metal-organic papers

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Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.009 Å Disorder in main residue R factor = 0.083 wR factor = 0.262 Data-to-parameter ratio = 13.2

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

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The molecular structure of the tetrahydrofuran adduct of the lithium salt of the primary phosphane DnpPH₂ (Dnp = 2,6-di-1-naphthylphenyl) of composition [Li(THF){P(H)Dnp}-(THF)]₂ is reported. The dimeric and centrosymmetric complex, [Li₂(C₄H₈O)₂(C₂₆H₂₂OP)₂], features a three-coordinate Li atom in a slightly pyramidal coordination environment, with the sum of angles around it equal to 352.0° and a P atom in a highly distorted tetrahedral coordination environ-

bis[(tetrahydrofuran)lithium(I)]

Bis[µ-bis(2,6-di-1-naphthylphenyl)phosphanido]-

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Comment

ment.

Lithium phosphanide compounds have been known for decades and have been used extensively for the synthesis of a variety of metal complexes (Izod, 2000; Rabe *et al.*, 2001). Thus nowadays both primary and secondary lithium phosphanides are a well investigated class of compounds and several examples have been structurally characterized. We here report the first example of an alkali metal complex of the novel chiral (racemic) terphenyl-based phosphane DnpH₂ (DnpPH₂ = 2,6-di-1-naphthylphenyl). The terphenyl-like moiety adopts the chiral D,L form.



The central four-membered ring is planar, with Li···Li and P···P distances of 3.24 (2) and 3.778 (3) Å, and angles at P1 and Li1 of 81.2 (4) and 98.8 (4)°, respectively. Atom Li1 was found to be close to atoms C18ⁱ, C19ⁱ and C20ⁱ with distances of 3.04 (1), 2.81 (1) and 3.07 (1) Å, respectively [symmetry code: (i) 1 - x, 1 - y, 1 - z]. This may indicate the presence of weak allyl-like electrostatic interactions between the Li atom and these atoms. The naphthyl ring systems are tilted with respect to the central benzene ring, with dihedral angles of 78.2 (2)° for C7–C16 and 75.5 (2)° for C17–C26. Compound (I) was found to be isostructural with its sodium analogue (Rabe, 2004).

Experimental

DnpPH₂ (Rabe, 2004) was prepared following the published procedures for ${}^{t}Bu_{3}C_{6}H_{2}PH_{2}$ (Cowley *et al.*, 1990) and DmpPH₂ (Urnezius & Protasiewicz, 1996). Li(THF)P(H)Dnp was prepared by reaction of equimolar amounts of DnpPH₂ and *n*-BuLi in tetrahydrofuran at

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Figure 1

The molecular structure of the title compound, showing the atomlabelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms, except for those of the the P-H groups, have been omitted for clarity. Dashed lines are used to highlight the disorder with respect to C28, C29 and C30. Except for THF atoms, all atoms labelled as A were generated using the symmetry code (1 - x, 1 - y, 1 - z).

195 K and worked up following standard laboratory procedures. Single crystalline material of the title compound was obtained from a toluene solution at 248 K.

Crystal data

$[\text{Li}_{2}(\text{C}_{4}\text{H}_{8}\text{O})_{2}(\text{C}_{26}\text{H}_{22}\text{OP})_{2}]$ $M_{r} = 880.84$	Z = 1 $D_x = 1.247 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 10.4586 (19) Å	Cell parameters from 700
b = 11.3387 (19) Å	reflections
c = 11.5813 (19) Å	$\theta = 2.0-25.0^{\circ}$
$\alpha = 62.710 \ (10)^{\circ}$	$\mu = 0.14 \text{ mm}^{-1}$
$\beta = 74.360 \ (11)^{\circ}$	T = 173 (2) K
$\gamma = 86.041 \ (11)^{\circ}$	Block, red
$V = 1172.9 (4) \text{ Å}^3$	$0.30\times0.15\times0.15~\text{mm}$

Data collection

diffractometer2126 reflections with $I > 2\sigma(I)$ str φ and ω scans $R_{int} = 0.067$ SH Absorption correction: multi-scan $\theta_{max} = 25.0^{\circ}$ SH $(SADABS; Bruker, 2001)$ $h = -12 \rightarrow 12$ SH $T_{min} = 0.940, T_{max} = 0.980$ $k = -13 \rightarrow 13$ $I = -12 \rightarrow 13$	Bruker SMART CCD area-detector	4117 independent reflections	stru
$ \begin{array}{ll} \varphi \mbox{ and } \omega \mbox{ scans } & R_{\rm int} = 0.067 & SH \\ \mbox{ Absorption correction: multi-scan } & \theta_{\rm max} = 25.0^{\circ} & SH \\ \mbox{ (SADABS; Bruker, 2001) } & h = -12 \rightarrow 12 \\ T_{\rm min} = 0.940, \ T_{\rm max} = 0.980 & k = -13 \rightarrow 13 \\ \mbox{ 9754 measured reflections } & l = -12 \rightarrow 13 \end{array} $	diffractometer	2126 reflections with $I > 2\sigma(I)$	stru
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$(SADABS; Bruker, 2001) h = -12 \rightarrow 12 T_{min} = 0.940, T_{max} = 0.980 k = -13 \rightarrow 13 9754 measured reflections l = -12 \rightarrow 13 $	Absorption correction: multi-scan	$\theta_{\rm max} = 25.0^{\circ}$	SH
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IN	9754 measured reflections	$l = -12 \rightarrow 13$	D (

Refinement

 $\begin{array}{ll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.1225P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.083 & w e 1/[\sigma^2(F_o^2) + (0.1225P)^2 \\ + 0.6192P] & where \ P = (F_o^2 + 2F_c^2)/3 \\ S = 1.06 & (\Delta/\sigma)_{\rm max} < 0.001 \\ 4117 \ reflections & \Delta\rho_{\rm max} = 0.67 \ e \ {\rm \AA}^{-3} \\ 312 \ parameters & \Delta\rho_{\rm min} = -0.26 \ e \ {\rm \AA}^{-3} \\ H \ atoms \ treated \ by \ a \ mixture \ of \\ independent \ and \ constrained \\ refinement & \end{array}$



Figure 2

The contents of the unit cell, viewed along the a axis. All H atoms have been omitted for clarity.

Table 1

Selected geometric parameters (Å, °).

Li1-01	1.867 (11)	$P1-P1^{i}$	3.778 (3)
Li1-P1	2.483 (11)	P1-C1	1.795 (5)
Li1–P1 ⁱ	2.493 (9)	P1-H1	1.34 (6)
Li1···Li1 ⁱ	3.24 (2)		
O1-Li1-P1	131.0 (4)	$P1-Li1-P1^{i}$	98.8 (4)
O1-Li1-P1 ⁱ	122.2 (5)	Li1-P1-Li1 ⁱ	81.2 (4)

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

The high values of the residuals *R* and *wR* can be attributed to the poor quality of the crystal that was used. Atoms C28, C29 and C30 of the THF ring were found to be disordered over two positions with occupancy 0.62 (2)/0.38 (3). The C-C distances in the THF ring were restrained and the displacement parameters for each disordered pair (C28/C28A, C29/C29A and C30/C30A) were set equal to each other. The phosphorus H atom was found in a Fourier difference map and was refined freely. All other H atoms were placed in calculated positions (C-H = 0.95 and 0.99 Å), with isotropic displacement parameters fixed at 1.2 or 1.5 times U_{eq} of the parent atom and were refined as riding atoms.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SMART*; data reduction: *SHELXTL* (Bruker, 2001); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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