Acta Crystallographica Section C Crystal Structure Communications ISSN 0108-2701

Secondary interactions in the isomorphous compounds 2,6-bis(chloromethyl)pyridinium chloride and 2,6bis(bromomethyl)pyridinium bromide

Virginia Lozano and Peter G. Jones*

Institut für Anorganische und Analytische Chemie, Technische Universität Braunschweig, Postfach 3329, 38023 Braunschweig, Germany Correspondence e-mail: p.jones@tu-bs.de

Received 25 June 2004 Accepted 12 July 2004 Online 11 August 2004

The title compounds, $C_7H_8Cl_2N^+\cdot Cl^-$ and $C_7H_8Br_2N^+\cdot Br^-$, are isomorphous. In the crystal packing, layers parallel to the *ac* plane are formed by a classical $N^+ - H \cdots X^-$ hydrogen bond (X = halogen) and two $X \cdots X$ contacts. A third $X \cdots X$ contact links the layers, and a fourth, which is however very long, completes a ladder-like motif of halogen atoms. Hydrogen bonds of the form $C - H \cdots X$ play at best a subordinate role in the packing.

Comment

We are interested in secondary interactions (hydrogen bonds and halogen–halogen contacts) in halides of simple halogenated derivatives of anilines (Gray & Jones, 2002, and references therein) and pyridines [halopyridines (Freytag & Jones, 2001, and references therein) and halomethylpyridines (Jones & Vancea, 2003, and references therein)]. We report here the structures of the isomorphous pair of compounds 2,6-bis-(chloromethyl)pyridinium chloride, (I), and 2,6-bis(bromomethyl)pyridinium bromide, (II). It is common for such pairs to be isomorphous, *e.g.* 4-chloropyridinium chloride and its bromine analogue (Freytag *et al.*, 1999).



The asymmetric units of (I) and (II) are shown in Figs. 1 and 2, respectively. Bond lengths and angles may be regarded as normal, *e.g.* the widened angles at the ring N atom (Tables 1 and 3). The rings are essentially planar [r.m.s. deviations of 0.006 Å for (I) and (II)], with the substituent C atoms lying slightly outside the plane [in (I): C7 -0.105 (3) Å and C8 0.065 (2) Å; in (II): C7 -0.102 (4) Å and C8 0.063 (3) Å]. The



Figure 1

The asymmetric unit of (I) in the crystal. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. The classical hydrogen bond is indicated as a broken line.





The asymmetric unit of (II) in the crystal. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. The classical hydrogen bond is indicated as a broken line.

C-X vectors (X = halogen) of the halomethyl groups extend almost perpendicularly from, and to opposite sides of, the ring (for torsion angles see Tables 1 and 3).

Both compounds form the expected classical hydrogen bond from the N⁺-H group to the halide ion (Tables 2 and 4). Non-classical hydrogen-bond contacts of the form $C-H\cdots X$ are observed, but are all either long (uncorrected $H\cdots X >$ 2.9 Å) and/or markedly non-linear. Three independent halogen-halogen contacts in each structure provide more striking examples of secondary interactions (Table 5). The contact to the anion is, in each case, the shortest (because it is charge-assisted) and essentially linear, as would be expected from the concept of a small positive region in the extension of the C-Cl vector. The other two contacts, between cations, may be classified as type I ($C-X\cdots X$ angles approximately equal) and type II (one $C-X\cdots X$ angle *ca* 90° and the other *ca* 180°) according to the classification of Pedireddi *et al.* (1994).

The net effect of the classical hydrogen bond and the two shorter halogen-halogen interactions is to connect the resi-



Figure 3

A packing diagram for (I), viewed approximately parallel to the b axis, showing one layer at $b \simeq \frac{1}{4}$. Hydrogen bonds and Cl···Cl interactions are indicated by dashed lines. H atoms not involved in hydrogen bonding have been omitted.



Figure 4

A packing diagram for (I) as a projection parallel to the c axis. Cl···Cl interactions are indicated by dashed lines. Note that the chlorine parallelograms (e.g. at the left-hand cell margin), which are not an artefact of projection, can themselves be linked via Cl atoms (see text).

dues, via the glide-plane operators, to form layers parallel to the *ac* plane at $y \simeq \frac{1}{4}, \frac{3}{4}$. One such layer is shown in Fig. 3. The third and longest halogen-halogen interaction then links the layers, in the process forming halogen parallelograms (Fig. 4), with angles 117.54 (1) and 62.46 (1)° for (I), and 117.39 (1) and $62.61 (1)^{\circ}$ for (II). It is noteworthy that the parallelograms are themselves linked into ladder-like tapes via further halogenhalogen contacts [3.8840 (6) Å for (I) and 3.9610 (4) Å for (II); symmetry code: x, $\frac{1}{2} - y$, $\frac{1}{2} + z$], which are much longer than the sum of the van der Waals radii but may still be structurally significant.

Experimental

Compound (I) was obtained as a hygroscopic white solid by bubbling HCl gas through a solution of the corresponding pyridine (0.352 g,2 mmol) in dichloromethane (10 ml), and was recrystallized from dichloromethane-petroleum ether. Compound (II) was obtained in an analogous fashion, but is insoluble in dichloromethane and was recrystallized from ethanol-diisopropyl ether.

Crystal data

$C_7H_8Cl_2N^+ \cdot Cl^-$	Mo $K\alpha$ radiation
$M_r = 212.49$	Cell parameters from 4471
Monoclinic, $P2_1/c$	reflections
a = 7.2167 (6) Å	$\theta = 2.8 - 30.5^{\circ}$
b = 14.6054 (14) Å	$\mu = 0.97 \text{ mm}^{-1}$
c = 8.4990 (8) Å	T = 173 (2) K
$\beta = 98.716 \ (5)^{\circ}$	Tapering prism, colourless
$V = 885.47 (14) \text{ Å}^3$	$0.38 \times 0.07 \times 0.07 \text{ mm}$
Z = 4	
$D_x = 1.594 \text{ Mg m}^{-3}$	

Data collection

Bruker SMART1000 CCD areadetector diffractometer ω and φ scans Absorption correction: multi-scan (SADABS; Bruker, 1998) $T_{\min} = 0.722, \ T_{\max} = 0.942$ 13 915 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.031$ wR(F²) = 0.081 S = 0.992590 reflections 104 parameters

colourless 0.07 mm 2590 independent reflections

1991 reflections with $I > 2\sigma(I)$ $R_{\rm int}=0.038$ $\theta_{\rm max} = 30.0^{\circ}$ $h = -10 \rightarrow 10$ $k=-20\rightarrow 20$ $l = -11 \rightarrow 11$

H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0463P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.53 \ {\rm e} \ {\rm \AA}^{-3}$ $\Delta \rho_{\rm min} = -0.29 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Selected geometric parameters (°) for (I).

C2-N-C6	122.99 (13)		
N-C2-C7-Cl1	-98.77 (14)	N-C6-C8-Cl2	-87.88 (14)
C3-C2-C7-Cl1	78.07 (17)	C5-C6-C8-Cl2	89.62 (16)

Table 2

Hydrogen-bonding geometry (Å, °) for (I).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
N-H1···Cl3	0.95 (2)	2.10(2)	3.0516 (13)	174.4 (17)
$C7 - H7A \cdots Cl3^{i}$	0.99	2.99	3.9632 (17)	169
$C7 - H7B \cdot \cdot \cdot Cl3^{ii}$	0.99	2.94	3.6572 (16)	130
$C8 - H8A \cdots Cl3^{iii}$	0.99	2.95	3.6085 (16)	125
$C8 - H8B \cdot \cdot \cdot Cl3$	0.99	2.79	3.5322 (17)	133
$C3 - H3 \cdot \cdot \cdot Cl2^{iv}$	0.95	2.94	3.7727 (16)	147
$C7 - H7A \cdots Cl2^{iv}$	0.99	2.98	3.6178 (16)	123
$C8-H8A\cdots Cl1^{v}$	0.99	2.72	3.4758 (17)	134
Symmetry codes:	(i) $1 + x, y, z$:	(ii) $1 - x_{i} - x_{i}$	$y_{1} = -7$; (iii) $x_{1} = -7$	$-v_{1}\frac{1}{z}+z$; (iv)

 $1 + x, \frac{1}{2} - y, \frac{1}{2} + z; (v) \quad 1 - x, \frac{1}{2} + y, \frac{1}{2} - z.$

Compound (II)

Crystal data	
$C_7H_8Br_2N^+\cdot Br^-$	Mo <i>Kα</i> radiation
$M_r = 345.87$	Cell parameters from 4
Monoclinic, $P2_1/c$	reflections
a = 7.4633 (6) Å	$\theta = 2.7 - 30.5^{\circ}$
b = 15.0136 (12) Å	$\mu = 12.46 \text{ mm}^{-1}$
c = 8.7377 (6) Å	T = 133 (2) K
$\beta = 99.146 \ (4)^{\circ}$	Plate, colourless
$V = 966.62 (13) \text{ Å}^3$	$0.26 \times 0.18 \times 0.04 \text{ mm}$
Z = 4	
$D_x = 2.377 \text{ Mg m}^{-3}$	

4337

Data collection

Bruker SMART1000 CCD area- detector diffractometer ω and φ scans Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 1998) $T_{min} = 0.282, T_{max} = 0.608$ 17 828 measured reflections	2825 independent reflections 2503 reflections with $I > 2\sigma(I)$ $R_{int} = 0.043$ $\theta_{max} = 30.0^{\circ}$ $h = -10 \rightarrow 10$ $k = -20 \rightarrow 21$ $l = -12 \rightarrow 12$
Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.022$ $wR(F^2) = 0.057$ S = 1.02 2825 reflections 104 parameters H atoms treated by a mixture of independent and constrained refinement	$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0299P)^2 \\ &+ 0.6042P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.001 \\ \Delta\rho_{\text{max}} &= 0.61 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.68 \text{ e } \text{\AA}^{-3} \end{split}$

Table 3

Selected geometric parameters (°) for (II).

C2-N-C6	123.49 (18)		
N-C2-C7-Br1	-96.15 (19)	N-C6-C8-Br2	-87.12 (19)
C3-C2-C7-Br1	81.1 (2)	C5-C6-C8-Br2	90.7 (2)

Table 4

Hydrogen-bonding geometry (Å, °) for (II).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N-H1\cdots Br3$ $C7-H7A\cdots Br3^{i}$ $C7-H7B\cdots Br3^{ii}$ $C8-H8A\cdots Br3^{iii}$ $C8-H8B\cdots Br3$ $C3-H3\cdots Br2^{iv}$	0.88 (3) 0.99 0.99 0.99 0.99 0.99 0.95	2.37 (3) 3.08 3.07 2.99 2.92 3.10	3.2417 (17) 4.056 (2) 3.768 (2) 3.694 (2) 3.697 (2) 3.935 (2)	174 (2) 167 129 129 134 147
$C7 - H7A \cdots Br2^{v}$ $C8 - H8A \cdots Br1^{v}$	0.99 0.99	3.09 2.86	3.753 (2) 3.599 (2)	126 132

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

Crystals of compound (I) cracked badly at 133 K, presumably because of a phase transition, and were therefore measured at the slightly higher temperature of 173 K. The N-bound H atoms were

Table 5 Cl···Cl contacts for (I) and Br···Br contacts for (II) (Å, °).

$C = X \cdots X = C^{\dagger}$	$X \dots X$	$C = X \cdots X$	$X \cdots X = C$
	<i>A A</i>	0 / /	<u> </u>
$C8 - Cl2 \cdot \cdot \cdot Cl3^{vi}$	3.3085 (6)	173.02 (6)	
$C7-Cl1\cdots Cl2^{iv}-C8^{iv}$	3.5695 (6)	77.01 (5)	88.82 (5)
$C7 - Cl1 \cdots Cl2^{vii} - C8^{vii}$	3.5886 (6)	147.05 (5)	71.80 (5)
C8-Br2···Br3 ^{vi}	3.3550 (4)	174.19 (6)	
$C7-Br1\cdots Br2^{iv}-C8^{iv}$	3.7482 (4)	75.06 (7)	89.16 (6)
$C7-Br1\cdots Br2^{vii}-C8^{vii}$	3.6959 (4)	149.72 (6)	71.68 (6)

† Final C atom not applicable for chloride and bromide acceptors. Symmetry codes: (iv) $1 + x, \frac{1}{2} - y, \frac{1}{2} + z$; (vi) $x, \frac{1}{2} - y, z - \frac{1}{2}$; (vii) $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$.

refined freely. Other H atoms were introduced at geometrically calculated positions and refined using a riding model, with fixed C-H distances of 0.95 (sp^2 C-H) or 0.99 Å (methylene H), and with $U_{iso}(H) = 1.2U_{ea}(C)$.

For both compounds, data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL*97.

The authors thank Mr A. Weinkauf for technical assistance.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: BM1577). Services for accessing these data are described at the back of the journal.

References

Bruker (1998). *SMART* (Version 5.0), *SAINT* (Version 4.0) and *SADABS* (Version 2.0). Bruker AXS Inc., Madison, Wisconsin, USA.

Freytag, M. & Jones, P. G. (2001). Z. Naturforsch. Teil B, 56, 889-896.

Freytag, M., Jones, P. G., Ahrens, B. & Fischer, A. K. (1999). New J. Chem. 23, 1137–1138.

Gray, L. & Jones, P. G. (2002). Z. Naturforsch. Teil B, 57, 73-82.

Jones, P. G. & Vancea, F. (2003). CrystEngComm, 5, 303-304.

Pedireddi, V. R., Reddy, D. S., Goud, B. S., Craig, D. C., Rae, A. D. & Desiraju, G. R. (1994). J. Chem. Soc. Perkin Trans. 2, pp. 2353–2360.

Sheldrick, G. M. (1990). Acta Cryst. A46, 467-473.

Sheldrick, G. M. (1997). SHELXL97. University of Göttingen, Germany.

Siemens (1994). XP. Version 5.03. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.