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Isopropylammonium (isopropylamino)-oxoacetate monohydrate

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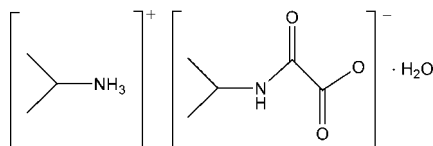
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.045; wR factor = 0.106; data-to-parameter ratio = 16.1.

The title compound, $\text{C}_3\text{H}_{10}\text{N}^+\cdot\text{C}_5\text{H}_8\text{NO}_3^-\cdot\text{H}_2\text{O}$, crystallizes as a salt from water. Intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds lead to two-dimensional layers. The layers are stacked in the c -axis direction with hydrophobic interactions between the methyl groups.

Related literature

For related literature, see: Chena *et al.* (2005); Bellouard *et al.* (1999); Dumonateil (1999).



Experimental

Crystal data

 $\text{C}_3\text{H}_{10}\text{N}^+\cdot\text{C}_5\text{H}_8\text{NO}_3^-\cdot\text{H}_2\text{O}$ $M_r = 208.26$ Triclinic, $P\bar{1}$ $a = 7.172$ (4) Å $b = 9.154$ (5) Å $c = 9.472$ (5) Å $\alpha = 104.372$ (8)° $\beta = 105.944$ (8)° $\gamma = 94.731$ (8)° $V = 571.6$ (5) Å³ $Z = 2$ Mo $K\alpha$ radiation $\mu = 0.10$ mm⁻¹ $T = 100$ (2) K $0.38 \times 0.19 \times 0.08$ mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer

Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{\min} = 0.962$, $T_{\max} = 0.990$

6062 measured reflections

2324 independent reflections

2126 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.023$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.106$ $S = 1.16$

2324 reflections

144 parameters

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\text{max}} = 0.40$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|--|----------|-------------|-------------|---------------|
| $\text{O4}-\text{H9}\cdots\text{O2}^{\text{i}}$ | 0.82 (3) | 2.06 (3) | 2.864 (2) | 170 (3) |
| $\text{O4}-\text{H10}\cdots\text{O3}$ | 0.88 (2) | 1.96 (2) | 2.8404 (19) | 173.5 (19) |
| $\text{N1}-\text{H1}\cdots\text{O4}^{\text{ii}}$ | 0.84 (2) | 2.14 (2) | 2.918 (2) | 153.7 (17) |
| $\text{N2}-\text{H2A}\cdots\text{O2}$ | 0.91 | 1.89 | 2.7957 (19) | 175 |
| $\text{N2}-\text{H2B}\cdots\text{O3}^{\text{ii}}$ | 0.91 | 2.02 | 2.8808 (19) | 157 |
| $\text{N2}-\text{H2C}\cdots\text{O1}^{\text{iii}}$ | 0.91 | 1.91 | 2.8182 (18) | 172 |

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

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001; Atwood & Barbour, 2003); software used to prepare material for publication: *X-SEED*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HY2088).

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supplementary materials

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Isopropylammonium (isopropylamino)oxoacetate monohydrate

X. Sheng, C. E. Strasser, H. G. Raubenheimer and R. C. Luckay

Comment

Chagas disease, which is caused by *Trypanosoma cruzi*, is an endemic parasitic disease in Latin America, specially in the Southern part of Mexico (Dumonateil, 1999). The *Trypanosoma cruzi* enzyme designated α -hydroxyacid dehydrogenase (HADH) exhibits two molecular forms (I and II). The trypanocidal activity of *N*-isopropylloxamate (NIPOx) on cultured epimastigotes (*in vitro*) and marine trypanosomiasis (*in vitro*) is used in different *Trypanosoma cruzi* strains. It is an effective and selective inhibitor of HADH-isozyme II (Chena *et al.*, 2005).

It was serendipitously synthesized during a chromatographic partition of cyclenoxamide. The cyclen and diethyl oxalate were added in an equal molar ratio as starting materials to form the cyclenoxamide at room temperature with stirring for 48 h (Bellouard *et al.*, 1999). The reaction stoichiometry should be one to one, so the excess diethyl oxalate was separated by chromatography on silica (CHCl₃:isopropylamine, 5:1). The diethyl oxalate reacted with isopropylamine to produce the title compound at room temperature.

The title compound is a salt, consisting of an *N*-isopropylloxamate anion, an isopropylammonium cation and a water molecule (Fig. 1). The C1—C2 bond was 1.549 (2) Å which is longer than the typical C(*sp*²)—C(*sp*²) bond length. This is most likely due to the stretching by hydrogen bonds at the two sides of the bond. There is a partial double bond between the C2 and N1 [1.326 (2) Å] in the amide group. The cation, anion and water molecule are connected by intermolecular N—H \cdots O and O—H \cdots O hydrogen bonds, as detailed in Table 1 and Figs. 1 and 2. This hydrogen-bonding network is extended in the *ab* plane to form a layer. The molecular packing consists of the hydrogen-bonding layers alternating with hydrophobic domains along the *c* direction.

Experimental

The title compound was synthesized by reacting excess diethyl oxalate with isopropylamine in chloroform at room temperature. Crystals were obtained by the slow evaporation method. After two weeks colourless plate crystals were deposited. The compound was characterized by proton NMR in deuterated chloroform as follows: 1.12 p.p.m. (6H, d, CH₃ for anion); 1.13 p.p.m. (6H, m, CH₃ for cation); 3.40 p.p.m. (1H, m, CH for cation); 4.18 p.p.m. (1H, m, CH for anion); 7.35 p.p.m. (1H, d, NH for anion); 7.97(1H, br, NH for cation).

Refinement

H atoms of CH, CH₃ and NH₃ groups were positioned geometrically and refined as riding, with C—H = 1.00 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for CH groups, and with C—H = 0.98 and 0.91 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C}, \text{N})$ for CH₃ and NH₃ groups. H atoms on N1 and on the water molecule were refined isotropically.

Figures

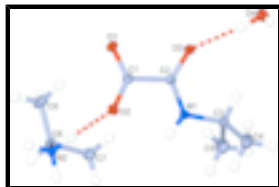


Fig. 1. Molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. The red dashed lines represent hydrogen bonds.

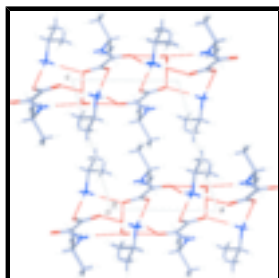


Fig. 2. The molecular packing in the title compound with hydrogen bonds shown as red dashed lines.

Isopropylammonium (isopropylamino)oxoacetate monohydrate

Crystal data

$C_3H_{10}N^+ \cdot C_5H_8NO_3^- \cdot H_2O$

$M_r = 208.26$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 7.172\ (4)\ \text{\AA}$

$b = 9.154\ (5)\ \text{\AA}$

$c = 9.472\ (5)\ \text{\AA}$

$\alpha = 104.372\ (8)^\circ$

$\beta = 105.944\ (8)^\circ$

$\gamma = 94.731\ (8)^\circ$

$V = 571.6\ (5)\ \text{\AA}^3$

$Z = 2$

$F_{000} = 228$

$D_x = 1.210\ \text{Mg m}^{-3}$

Melting point: 56 K

Mo $K\alpha$ radiation

$\lambda = 0.71073\ \text{\AA}$

Cell parameters from 3071 reflections

$\theta = 2.3\text{--}26.5^\circ$

$\mu = 0.10\ \text{mm}^{-1}$

$T = 100\ (2)\ \text{K}$

Plate, colourless

$0.38 \times 0.19 \times 0.08\ \text{mm}$

Data collection

Bruker SMART APEX CCD area-detector diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 100\ (2)\ \text{K}$

φ and ω scans

Absorption correction: multi-scan

(SADABS; Bruker, 2002)

$T_{\min} = 0.962$, $T_{\max} = 0.990$

6062 measured reflections

2324 independent reflections

2126 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.023$

$\theta_{\max} = 26.4^\circ$

$\theta_{\min} = 2.3^\circ$

$h = -9 \rightarrow 8$

$k = -11 \rightarrow 11$

$l = -11 \rightarrow 11$

Refinement

| | |
|--|--|
| Refinement on F^2 | Secondary atom site location: difference Fourier map |
| Least-squares matrix: full | Hydrogen site location: inferred from neighbouring sites |
| $R[F^2 > 2\sigma(F^2)] = 0.045$ | H atoms treated by a mixture of independent and constrained refinement |
| $wR(F^2) = 0.106$ | $w = 1/[\sigma^2(F_o^2) + (0.0443P)^2 + 0.2432P]$ |
| $S = 1.16$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 2324 reflections | $(\Delta/\sigma)_{\max} < 0.001$ |
| 144 parameters | $\Delta\rho_{\max} = 0.40 \text{ e } \text{\AA}^{-3}$ |
| Primary atom site location: structure-invariant direct methods | $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$ |
| | Extinction correction: none |

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|---------------|--------------|--------------|----------------------------------|
| O1 | 0.38400 (15) | 0.13612 (11) | 0.08003 (12) | 0.0186 (2) |
| N1 | 0.36897 (19) | 0.52457 (13) | 0.22589 (14) | 0.0147 (3) |
| C1 | 0.3063 (2) | 0.25345 (16) | 0.09391 (15) | 0.0131 (3) |
| H1 | 0.246 (3) | 0.512 (2) | 0.205 (2) | 0.021 (5)* |
| O2 | 0.12699 (14) | 0.26313 (11) | 0.05100 (11) | 0.0167 (2) |
| N2 | -0.18564 (17) | 0.14413 (13) | 0.13379 (13) | 0.0140 (3) |
| H2A | -0.0868 | 0.1780 | 0.1010 | 0.021* |
| H2B | -0.2743 | 0.2098 | 0.1311 | 0.021* |
| H2C | -0.2463 | 0.0496 | 0.0717 | 0.021* |
| C2 | 0.4503 (2) | 0.40621 (16) | 0.17327 (15) | 0.0131 (3) |
| O3 | 0.62792 (14) | 0.41075 (11) | 0.18756 (12) | 0.0175 (2) |
| C3 | 0.4797 (2) | 0.67704 (16) | 0.31163 (17) | 0.0169 (3) |
| H3 | 0.6232 | 0.6711 | 0.3369 | 0.020* |
| O4 | 0.97028 (16) | 0.59125 (13) | 0.18861 (14) | 0.0211 (3) |
| C4 | 0.4348 (3) | 0.78700 (18) | 0.21497 (19) | 0.0257 (4) |
| H4A | 0.4721 | 0.7511 | 0.1217 | 0.039* |
| H4B | 0.5093 | 0.8888 | 0.2735 | 0.039* |
| H4C | 0.2938 | 0.7921 | 0.1874 | 0.039* |
| C5 | 0.4297 (3) | 0.73017 (19) | 0.45995 (18) | 0.0254 (4) |
| H5A | 0.2892 | 0.7361 | 0.4367 | 0.038* |
| H5B | 0.5061 | 0.8313 | 0.5188 | 0.038* |
| H5C | 0.4616 | 0.6574 | 0.5200 | 0.038* |
| C6 | -0.1027 (2) | 0.13576 (16) | 0.29428 (16) | 0.0161 (3) |
| H6 | -0.2136 | 0.1022 | 0.3300 | 0.019* |
| C7 | -0.0006 (2) | 0.29326 (17) | 0.39773 (17) | 0.0201 (3) |
| H7A | 0.1131 | 0.3254 | 0.3678 | 0.030* |
| H7B | 0.0436 | 0.2897 | 0.5042 | 0.030* |
| H7C | -0.0924 | 0.3664 | 0.3880 | 0.030* |
| C8 | 0.0341 (2) | 0.01765 (18) | 0.29648 (18) | 0.0234 (4) |

supplementary materials

| | | | | |
|-----|-----------|-----------|-----------|------------|
| H8A | -0.0399 | -0.0829 | 0.2328 | 0.035* |
| H8B | 0.0909 | 0.0136 | 0.4018 | 0.035* |
| H8C | 0.1397 | 0.0460 | 0.2563 | 0.035* |
| H9 | 0.939 (3) | 0.622 (3) | 0.113 (3) | 0.041 (6)* |
| H10 | 0.859 (3) | 0.539 (2) | 0.184 (2) | 0.033 (5)* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|----|------------|------------|------------|------------|------------|-------------|
| O1 | 0.0177 (5) | 0.0116 (5) | 0.0241 (6) | 0.0033 (4) | 0.0055 (4) | 0.0013 (4) |
| N1 | 0.0106 (6) | 0.0132 (6) | 0.0187 (6) | 0.0023 (5) | 0.0038 (5) | 0.0022 (5) |
| C1 | 0.0147 (7) | 0.0148 (7) | 0.0106 (6) | 0.0025 (5) | 0.0053 (5) | 0.0032 (5) |
| O2 | 0.0127 (5) | 0.0166 (5) | 0.0203 (5) | 0.0014 (4) | 0.0049 (4) | 0.0046 (4) |
| N2 | 0.0122 (6) | 0.0107 (6) | 0.0170 (6) | 0.0016 (4) | 0.0041 (5) | 0.0007 (5) |
| C2 | 0.0140 (7) | 0.0142 (7) | 0.0122 (6) | 0.0033 (5) | 0.0040 (5) | 0.0050 (5) |
| O3 | 0.0128 (5) | 0.0138 (5) | 0.0248 (6) | 0.0026 (4) | 0.0067 (4) | 0.0021 (4) |
| C3 | 0.0131 (7) | 0.0129 (7) | 0.0201 (7) | 0.0006 (5) | 0.0040 (6) | -0.0019 (6) |
| O4 | 0.0152 (6) | 0.0239 (6) | 0.0258 (6) | 0.0026 (5) | 0.0048 (5) | 0.0116 (5) |
| C4 | 0.0331 (9) | 0.0181 (8) | 0.0276 (9) | 0.0010 (7) | 0.0137 (7) | 0.0056 (7) |
| C5 | 0.0281 (9) | 0.0239 (8) | 0.0186 (8) | 0.0001 (7) | 0.0055 (7) | -0.0009 (6) |
| C6 | 0.0167 (7) | 0.0168 (7) | 0.0148 (7) | 0.0029 (6) | 0.0060 (6) | 0.0031 (5) |
| C7 | 0.0199 (8) | 0.0185 (8) | 0.0178 (7) | 0.0034 (6) | 0.0037 (6) | 0.0001 (6) |
| C8 | 0.0288 (9) | 0.0191 (8) | 0.0214 (8) | 0.0092 (6) | 0.0046 (7) | 0.0056 (6) |

Geometric parameters (\AA , $^\circ$)

| | | | |
|----------|-------------|------------|-----------|
| O1—C1 | 1.2451 (18) | C4—H4A | 0.9800 |
| N1—C2 | 1.3262 (19) | C4—H4B | 0.9800 |
| N1—C3 | 1.4623 (19) | C4—H4C | 0.9800 |
| N1—H1 | 0.84 (2) | C5—H5A | 0.9800 |
| C1—O2 | 1.2554 (18) | C5—H5B | 0.9800 |
| C1—C2 | 1.549 (2) | C5—H5C | 0.9800 |
| N2—C6 | 1.4963 (19) | C6—C7 | 1.518 (2) |
| N2—H2A | 0.9100 | C6—C8 | 1.519 (2) |
| N2—H2B | 0.9100 | C6—H6 | 1.0000 |
| N2—H2C | 0.9100 | C7—H7A | 0.9800 |
| C2—O3 | 1.2394 (18) | C7—H7B | 0.9800 |
| C3—C4 | 1.518 (2) | C7—H7C | 0.9800 |
| C3—C5 | 1.518 (2) | C8—H8A | 0.9800 |
| C3—H3 | 1.0000 | C8—H8B | 0.9800 |
| O4—H9 | 0.81 (2) | C8—H8C | 0.9800 |
| O4—H10 | 0.88 (2) | | |
| C2—N1—C3 | 123.84 (13) | H4A—C4—H4C | 109.5 |
| C2—N1—H1 | 118.0 (13) | H4B—C4—H4C | 109.5 |
| C3—N1—H1 | 118.1 (13) | C3—C5—H5A | 109.5 |
| O1—C1—O2 | 128.18 (13) | C3—C5—H5B | 109.5 |
| O1—C1—C2 | 115.46 (12) | H5A—C5—H5B | 109.5 |
| O2—C1—C2 | 116.36 (12) | C3—C5—H5C | 109.5 |

| | | | |
|-------------|--------------|-------------|--------------|
| C6—N2—H2A | 109.5 | H5A—C5—H5C | 109.5 |
| C6—N2—H2B | 109.5 | H5B—C5—H5C | 109.5 |
| H2A—N2—H2B | 109.5 | N2—C6—C7 | 109.47 (12) |
| C6—N2—H2C | 109.5 | N2—C6—C8 | 109.33 (12) |
| H2A—N2—H2C | 109.5 | C7—C6—C8 | 112.50 (13) |
| H2B—N2—H2C | 109.5 | N2—C6—H6 | 108.5 |
| O3—C2—N1 | 124.74 (13) | C7—C6—H6 | 108.5 |
| O3—C2—C1 | 120.31 (12) | C8—C6—H6 | 108.5 |
| N1—C2—C1 | 114.92 (12) | C6—C7—H7A | 109.5 |
| N1—C3—C4 | 110.08 (13) | C6—C7—H7B | 109.5 |
| N1—C3—C5 | 109.27 (12) | H7A—C7—H7B | 109.5 |
| C4—C3—C5 | 111.50 (13) | C6—C7—H7C | 109.5 |
| N1—C3—H3 | 108.6 | H7A—C7—H7C | 109.5 |
| C4—C3—H3 | 108.6 | H7B—C7—H7C | 109.5 |
| C5—C3—H3 | 108.6 | C6—C8—H8A | 109.5 |
| H9—O4—H10 | 102 (2) | C6—C8—H8B | 109.5 |
| C3—C4—H4A | 109.5 | H8A—C8—H8B | 109.5 |
| C3—C4—H4B | 109.5 | C6—C8—H8C | 109.5 |
| H4A—C4—H4B | 109.5 | H8A—C8—H8C | 109.5 |
| C3—C4—H4C | 109.5 | H8B—C8—H8C | 109.5 |
| C3—N1—C2—O3 | -1.4 (2) | O1—C1—C2—N1 | -162.00 (12) |
| C3—N1—C2—C1 | 176.57 (12) | O2—C1—C2—N1 | 18.40 (17) |
| O1—C1—C2—O3 | 16.08 (19) | C2—N1—C3—C4 | 109.88 (15) |
| O2—C1—C2—O3 | -163.51 (12) | C2—N1—C3—C5 | -127.36 (14) |

Hydrogen-bond geometry (\AA , $^\circ$)

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-----------------------------------|----------|-------------|-------------|---------------|
| O4—H9 \cdots O2 ⁱ | 0.82 (3) | 2.06 (3) | 2.864 (2) | 170 (3) |
| O4—H10 \cdots O3 | 0.88 (2) | 1.96 (2) | 2.8404 (19) | 173.5 (19) |
| N1—H1 \cdots O4 ⁱⁱ | 0.84 (2) | 2.14 (2) | 2.918 (2) | 153.7 (17) |
| N2—H2A \cdots O2 | 0.91 | 1.89 | 2.7957 (19) | 175 |
| N2—H2B \cdots O3 ⁱⁱⁱ | 0.91 | 2.02 | 2.8808 (19) | 157 |
| N2—H2C \cdots O1 ⁱⁱⁱ | 0.91 | 1.91 | 2.8182 (18) | 172 |

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

Fig. 1

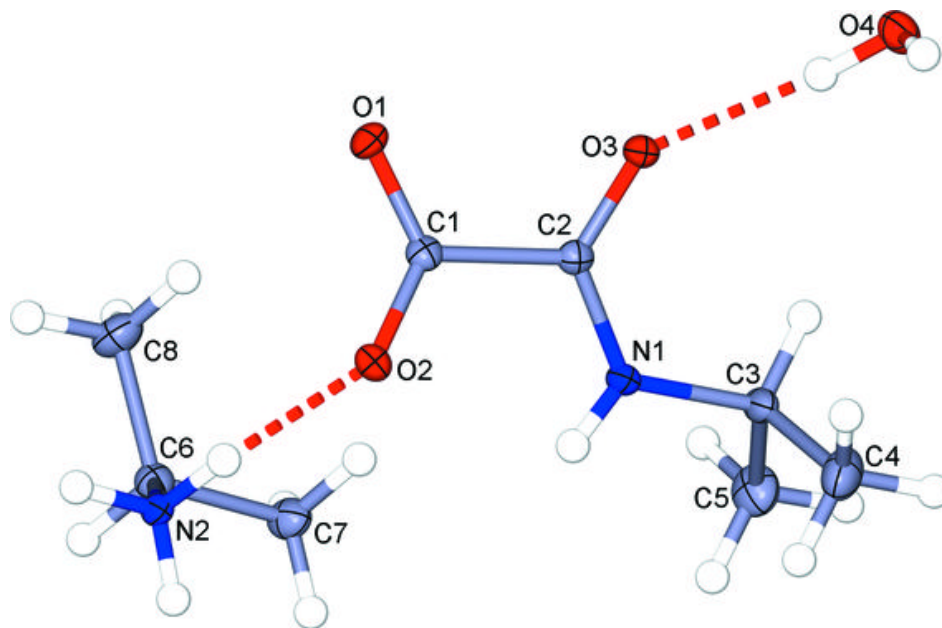


Fig. 2

