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Bis(adamantan-1-aminium) carbonate

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Key indicators: single-crystal X-ray study; T = 173 K; mean σ (C–C) = 0.002 Å; R factor = 0.042; wR factor = 0.107; data-to-parameter ratio = 14.0.

In the title compound, $2C_{10}H_{18}N^+ \cdot CO_3^{2-}$, the adamantan-1aminium cation forms three $N-H \cdot \cdot \cdot O$ hydrogen bonds to three carbonate ions, resulting in a layer parallel to (001) with the adamantane groups located on its surface so that adjacent layers form only $C-H \cdot \cdot \cdot H - C$ contacts. The carbonate anions occupy special positions of 32 symmetry, whereas the adamantan-1-aminium cations occupy special positions of 3 symmetry.

Related literature

For related structures, see: de Vries *et al.* (2011); Mullica *et al.* (1999); He & Wen (2006); Liu *et al.* (2009); Zhao *et al.* (2003). For applications of adamantane–ammonium salts in virology, see: Hoffmann (1973); Dolin *et al.* (1982); Bright *et al.* (2005); Betakova (2007). For applications of amines for the capture of CO_2 from the atmosphere, see: Yang *et al.* (2008).

 H_3N^+ CO_3^2

Experimental

Crystal data

 $2C_{10}H_{18}N^+ \cdot CO_3^{2-}$ $M_r = 364.52$ Trigonal, $P\overline{3}c1$ a = 6.4340 (6) Å c = 25.474 (2) Å V = 913.25 (14) Å³ Z = 2 Mo Kα radiation μ = 0.09 mm⁻¹ T = 173 K 0.30 × 0.22 × 0.08 mm organic compounds

629 independent reflections

 $R_{\rm int} = 0.062$

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

Data collection

Bruker APEXII CCD

dı	fractomete	er
3187	measured	reflections

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$	H atoms treated by a mixture of
$vR(F^2) = 0.107$	independent and constrained
S = 1.08	refinement
29 reflections	$\Delta \rho_{\rm max} = 0.22 \ {\rm e} \ {\rm \AA}^{-3}$
5 parameters	$\Delta \rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1 \cdots O1^i$	0.999 (16)	1.778 (15)	2.764 (1)	168.7 (18)
Symmetry code: (i)	-y + 1, x - y + 1	l, z.		

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT-Plus* (Bruker, 2005); data reduction: *SAINT-Plus* and *XPREP* (Bruker 2005); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997), *Mercury* (Macrae *et al.*, 2008) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2467).

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S1. Comment

It has been reported that 1-aminoadamantane hydrochloride (marketed as Symmetrel) is effective in the prevention and treatment of the influenza (A) virus (Hoffmann, 1973; Dolin *et al.*, 1982; Bright *et al.*, 2005). However recent studies suggest that the virus is becoming increasingly resistant to this anti-influenza drug (Betakova, 2007).

In an attempt to crystallize pure 1-aminoadamantane from ethanol we obtained instead adamantan-1-aminium carbonate, illustrated in Fig. 1, suggesting that the amine had captured atmospheric CO_2 . We report the structure here. It is known that organic amines can trap CO_2 as the ammonium carbonate salt and this property is being explored as a way to capture carbon dioxide from the atmosphere (Yang *et al.*, 2008).

Each carbonate ion of the title compound forms hydrogen bonds to six adamantane-ammonium ions, as shown in Fig. 2, forming a two-dimensional layer of adamantan-1-aminium carbonates parallel to (001). The hydrophobic adamantane layers interact with the neighbouring layers of adamantane-ammonium molecules *via* C—H···H–C contacts (see Fig. 3).

It is noted here that the structure of adamantan-1-aminium bicarbonate (Liu *et al.*, 2009) reported in the literature is isomorphous to adamantan-1-aminium nitrate (Zhao *et al.*, 2003). The former structure has unusually short H···H intermolecular contacts between NH_3^+ group H atom and bicarbonate H atom of 1.50 Å In addition the geometry of the hydrogen carbonate ion is very similar to that of the nitrate ion. A re-investigation of these structures is warranted.

S2. Experimental

Crystals were grown by slow evaporation of an ethanol solution of the title compound, 0.500 g in 10 ml of ethanol, and afforded colourless plates after three days under ambient conditions. Crystals decompose, with an emission of gas bubbles (presumably CO_2), at 423–428 K.

S3. Refinement

The N-bound H atom was placed according to the observed electron density and was allowed to refine freely. The remaining H atoms were positioned geometrically and allowed to ride on their respective parent atoms, with C—H bond lengths of 1.00 (methine) and 0.99 Å (methylene CH₂) and with $U_{iso}(H) = 1.2$ times $U_{eq}(C)$.



Figure 1

The molecular structure of the title compound. Displacement ellipsoids are shown at the 50% probability level. The atoms C2f to C4f are generated by the symmetry (1-y,x-y,z); C2g to C4g by (1-x+y, 1-x,z); O1a by (-y,x-y,z) and O2b by (-x+y,-x,z).



Figure 2

Intermolecular N—H…O hydrogen bonded (dashed lines) layers along [001] showing only the C-NH₃ and CO₃ groups for clarity.



Figure 3

A view down the b axis of the unit cell of the title compound showing the hydrogen bonded layers. Notice that the carbonate ions occupy sites with 32 symmetry whereas cations occupy the sites of 3 symmetry.

Bis(adamantan-1-aminium) carbonate

Crystal data

 $2C_{10}H_{18}N^{+}CO_{3}^{2-}$ $M_{r} = 364.52$ Trigonal, P3c1Hall symbol: -P 3 2"c a = 6.4340 (6) Å c = 25.474 (2) Å V = 913.25 (14) Å³ Z = 2F(000) = 400

Data collection

Bruker APEXII CCD diffractometer Graphite monochromator φ and ω scans 3187 measured reflections 629 independent reflections

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.042$ $wR(F^2) = 0.107$ S = 1.08629 reflections 45 parameters 0 restraints $D_{\rm x} = 1.326 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation, $\lambda = 0.71069 \text{ Å}$ Cell parameters from 819 reflections $\theta = 3.2-25.8^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 173 KPrism, colourless $0.30 \times 0.22 \times 0.08 \text{ mm}$

493 reflections with $I > 2\sigma(I)$ $R_{int} = 0.062$ $\theta_{max} = 26.4^{\circ}, \ \theta_{min} = 1.6^{\circ}$ $h = -8 \rightarrow 5$ $k = -2 \rightarrow 8$ $l = -31 \rightarrow 31$

Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H atoms treated by a mixture of independent and constrained refinement

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0587P)^{2} + 0.010P] \qquad \Delta \rho_{max} = 0.22 \text{ e } \text{\AA}^{-3}$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3 \qquad \Delta \rho_{min} = -0.18 \text{ e } \text{\AA}^{-3}$ $(\Delta/\sigma)_{max} < 0.001$

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.

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	$U_{ m iso}$ */ $U_{ m eq}$	
C1	0.6667	0.3333	0.16486 (9)	0.0236 (6)	
C2	0.9213 (2)	0.5034 (2)	0.14518 (6)	0.0278 (4)	
H2A	1.032	0.4505	0.1584	0.033*	
H2B	0.977	0.6678	0.1583	0.033*	
C3	0.9215 (2)	0.5033 (3)	0.08498 (6)	0.0309 (4)	
H3	1.0877	0.614	0.0719	0.037*	
C4	0.8365 (3)	0.2482 (3)	0.06502 (6)	0.0346 (4)	
H4A	0.838	0.2474	0.0262	0.042*	
H4B	0.9464	0.1932	0.0777	0.042*	
C5	0	0	0.25	0.0219 (7)	
N1	0.6667	0.3333	0.22372 (8)	0.0278 (5)	
01	0	0.1993 (2)	0.25	0.0343 (4)	
H1	0.732 (4)	0.502 (3)	0.2358 (7)	0.060 (5)*	

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0198 (8)	0.0198 (8)	0.0312 (12)	0.0099 (4)	0	0
C2	0.0195 (8)	0.0205 (7)	0.0415 (9)	0.0087 (6)	-0.0018 (6)	-0.0014 (6)
C3	0.0217 (8)	0.0262 (8)	0.0404 (9)	0.0085 (6)	0.0061 (6)	0.0039 (6)
C4	0.0317 (9)	0.0350 (9)	0.0397 (8)	0.0185 (8)	0.0055 (6)	-0.0016 (7)
C5	0.0214 (10)	0.0214 (10)	0.0228 (15)	0.0107 (5)	0	0
N1	0.0253 (7)	0.0253 (7)	0.0328 (11)	0.0127 (3)	0	0
01	0.0299 (8)	0.0221 (6)	0.0536 (10)	0.0149 (4)	-0.0091 (7)	-0.0045 (3)

Geometric parameters (Å, °)

C1—N1	1.500 (3)	C3—C4	1.534 (2)
C1—C2	1.5295 (14)	С3—Н3	1
C2—C3	1.5335 (19)	C4—H4A	0.99
C2—H2A	0.99	C4—H4B	0.99
C2—H2B	0.99	C5—O1	1.2820 (13)

C3—C4 ⁱ	1.532 (2)	N1—H1	0.999 (16)
N1—C1—C2	109.13 (9)	C2—C3—C4	109.40 (12)
C2 ⁱⁱ —C1—C2	109.81 (9)	С2—С3—Н3	109.5
C1—C2—C3	109.18 (12)	C4—C3—H3	109.5
C1—C2—H2A	109.8	C3 ⁱⁱ —C4—C3	109.57 (13)
C3—C2—H2A	109.8	C3—C4—H4A	109.8
C1—C2—H2B	109.8	C3—C4—H4B	109.8
C3—C2—H2B	109.8	H4A—C4—H4B	108.2
H2A—C2—H2B	108.3	O1 ⁱⁱⁱ —C5—O1	120
C4 ⁱ —C3—C2	109.29 (11)	C1—N1—H1	107.9 (11)
C4 ⁱ —C3—C4	109.58 (14)		
N1—C1—C2—C3	179.92 (8)	C1—C2—C3—C4	-59.84 (12)
C2 ⁱⁱ —C1—C2—C3	60.34 (13)	C4 ⁱ —C3—C4—C3 ⁱⁱ	-59.76 (18)
C2 ⁱ —C1—C2—C3	-60.50 (13)	C2C3C4C3 ⁱⁱ	60.04 (15)
C1-C2-C3-C4 ⁱ	60.13 (13)		

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

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	Н…А	D····A	D—H···A
N1—H1···O1 ^{iv}	0.999 (16)	1.778 (15)	2.7644 (11)	168.7 (18)

Symmetry code: (iv) -y+1, x-y+1, z.