

1-(Diphenylphosphinoyl)-3,3-dimethylbut-1-yne

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Key indicators

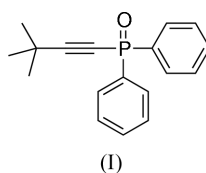
Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.036
 wR factor = 0.088
Data-to-parameter ratio = 12.8For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.The title compound, $\text{C}_{18}\text{H}_{19}\text{OP}$, has a mirror plane passing through the phosphinoyl and acetylene groups. There is a weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ interaction.

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Comment

Acetylenic phosphine oxides have attracted considerable interest because they are versatile intermediates for the synthesis of complex natural products and biologically active compounds (Braga *et al.*, 2002). The asymmetric unit of the title compound, (I), consists of a half molecule. The molecule contains *tert*-butyl and $-\text{O}=\text{P}(\text{Ph})_2$ moieties linked by an acetylene fragment. Atoms C9 and P1 and the acetylene group form a linear moiety and are located on a mirror plane. Atom P1 has a slightly distorted tetrahedral configuration. Bond angles around P1 are in the range 104.5 (1)– 114.2 (2) $^\circ$ (Table 1). There are two benzene-ring planes, forming a dihedral angle of 119.15 (9) $^\circ$. The three-dimensional framework of (I) is stabilized by a weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ interaction (Table 1).

Experimental

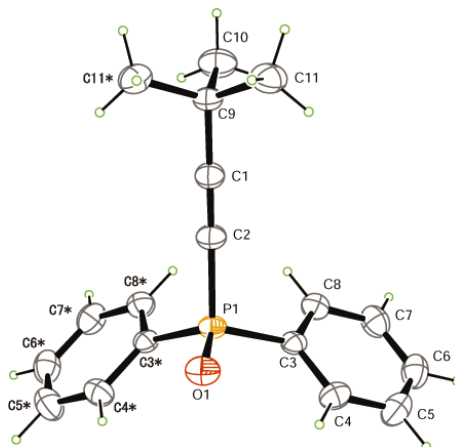
tert-Butylethynylphosphine (50 mmol) was dissolved in acetone (50 ml), the mixture cooled to 273 K and 30% H_2O_2 (6 ml) in acetone

Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids. [Symmetry code: (*) $1 - x, y, z$.]

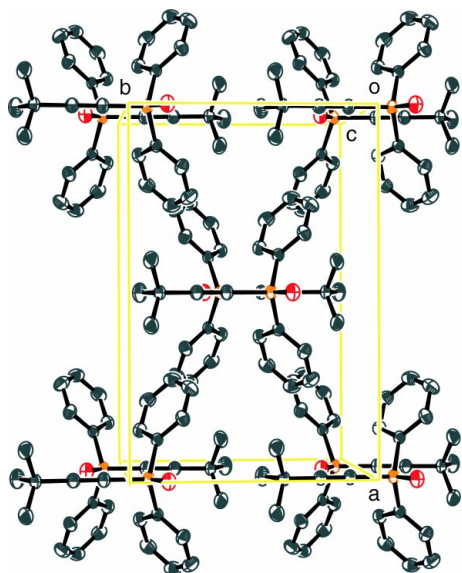


Figure 2
The crystal structure of (I). H atoms have been omitted.

(30 ml) added dropwise in 1 h. The mixture was extracted with ethyl acetate, dried with MgSO_4 and crystallized from diethyl ether and petroleum ether to obtain the title compound, (I), as a colorless solid (yield 13.5 g, 96%; Charrier *et al.*, 1966).

Crystal data

$\text{C}_{18}\text{H}_{19}\text{OP}$
 $M_r = 282.32$
 Orthorhombic, $\text{Cmc}2_1$
 $a = 14.4742$ (7) Å
 $b = 9.5767$ (7) Å
 $c = 11.2075$ (6) Å
 $V = 1553.5$ (2) Å³
 $Z = 4$
 $D_x = 1.207$ Mg m⁻³

Data collection

Rigaku R-AXIS RAPID
 diffractometer
 ω scans
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.873$, $T_{\max} = 0.983$
 7260 measured reflections

Mo $K\alpha$ radiation
 Cell parameters from 8920
 reflections
 $\theta = 2.5$ – 27.5°
 $\mu = 0.17$ mm⁻¹
 $T = 296$ (1) K
 Block, colorless
 $0.60 \times 0.20 \times 0.10$ mm

 1688 independent reflections
 1309 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.036$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -18 \rightarrow 18$
 $k = -12 \rightarrow 12$
 $l = -13 \rightarrow 14$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.088$
 $S = 1.00$
 1309 reflections
 102 parameters
 H-atom parameters constrained

$w = 1/[0.001F_o^2 + 1.1\sigma(F_o^2)]/(4F_o^2)$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.29$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.33$ e Å⁻³
 Absolute structure: (Flack, 1983),
 632 Friedel pairs
 Flack parameter = 0.01 (1)

Table 1

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{C8}-\text{H8} \cdots \text{O1}^i$	0.97	2.55	3.464 (3)	156

Symmetry code: (i) $x, 1 - y, z - \frac{1}{2}$.

The H atoms were placed in calculated positions, with $\text{C}-\text{H} = 0.97$ Å, and included in the refinement using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *CRYSTALS* (Watkin *et al.*, 1996); molecular graphics: *ORTEP-3* for Windows (Farrugia, 1997); software used to prepare material for publication: *CrystalStructure*.

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