# metal-organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 273 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.024 wR factor = 0.056 Data-to-parameter ratio = 20.3

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

## Chloro(triphenylphosphine)(tropolonato)palladium(II)

The title compound,  $[Pd(C_7H_5O_2)Cl(C_{18}H_{15}P)]$ , is a new type of  $\alpha$ -diketone palladium(II) complex containing a tertiary arylphosphine. It crystallizes with a distorted square-planar geometry about the palladium(II) metal centre. The most important bond distances and angles include Pd-O (*trans* P) = 2.0481 (12) Å, Pd-O (*trans* Cl) = 2.0016 (12) Å, Pd-P = 2.2268 (4) Å, Pd-Cl = 2.2770 (5) Å, O-Pd-O = 80.22 (5)° and O-C-C-O = 3.7 (2)°. A  $\pi$ - $\pi$  stacking interaction is observed between neighbouring tropolonate groups, with an interplanar distance of 3.377 (6) Å.

#### Comment

A variety of bis- $\beta$ -diketonato-palladium(II) complexes have been characterized to date for acetylacetonate-type ligands (Cambridge Structural Database; Version 5.27; Allen, 2002). The addition of tertiary aryl phosphines to these types of complexes has resulted in a limited number of reported structures (Okeya *et al.*, 1984; Ooi *et al.*, 1983; Siedle *et al.*, 1982). The inclusion of a chloro derivative with an  $\alpha$ - (Lang *et al.*, 1999) or  $\beta$ -diketone (Woisetschlager *et al.*, 2000; Navarro *et al.*, 2005) has attracted less attention. In a previous study (Steyl, 2006), we reported the structure of [(PCy<sub>3</sub>)(Trop)<sub>2</sub>Pd] (Cy = cyclohexyl and Trop = 2-hydroxy-2,4,6-cycloheptatrienone). The title compound, (I), is presented as an example of a monotropolonate-palladium(II) complex containing chloro and triphenylphosphine ligands.



The molecule of (I) crystallizes with a slightly distorted square-planar geometry about the palladium(II) metal centre. The Pd<sup>II</sup> atom is elevated by 0.0068 (1) Å above the plane defined by the four coordinated atoms (O11, O12, P, Cl). The Pd–O bond distances (Table 1) do not differ from those observed for the bis(tropolonato)palladium(II) (Steyl, 2005) or the tricyclohexyl (Steyl, 2006) derivatives. The bidentate bite angle of 80.22 (5)° (Table 1) does not differ significantly from that in the previously reported tropolonate structures of palladium(II). A torsional twist is observed for the tropolonate ligand (see values of torsion angles in Table 1).

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#### Figure 1

The molecular structure of (I), showing the numbering scheme and displacement ellipsoids drawn at the 50% probability level. H atoms have been omitted for clarity.



#### Figure 2

A pair of moleculesl, showing the stacking [symmetry code: (i) -x, 2 - y, -z]. H atoms have been omitted.

A weak intramolecular C—H···Cl hydrogen bond (Table 2) is found in (I).  $\pi$ - $\pi$  Stacking between parallel tropolonate ring systems (Fig. 2) is observed, with an interplanar distance of 3.377 (6) Å, which is significantly shorter than that [3.51 (1)–3.80 (2) Å] in the previously published tropolonate–palladium(II) complexes (Steyl, 2005, 2006). This close interaction can be attributed to the shift of the tropolonate ring system to form an eclipsed conformation of the C11–C17 sevenmembered ring.

In conclusion, the substitution of a chloro group for a tropolonate  $\{[PCy_3(Trop)_2Pd^{II}]; Steyl, 2006\}$  ligand does not significantly alter the bonding mode of the remaining tropolonate ligand to the Pd<sup>II</sup> atom.

## **Experimental**

The title complex was synthesized by the addition of  $PPh_3$  (83 mg, 0.316 mmol) to an acetone solution (10 ml) of the bis-tropolonato-

palladium(II) complex (100 mg, 0.287 mmol) containing hydrochloric acid (0.5 ml). The suspension dissolved and gave an orange solution; on evaporation of the solvent, crystals suitable for X-ray crystallography were obtained. Yield 45 mg (30%).

### Crystal data

 $\begin{bmatrix} Pd(C_7H_5O_2)Cl(C_{18}H_{15}P) \end{bmatrix} \\ M_r = 525.23 \\ Triclinic, P\overline{1} \\ a = 9.8561 (3) \text{ Å} \\ b = 9.9719 (2) \text{ Å} \\ c = 13.2792 (3) \text{ Å} \\ \alpha = 85.851 (1)^{\circ} \\ \beta = 72.262 (1)^{\circ} \\ \gamma = 63.317 (1)^{\circ} \end{bmatrix}$ 

#### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Bruker, 1998)  $T_{\min} = 0.782, T_{\max} = 0.897$ 24655 measured reflections

### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.024$   $wR(F^2) = 0.057$  S = 1.045506 reflections 271 parameters H-atom parameters constrained Prism, red  $0.25 \times 0.14 \times 0.11 \text{ mm}$ 

V = 1107.51 (5) Å<sup>3</sup>

 $D_x = 1.575 \text{ Mg m}^{-3}$ 

Mo  $K\alpha$  radiation  $\mu = 1.05 \text{ mm}^{-1}$ 

T = 273 (2) K

Z = 2

5506 independent reflections 4981 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.026$  $\theta_{\text{max}} = 28.4^{\circ}$ 

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.039P)^2 \\ &+ 3.4366P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.002 \\ \Delta\rho_{\text{max}} &= 0.37 \text{ e } \text{ Å}^{-3} \\ \Delta\rho_{\text{min}} &= -0.45 \text{ e } \text{ Å}^{-3} \end{split}$$

### Table 1

Selected geometric parameters (Å, °).

Pd-O11	2.0481 (12)	C11-O11	1.291 (2)
Pd-O12	2.0016 (12)	C12-O12	1.295 (2)
Pd-P	2.2268 (4)	C12-C11	1.462 (2)
Pd-Cl	2.2770 (5)		
O11-Pd-Cl	93 10 (4)	O12 - Pd - P	95.01 (4)
O12-Pd-O11	80.22 (5)	P-Pd-Cl	91.818 (18)
O12-C12-C11-O11	3.7 (2)	C13-C12-C11-C17	5.2 (3)
	(=)		(-)

## Table 2

Hydrogen-bond	geometry	(A, °	).
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$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
C32—H32···Cl	0.93	2.67	3.496 (2)	148
$C25 - H25 \cdots O11^{ii}$	0.93	2.46	3.333 (3)	157

Symmetry code: (ii) -x + 1, -y + 1, -z.

All H atoms were positioned geometrically and allowed to ride on their parent atoms, with C-H = 0.93 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$ .

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT-Plus* (Bruker, 2004); data reduction: *SAINT-Plus* and *XPREP* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2004); software used to prepare material for publication: *SHELXL97*.

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