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trans-Carbonylchloridobis(tri-*o*-tolylphosphane- κ P)rhodium(I)

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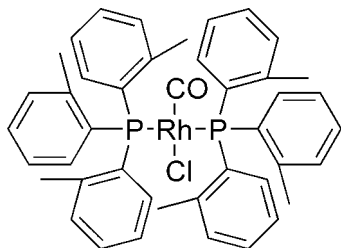
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; disorder in main residue; R factor = 0.049; wR factor = 0.143; data-to-parameter ratio = 19.1.

In the title compound, $[\text{RhCl}(\text{C}_{21}\text{H}_{21}\text{P})_2(\text{CO})]$, the coordination geometry around the Rh^{I} atom is slightly distorted square-planar with the phosphane ligands in *trans* positions with respect to each other. The chloride and carbonyl ligands show positional disorder, and the Rh^{I} atom lies on a center of inversion. The effective cone angle Θ_{E} for the title compound is $169.0(3)^\circ$. There are no significant intermolecular interactions.

Related literature

For background information, see: Angoletta (1959); Vaska & Di Luzio (1961); For a review of related compounds, see: Roodt *et al.* (2003). For related structures, see: Meijboom *et al.* (2005); Otto *et al.* (1999).



Experimental

Crystal data

 $[\text{RhCl}(\text{C}_{21}\text{H}_{21}\text{P})_2(\text{CO})]$
 $M_r = 775.07$

 Monoclinic, $P2_1/n$
 $a = 10.6440(14)$ Å
 $b = 10.9464(15)$ Å
 $c = 15.605(2)$ Å
 $\beta = 93.102(5)^\circ$
 $V = 1815.5(4)$ Å³
 $Z = 2$

 Mo $K\alpha$ radiation
 $\mu = 0.67$ mm⁻¹
 $T = 100$ K
 $0.30 \times 0.17 \times 0.12$ mm

Data collection

 Bruker APEXII CCD
 diffractometer
 Absorption correction: multi-scan
 (SADABS; Bruker, 2007)
 $T_{\text{min}} = 0.872$, $T_{\text{max}} = 0.921$

 21616 measured reflections
 4481 independent reflections
 3344 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.078$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.143$
 $S = 1.03$
 4481 reflections

 235 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 2.35$ e Å⁻³
 $\Delta\rho_{\text{min}} = -1.13$ e Å⁻³

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT-Plus (Bruker, 2007); data reduction: SAINT-Plus and XPREP (Bruker, 2007); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: WinGX (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2465).

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supporting information

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***trans*-Carbonylchloridobis(tri-*o*-tolylphosphane- κ P)rhodium(I)**

Stefan Warsink, Renier Koen and Andreas Roodt

S1. Comment

The complex first synthesized by Angoletta (1959) and later correctly formulated by Vaska (Vaska & Di Luzio, 1961), *trans*-[RhCl(CO)(PPh₃)₂] has become known under the name of the latter. This complex and its analogues have been extensively used as catalysts and model compounds (Roodt *et al.*, 2003).

Here we report a rhodium analogue bearing *o*-tolyl substituents on the phosphane ligands (I). As with many of these 'Vaska complexes', compound (I) crystallized with the metal on a crystallographic centre of symmetry, leading to a 50/50 disorder for the chloride and carbonyl ligands. With this report, Vaska complexes bearing all isomers of tritolylphosphane have been described. Compound (I) crystallizes in a slightly distorted square planar geometry with the phosphane ligands in *trans*-position to each other (Fig. 1).

The Rh—C and Rh—Cl bonds do not show large deviations from similar complexes, showing that the steric influence of the *o*-methyl substituents is not so significant as to distort the coordination geometry to a large degree. However, the Rh—P bond is longer than in similar complexes, indicating that the bulky *ortho*-aryl substituents force the phosphane ligands away from the rhodium. A useful indicator for the steric influence of phosphane ligands is the effective cone angle Θ_E . For compound (I) this angle was found to be 169.0 (3)°, significantly larger than differently substituted triarylphosphanes like tri(*m*-tolyl)-phosphane, for which values of 155° and 160° were reported (Meijboom *et al.*, 2005).

S2. Experimental

Compound (I) was synthesized by slow addition of 4 equivalents of tri(*o*-tolyl)phosphane to a dimethyl formamide solution of [RhCl(CO)₂]₂. After precipitation with ice water and separation of the product, it was recrystallized by slow evaporation from a 1:5 dichloromethane/hexane mixture. Analysis of the compound showed a CO stretching signal in IR at 1972 cm⁻¹, and a signal for the phosphane ligands in ³¹P NMR at 25.7 p.p.m. with a J_{Rh-P} of 125.9 Hz. These signals are in good agreement with various other rhodium Vaska's complexes.

S3. Refinement

The aromatic and methyl H atoms were placed in geometrically idealized positions (C—H = 0.93 and 0.96 Å, respectively) and constrained to ride on their parent atoms with $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic, and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H-atoms. The highest residual electron density was located 0.87 Å from Rh1 and was essentially meaningless.

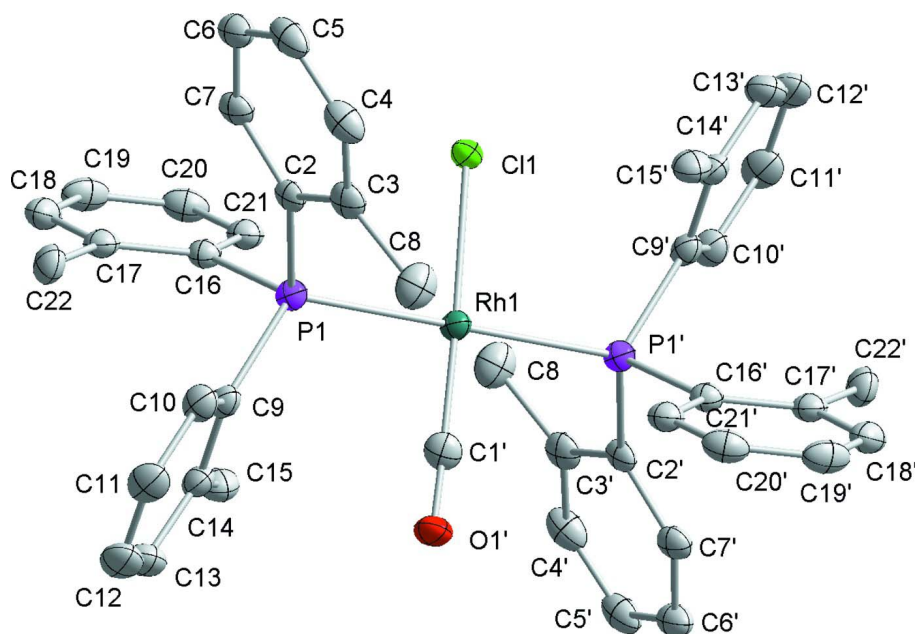


Figure 1

Molecular Structure of (I). Displacement ellipsoids are drawn at the 50% probability level [symmetry code: (i) $-x, -y, 1 - z$]. H atoms and disordered Chlorido and carbonyl ligands have been omitted for clarity.

trans-Carbonylchloridobis(tri-*o*-tolylphosphane- κ P)rhodium(I)

Crystal data

[RhCl(C₂₁H₂₁P)₂(CO)]

$M_r = 775.07$

Monoclinic, $P2_1/n$

Hall symbol: $-P 2_1n$

$a = 10.6440$ (14) Å

$b = 10.9464$ (15) Å

$c = 15.605$ (2) Å

$\beta = 93.102$ (5)°

$V = 1815.5$ (4) Å³

$Z = 2$

$F(000) = 800$

$D_x = 1.418$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71069$ Å

Cell parameters from 6713 reflections

$\theta = 2.3$ – 28.1 °

$\mu = 0.67$ mm⁻¹

$T = 100$ K

Cuboid, yellow

$0.30 \times 0.17 \times 0.12$ mm

Data collection

Bruker X8 APEXII 4K KappaCCD
diffractometer

Radiation source: sealed tube

Graphite monochromator

Detector resolution: 512 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2007)

$T_{\min} = 0.872$, $T_{\max} = 0.921$

21616 measured reflections

4481 independent reflections

3344 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.078$

$\theta_{\max} = 28.3$ °, $\theta_{\min} = 2.3$ °

$h = -12 \rightarrow 14$

$k = -13 \rightarrow 14$

$l = -20 \rightarrow 19$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.049$	H-atom parameters constrained
$wR(F^2) = 0.143$	$w = 1/[\sigma^2(F_o^2) + (0.0845P)^2 + 0.2003P]$
$S = 1.03$	where $P = (F_o^2 + 2F_c^2)/3$
4481 reflections	$(\Delta/\sigma)_{\max} = 0.001$
235 parameters	$\Delta\rho_{\max} = 2.35 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\min} = -1.13 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

Experimental. The intensity data was collected on a Bruker X8 Apex II 4 K Kappa CCD diffractometer using an exposure time of 60 s/frame. A total of 1376 frames was collected with a frame width of 0.5° covering up to $\theta=28.27^\circ$ with 99.5% completeness accomplished.

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	0.1009 (10)	-0.1237 (10)	0.4794 (7)	0.025 (2)	0.50
C2	0.2646 (3)	0.1270 (3)	0.4195 (2)	0.0193 (7)	
C3	0.3138 (3)	0.1696 (3)	0.4992 (2)	0.0249 (8)	
C4	0.4438 (4)	0.1739 (3)	0.5146 (3)	0.0294 (8)	
H10	0.4765	0.2061	0.5663	0.035*	
C5	0.5256 (4)	0.1321 (3)	0.4558 (3)	0.0312 (9)	
H6	0.6121	0.1359	0.4676	0.037*	
C6	0.4771 (3)	0.0841 (3)	0.3786 (2)	0.0279 (8)	
H15	0.5311	0.0537	0.3389	0.033*	
C7	0.3484 (3)	0.0814 (3)	0.3608 (2)	0.0230 (7)	
H9	0.3168	0.0488	0.3090	0.028*	
C8	0.2326 (4)	0.2095 (4)	0.5707 (2)	0.0345 (9)	
H24A	0.1527	0.2374	0.5467	0.052*	
H24B	0.2736	0.2747	0.6024	0.052*	
H24C	0.2198	0.1418	0.6083	0.052*	
C9	0.0342 (3)	0.2733 (3)	0.3794 (2)	0.0211 (7)	
C10	0.1102 (4)	0.3733 (3)	0.4027 (2)	0.0263 (8)	
H22	0.1910	0.3603	0.4266	0.032*	
C11	0.0664 (4)	0.4919 (3)	0.3905 (3)	0.0327 (9)	
H18	0.1174	0.5581	0.4062	0.039*	
C12	-0.0537 (4)	0.5105 (3)	0.3547 (3)	0.0362 (10)	
H4	-0.0836	0.5897	0.3462	0.043*	

C13	-0.1299 (4)	0.4121 (4)	0.3314 (3)	0.0319 (9)	
H19	-0.2102	0.4264	0.3070	0.038*	
C14	-0.0886 (3)	0.2918 (3)	0.3437 (2)	0.0238 (7)	
C15	-0.1760 (3)	0.1889 (4)	0.3142 (2)	0.0285 (8)	
H14A	-0.1528	0.1600	0.2592	0.043*	
H14B	-0.2611	0.2184	0.3097	0.043*	
H14C	-0.1695	0.1233	0.3550	0.043*	
C16	0.0791 (3)	0.0561 (3)	0.2837 (2)	0.0188 (7)	
C17	0.1169 (3)	0.1212 (3)	0.2114 (2)	0.0222 (7)	
C18	0.0976 (3)	0.0660 (4)	0.1308 (2)	0.0271 (8)	
H23	0.1215	0.1074	0.0822	0.033*	
C19	0.0436 (4)	-0.0488 (4)	0.1218 (2)	0.0305 (8)	
H17	0.0326	-0.0838	0.0676	0.037*	
C20	0.0061 (3)	-0.1115 (3)	0.1923 (2)	0.0286 (8)	
H13	-0.0311	-0.1880	0.1858	0.034*	
C21	0.0239 (3)	-0.0601 (3)	0.2729 (2)	0.0235 (7)	
H5	-0.0008	-0.1029	0.3206	0.028*	
C22	0.1748 (4)	0.2474 (3)	0.2150 (2)	0.0273 (8)	
H16A	0.2179	0.2619	0.1635	0.041*	
H16B	0.2334	0.2529	0.2638	0.041*	
H16C	0.1098	0.3073	0.2200	0.041*	
O1	0.1652 (12)	-0.2035 (10)	0.4677 (9)	0.030 (3)	0.50
P1	0.09397 (8)	0.11591 (8)	0.39409 (6)	0.0182 (2)	
Cl1	0.1376 (3)	-0.1671 (3)	0.4666 (2)	0.0226 (7)	0.50
Rh1	0.0000	0.0000	0.5000	0.01825 (13)	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.029 (6)	0.018 (6)	0.027 (5)	-0.004 (4)	0.001 (4)	-0.004 (4)
C2	0.0171 (15)	0.0122 (15)	0.0287 (17)	-0.0030 (12)	0.0015 (13)	0.0071 (13)
C3	0.0302 (19)	0.0125 (16)	0.0319 (19)	-0.0026 (14)	0.0020 (15)	0.0026 (14)
C4	0.0285 (19)	0.0198 (19)	0.039 (2)	-0.0063 (15)	-0.0085 (16)	0.0042 (16)
C5	0.0226 (18)	0.023 (2)	0.048 (2)	-0.0048 (15)	-0.0030 (16)	0.0143 (17)
C6	0.0220 (17)	0.0217 (19)	0.040 (2)	0.0005 (14)	0.0065 (15)	0.0088 (16)
C7	0.0245 (17)	0.0132 (16)	0.0314 (19)	-0.0003 (13)	0.0024 (14)	0.0032 (14)
C8	0.037 (2)	0.037 (2)	0.029 (2)	-0.0065 (18)	0.0011 (16)	-0.0055 (17)
C9	0.0238 (17)	0.0160 (16)	0.0244 (17)	-0.0018 (13)	0.0089 (13)	0.0005 (14)
C10	0.0303 (19)	0.0183 (18)	0.0309 (19)	-0.0033 (15)	0.0083 (15)	-0.0002 (15)
C11	0.041 (2)	0.0162 (18)	0.042 (2)	-0.0027 (16)	0.0131 (18)	-0.0035 (16)
C12	0.045 (2)	0.020 (2)	0.046 (2)	0.0091 (17)	0.016 (2)	0.0019 (17)
C13	0.032 (2)	0.025 (2)	0.040 (2)	0.0087 (16)	0.0080 (16)	0.0019 (17)
C14	0.0231 (17)	0.0233 (18)	0.0259 (18)	0.0018 (14)	0.0097 (14)	0.0017 (14)
C15	0.0215 (17)	0.031 (2)	0.033 (2)	0.0027 (15)	0.0019 (14)	0.0020 (16)
C16	0.0165 (15)	0.0171 (17)	0.0228 (16)	0.0001 (13)	0.0013 (12)	-0.0004 (14)
C17	0.0179 (16)	0.0210 (18)	0.0279 (18)	0.0032 (13)	0.0036 (13)	0.0002 (14)
C18	0.0267 (18)	0.029 (2)	0.0256 (18)	0.0086 (16)	0.0012 (14)	0.0007 (15)
C19	0.030 (2)	0.031 (2)	0.030 (2)	0.0074 (17)	-0.0033 (15)	-0.0135 (17)

C20	0.0250 (18)	0.0208 (18)	0.040 (2)	0.0014 (15)	-0.0034 (15)	-0.0072 (16)
C21	0.0197 (16)	0.0218 (19)	0.0292 (18)	0.0033 (14)	0.0028 (13)	0.0007 (15)
C22	0.0302 (19)	0.0255 (19)	0.0268 (19)	-0.0044 (15)	0.0079 (15)	0.0041 (15)
O1	0.028 (5)	0.020 (6)	0.041 (4)	0.007 (4)	0.011 (4)	0.006 (4)
P1	0.0184 (4)	0.0132 (4)	0.0233 (4)	-0.0029 (3)	0.0037 (3)	-0.0007 (3)
Cl1	0.025 (2)	0.016 (2)	0.0269 (13)	0.0042 (13)	0.0070 (13)	0.0036 (16)
Rh1	0.01767 (19)	0.0145 (2)	0.0230 (2)	-0.00172 (14)	0.00487 (14)	0.00057 (14)

Geometric parameters (Å, °)

C1—C11	0.654 (9)	C13—C14	1.399 (5)
C1—O1	1.131 (11)	C13—H19	0.9300
C1—Rh1	1.769 (11)	C14—C15	1.516 (5)
C2—C3	1.402 (5)	C15—H14A	0.9600
C2—C7	1.405 (5)	C15—H14B	0.9600
C2—P1	1.842 (3)	C15—H14C	0.9600
C3—C4	1.393 (5)	C16—C21	1.408 (5)
C3—C8	1.513 (5)	C16—C17	1.411 (5)
C4—C5	1.377 (6)	C16—P1	1.841 (3)
C4—H10	0.9300	C17—C18	1.401 (5)
C5—C6	1.387 (5)	C17—C22	1.513 (5)
C5—H6	0.9300	C18—C19	1.386 (6)
C6—C7	1.383 (5)	C18—H23	0.9300
C6—H15	0.9300	C19—C20	1.373 (6)
C7—H9	0.9300	C19—H17	0.9300
C8—H24A	0.9600	C20—C21	1.382 (5)
C8—H24B	0.9600	C20—H13	0.9300
C8—H24C	0.9600	C21—H5	0.9300
C9—C10	1.398 (5)	C22—H16A	0.9600
C9—C14	1.407 (5)	C22—H16B	0.9600
C9—P1	1.846 (4)	C22—H16C	0.9600
C10—C11	1.389 (5)	P1—Rh1	2.3496 (10)
C10—H22	0.9300	Cl1—Rh1	2.418 (3)
C11—C12	1.383 (7)	Rh1—C1 ⁱ	1.769 (11)
C11—H18	0.9300	Rh1—P1 ⁱ	2.3496 (10)
C12—C13	1.385 (6)	Rh1—Cl1 ⁱ	2.418 (3)
C12—H4	0.9300		
Cl1—C1—Rh1	172.6 (14)	H14A—C15—H14C	109.5
O1—C1—Rh1	178.8 (14)	H14B—C15—H14C	109.5
C3—C2—C7	118.4 (3)	C21—C16—C17	119.7 (3)
C3—C2—P1	122.0 (3)	C21—C16—P1	116.6 (2)
C7—C2—P1	119.3 (3)	C17—C16—P1	123.7 (3)
C4—C3—C2	119.0 (3)	C18—C17—C16	117.8 (3)
C4—C3—C8	117.7 (3)	C18—C17—C22	117.8 (3)
C2—C3—C8	123.3 (3)	C16—C17—C22	124.4 (3)
C5—C4—C3	122.1 (4)	C19—C18—C17	121.4 (3)
C5—C4—H10	119.0	C19—C18—H23	119.3

C3—C4—H10	119.0	C17—C18—H23	119.3
C4—C5—C6	119.0 (3)	C20—C19—C18	120.6 (3)
C4—C5—H6	120.5	C20—C19—H17	119.7
C6—C5—H6	120.5	C18—C19—H17	119.7
C7—C6—C5	120.0 (4)	C19—C20—C21	119.6 (4)
C7—C6—H15	120.0	C19—C20—H13	120.2
C5—C6—H15	120.0	C21—C20—H13	120.2
C6—C7—C2	121.2 (3)	C20—C21—C16	120.9 (3)
C6—C7—H9	119.4	C20—C21—H5	119.6
C2—C7—H9	119.4	C16—C21—H5	119.6
C3—C8—H24A	109.5	C17—C22—H16A	109.5
C3—C8—H24B	109.5	C17—C22—H16B	109.5
H24A—C8—H24B	109.5	H16A—C22—H16B	109.5
C3—C8—H24C	109.5	C17—C22—H16C	109.5
H24A—C8—H24C	109.5	H16A—C22—H16C	109.5
H24B—C8—H24C	109.5	H16B—C22—H16C	109.5
C10—C9—C14	120.2 (3)	C16—P1—C2	105.01 (15)
C10—C9—P1	120.5 (3)	C16—P1—C9	101.78 (16)
C14—C9—P1	119.3 (3)	C2—P1—C9	107.12 (15)
C11—C10—C9	120.7 (4)	C16—P1—Rh1	116.59 (11)
C11—C10—H22	119.6	C2—P1—Rh1	109.66 (11)
C9—C10—H22	119.6	C9—P1—Rh1	115.72 (11)
C12—C11—C10	119.3 (4)	C1—Rh1—C1 ⁱ	180.000 (2)
C12—C11—H18	120.4	C1—Rh1—P1 ⁱ	90.0 (4)
C10—C11—H18	120.4	C1 ⁱ —Rh1—P1 ⁱ	90.0 (4)
C11—C12—C13	120.5 (4)	C1—Rh1—P1	90.0 (4)
C11—C12—H4	119.8	C1 ⁱ —Rh1—P1	90.0 (4)
C13—C12—H4	119.8	P1 ⁱ —Rh1—P1	180.0
C12—C13—C14	121.4 (4)	C1—Rh1—Cl1	2.0 (4)
C12—C13—H19	119.3	C1 ⁱ —Rh1—Cl1	178.0 (4)
C14—C13—H19	119.3	P1 ⁱ —Rh1—Cl1	91.66 (10)
C13—C14—C9	117.9 (3)	P1—Rh1—Cl1	88.34 (10)
C13—C14—C15	118.3 (3)	C1—Rh1—Cl1 ⁱ	178.0 (4)
C9—C14—C15	123.7 (3)	C1 ⁱ —Rh1—Cl1 ⁱ	2.0 (4)
C14—C15—H14A	109.5	P1 ⁱ —Rh1—Cl1 ⁱ	88.34 (10)
C14—C15—H14B	109.5	P1—Rh1—Cl1 ⁱ	91.66 (10)
H14A—C15—H14B	109.5	Cl1—Rh1—Cl1 ⁱ	180.00 (18)
C14—C15—H14C	109.5		

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