metal-organic papers

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.024 wR factor = 0.078 Data-to-parameter ratio = 21.6

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

(Tricyclohexylphosphine)bis(tropolonato)palladium(II)

A new type of α -diketone palladium(II) complex containing a tertiary phosphine, namely the title compound, $[Pd(C_7H_5O_2)_2(C_{18}H_{33}P)]$, crystallizes with a square-planar geometry about the palladium(II) metal centre. One tropolonate ligand is bonded in a bidentate chelating mode, while the second one is bonded through a single O atom to the palladium(II) metal centre. The Pd-O distances range from 1.9907 (13) to 2.0702 (12) Å, and the O-Pd-O bite angle for the bidentate tropolonate ligand is 80.30 (5)°.

Comment

A variety of bis- β -diketonate palladium(II) complexes have been characterized to date for acetylacetonato type ligands (Cambridge Structural Database; Version 5.27; Allen, 2002). The addition of tertiary phosphines to these types of complex has resulted in only three reported structures (Okeya *et al.*, 1984; Ooi *et al.*, 1983; Siedle *et al.*, 1982). To date, no α -diketone palladium(II) complexes containing a phosphine ligand have been reported. In the current paper, a tricyclohexylphosphine derivative of bis(tropolonato)palladium(II) (Steyl, 2005) is presented as an example of a mono/bidentate tropolonate complex.



The title compound, (I), crystallizes with a slightly distorted square-planar geometry about the palladium(II) metal centre. The palladium atom is elevated by 0.0501 (1) Å above the plane defined by the four coordinated atoms, *viz*. O11, O12, O22 and P. The Pd–O distances do not differ from those observed for the bis(tropolonato)palladium(II) complex (Steyl, 2005). The O12–Pd–O11 bidentate bite angle [80.30 (5)°; Table 1] is less than the value of 81.8 (1)° observed in the bis(tropolonato)palladium(II) complex.

An interesting feature in (I) is the rotation of one tropolonato group to form a monodentate ligand (Fig. 1). A similar situation was observed in [Rh(tropBr₃)CO(PPh₃)] (trop is 2hydroxy-2,4,6-cycloheptatrienone) and [Rh(tropBr₃)CO-

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Figure 1

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



A fraction of the unit cell contents, showing the intermolecular hydrogen bonding and $\pi - \pi$ stacking as dashed lines [symmetry codes: (i) x, -y, z – $\frac{1}{2}$; (ii) -x, -y, -z].

(AsPh₃)] (Roodt et al., 2003), although a formal monodentate ligand system was not apparent. The monodentate tropolonate ligand is nearly perpendicular $[78.4 (1)^{\circ}]$ to the mean plane defined by the four atoms coordinating the palladium metal centre. Uncoordinated atom O21 lies 2.812 (2) Å from the Pd atom and is displaced from the apical position with a $Pd-O22 \cdots O21$ angle of 72.4 (2)°. The short interplanar distance of 3.51 (1) Å between the two bidentate tropolonate ligands related by the symmetry operation (-x, -y, -z)indicates the presence of π - π interactions.

In conclusion, the addition of tertiary phosphine to the bis(tropolonato)palladium(II) complex resulted in the weakening of the bidentate character of one tropolonato ligand. However, complete dissociation or the formation of a Pd-C(tropolonate) bond as found in acetylacetonate-type complexes (Ooi et al., 1983) were not observed.

Experimental

The title complex was synthesized by the addition of PCy₃ (88 mg, 0.316 mmol) to an acetone solution of the bis(tropolonato)palladium(II) complex (100 mg, 0.287 mmol). The solution changed colour from orange to red. On evaporation of the solvent, crystals suitable for X-ray diffraction were obtained (yield: 60 mg, 33%). ¹H NMR (CDCl₃, 600.28 MHz): δ 7.24 (d, 4), 7.17 (dd, 4), 6.74 (t, 2), 2.05 $(m, 3), 1.75 (m, 24), 1.25 (m, 6); {}^{13}C NMR (CDCl_3, 150.97 MHz); \delta$ 183.2 (C11, C12, C21, C22), 136.0 (C14, C16, C24, C26), 126.3 (C13, C17, C23, C27), 124.3 (C15, C25), 32.01 (d, C31, C41, C51), 28.27 (C32, C36, C42, C46, C52, C56), 27.45 (C33, C35, C43, C45, C53, C55), 26.25 (C34, C44, C54) with positions of C atoms confirmed by HSQC and HMBC data. Exchange is observed in CDCl₃ of tropolonate moieties. ³¹P NMR (CDCl₃, 242.98 MHz): δ 42.31 (s).

Crystal data

$[Pd(C_7H_5O_2)_2(C_{18}H_{33}P)]$	$D_x = 1.396 \text{ Mg m}^{-3}$
$M_r = 629.05$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 7765
a = 17.6699 (4) Å	reflections
b = 27.7962 (6) Å	$\theta = 2.5 - 28.2^{\circ}$
c = 12.2289 (2) Å	$\mu = 0.71 \text{ mm}^{-1}$
$\beta = 94.523 \ (1)^{\circ}$	T = 293 (2) K
V = 5987.6 (2) Å ³	Plate, red
Z = 8	$0.37 \times 0.31 \times 0.10 \text{ mm}$

Data collection

Bruker SMART CCD area-detector	7406 independent reflections
diffractometer	6361 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.030$
Absorption correction: multi-scan	$\theta_{\rm max} = 28.3^{\circ}$
(SADABS; Bruker, 1998)	$h = -23 \rightarrow 23$
$T_{\min} = 0.776, T_{\max} = 0.936$	$k = -37 \rightarrow 37$
81120 measured reflections	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.039P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.024$	+ 3.4366 <i>P</i>]
$wR(F^2) = 0.078$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.12	$(\Delta/\sigma)_{\rm max} = 0.001$
7406 reflections	$\Delta \rho_{\rm max} = 0.40 \ {\rm e} \ {\rm \AA}^{-3}$
343 parameters	$\Delta \rho_{\rm min} = -0.29 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

Pd-O12	1.9907 (13)	Pd-O11	2.0702 (12)
Pd-O22	2.0132 (13)	Pd-P	2.2342 (4)
O12-Pd-O11	80.30 (5)	O12-Pd-P	93.23 (4)
O22-Pd-O11	95.13 (5)	O22-Pd-P	91.32 (4)
011-C11-C12-O12	-6.6 (3)	O21-C21-C22-O22	5.3 (3)

Та	ble	2					
Hy	drog	gen-ł	oond ge	eometry	(Å, '	°).	
-							

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
C46-H46A···O21	0.97	2.43	3.348 (3)	159
$C13-H13\cdots O11^{i}$	0.93	2.56	3.456 (2)	161
$C15-H15\cdots O21^{ii}$	0.93	2.44	3.309 (3)	155

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

All H atoms were positioned geometrically and allowed to ride on their parent atoms, with C-H = 0.93–0.98 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$ of the parent atom.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT-Plus* (Bruker, 2004); data reduction: *SAINT-Plus*; 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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