

2,3-Dimethylquinoxalinediium hexa-bromidostannate(IV) trihydrate

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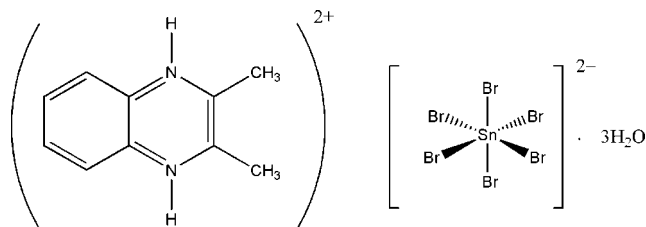
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 Key indicators: single-crystal X-ray study; $T = 84$ K; mean $\sigma(\text{C}-\text{C}) = 0.013$ Å; R factor = 0.062; wR factor = 0.132; data-to-parameter ratio = 23.0.

In the title compound, $(\text{C}_{10}\text{H}_{12}\text{N}_2)[\text{SnBr}_6]\cdot 3\text{H}_2\text{O}$, the asymmetric unit contains one cation, one anion and three water molecules, and the Sn atom has a distorted octahedral environment. In the crystal structure, intra- and intermolecular hydrogen bonds and $\text{Br}\cdots\text{Br}$ interactions [$\text{Br}\cdots\text{Br} = 3.6726$ (19), 3.6913 (18) and 3.6517 (21) Å] lead to the formation of a supramolecular architecture.

Related literature

For related literature, see: Desiraju (1997); Ali *et al.* (2007); Ali & Al-Far (2007); Al-Far & Ali (2007*a,b*); Al-Far *et al.* (2007); Tudela & Khan (1991); Willey *et al.* (1998). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$(\text{C}_{10}\text{H}_{12}\text{N}_2)[\text{SnBr}_6]\cdot 3\text{H}_2\text{O}$
 $M_r = 812.37$
 Monoclinic, $P2_1/c$
 $a = 9.898$ (3) Å
 $b = 13.620$ (5) Å
 $c = 16.098$ (6) Å
 $\beta = 103.599$ (6)°

$V = 2109.3$ (13) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 12.59$ mm⁻¹
 $T = 84$ (2) K
 $0.30 \times 0.20 \times 0.10$ mm

Data collection

Rigaku Mercury CCD diffractometer

Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2000)
 $T_{\min} = 0.055$, $T_{\max} = 0.277$

27021 measured reflections
 4633 independent reflections

4210 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.045$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.132$
 $S = 1.20$
 4633 reflections

201 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 1.41$ e Å⁻³
 $\Delta\rho_{\text{min}} = -1.11$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Sn1—Br1	2.4405 (12)	Sn1—Br6	2.6570 (12)
Sn1—Br3	2.4603 (12)	Sn1—Br5	2.7026 (12)
Sn1—Br4	2.6534 (11)	Sn1—Br2	2.7114 (11)
Br1—Sn1—Br3	177.29 (4)	Br4—Sn1—Br5	85.87 (4)
Br1—Sn1—Br4	93.48 (4)	Br6—Sn1—Br5	178.17 (3)
Br3—Sn1—Br4	88.41 (4)	Br1—Sn1—Br2	87.41 (4)
Br1—Sn1—Br6	86.37 (4)	Br3—Sn1—Br2	90.69 (4)
Br3—Sn1—Br6	95.38 (4)	Br4—Sn1—Br2	179.08 (4)
Br4—Sn1—Br6	95.87 (4)	Br6—Sn1—Br2	84.41 (4)
Br1—Sn1—Br5	92.93 (3)	Br5—Sn1—Br2	93.86 (4)
Br3—Sn1—Br5	85.26 (4)		

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1W—H1W1 \cdots O2W	0.86	1.78	2.640 (14)	170.00
O1W—H1W2 \cdots Br3 ⁱ	0.87	2.79	3.348 (9)	123.1 (5)
O2W—H2W1 \cdots Br2 ⁱⁱ	0.84	2.57	3.401 (12)	167.4 (7)
O2W—H2W2 \cdots Br3 ⁱⁱ	0.87	2.83	3.482 (11)	132.9 (7)
O3W—H3W2 \cdots Br1	0.86	2.83	3.349 (11)	120.00
N1—H1N \cdots O1W	0.86	1.82	2.672 (10)	172
N2—H2N \cdots O3W ⁱⁱⁱ	0.86	1.81	2.660 (11)	172

Symmetry codes: (i) $x - 1, -y + \frac{3}{2}, z + \frac{1}{2}$; (ii) $-x + 2, y - \frac{1}{2}, -z + \frac{1}{2}$; (iii) $x, -y + \frac{3}{2}, z + \frac{1}{2}$.

Data collection: *CrystalClear* (Rigaku, 2000); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); 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: HK2278).

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supplementary materials

Acta Cryst. (2007). E63, m2102-m2103 [doi:10.1107/S1600536807032771]

2,3-Dimethylquinoxalinediium hexabromidostannate(IV) trihydrate

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Comment

Noncovalent interactions play an important role in organizing structural units in both natural and artificial systems (Desiraju, 1997). In connection with ongoing studies (Ali *et al.*, 2007; Ali & Al-Far, 2007; Al-Far & Ali, 2007a,b) of the structural aspects of bromo metal anions salts, we herein report the crystal structure of the title compound, (I).

The asymmetric unit of the title compound, (I), contains one cation, one anion and three water molecules, where the Sn atom has a distorted octahedral environment (Fig. 1, Table 1). The bond lengths and angles (Table 1) are generally within normal ranges (Allen *et al.*, 1987). In the anion, the Sn1—Br1 [2.4406 (12) Å] and Sn1—Br3 [2.4605 (12) Å] bonds are shorter than the other Sn—Br bonds, in which they are within the range of Sn—Br bonds reported previously for compounds containing [SnBr₆]²⁻ anions (Ali & Al-Far, 2007; Al-Far & Ali, 2007a,b; Al-Far *et al.*, 2007; Tudela & Khan, 1991; Willey *et al.*, 1998). In the cation, the bond lengths and angles are in accordance with the corresponding values (Al-Far & Ali, 2007a,b; Ali *et al.*, 2007). The cation is, of course, planar, in which C1 and C10 atoms are also coplanar.

The packing of the structure can be regarded as alternating layers of anions and cations (Fig. 2). The anions within each layer (Fig. 3) interact *via* Br1^{iv}⋯Br3^{iv} = 3.6726 (19) Å [symmetry code: (iv) $x, 3/2 + y, 1/2 + z$] interactions parallel to *c* axis. Each anionic layer further interacts *via* Br2^v⋯Br6^v = 3.6913 (18) Å [symmetry code: (v) $-x + 3, 1/2 + y, 1/2 - z$] and Br4^{vi}⋯Br4^{vi} = 3.6517 (21) Å [symmetry code: (vi) $-x + 2, -y + 1, -z$] interactions to form a two-dimensional anionic network parallel to *ac* plane (Fig. 3).

The crystal supramolecularity is represented in the significantly short hydrogen bonds (Table 2, Fig. 4) along with Br⋯Br interactions that allow the formation of supramolecular assembly of the anion, cation and water molecules in three-dimensional structure, in which they may be effective in the stabilization of the crystal structure.

Experimental

For the preparation of (I), tin metal (0.119 mg, 1 mmol) dissolved in absolute ethanol (10 ml), HBr (60%, 5 ml) and liquid Br₂ (60%, 2 ml), was added dropwise to a stirred hot solution of 2,3-dimethylquinoxaliniium (0.158 mg, 1 mmol) dissolved in ethanol (10 ml) and HBr (60%, 2 ml). After refluxing for 1 h, the mixture was filtered off, and then allowed to stand undisturbed at room temperature. The salt crystallized over 1 d as yellow crystals. Crystals were filtered off, washed with diethylether and dried under vacuum (yield; 0.750 mg; 92.4%).

Refinement

H atoms were positioned geometrically, with O—H = 0.84–0.90 Å (for H₂O), N—H = 0.86 Å (for NH) and C—H = 0.95 and 0.98 Å, for aromatic and methylene H atoms, and constrained to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C}, \text{O}, \text{N})$, where $x = 1.2$ for NH and aromatic H atoms, and $x = 1.5$ for all other H atoms.

Figures

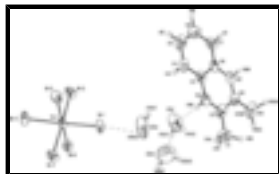


Fig. 1. The molecular structure of the title molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen bonds are shown as dashed lines.



Fig. 2. A partial packing diagram of (I). Hydrogen bonds and Br...Br interactions are shown as dashed lines. Viewed down *c* axis, where *a* and *b* axes are horizontal and vertical, respectively.

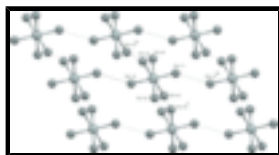


Fig. 3. Anion network of Br...Br interactions. Viewed down *b* axis, where *c* and *a* axes are horizontal and vertical, respectively [Symmetry codes: (iv) $x, 3/2 + y, 1/2 + z$; (v) $-x + 3, 1/2 + y, 1/2 - z$; (vi) $-x + 2, -y + 1, -z$].



Fig. 4. Anion, cation and water molecules intermolecular interactions. Sn—Br...H—O, H₂O...H—OH, H₂O...H—N and Br...Br (parallel to *c* axis) intermolecular interactions are shown as dashed lines. Viewed down *b* axis, where *c* and *a* axes are horizontal and vertical, respectively. H atoms are omitted for clarity. Oxygen atoms of water molecules shown as balls.

'2,3-Dimethylquinoxalinediium hexabromostannate(IV) trihydrate'

Crystal data

(C₁₀H₁₂N₂)[SnBr₆]·3H₂O

M_r = 812.37

Monoclinic, *P*2₁/*c*

Hall symbol: -*P* 2ybc

a = 9.898 (3) Å

b = 13.620 (5) Å

c = 16.098 (6) Å

β = 103.599 (6)°

V = 2109.3 (13) Å³

Z = 4

*F*₀₀₀ = 1504

D_x = 2.558 Mg m⁻³

Mo *K*α radiation

λ = 0.71070 Å

Cell parameters from 4210 reflections

θ = 1.3–27.9°

μ = 12.59 mm⁻¹

T = 84 (2) K

Needle, yellow

0.30 × 0.20 × 0.10 mm

Data collection

Rigaku Mercury CCD
diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

Detector resolution: 14.6306 pixels mm⁻¹

T = 84(2) K

4633 independent reflections

4210 reflections with *I* > 2σ(*I*)

*R*_{int} = 0.045

θ_{max} = 27.5°

θ_{min} = 2.1°

dtintegrate.ref scans $h = -12 \rightarrow 11$
 Absorption correction: multi-scan $k = -17 \rightarrow 17$
 Shape Tracing Software $l = -20 \rightarrow 20$
 $T_{\min} = 0.055$, $T_{\max} = 0.277$
 27021 measured reflections

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.062$	H-atom parameters constrained
$wR(F^2) = 0.132$	$w = 1/[\sigma^2(F_o^2) + (0.0478P)^2 + 7.030P]$
$S = 1.20$	where $P = (F_o^2 + 2F_c^2)/3$
4633 reflections	$(\Delta/\sigma)_{\max} = 0.001$
201 parameters	$\Delta\rho_{\max} = 1.41 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{\min} = -1.11 \text{ e } \text{\AA}^{-3}$
	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 $F^2 > 2\sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
Sn1	1.23031 (5)	0.73783 (4)	0.15141 (3)	0.03598 (16)
Br1	1.17497 (9)	0.68693 (8)	0.28495 (5)	0.0562 (3)
Br2	1.41637 (10)	0.86480 (7)	0.24513 (6)	0.0600 (3)
Br3	1.28560 (9)	0.79729 (8)	0.01892 (5)	0.0594 (3)
Br4	1.04854 (9)	0.61509 (7)	0.05744 (6)	0.0573 (3)
Br5	1.02270 (9)	0.87143 (7)	0.11762 (6)	0.0538 (2)
Br6	1.43853 (9)	0.60996 (7)	0.18886 (6)	0.0529 (2)
O1W	0.5390 (9)	0.6748 (7)	0.4150 (5)	0.099 (3)
H1W1	0.5917	0.6271	0.4063	0.149*
H1W2	0.4568	0.6502	0.4107	0.149*
O2W	0.7208 (12)	0.5361 (8)	0.4048 (8)	0.150 (5)
H2W1	0.6941	0.4985	0.3624	0.226*
H2W2	0.7408	0.5002	0.4510	0.226*

supplementary materials

O3W	0.9249 (12)	0.6547 (9)	0.3909 (5)	0.134 (4)
H3W1	0.9395	0.7059	0.4277	0.201*
H3W2	1.0060	0.6290	0.3961	0.201*
N1	0.6611 (6)	0.7296 (5)	0.5744 (4)	0.0432 (15)
H1N	0.6144	0.7109	0.5249	0.052*
N2	0.8084 (7)	0.7870 (5)	0.7326 (4)	0.0457 (16)
H2N	0.8534	0.8050	0.7827	0.055*
C1	0.6530 (10)	0.5601 (7)	0.6141 (7)	0.067 (3)
H1A	0.5879	0.5584	0.5579	0.101*
H1B	0.6076	0.5337	0.6572	0.101*
H1C	0.7349	0.5202	0.6126	0.101*
C2	0.6965 (8)	0.6629 (7)	0.6361 (6)	0.049 (2)
C3	0.6962 (7)	0.8271 (6)	0.5865 (5)	0.0399 (17)
C4	0.6556 (9)	0.8934 (7)	0.5197 (5)	0.050 (2)
H4	0.6045	0.8736	0.4646	0.060*
C5	0.6936 (10)	0.9891 (8)	0.5380 (7)	0.063 (3)
H5	0.6690	1.0367	0.4939	0.075*
C6	0.7682 (10)	1.0201 (7)	0.6201 (8)	0.067 (3)
H6	0.7899	1.0878	0.6289	0.080*
C8	0.7744 (7)	0.8569 (6)	0.6704 (5)	0.0399 (17)
C7	0.8110 (9)	0.9554 (7)	0.6884 (6)	0.055 (2)
H7	0.8613	0.9766	0.7432	0.066*
C9	0.7761 (8)	0.6942 (6)	0.7202 (5)	0.0453 (19)
C10	0.8231 (11)	0.6235 (8)	0.7927 (6)	0.073 (3)
H10A	0.7928	0.6469	0.8429	0.109*
H10B	0.9248	0.6186	0.8064	0.109*
H10C	0.7826	0.5587	0.7761	0.109*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Sn1	0.0316 (3)	0.0490 (3)	0.0301 (3)	-0.0024 (2)	0.0128 (2)	-0.0062 (2)
Br1	0.0512 (5)	0.0851 (7)	0.0392 (4)	0.0018 (4)	0.0247 (4)	0.0052 (4)
Br2	0.0516 (5)	0.0684 (6)	0.0626 (6)	-0.0193 (4)	0.0188 (4)	-0.0240 (5)
Br3	0.0540 (5)	0.0901 (7)	0.0412 (4)	-0.0031 (5)	0.0252 (4)	0.0099 (4)
Br4	0.0517 (5)	0.0687 (6)	0.0546 (5)	-0.0196 (4)	0.0184 (4)	-0.0223 (4)
Br5	0.0447 (5)	0.0622 (5)	0.0588 (5)	0.0063 (4)	0.0210 (4)	-0.0041 (4)
Br6	0.0473 (5)	0.0622 (5)	0.0540 (5)	0.0130 (4)	0.0217 (4)	0.0009 (4)
O1W	0.106 (6)	0.134 (7)	0.054 (4)	0.010 (6)	0.013 (4)	-0.019 (5)
O2W	0.175 (11)	0.107 (8)	0.209 (13)	-0.005 (7)	0.123 (10)	-0.016 (8)
O3W	0.151 (9)	0.206 (11)	0.043 (4)	0.017 (8)	0.016 (5)	0.017 (6)
N1	0.035 (3)	0.056 (4)	0.040 (3)	0.003 (3)	0.011 (3)	-0.004 (3)
N2	0.038 (4)	0.070 (5)	0.032 (3)	0.001 (3)	0.013 (3)	0.000 (3)
C1	0.059 (6)	0.056 (6)	0.094 (8)	-0.003 (5)	0.030 (5)	-0.005 (5)
C2	0.031 (4)	0.059 (5)	0.059 (5)	-0.001 (4)	0.016 (4)	0.011 (4)
C3	0.032 (4)	0.049 (4)	0.045 (4)	0.004 (3)	0.021 (3)	0.002 (4)
C4	0.046 (5)	0.065 (6)	0.042 (4)	0.007 (4)	0.017 (4)	0.012 (4)
C5	0.054 (6)	0.071 (7)	0.070 (6)	0.013 (5)	0.030 (5)	0.023 (5)

C6	0.065 (6)	0.047 (5)	0.103 (8)	0.003 (4)	0.049 (6)	0.001 (5)
C8	0.027 (4)	0.061 (5)	0.036 (4)	0.002 (3)	0.015 (3)	0.001 (3)
C7	0.049 (5)	0.061 (5)	0.065 (6)	-0.005 (4)	0.032 (4)	-0.015 (5)
C9	0.036 (4)	0.060 (5)	0.043 (4)	-0.001 (4)	0.017 (3)	0.007 (4)
C10	0.063 (6)	0.099 (8)	0.060 (6)	0.009 (6)	0.023 (5)	0.037 (6)

Geometric parameters (Å, °)

Sn1—Br1	2.4405 (12)	C1—H1A	0.9800
Sn1—Br3	2.4603 (12)	C1—H1B	0.9800
Sn1—Br4	2.6534 (11)	C1—H1C	0.9800
Sn1—Br6	2.6570 (12)	C2—C9	1.461 (12)
Sn1—Br5	2.7026 (12)	C3—C4	1.391 (11)
Sn1—Br2	2.7114 (11)	C3—C8	1.447 (11)
O1W—H1W1	0.86000	C4—C5	1.368 (13)
O1W—H1W2	0.87000	C4—H4	0.9500
O2W—H2W1	0.84000	C5—C6	1.418 (14)
O2W—H2W2	0.87000	C5—H5	0.9500
O3W—H3W1	0.90000	C6—C7	1.395 (14)
O3W—H3W2	0.86000	C6—H6	0.9500
N1—C2	1.330 (10)	C8—C7	1.403 (12)
N1—C3	1.375 (10)	C7—H7	0.9500
N1—H1N	0.8600	C9—C10	1.500 (11)
N2—C9	1.308 (11)	C10—H10A	0.9800
N2—C8	1.365 (10)	C10—H10B	0.9800
N2—H2N	0.8600	C10—H10C	0.9800
C1—C2	1.483 (13)		
Br1—Sn1—Br3	177.29 (4)	N1—C2—C9	118.9 (8)
Br1—Sn1—Br4	93.48 (4)	N1—C2—C1	117.1 (8)
Br3—Sn1—Br4	88.41 (4)	C9—C2—C1	124.0 (8)
Br1—Sn1—Br6	86.37 (4)	N1—C3—C4	120.0 (8)
Br3—Sn1—Br6	95.38 (4)	N1—C3—C8	117.8 (7)
Br4—Sn1—Br6	95.87 (4)	C4—C3—C8	122.2 (7)
Br1—Sn1—Br5	92.93 (3)	C5—C4—C3	115.9 (8)
Br3—Sn1—Br5	85.26 (4)	C5—C4—H4	122.0
Br4—Sn1—Br5	85.87 (4)	C3—C4—H4	122.0
Br6—Sn1—Br5	178.17 (3)	C4—C5—C6	122.7 (9)
Br1—Sn1—Br2	87.41 (4)	C4—C5—H5	118.6
Br3—Sn1—Br2	90.69 (4)	C6—C5—H5	118.6
Br4—Sn1—Br2	179.08 (4)	C7—C6—C5	122.8 (9)
Br6—Sn1—Br2	84.41 (4)	C7—C6—H6	118.6
Br5—Sn1—Br2	93.86 (4)	C5—C6—H6	118.6
H1W1—O1W—H1W2	106.70	N2—C8—C7	120.5 (8)
H2W1—O2W—H2W2	108.30	N2—C8—C3	118.3 (7)
H3W1—O3W—H3W2	104.40	C7—C8—C3	121.1 (8)
C2—N1—C3	122.6 (7)	C6—C7—C8	115.2 (9)
C2—N1—H1N	118.7	C6—C7—H7	122.4
C3—N1—H1N	118.7	C8—C7—H7	122.4
C9—N2—C8	123.5 (7)	N2—C9—C2	118.9 (7)

supplementary materials

C9—N2—H2N	118.2	N2—C9—C10	118.8 (8)
C8—N2—H2N	118.2	C2—C9—C10	122.4 (8)
C2—C1—H1A	109.5	C9—C10—H10A	109.5
C2—C1—H1B	109.5	C9—C10—H10B	109.5
H1A—C1—H1B	109.5	H10A—C10—H10B	109.5
C2—C1—H1C	109.5	C9—C10—H10C	109.5
H1A—C1—H1C	109.5	H10A—C10—H10C	109.5
H1B—C1—H1C	109.5	H10B—C10—H10C	109.5

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1W—H1W1 \cdots O2W	0.86	1.78	2.640 (14)	170.00
O1W—H1W2 \cdots Br3 ⁱ	0.87	2.79	3.348 (9)	123.1 (5)
O2W—H2W1 \cdots Br2 ⁱⁱ	0.84	2.57	3.401 (12)	167.4 (7)
O2W—H2W2 \cdots Br3 ⁱⁱ	0.87	2.83	3.482 (11)	132.9 (7)
O3W—H3W2 \cdots Br1	0.86	2.83	3.349 (11)	120.00
N1—H1N \cdots O1W	0.86	1.82	2.672 (10)	172
N2—H2N \cdots O3W ⁱⁱⁱ	0.86	1.81	2.660 (11)	172

Symmetry codes: (i) $x-1, -y+3/2, z+1/2$; (ii) $-x+2, y-1/2, -z+1/2$; (iii) $x, -y+3/2, z+1/2$.

Fig. 1

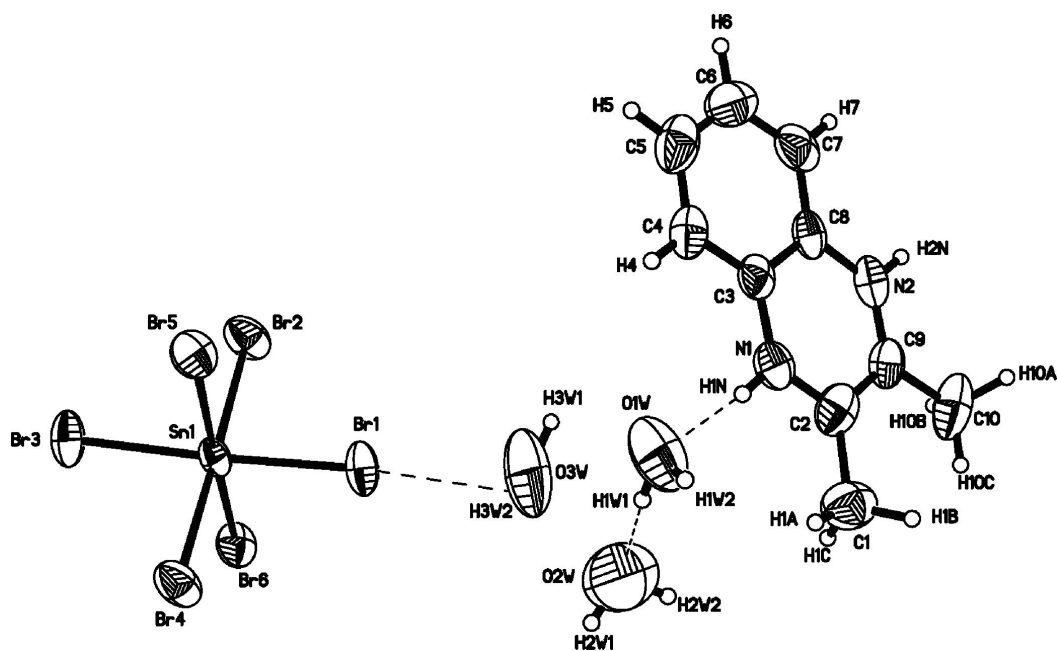


Fig. 2

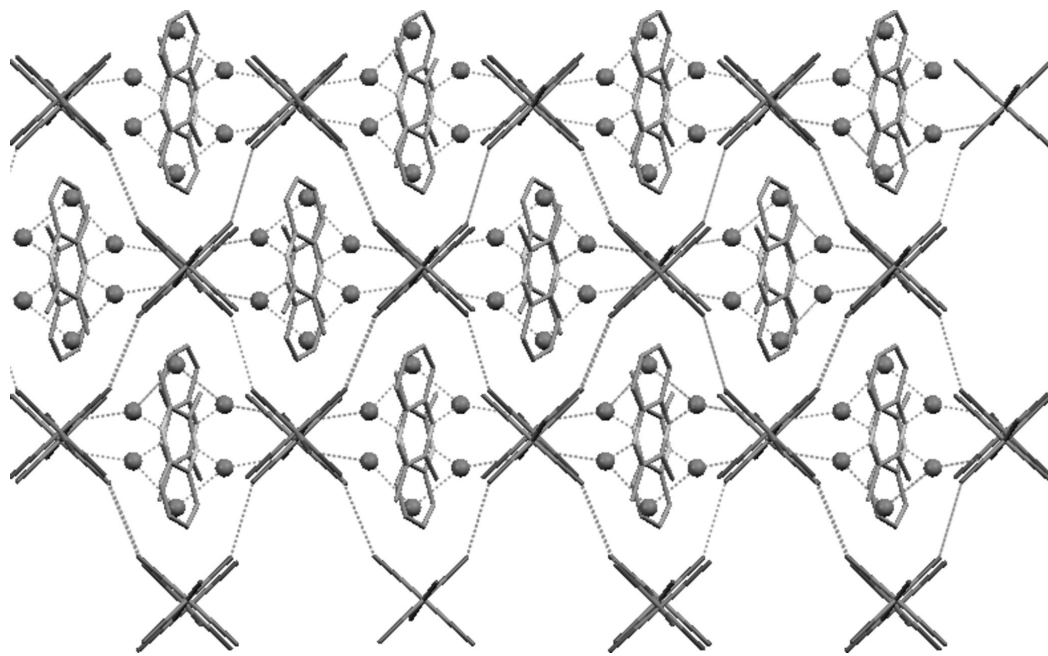


Fig. 3

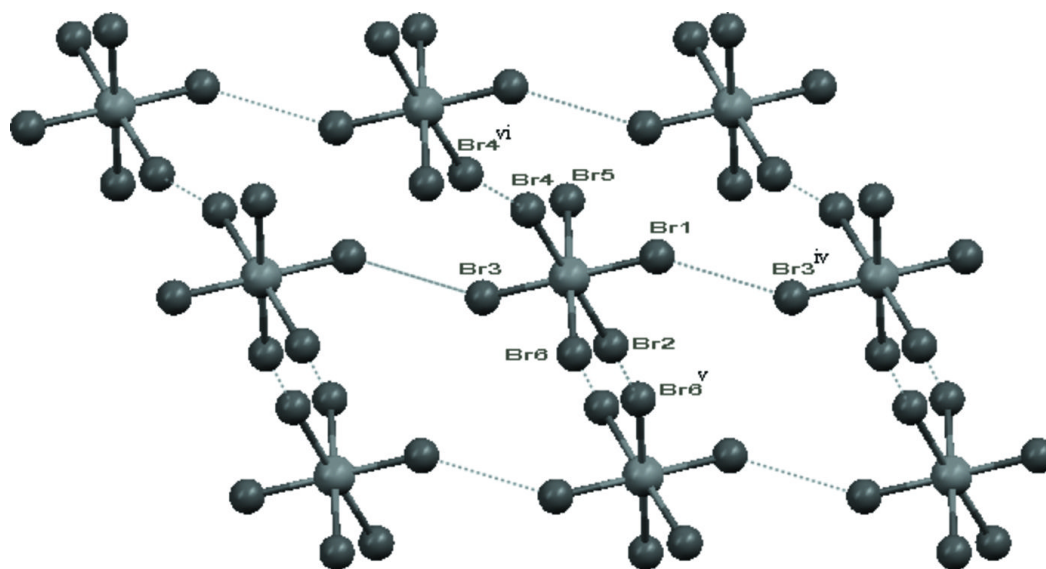


Fig. 4

