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Key indicators

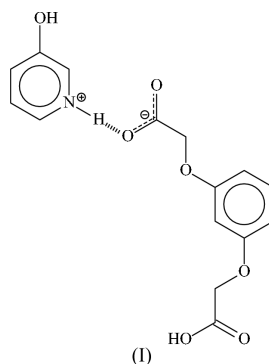
Single-crystal X-ray study
 $T = 295\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.057
 wR factor = 0.129
Data-to-parameter ratio = 15.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.3-Hydroxypyridinium 3-(carboxymethoxy)-
phenoxyacetate

The crystal structure of the title compound, $\text{C}_5\text{H}_6\text{NO}^+ \cdot \text{C}_{10}\text{H}_9\text{O}_6^-$, consists of pairs of 3-hydroxypyridinium cations and pairs of 3-(carboxymethoxy)phenoxyacetate anions that are connected *via* hydrogen bonding into linear ribbons, which are approximately planar. The asymmetric unit consists of one cation and one anion, each located in a general position.

Comment

Pyridine is not commonly used to form pyridinium carboxylates and examples of such salts are rare (Palmore & McBride-Wieser, 1997). The reaction of pyridine with 1,3-phenylenedi(oxyacetic acid) under hydrothermal conditions leads to the formation of the 3-hydroxypyridinium cation, which is formed *in situ* by oxidation of the pyridine at its 3-position.

In the crystal structure of the title salt, (I), anions are linked by $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonding into pairs, which are located around a center of inversion. These pairs are connected into ribbons by two 3-hydroxypyridinium cations, which are also located around a center of inversion, *via* $\text{N}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonding (Table 2 and Fig. 1).



A search of the Cambridge Structural Database (Version 5.25; Allen, 2002) yielded a number of 3-hydroxypyridinium carboxylates (Aakeroy & Nieuwenhuyen, 1994; Byriel *et al.*, 1992; Fukunaga *et al.*, 2003; Lynch *et al.*, 1992, 1999; Tafeenko *et al.*, 1990). In these compounds, as well as in the title compound, the cations form two hydrogen bonds.

Experimental

The title compound was prepared by the hydrothermal reaction of pyridine (3 ml) and 1,3-phenylenedi(oxyacetic acid) (4.52 g, 20 mmol) in water. The mixture was sealed in a 15 ml Teflon-lined stainless steel bomb. The bomb was heated at 403 K for 72 h and then cooled slowly to room temperature. Colorless crystals were obtained when the filtered solution was stored for a few days. Analysis calculated for $\text{C}_{15}\text{H}_{15}\text{NO}_7$: C 56.08, H 4.71, N 4.36%; found: C 56.20, H 4.65, N 4.29%.

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Crystal data

$C_5H_6NO^+ \cdot C_{10}H_9O_6^-$
 $M_r = 321.28$
 Monoclinic, $P2_1/n$
 $a = 15.960(2) \text{ \AA}$
 $b = 4.8404(4) \text{ \AA}$
 $c = 20.313(3) \text{ \AA}$
 $\beta = 110.58(4)^\circ$
 $V = 1469.1(3) \text{ \AA}^3$
 $Z = 4$

$D_x = 1.453 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 Cell parameters from 6106 reflections
 $\theta = 4.4\text{--}27.5^\circ$
 $\mu = 0.12 \text{ mm}^{-1}$
 $T = 295(2) \text{ K}$
 Prism, colorless
 $0.38 \times 0.25 \times 0.19 \text{ mm}$

Data collection

Rigaku R-Axis RAPID diffractometer
 ω scans
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.790$, $T_{\max} = 0.978$
 13 092 measured reflections

3367 independent reflections
 2121 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.063$
 $\theta_{\max} = 27.5^\circ$
 $h = -20 \rightarrow 20$
 $k = -6 \rightarrow 5$
 $l = -26 \rightarrow 26$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.129$
 $S = 1.02$
 3367 reflections
 220 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.06P)^2 + 0.1723P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.20 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

| | | | |
|------------|-----------|-------------|-----------|
| O1—C2 | 1.334 (3) | C2—C3 | 1.385 (3) |
| O2—C6 | 1.257 (3) | C3—C4 | 1.372 (3) |
| O3—C6 | 1.250 (2) | C4—C5 | 1.369 (4) |
| O4—C8 | 1.374 (2) | C6—C7 | 1.514 (3) |
| O4—C7 | 1.416 (2) | C8—C9 | 1.391 (3) |
| O5—C10 | 1.371 (2) | C8—C13 | 1.382 (3) |
| O5—C14 | 1.410 (2) | C9—C10 | 1.384 (3) |
| O6—C15 | 1.196 (2) | C10—C11 | 1.385 (3) |
| O7—C15 | 1.318 (3) | C11—C12 | 1.370 (3) |
| N1—C1 | 1.327 (3) | C12—C13 | 1.391 (3) |
| N1—C5 | 1.331 (3) | C14—C15 | 1.506 (3) |
| C1—C2 | 1.383 (3) | | |
| C7—O4—C8 | 118.7 (2) | O4—C8—C13 | 124.4 (2) |
| C10—O5—C14 | 116.7 (2) | O4—C8—C9 | 114.0 (2) |
| C1—N1—C5 | 123.4 (2) | C9—C8—C13 | 121.6 (2) |
| N1—C1—C2 | 119.7 (2) | C8—C9—C10 | 118.8 (2) |
| O1—C2—C1 | 122.7 (2) | O5—C10—C9 | 124.4 (2) |
| O1—C2—C3 | 119.1 (2) | O5—C10—C11 | 115.2 (2) |
| C1—C2—C3 | 118.2 (2) | C9—C10—C11 | 120.4 (2) |
| C2—C3—C4 | 120.0 (2) | C10—C11—C12 | 119.7 (2) |
| C3—C4—C5 | 120.0 (2) | C11—C12—C13 | 121.4 (2) |
| N1—C5—C4 | 118.8 (2) | C8—C13—C12 | 118.0 (2) |
| O3—C6—O2 | 124.0 (2) | O5—C14—C15 | 109.6 (2) |
| O3—C6—C7 | 116.8 (2) | O6—C15—O7 | 125.8 (2) |
| O2—C6—C7 | 119.2 (2) | O6—C15—C14 | 125.8 (2) |
| O4—C7—C6 | 108.5 (2) | O7—C15—C14 | 108.4 (2) |

Table 2

Hydrogen-bonding geometry (\AA , $^\circ$).

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|----------------------------------|----------|--------------|--------------|----------------|
| N1—H1N \cdots O2 | 0.87 (1) | 1.77 (1) | 2.638 (3) | 173 (3) |
| O1—H1O \cdots O3 ⁱ | 0.86 (1) | 1.72 (1) | 2.569 (2) | 172 (3) |
| O7—H7O \cdots O2 ⁱⁱ | 0.85 (1) | 1.82 (1) | 2.663 (2) | 170 (3) |

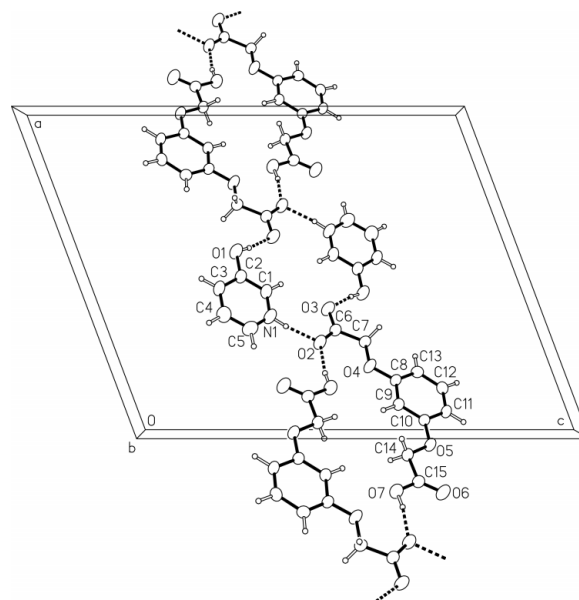
Symmetry codes: (i) $1-x, -y, 1-z$; (ii) $-x, 2-y, 1-z$.

Figure 1

ORTEP (Johnson, 1976) plot with the atom labeling, showing the ribbon structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii and hydrogen bonding is indicated by dotted lines.

Carbon-bound H atoms were placed in calculated positions [aromatic C—H = 0.93 \AA and aliphatic C—H = 0.97 \AA , and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$] and were refined isotropically using a riding model. The N—H, hydroxyl and carboxyl H atoms were located in a difference map and refined with individual isotropic displacement parameters and with a distance restraint of 0.85 (1) \AA for N—H and O—H.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP* (Johnson, 1976).

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