C1C9C10	112.1 (2)	C17-C25-C26	112.6 (3)
01-C10-C9	121.9 (4)	O2-C26-C25	121.5 (3)
01-C10-C11	119.1 (2)	O2-C26-C27	120.3 (3)
C9-C10-C11	119.0 (3)	C25-C26-C27	118.2 (3)
C10C11C12	122.1 (2)	C26-C27-C28	124.0 (3)
C10C11C16	117.5 (3)	C26-C27-C32	116.9 (4)
C12-C11-C16	120.4 (3)	C28-C27-C32	119.1 (3)
C11—C12—C13	119.3 (2)	C27-C28-C29	120.4 (4)
C12-C13-C14	120.2 (3)	C28-C29-C30	120.2 (3)
C13-C14-C15	120.5 (4)	C29-C30-C31	119.8 (4)
C14-C15-C16	120.1 (3)	C30-C31-C32	120.6 (4)
C11-C16-C15	119.5 (3)	C27—C32—C31	119.9 (3)

The structure was solved by direct methods and difference Fourier syntheses. Refinement was by full-matrix least-squares methods with anisotropic displacement parameters for all non-H atoms and isotropic displacement parameters for H atoms.

All calculations were performed on a PDP 11/44 computer with the SDP (Frenz, 1978) programs.

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: CR1131). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# 2,6-Dichlorophenylguanidinium Chloride

#### JAMES BARKER

The Associated Octel Company Ltd, PO Box 17, R & D Laboratories, Oil Sites Road, Ellesmere Port, South Wirral L65 4HF, England

HAROLD R. POWELL

Cambridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, England

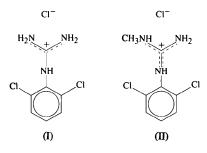
(Received 11 November 1994; accepted 13 February 1995)

## Abstract

The C—N bonds in  $C_7H_8Cl_2N_3^+.Cl^-$  were found to differ in length [1.304 (4), 1.308 (4) and 1.493 (4) Å]. There is not, therefore, the symmetrical delocalization around the central N<sub>3</sub>C expected for a guanidinium system, and the molecule is closer in nature to a substituted amidinium salt.

### Comment

Structural studies of guanidines and their salts continue to attract attention (Carpy, Leger, Wermuth & Leclerc, 1981; Hafelinger & Kuske, 1991) because of their biological and pharmaceutical, *e.g.* neuroleptic and antihypersensitive, activity (Rouot, Leclerc, Wermuth, Miesch & Schwarz, 1978; Bream, Lauener, Picard, Scholtysik & White, 1975). In the title compound, (I), the asymmetric unit contains one molecule, resolved as the *E* isomer (Fig. 1).



Two of the three C—N bond lengths of the N<sub>3</sub>C fragment are similar [N(14)—C(12) 1.304 (4) and N(13)— C(12) 1.308 (4) Å] and are only slightly longer than the mean of 277  $C_{sp^2}$ —N double bonds [1.295 Å] located in the Cambridge Structural Database (Allen *et al.*, 1987), indicating delocalization around the N—C—N fragment. The third C—N bond length [C(12)—N(11) 1.493 (4) Å] and the C(aryl)—N distance [C(1)—N(11) 1.519 (4) Å] are clearly single bond in character, and thus the delocalization is not across the whole of the N<sub>3</sub>C fragment, nor does it extend into the aromatic ring. The C(1)—N(11)—C(12) bond angle of  $115.6(3)^{\circ}$ . which is midway between that expected for an  $sp^2$ and for an  $sp^3$  N atom, and the C(12)-N(11)-C(1)-C(2) torsion angle of 82.3 (4)°, support this point of view. The structure has, therefore, more in common with a substituted amidinium salt than the closely related 2,6-dichlorophenyl-2-methyl-1-guanidinium chloride hydrate, (II) (Carpy, Leger, Wermuth & Leclerc, 1981), in which the C-N bond lengths of 1.312(6), 1.349 (6) and 1.388 (6) Å indicate delocalization around the N<sub>3</sub>C fragment, as expected for a guanidinium salt. The only chemical difference between the two compounds is the methyl substituent in (II). In the title compound, two of the N-C-N angles are similar: N(13)-C(12)-N(14) 120.8 (3) and N(13)-C(12)-N(11) 122.1 (3)°. The third, N(11)—C(12)—N(14), is smaller, at 117.1 (3)°, which is similar to that found for 2.6-dichlorophenyl-2-methyl-1-guanidinium chloride hydrate (Carpy, Leger, Wermuth & Leclerc, 1981).

There are weak intermolecular hydrogen bonds present extending through the cell, as indicated by the close Cl···N contacts [Cl(1)···N(13) 3.210 (3), Cl(1)···N(13)(1 -  $x, y - \frac{1}{2}, -\frac{1}{2} - z$ ) 3.269 (2), Cl(1)··· N(14)( $x - \frac{1}{2}, \frac{1}{2} - y, -z - 1$ ) 3.260 (3) and Cl(1)··· N(14)(1 -  $x, y - \frac{1}{2}, -\frac{1}{2} - z$ ) 3.209 (3) Å].

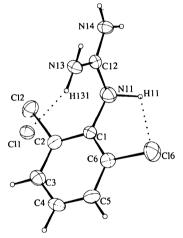


Fig. 1. Molecular structure and non-H atomic labels of (I). Displacement ellipsoids are shown at 50% probability levels. H atoms are drawn as small circles of arbitrary radii.

#### Experimental

## Crystal data

 $C_7H_8Cl_2N_3^+.Cl^ M_r = 240.52$ Orthorhombic  $P2_12_12_1$  a = 10.3302 (3) Å b = 10.6380 (2) Å c = 9.7044 (2) Å Cu  $K\alpha$  radiation  $\lambda = 1.5418$  Å Cell parameters from 16 reflections  $\theta = 28.74-34.80^{\circ}$   $\mu = 7.644$  mm<sup>-1</sup> T = 293 K

$$V = 1066.44 \text{ A}^{3}$$
  
 $Z = 4$   
 $D_{x} = 1.50 \text{ Mg m}^{-3}$ 

0 2

Data collection Rigaku AFC-7R diffractometer  $\theta/2\theta$  scans Absorption correction:  $\psi$  scans (North, Phillips & Mathews, 1968)  $T_{min} = 0.403, T_{max} =$  1.0001800 measured reflections 1744 independent reflections 1687 observed reflections  $[F > 3\sigma(F)]$ 

#### Refinement

Refinement on F Extinction correction: R = 0.0405Larson (1970) wR = 0.0429Extinction coefficient: S = 1.0991205.4 (69) 1687 reflections Atomic scattering factors 145 parameters from International Tables for X-ray Crystallography H atoms riding on attached C or N (1974, Vol. IV, Table Weighting scheme: 2.2B) Chebyshev Absolute configuration:  $(\Delta/\sigma)_{\rm max} = 0.056$ Flack (1983) enantiopole  $\Delta \rho_{\rm max} = 0.58 \ {\rm e} \ {\rm \AA}^{-3}$ parameter refined to  $\Delta \rho_{\rm min} = -0.51 \ {\rm e} \ {\rm \AA}^{-3}$ 1.02(3) for the opposite hand of the solution, and 0.02(3) for the correct

## Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

hand

 $U_{\text{eq}} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_i^* \mathbf{a}_i \cdot \mathbf{a}_j.$ 

	x	у	Z	$U_{eq}$
Cl(1)	0.40364 (8)	0.07692 (7)	-0.23293 (8)	0.0389 (4)
Cl(2)	0.41486 (9)	0.15515 (9)	-0.7043 (1)	0.0520 (5)
Cl(6)	0.8276 (1)	0.0082(1)	-0.4028 (1)	0.0665 (7)
N(11)	0.6806 (3)	0.1988 (3)	-0.5850 (4)	0.0534 (2)
N(13)	0.5681 (3)	0.2910 (2)	-0.3868 (3)	0.0386(1)
N(14)	0.6745 (3)	0.4188 (3)	-0.5361 (3)	0.0406(1)
C(1)	0.6171 (3)	0.0734 (3)	-0.5533 (3)	0.0334(1)
C(2)	0.4960 (3)	0.0432 (3)	-0.6065 (3)	0.0354 (1)
C(3)	0.4373 (4)	-0.0726 (3)	-0.5878 (4)	0.0419 (2)
C(4)	0.5005 (4)	-0.1605 (3)	-0.5081 (4)	0.0438 (2)
C(5)	0.6198 (4)	-0.1360 (3)	-0.4531 (4)	0.0458 (2)
C(6)	0.6762 (4)	-0.0191 (3)	-0.4770 (4)	0.0419 (2)
C(12)	0.6377 (3)	0.3068 (3)	-0.4980 (3)	0.0327 (1)

#### Table 2. Selected geometric parameters (Å, °)

Cl(2)—C(2)	1.738 (3)	C(1)—C(2)	1.391 (5)
Cl(6)—C(6)	1.746 (4)	C(1)—C(6)	1.374 (5)
N(11)C(1)	1.519 (4)	C(2)—C(3)	1.384 (5)
N(11)—C(12)	1.493 (4)	C(3)—C(4)	1.378 (5)
N(13)—C(12)	1.308 (4)	C(4)—C(5)	1.369 (5)
N(14)—C(12)	1.304 (4)	C(5)—C(6)	1.392 (5)
C(1)—N(11)—C(12)	115.6 (3)	C(3)—C(4)—C(5)	121.1 (3)
N(11) - C(1) - C(2)	121.1 (3)	C(4)—C(5)—C(6)	118.8 (3)

Irregular block  $0.40 \times 0.25 \times 0.22$  mm Colourless Crystal source: Associated Octel Company Ltd

 $R_{int} = 0.0411$   $\theta_{max} = 75^{\circ}$   $h = -1 \rightarrow 12$   $k = -1 \rightarrow 13$   $l = -1 \rightarrow 12$ 3 standard reflections monitored every 100 reflections intensity decay: none

N(11)-C(1)-C(6)	123.1 (3)	Cl(6) - C(6) - C(1)	120.1 (3)
C(2)—C(1)—C(6)	115.7 (3)	Cl(6)—C(6)—C(5)	117.0 (3)
Cl(2)—C(2)—C(1)	118.6 (2)	C(1)-C(6)-C(5)	122.9 (3)
Cl(2)C(2)C(3)	118.0 (3)	N(11)-C(12)-N(13)	122.1 (3)
C(1)—C(2)—C(3)	123.4 (3)	N(11)-C(12)-N(14)	117.1 (3)
C(2)C(3)-C(4)	118.0 (3)	N(13)C(12)N(14)	120.8 (3)

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1993). Cell refinement: MSC/AFC Diffractometer Control Software. Data reduction: MSC/AFC Diffractometer Control Software. Program(s) used to solve structure: SIR92 (Altomare, Cascarano, Giacovazzo & Guagliardi, 1993). Program(s) used to refine structure: CRYS-TALS (Watkin, Carruthers & Betteridge, 1985). Molecular graphics: CAMERON (Pearce & Watkin, 1993). Software used to prepare material for publication: CRYSTALS.

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: HA1140). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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