# metal-organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.032 wR factor = 0.083 Data-to-parameter ratio = 18.2

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

# (*Trans*-dichloro-*κCl*)bis[(dicyclohexylphosphino)methane-*κ*<sup>2</sup>*P*,*P*'] ruthenium(II)

In the title complex,  $[Ru(C_{25}H_{46}P_2)_2Cl_2]$  or  $RuCl_2(dcpm)_2$  [dcpm is 1,2-bis(dicyclohexylphosphino)methane,  $C_{25}H_{46}P_2$ ], the Ru atom (site symmetry  $\overline{1}$ ) is in a octahedral coordination environment with two chelating dcpm ligands and two *trans* chloro ligands.

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## Comment

Ruthenium complexes containing chelating phosphine ligands play a prominent role as active catalysts in many fields (Collman, 1987 White & Coville, 1994). Many of these complexes have been prepared and characterized (Shawkataly *et al.*, 1998; Lucas *et al.*, 2000; Hockless *et al.*, 1996), but complexes containing dcpm [dcpm is 1,2-bis(dicyclohexylphosphino)methane,  $C_{25}H_{46}P_2$ ] are uncommon (Joslin *et al.* 1991). Here, we report the synthesis and structure of the title compound, (I), (Fig. 1), which may serve as a useful precursor for the synthesis of further ruthenium complexes (Grocott & Wild, 1982; Nolan *et al.*, 1997; Volland *et al.*, 2004).



In (I) the central Ru(II) atom occupies an inversion centre and is coordinated by two *trans* chloro ligands and two *P*,*P*chelating dcpm ligands, resulting in octahedral geometry for the ruthenium atom (Table 1). The Ru–Cl distances in (I) are comparable to those in related compounds (Blake *et al.*, 1993; Hockless *et al.*, 1996). The two Ru–P distances in (I) differ by 0.056 Å. The packing in (I) is shown in Fig. 2.

### Experimental

Compound (I) was prepared from the reaction of the chelating phosphine ligand with ruthenium(III) chloride by a standard procedure (Grocott & Wild, 1982). Suitable crystals for single-crystal analysis were grown by diffusion of diethyl ether into a dichloromethane solution of (I) at room temperature. Yield 85%. Elemental analysis calcd for  $C_{50}H_{92}Cl_2P_4Ru$ : C 60.72%, H 9.37%, found: C 60.68%, H 9.35%.

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#### Crystal data

 $\begin{bmatrix} \text{Ru}(\text{C}_{25}\text{H}_{46}\text{P}_2)_2\text{Cl}_2 \end{bmatrix} \\ M_r = 989.09 \\ \text{Monoclinic, } C2/c \\ a = 24.120 (11) \text{ Å} \\ b = 10.732 (5) \text{ Å} \\ c = 21.780 (11) \text{ Å} \\ \beta = 108.098 (8)^{\circ} \\ V = 5359 (4) \text{ Å}^3 \\ Z = 4 \\ \end{bmatrix}$ 

#### Data collection

Bruker SMART CCD
diffractometer
$\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 1998)
$T_{\min} = 0.796, T_{\max} = 0.937$
13674 measured reflections

#### Refinement

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Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0366P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.032$	+ 1.6382P]
$wR(F^2) = 0.083$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.09	$(\Delta/\sigma)_{\rm max} = 0.001$
4717 reflections	$\Delta \rho_{\rm max} = 0.30 \ {\rm e} \ {\rm \AA}^{-3}$
259 parameters	$\Delta \rho_{\rm min} = -0.42 \text{ e} \text{ \AA}^{-3}$
H-atom parameters constrained	

 $D_x = 1.226 \text{ Mg m}^{-3}$ 

Cell parameters from 858

 $0.30 \times 0.18 \times 0.12 \text{ mm}$ 

4717 independent reflections

3644 reflections with  $I > 2\sigma(I)$ 

Mo  $K\alpha$  radiation

reflections

 $\mu = 0.54~\mathrm{mm}^{-1}$ 

T = 293 (2) K

Block, orange

 $R_{\rm int} = 0.040$ 

 $\theta_{\rm max} = 25.0^{\circ}$ 

 $h = -28 \rightarrow 24$ 

 $k = -12 \rightarrow 12$ 

 $l = -25 \rightarrow 21$ 

 $\theta = 2.7 - 26.3^{\circ}$ 

#### Table 1

Selected geometric parameters (Å, °).

Ru1-P2	2.3669 (12)	Ru1-Cl1	2.4334 (11)
Ru1-P1	2.3775 (10)		
P2-Ru1-P1	71.42 (4)	P2-C13-P1	97.10 (13)

H atoms were initially located in a difference Fourier map, then placed in ideal positions, with C-H = 0.97 Å, and refined as riding, with the constraint  $U_{iso}(H) = 1.2U_{eq}(C)$  applied.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SMART* or **SAINT**?; data reduction: *SAINT* (Bruker, 1998); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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## Figure 1

View of (I), with 30% displacement ellipsoids and H atoms omitted for clarity. Atoms with the suffix A are generated by the symmetry operation 1 - x, 2 - y, -z.



#### Figure 2

Unit-cell packing in (I), viewed along the a axis. H atoms have been omitted for clarity.

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