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2-Methylpiperidinium bromide

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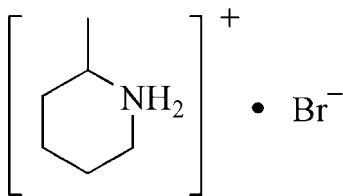
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.008$ Å; R factor = 0.049; wR factor = 0.118; data-to-parameter ratio = 25.4.

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).

References

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supplementary materials

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2-Methylpiperidinium bromide

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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).

Experimental

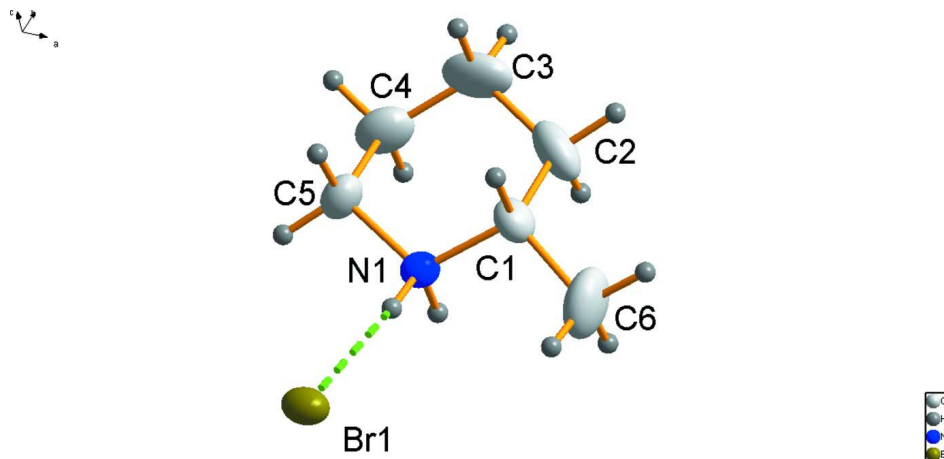
The title compound was obtained by the addition of hydrobromic acid (0.8 g, 0.01 mol) to a solution of 2-methylpiperidine (0.97 g, 0.01 mol) in water, *i.e.*, in the stoichiometric ratio of 1:1. Good quality single crystals were obtained by slow evaporation of water after two days (the chemical yield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

Refinement

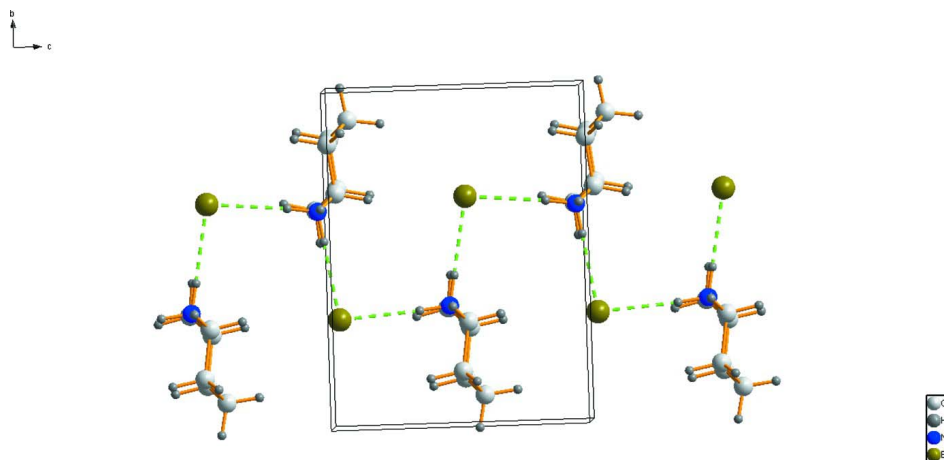
All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with C—H = 0.97–0.98 Å, N—H = 0.90 Å and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{iso}}(\text{C}, \text{N})$ and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{iso}}(\text{C})$ for methyl hydrogen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

Computing details

Data collection: *SCXmini* (Rigaku, 2006); cell refinement: *SCXmini* (Rigaku, 2006); data reduction: *SCXmini* (Rigaku, 2006); 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* (Sheldrick, 2008).


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 | 15678 measured reflections |
| Radiation source: fine-focus sealed tube | 1907 independent reflections |
| Graphite monochromator | 1142 reflections with $I > 2\sigma(I)$ |
| Detector resolution: 13.6612 pixels mm ⁻¹ | $R_{\text{int}} = 0.109$ |
| ω scans | $\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.4^\circ$ |
| Absorption correction: multi-scan (<i>CrystalClear</i> ; Rigaku, 2005) | $h = -28 \rightarrow 28$ |
| $T_{\text{min}} = 0.134$, $T_{\text{max}} = 0.223$ | $k = -12 \rightarrow 12$ |
| | $l = -9 \rightarrow 9$ |

Refinement

| | |
|--|---|
| Refinement on F^2 | Hydrogen site location: inferred from neighbouring sites |
| Least-squares matrix: full | H-atom parameters constrained |
| $R[F^2 > 2\sigma(F^2)] = 0.049$ | $w = 1/[\sigma^2(F_o^2) + (0.0407P)^2]$ |
| $wR(F^2) = 0.118$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| $S = 1.05$ | $(\Delta/\sigma)_{\text{max}} = 0.001$ |
| 1907 reflections | $\Delta\rho_{\text{max}} = 0.38 \text{ e } \text{\AA}^{-3}$ |
| 75 parameters | $\Delta\rho_{\text{min}} = -0.48 \text{ e } \text{\AA}^{-3}$ |
| 0 restraints | Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$ |
| Primary atom site location: structure-invariant direct methods | Extinction coefficient: 0.0022 (5) |
| Secondary atom site location: difference Fourier map | |

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)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|------------|------------|-------------|----------------------------------|
| 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 (\AA , $^\circ$)

| | | | |
|------------|-----------|------------|-----------|
| 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—H...A</i> | <i>D—H</i> | <i>H...A</i> | <i>D...A</i> | <i>D—H...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) $x, -y+1, z-1/2$.