

(Acetylacetonato)carbonyl[diallyl(*tert*-butyl)-phosphine]rhodium(I)

Edgar Ocando-Mavarez,^{a*} Julián Ascanio,^a Teresa González,^a Reinaldo Atencio,^a Merlín Rosales^b and Ninoska Silva^a

^aCentro de Química, Instituto Venezolano de Investigaciones Científicas (IVIC), Apartado 21827, Caracas 1020-A, Venezuela, and ^bLa Universidad del Zulia, Facultad Experimental de Ciencias, Departamento de Química, Laboratorio de Química Inorgánica, Apartado 526, Maracaibo, Venezuela

Correspondence e-mail: eocando@ivic.ve

Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$

R factor = 0.022

wR factor = 0.049

Data-to-parameter ratio = 14.7

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

The crystal structure of the title compound, $[\text{Rh}(\text{C}_5\text{H}_7\text{O}_2)\text{-force}]\text{b}(\text{C}_{10}\text{H}_{19}\text{P})(\text{CO})$, consists of discrete chiral molecules. The square-planar coordination geometry around Rh^{I} involves two O atoms of one acetylacetonate ligand, one carbonyl group and one P atom of a monodentate diallylphosphine. The double bonds of the allyl groups have a mutual $\pi-\pi$ interaction.

Comment

Transition metal complexes bearing functionalized phosphines are of interest because of their potential catalytic properties (Pruchnik *et al.*, 1998; Braunstein *et al.*, 2001; Appleby & Woollins, 2002; Jones *et al.*, 2002). In 1:1 complexes, β -functionalized phosphines can interact with the metal centre in three principal ways, as indicated in the *Scheme*.



In all these cases, the electronic and steric effects of the phosphine ligands may lead to transition metal complexes with useful new catalytic behaviour. As part of our work on the synthesis and reactivity of di- β -functionalized phosphines (Martín & Ocando-Mavarez, 1991; Ocando-Mavarez *et al.*, 1998), we have initiated studies on the use of diallyl-, di-cyanomethyl- and dipropargylphosphines and their interactions with metal centres. Interestingly, a considerable group of 32 metal compounds containing monoallylphosphines have been structurally characterized, according to the Cambridge Structural Database (Version 5.24, with updates to April 2003; Allen, 2002). In contrast, only one crystal structure of a

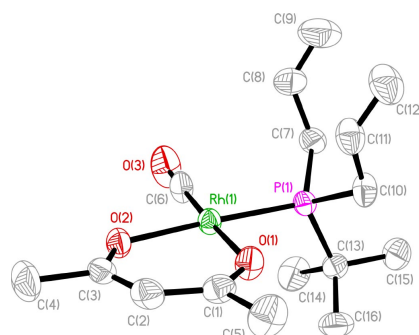
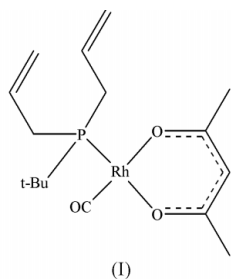


Figure 1

View of the title complex, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 35% probability level.

ruthenium complex (Jensen & Spicer, 2001) bearing a diallylphosphine is known so far. We report here the synthesis and crystal structure of the rhodium complex [(acac)Rh(CO)-forcebl][^tBuP(CH₂CH=CH₂)₂] (acac is acetylacetonate), (I).



The crystal structure of (I) has one molecule in the asymmetric unit (Fig. 1). The complex has no molecular or crystallographic symmetry, and the chiral molecule crystallizes in space group $P2_12_12_1$. The square-planar coordination environment around the metal centre consists of two O atoms of one acetylacetonate ligand, one carbonyl group and one P atom of a monodentate diallylphosphine. All non-H atoms of the acac ligand, together with atoms Rh, P and C6, lie in a plane, with a mean deviation of only 0.0247 Å. The Rh1—P1 distance [2.2387 (8) Å] is shorter than that observed in the ruthenium complex (2.326 Å) reported previously (Jensen & Spicer, 2001). A slightly asymmetric coordination of the acac ligand is observed, in which the Rh1—O2 distance [2.082 (2) Å] is longer than that for Rh1—O1 [2.032 (2) Å]. This asymmetry may be attributed to a *trans*-influence of the phosphine ligand, as has previously been invoked by Jensen & Spicer (2001). The conformation adopted by both allyl groups places the double bonds far away from the metal centre, precluding any intramolecular interaction between these bonds and Rh^I. It is noted, however, that the two allyl groups are approximately parallel to each other, with a centroid-to-centroid distance of only 3.569 Å, indicative of a π - π interaction.

Experimental

The synthesis of the diallylphosphine was carried out according to the procedure of Martín *et al.* (1992). Under vacuum (Burford *et al.*, 1994), 155 mg (0.911 mmol) of ^tBuP(CH₂CH=CH₂)₂ was distilled over a suspension of [(acac)Rh(CO)₂] (230 mg, 0.893 mmol) in CH₂Cl₂ (15 ml) and maintained under agitation at 195 K for 10 min. The cold bath was removed and the reaction mixture was stirred at room temperature for 30 min. The solvent was removed under vacuum and the solid residue was washed with pentane (5 ml) at 243 K, affording 310 mg [yield 87%; m.p. 351–353 K (with decomposition)] of a dark yellow solid. ¹H NMR (CDCl₃): 1.27 (*d*, ³J_{HP} = 14.19 Hz, 9H, ^tBu), 1.86 (*s*, 3H, CH₃—acac), 2.04 (*s*, 3H, CH₃—acac), 2.67 (*m*, 2H, P—CH₂), 2.83 (*m*, 2H, P—CH₂), 5.14 (*m*, 4H, C=CH₂), 5.43 (*s*, H—acac), 6.01 (*m*, 2H, CH=C); ¹³C NMR (CDCl₃): 27.13 (*s*, CH₃—acac), 27.64 (*bs*, CH₃—^tBu), 27.92 (*d*, ¹J_{CP} = 24.55 Hz, PCH₂), 32.99 (*d*, ¹J_{CP} = 26.42 Hz, C—^tBu), 100.53 (*s*, CH—acac), 118.52 (*s*, =CH₂), 132.20 (*s*, CH=), 184.88 (*s*, C—O—acac), 187.90 (*s*, C—O—acac), 189.80 (*dd*, ¹J_{CRh} = 75.96 Hz, ²J_{CP} = 24.27 Hz, CO); ³¹P NMR CDCl₃: 55.68 (*d*, ¹J_{PRh} = 170.00 Hz);

IR (cm⁻¹): 1962, 1579, 1520, 1378.

Crystal data

[Rh(C₅H₇O₂)(C₁₀H₁₉P)(CO)]
*M*_r = 400.25
 Orthorhombic, $P2_12_12_1$
a = 11.970 (2) Å
b = 12.270 (3) Å
c = 12.758 (3) Å
V = 1873.7 (7) Å³
Z = 4
*D*_x = 1.419 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 25 reflections
 θ = 26.4–31.7°
 μ = 1.00 mm⁻¹
T = 293 (2) K
 Prism, yellow
 0.36 × 0.30 × 0.15 mm

Data collection

Rigaku AFC-7S diffractometer
 ω -2 θ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
*T*_{min} = 0.766, *T*_{max} = 0.860
 3098 measured reflections
 2795 independent reflections
 2580 reflections with *I* > 2 σ (*I*)

*R*_{int} = 0.013
 θ _{max} = 25.0°
h = -10 → 14
k = 0 → 14
l = -1 → 15
 3 standard reflections
 every 150 reflections
 intensity decay: none

Refinement

Refinement on *F*²
R[*F*² > 2 σ (*F*²)] = 0.022
wR(*F*²) = 0.049
S = 1.05
 2795 reflections
 190 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0248P)^2 + 0.2043P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.25 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.21 \text{ e \AA}^{-3}$
 Absolute structure: Flack (1983),
 956 Friedel pairs
 Flack parameter = -0.05 (3)

Table 1

Selected geometric parameters (Å, °).

Rh1—C6	1.797 (4)	C1—C5	1.490 (5)
Rh1—O1	2.036 (2)	C2—C3	1.395 (5)
Rh1—O2	2.082 (2)	C3—C4	1.500 (5)
Rh1—P1	2.2387 (8)	C7—C8	1.493 (5)
P1—C10	1.826 (3)	C8—C9	1.267 (7)
P1—C7	1.834 (3)	C10—C11	1.487 (5)
P1—C13	1.869 (3)	C11—C12	1.270 (6)
O1—C1	1.272 (4)	C13—C16	1.528 (5)
O2—C3	1.264 (4)	C13—C15	1.529 (5)
O3—C6	1.141 (4)	C13—C14	1.536 (5)
C1—C2	1.377 (5)		
C6—Rh1—O1	177.45 (14)	C7—P1—C13	103.08 (17)
C6—Rh1—O2	94.38 (13)	C10—P1—Rh1	111.84 (13)
O1—Rh1—O2	88.03 (10)	C7—P1—Rh1	117.82 (11)
C6—Rh1—P1	89.44 (12)	O3—C6—Rh1	178.8 (3)
O1—Rh1—P1	88.15 (7)	C8—C7—P1	115.5 (3)
O2—Rh1—P1	176.15 (7)	C9—C8—C7	125.3 (5)
C10—P1—C7	104.8 (2)	C11—C10—P1	115.2 (2)
C10—P1—C13	104.03 (16)	C12—C11—C10	126.0 (4)

All H atoms were placed in idealized positions and constrained to ride on their parent atoms, with C—H distances of 0.96 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C).

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1991); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation, 1999); program(s) used to solve structure: *SHELXTL-NT* (Bruker, 1998); program(s) used to refine structure: *SHELXTL-NT*; molecular graphics: *SHELXTL-NT*; software used to prepare material for publication: *SHELXTL-NT*.

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