# metal-organic papers

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#### Key indicators

Single-crystal X-ray study T = 293 KMean  $\sigma(\text{C-C}) = 0.005 \text{ Å}$  R factor = 0.033 wR factor = 0.083 Data-to-parameter ratio = 19.7

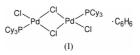
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# *trans*-Di-*µ*-chloro-bis[chloro(tricyclohexyl-phosphine)palladium] benzene solvate

The title compound, *trans*- $[Pd_2Cl_4(PCy_3)_2]\cdot C_6H_6$ , where PCy<sub>3</sub> is tricyclohexylphosphine (C<sub>18</sub>H<sub>33</sub>P), cocrystallizes with one benzene molecule in the centrosymmetric space group  $P2_1/c$  and both molecules lie on inversion centers. Each Pd atom adopts a distorted square-planar geometry [angles vary from 84.23 (3) to 95.82 (3)°], and is bonded to a terminal chloride, a phosphine and two bridging chlorides. Bond lengths are typical of related species.

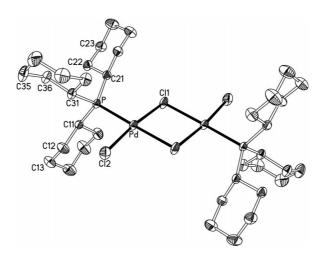
# Comment

The general class of  $[PdCl_2P]_2$  complexes has been known since the early studies of Mann & Purdie (1935) and Mann & Wells (1938). The title complex, (I), was prepared as a precursor to other palladium complexes. The structures of a range of related palladium complexes have been determined (Chaloner *et al.*, 1995; Coles *et al.*, 1999; Grigsby & Nicholson, 1992; Vicente *et al.*, 1997), all presenting virtually the same structural parameters as those of the complex reported here.



# **Experimental**

The title compound was isolated as a side-product of the reaction of (PhCN)<sub>2</sub>PdCl<sub>2</sub> with indenyllithium and PCy<sub>3</sub>. Single crystals suitable for X-ray diffraction study were obtained from a benzene solution.



#### Figure 1

© 2003 International Union of Crystallography Printed in Great Britain – all rights reserved View of the title molecule. Ellipsoids correspond to 30% probability. The unlabeled part of the molecule is related by the symmetry transformation (1 - x, -y, 2 - z).

Received 30 June 2003 Accepted 10 July 2003 Online 24 July 2003 Crystal data

$[Pd_2Cl_4(C_{18}H_{33}P)_2] \cdot C_6H_6$
$M_r = 993.58$
Monoclinic, $P2_1/c$
a = 9.9954 (1)  Å
b = 16.463 (1)  Å
c = 14.095 (1)  Å
$\beta = 95.657 \ (1)^{\circ}$
$V = 2308.1 (2) \text{ Å}^3$
Z = 2

## Data collection

Bruker SMART 2K diffractometer
$\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996a)
$T_{\min} = 0.266, T_{\max} = 0.475$
27 337 measured reflections
4457 independent reflections

#### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.033$	$w = 1/[\sigma^2(F_o^2) + (0.0525P)^2]$
$wR(F^2) = 0.083$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.001$
4457 reflections	$\Delta \rho_{\rm max} = 0.42 \ {\rm e} \ {\rm \AA}^{-3}$
226 parameters	$\Delta \rho_{\rm min} = -0.57 \text{ e } \text{\AA}^{-3}$

 $D_x = 1.430 \text{ Mg m}^{-3}$ 

Cell parameters from 6051

Block, light orange-red

 $0.16 \times 0.14 \times 0.08 \ \mathrm{mm}$ 

3791 reflections with  $I > 2\sigma(I)$ 

Cu  $K\alpha$  radiation

reflections

 $\theta = 2.7-72.5^{\circ}$  $\mu = 9.27 \text{ mm}^{-1}$ 

T = 293 (2) K

 $R_{\rm int} = 0.063$ 

 $\theta_{\rm max} = 72.7^{\circ}$ 

 $h = -9 \rightarrow 11$  $k = -20 \rightarrow 20$ 

 $l = -17 \rightarrow 17$ 

#### Table 1

Selected geometric parameters (Å, °).

Pd-P	2.2495 (7)	Pd-Cl1	2.3217 (8)
Pd-Cl2	2.2862 (8)	Pd-Cl1 <sup>i</sup>	2.4370 (7)
P-Pd-Cl2	90.16 (3)	Cl2-Pd-Cl1 <sup>i</sup>	89.86 (3)
P-Pd-Cl1	95.82 (3)	Cl1-Pd-Cl1 <sup>i</sup>	84.23 (3)
Cl2-Pd-Cl1	173.53 (3)	Pd-Cl1-Pd <sup>i</sup>	95.77 (3)
P-Pd-Cl1 <sup>i</sup>	178.32 (3)		
$P-Pd-Cl1-Pd^i$	178.31 (3)	Cl1-Pd-P-C31	-136.92 (11)
Cl1 <sup>i</sup> -Pd-Cl1-Pd <sup>i</sup>	0	Cl2-Pd-P-C21	169.53 (11)
Cl2-Pd-P-C11	-73.97(10)	Cl1-Pd-P-C21	-12.94(11)
Cl1-Pd-P-C11	103.56 (10)	C21-P-C31-C32	-78.0(3)
Cl2-Pd-P-C31	45.55 (11)	Pd-P-C31-C32	45.7 (3)

Symmetry code: (i) 1 - x, -y, 2 - z.

H atoms were constrained to the parent site using a riding model, with *SHELXL*96 (Sheldrick, 1996b) defaults (C-H = 0.94 Å).  $U_{iso}$ (H) values were set at  $1.2U_{eq}$  of the parent C atoms. A final verification of possible voids was performed using the VOID routine of the *PLATON* program (Spek, 1995).

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*96 (Sheldrick, 1996b); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXL*96.

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