

1-Hydroxymethyl-1-methylethan-aminium chloride

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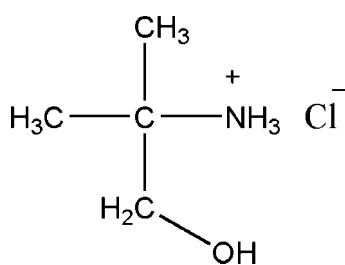
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.041; wR factor = 0.154; data-to-parameter ratio = 18.5.

The asymmetric unit of the title compound, $\text{C}_4\text{H}_{12}\text{NO}^+\cdot\text{Cl}^-$, contains two independent ion pairs. Weak intramolecular $\text{C}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds result in the formation of three five-membered rings, which have envelope conformations. The crystal structure contains intermolecular $\text{O}-\text{H}\cdots\text{Cl}$, $\text{N}-\text{H}\cdots\text{O}$, $\text{N}-\text{H}\cdots\text{Cl}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For related literature, see: Senkus (1948). For general background, see: Pazenok (2007). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_4\text{H}_{12}\text{NO}^+\cdot\text{Cl}^-$	$V = 1353.8(5)\text{ \AA}^3$
$M_r = 125.60$	$Z = 8$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 6.4940(13)\text{ \AA}$	$\mu = 0.46\text{ mm}^{-1}$
$b = 9.5230(19)\text{ \AA}$	$T = 298(2)\text{ K}$
$c = 21.903(4)\text{ \AA}$	$0.30 \times 0.20 \times 0.20\text{ mm}$
$\beta = 91.88(3)^\circ$	

Data collection

Enraf-Nonius CAD-4
diffractometer
Absorption correction: ψ scan
(North *et al.*, 1968)
 $T_{\min} = 0.874$, $T_{\max} = 0.913$
2651 measured reflections

2421 independent reflections
1857 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$
3 standard reflections
frequency: 120 min
intensity decay: none

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.153$
 $S = 1.01$
2421 reflections
131 parameters
1 restraint

H atoms treated by a mixture of
independent and constrained
refinement
 $\Delta\rho_{\max} = 0.26\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.28\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1A \cdots Cl1	0.82 (3)	2.25 (4)	3.067 (2)	174 (3)
N1—H1B \cdots O2	0.89	2.02	2.876 (3)	161
N1—H1C \cdots Cl1 ⁱ	0.89	2.33	3.216 (3)	171
N1—H1G \cdots Cl1 ⁱⁱ	0.89	2.42	3.261 (2)	157
O2—H2D \cdots O1 ⁱⁱⁱ	0.82	1.91	2.731 (3)	175
N2—H2E \cdots Cl1	0.89	2.53	3.391 (2)	163
N2—H2E \cdots O2	0.89	2.49	2.811 (3)	102
N2—H2F \cdots Cl2 ^{iv}	0.89	2.24	3.130 (2)	177
N2—H2G \cdots Cl2	0.89	2.27	3.144 (2)	169
C2—H2A \cdots O1	0.96	2.53	2.880 (4)	102
C6—H6A \cdots O2	0.96	2.59	2.933 (4)	101

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

Data collection: *CAD-4 Software* (Enraf-Nonius, 1985); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); 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: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HK2463).

References

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supplementary materials

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Comment

The title compound, (I), is an important intermediate for the synthesis of 2-amino-2-methylpropylsulfate, which can be used to synthesize 2-methyl-1-(methyl-thio)propane-2-amine (Pazenok, 2007). We report herein the crystal structure of the title compound, (I).

The asymmetric unit of (I) contains two independent molecules (Fig. 1). The bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. The weak intramolecular C-H···O and N-H···O hydrogen bonds (Table 1) result in the formation of three five-membered rings A (C2-C4/O1/H2A), B (C6-C8/O2/H6A) and C (C7/C8/O2/N2/H2E). They adopt envelope conformations, with C3 and C7 atoms displaced by -0.637 (3), -0.686 (4) and 0.711 (3) Å from the planes of the other ring atoms, respectively.

In the crystal structure, intramolecular O-H···Cl, N-H···O and N-H···Cl and intermolecular N-H···Cl and O-H···O hydrogen bonds (Table 1) link the molecules to form a three dimensional network (Fig. 2), in which they may be effective in the stabilization of the structure.

Experimental

The title compound, (I), was synthesized according to the literature method (Senkus, 1948). Crystals suitable for X-ray analysis were obtained by dissolving (I) (0.30 g, 2.4 mmol) in methanol (25 ml) and evaporating the solvent slowly at room temperature for about 4 d.

Refinement

H1A atom (for OH) was located in a difference syntheses and refined [$O1-H1A = 0.82$ (3) Å and $U_{iso}(H) = 0.070 \text{ \AA}^2$]. The remaining H atoms were positioned geometrically, with O-H = 0.82 Å (for OH), N-H = 0.89 Å (for NH₃) and C-H = 0.97 and 0.96 Å for methylene and methyl H, respectively, and constrained to ride on their parent atoms with $U_{iso}(H) = xU_{eq}(C,O,N)$, where $x = 1.2$ for methylene H, and $x = 1.5$ for all other H atoms.

Figures

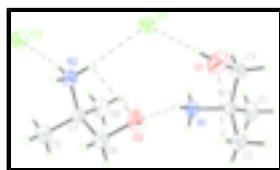


Fig. 1. The molecular structure of the title molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 40% probability level. Hydrogen bonds are shown as dashed lines.

supplementary materials

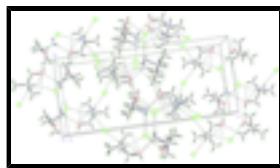


Fig. 2. A packing diagram of (I). Hydrogen bonds are shown as dashed lines.

1-Hydroxymethyl-1-methylethanaminium chloride

Crystal data

$\text{C}_4\text{H}_{12}\text{NO}^+\cdot\text{Cl}^-$	$F_{000} = 544$
$M_r = 125.60$	$D_x = 1.232 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
Hall symbol: -P 2yn	$\lambda = 0.71073 \text{ \AA}$
$a = 6.4940 (13) \text{ \AA}$	Cell parameters from 25 reflections
$b = 9.5230 (19) \text{ \AA}$	$\theta = 11\text{--}14^\circ$
$c = 21.903 (4) \text{ \AA}$	$\mu = 0.46 \text{ mm}^{-1}$
$\beta = 91.88 (3)^\circ$	$T = 298 (2) \text{ K}$
$V = 1353.8 (5) \text{ \AA}^3$	Block, colorless
$Z = 8$	$0.30 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Enraf-Nonius CAD-4 diffractometer	$R_{\text{int}} = 0.020$
Radiation source: fine-focus sealed tube	$\theta_{\text{max}} = 25.2^\circ$
Monochromator: graphite	$\theta_{\text{min}} = 1.9^\circ$
$T = 298(2) \text{ K}$	$h = 0 \rightarrow 7$
$\omega/2\theta$ scans	$k = 0 \rightarrow 11$
Absorption correction: ψ scan (North <i>et al.</i> , 1968)	$l = -26 \rightarrow 26$
$T_{\text{min}} = 0.874$, $T_{\text{max}} = 0.913$	3 standard reflections
2651 measured reflections	every 120 min
2421 independent reflections	intensity decay: none
1857 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.040$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.154$	$w = 1/[\sigma^2(F_o^2) + (0.1P)^2 + 0.25P]$
$S = 1.01$	where $P = (F_o^2 + 2F_c^2)/3$
2421 reflections	$(\Delta/\sigma)_{\text{max}} = 0.001$
	$\Delta\rho_{\text{max}} = 0.26 \text{ e \AA}^{-3}$

131 parameters	$\Delta\rho_{\min} = -0.28 \text{ e } \text{\AA}^{-3}$
1 restraint	Extinction correction: SHELXL97 (Sheldrick, 2008), $F_c^* = k F_c [1 + 0.001 x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.018 (4)

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 $F^2 > 2\sigma(F^2)$ 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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	-0.24225 (12)	0.14202 (8)	0.53408 (3)	0.0414 (3)
Cl2	-0.24989 (15)	0.01158 (9)	0.76365 (4)	0.0527 (3)
O1	-0.0723 (3)	0.3564 (2)	0.44418 (11)	0.0468 (6)
H1A	-0.121 (6)	0.304 (4)	0.4694 (14)	0.070*
O2	0.1531 (3)	0.3696 (2)	0.58458 (8)	0.0383 (5)
H2D	0.1269	0.4503	0.5738	0.057*
N1	0.2905 (4)	0.1957 (2)	0.48659 (10)	0.0313 (6)
H1B	0.2196	0.2435	0.5138	0.047*
H1C	0.4221	0.1910	0.4990	0.047*
H1G	0.2393	0.1093	0.4830	0.047*
N2	-0.0750 (4)	0.2407 (2)	0.67597 (10)	0.0304 (6)
H2E	-0.1178	0.2352	0.6370	0.046*
H2F	-0.1221	0.3195	0.6923	0.046*
H2G	-0.1221	0.1672	0.6964	0.046*
C1	0.3962 (6)	0.1829 (4)	0.38056 (15)	0.0485 (9)
H1D	0.3885	0.2266	0.3411	0.073*
H1E	0.3397	0.0899	0.3777	0.073*
H1F	0.5376	0.1778	0.3947	0.073*
C2	0.3599 (5)	0.4154 (3)	0.43319 (14)	0.0410 (7)
H2A	0.2787	0.4665	0.4615	0.061*
H2B	0.3559	0.4628	0.3945	0.061*
H2C	0.4998	0.4101	0.4486	0.061*
C3	0.2741 (4)	0.2692 (3)	0.42531 (12)	0.0300 (6)
C4	0.0495 (5)	0.2723 (3)	0.40505 (14)	0.0383 (7)
H4A	0.0375	0.3090	0.3638	0.046*
H4B	-0.0039	0.1771	0.4044	0.046*
C5	0.2240 (5)	0.2504 (4)	0.74660 (14)	0.0467 (8)
H5A	0.1744	0.3366	0.7635	0.070*

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H5B	0.3717	0.2483	0.7502	0.070*
H5C	0.1685	0.1724	0.7684	0.070*
C6	0.2298 (5)	0.1056 (3)	0.65041 (15)	0.0464 (8)
H6A	0.1841	0.1034	0.6083	0.070*
H6B	0.1740	0.0265	0.6714	0.070*
H6C	0.3776	0.1016	0.6530	0.070*
C7	0.1566 (4)	0.2409 (3)	0.67977 (12)	0.0305 (6)
C8	0.2295 (5)	0.3699 (3)	0.64625 (13)	0.0369 (7)
H8A	0.1820	0.4536	0.6667	0.044*
H8B	0.3789	0.3717	0.6470	0.044*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0418 (5)	0.0385 (5)	0.0435 (5)	-0.0061 (3)	-0.0065 (3)	0.0059 (3)
Cl2	0.0708 (6)	0.0471 (5)	0.0406 (5)	-0.0228 (4)	0.0074 (4)	0.0050 (3)
O1	0.0390 (12)	0.0394 (13)	0.0628 (15)	0.0087 (10)	0.0117 (11)	0.0180 (10)
O2	0.0545 (13)	0.0310 (11)	0.0296 (11)	0.0003 (10)	0.0050 (9)	0.0044 (8)
N1	0.0321 (13)	0.0338 (12)	0.0281 (12)	0.0047 (10)	0.0007 (10)	0.0042 (10)
N2	0.0340 (13)	0.0274 (12)	0.0299 (12)	0.0009 (10)	0.0042 (10)	-0.0003 (9)
C1	0.059 (2)	0.053 (2)	0.0340 (16)	0.0105 (17)	0.0127 (15)	-0.0028 (14)
C2	0.0424 (18)	0.0367 (17)	0.0435 (17)	-0.0089 (14)	-0.0020 (14)	0.0048 (13)
C3	0.0355 (15)	0.0314 (15)	0.0230 (13)	-0.0019 (12)	0.0018 (11)	0.0026 (11)
C4	0.0381 (17)	0.0379 (16)	0.0385 (16)	-0.0033 (14)	-0.0065 (13)	0.0047 (13)
C5	0.052 (2)	0.057 (2)	0.0313 (16)	-0.0046 (17)	-0.0061 (14)	0.0075 (14)
C6	0.056 (2)	0.0366 (17)	0.0473 (19)	0.0154 (15)	0.0069 (16)	0.0066 (14)
C7	0.0299 (14)	0.0332 (15)	0.0284 (14)	0.0004 (12)	0.0029 (11)	0.0015 (11)
C8	0.0416 (17)	0.0365 (16)	0.0328 (15)	-0.0097 (14)	0.0020 (13)	0.0024 (12)

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

O1—C4	1.430 (4)	O2—C8	1.424 (3)
O1—H1A	0.82 (3)	O2—H2D	0.8200
N1—C3	1.515 (3)	N2—C7	1.503 (4)
N1—H1B	0.8900	N2—H2E	0.8900
N1—H1C	0.8900	N2—H2F	0.8900
N1—H1G	0.8900	N2—H2G	0.8900
C1—C3	1.522 (4)	C5—C7	1.517 (4)
C1—H1D	0.9600	C5—H5A	0.9600
C1—H1E	0.9600	C5—H5B	0.9600
C1—H1F	0.9600	C5—H5C	0.9600
C2—C3	1.507 (4)	C6—C7	1.523 (4)
C2—H2A	0.9600	C6—H6A	0.9600
C2—H2B	0.9600	C6—H6B	0.9600
C2—H2C	0.9600	C6—H6C	0.9600
C3—C4	1.511 (4)	C7—C8	1.515 (4)
C4—H4A	0.9700	C8—H8A	0.9700
C4—H4B	0.9700	C8—H8B	0.9700

C4—O1—H1A	107 (3)	C8—O2—H2D	109.5
C3—N1—H1B	109.5	C7—N2—H2E	109.5
C3—N1—H1C	109.5	C7—N2—H2F	109.5
H1B—N1—H1C	109.5	H2E—N2—H2F	109.5
C3—N1—H1G	109.5	C7—N2—H2G	109.5
H1B—N1—H1G	109.5	H2E—N2—H2G	109.5
H1C—N1—H1G	109.5	H2F—N2—H2G	109.5
C3—C1—H1D	109.5	C7—C5—H5A	109.5
C3—C1—H1E	109.5	C7—C5—H5B	109.5
H1D—C1—H1E	109.5	H5A—C5—H5B	109.5
C3—C1—H1F	109.5	C7—C5—H5C	109.5
H1D—C1—H1F	109.5	H5A—C5—H5C	109.5
H1E—C1—H1F	109.5	H5B—C5—H5C	109.5
C3—C2—H2A	109.5	C7—C6—H6A	109.5
C3—C2—H2B	109.5	C7—C6—H6B	109.5
H2A—C2—H2B	109.5	H6A—C6—H6B	109.5
C3—C2—H2C	109.5	C7—C6—H6C	109.5
H2A—C2—H2C	109.5	H6A—C6—H6C	109.5
H2B—C2—H2C	109.5	H6B—C6—H6C	109.5
C2—C3—C4	111.4 (2)	N2—C7—C8	107.6 (2)
C2—C3—N1	108.1 (2)	N2—C7—C5	108.1 (2)
C4—C3—N1	107.9 (2)	C8—C7—C5	109.5 (2)
C2—C3—C1	112.0 (3)	N2—C7—C6	107.5 (2)
C4—C3—C1	109.9 (3)	C8—C7—C6	112.0 (2)
N1—C3—C1	107.3 (2)	C5—C7—C6	112.0 (2)
O1—C4—C3	112.5 (2)	O2—C8—C7	110.7 (2)
O1—C4—H4A	109.1	O2—C8—H8A	109.5
C3—C4—H4A	109.1	C7—C8—H8A	109.5
O1—C4—H4B	109.1	O2—C8—H8B	109.5
C3—C4—H4B	109.1	C7—C8—H8B	109.5
H4A—C4—H4B	107.8	H8A—C8—H8B	108.1
C2—C3—C4—O1	−53.1 (3)	N2—C7—C8—O2	−57.7 (3)
N1—C3—C4—O1	65.4 (3)	C5—C7—C8—O2	−174.9 (2)
C1—C3—C4—O1	−177.8 (2)	C6—C7—C8—O2	60.2 (3)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
O1—H1A···Cl1	0.82 (3)	2.25 (4)	3.067 (2)	174 (3)
N1—H1B···O2	0.89	2.02	2.876 (3)	161
N1—H1C···Cl1 ⁱ	0.89	2.33	3.216 (3)	171
N1—H1G···Cl1 ⁱⁱ	0.89	2.42	3.261 (2)	157
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N2—H2E···Cl1	0.89	2.53	3.391 (2)	163
N2—H2E···O2	0.89	2.49	2.811 (3)	102
N2—H2F···Cl2 ^{iv}	0.89	2.24	3.130 (2)	177
N2—H2G···Cl2	0.89	2.27	3.144 (2)	169
C2—H2A···O1	0.96	2.53	2.880 (4)	102

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C6—H6A···O2 0.96 2.59 2.933 (4) 101
Symmetry codes: (i) $x+1, y, z$; (ii) $-x, -y, -z+1$; (iii) $-x, -y+1, -z+1$; (iv) $-x-1/2, y+1/2, -z+3/2$.

Fig. 1

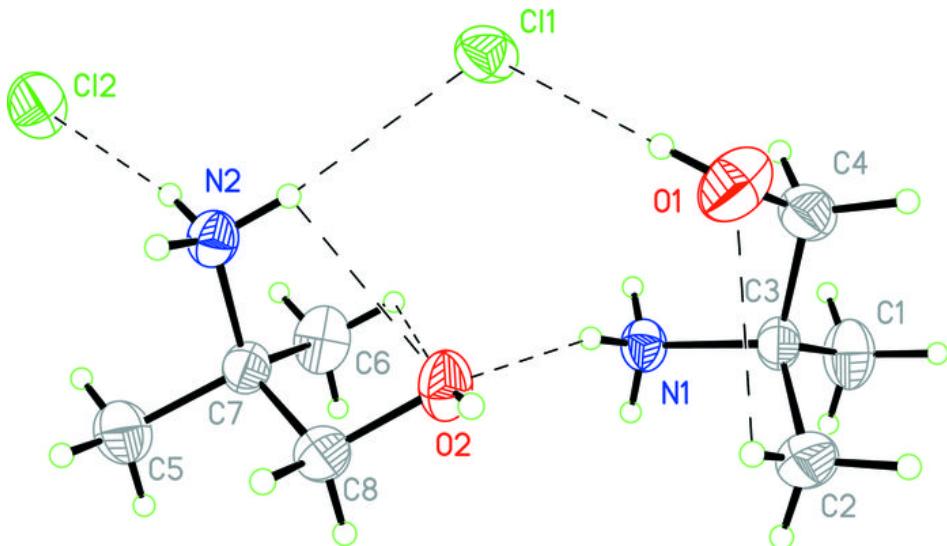


Fig. 2

