

2,2'-Diamino-4,4'-bi-1,3-thiazolium dichloride

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Key indicators

Single-crystal X-ray study
T = 298 K
Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$
R factor = 0.058
wR factor = 0.180
Data-to-parameter ratio = 13.6

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $\text{C}_6\text{H}_8\text{N}_4\text{S}_2^{2+} \cdot 2\text{Cl}^-$, includes a *trans*, planar bithiazole moiety occupying a special position on a crystallographic inversion centre. Both symmetry-independent N atoms participate in hydrogen-bonded links with the chloride anion [$\text{N} \cdots \text{Cl}$ 3.094 (5), 3.222 (5) and 3.362 (5) Å]. A short contact of 3.3201 (18) Å is also observed between the S atom of the thiazolium cation and the chloride anion.

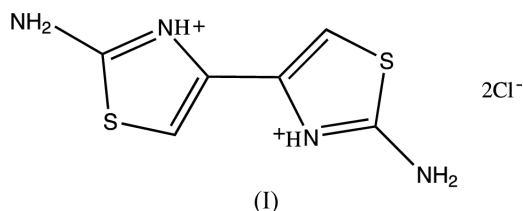
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Comment

Transition-metal complexes with 2,2'-diamino-4,4'-bithiazole (DABT) or its derivatives have shown interesting properties and promise potential applications in a number of fields (Waring, 1981; Fisher *et al.*, 1985; Sun *et al.*, 1997). A series of metal complexes, with DABT chelating to the metal atom, have recently been prepared in this laboratory and crystal structure determinations have been reported for several of them (Liu *et al.*, 2001). We present here the structure of DABT chloride, (I), thereby allowing comparison of the DABT structures in metal complexes and in the free state.



The molecular structure of (I) and a fragment of the crystal packing are shown in Fig. 1. The centrosymmetric molecule displays a *trans* structure, which differs from the *cis* configuration found in metal complexes. The two thiazole rings are coplanar, the maximum atomic deviation being 0.016 (3) Å (N3). The structure is also different from the twisted structure found in the metal complexes (Liu *et al.*, 2001).

The N—H groups of the heterocyclic protonated atom N3, as well as the N6 amino atom, are engaged in hydrogen bonds with the chloride anion (Table 1). The rather short $\text{Cl} \cdots \text{S1}^{\text{ii}}$ [symmetry code: (ii) $1 + x, \frac{1}{2} - y, -\frac{1}{2} + z$] contact of 3.3201 (18) Å is also noteworthy. This contact is significantly shorter than the sum of the van der Waals radii (3.55 Å ; Rodgers, 1994); it is comparable, however, with the $\text{S} \cdots \text{Cl}$ contact of 3.3175 (15) Å found in bis(2-amino-4-methylthiazolium) tetrachlorocopper (Fernández *et al.*, 1996).

Experimental

DABT was prepared according to the literature procedure of Erlenmeyer (1948). The title compound was obtained from an aqueous solution as a by-product in the course of the synthesis of an Ni^{II} complex with DABT, using NiCl_2 as a starting reagent.

Crystal data

$C_6H_8N_4S_2^{2+} \cdot 2Cl^-$
 $M_r = 271.18$
 Monoclinic, $P2_1/c$
 $a = 4.8211$ (8) Å
 $b = 9.2441$ (14) Å
 $c = 12.0228$ (10) Å
 $\beta = 98.224$ (12)°
 $V = 530.31$ (13) Å³
 $Z = 2$

$D_x = 1.698$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 22 reflections
 $\theta = 7.6$ – 12.6 °
 $\mu = 0.97$ mm⁻¹
 $T = 298$ (2) K
 Prism, orange
 $0.50 \times 0.26 \times 0.18$ mm

Data collection

Rigaku AFC-7S diffractometer
 $\omega/2\theta$ scans
 Absorption correction: ψ scan
 (North *et al.*, 1968)
 $T_{min} = 0.616$, $T_{max} = 0.840$
 1163 measured reflections
 1037 independent reflections
 886 reflections with $I > 2\sigma(I)$

$R_{int} = 0.049$
 $\theta_{max} = 26.0$ °
 $h = 0 \rightarrow 5$
 $k = 0 \rightarrow 11$
 $l = -14 \rightarrow 14$
 3 standard reflections
 every 150 reflections
 intensity decay: 0.3%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.180$
 $S = 1.12$
 1037 reflections
 76 parameters
 Only coordinates of H atoms
 refined

$w = 1/[\sigma^2(F_o^2) + (0.0944P)^2 + 1.6P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.86$ e Å⁻³
 $\Delta\rho_{min} = -0.83$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

S1—C2	1.718 (5)	N3—C4	1.393 (6)
S1—C5	1.733 (5)	C4—C5	1.347 (7)
C2—N3	1.337 (6)	C4—C4 ⁱ	1.446 (10)
C2—N6	1.317 (7)		
C2—S1—C5	90.1 (2)	C5—C4—N3	112.3 (5)
N6—C2—N3	122.5 (5)	C5—C4—C4 ⁱ	128.7 (6)
N6—C2—S1	125.5 (4)	N3—C4—C4 ⁱ	119.1 (5)
N3—C2—S1	111.9 (4)	C4—C5—S1	111.9 (4)
C2—N3—C4	113.8 (4)		

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

Table 2

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N3—H3 \cdots Cl	1.03 (8)	2.10 (8)	3.094 (5)	159 (6)
N6—H6A \cdots Cl ⁱ	0.94 (8)	2.40 (8)	3.222 (5)	146 (7)
N6—H6B \cdots Cl	0.97 (8)	2.50 (8)	3.362 (5)	148 (7)

Symmetry code: (i) $1 - x, \frac{1}{2} + y, \frac{1}{2} - z$.

H atoms were located in a difference Fourier map and refined isotropically with a fixed displacement parameter of 0.08 Å², N—H = 0.95–1.03 Å and C5—H5 = 1.10 (8) Å.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1992); cell refinement: *MSC/AFC*

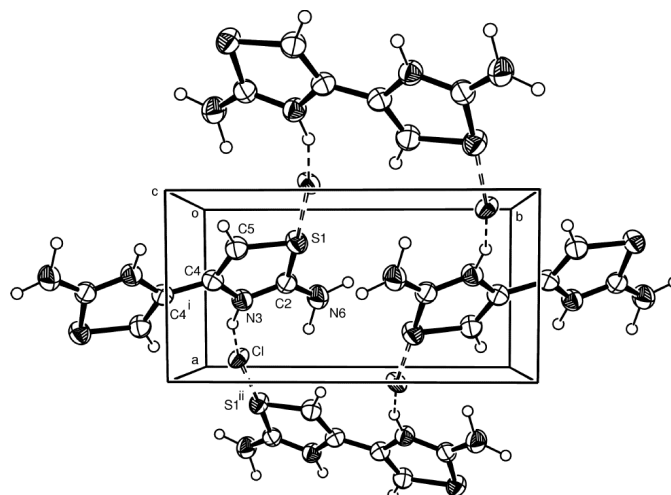


Figure 1

The molecular packing of (I) in the unit cell, shown with 30% probability displacement ellipsoids for non-H atoms. The dashed lines show the hydrogen bonds while the double dashed lines show the short contact between Cl and S atoms. [Symmetry codes: (i) $1 - x, -y, 1 - z$; (ii) $1 + x, \frac{1}{2} - y, -\frac{1}{2} + z$.]

Diffractometer Control Software; data reduction: *TEXSAN* (Molecular Structure Corporation, 1985); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994).

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