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Key indicators

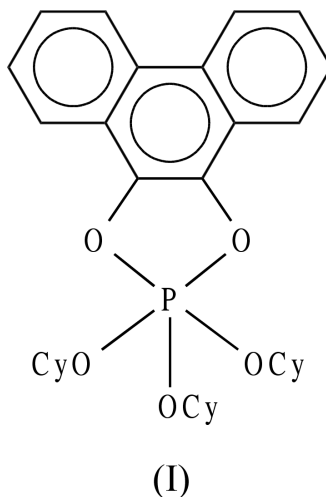
Single-crystal X-ray study
 $T = 143\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
Disorder in main residue
 R factor = 0.048
 wR factor = 0.119
Data-to-parameter ratio = 13.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

2,2,2-Tris(cyclohexyloxy)-4,5-(2',2''-biphenyl)-1,3,2-dioxaphospholene

The title compound (2,2,2-tris(cyclohexyloxy)phenanthro[9,10-d][1,3,2]dioxaphosphole), $\text{C}_{32}\text{H}_{41}\text{O}_5\text{P}$, displays trigonal-bipyramidal geometry at the central P atom. The P–O bond lengths (\AA) are: in the five-membered ring 1.652 (2) \AA equatorial and 1.744 (2) \AA axial, and to the cyclohexyloxy groups 1.581 (2) and 1.589 (2) \AA equatorial, and 1.624 (2) \AA axial.

Comment

We are interested in the correlation between structure and reactivity (see *e.g.* Amos *et al.*, 1992). The structure of the title compound, (I), the adduct of tricyclohexyl phosphite and phenanthraquinone, was determined as part of a combined crystallographic and theoretical investigation of pKa values of pentaoxyphosphoranes (Davies *et al.*, 2002).



The structure is shown in Fig. 1; a minor (11%) disorder component of the ring C11–16 is not shown. Of the three cyclohexyloxy groups, one is axial (O5, C31 *etc.*) and the other two are equatorial. This is reflected in the normal bond-length pattern for trigonal-bipyramidal geometry (axial bonds longer than equatorial), as was also observed in pentakis(cyclohexyloxy)phosphorane (Davies *et al.*, 2001; average values in four independent molecules 1.662 \AA for axial and 1.606 \AA for equatorial P–O bonds). The triisopropoxy analogue of the title compound was subjected to structure analysis in 1967. There were two modifications, of which the more precisely determined modification (Spratley *et al.*, 1967) displayed a similar P–O bond-length pattern to the title compound: ring apical 1.751 (7) \AA , equatorial 1.633 (7) \AA ; *i*Pr apical 1.638 (6) \AA , and equatorial 1.588 (7) and 1.574 (7) \AA .

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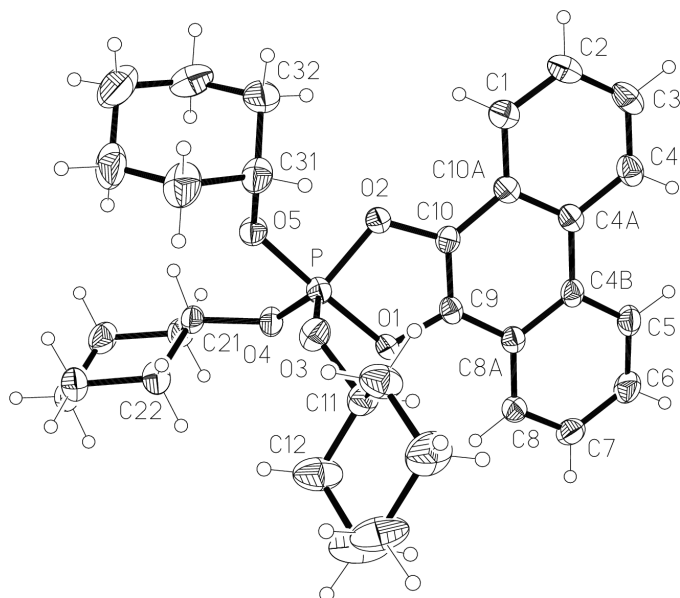


Figure 1
The molecule of the title compound in the crystal. The minor disorder component of the ring C11–C16 has been omitted. Ellipsoids represent 50% probability levels.

Experimental

Compound (I) was prepared by adding triisopropyl phosphite to one equivalent of phenanthraquinone at room temperature, according to the method of Hamilton *et al.* (1967).

Crystal data

$C_{32}H_{41}O_5P$
 $M_r = 536.62$
Monoclinic, $P2_1/n$
 $a = 9.665$ (2) Å
 $b = 15.394$ (3) Å
 $c = 19.200$ (4) Å
 $\beta = 93.92$ (2)°
 $V = 2850.1$ (10) Å³
 $Z = 4$

$D_x = 1.251$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 50 reflections
 $\theta = 10$ – 11.5°
 $\mu = 0.14$ mm⁻¹
 $T = 143$ (2) K
Prism, colourless
 $0.65 \times 0.30 \times 0.30$ mm

Data collection

Stoe Stadi-4 diffractometer
 ω/θ scans
Absorption correction: none
5588 measured reflections
5037 independent reflections
3708 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.024$

$\theta_{max} = 25.0^\circ$
 $h = -11 \rightarrow 0$
 $k = -18 \rightarrow 1$
 $l = -22 \rightarrow 22$
3 standard reflections
frequency: 60 min
intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.119$
 $S = 1.04$
5037 reflections
368 parameters
H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0447P)^2 + 1.4014P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.21$ e Å⁻³
 $\Delta\rho_{min} = -0.31$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

P–O3	1.581 (2)	P–O2	1.652 (2)
P–O4	1.589 (2)	P–O1	1.744 (2)
P–O5	1.624 (2)		
O3–P–O4	117.35 (9)	O5–P–O2	88.52 (8)
O3–P–O5	92.73 (9)	O3–P–O1	94.57 (8)
O4–P–O5	91.98 (8)	O4–P–O1	84.38 (8)
O3–P–O2	115.31 (9)	O5–P–O1	172.69 (9)
O4–P–O2	127.24 (9)	O2–P–O1	88.66 (8)

H atoms were positioned geometrically and refined using a riding model. Cyclohexyl ring C11–16 is disordered over two positions with a relative occupancy of *ca* 9:1. The minor component was refined isotropically with an appropriate system of restraints.

Data collection: *DIF4* (Stoe & Cie, 1992); cell refinement: *DIF4*; data reduction: *REDU4* (Stoe & Cie, 1992); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL97*.

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