

CRYSTAL STRUCTURE OF TRIBUTYLTIN *N*-PHTHALOYLLEUCINATEKong Mun Lo¹ and Seik Weng Ng²¹Department of Chemistry ²Institute of Postgraduate Studies and Research
University of Malaya, 50603 Kuala Lumpur, Malaysia**Abstract**

The two symmetry-independent molecules of tributyltin *N*-phthaloylleucinate are linked by long tin-amide bridges [Sn-O = 2.897(7), 2.960(7) Å] into a linear chain that propagates along the *a*-*c* diagonal of the monoclinic cell. The tin atoms are five-coordinate in *trans*-C₃SnO₂ trigonal bipyramidal geometries.

Discussion

The tributyltin derivative of *N*-phthaloylglycine exists as two symmetry-independent hydrates, and the coordinated water molecules engage in hydrogen bonding interactions that consolidate the crystal structure [1]. The incorporation of an *i*-propyl group in the protected amino acid appears to render the tin atom less Lewis acidic in the resulting carboxylate; unlike trialkyltin carboxylates that generally are linked by strong carboxylate bridges [2,3], the molecules of tributyltin *N*-phthaloylleucinate are connected by a long tin-amide interaction into a linear chain. A similar long interaction is also found in tricyclohexyltin *N*-phthaloylglycinate [4].

Table 1. Crystal data for tributyltin *N*-phthaloylleucinate.

Formula	C ₂₆ H ₄₁ NO ₄ Sn
Crystal system	Monoclinic
Formula weight	550.29
Space group	<i>P</i> 2 ₁
Cell constants	<i>a</i> = 10.609(1), <i>b</i> = 23.023(1), <i>c</i> = 11.951(1) Å; β = 101.73(1)°
<i>Z</i>	4
Temperature	25°C
Density (calculated)	1.279 g cm ⁻³
μ	0.921 mm ⁻¹
<i>F</i> (000)	1144
Diffractionmeter	CAD-4 (λ = 0.71073 Å)
Theta range for data collection	1.74 to 24.98°
Index ranges	0 ≤ <i>h</i> ≤ 12, -27 ≤ <i>k</i> ≤ 0, -14 ≤ <i>l</i> ≤ 13
Total reflections measured	5455
Unique reflections measured	5166 (<i>R</i> _{int} = 0.028)
Transmission factors	0.724 - 0.803
Data / restraints / parameters	5166 / 246 / 579
Reflections with <i>I</i> > 2σ(<i>I</i>)	4278
<i>R</i> (<i>wR</i>) for <i>I</i> > 2σ(<i>I</i>) data	0.042 (0.110)
<i>R</i> (<i>wR</i>) for all data	0.057 (0.121)
Goodness-of-fit	1.039
ρ, e Å ⁻³	- 0.371 to 495
Programs used	SHELXS-97 [6] SHELXL-97 [7]
Deposition number	CCDC

Experimental

N-Phthaloylleucinate was synthesized by melting phthalic anhydride with leucine according to the procedure for preparing *N*-phthaloylglycine [5]. The acid and bis(tributyltin) oxide (2:1 molar stoichiometry) were heated in a small volume of ethanol to afford the organotin carboxylate, which was isolated and recrystallized from ethanol. A specimen, 0.22 x 0.14 x 0.14 mm, was used in the diffraction measurements. The refinement of the crystal structure necessitated imposing several restraints as the butyl groups in molecule *a* were disordered. Each butyl group was refined as two butyl groups sharing a common C_α atom. Distances in all butyl groups and in the two *i*-propyl groups were restrained (C-C = 1.54±0.01; C[⋯]C[⋯]C = 2.52±0.02 Å, and the temperature factors in these groups were restrained to be nearly isotropic by an *ISOR* 0.02 command.

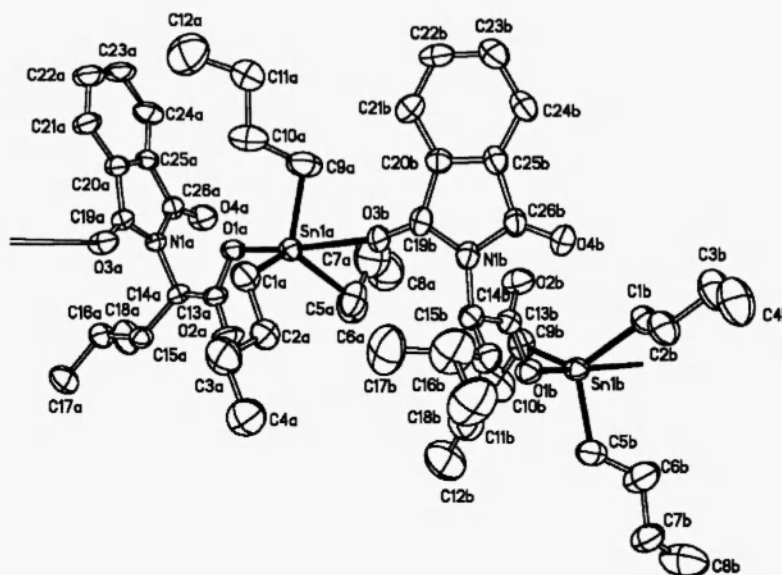


Figure 1. ORTEP [8] plot of tributyltin *N*-phthaloylleucinate. Selected bond distances and angles: Sn1a-C1a = 2.14(1), Sn1a-C5a = 2.13(1), Sn1a-C9a = 2.14(1), Sn1a-O1a = 2.068(6), Sn1a-O3b = 2.960(7), Sn1b-C1b = 2.14(1), Sn1b-C5b = 2.10(1), Sn1b-C9b = 2.12(1), Sn1b-O1b = 2.139(7), Sn1b-O3a' = 2.897(7) Å; C1a-Sn1a-C5a = 119.0(5), C1a-Sn1a-C9a = 115.7(6), C1a-Sn1a-O1a = 102.7(4), C1a-Sn1a-O3b = 79.2(4), C5a-Sn1a-C9a = 115.3(6), C5a-Sn1a-O1a = 105.2(4), C5a-Sn1a-O3b = 77.2(4), C9a-Sn1a-O1a = 93.4(4), C9a-Sn1a-O3b = 81.9(4), O1a-Sn1a-O3b = 175.3(2), C1b-Sn1b-C5b = 114.3(6), C1b-Sn1b-C9b = 123.5(5), C1b-Sn1b-O1b = 100.7(4), C1b-Sn1b-O3a' = 76.9(3), C5b-Sn1b-C9b = 115.7(7), C5b-Sn1b-O1b = 91.6(5), C5b-Sn1b-O3a' = 89.0(4), C9b-Sn1b-O1b = 102.4(4), C9b-Sn1b-O3a' = 79.4(4), O1b-Sn1b-O3a' = 177.5(3)°. Translational code: $i = 1 + x, y, 1 + z$.

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