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#### Key indicators

Single-crystal X-ray study T = 293 KMean  $\sigma(C-C) = 0.003 \text{ Å}$  R factor = 0.042 wR factor = 0.102 Data-to-parameter ratio = 13.2

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

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# Strong symmetrical intramolecular hydrogen bonds in diprotonated piperazinium bis(hydrogenmaleate)

In the title salt,  $C_4H_{12}N_2^{2+}\cdot 2C_4H_3O_4^{-}$ , the diprotonated piperazine cation links to two hydrogen maleate anions through N-H···O hydrogen bonds. An H atom is approximately centered in an extremely short intramolecular hydrogen bond of the O···H···O type in each of the hydrogen maleate anions.

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### Comment

Maleic acid may bind to cations in either the monoanionic or dianionic form, displaying a plethora of interesting structures. Piperazine is a strongly basic amine, able to form a dication, in which all four N-H bonds are generally active in hydrogenbond formation (Coupar *et al.*, 1996; Ferguson *et al.*, 1998; Loehlin *et al.*, 1994). Many cases of symmetrical intramolecular hydrogen bonds have been observed in the hydrogen maleate anion, with an H atom centered in an extremely short  $O \cdots H \cdots O$  intramolecular hydrogen bond (Darlow, 1961; Darlow & Cochran, 1961; Fillaux *et al.*, 1999). In this work, the combination of piperazine and maleic acid has been chosen to explore the special hydrogen-bond pattern.



The crystal structure shows a local pseudocenter of symmetry at x = 0.887 (3), y = 0.757 (9), z = 0.384 (5). Adopting a normal chair conformation, as reported previously (Bois, 1970; Kuppayee *et al.*, 1999; Jin *et al.*, 2001), the dication of the title salt, (I), links to two anions through N1-H···O1 and N2-H···O8 hydrogen bonds, where the H atoms are equatorial.

In (I), two hydrogen maleate anions each hold an H atom at the approximate center of the extremely short intramolecular hydrogen bonds  $O_2 \cdots H1' \cdots O_4$  and  $O_7 \cdots H2' \cdots O_5$ . With the centered H atoms in the intramolecular hydrogen bonds, atoms O<sub>2</sub>, O<sub>4</sub>, O<sub>5</sub> and O<sub>7</sub> are both hydrogen donors and acceptors. The O<sub>2</sub>-C<sub>1</sub>, O<sub>4</sub>-C<sub>4</sub>, O<sub>5</sub>-C<sub>5</sub> and O<sub>7</sub>-C<sub>8</sub> distances are longer than those of O<sub>1</sub>-C<sub>1</sub>, O<sub>3</sub>-C<sub>4</sub>, O<sub>6</sub>-C<sub>5</sub> and O<sub>8</sub>-C<sub>8</sub>, showing differences between the C-O and C=O bond types of the carboxylic acid group (Borthwick, 1980).

Supramolecular layers of the title salt (Fig. 2) are formed parallel to the *ac* plane by N1-H···O1, N2-H···O8, N1-H···O3<sup>i</sup> and N2-H···O6<sup>ii</sup> hydrogen bonds (see Table 2).

# organic papers



Figure 1

The molecular structure of the title salt, showing 35% probability displacement ellipsoids; hydrogen bonds are illustrated by dashed lines.

Furthermore, these layers are stabilized by  $C10-H \cdot \cdot \cdot O4^{i}$  and  $C12-H \cdot \cdot \cdot O5^{ii}$  hydrogen bonds. There are no hydrogen bonds between these layers.

### **Experimental**

Crystals of the title salt were obtained from a 1:2 aqueous solution of piperazine and maleic acid by slow concentration for a week.

 $D_x = 1.490 \text{ Mg m}^{-3}$ 

Cell parameters from 22

 $0.46 \times 0.30 \times 0.20 \text{ mm}$ 

3 standard reflections

every 97 reflections intensity decay: 1.3%

Mo  $K\alpha$  radiation

reflections

 $\mu = 0.13 \text{ mm}^{-1}$ 

T = 293 (2) K

 $\begin{array}{l} h = 0 \rightarrow 13 \\ k = 0 \rightarrow 14 \end{array}$ 

 $l = -14 \rightarrow 13$ 

Prism, colorless

 $\theta = 4.0 - 12.6^{\circ}$ 

Crystal data

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\begin{array}{l} C_4 H_{12} N_2^{2+} \cdot 2 C_4 H_3 O_4^{-} \\ M_r = 318.28 \\ \text{Monoclinic, } P_2_1/n \\ a = 10.735 \ (2) \ \text{\AA} \\ b = 11.603 \ (2) \ \text{\AA} \\ c = 11.843 \ (2) \ \text{\AA} \\ \beta = 105.87 \ (1)^{\circ} \\ V = 1418.9 \ (4) \ \text{\AA}^3 \\ Z = 4 \end{array}
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#### Data collection

Siemens *P*4 diffractometer  $\omega$  scans 2775 measured reflections 2631 independent reflections 1500 reflections with *I* > 2 $\sigma$ (*I*) *R*<sub>int</sub> = 0.028  $\theta_{max} = 25.5^{\circ}$ 

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0698P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.102$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 0.91	$\Delta \rho_{\rm max} = 0.14 \text{ e } \text{\AA}^{-3}$
2631 reflections	$\Delta \rho_{\rm min} = -0.17 \text{ e} \text{ Å}^{-3}$
200 parameters	Extinction correction: SHELXTI
H atoms: see below	Extinction coefficient: 0.0156 (16)

#### Table 1

Selected geometric parameters (Å, °).

O1-C1	1.232 (3)	O2-C1	1.281 (3)
O3-C4	1.226 (3)	O4-C4	1.281 (3)
O6-C5	1.230 (3)	O5-C5	1.285 (3)
O8-C8	1.230 (3)	O7-C8	1.279 (3)
01 C1 02	122.0(2)	04 C4 C3	110.8(2)
01-C1-C2	118.0 (2)	04 - C4 - C5 06 - C5 - O5	119.8 (2) 122.4 (2)
O2-C1-C2	119.9 (2)	O6-C5-C6	117.9 (2)
08-C8-07	122.6 (2)	O5-C5-C6	119.6 (2)
O8-C8-C7	117.8 (2)	C3-C2-C1	130.2 (2)
07-C8-C7	119.7 (2)	C6-C7-C8	130.8 (2)
O3-C4-O4	122.5 (2)	C2-C3-C4	130.5 (2)
O3-C4-C3	117.6 (2)	C7-C6-C5	130.9 (2)
03-C4-O4 03-C4-C3	122.5 (2) 117.6 (2)	$C_2 = C_3 = C_4$ $C_7 = C_6 = C_5$	130. 130.



#### Figure 2

A three-dimensional view of the supramolecular layer of the title salt; hydrogen bonds are illustrated by dashed lines.

#### Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
O2-H1'···O4	1.21	1.22	2.419 (2)	171
$O7-H2'\cdots O5$	1.19	1.23	2.420 (2)	171
$N1 - H1B \cdots O1$	0.98	1.90	2.842 (2)	161
$N2 - H2B \cdots O8$	1.04	1.73	2.757 (2)	169
$N1-H1A\cdots O3^{i}$	0.92	1.94	2.813 (3)	157
$N2-H2A\cdots O6^{ii}$	0.96	1.84	2.770 (3)	162
$C10-H10B\cdots O4^{i}$	0.97	2.46	3.356 (3)	153
$C12-H12A\cdots O5^{ii}$	0.97	2.50	3.357 (3)	148

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

The H atoms involved in O–H···O and N–H···O hydrogen bonds were found in difference Fourier maps and were not incorporated in the refinement. All other H atoms were positioned geometrically and included in the refinement as riding, with  $U_{\rm iso}$ values equal to  $1.2U_{\rm eq}$  of the carrier atoms.

Data collection: XSCANS (Siemens, 1996); cell refinement: XSCANS; data reduction: SHELXTL (Siemens, 1996); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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