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Dipyridinium tribromidochloridobis(4-chlorophenyl)stannate(IV)

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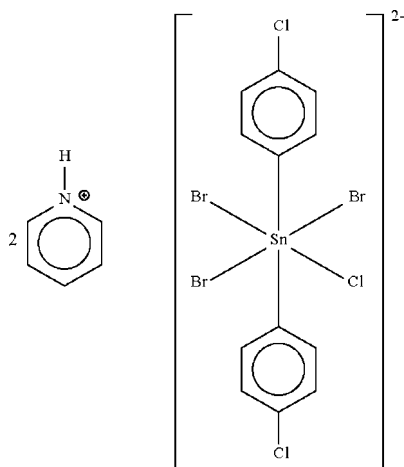
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.018; wR factor = 0.047; data-to-parameter ratio = 19.9.

The tin atom in the substituted ammonium stannate(IV), $(\text{C}_5\text{H}_6\text{N})_2[\text{SnBr}_3(\text{C}_6\text{H}_4\text{Cl})_2\text{Cl}]$, lies on a center of symmetry in a distorted octahedral coordination geometry. Each independent halogen site is occupied by bromine and chlorine anions in an approximate 3:1 ratio. The pyridinium cation forms a hydrogen bond to only one of the halogen atoms.

Related literature

For bis(4-dimethylaminopyridinium) tetrahalidodiorganostannates, see: Lo & Ng (2008*a,b*); Yap *et al.* (2008).



Experimental

Crystal data

$(\text{C}_5\text{H}_6\text{N})_2[\text{SnBr}_3(\text{C}_6\text{H}_4\text{Cl})_2\text{Cl}]$
 $M_r = 777.17$
Monoclinic, $C2/c$
 $a = 11.5130$ (2) Å
 $b = 11.7139$ (2) Å
 $c = 18.7748$ (3) Å
 $\beta = 93.230$ (1)°

$V = 2527.99$ (7) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 6.08$ mm⁻¹
 $T = 100$ K
 $0.27 \times 0.19 \times 0.12$ mm

Data collection

Bruker SMART APEX
diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.327$, $T_{\max} = 0.529$
(expected range = 0.298–0.482)

11728 measured reflections
2903 independent reflections
2668 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.018$
 $wR(F^2) = 0.047$
 $S = 1.02$
2903 reflections
146 parameters

4 restraints
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.39$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.84$ e Å⁻³

Table 1

Selected bond lengths (Å) ($X = \text{Br}, \text{Cl}$).

Sn1—C1	2.149 (2)	Sn1—X2	2.7060 (2)
Sn1—X1	2.7166 (2)		

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINTE* (Bruker, 2007); data reduction: *SAINTE*; 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, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT2945).

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supporting information

Acta Cryst. (2009). E65, m630 [doi:10.1107/S1600536809016687]

Dipyridinium tribromidochloridobis(4-chlorophenyl)stannate(IV)

Kong Mun Lo and Seik Weng Ng

S1. Experimental

Bis(4-chlorophenyl)tin dichloride (0.40 g, 1 mol) and pyridine hydrobromide perbromide (0.64 g, 2 mmol) were heated in chloroform for 3 h. Crystals separated from the cool solution after a day.

S2. Refinement

Hydrogen atoms were placed in calculated positions (C—H 0.95, N—H 0.88 Å) and were included in the refinement in the riding model approximation, with $U(H)$ set to $1.2U(C,N)$.

Each of the two independent tin-bound halogen atoms is a mixture of chlorine and bromine; as the total occupancy of chlorine refined to nearly 0.5 and that of bromine to nearly 1.5, these values were fixed as 0.5 and 1.5. Furthermore, the different halogen atoms sharing the same site were constrained to have the same coordinates and the same anisotropic displacement parameters. The final difference Fourier map did not have large peaks/deep holes near the disordered atoms.

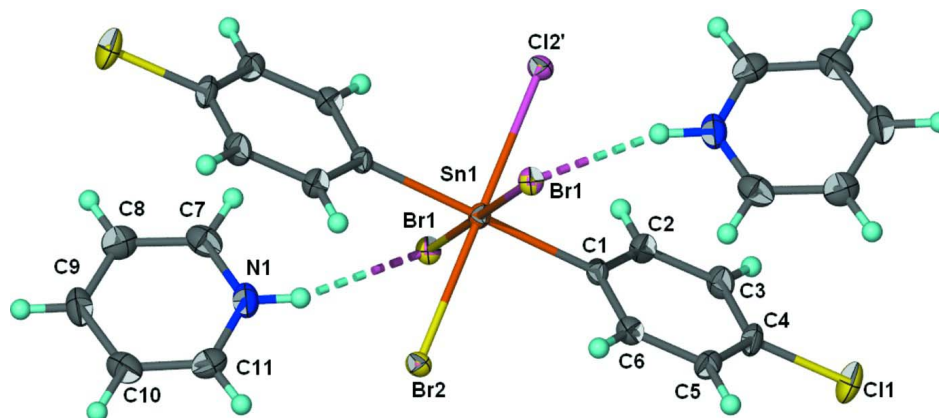


Figure 1

70% Probability anisotropic displacement ellipsoid plot of the ion-pair $2(C_5H_6N)[SnBr_3Cl(C_6H_4Cl)_2]$. Hydrogen atoms are drawn as spheres of arbitrary radius. Dashed lines denote hydrogen bonds. The tin-bound halogen atoms are disordered.

Dipyridinium tribromidochloridobis(4-chlorophenyl)stannate(IV)

Crystal data

$(C_5H_6N)_2[SnBr_3(C_6H_4Cl)_2Cl]$

$M_r = 777.17$

Monoclinic, $C2/c$

Hall symbol: $-C 2yc$

$a = 11.5130(2) \text{ \AA}$

$b = 11.7139(2) \text{ \AA}$

$c = 18.7748(3) \text{ \AA}$

$\beta = 93.230(1)^\circ$

$V = 2527.99(7) \text{ \AA}^3$

$Z = 4$

$F(000) = 1488$
 $D_x = 2.042 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 6664 reflections
 $\theta = 2.5\text{--}28.3^\circ$

$\mu = 6.08 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
 Prism, brown
 $0.27 \times 0.19 \times 0.12 \text{ mm}$

Data collection

Bruker SMART APEX
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.327$, $T_{\max} = 0.529$

11728 measured reflections
 2903 independent reflections
 2668 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.022$
 $\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 2.2^\circ$
 $h = -14 \rightarrow 14$
 $k = -15 \rightarrow 15$
 $l = -23 \rightarrow 24$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.018$
 $wR(F^2) = 0.047$
 $S = 1.02$
 2903 reflections
 146 parameters
 4 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
 $w = 1/[\sigma^2(F_o^2) + (0.0246P)^2 + 3.4843P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.39 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.84 \text{ e \AA}^{-3}$

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Sn1	0.5000	0.5000	0.5000	0.01196 (6)	
Br1	0.308690 (19)	0.509165 (19)	0.578153 (12)	0.01383 (8)	0.7365 (11)
Br2	0.551744 (19)	0.288797 (18)	0.550827 (12)	0.01450 (7)	0.7635 (11)
C11'	0.308690 (19)	0.509165 (19)	0.578153 (12)	0.01383 (8)	0.2635 (11)
C12'	0.551744 (19)	0.288797 (18)	0.550827 (12)	0.01450 (7)	0.2365 (11)
C11	0.83429 (5)	0.71755 (5)	0.76748 (3)	0.02688 (12)	
N1	0.15868 (16)	0.59124 (16)	0.43101 (10)	0.0217 (4)	
H1	0.2206	0.5891	0.4607	0.026*	
C1	0.60325 (16)	0.57206 (16)	0.58772 (10)	0.0129 (4)	
C2	0.60187 (18)	0.52532 (17)	0.65598 (11)	0.0165 (4)	
H2	0.5522	0.4626	0.6644	0.020*	
C3	0.67258 (18)	0.56972 (18)	0.71184 (11)	0.0188 (4)	
H3	0.6716	0.5379	0.7584	0.023*	
C4	0.74442 (17)	0.66121 (18)	0.69830 (11)	0.0182 (4)	
C5	0.74654 (17)	0.70998 (17)	0.63157 (11)	0.0171 (4)	
H5	0.7961	0.7729	0.6234	0.021*	
C6	0.67469 (17)	0.66515 (17)	0.57641 (11)	0.0155 (4)	
H6	0.6745	0.6987	0.5303	0.019*	
C7	0.0884 (2)	0.50083 (18)	0.42776 (13)	0.0225 (5)	
H7	0.1048	0.4366	0.4575	0.027*	
C8	-0.0077 (2)	0.50100 (18)	0.38125 (13)	0.0235 (5)	

H8	-0.0579	0.4366	0.3777	0.028*
C9	-0.03042 (19)	0.5968 (2)	0.33953 (12)	0.0238 (5)
H9	-0.0971	0.5988	0.3074	0.029*
C10	0.0440 (2)	0.68937 (19)	0.34465 (12)	0.0231 (5)
H10	0.0290	0.7552	0.3161	0.028*
C11	0.13982 (19)	0.68517 (19)	0.39135 (12)	0.0224 (5)
H11	0.1920	0.7479	0.3955	0.027*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Sn1	0.01296 (10)	0.01329 (10)	0.00948 (10)	-0.00098 (6)	-0.00070 (7)	0.00034 (7)
Br1	0.01297 (12)	0.01610 (12)	0.01249 (13)	-0.00044 (8)	0.00142 (9)	0.00093 (8)
Br2	0.01749 (12)	0.01242 (11)	0.01334 (12)	0.00098 (8)	-0.00136 (8)	0.00188 (8)
Cl1'	0.01297 (12)	0.01610 (12)	0.01249 (13)	-0.00044 (8)	0.00142 (9)	0.00093 (8)
Cl2'	0.01749 (12)	0.01242 (11)	0.01334 (12)	0.00098 (8)	-0.00136 (8)	0.00188 (8)
Cl1	0.0254 (3)	0.0349 (3)	0.0193 (3)	-0.0073 (2)	-0.0073 (2)	-0.0056 (2)
N1	0.0177 (9)	0.0277 (10)	0.0190 (9)	0.0050 (7)	-0.0038 (7)	-0.0047 (8)
C1	0.0126 (9)	0.0155 (9)	0.0103 (9)	0.0020 (7)	-0.0011 (7)	-0.0014 (7)
C2	0.0160 (9)	0.0177 (9)	0.0156 (10)	-0.0017 (7)	0.0004 (8)	-0.0001 (8)
C3	0.0221 (10)	0.0222 (10)	0.0119 (9)	0.0010 (8)	-0.0016 (8)	0.0022 (8)
C4	0.0153 (10)	0.0237 (10)	0.0150 (10)	0.0002 (8)	-0.0041 (8)	-0.0054 (8)
C5	0.0155 (9)	0.0170 (9)	0.0188 (10)	-0.0028 (7)	0.0002 (8)	-0.0013 (8)
C6	0.0154 (9)	0.0170 (9)	0.0142 (10)	0.0012 (7)	0.0018 (7)	0.0010 (8)
C7	0.0251 (12)	0.0223 (11)	0.0204 (12)	0.0064 (8)	0.0055 (9)	0.0023 (9)
C8	0.0194 (11)	0.0239 (11)	0.0276 (13)	-0.0015 (8)	0.0062 (9)	-0.0024 (9)
C9	0.0173 (10)	0.0332 (12)	0.0204 (11)	0.0049 (9)	-0.0028 (8)	-0.0028 (9)
C10	0.0280 (11)	0.0214 (10)	0.0201 (11)	0.0058 (9)	0.0029 (9)	0.0019 (9)
C11	0.0251 (11)	0.0196 (10)	0.0225 (11)	-0.0018 (8)	0.0034 (9)	-0.0048 (9)

Geometric parameters (Å, °)

Sn1—C1 ⁱ	2.149 (2)	C3—C4	1.386 (3)
Sn1—C1	2.149 (2)	C3—H3	0.9500
Sn1—Br1	2.7166 (2)	C4—C5	1.378 (3)
Sn1—Cl2 ^{ri}	2.7060 (2)	C5—C6	1.391 (3)
Sn1—Br2 ⁱ	2.7060 (2)	C5—H5	0.9500
Sn1—Br2	2.7060 (2)	C6—H6	0.9500
Sn1—Cl1 ^{ri}	2.7166 (2)	C7—C8	1.370 (3)
Sn1—Br1 ⁱ	2.7166 (2)	C7—H7	0.9500
Cl1—C4	1.744 (2)	C8—C9	1.384 (3)
N1—C7	1.332 (3)	C8—H8	0.9500
N1—C11	1.339 (3)	C9—C10	1.383 (3)
N1—H1	0.8800	C9—H9	0.9500
C1—C6	1.389 (3)	C10—C11	1.371 (3)
C1—C2	1.394 (3)	C10—H10	0.9500
C2—C3	1.392 (3)	C11—H11	0.9500
C2—H2	0.9500		

C1 ⁱ —Sn1—C1	180.000 (1)	C6—C1—Sn1	119.86 (14)
C1 ⁱ —Sn1—Cl2 ⁱ	89.29 (5)	C2—C1—Sn1	121.05 (14)
C1—Sn1—Cl2 ⁱ	90.71 (5)	C3—C2—C1	120.63 (19)
C1 ⁱ —Sn1—Br2 ⁱ	89.29 (5)	C3—C2—H2	119.7
C1—Sn1—Br2 ⁱ	90.71 (5)	C1—C2—H2	119.7
Cl2 ⁱ —Sn1—Br2 ⁱ	0.000 (13)	C4—C3—C2	118.71 (19)
C1 ⁱ —Sn1—Br2	90.71 (5)	C4—C3—H3	120.6
C1—Sn1—Br2	89.29 (5)	C2—C3—H3	120.6
Cl2 ⁱ —Sn1—Br2	180.0	C5—C4—C3	121.90 (19)
Br2 ⁱ —Sn1—Br2	180.0	C5—C4—Cl1	118.68 (16)
C1 ⁱ —Sn1—Cl1 ⁱ	90.05 (5)	C3—C4—Cl1	119.41 (17)
C1—Sn1—Cl1 ⁱ	89.95 (5)	C4—C5—C6	118.67 (19)
Cl2 ⁱ —Sn1—Cl1 ⁱ	90.845 (7)	C4—C5—H5	120.7
Br2 ⁱ —Sn1—Cl1 ⁱ	90.845 (7)	C6—C5—H5	120.7
Br2—Sn1—Cl1 ⁱ	89.155 (7)	C5—C6—C1	120.98 (19)
C1 ⁱ —Sn1—Br1 ⁱ	90.05 (5)	C5—C6—H6	119.5
C1—Sn1—Br1 ⁱ	89.95 (5)	C1—C6—H6	119.5
Cl2 ⁱ —Sn1—Br1 ⁱ	90.845 (7)	N1—C7—C8	119.7 (2)
Br2 ⁱ —Sn1—Br1 ⁱ	90.845 (7)	N1—C7—H7	120.2
Br2—Sn1—Br1 ⁱ	89.155 (7)	C8—C7—H7	120.2
Cl1 ⁱ —Sn1—Br1 ⁱ	0.000 (8)	C7—C8—C9	118.8 (2)
C1 ⁱ —Sn1—Br1	89.95 (5)	C7—C8—H8	120.6
C1—Sn1—Br1	90.05 (5)	C9—C8—H8	120.6
Cl2 ⁱ —Sn1—Br1	89.155 (7)	C8—C9—C10	120.0 (2)
Br2 ⁱ —Sn1—Br1	89.155 (7)	C8—C9—H9	120.0
Br2—Sn1—Br1	90.845 (7)	C10—C9—H9	120.0
Cl1 ⁱ —Sn1—Br1	180.0	C11—C10—C9	119.3 (2)
Br1 ⁱ —Sn1—Br1	180.0	C11—C10—H10	120.3
C7—N1—C11	123.26 (19)	C9—C10—H10	120.3
C7—N1—H1	118.4	N1—C11—C10	119.0 (2)
C11—N1—H1	118.4	N1—C11—H11	120.5
C6—C1—C2	119.08 (18)	C10—C11—H11	120.5
Cl2 ⁱ —Sn1—C1—C6	-40.72 (15)	C1—C2—C3—C4	-0.1 (3)
Br2 ⁱ —Sn1—C1—C6	-40.72 (15)	C2—C3—C4—C5	0.9 (3)
Br2—Sn1—C1—C6	139.28 (15)	C2—C3—C4—Cl1	-179.64 (16)
Cl1 ⁱ —Sn1—C1—C6	50.12 (15)	C3—C4—C5—C6	-0.4 (3)
Br1 ⁱ —Sn1—C1—C6	50.12 (15)	Cl1—C4—C5—C6	-179.89 (15)
Br1—Sn1—C1—C6	-129.88 (15)	C4—C5—C6—C1	-0.9 (3)
Cl2 ⁱ —Sn1—C1—C2	140.70 (15)	C2—C1—C6—C5	1.7 (3)
Br2 ⁱ —Sn1—C1—C2	140.70 (15)	Sn1—C1—C6—C5	-176.91 (15)
Br2—Sn1—C1—C2	-39.30 (15)	C11—N1—C7—C8	1.0 (3)
Cl1 ⁱ —Sn1—C1—C2	-128.46 (15)	N1—C7—C8—C9	-1.2 (3)
Br1 ⁱ —Sn1—C1—C2	-128.46 (15)	C7—C8—C9—C10	0.8 (3)
Br1—Sn1—C1—C2	51.54 (15)	C8—C9—C10—C11	-0.1 (3)

C6—C1—C2—C3	-1.2 (3)	C7—N1—C11—C10	-0.3 (3)
Sn1—C1—C2—C3	177.39 (15)	C9—C10—C11—N1	-0.1 (3)

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

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—H1...Br1	0.88	2.55	3.317 (2)	146