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Key indicators

Single-crystal X-ray study

T = 100 K

Mean $\sigma(C-C) = 0.002 \text{ \AA}$

R factor = 0.033

wR factor = 0.071

Data-to-parameter ratio = 33.4

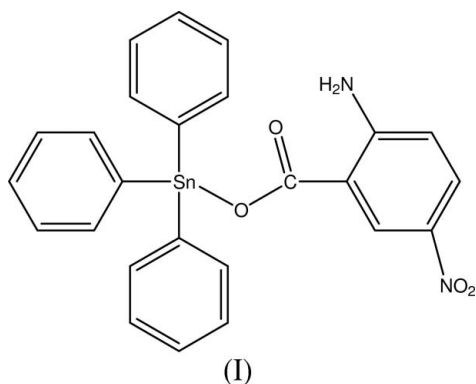
For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

(2-Amino-5-nitrobenzoato)triphenyltin(IV)

In the title compound, $[\text{Sn}(\text{C}_6\text{H}_5)_3(\text{C}_7\text{H}_5\text{N}_2\text{O}_4)]$, the nitro and amino groups are coplanar with the attached benzene ring. The molecules are linked by intermolecular $\text{N}-\text{H}\cdots\text{O}$ interactions, forming one-dimensional chains along $[010]$. In addition, the crystal structure is further stabilized by intermolecular $\text{C}-\text{H}\cdots\pi$ and $\pi-\pi$ interactions.

Comment

Triorganotin(IV) complexes are reported to exhibit antimicrobial and antitumour properties as well as being noted for their significance in industrial applications (Willem *et al.*, 1997; Novelli *et al.*, 1999; Gielen *et al.*, 2000). In general, triphenyltin(IV) carboxylate complexes exist as monomeric structures with four-coordinate distorted tetrahedral or five-coordinate trigonal-bipyramidal geometries (Baul *et al.*, 2001; Yeap & Teoh, 2003; Win, Guan, Ismail *et al.*, 2006). However, (2-chloro-5-aminobenzoato)triphenyltin(IV) was found to exist as a five-coordinate polymeric structure with a linkage of the amino group to an adjacent tin atom and bridging of the carboxylate anion to two neighbouring Sn atoms (Yeap & Teoh, 2003). However, the title compound, (2-amino-5-nitrobenzoato)triphenyltin(IV), (I), is found to exist as a monomeric structure even though the amino group is attached in the *ortho*-position to the benzoate group, as is the case in (2-amino-5-nitrobenzoato)triphenyltin(IV).



Bond lengths and angles in (I) (Fig. 1) have normal values (Allen *et al.*, 1987) and agree well with those found in related structures (Ng *et al.*, 1991; Win, Guan, Ismail *et al.*, 2006; Win, Guan & Yamin, 2006). The nitro group at C4 and amino group at C1 are coplanar with the attached ring, with torsion angles $\text{O}3-\text{N}2-\text{C}4-\text{C}3 = 2.0 (2)^\circ$ and $\text{N}1-\text{C}1-\text{C}6-\text{C}7 = -0.2 (2)^\circ$.

The intramolecular $\text{C}5-\text{H}5\text{A}\cdots\text{O}1$, $\text{N}1-\text{H}1\text{B}\cdots\text{O}2$ and $\text{C}9-\text{H}9\text{A}\cdots\text{O}2$ interactions (Table 2 and Fig. 1) generate

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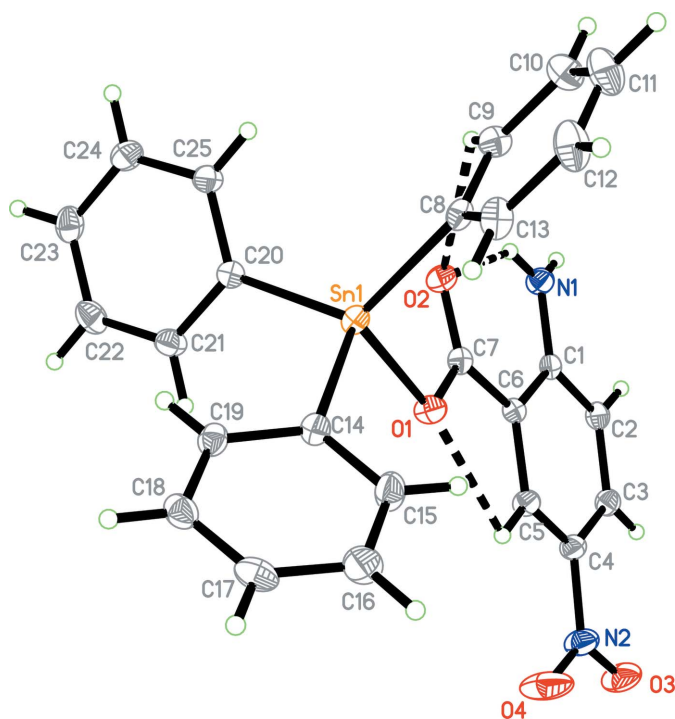


Figure 1
The molecular structure of (I), showing 50% probability displacement ellipsoids and the atomic numbering. The dashed lines indicate intramolecular hydrogen bonds.

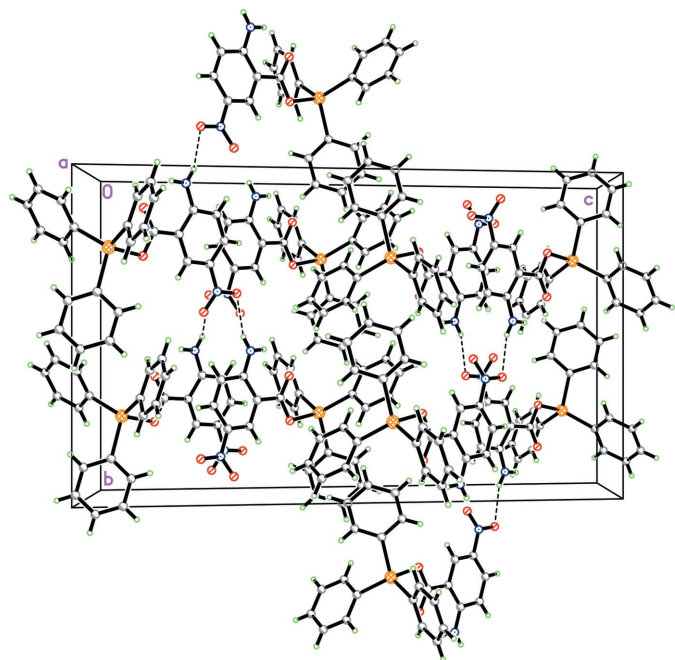


Figure 2
The crystal packing of (I), viewed down the *a* axis. Hydrogen bonds are shown as dashed lines.

S(5), *S*(6) and *S*(7) ring motifs, respectively (Bernstein *et al.*, 1995). In the crystal structure, the molecules form a one-dimensional chain along the [010] direction by intermolecular N1—H1A···O3ⁱ interactions (Fig. 2). The crystal packing is

stabilized by intermolecular C—H··· π interactions involving the C1—C6 (centroid *Cg*1), C8—C13 (centroid *Cg*2), C14—C19 (centroid *Cg*3) and C20—C25 (centroid *Cg*4) benzene rings (Table 2). In addition, the crystal packing is further stabilized by weak intermolecular π — π interactions involving the C1—C6 ring at (*x*, *y*, *z*) and its symmetry equivalent at ($-x$, *y*, $\frac{1}{2} - z$); the centroid—centroid distance is 3.7260 (9) Å.

Experimental

The complex (2-amino-5-nitrobenzoato)triphenyltin(IV) was obtained by heating under reflux a 1:1 molar mixture of triphenyltin(IV) hydroxide (0.73 g, 2 mmol) and 2-amino-5-nitrobenzoic acid (0.36 g, 2 mmol) in methanol (60 ml) for an hour. A clear yellowish transparent solution was isolated by filtration and kept in a bottle. After 6 days, some yellowish crystals (0.51 g, 96% yield) were obtained (m.p. 481.5–481.9 K. Analysis found for C₂₅H₂₀N₂O₄Sn₁: C, 56.41; H, 3.48; N, 5.23; Sn, 22.03%. Calculated for C₂₅H₂₀N₂O₄Sn₁: C, 56.53; H, 3.80; N, 5.27; Sn, 22.35%. FTIR from a KBr disc (cm⁻¹): $\nu(\text{COO})_{\text{as}}$ 1618, $\nu(\text{COO})_{\text{s}}$ 1310, $\nu(\text{O—Sn—O})$ 641, $\nu(\text{Sn—O})$ 443. ¹³C-NMR: δ : phenyl carbons *C*_{ipso} 142.87 (839.31 Hz), *C*_{ortho} 136.11 (45.60 Hz), *C*_{meta} 128.43, *C*_{para} 129.03 (18.37 Hz); benzene 112.77, 115.75, 127.72, 127.97, 134.89, 155.92; COO 169.52 p.p.m.. ¹¹⁹Sn-NMR: δ : -265.89 p.p.m..

Crystal data

[Sn(C₆H₅)₃(C₇H₅N₂O₄)]
M_r = 531.12
 Monoclinic, *C*2/*c*
a = 11.0062 (4) Å
b = 15.8022 (4) Å
c = 26.5645 (8) Å
 β = 101.415 (2)°
V = 4528.8 (2) Å³

Z = 8
D_x = 1.558 Mg m⁻³
 Mo *K* α radiation
 μ = 1.16 mm⁻¹
T = 100.0 (1) K
 Block, yellow
 0.76 × 0.73 × 0.40 mm

Data collection

Bruker SMART APEX2 CCD area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2005)
*T*_{min} = 0.482, *T*_{max} = 0.657

47389 measured reflections
 9913 independent reflections
 9343 reflections with *I* > 2 σ (*I*)
*R*_{int} = 0.029
 θ_{max} = 35.0°

Refinement

Refinement on *F*²
R [*F*² > 2 σ (*F*²)] = 0.033
 wR (*F*²) = 0.071
S = 1.24
 9913 reflections
 297 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0165P)^2 + 9.0039P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.50 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -1.18 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Sn1—O1	2.0774 (11)	Sn1—C8	2.1230 (15)
Sn1—C20	2.1224 (15)	Sn1—C14	2.1264 (15)
O1—Sn1—C20	108.88 (5)	O1—Sn1—C14	97.53 (5)
O1—Sn1—C8	106.21 (5)	C20—Sn1—C14	109.30 (6)
C20—Sn1—C8	120.75 (5)		

Table 2
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—H1B···O2	0.84 (2)	2.03 (2)	2.681 (2)	134 (2)
C5—H5A···O1	0.93	2.40	2.725 (2)	101
C9—H9A···O2	0.93	2.46	3.073 (2)	124
N1—H1A···O3 ⁱ	0.83 (3)	2.25 (3)	3.058 (2)	162 (2)
C3—H3A···Cg2 ⁱⁱ	0.93	2.77	3.616 (2)	152
C12—H12A···Cg1 ⁱⁱⁱ	0.93	3.13	3.823 (2)	133
C13—H13A···Cg4 ^{iv}	0.93	3.10	3.661 (2)	120
C17—H17A···Cg4 ^v	0.93	3.15	3.804 (2)	129
C19—H19A···Cg2 ^{iv}	0.93	3.00	3.681 (2)	131
C25—H25A···Cg3 ^{iv}	0.93	2.88	3.544 (2)	129

Symmetry codes: (i) $-x - \frac{1}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$; (ii) $-x, y, -z + \frac{1}{2}$; (iii) $x + 1, y, z$; (iv) $-x + \frac{1}{2}, -y + \frac{1}{2}, -z$; (v) $x + \frac{1}{2}, y + \frac{1}{2}, z$. Cg1, Cg2, Cg3 and Cg4 are the centroids of the C1–C6, C8–C13, C14–C19 and C20–C25 rings, respectively.

The H atoms on N were located in a difference map and refined isotropically, with N—H = 0.84 (3) Å. The remaining H atoms were positional geometrically and refined as riding, with C—H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The deepest hole in the final difference map is 0.61 Å from atom Sn1.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINTE* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

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