure, formed as a result of these N-H···O hydrogen bonds. Furthermore, there is also an intermolecular hydrogen bond in (I) (Table 2).



The average C-O distances in carboxylate groups that form intermolecular hydrogen bonds are 1.32 (2) Å for the

hydroxyl C–OH bond and 1.21 (3) Å for the carbonyl C=O bond (Borthwick, 1980). The value for the carboxylate anion is also reported as 1.25 Å (Borthwick, 1980). The O1-C11 and O2-C11 bond lengths in (I) thus fall into the category of a normal COO⁻ group (Table 1).

The C-N-C angle of pyridines is very sensitive to protonation (Jin, Pan, Xu & Xu, 2000; Jin et al., 2002). In comparison with 2-amino-6-methylpyridinium neoabietate (Jin, Pan, Liu & Xu, 2000), the complete protonation of the heterocycle in (I) is indicated by the enlarged C1-N1-C5 angle $[122.69 (15)^{\circ}]$ and the reduced N1-C1-C2 angle [117.47 (16)°]. The 2-aminopyridine-carboxylic acid system has been the subject of theoretical (Inuzuka & Fujimoto, 1990) and spectroscopic (Inuzuka & Fujimoto, 1986) amino-imino tautomerization studies. The main features of amino-imino tautomerization (see Scheme 1) are demonstrated in the structure of (I) by the bond lengths and angles of the



Figure 1

A view of the moieties of (I), showing the atom-numbering scheme and 50% probability displacement ellipsoids [symmetry code: (i) $\frac{1}{2} - x$, $\frac{1}{2} - y, z$].



Figure 2 A packing diagram for (I), viewed along the c axis.

organic compounds

heterocycle and the maleate anion, respectively. The present investigation, like our previous work (Büyükgüngör & Odabaşoğlu, 2002; Odabaşoğlu *et al.*, 2003), clearly shows that the positive charge in the 2-aminopyridinium ions of (I) is on the amino group.

Experimental

The title compound was prepared by dissolving 2-aminopyridine and maleic acid in a 2:1 molar ratio in water at 373 K. Crystals of (I) were obtained by slow evaporation of the solvent at the room temperature.

Crystal data

$2C_{5}H_{7}N_{2}^{+} \cdot C_{4}H_{2}O_{4}^{2-}$ $M_{r} = 304.30$ Orthorhombic, <i>Fdd2</i> a = 21.756 (5) Å b = 23.531 (5) Å c = 5.6280 (11) Å V = 2881.2 (11) Å ³ Z = 8 $D_{x} = 1.403$ Mg m ⁻³	Mo K α radiation Cell parameters from 161 reflections $\theta = 2.6-12.7^{\circ}$ $\mu = 0.11 \text{ mm}^{-1}$ T = 153 (2) K Rectangular, light yellow 0.20 × 0.15 × 0.10 mm
Data collection	
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Siemens P4 diffractometer	$h = -27 \rightarrow 5$
ω scans	$k = -30 \rightarrow 0$
1564 measured reflections	$l = -6 \rightarrow 7$
873 independent reflections	1 standard reflection
814 reflections with $I > 2\sigma(I)$	frequency: 120 min
$R_{\rm int} = 0.016$	intensity decay: <2%
$\theta_{\rm max} = 27^{\circ}$	

Table 1

Selected geometric parameters (Å, °).

N1-C1	1.3434 (19) 1.354 (2)	01-C11 02 C11	1.264 (2)
N1-C3 N2-C1	1.318 (2)	C11-C12	1.500 (2)
C1-C2	1.412 (3)	$C12 - C12^{i}$	1.323 (3)
C1-N1-C5	122.69 (15)	N1-C1-C2	117.47 (16)
Symmetry code: (i) $\frac{1}{2}$	$-x, \frac{1}{2}-y, z.$		

Table 2

Hydrogen-bonding geometry (Å, $^{\circ}$).

D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
0.88	1.78	2.6557 (18)	172
0.88	1.96	2.837 (2)	177
0.88	1.97	2.832 (2)	168
	<i>D</i> -H 0.88 0.88 0.88	D−H H···A 0.88 1.78 0.88 1.96 0.88 1.97	$D-H$ $H\cdots A$ $D\cdots A$ 0.881.782.6557 (18)0.881.962.837 (2)0.881.972.832 (2)

Symmetry code: (i) $x - \frac{1}{4}, \frac{1}{4} - y, z - \frac{1}{4}$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0451P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.026$	+ 0.5722P]
$wR(F^2) = 0.072$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} < 0.001$
873 reflections	$\Delta \rho_{\rm max} = 0.11 \ {\rm e} \ {\rm \AA}^{-3}$
101 parameters	$\Delta \rho_{\rm min} = -0.11 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
	Extinction coefficient: 0.0048 (6)

Although space group Cc gave a chemically reasonable and computationally stable refinement, the correct space group was found to be *Fdd2*. Refinement of the absolute structure parameter was meaningless because of its large s.u. (1.2), and so Friedel-pair reflections were averaged before the final refinement. All H atoms were treated using a riding model, with C–H distances of 0.95 Å and N–H distances of 0.88 Å, and with $U_{iso}(H) = 1.2U_{eq}(parent)$.

Data collection: *XSCANS* (Siemens, 1991); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Sheldrick, 1990); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1998).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1408). Services for accessing these data are described at the back of the journal.

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