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**(Acetonitrile-*N*)(*o*-benzoquinone
diimine-*N,N'*)chloro-*trans*-bis(triphenylphosphine-*P*)ruthenium(II)
hexafluorophosphate 0.25-hydrate**

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(Acetonitrile-*N*)(*o*-benzoquinone diimine-*N,N'*)chloro-*trans*-bis(triphenylphosphine-*P*)ruthenium(II) hexafluorophosphate 0.25-hydrate

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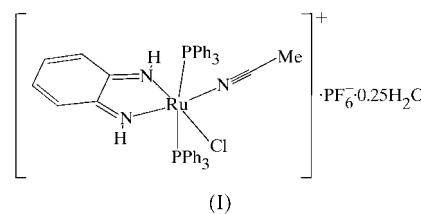
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The Ru atom in the title compound, $[\text{RuCl}(\text{CH}_3\text{CN})-\{\text{P}(\text{C}_6\text{H}_5)_3\}_2\{\text{C}_6\text{H}_4(\text{NH})_2\}]\text{PF}_6 \cdot 0.25\text{H}_2\text{O}$, has a six-coordinate octahedral geometry, with a *trans* arrangement of the triphenylphosphine groups. The asymmetric unit contains two complex molecules and a partially occupied water site. Principal dimensions include Ru—N 1.958 (4)—2.044 (5), Ru—P 2.3897 (16)—2.4092 (15), and Ru—Cl 2.4280 (15) and 2.4295 (16) Å.

Comment

In order to understand the mixing between ruthenium and redox-active (non-innocent) ligands and its implication on the electrochemical parameter (Lever, 1990; Masui *et al.*, 2000; Venegas-Yazigi *et al.*, 2000, and references therein), we synthesized a family of ruthenium complexes containing *o*-benzoquinone diimine (BQDI), two triphenylphosphine ligands in a *trans* arrangement and two other non-redox-active ligands such as chlorine, acetonitrile, diethyldithiocarbamate *etc*. The unit cell of the title compound, (I), contains two molecules of complex, two hexafluorophosphate anions and a half molecule of water. The Ru atom shows a six-coordinate octahedral geometry, with a *trans* arrangement of the triphenylphosphine groups. C—C distances in the quinone ligand have been very important in confirming the quinoid form of BQDI.



Experimental

Crystal data

$[\text{RuCl}(\text{C}_2\text{H}_3\text{N})(\text{C}_6\text{H}_6\text{N}_2)-(\text{C}_18\text{H}_{15}\text{P})_2]\text{PF}_6 \cdot 0.25\text{H}_2\text{O}$

$M_r = 957.72$

Monoclinic, $P2_1/c$

$a = 24.2211 (8)$ Å

$b = 10.6752 (4)$ Å

$c = 35.1913 (12)$ Å

$\beta = 107.747 (2)$ °

$V = 8666.2 (5)$ Å³

$Z = 8$

$D_x = 1.468$ Mg m⁻³

Mo $K\alpha$ radiation

Cell parameters from 36022 reflections

$\theta = 3.41\text{--}25.02^\circ$

$\mu = 0.596$ mm⁻¹

$T = 150$ (2) K

Very fine needle, purple
0.17 × 0.09 × 0.04 mm

Data collection

Nonius KappaCCD diffractometer

φ and ω scans with κ offsets

Absorption correction: multi-scan (*DENZOSMN*; Otwinowski & Minor, 1997)

$T_{\min} = 0.906$, $T_{\max} = 0.977$

36 022 measured reflections

15 108 independent reflections

6785 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.108$

$\theta_{\max} = 25.02^\circ$

$h = -29 \rightarrow 28$

$k = -13 \rightarrow 12$

$l = -41 \rightarrow 39$

Refinement

Refinement on F^2

$R(F) = 0.056$

$wR(F^2) = 0.118$

$S = 0.875$

15 108 reflections

1056 parameters

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0356P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.012$

$\Delta\rho_{\max} = 0.62$ e Å⁻³

$\Delta\rho_{\min} = -0.43$ e Å⁻³

Table 1
Selected geometric parameters (Å, °).

Ru1A—N3A	1.971 (4)	Ru1B—N1B	2.029 (5)
Ru1A—N2A	2.015 (4)	Ru1B—N2B	2.030 (4)
Ru1A—N1A	2.044 (5)	Ru1B—P2B	2.3897 (16)
Ru1A—P2A	2.3937 (15)	Ru1B—P1B	2.3899 (15)
Ru1A—P1A	2.4092 (15)	Ru1B—Cl1B	2.4295 (16)
Ru1B—N3B	1.958 (4)		
N3A—Ru1A—N2A	77.08 (18)	N3B—Ru1B—N1B	91.64 (19)
N3A—Ru1A—N1A	92.10 (18)	N3B—Ru1B—N2B	76.9 (2)
N2A—Ru1A—N1A	169.18 (18)	N1B—Ru1B—N2B	168.5 (2)
N3A—Ru1A—P2A	91.65 (12)	N3B—Ru1B—P2B	92.39 (12)
N2A—Ru1A—P2A	92.84 (12)	N1B—Ru1B—P2B	88.95 (14)
N1A—Ru1A—P2A	87.56 (12)	N2B—Ru1B—P2B	91.62 (12)
N3A—Ru1A—P1A	93.09 (12)	N3B—Ru1B—P1B	91.61 (12)
N2A—Ru1A—P1A	92.36 (12)	N1B—Ru1B—P1B	89.03 (13)
N1A—Ru1A—P1A	88.02 (13)	N2B—Ru1B—P1B	91.15 (12)
P2A—Ru1A—P1A	173.64 (6)	P2B—Ru1B—P1B	175.57 (6)

All H atoms were visible from difference Fourier maps (except for those bonded to the partial-occupancy water molecule) and were included in calculated positions and refined as riding atoms with C—H = 0.95–0.98 Å and N—H = 0.88 Å. The H atoms of the partial-occupancy water molecule were not included in the calculations, but are included in the molecular formula.

Data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SHELXTL/PC* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL/PC*; software used to prepare material for publication: *SHELXTL/PC*.

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