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Key indicators

Single-crystal X-ray study T = 297 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ Disorder in main residue R factor = 0.058 wR factor = 0.183 Data-to-parameter ratio = 13.8

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

Ethylenediaminium pyridine-2,5-dicarboxylate dihydrate

The title compound, $C_2H_{10}N_2^{2+}C_7H_3NO_4^{2-}H_2O$, forms a three-dimensional hydrogen-bonded framework structure in which both of the amine groups of ethylenediamine are protonated and participate in a total of six hydrogen-bonding interactions with carboxylate O-atom acceptors of the anions as well as the water molecules. The cations lie on crystal-lographic inversion centres and adopt extended conformations, while the anions have twofold rotational symmetry. This results in the pyridine N atom and its *ortho*-related CH group being disordered over two 50% occupancy sites.

Comment

Ethylenediamine (1,2-ethanediamine, EN) has relatively close dissociation constants ($pK_{a1} = 7.3$ and $pK_{a2} = 10.1$), and in reaction with organic acids most commonly forms dications. Therefore with monoprotic acids, 1:2 salts are usual, e.g. with the aromatic acids 4-chlorobenzoic acid (Xia et al., 2003), 4-nitrobenzoic acid (Lynch & Healy, 2003), 4-nitroanthranilic acid (Smith et al., 2002), 5-nitrosalicylic acid (Smith et al., 2005), 3,5-dinitrobenzoic acid (Nethaji et al., 1992; Lynch et al., 1994; Jones et al., 2005) and 5-sulfosalicylic acid (Gao et al., 2004). Even with diprotic acids, these 1:2 salts are commonly found, forming hydrogenearboxylates, e.g. with oxalic acid (Barnes et al., 1998), malonic acid (Barnes & Weakley, 2000) and succinic acid (Schreuer & Muench, 1994). The six aminium protons on the EN²⁺ cations in these compounds may be involved in six or more intermolecular hydrogen-bonding interactions with suitable acceptor atoms, thus giving, in most cases, hydrogen-bonded framework structures which often incorporate water molecules of solvation (the 5-nitrosalicylate, the hydrogen malonate and the hydrogen oxalate are all monohydrates, the 4-nitrobenzoate and 4-nitroanthanilate are dihydrates, while the 5-sulfosalicylate is a tetrahydrate). However, the 1:1 stoichiometric reaction of ethylenediamine with diprotic pyridine-2,5-dicarboxylic acid (PYCA) gave a 1:1 dihydrate salt EN²⁺·PYCA²⁻·2H₂O, (I), the structure of which is reported here.



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Figure 1

Molecular configuration and atom labelling scheme for the EN²⁺ cation, $PYCA^{2-}$ anion and the water molecule in (I). The symmetry code for the inversion-related portion of the cation (v) is $\left(-x+\frac{1}{2},-y+\frac{1}{2},-z+1\right)$, while the anion has twofold rotational symmetry [symmetry code: (vi) $-x + 2, y, -z + \frac{3}{2}$]. The inversion-related water molecule O1W^{iv} (see Table 1 for symmetry code) is also shown. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

Perspective view of the packing in the unit cell, showing hydrogenbonding associations as dashed lines. Symmetry code (vii) for O212^{vii} is $\left(-x+\frac{3}{2},y+\frac{1}{2},-z+\frac{1}{2}\right)$. Symmetry code (i) as in Fig. 1; the other symmetry codes are listed in Table 1.

Compound (I) comprises an EN²⁺ cation which lies across a crystallographic inversion centre [symmetry code: (v) $-x + \frac{1}{2}$, $-y + \frac{1}{2}$, -z + 1], a PYCA²⁻ anion which lies across a twofold rotation axis [symmetry code: (vi) -x + 2, y, $-z + \frac{3}{2}$] and two water molecules (Fig. 1). The EN^{2+} cations adopt an extended conformation, with torsion angle $N11-C11-C11^{v}-N11^{v} =$ 180 Å. The occurrence of centrosymmetric EN²⁺ cations in the structures of EN salts is quite common, e.g. in ethylenediaminium terephthalate (Moritani & Kashino, 1991), which is structurally analogous to (I). The carboxylate groups in the PYCA²⁻ anions are related by twofold rotation symmetry, which means that the pyridine N (N1) and the ortho-related C1^{vi}/H1^{vi} group are disordered over two 50% occupancy sites. The carboxylate groups of the anion are rotated slightly out of the plane of the benzene ring [C3-C2-C21-O212 =

 $-159.0 (4)^{\circ}$]. All six aminium protons are involved in hydrogen-bonding interactions (Table 1), two to water molecules of solvation and the remainder to separate carboxylate-O-atom acceptors. The water molecules also act as donors to carboxylate O atoms, leading to a three-dimensional layered framework structure (Fig. 2).

Experimental

The title compound was synthesized by heating together 1 mmol quantities of PYCA and EN in 50 ml of 50% ethanol-water under reflux for 10 min. After concentration to ca 30 ml, total roomtemperature evaporation of the hot-filtered solution gave colourless crystal plates (m.p. 527 K) of (I).

 $R_{\rm int} = 0.016$

 $\theta_{\rm max} = 27.5^\circ$

3 standard reflections

frequency: 150 min

 $w = 1/[\sigma^2(F_0^2) + (0.1P)^2]$

where $P = (F_0^2 + 2F_c^2)/3$

-3

+ 9.335P]

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta\rho_{\rm max} = 0.34 \text{ e Å}$

 $\Delta \rho_{\rm min} = -0.28 \text{ e } \text{\AA}^{-3}$

intensity decay: 2.5%

Crystal data

$C_2H_{10}N_2^{2+} \cdot C_7H_3NO_4^{2-} \cdot 2H_2O$	Z = 4
$M_r = 263.26$	$D_x = 1.428 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
a = 11.4733 (16) Å	$\mu = 0.12 \text{ mm}^{-1}$
b = 12.1572 (13) Å	T = 297 (2) K
c = 10.0129 (13) Å	Plate, colourless
$\beta = 118.750 \ (9)^{\circ}$	$0.30 \times 0.30 \times 0.12 \text{ mm}$
V = 1224.5 (3) Å ³	

Data collection

Rigaku AFC-7R diffractometer ω -2 θ scans Absorption correction: none 1699 measured reflections 1408 independent reflections 1017 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.059$ $wR(F^2) = 0.183$ S = 0.891408 reflections 102 parameters H atoms treated by a mixture of

independent and constrained refinement

Table T			
Hydrogen-bond	geometry	(Å,	°).

Hydrogen-bond geom	etry (\mathbf{A}, \mathbf{r}) .			
$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1W - H1W \cdots O211^{i}$	0.91 (5)	1.86 (5)	2.769 (3)	179 (7)
$O1W - H2W \cdot \cdot \cdot O212^{ii}$	0.91 (5)	1.83 (5)	2.740 (4)	176 (5)
N11−H111···O211 ⁱⁱⁱ	0.98 (4)	1.85 (4)	2.811 (3)	168 (4)
$N11 - H112 \cdots O212^{iv}$	0.85 (5)	1.96 (5)	2.786 (4)	164 (4)
$M11 = H113 \dots O1W$	0.92(6)	1.90 (6)	2.796(4)	166(3)

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

H atoms involved in hydrogen-bonding interactions were located by difference methods and their positional and isotropic displacement parameters were refined. Other H atoms were included in the refinement in calculated positions (C-H = 0.93 and 0.95 Å) using a riding-model approximation, with $U_{iso}(H) = 1.2U_{eq}(C)$ with the exception of C1, which was set invariant with U_{iso} arbitrarily as *ca*. U_{eq} (C1). The pyridine N atom and its *ortho*-related CH group, which are related by crystallographic twofold symmetry, were refined with 50% N,C-H site occupancy.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1999); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN* for Windows (Molecular Structure Corporation, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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