metal-organic papers

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Key indicators

Single-crystal X-ray study T = 113 K Mean σ (C–C) = 0.006 Å R factor = 0.044 wR factor = 0.119 Data-to-parameter ratio = 17.2

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

{N-[Bis(2-pyridyl)methyl]-N,N-bis(2-pyridylmethyl)amine- $\kappa^5 N$ }chlororuthenium(II) perchlorate methanol solvate

The structure of the title compound, $[RuCl(C_{23}H_{21}N_5)]$ -ClO₄·CH₄O, has been determined at 113 K, revealing a mononuclear six-coordinate structure with a 0.263 (2) Å displacement of the Ru^{II} atom from the N₄ equatorial plane. Received 30 November 2004 Accepted 6 December 2004 Online 8 January 2005

Comment

Ruthenium complexes with mulidentate pyridylamine derivatives as ligands have been shown to perform catalytic hydrocarbon oxygenation with use of various terminal oxidants. Among these, we and other groups have reported catalytic alkane oxygenation by ruthenium complexes with tris(2-pyridylmethyl)amine (TPA) and its derivatives as ligands (Kojima, 1996; Yamaguchi et al., 1997; Kojima & Matsuda, 1999; Kojima, Matsuo & Matsuda, 2000; Jitsukawa et al., 2001; Yamaguchi et al., 2004). Recently, Yamaguchi and coworkers have reported on those reactions of a Ru^{II} complex N-[tris(2-pyridyl)methyl]-N,N-tris(2-pyridyl)methylwith amine (N₄py) (Lubben et al., 1995; Roelfes et al., 1997) in CHCl₃ with *m*-chloroperbenzoic acid (Yamaguchi et al., 2002). However, no crystal structure has been reported of the catalyst, $[RuCl(N_4py)]^+$. In order to set up a baseline to consider the reactivity and properties of the title complex, (I), we determined the crystal structure of its perchlorate salt.



Compound (I) is depicted in Fig. 1, showing the atomic numbering scheme, and selected bond lengths and angles are given in Table 1. The Ru^{II} center in (I) exhibits a distorted octahedral geometry, with a displacement of 0.263 (2) Å toward the chloro ligand from the equatorial least-squares plane consisting of atoms N2–N5. This value is smaller than that in [ZnCl(N₄py)]ClO₄·H₂O [0.502 (2) Å; Kojima, Weber & Choma, 2004]. The bond lengths around the Ru^{II} center are comparable to those observed for other Ru^{II}–TPA complexes reported so far (Kojima *et al.*, 1998, 2003; Kojima, Hayashi & Matsuda, 2000, 2004; Kojima & Matsuda, 2001; Kojima, Sakamoto & Matsuda, 2004; Kojima, Miyazaki *et al.*, 2004).

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Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

The dihedral angles of the planes of the pyridine rings relative to the equatorial mean plane were estimated to be 96.0 $(1)^{\circ}$ for the pyridine ring containing N2, 83.0 $(1)^{\circ}$ for that with N3, 110.3 (1)° for that with N4, and 64.4° for that with N5. The pyridine ring containing N5, which is connected to a methine C atom of N₄py, undergoes severe distortion from an ideal orientation (ca 90°) toward the equatorial plane. In the crystal structure, an intermolecular hydrogen bond was observed between the chloro ligand and the OH group of the methanol molecule [3.158 (3) Å; Table 2].

Experimental

[RuCl(N₄py)]ClO₄·CH₃OH refluxing was prepared by $[RuCl_2(DMSO)_4]$ (90 mg, 1.9 × 10⁻⁴ mol) with N₄py·HClO₄ (87 mg, 1.9 X 10⁻⁴ mol) in methanol (10 ml) for 24 h. After filtration, the solvent was removed by a rotary evaporator. The crude product was recrystallized from methanol with vapor diffusion of diethyl ether to give red crystals of the title compound.

Crystal data

[RuCl(C ₂₃ H ₂₁ N ₅)]ClO ₄ ·CH ₄ O	$D_x = 1.650 \text{ Mg m}^{-3}$
$M_r = 635.47$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 31 496
a = 14.768 (2) Å	reflections
b = 13.048 (2) Å	$\theta = 2.2 - 27.5^{\circ}$
c = 15.084 (2) Å	$\mu = 0.87 \text{ mm}^{-1}$
$\beta = 118.349 \ (3)^{\circ}$	T = 113.2 K
V = 2558.0 (5) Å ³	Block, red
Z = 4	$0.48 \times 0.45 \times 0.20 \ \mathrm{mm}$
Data collection	
Rigaku R-AXIS RAPID imaging-	5745 independent reflections
0 0	
plate diffractometer	4780 reflections with $F^2 > 2\sigma(F^2)$
plate diffractometer ω scans	4780 reflections with $F^2 > 2\sigma(F^2)$ $R_{\text{int}} = 0.042$
plate diffractometer ω scans Absorption correction: multi-scan	4780 reflections with $F^2 > 2\sigma(F^2)$ $R_{\text{int}} = 0.042$ $\theta_{\text{max}} = 27.5^{\circ}$
plate diffractometer ω scans Absorption correction: multi-scan (<i>ABSCOR</i> ; Higashi, 1995)	4780 reflections with $F^2 > 2\sigma(F^2)$ $R_{\text{int}} = 0.042$ $\theta_{\text{max}} = 27.5^{\circ}$ $h = -19 \rightarrow 19$
plate diffractometer ω scans Absorption correction: multi-scan (<i>ABSCOR</i> ; Higashi, 1995) $T_{min} = 0.444, T_{max} = 0.841$	4780 reflections with $F^2 > 2\sigma(F^2)$ $R_{int} = 0.042$ $\theta_{max} = 27.5^{\circ}$ $h = -19 \rightarrow 19$ $k = -16 \rightarrow 16$

Ret	finement
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Refinement on F^2	H-atom parameters not refined
$R[F^2 > 2\sigma(F^2)] = 0.044$	$w = 1/{\sigma^2(F_o^2)} + [0.06(\max(F_o^2, 0))]$
$wR(F^2) = 0.120$	$+2F_{c}^{2}/3]^{2}$
S = 1.32	$(\Delta/\sigma)_{\rm max} < 0.001$
5743 reflections	$\Delta \rho_{\rm max} = 1.18 \text{ e } \text{\AA}^{-3}$
334 parameters	$\Delta \rho_{\rm min} = -1.24 \text{ e} \text{ Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Ru1-Cl1	2.4162 (8)	Ru1-N3	2.062 (3)
Ru1-N1	2.036 (3)	Ru1-N4	2.059 (3)
Ru1-N2	2.060 (3)	Ru1-N5	2.047 (3)
Cl1-Ru1-N1	179.28 (8)	N1-Ru1-N5	82.0 (1)
Cl1-Ru1-N2	97.27 (8)	N2-Ru1-N3	85.0 (1)
Cl1-Ru1-N3	96.94 (8)	N2-Ru1-N4	165.6 (1)
Cl1-Ru1-N4	97.16 (8)	N2-Ru1-N5	93.3 (1)
Cl1-Ru1-N5	97.85 (7)	N3-Ru1-N4	93.7 (1)
N1-Ru1-N2	83.4 (1)	N3-Ru1-N5	165.2 (1)
N1-Ru1-N3	83.2 (1)	N4-Ru1-N5	84.25 (10)
N1-Ru1-N4	82.1 (1)		

Table 2 Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O5-H25\cdots Cl1^i$	0.93	2.23	3.158 (3)	174
	1	. 3		

Symmetry code: (i) $-x + 1, +y + \frac{1}{2}, -z + \frac{3}{2}$.

All H atoms were placed at ideal positions (C-H = 0.95 Å), except H25, which was located in a difference Fourier map, and were refined as riding $[U_{iso} = 1.2U_{eq}(C)]$. The maximum residual peak in the final Fourier map was found in the vicinity of the Ru ion.

Data collection: PROCESS-AUTO (Rigaku Corporation, 1998); cell refinement: PROCESS-AUTO; data reduction: TEXSAN Version 1.11 (Molecular Structure Corporation & Rigaku Corporation, 2000); program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: TEXSAN Version 1.10 (Molecular Structure Corporation, 1985–1999); molecular graphics: TEXSAN Version 1.11; software used to prepare material for publication: TEXSAN Version 1.11.

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