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2-Methylpiperidinium bromide

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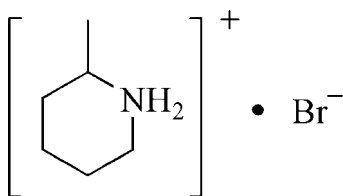
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.008$ Å; R factor = 0.049; wR factor = 0.118; data-to-parameter ratio = 25.4.

In the title organic–inorganic hybrid salt, $\text{C}_6\text{H}_{14}\text{N}^+\cdot\text{Br}^-$, $\text{N}-\text{H}\cdots\text{Br}$ hydrogen bonds link the cations and anions, forming extended hydrogen-bonded chains along the c axis.

Related literature

For general background to ferroelectric organic frameworks, see: Ye *et al.* (2006); Zhang *et al.* (2008, 2010).



Experimental

Crystal data

 $\text{C}_6\text{H}_{14}\text{N}^+\cdot\text{Br}^-$ $M_r = 180.09$ Orthorhombic, $Pbcn$ $a = 22.137$ (4) Å $b = 9.918$ (2) Å $c = 7.5853$ (15) Å $V = 1665.5$ (6) Å³ $Z = 8$ Mo $K\alpha$ radiation $\mu = 4.85$ mm⁻¹ $T = 293$ K $0.55 \times 0.44 \times 0.36$ mm

Data collection

Rigaku SCXmini diffractometer

Absorption correction: multi-scan

(CrystalClear; Rigaku, 2005) $T_{\min} = 0.134$, $T_{\max} = 0.223$

15678 measured reflections

1907 independent reflections

1142 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.109$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.049$ $wR(F^2) = 0.118$ $S = 1.05$

1907 reflections

75 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.38$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.48$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{Br1}$	0.90	2.34	3.238 (4)	176
$\text{N1}-\text{H1B}\cdots\text{Br1}^i$	0.90	2.36	3.262 (3)	176

Symmetry code: (i) $x, -y + 1, z - \frac{1}{2}$.

Data collection: *SCXmini* (Rigaku, 2006); cell refinement: *SCXmini*; data reduction: *SCXmini*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2005); software used to prepare material for publication: *SHELXL97*.

The author is grateful to the starter fund of Southeast University for the purchase of the diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FY2056).

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supporting information

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2-Methylpiperidinium bromide

Qian Xu

S1. Comment

Dielectric-ferroelectrics constitute an interesting class of materials, comprising organic ligands, metal-organic coordination compounds and organic-inorganic hybrids. (Zhang *et al.*, 2010; Zhang *et al.*, 2008; Ye *et al.*, 2006). Unfortunately, the dielectric constant of the title compound as a function of temperature indicates that the permittivity is basically temperature-independent below the melting point of the compound (428–429 K). We have found that title compound has no dielectric disuniformity from 80 K to 405 K. Herein we describe the crystal structure of this compound.

Regarding its crystal structure, the asymmetric unit of the title compound consists of a 2-methylpiperidinium cation and a bromide anion (Fig. 1). The cations and anions are connected by N—H \cdots Br hydrogen bonds, which make a great contribution to the stability of the crystal structure (Fig. 2 and Table 1).

S2. Experimental

The title compound was obtained by the addition of hydrobromic acid (0.8 g, 0.01 mol) to a solution of 2-methylpiperidine (0.97 g, 0.01 mol) in water, *i.e.*, in the stoichiometric ratio of 1:1. Good quality single crystals were obtained by slow evaporation of water after two days (the chemical yield is 65%).

S3. Refinement

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with C—H = 0.97–0.98 Å, N—H = 0.90 Å and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{iso}}(\text{C}, \text{N})$ and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{iso}}(\text{C})$ for methyl hydrogen atoms.

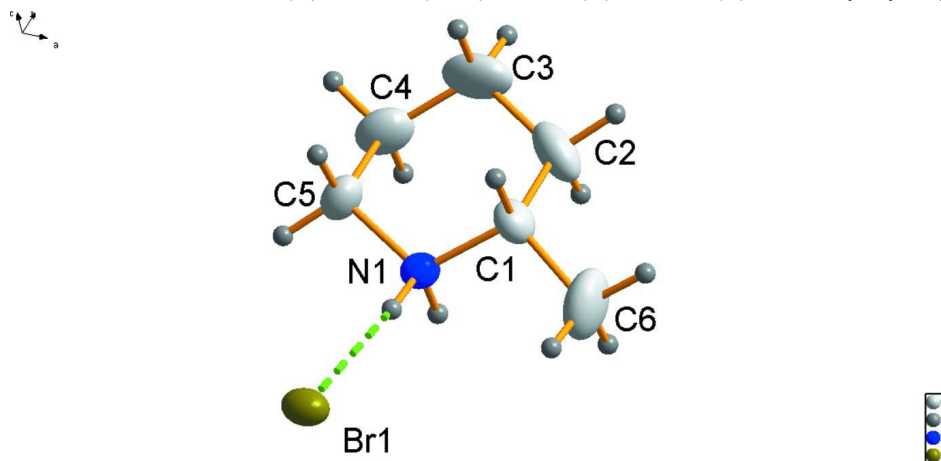
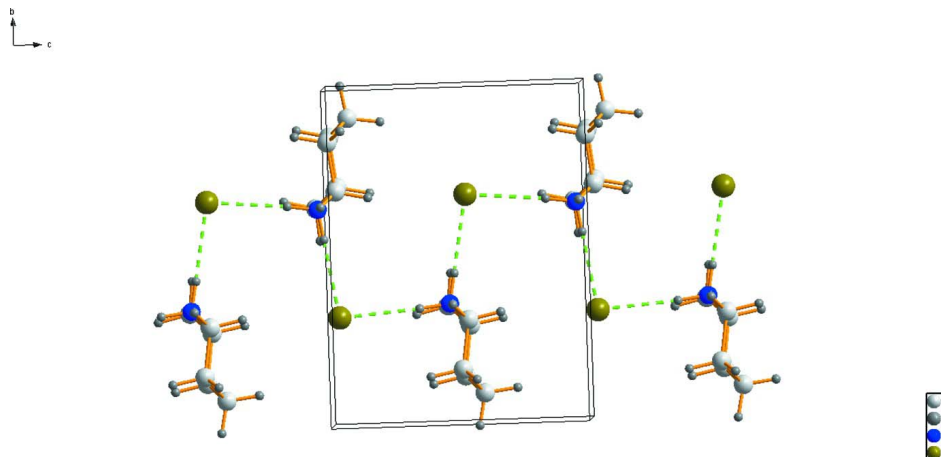


Figure 1

Molecular structure of the title compound, with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. The dashed line indicates a hydrogen bond.

**Figure 2**

A view of the packing of the title compound along the *a* axis. Dashed lines indicate hydrogen bonds.

2-Methylpiperidinium bromide

Crystal data

$C_6H_{14}N^+ \cdot Br^-$

$M_r = 180.09$

Orthorhombic, *Pbcn*

Hall symbol: -P 2n 2ab

$a = 22.137(4) \text{ \AA}$

$b = 9.918(2) \text{ \AA}$

$c = 7.5853(15) \text{ \AA}$

$V = 1665.5(6) \text{ \AA}^3$

$Z = 8$

$F(000) = 736$

$D_x = 1.436 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 3638 reflections

$\theta = 3.0\text{--}27.5^\circ$

$\mu = 4.85 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Block, colorless

$0.55 \times 0.44 \times 0.36 \text{ mm}$

Data collection

Rigaku SCXmini
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: $13.6612 \text{ pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*CrystalClear*; Rigaku, 2005)

$T_{\min} = 0.134$, $T_{\max} = 0.223$

15678 measured reflections

1907 independent reflections

1142 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.109$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 3.4^\circ$

$h = -28 \rightarrow 28$

$k = -12 \rightarrow 12$

$l = -9 \rightarrow 9$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.049$

$wR(F^2) = 0.118$

$S = 1.05$

1907 reflections

75 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0407P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.38 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.48 \text{ e \AA}^{-3}$

Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: $0.0022(5)$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.6672 (2)	0.7098 (5)	0.0449 (6)	0.0546 (13)
H1	0.6680	0.6902	0.1715	0.065*
C2	0.6608 (3)	0.8597 (6)	0.0205 (7)	0.0817 (19)
H2A	0.6633	0.8808	-0.1041	0.098*
H2B	0.6939	0.9049	0.0797	0.098*
C3	0.6018 (3)	0.9124 (6)	0.0922 (8)	0.097 (2)
H3A	0.6007	0.8994	0.2189	0.117*
H3B	0.5988	1.0082	0.0686	0.117*
C4	0.5503 (3)	0.8414 (6)	0.0099 (7)	0.0799 (18)
H4A	0.5128	0.8733	0.0615	0.096*
H4B	0.5494	0.8611	-0.1153	0.096*
C5	0.5556 (2)	0.6950 (5)	0.0365 (6)	0.0594 (13)
H5A	0.5226	0.6497	-0.0231	0.071*
H5B	0.5528	0.6747	0.1613	0.071*
C6	0.7220 (2)	0.6499 (6)	-0.0376 (8)	0.104 (2)
H6A	0.7210	0.5536	-0.0242	0.156*
H6B	0.7575	0.6850	0.0189	0.156*
H6C	0.7229	0.6722	-0.1607	0.156*
N1	0.61363 (14)	0.6448 (4)	-0.0327 (4)	0.0436 (9)
H1A	0.6158	0.5554	-0.0128	0.052*
H1B	0.6143	0.6574	-0.1502	0.052*
Br1	0.61324 (2)	0.32302 (5)	0.03937 (6)	0.0540 (2)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.055 (3)	0.060 (3)	0.049 (3)	-0.010 (2)	-0.014 (2)	0.001 (2)
C2	0.100 (5)	0.071 (4)	0.075 (4)	-0.040 (4)	-0.030 (4)	0.017 (3)
C3	0.160 (7)	0.043 (3)	0.088 (5)	0.015 (4)	0.003 (5)	-0.002 (3)
C4	0.095 (5)	0.062 (4)	0.082 (4)	0.027 (3)	0.012 (3)	0.004 (3)
C5	0.050 (3)	0.060 (3)	0.068 (3)	0.008 (2)	0.013 (2)	0.003 (3)
C6	0.046 (4)	0.143 (6)	0.121 (6)	0.003 (3)	0.000 (3)	0.027 (4)
N1	0.048 (2)	0.041 (2)	0.042 (2)	0.0030 (16)	0.0024 (18)	0.0000 (16)
Br1	0.0752 (4)	0.0440 (3)	0.0427 (3)	0.0012 (2)	-0.0015 (2)	-0.0005 (2)

Geometric parameters (Å, °)

C1—N1	1.473 (5)	C4—H4A	0.9700
C1—C6	1.488 (7)	C4—H4B	0.9700
C1—C2	1.504 (7)	C5—N1	1.475 (5)
C1—H1	0.9800	C5—H5A	0.9700
C2—C3	1.507 (8)	C5—H5B	0.9700
C2—H2A	0.9700	C6—H6A	0.9600
C2—H2B	0.9700	C6—H6B	0.9600
C3—C4	1.478 (8)	C6—H6C	0.9600
C3—H3A	0.9700	N1—H1A	0.9000
C3—H3B	0.9700	N1—H1B	0.9000
C4—C5	1.471 (6)		
N1—C1—C6	108.2 (4)	C5—C4—H4B	109.5
N1—C1—C2	107.9 (4)	C3—C4—H4B	109.5
C6—C1—C2	114.9 (4)	H4A—C4—H4B	108.1
N1—C1—H1	108.6	C4—C5—N1	110.7 (4)
C6—C1—H1	108.6	C4—C5—H5A	109.5
C2—C1—H1	108.6	N1—C5—H5A	109.5
C1—C2—C3	112.4 (4)	C4—C5—H5B	109.5
C1—C2—H2A	109.1	N1—C5—H5B	109.5
C3—C2—H2A	109.1	H5A—C5—H5B	108.1
C1—C2—H2B	109.1	C1—C6—H6A	109.5
C3—C2—H2B	109.1	C1—C6—H6B	109.5
H2A—C2—H2B	107.9	H6A—C6—H6B	109.5
C4—C3—C2	110.5 (5)	C1—C6—H6C	109.5
C4—C3—H3A	109.5	H6A—C6—H6C	109.5
C2—C3—H3A	109.5	H6B—C6—H6C	109.5
C4—C3—H3B	109.5	C1—N1—C5	114.3 (4)
C2—C3—H3B	109.5	C1—N1—H1A	108.7
H3A—C3—H3B	108.1	C5—N1—H1A	108.7
C5—C4—C3	110.5 (5)	C1—N1—H1B	108.7
C5—C4—H4A	109.5	C5—N1—H1B	108.7
C3—C4—H4A	109.5	H1A—N1—H1B	107.6

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1A...Br1	0.90	2.34	3.238 (4)	176
N1—H1B...Br1 ⁱ	0.90	2.36	3.262 (3)	176

Symmetry code: (i) *x*, $-y+1$, $z-1/2$.