

Bis(2,6-dimethylpyridinium) tetrabromidozincate(II)

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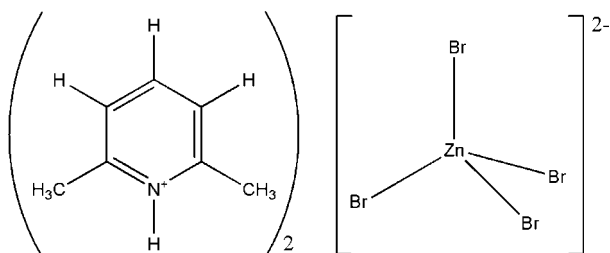
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.03$ Å; R factor = 0.088; wR factor = 0.177; data-to-parameter ratio = 20.2.

In the crystal structure of the title compound, $(\text{C}_7\text{H}_{10}\text{N})_2[\text{ZnBr}_4]$, the coordination geometry of the anion is approximately tetrahedral and a twofold rotation axis passes through the Zn atom. The Zn–Br bond lengths range from 2.400 (2) to 2.408 (3) Å and the Br–Zn–Br angles range from 108.14 (6) to 115.15 (15)°. In the crystal structure, the $[\text{ZnBr}_4]^{2-}$ anion is connected to two cations through $\text{N}-\text{H}\cdots\text{Br}$ and $\text{H}_2\text{C}-\text{H}\cdots\text{Br}$ hydrogen bonds, forming two-dimensional cation–anion–cation layers normal to the b axis. No significant $\text{Br}\cdots\text{Br}$ interactions [the shortest being 4.423 (4) Å] are observed in the structure.

Related literature

The title salt is isotopic with the Co-analogue, see: Ali *et al.* (2008). For non-covalent interactions and their influence on the organization and properties of materials, see: Desiraju (1997); Desiraju & Steiner (1999); Hunter (1994); Allen *et al.* (1997); Dolling *et al.* (2001); Panunto *et al.* (1987); Robinson *et al.* (2000). For the structures of related halo-metal anion salts, see: Ali & Al-Far (2007); Al-Far & Ali (2007); Al-Far & Ali (2009). For distances and angles in $[\text{ZnBr}_4]$ anions, see: Gao *et al.* (2007). For cation bond distances, see: Allen *et al.* (1987).



Experimental

Crystal data

| | |
|--|-----------------------------------|
| $(\text{C}_7\text{H}_{10}\text{N})_2[\text{ZnBr}_4]$ | $V = 2147.9$ (5) Å ³ |
| $M_r = 601.33$ | $Z = 4$ |
| Orthorhombic, $Pbcn$ | Mo $K\alpha$ radiation |
| $a = 17.237$ (2) Å | $\mu = 8.58$ mm ⁻¹ |
| $b = 9.0754$ (17) Å | $T = 293$ K |
| $c = 13.7302$ (14) Å | $0.30 \times 0.20 \times 0.20$ mm |

Data collection

| | |
|---|--|
| Bruker P4 diffractometer | 1850 reflections with $I > 2\sigma(I)$ |
| Absorption correction: numerical (<i>SADABS</i> ; Bruker 2001) | $R_{\text{int}} = 0.086$ |
| $T_{\text{min}} = 0.183$, $T_{\text{max}} = 0.279$ | 3 standard reflections |
| 2020 measured reflections | every 97 reflections |
| 1987 independent reflections | intensity decay: 0.01% |

Refinement

| | |
|---------------------------------|---|
| $R[F^2 > 2\sigma(F^2)] = 0.088$ | 98 parameters |
| $wR(F^2) = 0.177$ | H-atom parameters constrained |
| $S = 0.98$ | $\Delta\rho_{\text{max}} = 0.60$ e Å ⁻³ |
| 1980 reflections | $\Delta\rho_{\text{min}} = -0.48$ e Å ⁻³ |

Table 1

Hydrogen-bond geometry (Å, °).

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|--|--------------|--------------------|-------------|----------------------|
| $\text{N1}-\text{H1}\cdots\text{Br2}$ | 0.86 | 2.49 | 3.351 (12) | 175 |
| $\text{C7}-\text{H7C}\cdots\text{Br1}^{\text{ii}}$ | 0.96 | 2.91 | 3.861 (18) | 171 |

Symmetry code: (ii) $-x + 1, -y + 2, -z + 1$.

Data collection: *XSCANS* (Siemens, 1996); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Sheldrick, 2008); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: AT2771).

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supplementary materials

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Bis(2,6-dimethylpyridinium) tetrabromidozincate(II)

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Comment

Non-covalent interactions play an important role in organizing structural units in both natural and artificial systems (Desiraju, 1997). They exercise important effects on the organization and properties of many materials in areas such as biology (Hunter 1994; Desiraju & Steiner 1999), crystal engineering (see for example: Allen *et al.*, 1997; Dolling *et al.*, 2001) and material science (Panunto *et al.*, 1987; Robinson *et al.*, 2000). The interactions governing the crystal organization are expected to affect the packing and then the specific properties of solids. In connection with ongoing studies (Al-Far & Ali, 2007; Ali & Al-Far, 2007; Ali *et al.*, 2008; Al-Far & Ali, 2009) of the structural aspects of halo-metal anion salts, we herein report the crystal structure of title compound (I).

The asymmetric unit in (I), contains half an anion and one cation (Fig. 1). The geometry of ZnBr_4^{2-} anions is approximately tetrahedral and a twofold rotation axis passes through the Zn^{II} ion (Table 1). The Zn—Br bonds range from 2.400 (2) to 2.408 (3) Å and the Br—Zn—Br angles range from 108.14 (6) to 115.15 (15)°. The bond distances and angles fall in the range of those reported previously for compounds containing Zn—Br anions (Gao *et al.*, 2007). In the cation, the bond lengths and angles are within normal range (Allen *et al.*, 1987).

The packing of the structure (Fig. 2) can be regarded as alternating stacks of anions and stacks of cations. The anion stacks are parallel to the cation stacks, with no significant inter- and intra-stack halogen...halogen interactions [shortest Br...Br interactions being 4.4233 (35) Å]. The anions and cations are interacting significantly through extensive N—H...Br and C—H...Br hydrogen bonding involving Br^- anions and N—H and CH_3 groups (Table 2). These interactions link anions and cations into two-dimensional cation...anion...cation layers normal to the crystallographic *b* axis (Fig. 2).

The N—H...Br and C—H...Br hydrogen bonding are potential building blocks for this stable supramolecular lattice. The stability of this lattice is evident in the isostructurality with the reported analogue (Ali *et al.*, 2008).

Experimental

Warm solution of ZnCl_2 (1.0 mmol) dissolved in absolute ethanol (10 ml) and HBr (60%, 5 ml), was mixed with a stirred hot solution of 2,6-dimethylpyridine (2 mmol) dissolved in ethanol (10 ml). The mixture was then refluxed for 2 h, and then allowed to evaporate undisturbed at room temperature. The salt crystallized over 3 d as nice colourless crystals.

Refinement

H atoms bound to carbon and nitrogen were placed at idealized positions [C—H = 0.93 and 0.96 Å and N—H = 0.86 Å] and allowed to ride on their parent atoms with U_{iso} fixed at 1.2 or 1.5 $U_{\text{eq}}(\text{C}, \text{N})$.

Figures

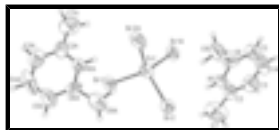


Fig. 1. A view of the asymmetric unit of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry operation A: $-x + 1, y, -z + 1/2$].

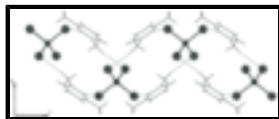


Fig. 2. A packing diagram of (I), shows alternating stacks of anions and cations. C₃N—H...Br—Zn interactions are shown as dashed lines.

Bis(2,6-dimethylpyridinium) tetrabromidozincate(II)

Crystal data

(C₇H₁₀N)₂[ZnBr₄]

$M_r = 601.33$

Orthorhombic, *Pbcn*

Hall symbol: $-P\ 2n\ 2ab$

$a = 17.237\ (2)\ \text{\AA}$

$b = 9.0754\ (17)\ \text{\AA}$

$c = 13.7302\ (14)\ \text{\AA}$

$V = 2147.9\ (5)\ \text{\AA}^3$

$Z = 4$

$F_{000} = 1152$

$D_x = 1.860\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation

$\lambda = 0.71073\ \text{\AA}$

Cell parameters from 250 reflections

$\theta = 3.2\text{--}18.0^\circ$

$\mu = 8.58\ \text{mm}^{-1}$

$T = 293\ \text{K}$

Plate, colourless

$0.30 \times 0.20 \times 0.20\ \text{mm}$

Data collection

Bruker P4
diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 293\ \text{K}$

ω scans

Absorption correction: numerical
(SADABS; Bruker 2001)

$T_{\min} = 0.183, T_{\max} = 0.279$

2020 measured reflections

1987 independent reflections

1850 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.086$

$\theta_{\max} = 25.5^\circ$

$\theta_{\min} = 2.5^\circ$

$h = -1 \rightarrow 20$

$k = -1 \rightarrow 10$

$l = -1 \rightarrow 16$

3 standard reflections

every 97 reflections

intensity decay: 0.01%

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.088$

$wR(F^2) = 0.177$

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0451P)^2]$

| | |
|--|--|
| $S = 0.98$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 1980 reflections | $(\Delta/\sigma)_{\max} < 0.001$ |
| 98 parameters | $\Delta\rho_{\max} = 0.60 \text{ e } \text{\AA}^{-3}$ |
| Primary atom site location: structure-invariant direct methods | $\Delta\rho_{\min} = -0.48 \text{ e } \text{\AA}^{-3}$ |
| | Extinction correction: none |

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.

7 reflections were rejected based on high deviation from observed ones

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

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|-------------|--------------|----------------------------------|
| Zn1 | 0.5000 | 0.7858 (3) | 0.2500 | 0.0644 (10) |
| Br1 | 0.60634 (9) | 0.9276 (2) | 0.18715 (12) | 0.0652 (6) |
| N1 | 0.3780 (7) | 0.6538 (15) | 0.4943 (10) | 0.057 (4) |
| H1 | 0.4205 | 0.6422 | 0.4626 | 0.068* |
| Br2 | 0.54864 (10) | 0.6305 (2) | 0.37844 (13) | 0.0812 (7) |
| C2 | 0.3794 (11) | 0.736 (2) | 0.5723 (14) | 0.062 (5) |
| C3 | 0.3122 (13) | 0.747 (2) | 0.6241 (14) | 0.087 (6) |
| H3 | 0.3123 | 0.8001 | 0.6818 | 0.105* |
| C4 | 0.2454 (14) | 0.683 (3) | 0.5953 (19) | 0.105 (9) |
| H4 | 0.1999 | 0.6943 | 0.6308 | 0.126* |
| C5 | 0.2470 (11) | 0.599 (2) | 0.5103 (15) | 0.093 (7) |
| H5 | 0.2025 | 0.5519 | 0.4879 | 0.112* |
| C6 | 0.3146 (12) | 0.586 (2) | 0.4611 (11) | 0.065 (5) |
| C7 | 0.4534 (12) | 0.819 (2) | 0.5998 (14) | 0.134 (9) |
| H7A | 0.4741 | 0.8668 | 0.5431 | 0.200* |
| H7B | 0.4908 | 0.7507 | 0.6250 | 0.200* |
| H7C | 0.4416 | 0.8915 | 0.6485 | 0.200* |
| C8 | 0.3264 (10) | 0.498 (2) | 0.3707 (13) | 0.102 (7) |
| H8A | 0.3733 | 0.4416 | 0.3764 | 0.153* |
| H8B | 0.3302 | 0.5628 | 0.3157 | 0.153* |
| H8C | 0.2833 | 0.4323 | 0.3618 | 0.153* |

supplementary materials

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|-------------|--------------|
| Zn1 | 0.0466 (16) | 0.079 (2) | 0.0675 (17) | 0.000 | 0.0036 (16) | 0.000 |
| Br1 | 0.0502 (10) | 0.0731 (13) | 0.0722 (11) | -0.0146 (10) | 0.0135 (10) | -0.0024 (12) |
| N1 | 0.041 (8) | 0.073 (11) | 0.057 (9) | -0.007 (8) | 0.001 (8) | -0.013 (9) |
| Br2 | 0.0493 (10) | 0.1111 (17) | 0.0831 (12) | 0.0147 (12) | 0.0050 (11) | 0.0377 (13) |
| C2 | 0.065 (13) | 0.060 (13) | 0.061 (12) | -0.004 (11) | 0.009 (11) | 0.010 (11) |
| C3 | 0.090 (15) | 0.097 (17) | 0.075 (13) | 0.007 (15) | 0.012 (15) | -0.012 (14) |
| C4 | 0.069 (15) | 0.11 (2) | 0.14 (2) | 0.034 (15) | 0.052 (16) | 0.023 (19) |
| C5 | 0.043 (11) | 0.12 (2) | 0.116 (17) | 0.005 (13) | 0.017 (13) | 0.018 (18) |
| C6 | 0.074 (13) | 0.085 (15) | 0.037 (10) | 0.014 (13) | -0.021 (10) | 0.011 (12) |
| C7 | 0.107 (18) | 0.17 (2) | 0.125 (18) | -0.016 (18) | -0.024 (15) | -0.080 (18) |
| C8 | 0.069 (12) | 0.14 (2) | 0.094 (15) | -0.031 (14) | 0.008 (13) | -0.011 (16) |

Geometric parameters (\AA , $^\circ$)

| | | | |
|--|-------------|------------|------------|
| Zn1—Br1 | 2.400 (2) | C4—C5 | 1.39 (3) |
| Zn1—Br1 ⁱ | 2.400 (2) | C4—H4 | 0.9300 |
| Zn1—Br2 ⁱ | 2.408 (3) | C5—C6 | 1.35 (2) |
| Zn1—Br2 | 2.408 (3) | C5—H5 | 0.9300 |
| N1—C2 | 1.31 (2) | C6—C8 | 1.49 (2) |
| N1—C6 | 1.333 (19) | C7—H7A | 0.9600 |
| N1—H1 | 0.8600 | C7—H7B | 0.9600 |
| C2—C3 | 1.36 (2) | C7—H7C | 0.9600 |
| C2—C7 | 1.53 (2) | C8—H8A | 0.9600 |
| C3—C4 | 1.35 (3) | C8—H8B | 0.9600 |
| C3—H3 | 0.9300 | C8—H8C | 0.9600 |
| Br1—Zn1—Br1 ⁱ | 115.15 (15) | C6—C5—C4 | 119 (2) |
| Br1—Zn1—Br2 ⁱ | 108.45 (6) | C6—C5—H5 | 120.6 |
| Br1 ⁱ —Zn1—Br2 ⁱ | 108.14 (6) | C4—C5—H5 | 120.6 |
| Br1—Zn1—Br2 | 108.14 (6) | N1—C6—C5 | 119.8 (17) |
| Br1 ⁱ —Zn1—Br2 | 108.45 (6) | N1—C6—C8 | 114.9 (17) |
| Br2 ⁱ —Zn1—Br2 | 108.35 (16) | C5—C6—C8 | 125 (2) |
| C2—N1—C6 | 124.1 (15) | C2—C7—H7A | 109.5 |
| C2—N1—H1 | 118.0 | C2—C7—H7B | 109.5 |
| C6—N1—H1 | 118.0 | H7A—C7—H7B | 109.5 |
| N1—C2—C3 | 116.7 (18) | C2—C7—H7C | 109.5 |
| N1—C2—C7 | 120.0 (16) | H7A—C7—H7C | 109.5 |
| C3—C2—C7 | 123 (2) | H7B—C7—H7C | 109.5 |
| C4—C3—C2 | 123 (2) | C6—C8—H8A | 109.5 |
| C4—C3—H3 | 118.5 | C6—C8—H8B | 109.5 |
| C2—C3—H3 | 118.5 | H8A—C8—H8B | 109.5 |
| C3—C4—C5 | 118 (2) | C6—C8—H8C | 109.5 |
| C3—C4—H4 | 121.2 | H8A—C8—H8C | 109.5 |
| C5—C4—H4 | 121.2 | H8B—C8—H8C | 109.5 |

| | | | |
|-------------|-------------|-------------|------------|
| C6—N1—C2—C3 | 3(2) | C3—C4—C5—C6 | 0(3) |
| C6—N1—C2—C7 | -175.0 (17) | C2—N1—C6—C5 | -2(3) |
| N1—C2—C3—C4 | -4(3) | C2—N1—C6—C8 | 179.9 (15) |
| C7—C2—C3—C4 | 175 (2) | C4—C5—C6—N1 | 0(3) |
| C2—C3—C4—C5 | 2(3) | C4—C5—C6—C8 | 178.4 (19) |

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

Hydrogen-bond geometry (\AA , $^\circ$)

| <i>D</i> —H \cdots <i>A</i> | <i>D</i> —H | H \cdots <i>A</i> | <i>D</i> \cdots <i>A</i> | <i>D</i> —H \cdots <i>A</i> |
|-----------------------------------|-------------|---------------------|----------------------------|-------------------------------|
| N1—H1 \cdots Br2 | 0.86 | 2.49 | 3.351 (12) | 175 |
| C7—H7C \cdots Br1 ⁱⁱ | 0.96 | 2.91 | 3.861 (18) | 171 |

Symmetry codes: (ii) $-x+1, -y+2, -z+1$.

Fig. 1

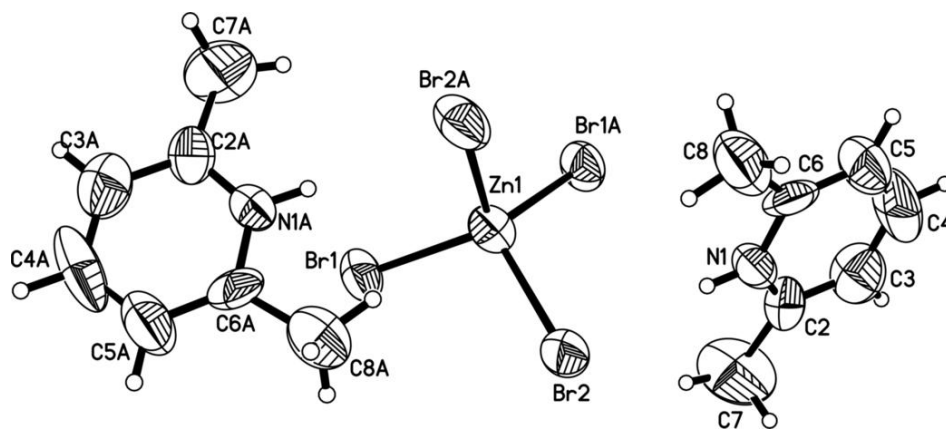


Fig. 2

