

Table 2. Selected bond distances (Å) and angles (°) for the bromide ion and the succinimide molecules

Br...N1	3.338 (11)	C5—O5	1.18 (2)
Br...N2	3.380 (10)	C5—C6	1.51 (2)
Br...H1	2.44	C6—C7	1.55 (2)
Br...H2	2.43	C7—C8	1.50 (2)
N1—C1	1.38 (3)	C8—O8	1.18 (2)
N1—C4	1.40 (2)	N3—C31	1.50 (2)
C1—O1	1.21 (2)	N3—C33	1.50 (2)
C1—C2	1.52 (2)	N3—C35	1.50 (2)
C2—C3	1.51 (2)	N3—C37	1.52 (2)
C3—C4	1.52 (2)	C31—C32	1.52 (2)
C4—O4	1.18 (2)	C33—C34	1.52 (2)
N2—C5	1.37 (2)	C35—C36	1.51 (2)
N2—C8	1.40 (2)	C37—C38	1.49 (2)
N1...Br...N2	102.8 (3)	N1—H1...Br	158
H1...Br...H2	96	N2—H2...Br	178

The structure was solved by direct methods (*MULTAN87*; Debaerdemaeker, Germain, Main, Tate & Woolfson, 1987). Some of the H atoms were found in  $\Delta\rho$  maps, the others were placed in calculated positions (C—H and N—H distances 0.95 Å). All were assigned an isotropic displacement parameter of 4.0 Å<sup>2</sup>, which is an approximate average of the displacement parameters of the corresponding C and N atoms. Br, N and O atoms were refined with anisotropic displacement parameters. The H atoms were not refined. The system of computer programs used for this study is described by Lundgren (1982).

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Lists of structure factors, anisotropic thermal parameters, H-atom coordinates and bond lengths and angles have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71314 (12 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: AL1042]

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## Refinement of the Molecular Structure of Benzhydrylamine Hydrochloride

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### Abstract

The molecular conformation of the title compound,  $\alpha$ -phenylbenzylammonium chloride, is helical  $C_2$  with ring twist angles of 57.8 (3) and 68.3 (2)°. Pairs of doubly hydrogen-bonded molecules, connected by hydrogen bonds, form infinite chains along *b*.

### Comment

Benzhydrylamine inhibits smooth-muscle spasms (Bruno Blanch, 1990); some of its derivatives obtained by aryl and/or aminic mono- and di-substitution present greater biological activity than the parent compound. Comparison of the molecular structure of the title compound and its derivatives would help towards understanding the structure–activity relationship. Consequently, a study of the structure of benzhydrylamine hydrochloride (BH), a salt used in activity determination, was performed (Rivero, Punte, Rigotti, Goeta, Quinteros Rivero, Vega & Bruno Blanch, 1991) but all attempts to obtain an *R* factor smaller than 0.07 failed. A thorough analysis of the information was conducted; an instability of the equipment was found that mimics crystal decay. Therefore, to provide a more accurate starting model to analyze the influence of substituents on molecular geometry, a freshly grown single crystal was used to perform the present study.

Intramolecular bond distances and angles in BH do not deviate notably from the expected values. The aromatic rings are planar within experimental error. The bridge angle,  $\Phi$ , is 113.7 (3)°, in agreement with that in diphenylmethane (DPM) [112.5 (6)° (Barnes, Paton, Damewood & Mislow, 1981)]. The dihedral angles subtended between the least-squares planes of the two rings and the central plane defined by C1A—C7—C1B,  $\Phi_1$  and  $\Phi_2$ , are 57.8 (3) and 68.3 (2)°, respectively. The angle between the rings,  $\Phi_{12}$ , is 97.7 (2)°. Barnes, Paton, Damewood & Mislow (1981) found, from data retrieved from the Cambridge Structural Database, that

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steric effects and/or hydrogen bonds produce changes in the molecular conformation when substituents at aryl H-atom positions are introduced in DPM. Comparison between the conformations of BH and DPM shows that the substitution of the ammonium group for a methane H atom decreases  $\Phi_1$  and  $\Phi_2$  [63.9 (6) and 71.1 (6) $^\circ$ , respectively, in DPM] and increases  $\Phi_{12}$  (80.9 $^\circ$ ) but does not alter the overall helical  $C_2$  conformation (Barnes, Paton, Damewood & Mislow, 1981).

The atomic displacement parameters have been analyzed using the program *THMA11* (Maverick & Trueblood, 1988). Orthogonalized  $U_{ij}$  values were calculated for each phenyl group separately and for the whole molecule (without H atoms), assuming a rigid-body model in each case:  $R_U = 0.055$  and  $0.063$  for the phenyl groups and  $0.13$  for the whole molecule. As expected, the whole molecule is less rigid than each phenyl group. A non-rigid model allowing internal torsions gave  $R_U = 0.081$ .

Examination of the packing, exhibited in Fig. 2, shows that the  $NH_3^+$  group is a donor in three hydrogen bonds involving chloride ions as acceptors. The average distance between the N atoms and the accepting chloride ions is 3.21 (3) Å, in agreement with the mean Cl—N distance of 3.19 (7) Å found by Fuller (1959). Pairs of doubly hydrogen-bonded molecules are related by the symmetry operation  $0.5 - x, 0.5 - y, 1 - z$ .

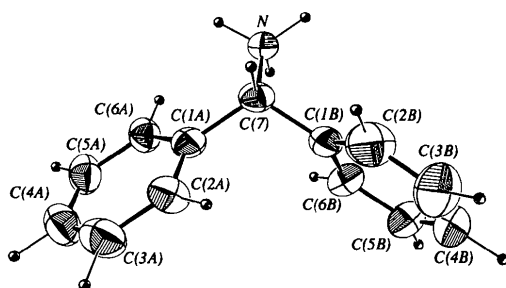


Fig. 1. ORTEP drawing of the molecule with non-H atom labels. The atoms are represented by their 50% probability ellipsoids for thermal motion. H atoms have been arbitrarily reduced.

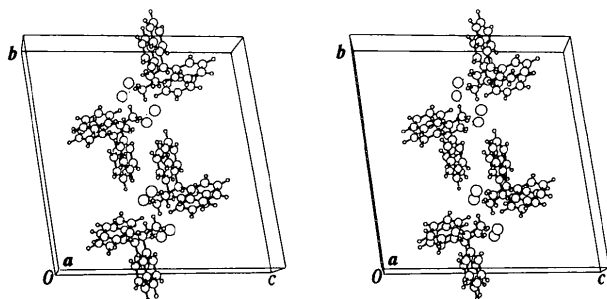


Fig. 2. Stereoscopic view of molecular packing.

## Experimental

### Crystal data

$C_{13}H_{14}N^+.Cl^-$   
 $M_r = 219.72$   
 Monoclinic  
 $C2/c$   
 $a = 21.474$  (4) Å  
 $b = 5.523$  (3) Å  
 $c = 20.161$  (2) Å  
 $\beta = 98.24$  (1) $^\circ$   
 $V = 2366$  (1) Å $^3$   
 $Z = 8$   
 $D_x = 1.233$  Mg m $^{-3}$   
 Mo  $K\alpha$  radiation  
 $\lambda = 0.7093$  Å

Cell parameters from 25 reflections

$\theta = 5-17^\circ$   
 $\mu = 0.288$  mm $^{-1}$   
 $T = 295$  K

Prism  
 $0.37 \times 0.15 \times 0.12$  mm  
 Colorless

Crystal source: the compound was synthesized by Leuckart's method (Moore, 1949)

### Data collection

Enraf-Nonius CAD-4 diffractometer  
 $\theta/2\theta$  scans  
 Absorption correction: none  
 2391 measured reflections  
 1779 independent reflections  
 1115 observed reflections  
 $[I > 3.0\sigma(I)]$

$R_{int} = 0.018$   
 $\theta_{max} = 25^\circ$   
 $h = 0 \rightarrow 25$   
 $k = 0 \rightarrow 6$   
 $l = -24 \rightarrow 24$   
 3 standard reflections  
 frequency: every 60 min  
 intensity variation: 3%

### Refinement

Refinement on  $F$   
 Final  $R = 0.044$   
 $wR = 0.045$   
 $S = 1.91$   
 1115 reflections  
 140 parameters  
 Only H-atom  $U$ 's refined  
 Calculated weights  
 $w = k/[\sigma^2(F) + 0.000177F^2]$   
 $(\Delta/\sigma)_{max} = 0.001$   
 $\Delta\rho_{max} = 0.21$  e Å $^{-3}$   
 $\Delta\rho_{min} = -0.20$  e Å $^{-3}$

Extinction correction:  
 $F' = F(1 - CF^2/\sin\theta)$   
 (Sheldrick, 1976)  
 Extinction coefficient:  
 $C = 7 \times 10^{-7}$

Atomic scattering factors from *International Tables for X-ray Crystallography* (1974, Vol. IV, Table 2.2B)

The H atoms were refined riding on their bound atoms. Overall temperature factors were refined for H atoms of the phenyl, CH $_3$  and ammonium groups. Data collection: *CAD-4 Software* (Enraf-Nonius, 1989). Cell refinement: *CAD-4 Software*. Data reduction: *MolEN* (Fair, 1990). Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELX76* (Sheldrick, 1976). Software used to prepare material for publication: *ORTEP* (Johnson, 1965), *MolEN*.

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

$$U_{eq} = \frac{1}{3} \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	$x$	$y$	$z$	$U_{eq}$
Cl	0.3230	0.0280 (2)	0.4608 (1)	0.0448 (4)
N	0.1759 (1)	-0.0230 (6)	0.4573 (2)	0.042 (1)
C7	0.1378 (2)	-0.0150 (7)	0.3893 (2)	0.039 (1)
C1A	0.1573 (2)	-0.2116 (7)	0.3435 (2)	0.040 (1)
C2A	0.1292 (2)	-0.2119 (9)	0.2773 (2)	0.054 (3)
C3A	0.1465 (2)	-0.384 (1)	0.2327 (2)	0.064 (3)
C4A	0.1920 (2)	-0.5516 (9)	0.2538 (2)	0.060 (3)
C5A	0.2201 (2)	-0.5527 (8)	0.3188 (2)	0.054 (3)

C6A	0.2031 (2)	-0.3841 (8)	0.3638 (2)	0.044 (1)
C1B	0.0683 (2)	-0.0155 (8)	0.3947 (2)	0.044 (1)
C2B	0.0310 (2)	0.1722 (9)	0.3672 (3)	0.073 (3)
C3B	-0.0337 (3)	0.169 (1)	0.3709 (3)	0.098 (4)
C4B	-0.0589 (2)	-0.018 (1)	0.4019 (3)	0.086 (3)
C5B	-0.0224 (2)	-0.203 (1)	0.4297 (3)	0.068 (3)
C6B	0.0410 (2)	-0.2036 (9)	0.4255 (2)	0.053 (3)

Rivero, B. E., Punte, G., Rigotti, G., Goeta, A. E., Quinteros Rivero, M. F., Vega, D. & Bruno Blanch, L. (1991). Proc. Am. Crystallogr. Assoc. Meet. p. 23, Abstract 19.  
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Table 2. Geometric parameters (Å, °)

N—C7	1.493 (5)	C6B—C5B	1.376 (6)
C7—C1A	1.522 (6)	C5B—C4B	1.359 (8)
C7—C1B	1.512 (6)	C4B—C3B	1.359 (9)
C1A—C2A	1.384 (5)	C3B—C2B	1.402 (8)
C1A—C6A	1.388 (6)	Cl...N	3.162 (2)
C2A—C3A	1.395 (7)	Cl...H1	2.2 (1)
C6A—C5A	1.385 (6)	Cl...N <sup>i</sup>	3.192 (4)
C5A—C4A	1.362 (5)	Cl...N <sup>ii</sup>	3.239 (4)
C4A—C3A	1.369 (7)	Cl...H2 <sup>i</sup>	2.3 (1)
C1B—C6B	1.383 (7)	Cl...H3 <sup>ii</sup>	2.3 (1)
C1B—C2B	1.377 (6)		
N—C7—C1A	111.9 (3)	C1A—C6A—C5A	120.7 (4)
N—C7—C1B	110.5 (3)	C6A—C5A—C4A	120.3 (4)
C1A—C7—C1B	113.7 (3)	C5A—C4A—C3A	120.0 (4)
C7—C1A—C2A	117.8 (4)	C2A—C3A—C4A	120.3 (4)
C6B—C5B—C4B	119.6 (5)	C7—C1B—C6B	121.3 (4)
C5B—C4B—C3B	121.1 (5)	C7—C1B—C2B	119.5 (4)
C4B—C3B—C2B	119.8 (5)	C6B—C1B—C2B	119.1 (4)
C1B—C2B—C3B	119.5 (5)	C1B—C6B—C5B	120.8 (5)
C7—C1A—C6A	123.7 (3)	Cl...H1—N	161.4 (4)
C2A—C1A—C6A	118.4 (4)	Cl...H2 <sup>i</sup> —N <sup>i</sup>	161.1 (2)
C1A—C2A—C3A	120.2 (4)	Cl...H3 <sup>ii</sup> —N <sup>ii</sup>	157.7 (3)

Symmetry codes: (i)  $\frac{1}{2} - x, \frac{1}{2} - y, 1 - z$ ; (ii)  $\frac{1}{2} - x, -\frac{1}{2} - y, 1 - z$ .

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Lists of structure factors, anisotropic thermal parameters, H-atom coordinates, complete geometry, least-squares-planes data and torsion angles have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71303 (15 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: CD1017]

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## Conformation of 4,4'-Dichloro-*N,N*-diethylbenzhydramine

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## Abstract

The molecular conformation of the title compound, 4-chloro- $\alpha$ -(4-chlorophenyl)-*N,N*-diethylbenzylamine hydrochloride (C<sub>17</sub>H<sub>20</sub>Cl<sub>2</sub>N<sup>+</sup>.Cl<sup>-</sup>), is helical C<sub>s</sub> with ring twist angles of -77.0 (3) and 62.9 (4)°. The C—H bridge group is involved in a short C—H...Cl contact [3.599 (5) Å].

## Comment

Benzhydramine (I) as well as its derived compounds, obtained by aminic and/or aryl substitution, displays inhibitive activity of the smooth musculature spasms. Some correlation between activity and hydrophobic effects has been observed (Bruno Blanch, 1990). A relationship between charge density localized at the bridge, as obtained by molecular-orbital calculations, and biological activity has been established for monoaminic substituted derivatives (Stiu, Cachau, Castro & Bruno Blanch, 1990).

The present study has been performed as part of a systematic study of derivatives of (I) to investigate the parameters governing the structure-activity relationship, especially in the case of diaminic substitution. Towards this goal the crystal structure of 4,4'-dichloro-*N,N*-diethylbenzhydramine hydrochloride (II) (the salt used

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