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Butyldiphenylphosphine oxide

Judy Caddy,^a E. Mabel Coyanis,^b Andreas Lemmerer,^b Setshaba D. Khanye^b and Bernard Omondi^b*

^aProject AuTEK, Mintek, Private Bag X3015, Randburg 2125, South Africa, and ^bMolecular Science Institute, School of Chemistry, University of the Witwatersrand, PO Wits, 2050 Johannesburg, South Africa

Correspondence e-mail: bernard@chem.wits.ac.za

Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.003 Å R factor = 0.031 wR factor = 0.084 Data-to-parameter ratio = 15.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The solid state structure of the title compound, $C_{16}H_{19}OP$, an organic tertiary phosphine oxide, is stabilized mainly by $C-H\cdots\pi$ intermolecular interactions.

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Comment

There has been considerable interest in studying phosphine oxides as ligands, particularly as chelating ligands for complex formation with transition metals. The title compound, (I) (Fig. 1), has the expected deformed tetrahedral arrangement around the phosphorus atom, with C-P-C angles smaller than tetrahedral and O-P-C angles larger than tetrahedral values (Table 1), and also with a normal P=O bond distance (Bye *et al.*, 1982). The dihedral angle between the mean weighted least-squares planes of the phenyl rings (atoms C5-C10 and C11-C16) is 70.62 (6)°.



In the crystal structure, molecules of (I) are connected through three $C-H\cdots\pi$ intermolecular interactions (Table 2), resulting in a two-dimensional network parallel to the *ac*



© 2007 International Union of Crystallography All rights reserved Figure 1 The molecular s

The molecular structure of (I), showing the atomic numbering scheme and 50% probability displacement ellipsoids.

plane. Atom C2 in the molecule at (x, y, z) acts a donor atom *via* H2A and H2B to the phenyl rings in the molecule at (1 + x, y, z), thereby generating a chain along the *a* axis. The third C-H··· π interaction is generated *via* atom H14 of the molecule at (x, y, z) to the C5-C10 phenyl ring of the molecule at $(-\frac{1}{2} + x, 1 - y, \frac{1}{2} + z)$ (Fig. 2).

Experimental

The title compound (the oxidized form of butyldiphenylphosphine) was serendipitously obtained upon exposure of the unoxidized ligand to air, during the attempted crystallization of an organometallic compound by slow evaporation from hexane.

Z = 2

Crystal data

 $C_{16}H_{19}OP$ $M_r = 258.28$ Monoclinic, *Pn* a = 8.1289 (12) Å b = 5.9814 (9) Å c = 14.600 (2) Å $\beta = 93.326$ (3)° V = 708.68 (18) Å³

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: none 4597 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.031$ $wR(F^2) = 0.084$ S = 1.062532 reflections 163 parameters H-atom parameters constrained $D_x = 1.210 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.18 \text{ mm}^{-1}$ T = 173 (2) K Block, colourless 0.44 \times 0.32 \times 0.22 mm

2532 independent reflections 2474 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.021$ $\theta_{\text{max}} = 28.0^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0541P)^2 \\ &+ 0.118P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.011 \\ \Delta\rho_{\text{max}} &= 0.34 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.19 \text{ e } \text{\AA}^{-3} \\ \text{Absolute structure: Flack (1983),} \\ & 825 \text{ Friedel pairs} \\ \text{Flack parameter: } -0.04 (8) \end{split}$$

Table 1

Selected bond angles (°).

O-P-C4	115.03 (8)	O-P-C10	110.95 (8)
O-P-C16	110.89 (8)	C4-P-C10	107.95 (9)
C4-P-C16	105.27 (9)	C16-P-C10	106.23 (8)
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Table 2

Hydrogen-bond geometry (Å, °).

Cg1 and Cg2 denote the centroids of the C5–C10 and C11–C16 rings, respectively.

D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
0.99	2.86	3.676 (3)	153
0.99	2.71	3.695 (3)	174
0.95	2.91	3.729 (3)	145
	<i>D</i> -H 0.99 0.99 0.95	D−H H···A 0.99 2.86 0.99 2.71 0.95 2.91	$D-H$ $H \cdots A$ $D \cdots A$ 0.992.863.676 (3)0.992.713.695 (3)0.952.913.729 (3)

Symmetry codes: (i) x + 1, y, z; (ii) $x - \frac{1}{2}, -y + 1, z + \frac{1}{2}$



Figure 2

Packing diagram of (I), viewed down the *b* axis. $C-H\cdots\pi$ interactions are shown as red dashed lines.

C-bound H atoms were positioned geometrically, with C-H = 0.93 Å, and allowed to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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