## metal-organic compounds

Acta Crystallographica Section C Crystal Structure Communications

ISSN 0108-2701

# An organometallic compound with Z' = 4: {2-[2-(benzylideneamino)phenyl]-1,2-bis(methoxycarbonyl)ethenyl- $\kappa^2 C^1$ ,*N*}iodo(triphenylphosphine- $\kappa P$ )palladium(II)

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Received 29 November 2006 Accepted 30 November 2006 Online 23 December 2006

The title compound,  $[Pd(C_{19}H_{17}NO_4)I(C_{18}H_{15}P)]$ , crystallizes with four independent molecules in the asymmetric unit. The main difference between the molecules is the disposition of the PPh<sub>3</sub> ligand, for which in each molecule one ring is perpendicular to the ligand plane, but may be directed in either direction away from the plane; of the four molecules, two represent each possible direction. The independent molecules are arranged to form a chain parallel to [101] with an approximate translation of (a+c)/4 between successive molecules, excluding the PPh<sub>3</sub> rings. This leads to a systematic weakness of the reflections with  $h + l \neq 4n$ .

## Comment

The title compound, (I), was synthesized during a study of aryl palladium complexes bearing functionalized subbituents on the *ortho* position. In these investigations, oxidative addition of Pd<sup>0</sup> precursors to aryl halides yielded organometallic compounds that are models of intermediates in palladium-catalysed organic syntheses (Vicente *et al.*, 2005) and may lead to new coordination modes for the ligands. There are a large number of known palladium–diimine complexes; a search of the Cambridge Structural Database (CSD, Version 5.27; Allen, 2002) revealed 106 hits for  $\alpha$ -diimine complexes of palladium, *e.g.* de Pater *et al.* (2005). However, the double bond in the imine group of (I) does not form part of the heterocycle.

Compound (I) crystallizes in the solvent-free form with four independent molecules in the asymmetric unit. Molecules 1 and 3 are shown in Figs. 1 and 2, respectively; a similar numbering scheme was used for all the molecules, replacing the first digit '1' in molecule 1 with the appropriate molecule number 2, 3 or 4. In all four molecules, the Pd atom shows the expected planar four-coordination, albeit with slight but



significant deviations from planarity (r.m.s. deviations of the five atoms Pd, I, P, N and C from their least-squares plane are 0.01, 0.04, 0.05 and 0.08 Å, respectively, for molecules 1-4). The bond lengths at Pd are consistent over all four molecules (see below for one possible exception). The bite of the C,Nbidentate ligand is constant at ca 82° (important molecular dimensions are given in Table 1). The chelate rings display a configuration whereby atoms Pdn, Cn01, Cn02 and Cn12 are coplanar (r.m.s. deviations  $\leq 0.02$  Å for all four molecules), and the other two atoms, Nn and Cn11, are displaced from this plane to the same side. The double-bond positions Cn01 = Cn02 are unambiguously established by the bond lengths. The carboxylate groups are disposed such that the absolute torsion angles Cn02=Cn01-Cn03=On01 lie in the range 9–17° and Cn01=Cn02-Cn04=On02 in the range 90– 97°.

Despite the above-mentioned similarities, the molecules differ in one important respect, namely the disposition of the triphenylphosphine ligand. For each molecule, one phenyl ring of PPh<sub>3</sub> is approximately perpendicular to the coordination plane (*cf.* torsion angles  $I-Pd-P-C_{ipso}$  in Table 1; in each



## Figure 1

The first independent molecule of compound (I), showing the atomnumbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.

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case, there is one ring with an absolute value of  $ca 90^{\circ}$ ). However, in molecules 1 and 4, the ring at Cn31 points out of the coordination plane to the same side as ring Cn11, whereas in molecules 2 and 3 (Fig. 2), the ring at Cn51 points to the other side of the plane from Cn11, viz. to the same side as Cn21. The differing Pd-P bond lengths (slightly longer for molecules 2 and 3) and the markedly irregular Pd-P-C bond angles for molecules 1 and 4, with wide angles of ca 120° to Cn41 to avoid closer contact between the phenyl ring and the methoxycarbonyl group at Cn01, also reflect these differences. Even within this loose classification as two pairs of similar molecules, there are considerable differences in the I-Pd-



### Figure 2

The third independent molecule of compound (I), showing the atomnumbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.



#### Figure 3

The four independent molecules of (I) in the unit cell, running in order from molecule 1 (bottom right) to molecule 4 (top left). H atoms have been omitted.

P-C and Pd-P-C-C torsion angles for either given pair.

In the CSD, there are 25 examples of palladium complexes with a coordination sphere of C/N(chelating)/P/I. However, the bond lengths at Pd vary over a very wide range (e.g. Pd-N = 2.09-2.32 Å), making generalization difficult, and the iodo ligand may be trans to C or P. There are no examples with P trans to C for this donor atom set; consistent with the great extent of P/C 'transphobia' (Vicente et al., 1997, 2002, 2006), the P- and C-atom donors are mutually cis in (I) and in the other literature examples.

The molecular packing of (I) is noteworthy. The molecules are associated into chains parallel to [101] (Fig. 3), in which neighbouring molecules, except for the PPh<sub>3</sub> phenyl groups as noted above, adopt positions related by the non-crystallographic translation of (a+c)/4. This pseudosymmetry is responsible for the fact that reflections with  $h + l \neq 4n$  are systematically weak. However, the seven non-classical ('weak') hydrogen-bonding interactions (Desiraju & Steiner, 1999; Table 2), of which five involve para-H atoms of the PPh<sub>3</sub> ligands, are not formed within the chains.

## **Experimental**

Under a dinitrogen atmosphere, dimethyl acetylenedicarboxylate (DMAD) (94 µl, 0.7 mmol) was added to a solution of trans- $[(C_6H_4N=CH_2Ph-2)I(PPh_3)_2Pd]$  (Vicente *et al.*, 1999) (125 mg, 0.13 mmol) in degassed CH<sub>2</sub>Cl<sub>2</sub> (5 ml) and the reaction mixture was stirred overnight (16 h). The solvent was removed and the residue was treated with diethyl ether (5 ml). The solid was collected by filtration, washed with diethyl ether  $(3 \times 3 \text{ ml})$  and dried in air to yield complex (I) as a yellow powder (yield 80 mg, 73%). Single crystals were obtained by slow diffusion of Et<sub>2</sub>O into a solution of complex (I) in CDCl<sub>3</sub>. IR (Nujol, v, cm<sup>-1</sup>): 1726 (CO); <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>):  $\delta$  8.70 (*d*,  ${}^{3}J_{H,H}$  = 7 Hz, 2H), 8.40 (*d*,  ${}^{1}J_{H,H}$  = 13.5 Hz, 1H), 7.68–7.33 (several m, 22H, 4Ph + 2H<sub>arom</sub>), 3.69 (s, 3H, OMe), 3.06 (s, 3H, OMe); <sup>31</sup>P{<sup>1</sup>H} NMR (121 MHz, CDCl<sub>3</sub>): δ 31.44 (PPh<sub>3</sub>). Analysis calculated for C<sub>37</sub>H<sub>31</sub>INO<sub>4</sub>PPd: C 54.33, H 3.82, N 1.71%; found: C 54.53, H 3.86, N 1.67%.

## Crystal data

$[Pd(C_{19}H_{17}NO_4)I(C_{18}H_{15}P)]$	Z = 16
$M_r = 817.90$	$D_x = 1.611 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
a = 16.6938 (11)  Å	$\mu = 1.55 \text{ mm}^{-1}$
<i>b</i> = 22.1074 (15) Å	T = 133 (2) K
c = 36.666 (3) Å	Prism, yellow
$\beta = 94.460 \ (4)^{\circ}$	$0.38 \times 0.22 \times 0.20$ m
$V = 13490.8 (17) \text{ Å}^3$	

## Data collection

Bruker SMART 1000 CCD areadetector diffractometer  $\omega$  and  $\varphi$  scans Absorption correction: multi-scan (SADABS; Bruker, 1998)  $T_{\min} = 0.653, T_{\max} = 0.733$ 

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.030$  $wR(F^2) = 0.070$ S = 0.9539449 reflections 1629 parameters

m 261441 measured reflections 39449 independent reflections

27970 reflections with  $I > 2\sigma(I)$  $R_{\rm int}=0.050$  $\theta_{\rm max} = 30.0^\circ$ 

H-atom parameters constrained  $w = 1/[\sigma^{\bar{2}}(F_{o}^{2}) + (0.033P)^{2}]$ where  $P = (F_0^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} = 0.006$  $\Delta \rho_{\rm max} = 1.26 \ {\rm e} \ {\rm \AA}^{-3}$  $\Delta \rho_{\rm min} = -0.90 \text{ e } \text{\AA}^{-3}$ 

## Table 1

Selected geometric parameters (Å, °).

Pd1-C101	2.019(2)	Pd3-C301	2.023(2)
Pd1 - N1	2.0744(18)	Pd3_N3	20727(17)
Pd1 P1	2.0711 (10)	Pd3 P3	2.2700 (6)
Dd1 I1	2.2324(0)	Dd2 12	2.2700(0)
	2.0/10 (3)	Fu3-15	2.0393 (3)
C101-C102	1.338 (3)	C301-C302	1.344 (3)
Pd2-C201	2.020 (2)	Pd4-C401	2.024 (2)
Pd2-N2	2.0813 (19)	Pd4-N4	2.0654 (18)
Pd2-P2	2.2833 (6)	Pd4-P4	2.2660 (6)
Pd2-I2	2.6683 (3)	Pd4-I4	2.6877 (3)
C201-C202	1.346 (3)	C401-C402	1.347 (3)
C101 DJ1 N1	97.16(9)	C201 D42 M2	97.01 (9)
Clui Pil Pi	82.10 (8)	$C_{301} - P_{03} - N_3$	82.01 (8)
C101-Pd1-P1	93.37 (6)	C301-Pd3-P3	91.08 (6)
N1-Pd1-P1	165.77 (5)	N3-Pd3-P3	170.16 (5)
C101-Pd1-I1	168.71 (6)	C301-Pd3-I3	168.52 (6)
N1-Pd1-I1	88.46 (5)	N3-Pd3-I3	87.45 (5)
P1-Pd1-I1	94.253 (17)	P3-Pd3-I3	98.796 (17)
C131-P1-Pd1	108.64 (8)	C331-P3-Pd3	109.98 (7)
C141-P1-Pd1	120.28 (7)	C341-P3-Pd3	118.36 (8)
C151 - P1 - Pd1	112.94 (7)	C351 - P3 - Pd3	114 94 (8)
$C_{201} = Pd_2 = N_2$	81.95 (8)	C401 - Pd4 - N4	82.05 (8)
$C_{201} = P_{102} = P_{201}$	03.81(7)	$C_{401} = P_{44} = P_{44}$	02.03 (0) 05.77 (6)
N2 D42 D2	175 10 (5)		169 12 (5)
$N_2 - Pd_2 - P_2$	175.10 (5)	N4 - P04 - P4	108.13 (5)
C201-Pd2-12	1/0.30 (7)	C401-Pd4-14	167.88 (7)
N2-Pd2-I2	89.52 (5)	N4-Pd4-14	87.77 (5)
P2-Pd2-I2	94.495 (17)	P4-Pd4-I4	92.847 (16)
C231-P2-Pd2	114.28 (8)	C431-P4-Pd4	108.49 (8)
C241-P2-Pd2	115.85 (8)	C441-P4-Pd4	122.76 (7)
C251-P2-Pd2	112.67 (8)	C451-P4-Pd4	113.26 (7)
I1-Pd1-P1-C131	-87.82 (8)	I3-Pd3-P3-C331	-148.06 (8)
I1-Pd1-P1-C141	155.74 (9)	I3-Pd3-P3-C341	-29.18(9)
I1-Pd1-P1-C151	29.24 (9)	I3-Pd3-P3-C351	88.14 (9)
Pd1-P1-C131-C132	-174.63(19)	Pd3-P3-C331-C332	-125.79 (19)
Pd1 - P1 - C141 - C142	-111 48 (19)	Pd3 - P3 - C341 - C342	-1080(2)
Pd1 = P1 = C151 = C152	-110.40(18)	Pd3 = P3 = C351 = C352	-179.46(19)
$C_{102}^{102} = C_{101}^{101} = C_{102}^{102} = O_{102}^{101}$	110.40(10) 1 172(2)	$C_{202}^{201}$ $C_{201}^{202}$ $C_{202}^{201}$ $C_{202}^{202}$ $O_{202}^{201}$	179.40(1)
C102 - C101 - C103 - O10	1 17.2(3) 2 055(2)	$C_{302} = C_{301} = C_{303} = C_{30}$	1 13.1(4)
C101 - C102 - C104 - O10	2 = 93.3(3)	$C_{501} = C_{502} = C_{504} = O_{50}$	2 -94.4(3)
12-Pd2-P2-C231	-165.59 (9)	14-Pd4-P4-C431	-80.69 (8)
12 - Pd2 - P2 - C241	-43.87 (9)	14 - Pd4 - P4 - C441	162.79 (9)
12 - Pd2 - P2 - C251	74.91 (9)	I4-Pd4-P4-C451	38.86 (8)
Pd2-P2-C231-C232	-128.6(2)	Pd4-P4-C431-C432	-174.12 (17)
Pd2-P2-C241-C242	-93.2 (2)	Pd4-P4-C441-C442	-129.76 (17)
Pd2-P2-C251-C252	-169.81(19)	Pd4-P4-C451-C452	-113.42 (18)
C202-C201-C203-O20	1 17.3 (4)	C402-C401-C403-O40	1 9.4 (4)
C201-C202-C204-O20	2 -97.0 (3)	C401-C402-C404-O40	2 - 90.4(3)
	(. )		

Because of the large number of parameters, full-matrix refinement was not possible, and the refinement was therefore split into two blocks consisting of molecules 1 and 2, and molecules 3 and 4, respectively. Methyl H atoms were identified in difference syntheses, idealized, and refined as rigid groups allowed to rotate but not tip. Other H atoms were included using a riding model. C–H bond lengths were fixed at 0.98 (methyl C) or 0.95 Å ( $Csp^2$ ), and methyl H–C–H angles were fixed at 109.5°.  $U_{iso}$ (H) values were fixed at 1.2

## Table 2 Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C154-H154\cdots O401^{i}$	0.95	2.44	3.325 (3)	155
$C254 - H254 \cdots O402^{i}$	0.95	2.49	3.334 (3)	148
C313-H313···O302 <sup>ii</sup>	0.95	2.42	3.170 (3)	136
C354-H354O102 <sup>iii</sup>	0.95	2.59	3.536 (3)	174
$C454 - H454 \cdots O301^{iii}$	0.95	2.45	3.088 (3)	124
$C133-H133\cdots I3^{iv}$	0.95	2.99	3.805 (3)	145
$C444{-}H444{\cdot}{\cdot}{\cdot}I1^v$	0.95	2.99	3.780 (2)	141

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

times  $U_{eq}$  of the parent atom. The maximum residual electron density of 1.3 e Å<sup>-3</sup> is associated with the methoxy group at C401, and may indicate slight disorder of this group. The next five difference peaks (down to 0.8 e Å<sup>-3</sup>) all lie close to Pd or I atoms.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL97*.

The authors thank Professor J. Vicente, University of Murcia, for his continuing support and encouragement.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD3068). Services for accessing these data are described at the back of the journal.

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