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Takeo Fukunaga^a* and Hiroyuki Ishida^b

 ^aGraduate School of Natural Science and Technology, Okayama University,
Tsushimanaka, Okayama 700-8530, Japan, and
^bDepartment of Chemistry, Faculty of Science,
Okayama University, Tsushimanaka, Okayama
700-8530, Japan

Correspondence e-mail: t_fuku@cc.okayama-u.ac.jp

Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.002 \text{ Å}$ R factor = 0.065 wR factor = 0.120 Data-to-parameter ratio = 17.3

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

Tartronic acid forms the title salt, $C_3H_5N_2^+ \cdot C_3H_3O_5^-$, with imidazole. Hydrogen bonds in the crystal link the anions, forming a hydrogen-bonded layer. The cations cross-link the layers, leading to a three-dimensional hydrogen-bonded network.

Imidazolium hydrogen tartronate

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Comment

The title compound, (I), was investigated as part of a structural study on $D - H \cdot \cdot A$ hydrogen bonding (D = N, O or C; A = N or O) in carboxylic acid and pyridine systems (Kashino et al., 2001; Ishida et al., 2001, 2002; Fukunaga et al., 2003). Tartronic acid is of potential interest in crystal engineering for the formation of specific two- or three-dimensional aggregates with organic bases, because the hydroxyl and two carboxyl groups of the acid may act as hydrogen-bond donors as well as acceptors (Taka et al., 1998; Moritani et al., 2001). Furthermore, the acid is an interesting candidate for the selective synthesis and crystallization of optically active substances, because its anion, the hydrogen tartronate ion, is optically active. We report here the packing and hydrogen-bonding patterns of the chiral molecules of hydrogen tartronate, induced by an acid-base interaction involving proton transfer in the crystal structure of (I).



In (I), the asymmetric unit is composed of $C_3H_5N_2^+ \cdot C_3H_3O_5^-$ and the crystal is racemic. The imidazolium cation and the hydrogen tartronate anion are held together by $N1-H1\cdots O5$ and $C3-H5\cdots O2$ hydrogen bonds (Fig. 1). In the anion, an intramolecular hydrogen bond, $O1-H7\cdots O2$, is



Figure 1

ORTEP-3 (Farrugia, 1997) drawing of the asymmetric unit of (I), with the atomic numbering. Displacement ellipsoids of non-H atoms are drawn at the 50% probability level. $N-H\cdots O$, $O-H\cdots O$ and $C-H\cdots O$ hydrogen bonds are indicated by dashed lines.

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Figure 2

The hydrogen-bonded layer of anions, formed parallel to the *ab* plane (symmetry codes as in Table 2).

observed. The packing of the anions of (I), viewed down the caxis, is shown in Fig. 2. The assembly of hydrogen tartronate ions, leading to layers parallel to the ab plane, is characterized by a short O3-H8···O4ⁱⁱ hydrogen bond between the carboxyl and carboxylate groups and a long $O1-H7\cdots O5^{i}$ hydrogen bond between the hydroxyl and carboxylate groups (Table 2). In the layer, adjacent tartronate anions, which are related by a 2_1 screw axis and linked by an O3-H8···O4ⁱⁱ hydrogen bond, form a homochiral helical chain running along the b direction. The adjacent anion chains related by the aglide plane are linked in the *a* direction by a weak O1- $H7 \cdots O5^{i}$ bond, resulting in the heterochiral layer. These layers are cross-linked in the c direction by the imidazolium cations, by way of N1-H1...O5, N2-H3...O4ⁱⁱⁱ, C2- $H4\cdots O2^{v}$, $C2-H4\cdots O3^{iii}$ and $C3-H5\cdots O2$ bonds, leading to a three-dimensional hydrogen-bonded network (Table 2 and Fig. 3).

Experimental

The title crystal was grown by slow evaporation of an ethanol solution of imidazole and tartronic acid in a 1:1 molar ratio.

Crystal data

$C_{3}H_{5}N_{2}^{+}\cdot C_{3}H_{3}O_{5}^{-}$	$D_x = 1.525 \text{ Mg m}^{-3}$
$M_r = 188.14$	Mo K α radiation
Monoclinic, $P2_1/a$	Cell parameters from 25
a = 8.5516 (15) Å	reflections
b = 10.2077 (17) Å	$\theta = 11.1 - 11.5^{\circ}$
c = 9.392 (3) Å	$\mu = 0.13 \text{ mm}^{-1}$
$\beta = 91.64 \ (2)^{\circ}$	T = 298 K
$V = 819.6 (3) \text{ Å}^3$	Prism, colourless
Z = 4	$0.50 \times 0.40 \times 0.35 \text{ mm}$
Data collection	
Rigaku AFC-5R diffractometer	$R_{\rm int} = 0.021$
ω –2 θ scans	$\theta_{\rm max} = 31.0^{\circ}$
Absorption correction: ψ scan	$h = -2 \rightarrow 12$
(North et al., 1968)	$k = -14 \rightarrow 14$
$T_{\min} = 0.88, T_{\max} = 0.95$	$l = -13 \rightarrow 13$
6785 measured reflections	3 standard reflections
2617 independent reflections	every 97 reflections

intensity decay: 4.2%



Figure 3

Packing diagram of (I), viewed along the *a* direction, showing $N-H\cdots O$, $O-H\cdots O$ and $C-H\cdots O$ hydrogen bonds (symmetry codes as in Table 2).

Refinement

Refinement on F^2	$(\Delta/\sigma)_{\rm max} = 0.001$
$R[F^2 > 2\sigma(F^2)] = 0.065$	$\Delta \rho_{\rm max} = 0.36 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.120$	$\Delta \rho_{\rm min} = -0.27 \ {\rm e} \ {\rm \AA}^{-3}$
S = 1.01	Extinction correction: Zachariasen
2617 reflections	(1967)
151 parameters	Extinction coefficient:
All H-atom parameters refined	$1.43(12) \times 10^{-5}$
$w = 1/[\sigma^2(F_o) + 0.00181 F_o ^2]$	

Table 1

Selected geometric parameters (Å, °).

O1-C4	1.4089 (12)	N1-C3	1.3659 (18)
O2-C5	1.2138 (12)	N2-C1	1.3169 (15)
O3-C5	1.3031 (12)	N2-C2	1.3651 (18)
O4-C6	1.2622 (12)	C2-C3	1.3420 (18)
O5-C6	1.2383 (12)	C4-C5	1.5279 (13)
N1-C1	1.3200 (15)	C4-C6	1.5384 (13)
C1 - N1 - C3	108 38 (10)	01 - C4 - C5	110 15 (8)
C1-N2-C2	108.80 (10)	02 - C5 - 03	126.13 (9)
N1-C1-N2	108.71 (11)	O3-C5-C4	112.94 (8)
N2-C2-C3	106.82 (12)	O4-C6-O5	124.77 (9)
N1-C3-C2	107.28 (12)		

lable 2			
Hydrogen-bonding geometry	(Å,	°).	

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O1-H7···O2	0.87 (2)	2.18 (2)	2.6552 (12)	113.8 (16)
$O1-H7\cdots O5^{i}$	0.87 (2)	2.48 (2)	3.1150 (16)	130.2 (17)
O3−H8···O4 ⁱⁱ	0.882 (17)	1.665 (17)	2.5361 (14)	169.0 (17)
$N1 - H1 \cdots O5$	0.901 (15)	1.825 (16)	2.7116 (15)	167.3 (17)
$N2-H3\cdots O4^{iii}$	0.920 (18)	1.875 (19)	2.7662 (14)	162.5 (19)
$C1-H2\cdots O1^{iv}$	0.928 (18)	2.516 (18)	3.3667 (18)	152.5 (13)
$C2-H4\cdots O2^{v}$	0.98 (2)	2.49 (2)	3.280 (2)	137.6 (19)
C2-H4···O3 ⁱⁱⁱ	0.98 (2)	2.524 (18)	3.1347 (18)	120.2 (14)
C3−H5···O2	0.953 (17)	2.462 (16)	3.3841 (19)	162.7 (15)

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

2032 reflections with $I > 2\sigma(I)$

All H atoms were located in difference Fourier maps and refined isotropically. Refined distances: C-H = 0.927 (18)–0.98 (2), N-H = 0.901 (15)–0.920 (18) and O-H = 0.87 (2)–0.883 (18) Å.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1990); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN for Windows* (Molecular Structure Corporation, 1997–1999); program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994); program(s) used to refine structure: *TEXSAN for Windows*; molecular graphics: *ORTEP*-3 (Farrugia, 1997); software used to prepare material for publication: *TEXSAN for Windows*.

X-ray measurements were made at the X-ray Laboratory of Okayama University.

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