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#### Key indicators

Single-crystal X-ray study T = 293 KMean  $\sigma(N-C) = 0.004 \text{ Å}$  R factor = 0.052 wR factor = 0.090 Data-to-parameter ratio = 19.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. 1-Amino-3-ammonioguanidinium dichloride

The structure of the title compound,  $CH_9N_5^{2+}\cdot 2Cl^-$ , has been determined. Strong N-H···Cl bonds are present, leading to a complex pattern of hydrogen-bonded rings.

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## Comment

It has been shown recently that guanidinium ions and various sulfonates assemble into crystalline host frameworks with exceptionally reproducible lamellar architectures (Holman et al., 2001). Interactions between ammonium cations and sulfonates lead to interesting crystalline inclusion architectures based upon second-sphere interactions (Reddy et al., 2003). Diaminoguanidinium sulfonate compounds could combine these interactions and could be used as interesting tectons in porous crystalline networks. Published structures of diaminoguanidinium salts are rare; the Cambridge Structural Database (Version 5.24, update 2; Allen 2002) contains only three (Cromer et al., 1988; Ritchie et al., 1990; Savel'eva et al., 1995). We attempted to synthesize diaminoguanidinium trifluoromethanesulfonate from diaminoguanidinium chloride and trifuoromethanesulfonic acid. The structure solution of the crystals present in the batch revealed a protonated diaminoguanidinium chloride salt, formally described as 1amino-3-ammonioguanidinium dichloride, (I).



The molecular structure of (I) is presented in Fig. 1. The asymmetric unit contains one protonated 1,3-diaminoguanidinium ion and two chloride counter-ions. Table 1 gives selected geometric parameters of the molecular structure. Hydrogen bonds are abundant in this structure, as could be expected from the chemical formula and the liability of amino groups to act as donor: ten different N-H···Cl bonds are found, with  $H \cdot \cdot \cdot Cl$  distances ranging from 2.14 to 2.75 Å. The two shortest, N8-H16···Cl2 (2.14 Å) and N7-H14···Cl1 (2.15 Å), have, in addition, nearly linear contact angles (167 and 165°, respectively). According to Brammer et al. (2001), the latter two bonds are among the strongest found for N-H···Cl contacts. Table 2 gives details of the hydrogen-bond geometry. Fig. 2 shows a projection of the structure on the bc plane; it gives an idea of the complex hydrogen-bond network formed. Among the more extended patterns formed are  $R_3^5(16)$  and two different  $R_2^2(8)$  rings (Bernstein *et al.*, 1995).

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One ring involves two amino groups of symmetry-related guanidinium ions, whereas the second involves two ammonium groups.

## Experimental

Single crystals of the title complex were obtained by slow evaporation of a methanol solution of diaminoguanidinium chloride (500 mg, 4 mmol) and trifluoromethanesulfonic acid (1800 mg, 12 mmol).

> $D_x = 1.627 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation Cell parameters from 21 reflections  $\theta = 10-12^\circ$

 $\mu = 0.89 \text{ mm}^{-1}$ 

Prism, colourless  $0.40 \times 0.10 \times 0.10$  mm

T = 293 K

 $\begin{array}{l} R_{\rm int} = 0.030 \\ \theta_{\rm max} = 30.0^{\circ} \\ h = -6 \rightarrow 6 \end{array}$ 

 $k = -29 \rightarrow 25$ 

2 standard reflections

frequency: 100 min

intensity decay: 6.0%

 $l = 0 \rightarrow 9$ 

#### Crystal data

$CH_9N_5^{2+}\cdot 2Cl^-$
$M_r = 162.02$
Monoclinic, $P2_1/c$
a = 4.503 (2)  Å
<i>b</i> = 21.234 (8) Å
c = 6.936 (2)  Å
$\beta = 94.36 \ (3)^{\circ}$
$V = 661.3 (4) \text{ Å}^3$
Z = 4

#### Data collection

Siemens P3 four-circle diffractometer  $\omega$  scans Absorption correction:  $\psi$  scan (PLATON; Spek, 2003) $T_{min} = 0.874, T_{max} = 0.912$ 3211 measured reflections 1840 independent reflections 1404 reflections with  $I > 2\sigma(I)$ 

#### Refinement

Refinement on  $F^2$ H-atom parameters constrained $R[F^2 > 2\sigma(F^2)] = 0.052$ Weighting scheme: see text $wR(F^2) = 0.090$  $(\Delta/\sigma)_{max} < 0.001$ S = 1.00 $\Delta \rho_{max} = 0.70 \text{ e Å}^{-3}$ 1404 reflections $\Delta \rho_{min} = -0.39 \text{ e Å}^{-3}$ 73 parameters $\Delta \rho_{min} = -0.39 \text{ e Å}^{-3}$ 

### Table 1

Selected geometric parameters (Å, °).

N3-C6	1.313 (4)	C6-N7	1.351 (4)
N4-N5	1.414 (4)	N7-N8	1.412 (4)
N5-C6	1.299 (4)		
C6-N5-N4	118.1 (3)	N5-C6-N3	122.2 (3)
N7-C6-N5	117.3 (3)	N8-N7-C6	117.1 (3)
N7-C6-N3	120.5 (3)		. ,

Та	ble	2
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Hydrogen-bonding geometry (Å, °).

$\overline{D-\mathrm{H}\cdots A}$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
N4-H9···Cl1 <sup>i</sup>	0.88	2.51	3.318 (3)	152
N4-H10···Cl1 <sup>ii</sup>	1.02	2.38	3.395 (3)	174
N5-H11···Cl1 <sup>iii</sup>	0.91	2.43	3.223 (3)	146
$N3-H12\cdots Cl2^{iv}$	0.86	2.59	3.223 (3)	131
$N3-H12\cdots N4$	0.86	2.21	2.655 (4)	112
$N3-H13\cdots Cl2^{v}$	0.86	2.36	3.196 (3)	164
$N7-H14\cdots Cl1^{vi}$	0.93	2.15	3.052 (3)	165
$N8-H15\cdots Cl1$	0.82	2.39	3.101 (3)	146
N8-H15···Cl2 <sup>vii</sup>	0.82	2.75	3.097 (3)	107
$N8-H16\cdots Cl2^{v}$	0.99	2.14	3.110 (3)	167
$N8-H17\cdots Cl2^{viii}$	0.86	2.28	3.096 (3)	160

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



#### Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level.



# Figure 2

Projection of the crystal structure of (I) on the bc plane.

A Chebychev polynomial (Watkin, 1994; Prince, 1982) was used in the weighting scheme, [weight] =  $1.0/[A_0T_0(x) + A_1T_1(x)... + A_{n-1}T_{n-1}(x)]$ , where  $A_i$  are the Chebychev coefficients listed below and  $x = F_{calc}/F_{max}$ . The robust weighting method (Prince, 1982) was used, with w = weight  $\times [1 - (\Delta F/6\sigma F)^2]^2$ .  $A_{0-2}$  are 10.8, 13.7, and 4.14, respectively. H atoms were found in Fourier difference maps and were allowed to ride on their parent N atoms. H atoms were located in Fourier difference maps and allowed to ride on their parent N atoms, with  $U_{iso}$  values fixed at 0.05 Å<sup>2</sup>.

Data collection: P3 (Siemens, 1993); cell refinement: P3; data reduction: XDISK in P3; program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994); program(s) used to refine structure: CRYS-TALS (Watkin *et al.*, 2001); molecular graphics: Mercury (CCDC, 2002) and PLATON (Spek, 2003); software used to prepare material for publication: CRYSTALS.

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