

Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

(Z)-N-[2-(Dimethylammonio)ethyl]thioacetamide chloride

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Received 28 November 2007; accepted 1 December 2007

Key indicators: single-crystal X-ray study; T = 292 K; mean σ (C–C) = 0.002 Å; R factor = 0.030; wR factor = 0.089; data-to-parameter ratio = 18.4.

The thioamide and quaternary amine parts of the title compound, $C_6H_{15}N_2S^+ \cdot Cl^-$, are mutually almost perpendicular, the dihedral angle being 80.6 $(7)^{\circ}$. The thioamide group is planar and adopts a Z conformation, whereas the amine end of the cation is in an extended conformation. In the supramolecular structure, molecules are linked into centrosymmetric dimers by two hydrogen bonds: N-H_{amine}····Cl and $N-H_{thioamide} \cdots Cl$.

Related literature

For details of the synthesis, see Spychała (2000, 2003). For bond-length data, see: Allen (2002).



 $\gamma = 87.863 \ (3)^{\circ}$

Z = 2

V = 486.32 (3) Å³

Mo $K\alpha$ radiation $\mu = 0.55 \text{ mm}^{-1}$ T = 292 (2) K $0.6 \times 0.2 \times 0.2$ mm

Experimental

Crystal data

$C_6H_{15}N_2S^+ \cdot Cl^-$
$M_r = 182.71$
Triclinic, P1
a = 5.9935 (2) Å
b = 7.7461 (3) Å
c = 10.8253 (4) Å
$\alpha = 79.489(3)^{\circ}$
$\beta = 79.796 (3)^{\circ}$

4065 measured reflections

 $R_{\rm int} = 0.008$

2332 independent reflections

1988 reflections with $I > 2\sigma(I)$

Data collection

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Kuma KM-4 CCD diffractometer
Absorption correction: multi-scan
  (CrvsAlis RED; Oxford
  Diffraction, 2007)
  T_{\min} = 0.785, T_{\max} = 1.000
  (expected range = 0.704-0.897)
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Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$	H atoms treated by a mixture of
$wR(F^2) = 0.089$	independent and constrained
S = 1.06	refinement
2332 reflections	$\Delta \rho_{\rm max} = 0.36 \text{ e} \text{ Å}^{-3}$
127 parameters	$\Delta \rho_{\rm min} = -0.27 \text{ e } \text{\AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1 \cdots Cl1$	0.883 (9)	2.174 (10)	3.0285 (11)	162.9 (16)
$N4 - H4 \cdots Cl1^{i}$	0.863 (9)	2.325 (10)	3.1801 (12)	171.0 (16)

Symmetry code: (i) -x + 1, -y + 1, -z + 2.

Data collection: CrysAlis CCD (Oxford Diffraction, 2007); cell refinement: CrysAlis RED (Oxford Diffraction, 2007); data reduction: CrysAlis RED; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: Stereochemical Workstation Operation Manual (Siemens 1989) and Mercury (Macrae et al., 2006); software used to prepare material for publication: SHELXL97.

This work was supported by funds from Adam Mickiewicz University, Faculty of Chemistry.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG2404).

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supporting information

Acta Cryst. (2008). E64, o229 [https://doi.org/10.1107/S1600536807064872] (Z)-N-[2-(Dimethylammonio)ethyl]thioacetamide chloride

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S1. Comment

The structure of the title compound (I), is shown below. Dimensions are available in the archived CIF.

The asymmetric unit of the title compound, (I), consists of one $[C_6H_{15}N_2S]^+$ cation and one chloride anion (Fig.1). The thioamide group is in *Z* configuration, its bond lengths are in agreement with the literature (Allen, 2002). The thioamide group is flat, the appropriate torsion angles adopt values close to 0° or 180° [S(5)—C(5)—N(4)—C(3) = τ_1 = -0.5 (2)°, C(51)—O(5)—N(4)—C(3) = τ_2 = -179.6 (1)°]. The quarternary amine end of the molecule is associated with conformational flexibility and it is apparent that the amine chain is in an almost extended conformation [C(12)—N(1)—C(2)—C(3) = τ_3 = 177.0 (1)°] and moreover, it lies in a plane that is nearly orthogonal to the thioamide plane, the dihedral angle between the least-squares planes is 80.6 (7)°. On the other hand, the amine and thioamide functions are mutually *gauche* oriented [N(1)—C(2)—C(3)—N(4) = τ_4 = 62.3 (2)°]. This conformation of the molecule causes some steric stress which is released by thioamide bond angles distortion from ideal values of 120°, *e.g.* the bond angles adopt values as follows: C(5)—N(4)—C(3) is 125.0 (1)°, S(5)—C(5)—N(4) 124.0 (1)°, C(3)—N(4)—H(4) 114.9 (1)°, C(51)—C(5)—N(4) 114.7 (1)°, although the sum of bond angles around C(5) and N(4) equals 360°. This diversity in bond angles seems to be the general property of amide and thioamide groups. The molecular conformation is stabilized by the intermolecular hydrogen bonds N—H…Cl which give rise to centrosymmetric dimers formation (Fig. 1). Each cation in the dimer participates in two hydrogen bonds to two chloride anions, *N*⁺—*H*(amine)…Cl⁻ and *N*—*H*(thioamide)…Cl⁻.

In supramolecular structure the dimers are ordered along the [010] direction one after the another thus forming alternate hydrophilic and hydrophobic segments. In the hydrophilic segments the chloride anions that interact with the cations *via* hydrogen bonds are located, whereas the hydrophobic segments are formed by two ribbons connected by van der Waals forces. Along the third direction, [100], the cations form stacks as the most efficient close packing motifs with the chloride anions forming columns in the channels of the close packing achieved by the cations.

S2. Experimental

Starting from thioacetamide and the appropriate diaminoalkane, the title compound was obtained by the transamination Wallach reaction by refluxing the reaction mixture in ethanol. The reaction was carried out under literature conditions described in the previous papers (Spychała 2000, 2003). Solvent: 2-propanol / diethyl ether. Single crystals were grown from the hot solution by slow cooling.

S3. Refinement

Hydrogen atoms were found from difference Fourier maps and refined except hydrogen atoms of three methyl groups, which were constrained to ride on their parent atom. The N—H distances were restrained to 0.88±0.01 Å.



Figure 1

Dimer of (I) generated by N^+ —H(amine)···Cl⁻ and N—H(thioamide)···Cl⁻ hydrogen bonds (dashed lines). A view along the [100] direction, (Macrae *et al.*, 2006).

(Z)-N-[2-(Dimethylammonio)ethyl]thioacetamide chloride

Crystal data $C_6H_{15}N_2S^+ \cdot CI^ M_r = 182.71$ Triclinic, $P\overline{1}$ a = 5.9935 (2) Å b = 7.7461 (3) Å c = 10.8253 (4) Å a = 79.489 (3)° $\beta = 79.796$ (3)° $\gamma = 87.863$ (3)° V = 486.32 (3) Å³

Z = 2 F(000) = 196 $D_x = 1.248 \text{ Mg m}^{-3}$ Mo K α radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 2792 reflections $\theta = 2.7-29.6^{\circ}$ $\mu = 0.55 \text{ mm}^{-1}$ T = 292 K Block, colourless $0.6 \times 0.2 \times 0.2 \text{ mm}$ Data collection

Kuma KM-4 CCD diffractometer Radiation source: fine-focus sealed tube Graphite monochromator Detector resolution: 8.1929 pixels mm ⁻¹ ω -scan Absorption correction: multi-scan <i>CrysAlis RED</i> (Oxford Diffraction, 2007) $T_{res} = 0.785$, $T_{res} = 1.000$	4065 measured reflections 2332 independent reflections 1988 reflections with $I > 2\sigma(I)$ $R_{int} = 0.008$ $\theta_{max} = 29.7^{\circ}, \theta_{min} = 3.0^{\circ}$ $h = -8 \rightarrow 6$ $k = -10 \rightarrow 10$ $l = -14 \rightarrow 11$
Refinement	
Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.030$ $wR(F^2) = 0.089$ S = 1.06 2332 reflections 127 parameters 0 restraints 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 $w = 1/[\sigma^2(F_o^2) + (0.0493P)^2 + 0.0911P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.36$ e Å ⁻³ $\Lambda \rho_{mix} = -0.27$ e Å ⁻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.

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 $(Å^2)$

	r	12	7	17. */17	
	<i>A</i>	y	2	U _{1SO} / U _{eq}	
Cl1	0.74439 (6)	0.68184 (5)	0.92798 (3)	0.04695 (12)	
N1	0.43953 (18)	0.72292 (15)	0.72918 (10)	0.0369 (2)	
C11	0.6186 (3)	0.7621 (2)	0.61349 (16)	0.0529 (4)	
H11A	0.7143	0.6609	0.6068	0.073 (6)*	
H11B	0.7081	0.8593	0.6201	0.077 (7)*	
H11C	0.5489	0.7915	0.5390	0.070 (6)*	
C12	0.3063 (3)	0.8834 (2)	0.75207 (18)	0.0564 (4)	
H12A	0.4021	0.9661	0.7733	0.066 (6)*	
H12C	0.1822	0.8526	0.8212	0.062 (5)*	
H12B	0.2485	0.9351	0.6764	0.062 (5)*	
C2	0.2817 (2)	0.5793 (2)	0.72500 (14)	0.0401 (3)	
C3	0.3998 (2)	0.4127 (2)	0.69463 (14)	0.0417 (3)	
N4	0.5321 (2)	0.33438 (15)	0.79089 (10)	0.0377 (2)	
C5	0.7308 (2)	0.25637 (17)	0.76729 (13)	0.0368 (3)	
C51	0.8327 (3)	0.1849 (2)	0.88345 (16)	0.0541 (4)	

supporting information

H51A	0.7944	0.0631	0.9114	0.134 (12)*	
H51B	0.7743	0.2491	0.9503	0.148 (13)*	
H51C	0.9945	0.1968	0.8630	0.127 (10)*	
S5	0.86290 (6)	0.23364 (6)	0.62085 (4)	0.05189 (14)	
H1	0.511 (3)	0.691 (2)	0.7946 (13)	0.052 (5)*	
H2A	0.180 (3)	0.564 (2)	0.8055 (18)	0.052 (5)*	
H2B	0.192 (3)	0.622 (2)	0.6625 (17)	0.046 (4)*	
H3A	0.290 (3)	0.329 (2)	0.6874 (17)	0.055 (5)*	
H3B	0.502 (3)	0.432 (2)	0.6142 (17)	0.043 (4)*	
H4	0.468 (3)	0.338 (2)	0.8684 (10)	0.048 (4)*	

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.03632 (18)	0.0692 (3)	0.03689 (19)	-0.00237 (15)	-0.00751 (13)	-0.01194 (16)
N1	0.0340 (5)	0.0447 (6)	0.0327 (5)	0.0040 (4)	-0.0078 (4)	-0.0072 (4)
C11	0.0460 (8)	0.0653 (10)	0.0444 (8)	-0.0118 (7)	0.0021 (6)	-0.0092 (7)
C12	0.0601 (10)	0.0474 (9)	0.0606 (10)	0.0114 (7)	-0.0097 (8)	-0.0103 (7)
C2	0.0302 (6)	0.0497 (8)	0.0408 (7)	0.0019 (5)	-0.0089(5)	-0.0064 (6)
C3	0.0405 (7)	0.0499 (8)	0.0383 (7)	-0.0007 (6)	-0.0128 (6)	-0.0113 (6)
N4	0.0425 (6)	0.0418 (6)	0.0291 (5)	0.0026 (5)	-0.0063 (4)	-0.0079 (4)
C5	0.0394 (6)	0.0379 (7)	0.0347 (6)	-0.0031 (5)	-0.0083 (5)	-0.0080(5)
C51	0.0602 (10)	0.0614 (11)	0.0432 (8)	0.0134 (8)	-0.0188 (7)	-0.0091 (7)
S5	0.0402 (2)	0.0778 (3)	0.0386 (2)	0.00595 (17)	-0.00335 (15)	-0.01740 (18)

Geometric parameters (Å, °)

N1—C12	1.488 (2)	C2—H2B	0.943 (18)
N1-C11	1.4908 (18)	C3—N4	1.4519 (18)
N1C2	1.4988 (18)	С3—НЗА	0.963 (19)
N1—H1	0.883 (9)	С3—Н3В	0.962 (18)
C11—H11A	0.9600	N4—C5	1.3206 (18)
C11—H11B	0.9600	N4—H4	0.863 (9)
C11—H11C	0.9600	C5—C51	1.501 (2)
C12—H12A	0.9600	C5—S5	1.6786 (14)
C12—H12C	0.9600	C51—H51A	0.9600
C12—H12B	0.9600	C51—H51B	0.9600
C2—C3	1.510(2)	C51—H51C	0.9600
C2—H2A	0.960 (19)		
C12—N1—C11	111.02 (13)	N1—C2—H2B	107.7 (10)
C12—N1—C2	109.51 (11)	C3—C2—H2B	109.2 (10)
C11—N1—C2	113.73 (11)	H2A—C2—H2B	106.1 (15)
C12—N1—H1	106.7 (12)	N4—C3—C2	112.06 (12)
C11—N1—H1	106.5 (12)	N4—C3—H3A	109.3 (11)
C2—N1—H1	109.1 (12)	С2—С3—НЗА	109.9 (11)
N1-C11-H11A	109.5	N4—C3—H3B	106.9 (10)
N1-C11-H11B	109.5	С2—С3—Н3В	112.5 (10)

H11A—C11—H11B	109.5	H3A—C3—H3B	105.9 (15)
N1—C11—H11C	109.5	C5—N4—C3	124.90 (12)
H11A—C11—H11C	109.5	C5—N4—H4	120.4 (12)
H11B—C11—H11C	109.5	C3—N4—H4	114.6 (12)
N1—C12—H12A	109.5	N4—C5—C51	114.66 (12)
N1—C12—H12C	109.5	N4—C5—S5	124.05 (10)
H12A—C12—H12C	109.5	C51—C5—S5	121.28 (11)
N1—C12—H12B	109.5	C5—C51—H51A	109.5
H12A—C12—H12B	109.5	C5—C51—H51B	109.5
H12C—C12—H12B	109.5	H51A—C51—H51B	109.5
N1—C2—H2A C3—C2—H2A	114.08 (11) 105.4 (11) 113.9 (11)	H51A-C51-H51C H51B-C51-H51C	109.5 109.5 109.5
C12—N1—C2—C3	176.67 (13)	C3—N4—C5—C51	-179.64 (14)
C11—N1—C2—C3	51.81 (16)	C3—N4—C5—S5	-0.5 (2)
N1—C2—C3—N4	62.34 (16)	S5—C5—N4—H4	177.1 (14)
C2—C3—N4—C5	-142.40 (13)	C51—C5—N4—H4	-2.1 (14)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H…A	D····A	<i>D</i> —H··· <i>A</i>
N1—H1…Cl1	0.88 (1)	2.17(1)	3.0285 (11)	163 (2)
N4—H4…Cl1 ⁱ	0.86(1)	2.33 (1)	3.1801 (12)	171 (2)

Symmetry code: (i) -x+1, -y+1, -z+2.