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4-(Dimethylamino)pyridinium dibromido(4-bromophenyl)dimethylstannate(IV)

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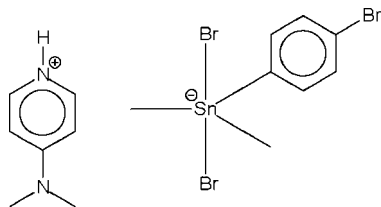
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 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.031; wR factor = 0.096; data-to-parameter ratio = 22.4.

The anion in the title compound, $(\text{C}_7\text{H}_{11}\text{N}_2)[\text{SnBr}_2(\text{CH}_3)_2(\text{C}_6\text{H}_4\text{Br})]$, is five-coordinate within a distorted *trans*- C_3SnBr_2 trigonal-bipyramidal geometry. The cation and anion are linked by an $\text{N}-\text{H}\cdots\text{Br}$ hydrogen bond.

Related literature

For the crystal structure of 4-(dimethylamino)pyridinium dibromidotriphenylstannate(IV), see: Norhafiza *et al.* (2008).



Experimental

Crystal data

 $(\text{C}_7\text{H}_{11}\text{N}_2)[\text{SnBr}_2(\text{CH}_3)_2(\text{C}_6\text{H}_4\text{Br})]$ $M_r = 587.76$

 Triclinic, $P\bar{1}$
 $a = 7.3397$ (1) Å
 $b = 11.1034$ (2) Å
 $c = 12.2270$ (2) Å
 $\alpha = 100.038$ (1)°
 $\beta = 102.472$ (1)°
 $\gamma = 94.679$ (1)°

 $V = 950.65$ (3) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 7.64$ mm⁻¹
 $T = 100$ (2) K
 $0.30 \times 0.20 \times 0.02$ mm

Data collection

 Bruker SMART APEX
 diffractometer
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.208$, $T_{\max} = 0.862$

 8826 measured reflections
 4346 independent reflections
 3718 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.096$
 $S = 1.13$
 4346 reflections

 194 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 1.07$ e Å⁻³
 $\Delta\rho_{\min} = -1.24$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{Br1}$	0.88	2.56	3.319 (3)	146

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2315).

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supporting information

Acta Cryst. (2008). E64, m1391 [doi:10.1107/S1600536808032248]

4-(Dimethylamino)pyridinium dibromido(4-bromophenyl)dimethylstannate(IV)

Chin Koon Yau, Kong Mun Lo and Seik Weng Ng

S1. Experimental

Bis(4-bromophenyl)dimethyltin (0.10 g, 0.2 mmol) [which was prepared by the reaction between dimethyltin dichloride and 4-bromophenylmagnesium bromide] and 4-dimethylaminopyridine hydrobromide perbromide (0.08 g, 0.2 mmol) were heated in chloroform (100 ml) for 3 h. The solution was filtered and the solvent allowed to evaporate to give colorless crystals.

S2. Refinement

Carbon-bound H-atoms were placed in calculated positions (C—H 0.95 to 0.98 Å) and were included in the refinement in the riding model approximation, with $U(\text{H})$ set to $1.2\text{--}1.5U_{\text{eq}}(\text{C})$. The ammonium H-atom was similarly treated [N—H 0.88 Å; $U(\text{H}) = 1.2U_{\text{eq}}(\text{N})$]. The final difference Fourier map had a large peak at 1 Å and a deep hole at about 1 Å from the Sn1 atom.

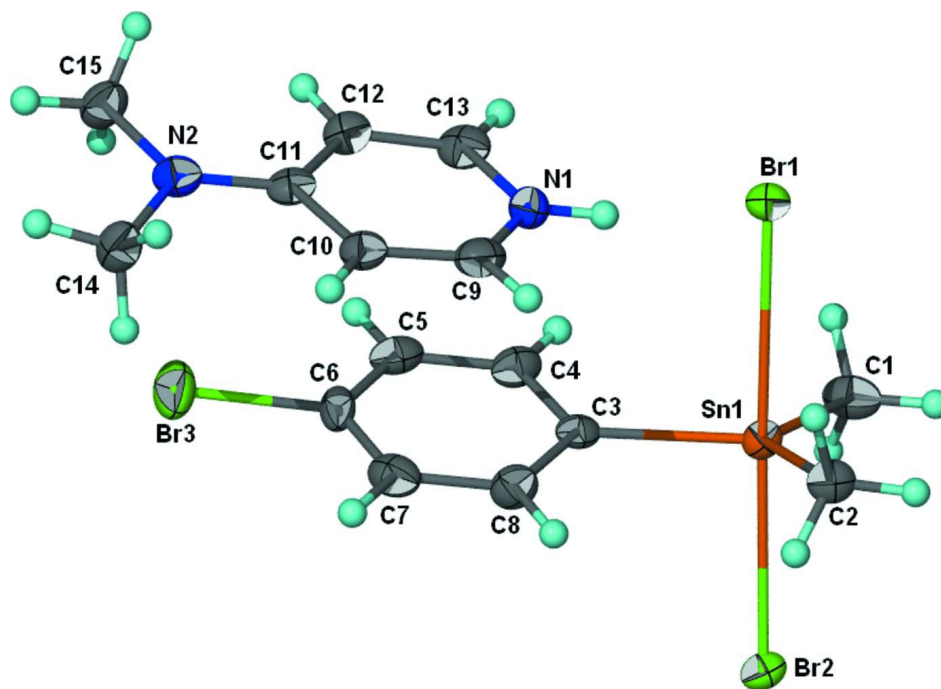


Figure 1

Thermal ellipsoid plot (Barbour, 2001) of $[\text{C}_7\text{H}_{11}\text{N}_2][\text{SnBr}_2(\text{CH}_3)_2(\text{C}_6\text{H}_4\text{Br})]$ at the 70% probability level. Hydrogen atoms are drawn as spheres of arbitrary radius.

4-(Dimethylamino)pyridinium dibromido(4-bromophenyl)dimethylstannate(IV)

Crystal data

(C₇H₁₁N₂)[SnBr₂(CH₃)₂(C₆H₄Br)]

M_r = 587.76

Triclinic, *P* $\bar{1}$

Hall symbol: -P 1

a = 7.3397 (1) Å

b = 11.1034 (2) Å

c = 12.2270 (2) Å

α = 100.038 (1)°

β = 102.472 (1)°

γ = 94.679 (1)°

V = 950.65 (3) Å³

Z = 2

F(000) = 560

D_x = 2.053 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 5297 reflections

θ = 2.3–28.3°

μ = 7.64 mm⁻¹

T = 100 K

Plate, colourless

0.30 × 0.20 × 0.02 mm

Data collection

Bruker SMART APEX

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

T_{min} = 0.208, *T_{max}* = 0.862

8826 measured reflections

4346 independent reflections

3718 reflections with *I* > 2σ(*I*)

R_{int} = 0.029

θ_{\max} = 27.5°, θ_{\min} = 1.7°

h = -9→9

k = -14→14

l = -15→15

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.031

wR(*F*²) = 0.096

S = 1.13

4346 reflections

194 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/[σ²(*F_o*²) + (0.0496*P*)² + 0.4605*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.001

Δρ_{max} = 1.07 e Å⁻³

Δρ_{min} = -1.24 e Å⁻³

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U_{iso}</i> [*] / <i>U_{eq}</i>
Sn1	0.38417 (3)	0.25338 (2)	0.773249 (19)	0.01548 (9)
Br1	0.74942 (5)	0.26874 (4)	0.91216 (3)	0.01853 (11)
Br2	0.02956 (5)	0.24575 (4)	0.63868 (3)	0.01814 (11)
Br3	0.36372 (6)	-0.36881 (4)	0.72001 (4)	0.03322 (13)
N1	0.8504 (5)	0.0064 (3)	0.7756 (3)	0.0221 (7)
H1	0.8438	0.0862	0.7872	0.027*
N2	0.8761 (5)	-0.3667 (3)	0.7213 (2)	0.0199 (7)
C1	0.3068 (6)	0.3599 (4)	0.9145 (3)	0.0251 (9)
H1A	0.1740	0.3357	0.9112	0.038*
H1B	0.3263	0.4475	0.9115	0.038*
H1C	0.3845	0.3456	0.9859	0.038*
C2	0.5034 (5)	0.3149 (4)	0.6446 (3)	0.0198 (8)

H2A	0.4313	0.2714	0.5690	0.030*
H2B	0.6343	0.2976	0.6553	0.030*
H2C	0.4994	0.4037	0.6507	0.030*
C3	0.3607 (5)	0.0584 (4)	0.7558 (3)	0.0164 (7)
C4	0.3542 (5)	0.0007 (4)	0.8482 (3)	0.0208 (8)
H4	0.3511	0.0496	0.9196	0.025*
C5	0.3520 (5)	-0.1260 (4)	0.8384 (3)	0.0227 (8)
H5	0.3462	-0.1637	0.9018	0.027*
C6	0.3585 (5)	-0.1962 (4)	0.7338 (3)	0.0201 (8)
C7	0.3613 (5)	-0.1437 (4)	0.6397 (3)	0.0238 (8)
H7	0.3632	-0.1933	0.5685	0.029*
C8	0.3611 (6)	-0.0170 (4)	0.6514 (3)	0.0217 (8)
H8	0.3611	0.0195	0.5866	0.026*
C9	0.8666 (6)	-0.0529 (4)	0.6733 (3)	0.0237 (8)
H9	0.8744	-0.0076	0.6150	0.028*
C10	0.8721 (5)	-0.1769 (4)	0.6517 (3)	0.0191 (8)
H10	0.8787	-0.2176	0.5778	0.023*
C11	0.8680 (5)	-0.2459 (4)	0.7390 (3)	0.0179 (8)
C12	0.8551 (5)	-0.1788 (4)	0.8470 (3)	0.0205 (8)
H12	0.8542	-0.2199	0.9089	0.025*
C13	0.8440 (5)	-0.0552 (4)	0.8610 (3)	0.0232 (8)
H13	0.8315	-0.0115	0.9325	0.028*
C14	0.8600 (6)	-0.4359 (4)	0.6050 (3)	0.0236 (9)
H14A	0.9550	-0.3986	0.5715	0.035*
H14B	0.7343	-0.4337	0.5581	0.035*
H14C	0.8799	-0.5217	0.6078	0.035*
C15	0.8821 (6)	-0.4367 (4)	0.8132 (3)	0.0244 (9)
H15A	0.9843	-0.3978	0.8788	0.037*
H15B	0.9039	-0.5215	0.7858	0.037*
H15C	0.7622	-0.4377	0.8362	0.037*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Sn1	0.01676 (15)	0.01707 (15)	0.01403 (14)	0.00355 (11)	0.00520 (10)	0.00423 (10)
Br1	0.0172 (2)	0.0208 (2)	0.01645 (19)	0.00236 (15)	0.00153 (14)	0.00369 (15)
Br2	0.0158 (2)	0.0221 (2)	0.01817 (19)	0.00498 (15)	0.00386 (14)	0.00728 (15)
Br3	0.0257 (2)	0.0189 (2)	0.0537 (3)	0.00577 (17)	0.0064 (2)	0.0061 (2)
N1	0.0209 (17)	0.0167 (16)	0.0251 (16)	0.0044 (14)	-0.0002 (14)	0.0005 (13)
N2	0.0215 (17)	0.0251 (18)	0.0133 (14)	0.0041 (14)	0.0044 (13)	0.0033 (13)
C1	0.021 (2)	0.035 (2)	0.0181 (18)	0.0055 (18)	0.0048 (16)	0.0007 (17)
C2	0.0186 (19)	0.025 (2)	0.0178 (17)	0.0023 (16)	0.0069 (15)	0.0064 (15)
C3	0.0104 (17)	0.0211 (19)	0.0173 (17)	0.0010 (14)	0.0017 (14)	0.0047 (15)
C4	0.0190 (19)	0.027 (2)	0.0184 (18)	0.0039 (16)	0.0047 (15)	0.0083 (16)
C5	0.0160 (19)	0.031 (2)	0.0216 (19)	0.0047 (17)	0.0023 (15)	0.0092 (17)
C6	0.0133 (18)	0.0131 (18)	0.033 (2)	0.0031 (14)	0.0044 (16)	0.0037 (16)
C7	0.018 (2)	0.028 (2)	0.0228 (19)	0.0056 (17)	0.0033 (16)	-0.0019 (17)
C8	0.024 (2)	0.024 (2)	0.0183 (17)	0.0051 (16)	0.0055 (15)	0.0061 (16)

C9	0.020 (2)	0.028 (2)	0.0218 (19)	0.0023 (17)	0.0000 (16)	0.0076 (17)
C10	0.0155 (18)	0.024 (2)	0.0183 (17)	0.0041 (15)	0.0034 (15)	0.0038 (15)
C11	0.0113 (17)	0.025 (2)	0.0187 (17)	0.0056 (15)	0.0047 (14)	0.0048 (15)
C12	0.020 (2)	0.024 (2)	0.0188 (18)	0.0052 (16)	0.0057 (15)	0.0048 (16)
C13	0.0167 (19)	0.029 (2)	0.0241 (19)	0.0082 (17)	0.0055 (16)	0.0022 (17)
C14	0.033 (2)	0.022 (2)	0.0182 (18)	0.0088 (18)	0.0084 (17)	0.0027 (16)
C15	0.030 (2)	0.024 (2)	0.0202 (18)	0.0065 (18)	0.0052 (17)	0.0087 (16)

Geometric parameters (Å, °)

Sn1—C1	2.127 (4)	C4—H4	0.9500
Sn1—C2	2.139 (3)	C5—C6	1.390 (6)
Sn1—C3	2.127 (4)	C5—H5	0.9500
Sn1—Br1	2.8211 (4)	C6—C7	1.380 (6)
Sn1—Br2	2.7486 (4)	C7—C8	1.390 (6)
Br3—C6	1.899 (4)	C7—H7	0.9500
N1—C9	1.343 (5)	C8—H8	0.9500
N1—C13	1.351 (5)	C9—C10	1.361 (6)
N1—H1	0.8800	C9—H9	0.9500
N2—C11	1.329 (5)	C10—C11	1.422 (5)
N2—C15	1.469 (5)	C10—H10	0.9500
N2—C14	1.469 (5)	C11—C12	1.426 (5)
C1—H1A	0.9800	C12—C13	1.364 (6)
C1—H1B	0.9800	C12—H12	0.9500
C1—H1C	0.9800	C13—H13	0.9500
C2—H2A	0.9800	C14—H14A	0.9800
C2—H2B	0.9800	C14—H14B	0.9800
C2—H2C	0.9800	C14—H14C	0.9800
C3—C4	1.400 (5)	C15—H15A	0.9800
C3—C8	1.400 (5)	C15—H15B	0.9800
C4—C5	1.390 (6)	C15—H15C	0.9800
C1—Sn1—C2	128.9 (2)	C7—C6—C5	121.7 (4)
C1—Sn1—C3	118.8 (2)	C7—C6—Br3	119.2 (3)
C2—Sn1—C3	112.1 (1)	C5—C6—Br3	119.1 (3)
C1—Sn1—Br2	90.82 (11)	C6—C7—C8	118.7 (3)
C3—Sn1—Br2	92.78 (10)	C6—C7—H7	120.7
C2—Sn1—Br2	89.98 (10)	C8—C7—H7	120.7
C1—Sn1—Br1	88.38 (11)	C7—C8—C3	121.8 (4)
C3—Sn1—Br1	88.88 (10)	C7—C8—H8	119.1
C2—Sn1—Br1	89.39 (10)	C3—C8—H8	119.1
Br1—Sn1—Br2	178.33 (1)	N1—C9—C10	121.1 (4)
C9—N1—C13	120.7 (3)	N1—C9—H9	119.4
C9—N1—H1	119.7	C10—C9—H9	119.4
C13—N1—H1	119.7	C9—C10—C11	120.4 (3)
C11—N2—C15	121.9 (3)	C9—C10—H10	119.8
C11—N2—C14	119.9 (3)	C11—C10—H10	119.8
C15—N2—C14	117.9 (3)	N2—C11—C10	122.0 (3)

Sn1—C1—H1A	109.5	N2—C11—C12	121.5 (3)
Sn1—C1—H1B	109.5	C10—C11—C12	116.5 (4)
H1A—C1—H1B	109.5	C13—C12—C11	119.6 (3)
Sn1—C1—H1C	109.5	C13—C12—H12	120.2
H1A—C1—H1C	109.5	C11—C12—H12	120.2
H1B—C1—H1C	109.5	N1—C13—C12	121.6 (3)
Sn1—C2—H2A	109.5	N1—C13—H13	119.2
Sn1—C2—H2B	109.5	C12—C13—H13	119.2
H2A—C2—H2B	109.5	N2—C14—H14A	109.5
Sn1—C2—H2C	109.5	N2—C14—H14B	109.5
H2A—C2—H2C	109.5	H14A—C14—H14B	109.5
H2B—C2—H2C	109.5	N2—C14—H14C	109.5
C4—C3—C8	117.4 (4)	H14A—C14—H14C	109.5
C4—C3—Sn1	122.0 (3)	H14B—C14—H14C	109.5
C8—C3—Sn1	120.5 (3)	N2—C15—H15A	109.5
C5—C4—C3	121.9 (4)	N2—C15—H15B	109.5
C5—C4—H4	119.1	H15A—C15—H15B	109.5
C3—C4—H4	119.1	N2—C15—H15C	109.5
C4—C5—C6	118.4 (4)	H15A—C15—H15C	109.5
C4—C5—H5	120.8	H15B—C15—H15C	109.5
C6—C5—H5	120.8		
C1—Sn1—C3—C4	-20.4 (4)	C6—C7—C8—C3	0.9 (6)
C2—Sn1—C3—C4	156.0 (3)	C4—C3—C8—C7	-2.1 (6)
Br2—Sn1—C3—C4	-112.9 (3)	Sn1—C3—C8—C7	174.7 (3)
Br1—Sn1—C3—C4	67.2 (3)	C13—N1—C9—C10	1.7 (6)
C1—Sn1—C3—C8	162.9 (3)	N1—C9—C10—C11	-2.4 (6)
C2—Sn1—C3—C8	-20.6 (3)	C15—N2—C11—C10	176.6 (4)
Br2—Sn1—C3—C8	70.5 (3)	C14—N2—C11—C10	-9.5 (6)
Br1—Sn1—C3—C8	-109.5 (3)	C15—N2—C11—C12	-3.3 (6)
C8—C3—C4—C5	1.3 (6)	C14—N2—C11—C12	170.6 (3)
Sn1—C3—C4—C5	-175.5 (3)	C9—C10—C11—N2	-179.0 (4)
C3—C4—C5—C6	0.7 (6)	C9—C10—C11—C12	0.9 (5)
C4—C5—C6—C7	-2.0 (6)	N2—C11—C12—C13	-178.9 (4)
C4—C5—C6—Br3	178.0 (3)	C10—C11—C12—C13	1.2 (5)
C5—C6—C7—C8	1.2 (6)	C9—N1—C13—C12	0.5 (6)
Br3—C6—C7—C8	-178.8 (3)	C11—C12—C13—N1	-1.9 (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—H1...Br1	0.88	2.56	3.319 (3)	146