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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.055 wR factor = 0.143 Data-to-parameter ratio = 11.0

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

Morpholinium dihydrogen citrate hydrate

In the crystal structure of the title compound, $[O(CH_2CH_2)_2NH_2][C_6H_7O_7] \cdot H_2O$, the cation interacts with the negatively charged carboxylato group of adjacent anions $[N \cdots O \ 2.847 \ (3)$ and $2.942 \ (3)$ Å], forming a linear chain running along the *b* axis of the crystal. Adjacent chains are linked through the carboxylic $-CO_2H$ groups and the water molecule into a layer structure.

Comment

A previous report described the dicyclohexylammonium salts of monocarboxylic and dicarboxylic acids (Ng, 2003). For some of the acids, their reaction with dicyclohexylamine afforded the corresponding ammonium carboxylate as a hydrate (Ng, 1992; Yang et al., 2002). On the other hand, others reacted with an excess of this secondary amine to yield only the mono(ammonium) salt, e.g., dicyclohexylammonium hydrogen oxalate (Ng, 1994). The reaction with the tribasic acid, citric acid, gave a compound that rapidly turned opaque when the crystals were removed from its solution in ethanol; on the other hand, the use of diisopropylamine in place of dicyclohexylamine led to a syrupy material after the solvent had evaporated. A cursory check through the Cambridge Structural Database (Allen, 2002) on the sec-ammonium derivatives of carboxylic acids suggested the use of morpholine, as this amine furnishes a large number of isolable carboxylates, such as chloronitrobenzoates (Ishida et al., 2001a,b,c) and morpholinoformate (Brown & Gray, 1982). Citric acid has also been characterized as an N-substituted piperidinylium salt (Peeters et al., 1979). In the present study, the reaction of citric acid with either one or two molar equivalents of morpholine gave the air-stable title compound, (I) (Fig. 1).



In the crystal structure of (I), the cation interacts with the carboxyl O atoms of two citrato anions $[N \cdots O 2.847 (3) \text{ and } 2.942 (3) \text{ Å}]$, resulting in the formation of a linear chain running along the *b* axis of the crystal. Such a hydrogenbonded motif has previously been noted in dicyclohexyl-ammonium trifluoroacetate (Ng, 1999). Adjacent chains are

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2103 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_o^2) + (0.0711P)^2]$

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

+ 1.4573P]

 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.29 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.19 \text{ e } \text{\AA}^{-3}$

$$\begin{split} R_{\rm int} &= 0.028\\ \theta_{\rm max} &= 25.0^\circ\\ h &= -13 \rightarrow 13\\ k &= -14 \rightarrow 10\\ l &= -20 \rightarrow 21 \end{split}$$



Figure 1

ORTEPII (Johnson, 1976) plot of (I), with displacement ellipsoids drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.



Figure 2

ORTEPII (Johnson, 1976) plot, illustrating the hydrogen-bonded layer structure. H atoms are omitted.

linked through the carboxylic $-CO_2H$ groups and water molecule into a layer structure (Fig. 2 and Table 2).

Experimental

Morpholine (0.87 g, 10 mmol) and citric acid monohydrate (0.21 g, 10 mol) were heated in a small volume of ethanol (25 ml); a few drops of water were added to dissolve the acid completely. Slow evaporation of the solvent yielded the title compound. The use of twice the quantity of the amine gave the identical compound. Elemental analysis, found (calc. for $C_{10}H_{19}O_9N$): 40.8 (40.4); H 6.5 (6.4); N 4.5% (4.7%). IR (KBr pellet): ν_{COOH} 1717, $\nu_{as(COO)}$ 1414, $\nu_{s(COO)}$ 1414, 1396; $\nu_{C=N}$ 1107 cm⁻¹.

Crystal data

$C_4H_{10}NO^+ \cdot C_6H_7O_7^- \cdot H_2O$	Mo Ka radiation
$M_r = 297.26$	Cell parameters from 4726
Orthorhombic, Pbca	reflections
a = 11.3774 (6) Å	$\theta = 2.4-27.3^{\circ}$
b = 12.4823 (7) Å	$\mu = 0.13 \text{ mm}^{-1}$
c = 18.437(1) Å	T = 298 (2) K
$V = 2618.3 (2) \text{ Å}^3$	Block, colorless
Z = 8	$0.28 \times 0.18 \times 0.18 \text{ mm}$
$D_x = 1.508 \text{ Mg m}^{-3}$	

Data collection

Bruker AXS area-detector
diffractometer
ω and ω scans
Absorption correction: none
2407 measured reflections
2305 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.055$ $wR(F^2) = 0.143$ S = 1.152305 reflections 209 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected geometric parameters (Å, °).

O1-C1	1.243 (3)	N1-C7	1.482 (3)
O2-C1	1.253 (3)	N1-C10	1.484 (3)
O3-C5	1.304 (3)	C1-C2	1.524 (3)
O4-C5	1.207 (3)	C2-C3	1.536 (3)
O5-C6	1.316 (3)	C3-C4	1.525 (3)
O6-C6	1.203 (3)	C3-C6	1.536 (3)
O7-C3	1.421 (3)	C4-C5	1.505 (3)
O8-C8	1.418 (3)	C7-C8	1.489 (4)
O8-C9	1.421 (3)	C9-C10	1.496 (4)
C8-O8-C9	109.9 (2)	C3-C4-C5	113.6 (2)
O1-C1-O2	124.2 (2)	O3-C5-O4	123.1 (2)
O1-C1-C2	118.7 (2)	O3-C5-C4	112.8 (2)
O2-C1-C2	117.2 (2)	O4-C5-C4	124.1 (2)
C1-C2-C3	114.6 (2)	O5-C6-O6	124.4 (2)
O7-C3-C2	109.6 (2)	O5-C6-C3	111.1 (2)
O7-C3-C4	107.7 (2)	O6-C6-C3	124.4 (2)
C2-C3-C4	110.9 (2)	N1-C7-C8	109.8 (2)
O7-C3-C6	109.4 (2)	O8-C8-C7	111.1 (2)
C2-C3-C6	109.0 (2)	O8-C9-C10	111.4 (2)
C4-C3-C6	110.3 (2)	N1-C10-C9	109.0 (2)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O3-H3o···O2 ⁱ	0.85(1)	1.76 (1)	2.612 (2)	176 (4)
$O5-H5o\cdots O1w$	0.86(1)	1.68 (1)	2.529 (3)	172 (4)
$O7-H7o\cdots O1$	0.85(1)	1.89 (2)	2.625 (2)	144 (3)
$N1-H1n2\cdots O1$	0.85(1)	2.13 (2)	2.847 (3)	143 (3)
$N1-H1n1\cdots O2^{ii}$	0.85(1)	2.12 (2)	2.942 (3)	162 (3)
$O1w - H1w1 \cdots O6^{iii}$	0.85 (1)	2.16 (2)	2.932 (3)	152 (3)
$O1w - H1w2 \cdots O7^{ii}$	0.85 (1)	1.93 (1)	2.782 (3)	175 (4)

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

The water, hydroxyl and ammonium H atoms were located and refined. The aliphatic H atoms were positioned geometrically (C–H 0.97 Å), and were included in the refinement in the riding model approximation; the displacement factors were set to 1.5 (methyl) or 1.2 (other H atoms) times $U_{\rm eq}$ of the parent atom.

The diffraction measurements were performed up to a 2θ angle of 56°; however, as the inclusion of reflections above 50° led to an *R* factor greater than 7%, these were excluded from the refinement.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine

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structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP*II (Johnson, 1976); software used to prepare material for publication: *SHELXL*97.

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