O3—P1—C6	110.9 (2)	O3-C12-C13	106.9 (4)
04P1C6	107.1 (2)	O4-C14-C15	106.8 (3)
05-P2-C1	114.6 (2)	P2-C16-C17	125.5 (3)
O5-P2-C16	111.2 (2)	P2-C16-C21	114.5 (3)
O5-P2-C22	112.0 (2)	C17-C16-C21	119.8 (4)
C1-P2-C16	106.3 (2)	C16-C17-C18	119.7 (4)
C1-P2-C22	108.4 (2)	C17C18C19	120.1 (4)
C16—P2—C22	103.6 (2)	C18-C19-C20	120.6 (4)
06-P3-C1	114.0 (2)	C19-C20-C21	119.5 (4)
O6-P3-C28	112.0 (2)	C16-C21-C20	120.2 (4)
O6-P3-C34	110.9 (2)	P2-C22-C23	121.7 (3)
C1-P3-C28	107.7 (2)	P2-C22-C27	118.9 (3)
C1-P3-C34	104.4 (2)	C23—C22—C27	119.4 (3)
C28-P3-C34	107.4 (2)	C22-C23-C24	120.3 (4)
P1	123.8 (3)	C23-C24-C25	118.7 (5)
P1-04-C14	120.5 (3)	C24—C25—C26	121.4 (4)
P2-C1-P3	112.4 (2)	C25C26C27	120.2 (4)
P2-C1-C2	111.2 (2)	C22-C27-C26	120.0 (4)
P3-C1-C2	109.3 (2)	P3-C28-C29	123.1 (3)
C1C2C3	120.1 (3)	P3-C28-C33	117.6 (3)
C1C2C7	121.3 (3)	C29-C28-C33	119.4 (4)
C3—C2—C7	118.3 (3)	C28—C29—C30	119.6 (4)
C2-C3-C4	124.0 (3)	C29-C30-C31	120.0 (5)
C3-C4-C5	116.7 (3)	C30-C31-C32	119.9 (5)
C3-C4-C8	122.9 (3)	C31—C32—C33	121.1 (4)
C5-C4-C8	120.4 (3)	C28—C33—C32	120.0 (4)
O1-C5-C4	118.0(3)	P3-C34-C35	116.6 (3)
01-C5-C6	121.9 (3)	P3—C34—C39	123.9 (3)
C4-C5-C6	120.1 (3)	C35-C34-C39	119.6 (4)
P1-C6-C5	120.8 (3)	C34—C35—C36	121.1 (4)
P1C6C7	118.6 (3)	C35—C36—C37	118.5 (4)
C5-C6-C7	120.6 (3)	C36—C37—C38	120.6 (4)
C2-C7C6	120.2 (3)	C37—C38—C39	120.8 (4)
C4—C8—C9	109.3 (3)	C34—C39—C38	119.4 (4)
C4-C8-C10	110.3 (3)		

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: PROCESS in MolEN (Fair, 1990). Program(s) used to solve structure: MULTAN11/82 (Main et al., 1982). Program(s) used to refine structure: LSFM in MolEN. Software used to prepare material for publication: PLATON (Spek, 1990).

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates, complete geometry and torsion angles have been deposited with the IUCr (Reference: KA1188). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# Isopropylamidinium Chloride

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### Abstract

The amidine fragment in each of the two independent molecules of  $C_4H_{11}N_2^+$ .  $Cl^-$  is essentially planar; the bond angles around the central C atom sum to 360.0(4)° in each case. This, and the almost equal C-N bond lengths [1.299 (6) to 1.320 (5) Å], indicate that an amidinium salt has been formed.

## Comment

A number of papers concerning the structures of amidines have appeared (Tykarska, Jaskolski & Kosturkiewicz, 1986; Thailambal, Pattabhi & Guru Row, 1986; Barker, Gould & Kilner, 1987; Barker & Kilner, 1993; Alcock, Barker & Kilner, 1988; Alcock, Barker, Blacker, Errington & Wallbridge, 1994; Dehnicke, 1990), reflecting the widespread interest in such compounds because of their pharmaceutical and biological importance, bonding characteristics and ligand properties. The structural investigation of the title compound, (I), was undertaken to investigate the effect of alkyl substitution on the central amidine fragment.



The asymmetric unit consists of two molecules with N-H. Cl hydrogen bonding present (Fig. 1), as it is throughout the structure (Fig. 2). The skeletal C-N bond lengths are clearly not differentiated in terms of single- or double-bond character, with mean C-N bond lengths of 1.317(5) and 1.300(6) Å for each molecule, respectively. They are similar to those found for acetamidinium chloride [1.305(2) and 1.310(3) Å; Cannon, White & Willis, 1976] and bis(acetamidinium)carbonate monohydrate [1.308(5) and 1.315 Å; Norrestam, 1984]. They differ somewhat to those found for acetamidine [1.298(1) and 1.344(1) Å; Norrestam, Mertz & Crossland, 1983], N,N-diphenylbenzamidine [1.302(7) and 1.360(8) Å; Alcock et al., 1988] and benzamidine hydrochloride monohydrate [1.293 (7) and 1.328 (7) Å; Thailambal et al., 1986]. The almost equal C-N bond lengths in the title compound and the planarity of the CCN<sub>2</sub> fragment are indicative of an amidinium salt configuration. Further, the mean C-N distance of 1.309 (6) Å is, as expected, longer than that found in [CH<sub>3</sub>ClCNH<sub>2</sub>]<sup>+</sup>Cl<sup>-</sup> (1.265 Å) which contains only one C-NH<sub>2</sub> linkage. Closer comparison of the C-N bond lengths in benzamidine hydrochloride (Thailambal et al., 1986) and isopropylamidinium chloride indicate the aryl substituent to be more restrictive toward delocalization than the alkyl group. The N-C-N angle [mean 120.4(4)°] is similar to that of acetamidinium chloride [120.5 (2)°; Cannon et al., 1976], but deviates slightly from those found for bisacetamidinium carbonate monohydrate [121.6 (3)°; Norrestam, 1984] and benzamidine hydrochloride hydrate [121.6 (4)°; Thailambal et al., 1987], and deviates considerably from that found for acetamidine  $[125.5(1)^{\circ}]$ ; Norrestram et al., 1983]. This indicates the amidinium



Fig. 1. Molecular structure and non-H atomic labels of isopropylamidinium chloride. Displacement ellipsoids are shown at 50% probability levels; H atoms are drawn as small circles of arbitrary radius.



Fig. 2. Projection down the c axis.



to chloride interaction to be stronger in acetamidinium chloride and isopropylamidinium chloride than in benzamidine hydrochloride. Again this may be due to the influence of the aryl substituent. Further, the amidinium oxygen interaction is not as strong as the amidinium chloride interaction. The delocalization around the N— C—N fragment is also responsible for a short C—CNN bond [mean 1.491 (6) Å]. There are a number of close Cl<sup>-</sup> ion to amino-H atom contacts as evidenced by short N···Cl distances (3.15–3.26 Å), indicating strong interactions. Consideration of the relationship of the methyl groups to the planar N—C—N fragment shows that they adopt an *anti* position to it (Fig. 3).

# Experimental

The title compound was obtained from the Associated Octel Company Ltd.

Crystal data

 $C_4H_{11}N_2^+.Cl^ M_r = 122.60$ Monoclinic  $P2_1/n$  a = 10.6713 (7) Å b = 6.6936 (6) Å c = 21.0352 (7) Å  $\beta = 100.776 (4)^\circ$   $V = 1476.04 Å^3$  Z = 8  $D_x = 1.103 \text{ Mg m}^{-3}$  $D_m$  not measured

Data collection

Rigaku AFC-7*R* diffractometer  $\theta/2\theta$  scans Absorption correction: empirical,  $\psi$  scan (Molecular Structure Corporation, 1985, 1992)  $T_{min} = 0.92$ ,  $T_{max} = 1.00$ 2435 measured reflections 1521 independent reflections

#### Refinement

Refinement on *F* R = 0.0368wR = 0.0443S = 4.976 Cu  $K\alpha$  radiation  $\lambda = 1.5418$  Å Cell parameters from 24 reflections  $\theta = 37.1-37.8^{\circ}$   $\mu = 3.85 \text{ mm}^{-1}$  T = 153 KPrism  $0.30 \times 0.24 \times 0.20 \text{ mm}$ Colourless

1366 observed reflections  $[F > 2.5\sigma(F)]$   $R_{int} = 0.0196$   $\theta_{max} = 50.78^{\circ}$   $h = -10 \rightarrow 10$   $k = -1 \rightarrow 6$   $l = -1 \rightarrow 20$ 3 standard reflections monitored every 100 reflections intensity decay: -0.015%

 $\Delta \rho_{\text{max}} = 0.488 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\text{min}} = -0.325 \text{ e } \text{\AA}^{-3}$ Extinction correction: Larson (1970)

1366 reflections	Extinction coefficient:
216 parameters	32.7 (22)
H atoms riding on attached	Atomic scattering factors
C or N	from International Tables
$w = 1/[\sigma^2(F) + 0.0004F^2]$	for X-ray Crystallography
$(\Delta/\sigma)_{\rm max} = 0.0181$	(1974, Vol. IV)

# Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters $(Å^2)$

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

	x	у	z	$U_{eq}$
<b>C</b> I(1)	-0.11100 (9)	0.7296 (2)	0.46747 (5)	0.0349 (6)
C1(2)	-0.52314(9)	0.5846 (2)	0.27248 (5)	0.0374 (6)
NÌÌÌ	0.0557 (4)	0.8273 (6)	0.3595 (2)	0.0455 (2)
N(12)	0.2237 (4)	0.7725 (6)	0.3083 (2)	0.0429 (2)
N(21)	-0.2828 (4)	0.4288 (6)	0.3745 (2)	0.0427 (2)
N(22)	-0.1522 (3)	0.2014 (6)	0.4318 (2)	0.0408 (2)
$\mathbf{C}(11)$	0.1618 (4)	0.7311 (7)	0.3551 (2)	0.0366 (3)
C(12)	0.2060 (4)	0.5734 (7)	0.4039 (2)	0.0395 (3)
C(13)	0.3382 (4)	0.6236 (9)	0.4410(2)	0.0557 (3)
C(14)	0.2051 (5)	0.3697 (8)	0.3713 (2)	0.0576 (3)
C(21)	-0.2525 (4)	0.2440 (7)	0.3884 (2)	0.0377 (3)
C(22)	-0.3310 (4)	0.0758 (7)	0.3553 (2)	0.0415 (3)
C(23)	-0.3302 (5)	0.0793 (8)	0.2834 (2)	0.0551 (3)
C(24)	-0.4628 (5)	0.0770 (9)	0.3722 (3)	0.0587 (4)

# Table 2. Geometric parameters (Å, °)

N(11) - C(11)	1.320 (5)	$Cl(1) \cdot \cdot \cdot N(11)$	3.20 (2)			
N(12) - C(11)	1.315 (5)	$Cl(1) \cdot \cdot \cdot N(21)$	3.15 (2)			
N(21) - C(21)	1.299 (6)	$Cl(1) \cdot \cdot \cdot N(22)$	3.62 (2)			
N(22) - C(21)	1.301 (5)	$Cl(1) \cdot \cdot \cdot N(22^{i})$	3.26 (2)			
C(11) - C(12)	1.486 (6)	$Cl(1) \cdot \cdot \cdot N(22^{ii})$	3.22 (2)			
C(12)—C(13)	1.517 (6)	$Cl(2) \cdots N(11^{iii})$	3.23 (2)			
C(12) - C(14)	1.526(6)	$Cl(2) \cdot \cdot \cdot N(12^{iv})$	3.19 (2)			
C(21)—C(22)	1.495 (6)	$Cl(2) \cdot \cdot \cdot N(12^{in})$	3.24 (2)			
C(22)—C(23)	1.514 (6)	$Cl(2) \cdot \cdot \cdot N(21)$	3.20 (2)			
C(22)—C(24)	1.513 (6)					
N(11) - C(11) - N(12)	120.4 (4)	N(21)—C(21)—N(22)	120.3 (4)			
N(11) - C(11) - C(12)	117.7 (4)	N(21)—C(21)—C(22)	121.2 (4)			
N(12) - C(11) - C(12)	121.9 (4)	N(22)—C(21)—C(22)	118.5 (4)			
C(11) - C(12) - C(13)	110.0 (4)	C(21)—C(22)—C(23)	110.3 (4)			
C(11) - C(12) - C(14)	110.4 (4)	C(21)—C(22)—C(24)	110.6 (4)			
C(13)—C(12)—C(14)	110.9 (4)	C(23)—C(22)—C(24)	114.4 (4)			
Symmetry codes: (i) x, 1 + y, z; (ii) $-x$ , 1 - y, 1 - z; (iii) $-\frac{1}{2} - x$ , y -						
$\frac{1}{2}, \frac{1}{2} - z;$ (iv) $x - 1, y, z.$						

Data collection: Rigaku AFC-7R software. Cell refinement: Rigaku AFC-7R software. Data reduction: Rigaku AFC-7R software. Program(s) used to solve structure: SIR92 (Altomare et al., 1994). Program(s) used to refine structure: CRYSTALS (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: HU1140). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# Diisopropylammonium Bromide

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#### Abstract

In the title compound,  $C_6H_{16}N^+$ .Br<sup>-</sup>, each diisopropylammonium cation interacts with the bromide anion *via* two hydrogen bonds leading to an infinite onedimensional chain of molecules in the crystal. The cation has an approximate twofold axis of symmetry.

#### Comment

Diisopropylammonium bromide (I) was obtained in the course of studying the reaction of  $BrRe(CO)_5$  with lithium dialkylamides with the intention of generating lithium carbamoylmetallates as precursors for amino-