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## Key indicators

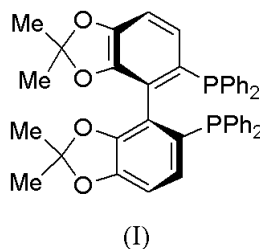
Single-crystal X-ray study  
 $T = 298$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.034  
 $wR$  factor = 0.081  
Data-to-parameter ratio = 8.5For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.**(S)-(-)-5,5'-Bis(diphenylphosphino)-2,2',2'-tetramethyl-4,4'-bi-1,3-benzodioxole**The title molecule,  $\text{C}_{42}\text{H}_{36}\text{O}_4\text{P}_2$ , is a new atropisomeric bis-phosphine ligand. All bond lengths and angles are normal. In the crystal structure, the molecule possesses a crystallographically imposed  $C_2$  axis. The two benzene rings in the biphenyl moiety make a dihedral angle of  $84.23(6)^\circ$ .

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## Comment

Since 1968, when a chiral phosphine was first utilized in asymmetric hydrogenation (Knowles & Sabacky, 1968; Horner *et al.*, 1968), much effort has been devoted to the design and synthesis of chiral phosphine ligands. The atropisomeric  $C_2$ -symmetric bis-phosphine ligands play an important role in asymmetric hydrogenation (Zhang *et al.*, 2000; Saito *et al.*, 2001). We report here the crystal structure of the title compound, (I), a new atropisomeric  $C_2$ -symmetric bis-phosphine ligand.In the crystal structure, molecules of (I) possess a crystallographically imposed  $C_2$  axis (Fig. 1). All bond lengths and angles are normal. The two benzene rings in the biphenyl moiety make a dihedral angle of  $84.23(6)^\circ$ . All rings are planar.

## Experimental

Under argon, a 100 ml three-necked flask was charged with (S)-(-)-5,5'-bis(diphenylphosphino)-2,2',2'-tetramethyl-4,4'-bi-1,3-benzodioxole (0.0698 g, 1 mmol), toluene (11 ml), *N,N*-dimethylaniline (1.4 ml, 10 mmol) and trichlorosilane (1.01 ml, 10 mmol). The mixture was refluxed for 10 h. After the mixture had cooled to 273 K, a degassed 50% NaOH solution (50 ml) was added carefully. The product was extracted with toluene (30 ml) twice. The extract was washed successively with 10% HCl, water and brine, then dried with anhydrous  $\text{Na}_2\text{SO}_4$  and concentrated to give the crude product. Recrystallization from MeOH afforded a white solid.  $^1\text{H}$  NMR (chloroform-*d*):  $\delta$  1.12 (6H, *s*), 1.58 (6H, *s*), 6.56 (2H, *d*,  $J = 7.8$  Hz), 6.67 (2H, *d*,  $J = 7.8$  Hz), 7.11–7.25 (20H, *m*).  $^{13}\text{C}$  NMR (chloroform-*d*): 25.27, 25.93, 108.46, 118.27, 122.77–147.86.  $^{31}\text{P}$  NMR (chloroform-*d*): -14.44. MS (ESI): 667 HRMS (ESI). Calculated for  $\text{C}_{42}\text{H}_{36}\text{NaO}_4\text{P}_2$  [ $M + \text{Na}$ ] $^+$ : 689.1891; found: 689.1966.

Crystal data

$C_{42}H_{36}O_4P_2$   
 $M_r = 666.65$   
 Orthorhombic,  $P2_12_12$   
 $a = 9.7007(9) \text{ \AA}$   
 $b = 20.774(2) \text{ \AA}$   
 $c = 8.9149(9) \text{ \AA}$   
 $V = 1796.6(3) \text{ \AA}^3$   
 $Z = 2$   
 $D_x = 1.232 \text{ Mg m}^{-3}$

Mo  $K\alpha$  radiation  
 Cell parameters from 2896 reflections  
 $\theta = 4.6\text{--}22.3^\circ$   
 $\mu = 0.16 \text{ mm}^{-1}$   
 $T = 298(2) \text{ K}$   
 Block, colorless  
 $0.50 \times 0.41 \times 0.35 \text{ mm}$

Data collection

Bruker SMART APEX area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2002)  
 $T_{\min} = 0.924, T_{\max} = 0.946$   
 9313 measured reflections

1877 independent reflections  
 1677 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.046$   
 $\theta_{\text{max}} = 25.2^\circ$   
 $h = -11 \rightarrow 10$   
 $k = -22 \rightarrow 24$   
 $l = -10 \rightarrow 10$

Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.081$   
 $S = 0.99$   
 1877 reflections  
 220 parameters  
 H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0487P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.19 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.17 \text{ e \AA}^{-3}$   
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.0085 (14)  
 Absolute structure: Flack (1983), 1357 Friedel pairs  
 Flack parameter = 0.00 (9)

H atoms were placed in calculated positions ( $C-H = 0.93\text{--}0.96 \text{ \AA}$ ) and refined as riding, with  $U_{\text{iso}}(H) = 1.2$  or  $1.5$  times  $U_{\text{eq}}(C)$ .

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2002); software used to prepare material for publication: SHELXL97.

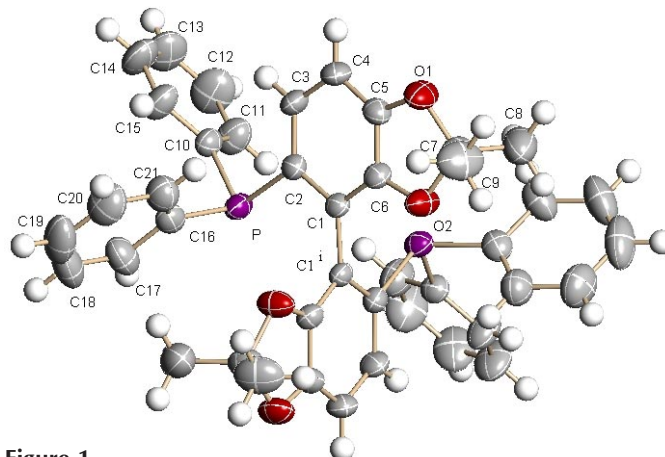


Figure 1 The formula unit of (I), with the atom numbering, showing displacement ellipsoids at the 50% probability level [symmetry code: (i)  $-x, 1 - y, z$ ].

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