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N-(2-Azaniumylethyl)carbamate monohydrate

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Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.002 Å; R factor = 0.034; wR factor = 0.093; data-to-parameter ratio = 12.2.

In the crystal structure of the title compound, $C_3H_8N_2O_2 \cdot H_2O$, the organic molecule exists as zwitterion with the carboxyl group deprotonated and the amino group protonated. In the crystal, the components are linked by $O-H \cdots O$ and $N-H \cdots O$ hydrogen bonds.

Related literature

 CO_2 readily reacts with amines to yied carbamates, see: Brown & Gray (1982); Dell'Amico *et al.* (2003); Jing *et al.* (2007). For *N*-(2-ammonioethyl)carbamate (AECM), a reactive product of ethylenediamine with CO_2 , see: Garbauskas *et al.* (1983); Antsyshkina *et al.* (2007). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\begin{array}{l} C_{3}H_{8}N_{2}O_{2} \cdot H_{2}O\\ M_{r} = 122.13\\ \text{Monoclinic, } P2_{1}/c\\ a = 8.0301 \ (6) \ \text{\AA}\\ b = 8.7842 \ (7) \ \text{\AA}\\ c = 8.1748 \ (6) \ \text{\AA}\\ \beta = 98.889 \ (1)^{\circ} \end{array}$

 $V = 569.71 (7) Å^{3}$ Z = 4Mo K\alpha radiation $\mu = 0.12 \text{ mm}^{-1}$ T = 293 K $0.35 \times 0.34 \times 0.30 \text{ mm}$ 2877 measured reflections

 $R_{\rm int} = 0.016$

1002 independent reflections

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

Data collection

Bruker APEX area-detector diffractometer Absorption correction: multi-scan (SADABS; Bruker, 2001) $T_{min} = 0.945, T_{max} = 0.966$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$	H atoms treated by a mixture of
$wR(F^2) = 0.093$	independent and constrained
S = 1.04	refinement
1002 reflections	$\Delta \rho_{\rm max} = 0.21 \text{ e } \text{\AA}^{-3}$
82 parameters	$\Delta \rho_{\rm min} = -0.27 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
$O3-H3A\cdots O1$	0.80(3)	1.92 (3)	2.708(2)	170 (3)
$N1 - H1C \cdots O3^{ii}$	0.86 (3) 0.89	1.92 (3) 1.89	2.767 (2)	171 (3) 167
$N1 - H1D \cdots O2^{m}$ $N1 - H1E \cdots O1^{iv}$	0.89 0.89	1.91 1.95	2.775 (2) 2.798 (2)	163 158
$N2 - H2 \cdot \cdot \cdot O2^{v}$	0.86	2.43	3.278 (2)	167 162
C2=H2A···01	0.97	2.30	5.499 (2)	105

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

Data collection: *SMART* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NC2244).

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S1. Comment

It has been known for decades that CO₂ readily reacts with amines to yied carbamates (Brown & Gray 1982; Dell'Amico *et al.* 2003; Jing *et al.* 2007). *N*-(2-ammonioethyl)carbamate (AECM), a reactive product of ethylenediamine with CO₂, was reported previously (Garbauskas *et al.* 1983; Antsyshkina *et al.* 2007). Recently, AECM hydrate, (I) (Scheme 1, Table 1), is prepared from ethylenediamine as starting material in our lab, and its structure is studied hereafter.

In (I), AECM molecule exists as zwitterion, the molecule is linked with the water molecule by an O3—H3A⁻⁻O1 hydrogen bond (Fig. 1, Table 2). The N1 atom is protonated, showing as the center of positive charge. The negative charge is concentrated on the O2 atom of the COO- fragment and is somewhat delocalized: the C3—O1 and C3—N2 bonds are slightly elongated, and the N2—C2 bond is shortened compared to standard values of 1.21, 1.334 and 1.454 Å, respectively (Allen *et al.* 1987). The torsion angle of N1—C1—C2—N2 [46.21 (18)%] is much smaller than that observed in the one of Garbauskas' polymorphs (175.6%), and is smaller than those observed in the second polymorph (66.6% in Antsyshkina's case, 65.5% in Garbauskas' case).

There are many hydrogen bonds in the crystal (Fig. 1, Table 2), playing important role in restraining the AECM comformation, and in building the crystal.

S2. Experimental

Ethylenediamine (10.1 ml) was dissolved in xylenol (25.2 ml), forming clear solution with stirring, afterwards, the resulting solution was exposed in the air for two month at room temperature. With the reaction deepened, the system separated into two layers gradually. Upper layer was yellowish and pasty, and lower layer was colorless and clear. Crystals of (I) (6.9 g) were at the bottom of the lower lay. Analysis: Cald. for (I) (%): C 29.50, H 8.25, N 22.94; found: C 29.45, H 8.31, N 22.90. IR Spectrum (KBr, cm⁻¹): 3289(*s*), 2964(*m*), 2214(w), 1673(*m*), 1601(*s*), 1492(*s*), 1381(*s*), 1332(*s*), 1210(w), 1146(*m*), 1050(w), 1029(w), 1010(w), 887(w), 861(w), 821(*m*), 725(*m*), 646(w), 555(*m*). ¹H NMR (500 MHz, D2O) δ /p.p.m.: 3.20 (t, 2 H, J = 5.95), 2.97 (t, 2H, J = 5.95).

S3. Refinement

H atoms of water melecule were deduced from Fourier Maps, and incoporated in refinement freely. The others were placed in calculated positions and allowed to ride on their parent atoms at distances of 0.86Å for acidamide N—H, 0.89Å for amonnium N—H and 0.97Å for ethylene C—H, respectively, with isotropic displacement parameters 1.2–1.5 times U_{eq} of the parent atoms.



Figure 1

Crystal structure of (I) with labeling and displacement ellipsoids drawn at the 40% probability level. Intermolecular hydrogen bonding is shown as a dashed line.





The crystal packing of (I) viewed down the b axis. Hydrogen bonds are drawn as dashed lines.

N-(2-Azaniumylethyl)carbamate monohydrate

Crystal data

C₃H₈N₂O₂·H₂O $M_r = 122.13$ Monoclinic, $P2_1/c$ Hall symbol: -P 2ybc a = 8.0301 (6) Å b = 8.7842 (7) Å c = 8.1748 (6) Å $\beta = 98.889$ (1)° V = 569.71 (7) Å³ Z = 4

Data collection

Bruker APEX area-detector diffractometer Radiation source: fine-focus sealed tube Graphite monochromator φ and ω scan Absorption correction: multi-scan (*SADABS*; Bruker, 2001) $T_{\min} = 0.945, T_{\max} = 0.966$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier
Least-squares matrix: full	map
$R[F^2 > 2\sigma(F^2)] = 0.034$	Hydrogen site location: inferred from
$wR(F^2) = 0.093$	neighbouring sites
S = 1.04	H atoms treated by a mixture of independent
1002 reflections	and constrained refinement
82 parameters	$w = 1/[\sigma^2(F_o^2) + (0.0545P)^2 + 0.1863P]$
0 restraints	where $P = (F_o^2 + 2F_c^2)/3$
Primary atom site location: structure-invariant	$(\Delta/\sigma)_{\rm max} < 0.001$
direct methods	$\Delta \rho_{\rm max} = 0.21 \ { m e} \ { m \AA}^{-3}$
	$\Delta \rho_{\rm min} = -0.27 \ {\rm e} \ {\rm \AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

F(000) = 264.0

 $\theta = 2.4 - 18.3^{\circ}$

 $\mu = 0.12 \text{ mm}^{-1}$

Block, colorless

 $0.35 \times 0.34 \times 0.30$ mm

2877 measured reflections

 $\theta_{\rm max} = 25.0^\circ, \ \theta_{\rm min} = 2.6^\circ$

1002 independent reflections

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

T = 293 K

 $R_{\rm int} = 0.016$

 $h = -9 \rightarrow 9$

 $l = -9 \rightarrow 6$

 $k = -10 \rightarrow 10$

 $D_{\rm x} = 1.424 {\rm Mg m^{-3}}$

Melting point: 358 K

Mo *K* α radiation, $\lambda = 0.71073$ Å

Cell parameters from 1358 reflections

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2 , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
03	0.79493 (17)	0.47441 (13)	0.42240 (14)	0.0480 (4)	
H3A	0.788 (2)	0.472 (2)	0.326 (3)	0.050 (5)*	
H3B	0.844 (3)	0.391 (3)	0.466 (3)	0.069 (6)*	

supporting information

O2	0.91742 (12)	0.29983 (10)	0.06003 (11)	0.0324 (3)	
01	0.75733 (13)	0.50450 (10)	0.08899 (11)	0.0325 (3)	
N1	0.82463 (13)	0.69023 (12)	-0.32825 (13)	0.0270 (3)	
H1D	0.9110	0.6736	-0.2480	0.041*	
H1E	0.8288	0.7855	-0.3644	0.041*	
H1C	0.8303	0.6260	-0.4114	0.041*	
N2	0.76650 (14)	0.40847 (12)	-0.16430 (13)	0.0265 (3)	
H2	0.8216	0.3547	-0.2257	0.032*	
C3	0.81591 (15)	0.40494 (13)	0.00256 (15)	0.0235 (3)	
C2	0.62537 (16)	0.49851 (15)	-0.24347 (16)	0.0280 (3)	
H2A	0.5340	0.4898	-0.1790	0.034*	
H2B	0.5862	0.4560	-0.3521	0.034*	
C1	0.66425 (16)	0.66607 (15)	-0.26316 (16)	0.0288 (3)	
H1A	0.5729	0.7126	-0.3380	0.035*	
H1B	0.6709	0.7161	-0.1566	0.035*	

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
03	0.0844 (9)	0.0333 (6)	0.0250 (6)	0.0148 (6)	0.0046 (6)	-0.0027 (5)
02	0.0397 (5)	0.0250 (5)	0.0294 (5)	0.0052 (4)	-0.0050 (4)	0.0000 (4)
01	0.0459 (6)	0.0274 (5)	0.0252 (5)	0.0038 (4)	0.0084 (4)	-0.0035 (4)
N1	0.0343 (6)	0.0222 (5)	0.0239 (5)	-0.0015 (4)	0.0022 (4)	0.0029 (4)
N2	0.0349 (6)	0.0236 (6)	0.0213 (6)	0.0056 (4)	0.0050 (4)	0.0001 (4)
C3	0.0279 (6)	0.0183 (6)	0.0243 (6)	-0.0041 (5)	0.0044 (5)	0.0003 (5)
C2	0.0273 (6)	0.0297 (7)	0.0263 (7)	-0.0017 (5)	0.0018 (5)	0.0038 (5)
C1	0.0316 (7)	0.0263 (7)	0.0283 (7)	0.0062 (5)	0.0036 (5)	0.0023 (5)

Geometric parameters (Å, °)

03—H3A	0.78 (2)	N2—C3	1.3608 (16)	
O3—H3B	0.88 (3)	N2—C2	1.4499 (16)	
O2—C3	1.2725 (15)	N2—H2	0.8600	
O1—C3	1.2603 (15)	C2—C1	1.5184 (18)	
N1-C1	1.4828 (16)	C2—H2A	0.9700	
N1—H1D	0.8900	C2—H2B	0.9700	
N1—H1E	0.8900	C1—H1A	0.9700	
N1—H1C	0.8900	C1—H1B	0.9700	
H3A—O3—H3B	110 (2)	N2—C2—C1	114.63 (10)	
C1—N1—H1D	109.5	N2—C2—H2A	108.6	
C1—N1—H1E	109.5	C1—C2—H2A	108.6	
H1D—N1—H1E	109.5	N2—C2—H2B	108.6	
C1—N1—H1C	109.5	C1—C2—H2B	108.6	
H1D—N1—H1C	109.5	H2A—C2—H2B	107.6	
H1E—N1—H1C	109.5	N1-C1-C2	112.39 (10)	
C3—N2—C2	123.13 (10)	N1—C1—H1A	109.1	
C3—N2—H2	118.4	C2—C1—H1A	109.1	

supporting information

C2—N2—H2 O1—C3—O2 O1—C3—N2 O2—C3—N2	118.4 124.74 (11) 118.03 (11) 117.23 (11)	N1—C1—H1B C2—C1—H1B H1A—C1—H1B	109.1 109.1 107.9
C2—N2—C3—O1	-13.44 (17)	C3—N2—C2—C1	79.98 (15)
C2—N2—C3—O2	165.99 (11)	N2—C2—C1—N1	46.09 (15)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	D—H	H···A	D···A	D—H··· A	
03—H3 <i>A</i> …O1	0.80(3)	1.92 (3)	2.708 (2)	170 (3)	
O3—H3 <i>B</i> ···O2 ⁱ	0.86 (3)	1.92 (3)	2.773 (2)	171 (3)	
N1—H1 <i>C</i> ···O3 ⁱⁱ	0.89	1.89	2.767 (2)	167	
N1—H1D····O2 ⁱⁱⁱ	0.89	1.91	2.775 (2)	163	
N1—H1 E ···O1 ^{iv}	0.89	1.95	2.798 (2)	158	
N2—H2···O2 ^v	0.86	2.43	3.278 (2)	167	
C2—H2A····O1 ^{vi}	0.97	2.56	3.499 (2)	163	

Symmetry codes: (i) x, -y+1/2, z+1/2; (ii) x, y, z-1; (iii) -x+2, -y+1, -z; (iv) x, -y+3/2, z-1/2; (v) x, -y+1/2, z-1/2; (vi) -x+1, -y+1, -z.