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# *cis*-Dichlorido[2-methyl-8-(pyridin-2-yl-methoxy)quinoline- $\kappa^3N,O,N'$ ](triphenylphosphane- $\kappa P$ )ruthenium(II) methanol monosolvate

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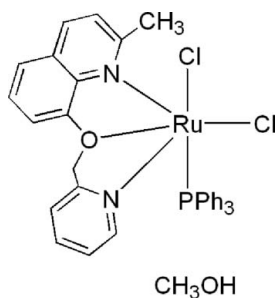
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Key indicators: single-crystal X-ray study;  $T = 291$  K; mean  $\sigma(C-C) = 0.008$  Å;  $R$  factor = 0.049;  $wR$  factor = 0.117; data-to-parameter ratio = 16.1.

In the structure of the title compound,  $[RuCl_2(C_{16}H_{14}N_2O)(C_{18}H_{15}P)] \cdot CH_3OH$ , the  $Ru^{II}$  ion shows a slightly distorted octahedral coordination by two N atoms and one O atom from the 2-methyl-8-(pyridin-2-ylmethoxy)quinoline acting as an  $N,O,N'$ -tridentate ligand, two Cl atoms, and one P atom from a  $PPh_3$  ligand. The two Cl atoms adopt a *cis* arrangement with the  $PPh_3$  ligand *trans* to one Cl atom. The  $N,O,N'$ -tridentate ligand occupies a *mer* position in the coordination sphere.

## Related literature

For related structures, see: Al-Mandhary & Steel (2003); Deng *et al.* (2005); Xu *et al.* (2009).



## Experimental

### Crystal data

$[RuCl_2(C_{16}H_{14}N_2O)(C_{18}H_{15}P)] \cdot CH_3O$   
 $M_r = 716.57$   
 Orthorhombic,  $P2_12_12_1$   
 $a = 8.868$  (2) Å  
 $b = 11.480$  (2) Å  
 $c = 31.6351$  (18) Å

$V = 3220.7$  (10) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.74$  mm<sup>-1</sup>  
 $T = 291$  K  
 $0.28 \times 0.24 \times 0.22$  mm

### Data collection

Bruker SMART APEX CCD diffractometer  
 Absorption correction: multi-scan (SADABS; Sheldrick, 2003)  
 $T_{min} = 0.820$ ,  $T_{max} = 0.855$

17669 measured reflections  
 6304 independent reflections  
 5620 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.030$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.049$   
 $wR(F^2) = 0.117$   
 $S = 1.10$   
 6304 reflections  
 391 parameters  
 H-atom parameters constrained

$\Delta\rho_{max} = 0.70$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.91$  e Å<sup>-3</sup>  
 Absolute structure: Flack (1983),  
 2722 Friedel pairs  
 Flack parameter: 0.06 (4)

Data collection: SMART (Bruker, 2003); cell refinement: SAINT-Plus (Bruker, 2003); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: JH2228).

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## supporting information

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## *cis*-Dichlorido[2-methyl-8-(pyridin-2-ylmethoxy)quinoline- $\kappa^3N,O,N'$ ](triphenylphosphane- $\kappa P$ )ruthenium(II) methanol monosolvate

Hui-Jun Xu, Yu Li and Qing-Yang Du

### S1. Comment

The *N,O,N'*-tridentate ligands with two pyridine-like donors and an ether donor are potentially a doubly chelating ligands in coordination chemistry. The presence of the flexible methylene and ether linkages allows the ligands to act as a folded *N,N'*-bidentate ligands or *N,O,N'*-tridentate ligands in meridional or facial arrangement, coordination to the metal coordinate (Al-Mandhary & Steel, 2003; Xu *et al.* 2009). Here, we report the synthesis and crystal structure of the title complex 1, [Ru(C<sub>16</sub>H<sub>14</sub>N<sub>2</sub>O)(PPh<sub>3</sub>)Cl<sub>2</sub>.CH<sub>3</sub>OH], which combines 2-methyl-8-(pyridineyl-2-methoxy)-quinoline and triphenylphosphane ligands. The molecular structure of the title compound is shown in Fig. 1. In the title complex 1, the ruthenium atom center is in a pseudooctahedral environment with the two nitrogen atoms and one oxygen atom which from 2-methyl-8-(pyridineyl-2-methoxy)-quinoline acts as a *mer N,O,N'*-tridentate ligand, two *cis* chlorine atoms and one phosphorus atom from PPh<sub>3</sub> ligand *trans* to one chloride. The N—Ru—N angle is 158.25 (17)°. The N1—Ru—O1 and N2—Ru—O1 angles are 79.47 (17)° and 79.66 (16)°, respectively. The coordination between *N,O,N'*-tridentate ligand and the Ru<sup>II</sup> yields two five-membered rings, RuN1C6C5O1 and RuO1C7C12N2. The Ru—Cl1 distance [2.4905 (14) Å], which is *trans* to the PPh<sub>3</sub> ligand, is longer than the Ru—Cl2 distance [2.4104 (14) Å], similar differences are in agreement with reported value (Deng *et al.* 2005; Xu *et al.* 2009). The Ru—*N*(pyridine) [2.067 (5) Å], Ru—*N*(quinoline) [2.163 (5) Å], Ru—O [2.060 (3) Å] and Ru—P [2.2931 (15) Å] are similar to reported value (Deng *et al.* 2005; Xu *et al.* 2009).

### S2. Experimental

The synthesis of the title compound 1 was carried out as follows. 2-methyl-8-(pyridineyl-2-methoxy)-quinoline (0.286 g, 0.55 mmol) was added to a solution of RuCl<sub>2</sub>(PPh<sub>3</sub>)<sub>3</sub> (0.491 g, 0.50 mmol) in CH<sub>3</sub>OH (25 ml). This mixture was refluxed for 8 h and then evaporated to dryness. The orange solid residue was dissolved in CH<sub>2</sub>Cl<sub>2</sub> (*ca* 3 ml) and the resulting solution was transferred to a silica gel chromatography column. Elution with CH<sub>2</sub>Cl<sub>2</sub>/CH<sub>3</sub>OH (40:1) gave a yellow-orange band, from which complex 1 was obtained after solvent removal. Crystal of 1 suitable for X-ray structure determination was grown from CH<sub>3</sub>OH solution of the complex layered with ethyl ether.

### S3. Refinement

H atoms on C atoms were placed in idealized positions (C—H = 0.93—0.97 Å) and refined as riding atoms, with the  $U_{\text{iso}}(\text{H}) = 1.2$  or  $1.5U_{\text{eq}}(\text{C})$ . H atom attached to O atom are located in a difference Fourier map and refined as riding in their 'as found' positions with the  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ .

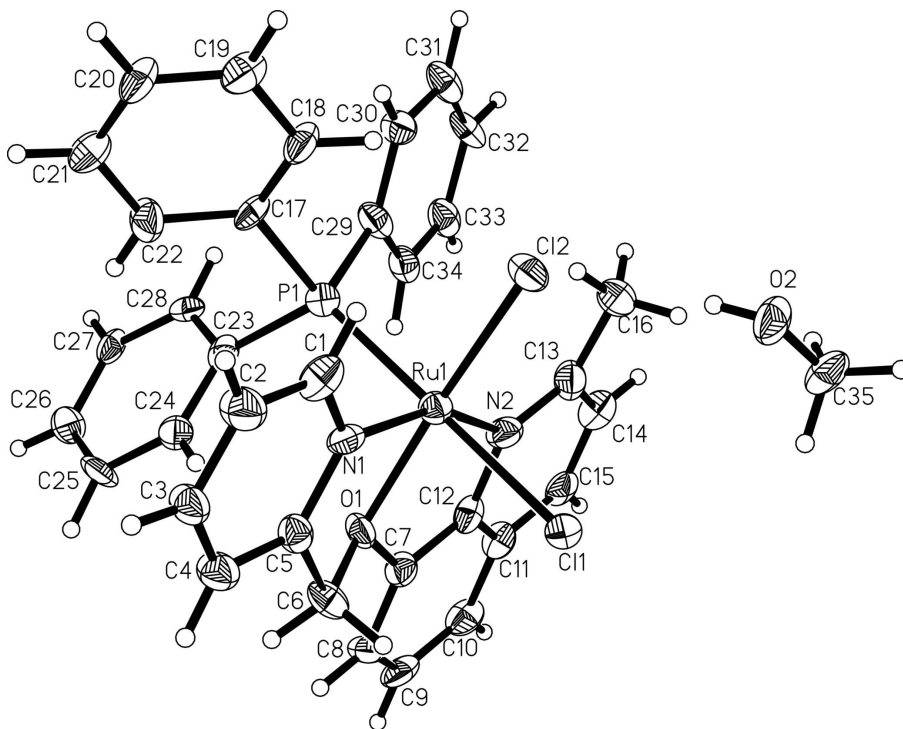


Figure 1

The structure of 1, with atom labels and 30% probability displacement ellipsoids.

***cis*-Dichlorido[2-methyl-8-(pyridin-2-ylmethoxy)quinoline- $\kappa^3N,O,N'$ ](triphenylphosphane- $\kappa P$ )ruthenium(II) methanol monosolvate**

*Crystal data*

[RuCl<sub>2</sub>(C<sub>16</sub>H<sub>14</sub>N<sub>2</sub>O)(C<sub>18</sub>H<sub>15</sub>P)]·CH<sub>4</sub>O

$M_r = 716.57$

Orthorhombic,  $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 8.868$  (2) Å

$b = 11.480$  (2) Å

$c = 31.6351$  (18) Å

$V = 3220.7$  (10) Å<sup>3</sup>

$Z = 4$

$F(000) = 1464$

$D_x = 1.478$  Mg m<sup>-3</sup>

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

Cell parameters from 592 reflections

$\theta = 2.4$ – $13.2^\circ$

$\mu = 0.74$  mm<sup>-1</sup>

$T = 291$  K

Block, orange

$0.28 \times 0.24 \times 0.22$  mm

*Data collection*

Bruker SMART APEX CCD  
diffractometer

Radiation source: sealed tube

Graphite monochromator

phi and  $\omega$  scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 2003)

$T_{\min} = 0.820$ ,  $T_{\max} = 0.855$

17669 measured reflections

6304 independent reflections

5620 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$

$\theta_{\max} = 26.0^\circ$ ,  $\theta_{\min} = 1.9^\circ$

$h = -10 \rightarrow 9$

$k = -14 \rightarrow 10$

$l = -38 \rightarrow 38$

*Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.049$  $wR(F^2) = 0.117$  $S = 1.10$ 

6304 reflections

391 parameters

0 restraints

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

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.06P)^2 + 1.99P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\max} < 0.001$  $\Delta\rho_{\max} = 0.70 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{\min} = -0.91 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), 2722 Friedel  
pairs

Absolute structure parameter: 0.06 (4)

*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.

**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  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.1967 (7)	0.4572 (6)	0.6458 (2)	0.0547 (15)
H1	0.1733	0.4993	0.6216	0.066*
C2	0.0977 (8)	0.3754 (6)	0.6601 (2)	0.0547 (16)
H2	0.0091	0.3628	0.6451	0.066*
C3	0.1231 (7)	0.3129 (5)	0.69486 (18)	0.0481 (14)
H3	0.0548	0.2561	0.7033	0.058*
C4	0.2514 (7)	0.3333 (5)	0.71807 (18)	0.0482 (14)
H4	0.2707	0.2934	0.7431	0.058*
C5	0.3511 (6)	0.4166 (5)	0.70246 (16)	0.0396 (11)
C6	0.4939 (7)	0.4460 (5)	0.72707 (17)	0.0455 (12)
H6A	0.4734	0.5064	0.7477	0.055*
H6B	0.5309	0.3776	0.7418	0.055*
C7	0.7414 (7)	0.5321 (5)	0.71092 (17)	0.0406 (12)
C8	0.8219 (7)	0.4838 (5)	0.74429 (15)	0.0424 (13)
H8	0.7791	0.4250	0.7606	0.051*
C9	0.9665 (7)	0.5233 (5)	0.75332 (18)	0.0512 (17)
H9	1.0204	0.4910	0.7757	0.061*
C10	1.0306 (6)	0.6112 (5)	0.72899 (18)	0.0499 (15)
H10	1.1273	0.6376	0.7350	0.060*
C11	0.9500 (6)	0.6594 (5)	0.69561 (19)	0.0416 (13)
C12	0.8054 (6)	0.6199 (5)	0.68658 (18)	0.0409 (12)
C13	0.7889 (7)	0.7560 (5)	0.6289 (2)	0.0498 (15)
C14	0.9335 (6)	0.7956 (5)	0.63790 (18)	0.0481 (14)

H14	0.9764	0.8543	0.6216	0.058*
C15	1.0141 (8)	0.7473 (4)	0.67127 (15)	0.0461 (13)
H15	1.1108	0.7737	0.6773	0.055*
C16	0.6974 (8)	0.8152 (6)	0.59660 (19)	0.0556 (16)
H16A	0.6168	0.7648	0.5878	0.083*
H16B	0.7595	0.8337	0.5727	0.083*
H16C	0.6561	0.8855	0.6082	0.083*
C17	0.4278 (6)	0.3833 (5)	0.56457 (16)	0.0389 (12)
C18	0.3214 (7)	0.4467 (5)	0.54310 (19)	0.0492 (15)
H18	0.3255	0.5276	0.5434	0.059*
C19	0.2062 (7)	0.3896 (6)	0.52066 (19)	0.0520 (16)
H19	0.1352	0.4319	0.5054	0.062*
C20	0.2012 (7)	0.2744 (5)	0.5218 (2)	0.0510 (15)
H20	0.1250	0.2367	0.5069	0.061*
C21	0.3002 (7)	0.2094 (6)	0.54320 (19)	0.0514 (15)
H21	0.2917	0.1287	0.5429	0.062*
C22	0.4163 (8)	0.2625 (5)	0.5660 (2)	0.0526 (15)
H22	0.4840	0.2183	0.5818	0.063*
C23	0.6972 (6)	0.3305 (5)	0.61146 (17)	0.0397 (12)
C24	0.6582 (7)	0.2702 (5)	0.64730 (19)	0.0469 (13)
H24	0.5781	0.2966	0.6637	0.056*
C25	0.7358 (8)	0.1700 (5)	0.65976 (17)	0.0490 (15)
H25	0.7136	0.1346	0.6855	0.059*
C26	0.8444 (8)	0.1244 (5)	0.63414 (19)	0.0529 (16)
H26	0.8935	0.0559	0.6416	0.063*
C27	0.8805 (6)	0.1802 (4)	0.59724 (17)	0.0401 (12)
H27	0.9554	0.1497	0.5799	0.048*
C28	0.8063 (7)	0.2826 (5)	0.58537 (16)	0.0407 (12)
H28	0.8302	0.3186	0.5599	0.049*
C29	0.6986 (7)	0.5251 (5)	0.55537 (18)	0.0456 (14)
C30	0.6518 (7)	0.5601 (5)	0.51666 (17)	0.0462 (14)
H30	0.5546	0.5422	0.5077	0.055*
C31	0.7457 (7)	0.6215 (5)	0.4905 (2)	0.0510 (16)
H31	0.7118	0.6441	0.4639	0.061*
C32	0.8927 (7)	0.6507 (5)	0.50324 (19)	0.0485 (15)
H32	0.9544	0.6944	0.4855	0.058*
C33	0.9447 (7)	0.6144 (5)	0.54189 (19)	0.0499 (15)
H33	1.0441	0.6272	0.5500	0.060*
C34	0.8443 (7)	0.5582 (5)	0.56832 (18)	0.0446 (14)
H34	0.8742	0.5415	0.5958	0.054*
C35	0.3573 (7)	1.0239 (6)	0.6490 (2)	0.0559 (16)
H35A	0.3218	1.1028	0.6508	0.084*
H35B	0.4463	1.0214	0.6317	0.084*
H35C	0.3806	0.9956	0.6768	0.084*
Cl1	0.43030 (16)	0.71863 (12)	0.70588 (4)	0.0412 (3)
Cl2	0.35481 (17)	0.69468 (12)	0.60054 (5)	0.0474 (3)
N1	0.3286 (5)	0.4787 (4)	0.66606 (13)	0.0385 (10)
N2	0.7249 (5)	0.6682 (4)	0.65321 (14)	0.0433 (11)

O1	0.6057 (4)	0.4866 (3)	0.69627 (11)	0.0388 (8)
O2	0.2468 (5)	0.9549 (4)	0.63110 (15)	0.0588 (12)
H2A	0.2852	0.8955	0.6216	0.071*
P1	0.57943 (18)	0.45692 (12)	0.59589 (4)	0.0401 (3)
Ru1	0.50826 (5)	0.58272 (3)	0.648543 (12)	0.03625 (11)

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.050 (3)	0.057 (4)	0.057 (4)	-0.008 (3)	-0.002 (3)	-0.018 (3)
C2	0.056 (4)	0.058 (4)	0.051 (4)	-0.014 (3)	-0.004 (3)	0.008 (3)
C3	0.054 (4)	0.044 (3)	0.047 (3)	-0.016 (3)	-0.014 (3)	0.011 (3)
C4	0.051 (3)	0.050 (3)	0.043 (3)	-0.005 (3)	-0.011 (3)	0.008 (3)
C5	0.041 (3)	0.035 (3)	0.044 (3)	0.004 (3)	0.004 (2)	0.001 (2)
C6	0.039 (3)	0.049 (3)	0.049 (3)	0.012 (3)	-0.001 (3)	0.018 (2)
C7	0.042 (3)	0.036 (3)	0.044 (3)	-0.002 (2)	-0.012 (2)	-0.004 (2)
C8	0.049 (3)	0.051 (3)	0.027 (3)	-0.012 (3)	-0.006 (2)	0.006 (2)
C9	0.057 (5)	0.058 (4)	0.038 (3)	0.015 (3)	-0.019 (3)	-0.021 (3)
C10	0.038 (3)	0.059 (4)	0.052 (3)	0.005 (3)	-0.015 (3)	-0.016 (3)
C11	0.030 (3)	0.043 (3)	0.051 (3)	-0.006 (2)	0.001 (2)	-0.009 (2)
C12	0.037 (3)	0.034 (3)	0.052 (3)	0.000 (2)	-0.006 (2)	-0.012 (2)
C13	0.057 (4)	0.036 (3)	0.056 (3)	-0.002 (3)	0.002 (3)	-0.001 (3)
C14	0.033 (3)	0.056 (3)	0.055 (3)	-0.005 (3)	0.011 (3)	-0.012 (3)
C15	0.053 (3)	0.044 (3)	0.041 (3)	0.004 (3)	0.019 (3)	-0.015 (2)
C16	0.069 (4)	0.051 (4)	0.047 (3)	-0.009 (3)	0.009 (3)	0.011 (3)
C17	0.043 (3)	0.040 (3)	0.034 (3)	0.005 (2)	0.004 (2)	-0.021 (2)
C18	0.051 (3)	0.039 (3)	0.058 (4)	-0.001 (3)	-0.018 (3)	-0.019 (3)
C19	0.043 (3)	0.059 (4)	0.054 (4)	0.015 (3)	-0.011 (3)	-0.019 (3)
C20	0.045 (3)	0.044 (3)	0.063 (4)	-0.003 (3)	-0.013 (3)	-0.022 (3)
C21	0.043 (3)	0.057 (4)	0.054 (3)	-0.011 (3)	-0.011 (3)	-0.014 (3)
C22	0.056 (4)	0.037 (3)	0.065 (4)	-0.015 (3)	-0.009 (3)	0.005 (3)
C23	0.037 (3)	0.035 (3)	0.047 (3)	-0.008 (2)	-0.009 (2)	-0.008 (2)
C24	0.057 (3)	0.039 (3)	0.044 (3)	0.001 (2)	-0.021 (3)	0.000 (3)
C25	0.067 (4)	0.045 (3)	0.035 (3)	0.005 (3)	0.001 (3)	0.024 (2)
C26	0.069 (4)	0.037 (3)	0.053 (3)	0.022 (3)	-0.009 (3)	0.003 (3)
C27	0.041 (3)	0.031 (3)	0.048 (3)	0.013 (2)	-0.012 (2)	-0.016 (2)
C28	0.051 (3)	0.046 (3)	0.025 (2)	0.000 (3)	-0.012 (2)	-0.004 (2)
C29	0.053 (3)	0.040 (3)	0.044 (3)	0.015 (3)	0.009 (3)	0.013 (3)
C30	0.055 (4)	0.045 (3)	0.039 (3)	0.004 (3)	0.001 (3)	0.003 (2)
C31	0.049 (3)	0.046 (3)	0.058 (4)	0.025 (3)	0.019 (3)	0.017 (3)
C32	0.048 (4)	0.047 (3)	0.051 (3)	0.016 (3)	0.013 (3)	0.026 (3)
C33	0.056 (4)	0.041 (3)	0.054 (3)	0.015 (2)	0.014 (3)	0.006 (3)
C34	0.055 (3)	0.034 (3)	0.044 (3)	0.007 (2)	0.020 (3)	0.006 (2)
C35	0.064 (4)	0.057 (4)	0.046 (3)	-0.013 (3)	0.023 (3)	-0.020 (3)
Cl1	0.0449 (7)	0.0410 (7)	0.0376 (6)	-0.0005 (6)	-0.0036 (6)	-0.0007 (5)
Cl2	0.0492 (8)	0.0461 (8)	0.0470 (7)	-0.0014 (6)	-0.0080 (6)	0.0143 (6)
N1	0.047 (3)	0.043 (3)	0.0254 (19)	-0.003 (2)	-0.0019 (19)	-0.0066 (18)
N2	0.047 (3)	0.048 (3)	0.035 (2)	-0.006 (2)	-0.007 (2)	-0.006 (2)

O1	0.045 (2)	0.0283 (18)	0.043 (2)	-0.0104 (15)	-0.0085 (17)	0.0118 (16)
O2	0.059 (3)	0.048 (3)	0.069 (3)	0.022 (2)	-0.015 (2)	-0.021 (2)
P1	0.0446 (8)	0.0387 (7)	0.0369 (7)	0.0004 (6)	-0.0016 (6)	-0.0002 (6)
Ru1	0.0366 (2)	0.03698 (19)	0.03518 (19)	-0.0014 (2)	-0.0037 (2)	0.00081 (16)

*Geometric parameters (Å, °)*

C1—N1	1.356 (8)	C19—H19	0.9300
C1—C2	1.362 (9)	C20—C21	1.337 (9)
C1—H1	0.9300	C20—H20	0.9300
C2—C3	1.332 (8)	C21—C22	1.398 (8)
C2—H2	0.9300	C21—H21	0.9300
C3—C4	1.374 (8)	C22—H22	0.9300
C3—H3	0.9300	C23—C24	1.373 (8)
C4—C5	1.393 (8)	C23—C28	1.385 (8)
C4—H4	0.9300	C23—P1	1.855 (6)
C5—N1	1.369 (7)	C24—C25	1.397 (8)
C5—C6	1.524 (8)	C24—H24	0.9300
C6—O1	1.466 (7)	C25—C26	1.363 (9)
C6—H6A	0.9700	C25—H25	0.9300
C6—H6B	0.9700	C26—C27	1.370 (8)
C7—C8	1.390 (8)	C26—H26	0.9300
C7—C12	1.390 (8)	C27—C28	1.398 (7)
C7—O1	1.391 (7)	C27—H27	0.9300
C8—C9	1.390 (9)	C28—H28	0.9300
C8—H8	0.9300	C29—C30	1.354 (8)
C9—C10	1.390 (9)	C29—C34	1.407 (9)
C9—H9	0.9300	C29—P1	1.837 (6)
C10—C11	1.390 (8)	C30—C31	1.369 (8)
C10—H10	0.9300	C30—H30	0.9300
C11—C12	1.390 (8)	C31—C32	1.404 (9)
C11—C15	1.390 (8)	C31—H31	0.9300
C12—N2	1.390 (7)	C32—C33	1.372 (8)
C13—N2	1.390 (8)	C32—H32	0.9300
C13—C14	1.390 (9)	C33—C34	1.382 (8)
C13—C16	1.470 (9)	C33—H33	0.9300
C14—C15	1.390 (9)	C34—H34	0.9300
C14—H14	0.9300	C35—O2	1.381 (7)
C15—H15	0.9300	C35—H35A	0.9600
C16—H16A	0.9600	C35—H35B	0.9600
C16—H16B	0.9600	C35—H35C	0.9600
C16—H16C	0.9600	Cl1—Ru1	2.4905 (14)
C17—C18	1.371 (8)	Cl2—Ru1	2.4104 (14)
C17—C22	1.392 (8)	N1—Ru1	2.067 (5)
C17—P1	1.872 (6)	N2—Ru1	2.162 (5)
C18—C19	1.406 (8)	O1—Ru1	2.060 (3)
C18—H18	0.9300	O2—H2A	0.8200
C19—C20	1.325 (9)	P1—Ru1	2.2931 (15)

N1—C1—C2	121.7 (7)	C24—C23—P1	118.2 (4)
N1—C1—H1	119.2	C28—C23—P1	123.1 (4)
C2—C1—H1	119.2	C23—C24—C25	121.6 (6)
C3—C2—C1	122.4 (7)	C23—C24—H24	119.2
C3—C2—H2	118.8	C25—C24—H24	119.2
C1—C2—H2	118.8	C26—C25—C24	119.8 (5)
C2—C3—C4	119.3 (6)	C26—C25—H25	120.1
C2—C3—H3	120.4	C24—C25—H25	120.1
C4—C3—H3	120.4	C25—C26—C27	119.5 (5)
C3—C4—C5	117.0 (5)	C25—C26—H26	120.2
C3—C4—H4	121.5	C27—C26—H26	120.2
C5—C4—H4	121.5	C26—C27—C28	120.8 (5)
N1—C5—C4	124.3 (5)	C26—C27—H27	119.6
N1—C5—C6	115.8 (5)	C28—C27—H27	119.6
C4—C5—C6	119.9 (5)	C23—C28—C27	120.2 (5)
O1—C6—C5	107.0 (4)	C23—C28—H28	119.9
O1—C6—H6A	110.3	C27—C28—H28	119.9
C5—C6—H6A	110.3	C30—C29—C34	117.7 (6)
O1—C6—H6B	110.3	C30—C29—P1	125.5 (5)
C5—C6—H6B	110.3	C34—C29—P1	116.1 (4)
H6A—C6—H6B	108.6	C29—C30—C31	120.8 (6)
C8—C7—C12	120.0 (5)	C29—C30—H30	119.6
C8—C7—O1	123.2 (5)	C31—C30—H30	119.6
C12—C7—O1	116.2 (5)	C30—C31—C32	120.9 (6)
C9—C8—C7	120.0 (5)	C30—C31—H31	119.5
C9—C8—H8	120.0	C32—C31—H31	119.5
C7—C8—H8	120.0	C33—C32—C31	119.7 (6)
C8—C9—C10	120.0 (5)	C33—C32—H32	120.1
C8—C9—H9	120.0	C31—C32—H32	120.1
C10—C9—H9	120.0	C32—C33—C34	117.7 (6)
C11—C10—C9	120.0 (5)	C32—C33—H33	121.2
C11—C10—H10	120.0	C34—C33—H33	121.2
C9—C10—H10	120.0	C33—C34—C29	122.8 (6)
C12—C11—C10	120.0 (5)	C33—C34—H34	118.6
C12—C11—C15	120.0 (6)	C29—C34—H34	118.6
C10—C11—C15	120.0 (5)	O2—C35—H35A	109.5
C11—C12—N2	120.0 (5)	O2—C35—H35B	109.5
C11—C12—C7	120.0 (5)	H35A—C35—H35B	109.5
N2—C12—C7	120.0 (5)	O2—C35—H35C	109.5
N2—C13—C14	120.0 (6)	H35A—C35—H35C	109.5
N2—C13—C16	119.6 (6)	H35B—C35—H35C	109.5
C14—C13—C16	120.1 (6)	C1—N1—C5	115.3 (5)
C13—C14—C15	120.0 (6)	C1—N1—Ru1	130.1 (4)
C13—C14—H14	120.0	C5—N1—Ru1	114.5 (4)
C15—C14—H14	120.0	C13—N2—C12	120.0 (5)
C14—C15—C11	120.0 (6)	C13—N2—Ru1	130.9 (4)
C14—C15—H15	120.0	C12—N2—Ru1	109.1 (4)



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C11—C15—H15	120.0	C7—O1—C6	118.9 (4)
C13—C16—H16A	109.5	C7—O1—Ru1	113.9 (3)
C13—C16—H16B	109.5	C6—O1—Ru1	111.9 (3)
H16A—C16—H16B	109.5	C35—O2—H2A	109.5
C13—C16—H16C	109.5	C29—P1—C23	101.2 (3)
H16A—C16—H16C	109.5	C29—P1—C17	103.7 (3)
H16B—C16—H16C	109.5	C23—P1—C17	101.1 (2)
C18—C17—C22	119.7 (5)	C29—P1—Ru1	113.4 (2)
C18—C17—P1	121.1 (4)	C23—P1—Ru1	117.07 (18)
C22—C17—P1	119.0 (5)	C17—P1—Ru1	118.10 (17)
C17—C18—C19	120.2 (5)	O1—Ru1—N1	79.47 (17)
C17—C18—H18	119.9	O1—Ru1—N2	79.66 (16)
C19—C18—H18	119.9	N1—Ru1—N2	158.25 (17)
C20—C19—C18	118.5 (6)	O1—Ru1—P1	94.55 (11)
C20—C19—H19	120.8	N1—Ru1—P1	92.46 (12)
C18—C19—H19	120.8	N2—Ru1—P1	95.22 (13)
C19—C20—C21	123.3 (6)	O1—Ru1—Cl2	169.80 (11)
C19—C20—H20	118.4	N1—Ru1—Cl2	92.40 (14)
C21—C20—H20	118.4	N2—Ru1—Cl2	107.61 (13)
C20—C21—C22	120.2 (6)	P1—Ru1—Cl2	91.93 (6)
C20—C21—H21	119.9	O1—Ru1—Cl1	85.32 (11)
C22—C21—H21	119.9	N1—Ru1—Cl1	87.29 (12)
C17—C22—C21	118.1 (6)	N2—Ru1—Cl1	84.98 (13)
C17—C22—H22	121.0	P1—Ru1—Cl1	179.74 (6)
C21—C22—H22	121.0	Cl2—Ru1—Cl1	88.17 (5)
C24—C23—C28	117.9 (5)		

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