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## Key indicators

Single-crystal X-ray study

T = 150 K

Mean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$ 

Disorder in main residue

R factor = 0.028

wR factor = 0.073

Data-to-parameter ratio = 12.1

For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.*O,O*-Bis(2-*tert*-butyl-4-methoxyphenyl)  
chlorothiophosphonate. Corrigendum.

In the original report by Odabaşoğlu, Büyükgüngör & Albayrak [*Acta Cryst.* (2005), E61, o2528–o2530], the structure was reported in the incorrect space group *Cc*. The structure is now reported as disordered in the correct space group *C2/c*. The P atom lies on a twofold rotation axis. A revised description of the hydrogen bonding is also given.

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## Comment

An *ORTEP*-3 (Farrugia, 1997) view of (I) and a packing diagram are shown in Figs. 1 and 2, respectively. The P atom lies on a twofold rotation axis, leading to disorder of the Cl and S atoms. Compound (I) has no classical hydrogen bonds, but there are two C–H... $\pi$  interactions: H8A...Cg1 = 3.12 (2) Å and C8–H8A...Cg1 = 122.8 (2)° (Cg1 is the centroid of the C1<sup>ii</sup>–C6<sup>ii</sup> ring), and H11A...Cg2 = 2.71 (2) Å and C11–H11A...Cg2 = 142.7 (1)° (Cg2 is the centroid of the C1<sup>iii</sup>–C6<sup>iii</sup>

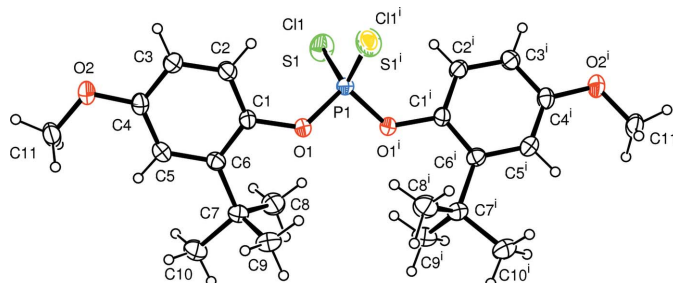


Figure 1

A view of (I), with the atom-numbering scheme and 50% probability displacement ellipsoids. [Symmetry code: (i)  $-x + 1, y, -z + \frac{1}{2}$ ]

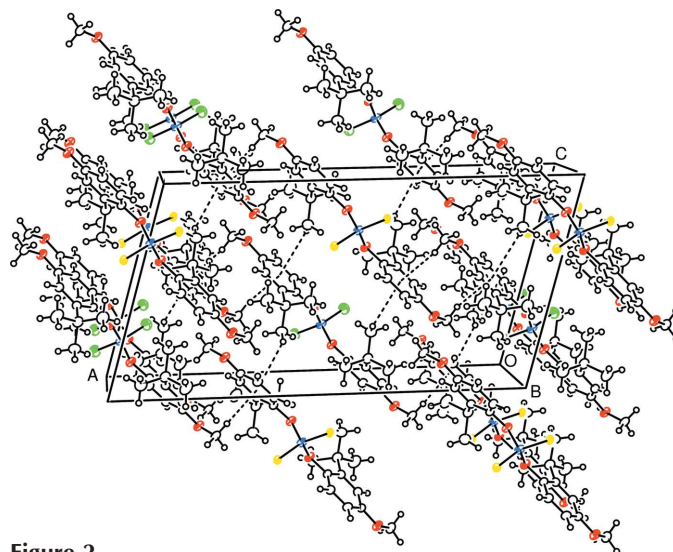


Figure 2

A view of the packing of (I); C–H... $\pi$  interactions are drawn as dashed lines.

ring) [symmetry codes: (ii)  $x, 1 - y, \frac{1}{2} + z$ ; (iii)  $\frac{1}{2} - x, \frac{1}{2} - y, -z$ ]. The dihedral angle between the symmetry-related benzene rings is  $41.2(2)^\circ$ . Selected bond distances and angles are given in Table 1.

## Experimental

### Crystal data

$C_{22}H_{30}ClO_4PS$	$D_x = 1.286 \text{ Mg m}^{-3}$
$M_r = 456.94$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 16485 reflections
$a = 23.592(3) \text{ \AA}$	$\theta = 1.7\text{--}26.7^\circ$
$b = 8.3111(6) \text{ \AA}$	$\mu = 0.34 \text{ mm}^{-1}$
$c = 12.5067(14) \text{ \AA}$	$T = 150 \text{ K}$
$\beta = 105.740(9)^\circ$	Prism, colorless
$V = 2360.3(4) \text{ \AA}^3$	$0.64 \times 0.59 \times 0.55 \text{ mm}$
$Z = 4$	

### Data collection

Stoe IPDS-II diffractometer	2146 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\text{int}} = 0.032$
Absorption correction: integration ( <i>X-RED32</i> ; Stoe & Cie, 2002)	$\theta_{\text{max}} = 26.0^\circ$
$T_{\text{min}} = 0.824, T_{\text{max}} = 0.869$	$h = -28 \rightarrow 28$
16485 measured reflections	$k = -10 \rightarrow 10$
2327 independent reflections	$l = -15 \rightarrow 15$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0362P)^2 + 1.6163P]$
$R[F^2 > 2\sigma(F^2)] = 0.028$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.073$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 1.08$	$\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$
2327 reflections	$\Delta\rho_{\text{min}} = -0.34 \text{ e \AA}^{-3}$
192 parameters	
All H-atom parameters refined	

**Table 1**

Selected geometric parameters ( $\text{\AA}, ^\circ$ ).

C1—C2	1.3946 (18)	C4—C5	1.4017 (19)
C1—C6	1.4086 (17)	C5—C6	1.4088 (18)
C1—O1	1.4226 (14)	O1—P1	1.5750 (9)
C2—C3	1.3860 (18)	P1—S1 <sup>i</sup>	1.9791 (4)
C3—C4	1.3979 (18)	P1—Cl1	1.9791 (4)
C2—C1—C6	123.25 (11)	C4—C5—C6	121.99 (12)
C2—C1—O1	119.13 (11)	C1—C6—C5	115.48 (11)
C6—C1—O1	117.58 (11)	O1—P1—O1 <sup>i</sup>	97.08 (7)
C3—C2—C1	119.77 (12)	O1—P1—S1 <sup>i</sup>	111.27 (4)
C2—C3—C4	119.11 (12)	O1—P1—Cl1	111.44 (3)
C3—C4—C5	120.39 (11)	S1 <sup>i</sup> —P1—Cl1	113.26 (3)

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

All H atoms were refined freely. Atoms S1 and Cl1 were assigned to the same atomic site and refined freely with the same atomic coordinates and with fixed site-occupancy factors of 0.5.

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* for Windows (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

## References

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