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# **Piperazinium L-Tartrate**

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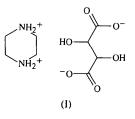
## Abstract

The structure of piperazinium L-tartrate,  $C_4H_{12}N_2^{2^+}$ .  $C_4H_4O_6^{2^-}$ , exhibits a complex three-dimensional network of hydrogen bonds. The divalent anions create puckered layers *via* two O—H···O hydrogen bonds and neighbouring layers are then crosslinked by four N—H···O interactions through the divalent cations.

## Comment

Recent years have seen considerable interest in the use of hydrogentartrate ions as a means of creating hydrogen-bonded two-dimensional aggregates (Aakeröv & Hitchcock, 1993; Zyss, Pecaut, Levy & Masse, 1993; Watanabe, Noritake, Hirose, Okada & Kurauchi, 1993; Dastidar, Row, Prasad, Subramanian & Bhattacharya, 1993). Such motifs have been employed as structural 'scaffolding' in the crystal engineering of many new materials for nonlinear optics. The cornerstone of the twodimensional sheet is a head-to-tail (from the carboxylic end to the carboxylate end) O-H···O interaction between adjacent anions. However, if we eliminate the chemical basis of this link (by deprotonating both ends of the acid), the strongest and most important hydrogen-bonding interaction between anions has been removed [observed  $O \cdot \cdot O$  distances for this interaction are commonly 2.50–2.55 Å (Aakeröy & Hitchcock, 1993)]. Consequently, this should have a considerable effect on the structure of the anionic aggregates in organic tartrate(2-) salts. The anions may still create a layered structure through other O-H···O interactions or, alternatively, the infinite two-dimensional aggregate may be broken up altogether in favour of another motif. In order to examine this scenario, we are currently performing a structural study of pairs of hydrogen-Ltartrate and L-tartrate salts with the same counterion. The crystallographic literature contains only two examples of resolved tartrates(2-) with organic counterions, ethylendiammonium L-tartrate (Fair & Schlemper, 1977; Palmer & Ladd, 1977) and bisguanidinium L-tartrate monohydrate (Krumbe, Haussuhl & Frohlich, 1989). The crystal structure of piperazinium(2+) bis(hydrogen-L-tartrate) has been determined previously (Aakeröy,

Hitchcock & Seddon, 1992), and the synthesis and X-ray single crystal structure determination of the corresponding L-tartrate(2-) salt (I) reported herein provides the basis for a valuable structural comparison.



The divalent cation of the title compound does not show any unusual geometric features (Fig. 1 and Table 1) and displays a chair conformation with expected endocyclic bond lengths and bond angles. The divalent anion exists in the expected trans configuration; the torsion angle of the C---C---C backbone is  $178.42(11)^{\circ}$ . There is some evidence to suggest that glycolic acid moieties normally exhibit a planar conformation in  $\alpha$ -hydroxy carboxylic acids (Stouten, Kroon-Batenburg & Kroon, 1989), an arrangement which is often the result of an intramolecular hydrogen-bond interaction. However, in the anion of the title compound, each carboxylate moiety shows considerable deviation from coplanarity with respect to the adjacent hydroxy functional group; O(12')—C(1')—C(2')—  $O(21') - 8.23(2)^{\circ}$  and O(31') - C(3') - C(4') - O(42') $-14.55(2)^{\circ}$ . This variation must be attributed to the specific intermolecular hydrogen-bond interactions between the anion and the strong hydrogen-bond donors of the cation.

Despite the fact that the possibility for a head-totail hydrogen bond between adjacent anions has been removed due to the double deprotonation of the acid,

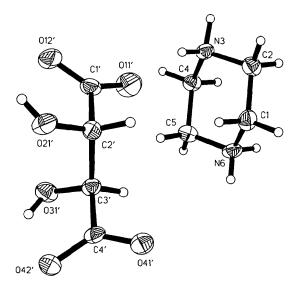


Fig. 1. Geometry, displacement ellipsoids (50% probability) and numbering scheme for the ion pair of the title compound.

the anions still create an infinite anionic layer, parallel to *ab* (Fig. 2). This motif is created through two  $O \cdots O$ interactions between the hydroxy groups (the donors) and the carboxylate moieties (the acceptors) (Table 2); the resulting layers display a puckered geometry.

The divalent cations, each with four strong hydrogenbond donors involved in hydrogen bonds to four different anions, provide crosslinks between neighbouring puckered anionic layers, through four  $N \cdots O$ hydrogen-bond interactions (Fig. 3). This complex threedimensional hydrogen-bonded network results in a densely packed architecture,  $D_x = 1.509 \,\mathrm{g \, cm^{-3}}$ .

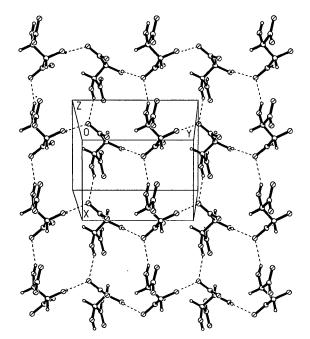


Fig. 2. A view of the infinite anionic layer, parallel with the ab plane. Hydrogen bonds are indicated by dashed lines.

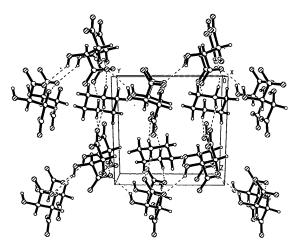


Fig. 3. The piperazinium(2+) crosslink between neighbouring (buckled) anionic layers. The layers are viewed edge on, only three cations are shown for clarity and hydrogen bonds are indicated by dashed lines.

The corresponding piperazinium(2+) bis(hydrogen-L-tartrate) salt (Aakeröy, Hitchcock & Seddon, 1992) also contains two-dimensional layers generated by two  $\mathbf{O} \cdots \mathbf{O}$  hydrogen bonds. In this case, neighbouring anions are linked in the expected head-to-tail fashion. The piperazinium cations form layers, halfway between the anion layers. Each cation is hydrogen bonded to two O atoms in the layer above and by symmetryrelated hydrogen bonds to two O atoms in the anion layer below. This compound, with its compact packing of ions throughout the structure, also has a high density,  $D_x = 1.63 \,\mathrm{g}\,\mathrm{cm}^{-3}$ .

A comparison with the other two resolved organic tartrates(2-), ethylendiammonium L-tartrate and bisguanidinium L-tartrate monohydrate, shows that in both cases the anions create infinite chains not layers. Consequently, based upon the three extant structures, the tartrate(2-) anions do not seem to be as structurally consistent as the corresponding hydrogentartrate anions (Aakeröy & Hitchcock, 1993). The reason for this is probably that whereas the hydrogentartrate anion contains a powerful hydrogen-bond donor (-COOH) which interacts strongly and preferentially with the (-COO<sup>-</sup>) acceptor, the only donor on the tartrate anion, (-OH), is not powerful enough to hold the layers together in the presence of cations with differing hydrogen-bond donor abilities.

## **Experimental**

Piperazinium(2+) L-tartrate(2-) was prepared by mixing equimolar aqueous solutions of piperazine and L-tartaric acid. Transparent colourless crystals were obtained by slow evaporation from water.

Crystal data

 $C_4H_{12}N_2^{2+}.C_4H_4O_6^{2-}$ Mo  $K\alpha$  radiation  $M_r = 236.23$  $\lambda = 0.71073$  Å Monoclinic  $P2_1$ reflections a = 6.4262 (10) Å $\theta = 5 - 12.5^{\circ}$ b = 9.1322(10) Å  $\mu = 0.13 \text{ mm}^{-1}$ c = 9.3627 (10) ÅT = 293 (2) K $\beta = 108.850 (10)^{\circ}$ Needle  $V = 519.98 (11) \text{ Å}^3$ Z = 2Colourless  $D_x = 1.509 \text{ Mg m}^{-3}$  $D_m$  not measured Data collection Siemens P4 diffractometer  $\theta_{\rm max} = 25^{\circ}$  $\omega$  scans  $h = 0 \rightarrow 7$  $k = -10 \rightarrow 10$ Absorption correction:

- none
- 1993 measured reflections
- 1823 independent reflections

1796 observed reflections  $[I > 2\sigma(I)]$ 

 $R_{\rm int} = 0.0124$ 

- Cell parameters from 38  $0.96 \times 0.36 \times 0.16$  mm
- $l = -11 \rightarrow 10$ 3 standard reflections monitored every 97 reflections intensity decay:  $\pm 1\%$

Refinement	
Refinement on $F^2$	$\Delta \rho_{\text{max}} = 0.157 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\text{min}} = -0.162 \text{ e } \text{\AA}^{-3}$
R(F) = 0.0248	$\Delta \rho_{\rm min} = -0.162 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.0658$	Extinction correction:
S = 1.085	$F_c^* = F_c k [1 + (0.001 \chi$
1823 reflections	$\times F_c^2 \lambda^3 / \sin 2\theta$ ]
146 parameters	Extinction coefficient:
$w = 1/[\sigma^2(F_o^2) + (g_1P)^2]$	$\chi = 0.177 (9)$
$+ g_2 P$ ]	Atomic scattering factors
where $P = [\max F_{q}^{2}, 0]$	from International Tables
$+ 2F_c^2$ ]/3	for X-ray Crystallography
$(\Delta/\sigma)_{\rm max} < 0.001$	(1974, Vol. IV)

## Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

# $U_{\text{eq}} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j.$

	x	у	z	$U_{eq}$
C(1)	-0.5779 (3)	0.4773 (2)	-0.6549 (2)	0.029(1)
C(2)	-0.5047 (3)	0.5182 (2)	-0.7866 (2)	0.029(1)
N(3)	-0.3463 (2)	0.4081(1)	-0.8054(1)	0.024(1)
C(4)	-0.4425 (3)	0.2579 (2)	-0.8263 (2)	0.026(1)
C(5)	-0.5191 (3)	0.2163 (2)	-0.6958 (2)	0.029(1)
N(6)	-0.6749 (2)	0.3277 (2)	-0.6748 (1)	0.028(1)
C(1')	0.1048 (2)	0.3610(2)	-0.4015 (2)	0.027(1)
O(11')	-0.0590 (2)	0.3588 (2)	0.5162 (1)	0.050(1)
O(12')	0.2969 (2)	0.3290 (2)	-0.3946(1)	0.038(1)
C(2')	0.0716 (2)	0.4078 (2)	-0.2529 (2)	0.024(1)
O(21')	0.2753 (2)	0.4254 (1)	-0.1377 (1)	0.031(1)
C(3')	-0.0698 (2)	0.2957 (2)	-0.2043 (2)	0.022(1)
O(31')	0.0288 (2)	0.1556 (1)	-0.1869 (1)	0.030(1)
C(4')	-0.1082 (2)	0.3450(2)	-0.0579 (1)	0.022(1)
O(41')	-0.2368 (2)	0.4501 (1)	-0.0665(1)	0.031(1)
O(42')	-0.0096 (2)	0.2744 (1)	0.0589(1)	0.031(1)

## Table 2. Hydrogen-bonding geometry (Å, °)

$D - H \cdots A$ N(3) - H \cdots O(41'') N(3) - H \cdots O(11') N(6) - H \cdots O(12''n) N(6) - H \cdots O(22''n) O(21') - H \cdot O(22''n) O(21') - H \cdot O(22''n)	H···A 1.760 (2) 1.936 (2) 1.907 (2) 1.859 (2) 2.105 (2)	D···A 2.780 (2) 2.777 (2) 2.688 (2) 2.759 (2) 2.632 (2)	$D - H \cdot \cdot \cdot A$ 170 (4) 153 (3) 156 (3) 160 (3) 125 (2)
$ \begin{array}{l} N(6) - H \cdots O(42'^{10}) \\ O(31') - H \cdots O(42'^{1v}) \\ O(21') - H \cdots O(12'^{v}) \end{array} $	1.859 (2)	2.759 (2)	160 (3)
	2.105 (2)	2.627 (2)	125 (3)
	2.207 (1)	2.607 (1)	110 (3)

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

Data collection: XSCANS (Fait, 1991). Cell refinement: XS-CANS. Data reduction: XSCANS. Program(s) used to solve structure: SHELXTL/PC (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: XP SHELXTL/PC.

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: PT1032). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# One Bicyclic $\beta$ -Lactam and Two Bicyclic $\gamma$ -Lactam Compounds

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### Abstract

The crystal structures of the bicyclic  $\beta$ -lactam, rac-(2R\*,4S\*,6S\*)-4-chloro-4-methyl-8-oxo-1-azabicyclo[4.2.0]octane-2-carboxylic acid methyl ester, C<sub>10</sub>H<sub>14</sub>ClNO<sub>3</sub> (1), and the bicyclic  $\gamma$ -lactams, rac-(5R\*,7S\*,9S\*)-7-chloro-7-methyl-3-oxooctahydroindolizine-5-carboxylic acid methyl ester, C<sub>11</sub>H<sub>16</sub>ClNO<sub>3</sub> (2), and rac-(5R\*,6S\*,10R\*)-3-oxo-6-vinyloctahydro-1Hpyrrolo[1,2-a]azepine-5-carboxylic acid methyl ester, C<sub>13</sub>H<sub>19</sub>NO<sub>3</sub> (3), have been established by X-ray crystallography. In (1) and (2) the  $\alpha$ -amino ester function and the Cl atom occupy an axial position in a chair-like piperidine ring. In (3) the methyl ester and the vinyl group are equatorially placed on the chair-like seven-

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