# organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### Key indicators

Single-crystal X-ray study T = 294 K Mean  $\sigma$ (C–C) = 0.005 Å Disorder in solvent or counterion R factor = 0.053 wR factor = 0.147 Data-to-parameter ratio = 18.4

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

# (S)-2,2',6,6'-Tetramethoxy-4,4'-bis(diphenylphosphino)-3,3'-bipyridine-(2S,3S)-(+)dibenzoyl-L-tartaric acid-chloroform (1/1/1)

Received 25 July 2003 Accepted 28 July 2003

Online 8 August 2003

The structure of the title cocrystal,  $C_{38}H_{34}N_2O_6P_2$ ·- $C_{18}H_{14}O_8$ ·CHCl<sub>3</sub>, (I), reveals that the crystals are built up of infinite chains in which (–)-2,2',6,6'-tetramethoxy-4,4'-bis(diphenylphosphino)-3,3'-bipyridine, (II), and (+)-dibenzoyl-L-tartaric acid, (III), are connected in an alternating fashion through two intermolecular hydrogen bonds between O atoms of the P=O groups in (II) and H atoms of the COOH groups in (III).

#### Comment

Transition-metal-catalysed asymmetric hydrogenation reactions are one of the most efficient methods for preparing a wide range of enantiomerically pure compounds. The possibility of discovering catalysts with improved utility, activity and selectivity by designing new ligands plays a crucial role in this research area. Consequently, the number of novel chiral ligands for catalytic asymmetric hydrogenation is growing rapidly. In this context, chiral bisphosphines (*e.g.* Dipamp, BINAP, DIOP, DuPhos *etc.*) are among the most useful and versatile ligands (Lin *et al.*, 2001). In contrast to the tremendous success achieved in the use of chiral arylphosphine ligands in Rh- and Ru-catalysed asymmetric reactions, chiral phosphine ligands containing heterocyclic moieties, such as pyridyl groups, have been relatively unexplored.

Recently, we have developed a chiral dipyridylphosphine ligand, namely 2,2',6,6'-tetramethoxy-4,4'-bis(diphenylphosphino)-3,3'-bipyridine, (II) (P-Phos; Pai *et al.*, 2000), which was found to be very effective in the Ru-catalysed asymmetric hydrogenation of 2-(6'-methoxy-2'-naphthyl)propenoic acid and  $\beta$ -ketoesters (Pai *et al.*, 2000).



The key step in the synthesis of P-Phos is the optical resolution of racemate (II) with optically active (+)-dibenzoyl-L-tartaric acid [(+)-DBT], (III), as a resolving reagent (Takaya *et al.*, 1988), which is remarkably efficient and highly selective. After several trials with our experimental modifications, the optically pure dipyridylphosphine oxides (R)-(II) and (S)-(II) were obtained in high enantiopurity. Thus, when a solution of

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Absolute structure: Flack (1983),

5987 Friedel pairs

Flack parameter = 0.05 (8)

(2S,3S)-(+)-DBT in ethyl acetate was added to a boiling solution of racemic (II) in dichloromethane, a 1:1 complex of (-)-(II) and (+)-DBT [(-)-(II)-(+)-DBT, (I)] precipitated as a crystalline solid. From internal comparison with (2S,3S)-(+)-DBT, the absolute configuration of (-)-II was determined to be S.

The crystal structure of (I) (Fig. 1) also contains a disordered chloroform molecule in the asymmetric unit. The dihedral angle between the least-squares planes of the two pyridyl rings is  $75.9(1)^{\circ}$ . Crystals are built up of infinite chains in which equimolar amounts of (-)-(II) and (+)-DBT are connected in an alternating manner through two intermolecular hydrogen bonds formed between O9 and O14 of the P=O groups of (-)-(II) and H atoms of the COOH groups of (+)-DBT. The O4···O14<sup>i</sup> and O6···O9<sup>ii</sup> distances are 2.554 (2) and 2.551 (2) Å, respectively [symmetry codes: (i)  $1 - x, y - \frac{1}{2}$ ,  $\frac{3}{2} - z$ ; (ii)  $\frac{1}{2} + x$ ,  $\frac{3}{2} - y$ , 1 - z; Fig. 2 and Table 1]. The low solubility of (I) is attributable to the observed polymeric structure.

## **Experimental**

To a boiling solution of racemic (II) (Pai et al., 2000) (1.87 g, 2.76 mmol) in chloroform (28 ml) was added a solution of (-)-DBT, (III), in ethyl acetate (28 ml). The mixture was refluxed until a white solid formed and then allowed to stand at ambient temperature overnight. The solid product was dried at room temperature under vacuum to give a 1:1 complex of (R)-(II) and (-)-DBT (93% yield). The mother liquor, which contained the (S)-(II)-(+)-DBT complex, was concentrated to dryness to give a solid material. This was dissolved in chloroform (10 ml) and then treated with 10% aqueous NaOH (10 ml). The mixture was extracted with two portions of chloroform (25 ml). Evaporation of the solvent gave crude (S)-(II), which was dissolved in refluxing chloroform (15 ml), and to this solution was added a solution of (+)-DBT in ethyl acetate (20 ml). The mixture was stirred at reflux temperature until the white precipitate formed and was then allowed to stand at room temperature overnight. The white precipitate was dried to give the pure (S)-(II)-(+)-DBT complex (92% yield). The crystal used for the data collection was obtained by slow evaporation from chloroform-ethyl acetate (1:1) saturated solution at room temperature.



### Figure 1

The molecular structure of (I), showing displacement ellipsoids at the 30% probability level (Bruker, 1995).

Crystal data

$\begin{array}{l} C_{38}H_{34}N_2O_6P_2 \cdot C_{18}H_{14}O_8 \cdot CHCl_3 \\ M_r = 1154.27 \\ \text{Orthorhombic, } P_{2_12_12_1} \\ a = 10.0967 \ (10) \text{ Å} \\ b = 21.000 \ (2) \text{ Å} \\ c = 27.390 \ (3) \text{ Å} \\ V = 5807.6 \ (10) \text{ Å}^3 \\ Z = 4 \\ D_x = 1.320 \text{ Mg m}^{-3} \\ \text{Data collection} \end{array}$	Mo $K\alpha$ radiation Cell parameters from 6132 reflections $\theta = 1-27.5^{\circ}$ $\mu = 0.28 \text{ mm}^{-1}$ T = 294 (2) K Block, colorless $0.28 \times 0.28 \times 0.20 \text{ mm}$
Siemens SMART CCD area- detector diffractometer $\varphi$ and $\omega$ scans Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 1996) $T_{min} = 0.926, T_{max} = 0.947$ 39 041 measured reflections	13 342 independent reflections 8576 reflections with $I > 2\sigma(I)$ $R_{int} = 0.043$ $\theta_{max} = 27.5^{\circ}$ $h = -13 \rightarrow 11$ $k = -23 \rightarrow 27$ $l = -30 \rightarrow 35$
Refinement	
Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.053$ $wR(F^2) = 0.147$ S = 1.01 13 342 reflections	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.08P)^{2}]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} = 0.014$ $\Delta\rho_{max} = 0.32 \text{ e} \text{ Å}^{-3}$ $\Lambda_{o} = -0.38 \text{ e} \text{ Å}^{-3}$

725 parameters H-atom parameters constrained

#### Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	<i>D</i> -H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O4-H4\cdots O14^{i}$	0.82	1.75	2.554 (2)	166
$O6-H6\cdots O9^{ii}$	0.82	1.76	2.551 (2)	162

Symmetry codes: (i)  $1 - x, y - \frac{1}{2}, \frac{3}{2} - z$ ; (ii)  $\frac{1}{2} + x, \frac{3}{2} - y, 1 - z$ .

H atoms were included in the riding model approximation, with C-H and O-H distances of 0.93–0.98 and 0.82 Å and a  $U_{iso}(H)$  of 1.2 or 1.5 times  $U_{eq}$  of the atom to which they were bound. Two of the Cl atoms in the solvent CHCl<sub>3</sub> molecule are disordered and were each



Figure 2 Packing diagram for (I). The chloroform molecules have been omitted.

modeled over two positions with site occupancies of 0.587(2) and 0.413(2).

Data collection: *SMART* (Siemens, 1995); cell refinement: *SMART*; data reduction: *SHELXTL-NT* (Siemens, 1995); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL-NT*; software used to prepare material for publication: *SHELXTL-NT*.

We thank The Hong Kong Polytechnic University ASD Fund for financial support of this study.

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