

Di- μ -carbonyl-bis[bis(triphenylphosphine)rhodium(0)]($Rh—Rh$) acetone disolvate¹

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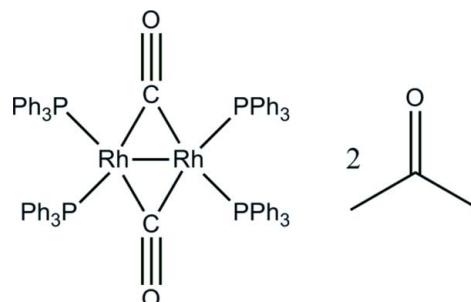
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(C—C) = 0.005$ Å; some non-H atoms missing; R factor = 0.032; wR factor = 0.082; data-to-parameter ratio = 18.2.

The dirhodium complex, $[Rh_2(C_{18}H_{15}P)_4(CO)_2] \cdot 2(CH_3)_2CO$, has crystallographic twofold symmetry and the $Rh—Rh$ distance is 2.6266 (8) Å. The four atoms proximate to each Rh atom [$Rh—P = 2.3222$ (7) and 2.3283 (8) Å, and $Rh—C = 1.961$ (3) and 2.045 (3) Å] form a distorted tetrahedron with large deviations from the putative tetrahedral angles [r.m.s. deviation = 23 (1)°]. The six angles more closely approximate those of a trigonal bipyramidal [r.m.s. deviation = 14 (1)°] with one missing equatorial ligand. The two bridging carbonyl ligands are much more linearly coordinated to one Rh [$Rh—C\equiv O = 151.0$ (2)°] than to the other [127.0 (2)°], and the two Rh_2CO planes form a dihedral angle of 45.43 (5)°. The two acetone solvent molecules are disordered, and their estimated scattering contribution was subtracted from the observed diffraction data using the SQUEEZE routine in PLATON [Spek (2009). *Acta Cryst. D* **65**, 148–155].

Related literature

For other dirhodium complex structures, see CCDC Refcode QAFHEM: Dzik *et al.* (2010), YOSMEZ: Okazaki *et al.* (2009), DEFJII: Douglas *et al.* (2005), TPCDRH10: Singh *et al.* (1973). For a description of the Cambridge Structural Database, see: Allen (2002). For the use of SQUEEZE, see: Spek (2009).



Experimental

Crystal data

$[Rh_2(C_{18}H_{15}P)_4(CO)_2] \cdot 2C_3H_6O$	$V = 6837.1$ (16) Å ³
$M_r = 1427.16$	$Z = 4$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 23.535$ (3) Å	$\mu = 0.63$ mm ⁻¹
$b = 13.0758$ (11) Å	$T = 298$ K
$c = 24.650$ (2) Å	$0.38 \times 0.38 \times 0.23$ mm
$\beta = 115.67$ (2)°	

Data collection

Enraf-Nonius CAD-4 diffractometer	6718 independent reflections
Absorption correction: ψ scan (North <i>et al.</i> , 1968)	5170 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.797$, $T_{\max} = 0.869$	$R_{\text{int}} = 0.026$
6874 measured reflections	3 standard reflections every 3 reflections

6718 independent reflections
5170 reflections with $I > 2\sigma(I)$
$R_{\text{int}} = 0.026$
3 standard reflections every 3 reflections
intensity decay: 4.0%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$	370 parameters
$wR(F^2) = 0.082$	H-atom parameters constrained
$S = 1.03$	$\Delta\rho_{\max} = 0.39$ e Å ⁻³
6718 reflections	$\Delta\rho_{\min} = -0.27$ e Å ⁻³

Data collection: *CAD-4 EXPRESS* (Enraf-Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008) and SQUEEZE in *PLATON* (Spek, 2009); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

The purchase of the diffractometer was made possible by a National Science Foundation chemical instrumentation grant, which we gratefully acknowledge. Improvements to the LSU X-ray Crystallography Facility were supported by grant No. LEQSF(1196–97)-ENH-TR-10, administered by the Louisiana Board of Regents.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5542).

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supporting information

Acta Cryst. (2012). E68, m1408–m1409 [doi:10.1107/S1600536812043528]

Di- μ -carbonyl-bis[bis(triphenylphosphane)rhodium(0)](*Rh*—*Rh*) acetone solvate

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S1. Comment

The dirhodium complex (I), $(\text{Rh}(\text{CO})(\text{PPh}_3)_2)_2$, is a precursor for $\text{HRh}(\text{CO})_2(\text{PPh}_3)_2$, a hydroformylation catalyst. A less precise crystal structure of I, as the dichloromethane solvate, was reported by Singh *et al.* (1973). In both solvates, the complex lies on a crystallographic twofold axis, with $\text{Rh}-\text{Rh} = 2.6266(8)$ Å (Singh *et al.*: 2.630(1) Å). The four atoms proximate to Rh ($\text{Rh}-\text{P}1 = 2.3222(7)$, $\text{Rh}-\text{P}2 = 2.3283(8)$, $\text{Rh}-\text{C}40 = 1.961(3)$, $\text{Rh}-\text{C}40' = 2.045(3)$ Å) form a distorted tetrahedron, but the six angles deviate markedly from the ideal tetrahedral angle ($\delta_{\text{r.m.s.}} = 23(1)^\circ$). The angles approximate more closely those of a trigonal bipyramidal ($\delta_{\text{r.m.s.}} = 14(1)^\circ$), with P1 and C40 in axial and P2 and C40' in equatorial positions, with one equatorial position vacant. The two bridging carbonyl ligands do not lie in the same plane, the two Rh_2CO planes forming a dihedral angle of 45.43(5)°. Furthermore, each carbonyl is asymmetrically coordinated to the two Rh atoms, with a $\text{Rh}-\text{C}$ distance of 1.961(3) Å and $\text{Rh}-\text{C}\equiv\text{O}$ angle 151.0(2)° to one Rh, 2.045(3) Å and 127.0(2)° to the other. This asymmetric carbonyl bridging is also seen in the DCM solvate structure (Singh *et al.*, 1973).

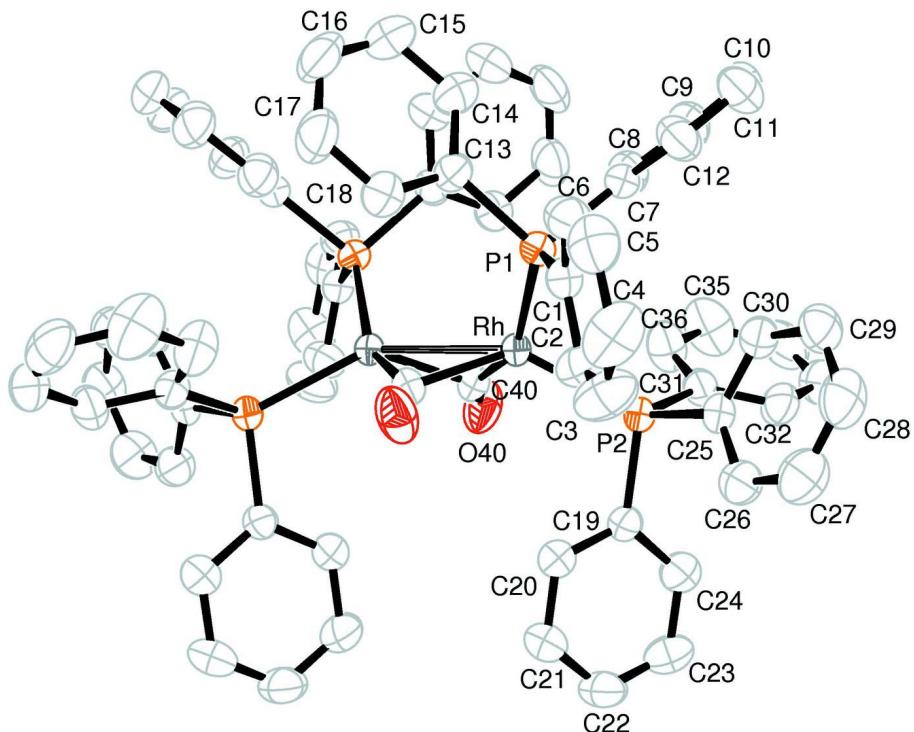
S2. Experimental

$\text{Rh}_2(\text{CO})_2(\text{PPh}_3)_4$ was synthesized from $\text{Rh}(\text{acac})(\text{CO})_2$ and excess triphenylphosphine under hydroformylation conditions, with acetone as solvent: A small autoclave was charged with a solution of $\text{Rh}(\text{acac})(\text{CO})_2$ (0.010 g), PPh_3 (9.750 g), and 1-hexene 7.883 g) in acetone (40 ml) while inside a glovebox. The autoclave was sealed, removed from the glovebox, and placed in a heating mantle. The headspace of the autoclave was purged with *syn* gas (1:1 $\text{H}_2:\text{CO}$, 3 x 60psig purges), and pressurized to 60psig with *syn* gas. The vessel was heated to 80°C and the *syn* gas pressure was adjusted to 80psig. After 18 h, the vessel was depressurized and purged with nitrogen (3x 60psig). The reaction mixture was analyzed with ^1H NMR: 91% aldehydes, 6.7% olefin isomerization, 2.2% 1-hexene. The n/i ratio of the aldehydes were 13.6. The reaction mixture slowly concentrated upon sitting in a glove box to afford a few red crystals of (I).

S3. Refinement

Each cavity (estimated volume 288 Å³) associated with a dirhodium molecule contains two disordered acetone molecules, for which no reasonable model could be developed. Therefore, the observed structure amplitudes were modified by PLATON/SQUEEZE (Spek, 2009) to subtract the scattering contribution of the electron density found in each cavity.

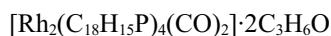
All H atoms were placed in calculated positions, with $\text{C}(sp^3)-\text{H} = 0.96$ Å, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$, and $\text{C}(sp^2)-\text{H} = 0.93$ Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$, and thereafter allowed to ride the attached C atom.

**Figure 1**

View of (I) (50% probability displacement ellipsoids). Unlabeled atoms are related by the symmetry operator ($-x+2, y, -z+3/2$).

Di- μ -carbonyl-bis[bis(triphenylphosphane)rhodium(0)]($Rh-Rh$) acetone disolvate

Crystal data



$M_r = 1427.16$

Monoclinic, $C2/c$

Hall symbol: -C 2yc

$a = 23.535 (3)$ Å

$b = 13.0758 (11)$ Å

$c = 24.650 (2)$ Å

$\beta = 115.67 (2)^\circ$

$V = 6837.1 (16)$ Å³

$Z = 4$

$F(000) = 2936$

$D_x = 1.386$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 25 reflections

$\theta = 2.6-27.5^\circ$

$\mu = 0.63$ mm⁻¹

$T = 298$ K

Prism, red

$0.38 \times 0.38 \times 0.23$ mm

Data collection

Enraf–Nonius CAD-4

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\theta/2\theta$ scans

Absorption correction: ψ scan

(North *et al.*, 1968)

$T_{\min} = 0.797$, $T_{\max} = 0.869$

6874 measured reflections

6718 independent reflections

5170 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.026$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 1.8^\circ$

$h = 0 \rightarrow 28$

$k = 0 \rightarrow 16$

$l = -30 \rightarrow 27$

3 standard reflections every 3 reflections

intensity decay: 4.0%

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.032$$

$$wR(F^2) = 0.082$$

$$S = 1.03$$

6718 reflections

370 parameters

0 restraints

0 constraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0423P)^2 + 0.3727P]$$
$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.003$$

$$\Delta\rho_{\max} = 0.39 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
P1	0.88187 (3)	-0.11053 (5)	0.72824 (3)	0.03726 (16)
C1	0.81798 (13)	-0.0719 (2)	0.74770 (11)	0.0410 (6)
C2	0.80127 (16)	0.0298 (2)	0.74235 (14)	0.0549 (8)
H2	0.8229	0.0768	0.73	0.066*
C3	0.7526 (2)	0.0623 (3)	0.75522 (17)	0.0752 (11)
H3	0.7419	0.1313	0.7518	0.09*
C4	0.71999 (19)	-0.0065 (3)	0.77297 (17)	0.0768 (11)
H4	0.6865	0.0155	0.7804	0.092*
C5	0.73669 (16)	-0.1068 (3)	0.77967 (14)	0.0621 (9)
H5	0.7148	-0.1534	0.792	0.075*
C6	0.78618 (15)	-0.1396 (2)	0.76812 (13)	0.0524 (7)
H6	0.7983	-0.2079	0.7741	0.063*
C7	0.84532 (13)	-0.19569 (19)	0.66282 (12)	0.0411 (6)
C8	0.87874 (15)	-0.2110 (2)	0.62867 (14)	0.0547 (8)
H8	0.9177	-0.1797	0.6398	0.066*
C9	0.85363 (17)	-0.2730 (3)	0.57799 (15)	0.0671 (10)
H9	0.8756	-0.2824	0.5548	0.081*
C10	0.79721 (18)	-0.3200 (3)	0.56208 (15)	0.0706 (10)
H10	0.7811	-0.3625	0.5285	0.085*
C11	0.76397 (18)	-0.3051 (3)	0.59513 (16)	0.0694 (10)
H11	0.7249	-0.3363	0.5835	0.083*
C12	0.78835 (15)	-0.2437 (2)	0.64586 (13)	0.0560 (8)
H12	0.7659	-0.2349	0.6686	0.067*
C13	0.92309 (13)	-0.2033 (2)	0.78900 (12)	0.0436 (6)
C14	0.93637 (16)	-0.3018 (2)	0.77782 (15)	0.0613 (9)
H14	0.9245	-0.3234	0.7384	0.074*
C15	0.96749 (18)	-0.3687 (3)	0.82533 (18)	0.0770 (11)
H15	0.9769	-0.4345	0.8174	0.092*

C16	0.98448 (17)	-0.3396 (3)	0.88318 (17)	0.0736 (11)
H16	1.0041	-0.3856	0.9146	0.088*
C17	0.97239 (17)	-0.2422 (3)	0.89466 (15)	0.0717 (10)
H17	0.9847	-0.2213	0.9342	0.086*
C18	0.94185 (15)	-0.1738 (3)	0.84792 (14)	0.0585 (8)
H18	0.934	-0.1074	0.8564	0.07*
P2	0.86831 (3)	0.09673 (5)	0.62360 (3)	0.03564 (16)
C19	0.88124 (12)	0.2359 (2)	0.63031 (12)	0.0401 (6)
C20	0.90876 (14)	0.2779 (2)	0.68686 (13)	0.0521 (8)
H20	0.9236	0.2352	0.7203	0.063*
C21	0.91480 (16)	0.3830 (2)	0.69503 (15)	0.0636 (9)
H21	0.9334	0.4101	0.7337	0.076*
C22	0.89343 (16)	0.4465 (2)	0.64633 (17)	0.0654 (9)
H22	0.8971	0.517	0.6518	0.079*
C23	0.86680 (18)	0.4065 (3)	0.58989 (17)	0.0717 (10)
H23	0.8526	0.4499	0.5567	0.086*
C24	0.86062 (16)	0.3016 (2)	0.58135 (14)	0.0593 (8)
H24	0.8425	0.275	0.5425	0.071*
C25	0.78347 (12)	0.0915 (2)	0.60350 (11)	0.0398 (6)
C26	0.74864 (14)	0.1729 (2)	0.60884 (13)	0.0523 (7)
H26	0.7679	0.2362	0.6214	0.063*
C27	0.68550 (16)	0.1614 (3)	0.59578 (16)	0.0680 (9)
H27	0.6628	0.2171	0.5995	0.082*
C28	0.65644 (16)	0.0694 (3)	0.57752 (16)	0.0685 (10)
H28	0.6143	0.0619	0.5697	0.082*
C29	0.68964 (15)	-0.0129 (3)	0.57065 (15)	0.0597 (8)
H29	0.6696	-0.0755	0.5574	0.072*
C30	0.75280 (14)	-0.0021 (2)	0.58356 (13)	0.0470 (7)
H30	0.775	-0.0577	0.5789	0.056*
C31	0.86728 (13)	0.0638 (2)	0.55073 (12)	0.0414 (6)
C32	0.81587 (15)	0.0855 (2)	0.49619 (12)	0.0545 (8)
H32	0.7809	0.1187	0.4962	0.065*
C33	0.81608 (19)	0.0587 (3)	0.44229 (14)	0.0676 (10)
H33	0.7815	0.0745	0.4063	0.081*
C34	0.8669 (2)	0.0088 (3)	0.44120 (16)	0.0728 (11)
H34	0.8673	-0.0083	0.4047	0.087*
C35	0.9171 (2)	-0.0156 (3)	0.49460 (19)	0.0835 (12)
H35	0.9514	-0.0503	0.4942	0.1*
C36	0.91725 (16)	0.0110 (3)	0.54926 (15)	0.0639 (9)
H36	0.9513	-0.007	0.5851	0.077*
Rh1	0.941218 (9)	0.013747 (14)	0.708963 (9)	0.03409 (7)
C40	1.01769 (13)	0.0593 (2)	0.70298 (12)	0.0435 (7)
O40	1.04318 (10)	0.09216 (19)	0.67507 (10)	0.0660 (6)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
P1	0.0398 (4)	0.0314 (3)	0.0343 (3)	-0.0018 (3)	0.0101 (3)	-0.0004 (3)

C1	0.0425 (15)	0.0420 (16)	0.0325 (13)	-0.0010 (12)	0.0106 (12)	-0.0010 (11)
C2	0.072 (2)	0.0469 (18)	0.0517 (18)	0.0104 (15)	0.0326 (16)	0.0081 (14)
C3	0.094 (3)	0.064 (2)	0.081 (3)	0.031 (2)	0.051 (2)	0.015 (2)
C4	0.075 (3)	0.101 (3)	0.071 (2)	0.027 (2)	0.048 (2)	0.016 (2)
C5	0.063 (2)	0.078 (2)	0.0519 (19)	-0.0085 (18)	0.0311 (17)	0.0064 (17)
C6	0.0587 (19)	0.0476 (17)	0.0506 (17)	-0.0074 (14)	0.0234 (15)	-0.0024 (14)
C7	0.0443 (15)	0.0309 (13)	0.0369 (14)	0.0001 (11)	0.0071 (12)	-0.0004 (11)
C8	0.0515 (18)	0.0554 (19)	0.0513 (18)	0.0051 (15)	0.0169 (15)	-0.0078 (14)
C9	0.074 (2)	0.068 (2)	0.0511 (19)	0.0114 (19)	0.0190 (17)	-0.0148 (17)
C10	0.080 (3)	0.056 (2)	0.0483 (19)	0.0008 (19)	0.0029 (18)	-0.0171 (16)
C11	0.072 (2)	0.059 (2)	0.063 (2)	-0.0199 (18)	0.0158 (19)	-0.0109 (17)
C12	0.061 (2)	0.0533 (19)	0.0499 (17)	-0.0183 (15)	0.0205 (15)	-0.0082 (14)
C13	0.0429 (15)	0.0394 (15)	0.0425 (15)	0.0003 (12)	0.0129 (13)	0.0071 (12)
C14	0.067 (2)	0.0473 (18)	0.0561 (19)	0.0123 (16)	0.0140 (16)	0.0040 (15)
C15	0.081 (3)	0.049 (2)	0.079 (3)	0.0132 (19)	0.014 (2)	0.0149 (19)
C16	0.063 (2)	0.067 (2)	0.073 (3)	0.0084 (18)	0.0131 (19)	0.035 (2)
C17	0.071 (2)	0.084 (3)	0.0462 (19)	0.000 (2)	0.0128 (17)	0.0166 (18)
C18	0.067 (2)	0.0502 (18)	0.0500 (18)	-0.0006 (16)	0.0172 (16)	0.0045 (14)
P2	0.0333 (3)	0.0350 (4)	0.0313 (3)	0.0018 (3)	0.0071 (3)	0.0008 (3)
C19	0.0349 (14)	0.0352 (14)	0.0426 (15)	0.0018 (11)	0.0097 (11)	0.0019 (11)
C20	0.0530 (18)	0.0422 (16)	0.0472 (17)	0.0042 (14)	0.0087 (14)	-0.0011 (13)
C21	0.067 (2)	0.0438 (18)	0.062 (2)	-0.0012 (16)	0.0106 (16)	-0.0124 (15)
C22	0.066 (2)	0.0346 (16)	0.087 (3)	-0.0006 (15)	0.025 (2)	-0.0032 (17)
C23	0.088 (3)	0.0410 (18)	0.075 (2)	0.0024 (17)	0.025 (2)	0.0164 (17)
C24	0.072 (2)	0.0459 (18)	0.0479 (18)	-0.0008 (16)	0.0146 (16)	0.0047 (14)
C25	0.0343 (14)	0.0432 (15)	0.0334 (13)	0.0033 (11)	0.0067 (11)	0.0032 (11)
C26	0.0454 (17)	0.0526 (18)	0.0557 (18)	0.0042 (14)	0.0188 (15)	-0.0013 (14)
C27	0.051 (2)	0.075 (2)	0.082 (2)	0.0142 (18)	0.0320 (19)	0.0062 (19)
C28	0.0387 (17)	0.086 (3)	0.076 (2)	-0.0043 (18)	0.0207 (17)	0.012 (2)
C29	0.0460 (17)	0.060 (2)	0.064 (2)	-0.0115 (15)	0.0156 (15)	0.0076 (16)
C30	0.0418 (15)	0.0487 (17)	0.0435 (15)	-0.0005 (13)	0.0118 (12)	0.0039 (12)
C31	0.0430 (15)	0.0404 (15)	0.0372 (14)	-0.0042 (12)	0.0141 (12)	-0.0014 (11)
C32	0.062 (2)	0.0548 (18)	0.0376 (15)	0.0089 (15)	0.0132 (14)	0.0042 (13)
C33	0.092 (3)	0.063 (2)	0.0361 (16)	-0.003 (2)	0.0164 (17)	0.0008 (15)
C34	0.096 (3)	0.081 (3)	0.051 (2)	-0.026 (2)	0.040 (2)	-0.0163 (18)
C35	0.071 (3)	0.111 (3)	0.080 (3)	-0.006 (2)	0.044 (2)	-0.027 (2)
C36	0.0499 (18)	0.085 (3)	0.0538 (19)	0.0008 (17)	0.0195 (15)	-0.0136 (17)
Rh1	0.03109 (11)	0.03049 (11)	0.03132 (11)	-0.00057 (8)	0.00472 (8)	0.00033 (8)
C40	0.0425 (16)	0.0366 (14)	0.0378 (14)	-0.0003 (12)	0.0047 (12)	0.0054 (12)
O40	0.0493 (13)	0.0884 (17)	0.0531 (13)	-0.0058 (12)	0.0154 (11)	0.0280 (12)

Geometric parameters (\AA , $^\circ$)

P1—C1	1.836 (3)	P2—Rh1	2.3283 (8)
P1—C7	1.838 (3)	C19—C20	1.372 (4)
P1—C13	1.843 (3)	C19—C24	1.386 (4)
P1—Rh1	2.3222 (7)	C20—C21	1.387 (4)
C1—C2	1.376 (4)	C20—H20	0.93

C1—C6	1.387 (4)	C21—C22	1.364 (5)
C2—C3	1.382 (5)	C21—H21	0.93
C2—H2	0.93	C22—C23	1.359 (5)
C3—C4	1.372 (5)	C22—H22	0.93
C3—H3	0.93	C23—C24	1.386 (4)
C4—C5	1.359 (5)	C23—H23	0.93
C4—H4	0.93	C24—H24	0.93
C5—C6	1.382 (4)	C25—C26	1.384 (4)
C5—H5	0.93	C25—C30	1.397 (4)
C6—H6	0.93	C26—C27	1.386 (4)
C7—C12	1.372 (4)	C26—H26	0.93
C7—C8	1.393 (4)	C27—C28	1.361 (5)
C8—C9	1.389 (4)	C27—H27	0.93
C8—H8	0.93	C28—C29	1.383 (5)
C9—C10	1.358 (5)	C28—H28	0.93
C9—H9	0.93	C29—C30	1.386 (4)
C10—C11	1.366 (5)	C29—H29	0.93
C10—H10	0.93	C30—H30	0.93
C11—C12	1.384 (4)	C31—C36	1.377 (4)
C11—H11	0.93	C31—C32	1.394 (4)
C12—H12	0.93	C32—C33	1.376 (4)
C13—C18	1.378 (4)	C32—H32	0.93
C13—C14	1.381 (4)	C33—C34	1.373 (5)
C14—C15	1.389 (4)	C33—H33	0.93
C14—H14	0.93	C34—C35	1.372 (6)
C15—C16	1.358 (5)	C34—H34	0.93
C15—H15	0.93	C35—C36	1.390 (5)
C16—C17	1.361 (5)	C35—H35	0.93
C16—H16	0.93	C36—H36	0.93
C17—C18	1.388 (4)	Rh1—C40	1.961 (3)
C17—H17	0.93	Rh1—C40 ⁱ	2.045 (3)
C18—H18	0.93	Rh1—Rh1 ⁱ	2.6266 (8)
P2—C31	1.837 (3)	C40—O40	1.173 (3)
P2—C25	1.838 (3)	C40—Rh1 ⁱ	2.045 (3)
P2—C19	1.841 (3)		
C1—P1—C7	105.72 (13)	C20—C19—P2	118.3 (2)
C1—P1—C13	99.86 (13)	C24—C19—P2	123.5 (2)
C7—P1—C13	101.28 (12)	C19—C20—C21	121.2 (3)
C1—P1—Rh1	119.62 (9)	C19—C20—H20	119.4
C7—P1—Rh1	109.80 (9)	C21—C20—H20	119.4
C13—P1—Rh1	118.38 (9)	C22—C21—C20	120.0 (3)
C2—C1—C6	118.3 (3)	C22—C21—H21	120
C2—C1—P1	118.2 (2)	C20—C21—H21	120
C6—C1—P1	123.4 (2)	C23—C22—C21	119.8 (3)
C1—C2—C3	120.4 (3)	C23—C22—H22	120.1
C1—C2—H2	119.8	C21—C22—H22	120.1
C3—C2—H2	119.8	C22—C23—C24	120.5 (3)

C4—C3—C2	120.3 (3)	C22—C23—H23	119.7
C4—C3—H3	119.8	C24—C23—H23	119.7
C2—C3—H3	119.8	C19—C24—C23	120.4 (3)
C5—C4—C3	120.0 (3)	C19—C24—H24	119.8
C5—C4—H4	120	C23—C24—H24	119.8
C3—C4—H4	120	C26—C25—C30	118.0 (3)
C4—C5—C6	120.0 (3)	C26—C25—P2	124.4 (2)
C4—C5—H5	120	C30—C25—P2	117.6 (2)
C6—C5—H5	120	C25—C26—C27	121.0 (3)
C5—C6—C1	120.8 (3)	C25—C26—H26	119.5
C5—C6—H6	119.6	C27—C26—H26	119.5
C1—C6—H6	119.6	C28—C27—C26	120.5 (3)
C12—C7—C8	118.9 (3)	C28—C27—H27	119.8
C12—C7—P1	124.7 (2)	C26—C27—H27	119.8
C8—C7—P1	116.4 (2)	C27—C28—C29	119.9 (3)
C9—C8—C7	119.7 (3)	C27—C28—H28	120
C9—C8—H8	120.1	C29—C28—H28	120
C7—C8—H8	120.1	C28—C29—C30	119.9 (3)
C10—C9—C8	120.4 (3)	C28—C29—H29	120
C10—C9—H9	119.8	C30—C29—H29	120
C8—C9—H9	119.8	C29—C30—C25	120.6 (3)
C9—C10—C11	120.3 (3)	C29—C30—H30	119.7
C9—C10—H10	119.9	C25—C30—H30	119.7
C11—C10—H10	119.9	C36—C31—C32	118.1 (3)
C10—C11—C12	120.1 (3)	C36—C31—P2	119.6 (2)
C10—C11—H11	120	C32—C31—P2	122.2 (2)
C12—C11—H11	120	C33—C32—C31	121.0 (3)
C7—C12—C11	120.6 (3)	C33—C32—H32	119.5
C7—C12—H12	119.7	C31—C32—H32	119.5
C11—C12—H12	119.7	C34—C33—C32	120.5 (3)
C18—C13—C14	118.4 (3)	C34—C33—H33	119.7
C18—C13—P1	119.2 (2)	C32—C33—H33	119.7
C14—C13—P1	122.5 (2)	C35—C34—C33	119.1 (3)
C13—C14—C15	120.1 (3)	C35—C34—H34	120.4
C13—C14—H14	120	C33—C34—H34	120.4
C15—C14—H14	120	C34—C35—C36	120.8 (4)
C16—C15—C14	121.1 (3)	C34—C35—H35	119.6
C16—C15—H15	119.4	C36—C35—H35	119.6
C14—C15—H15	119.4	C31—C36—C35	120.4 (3)
C15—C16—C17	119.2 (3)	C31—C36—H36	119.8
C15—C16—H16	120.4	C35—C36—H36	119.8
C17—C16—H16	120.4	C40—Rh1—C40 ⁱ	87.93 (14)
C16—C17—C18	120.7 (3)	C40—Rh1—P1	151.76 (8)
C16—C17—H17	119.7	C40 ⁱ —Rh1—P1	92.09 (9)
C18—C17—H17	119.7	C40—Rh1—P2	97.56 (8)
C13—C18—C17	120.6 (3)	C40 ⁱ —Rh1—P2	130.38 (8)
C13—C18—H18	119.7	P1—Rh1—P2	103.76 (3)
C17—C18—H18	119.7	C40—Rh1—Rh1 ⁱ	50.44 (8)

C31—P2—C25	100.08 (12)	C40 ⁱ —Rh1—Rh1 ⁱ	47.66 (8)
C31—P2—C19	104.56 (13)	P1—Rh1—Rh1 ⁱ	111.635 (19)
C25—P2—C19	100.32 (12)	P2—Rh1—Rh1 ⁱ	144.50 (2)
C31—P2—Rh1	117.82 (9)	O40—C40—Rh1	151.0 (2)
C25—P2—Rh1	121.02 (9)	O40—C40—Rh1 ⁱ	127.0 (2)
C19—P2—Rh1	110.55 (8)	Rh1—C40—Rh1 ⁱ	81.90 (11)
C20—C19—C24	118.0 (3)		
C7—P1—C1—C2	-115.7 (2)	Rh1—P2—C25—C26	107.8 (2)
C13—P1—C1—C2	139.5 (2)	C31—P2—C25—C30	61.5 (2)
Rh1—P1—C1—C2	8.7 (3)	C19—P2—C25—C30	168.5 (2)
C7—P1—C1—C6	65.1 (3)	Rh1—P2—C25—C30	-69.8 (2)
C13—P1—C1—C6	-39.7 (3)	C30—C25—C26—C27	1.1 (4)
Rh1—P1—C1—C6	-170.5 (2)	P2—C25—C26—C27	-176.5 (2)
C6—C1—C2—C3	-2.1 (5)	C25—C26—C27—C28	0.2 (5)
P1—C1—C2—C3	178.6 (3)	C26—C27—C28—C29	-1.4 (6)
C1—C2—C3—C4	-0.6 (6)	C27—C28—C29—C30	1.3 (5)
C2—C3—C4—C5	1.9 (6)	C28—C29—C30—C25	0.0 (5)
C3—C4—C5—C6	-0.5 (6)	C26—C25—C30—C29	-1.1 (4)
C4—C5—C6—C1	-2.2 (5)	P2—C25—C30—C29	176.6 (2)
C2—C1—C6—C5	3.5 (4)	C25—P2—C31—C36	-148.2 (3)
P1—C1—C6—C5	-177.3 (2)	C19—P2—C31—C36	108.3 (3)
C1—P1—C7—C12	-18.3 (3)	Rh1—P2—C31—C36	-14.9 (3)
C13—P1—C7—C12	85.5 (3)	C25—P2—C31—C32	28.3 (3)
Rh1—P1—C7—C12	-148.6 (2)	C19—P2—C31—C32	-75.2 (3)
C1—P1—C7—C8	161.5 (2)	Rh1—P2—C31—C32	161.6 (2)
C13—P1—C7—C8	-94.8 (2)	C36—C31—C32—C33	-2.6 (5)
Rh1—P1—C7—C8	31.2 (2)	P2—C31—C32—C33	-179.1 (3)
C12—C7—C8—C9	0.8 (4)	C31—C32—C33—C34	0.7 (5)
P1—C7—C8—C9	-179.0 (2)	C32—C33—C34—C35	1.1 (6)
C7—C8—C9—C10	-1.0 (5)	C33—C34—C35—C36	-1.0 (6)
C8—C9—C10—C11	1.3 (6)	C32—C31—C36—C35	2.7 (5)
C9—C10—C11—C12	-1.4 (6)	P2—C31—C36—C35	179.3 (3)
C8—C7—C12—C11	-1.0 (5)	C34—C35—C36—C31	-1.0 (6)
P1—C7—C12—C11	178.8 (2)	C1—P1—Rh1—C40	158.53 (19)
C10—C11—C12—C7	1.3 (5)	C7—P1—Rh1—C40	-79.1 (2)
C1—P1—C13—C18	-53.5 (3)	C13—P1—Rh1—C40	36.5 (2)
C7—P1—C13—C18	-161.9 (3)	C1—P1—Rh1—C40 ⁱ	69.02 (13)
Rh1—P1—C13—C18	78.1 (3)	C7—P1—Rh1—C40 ⁱ	-168.58 (12)
C1—P1—C13—C14	126.8 (3)	C13—P1—Rh1—C40 ⁱ	-53.06 (13)
C7—P1—C13—C14	18.5 (3)	C1—P1—Rh1—P2	-63.55 (10)
Rh1—P1—C13—C14	-101.5 (3)	C7—P1—Rh1—P2	58.85 (9)
C18—C13—C14—C15	0.5 (5)	C13—P1—Rh1—P2	174.37 (11)
P1—C13—C14—C15	-179.9 (3)	C1—P1—Rh1—Rh1 ⁱ	113.74 (9)
C13—C14—C15—C16	1.1 (6)	C7—P1—Rh1—Rh1 ⁱ	-123.86 (9)
C14—C15—C16—C17	-2.0 (6)	C13—P1—Rh1—Rh1 ⁱ	-8.34 (11)
C15—C16—C17—C18	1.4 (6)	C31—P2—Rh1—C40	59.88 (13)
C14—C13—C18—C17	-1.1 (5)	C25—P2—Rh1—C40	-176.87 (13)

P1—C13—C18—C17	179.3 (3)	C19—P2—Rh1—C40	−60.21 (13)
C16—C17—C18—C13	0.2 (5)	C31—P2—Rh1—C40 ⁱ	153.61 (15)
C31—P2—C19—C20	−154.4 (2)	C25—P2—Rh1—C40 ⁱ	−83.14 (15)
C25—P2—C19—C20	102.2 (2)	C19—P2—Rh1—C40 ⁱ	33.52 (15)
Rh1—P2—C19—C20	−26.6 (3)	C31—P2—Rh1—P1	−101.46 (10)
C31—P2—C19—C24	30.4 (3)	C25—P2—Rh1—P1	21.78 (11)
C25—P2—C19—C24	−73.0 (3)	C19—P2—Rh1—P1	138.45 (10)
Rh1—P2—C19—C24	158.1 (2)	C31—P2—Rh1—Rh1 ⁱ	82.89 (11)
C24—C19—C20—C21	1.2 (5)	C25—P2—Rh1—Rh1 ⁱ	−153.87 (10)
P2—C19—C20—C21	−174.3 (3)	C19—P2—Rh1—Rh1 ⁱ	−37.21 (11)
C19—C20—C21—C22	−0.3 (5)	C40 ⁱ —Rh1—C40—O40	−150.3 (5)
C20—C21—C22—C23	−0.6 (6)	P1—Rh1—C40—O40	119.1 (5)
C21—C22—C23—C24	0.6 (6)	P2—Rh1—C40—O40	−19.8 (5)
C20—C19—C24—C23	−1.2 (5)	Rh1 ⁱ —Rh1—C40—O40	177.3 (6)
P2—C19—C24—C23	174.0 (3)	C40 ⁱ —Rh1—C40—Rh1 ⁱ	32.40 (14)
C22—C23—C24—C19	0.3 (6)	P1—Rh1—C40—Rh1 ⁱ	−58.2 (2)
C31—P2—C25—C26	−120.9 (3)	P2—Rh1—C40—Rh1 ⁱ	162.88 (6)
C19—P2—C25—C26	−13.9 (3)		

Symmetry code: (i) $-x+2, y, -z+3/2$.