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#### **Key indicators**

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.039 wR factor = 0.114 Data-to-parameter ratio = 19.3

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

# Di-tert-butylphosphine oxide

The title compound,  $C_8H_{19}OP$ , is the oxidation product of  ${}^{t}Bu_2PH$ . The molecule is located on a crystallographic mirror plane. As a result, there is just one half-molecule in the asymmetric unit.

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## Comment

Transition metal complexes with phosphine ligands,  $PR_3$  (R = alkyl, aryl, H) play an important role in homogeneous catalysis (Holleman & Wiberg, 1995). A disadvantage of these ligands is their air sensitivity. Upon standing under atmospheric conditions, they are oxidized to the corresponding phosphine oxide  $R_3PO$  (R = alkyl, aryl, H). We report here the synthesis and the X-ray crystal structure analysis of di-*tert*-butyl-phosphine oxide, <sup>1</sup>Bu<sub>2</sub>HPO, (I). The synthesis of (I) was achieved by air oxidation of <sup>1</sup>Bu<sub>2</sub>PH, as indicated in the scheme below.



A perspective view of the title compound is shown in Fig. 1. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, Version 1.7; Mogul Version 1.0.1; Allen, 2002). A crystallographic mirror plane perpendicular to the *b* axis runs through HP=O. Therefore, there is just one half-molecule in the asymmetric unit.

## **Experimental**

Air oxidation of di-*tert*-butylphosphine (0.5 ml) for 2 d yields di-*tert*-butylphosphine oxide (yield 20%). Single crystals of di-*tert*-butylphosphine oxide were grown from the reaction mixture at ambient temperature. The NMR spectra were recorded on a Bruker DPX 250 and a Bruker Avance 400 spectrometer.<sup>1</sup>H NMR (C<sub>6</sub>D<sub>6</sub>, internal TMS):  $\delta$  0.97 (d, 2 × 'Bu, <sup>3</sup>J<sub>PH</sub> = 14.64 Hz), 5.82 (d, 1H, <sup>1</sup>J<sub>PH</sub> = 422.40 Hz). <sup>13</sup>C{<sup>1</sup>H} NMR (C<sub>6</sub>D<sub>6</sub>, internal TMS):  $\delta$  25.6 (d, 2 × CMe<sub>3</sub>, <sup>2</sup>J<sub>PC</sub> = 1.41 Hz), 33.5 (d, 2 × CMe<sub>3</sub>, <sup>1</sup>J<sub>PC</sub> = 59.29 Hz). <sup>31</sup>P NMR (C<sub>6</sub>D<sub>6</sub>, external H<sub>3</sub>PO<sub>4</sub>):  $\delta$  63.5 (dm, <sup>1</sup>J<sub>PH</sub> = 422.40 Hz, <sup>3</sup>J<sub>HP</sub> = 14.64 Hz).

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### Figure 1

Perspective view of the title compound with the atom numbering and displacement ellipsoids drawn at the 50% probability level. [Symmetry code: (i)  $x, -y + \frac{1}{2}, z$ .]

#### Crystal data

C<sub>8</sub>H<sub>19</sub>OP  $M_r = 162.20$ Orthorhombic, Pnma a = 10.4244 (10) Åb = 15.9819 (16) Å c = 6.0919 (7) Å V = 1014.92 (18) Å<sup>3</sup> Z = 4 $D_x = 1.062 \text{ Mg m}^{-3}$ 

#### Data collection

Stoe IPDS-II two-circle diffractometer  $\omega$  scans Absorption correction: multi-scan (MULABS; Spek, 2003; Blessing, 1995)

 $T_{\rm min}=0.915,\ T_{\rm max}=0.979$ 6207 measured reflections

Mo  $K\alpha$  radiation Cell parameters from 7754 reflections  $\theta = 3.8 - 25.6^{\circ}$  $\mu = 0.22 \text{ mm}^{-1}$ T = 173 (2) K Plate, colourless  $0.42\,\times\,0.18\,\times\,0.05$  mm

985 independent reflections 885 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.051$  $\theta_{\rm max} = 25.5^{\circ}$  $h = -12 \rightarrow 12$  $k = -17 \rightarrow 19$  $l = -7 \rightarrow 7$ 

Refinement

H atoms treated by a mixture of $\Delta \rho_{min} = -0.52 \text{ e A}$ Extinction correction: SHELXL9	S = 1.06 985 reflections 51 parameters H atoms treated by a mixture of	$\begin{array}{l} (\Delta/\sigma)_{max} < 0.001 \\ \Delta\rho_{max} = 0.49 \text{ e} \text{ Å}^{-3} \\ \Delta\rho_{min} = -0.32 \text{ e} \text{ Å}^{-3} \\ \text{Extinction correction: SHELXL97} \end{array}$
H atoms treated by a mixture of independent and constrained refinement Extinction coefficient: 0.011 (5)	H atoms treated by a mixture of independent and constrained refinement	Extinction correction: <i>SHELXL97</i> Extinction coefficient: 0.011 (5)

### Table 1 Selected bond lengths (Å).

P1-O1	1.4819 (19)	P1-H1	1.28 (3)
P1-C1	1.8484 (16)		

All H atoms were located in a difference map and were refined with fixed individual displacement parameters  $[U_{iso}(H) = 1.2U_{eq}(P)]$ or  $1.5U_{eq}(C)$ ] using a riding model (C-H = 0.98 Å). In addition, the P-H bond length was refined.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP in SHELXTL-Plus (Sheldrick, 1991); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2003).

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