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Key indicators

Single-crystal X-ray study T = 110 KMean σ (C–C) = 0.007 Å H-atom completeness 90% Disorder in solvent or counterion R factor = 0.027 wR factor = 0.076 Data-to-parameter ratio = 17.5

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

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μ -Hydroxo-bis[(4,4'-dimethyl-2,2'-bipyridine)tricarbonylrhenium(I)] perrhenate acetone hemisolvate

The title compound, $[\text{Re}_2(\text{OH})(\text{C}_{12}\text{H}_{12}\text{N}_2)_2(\text{CO})_6][\text{ReO}_4]$ -0.5C₃H₆O, is the first crystal structure reported for a hydroxo-bridged binuclear rhenium complex containing bipyridine or substituted bipyridine ligands. The cation has distorted octahedral coordination about both rhenium centers, resulting from geometric constraints imposed by the chelating 4,4'dimethyl-2,2'-bipyridine ligands, which are involved in an intermolecular π - π stacking interaction. The distance between the rhenium atoms is 4.0185 (3) Å, with no metalmetal bond. The H atom of the bridging hydroxo ligand was directly located in the crystallographic analysis, and is hydrogen-bonded to the perrhenate. The acetone solvent molecule is disordered.

Comment

The title compound has been prepared as part of a general study of rhenium complexes bearing polypyridine ligands, which may be intermediates in photocatalytic reductions of CO₂ (Gibson & Yin, 1998). Surprisingly, only been four crystal structures have been reported for hydroxo-bridged rhenium complexes containing nitrogen donor atoms, and none of these contain the bipyridyl ligand or its substituted derivatives. Böhm et al. (1991) observed a metal-metal bond [2.528 (1) Å] in the doubly-bridged $[(tacn)_2 Re_2 Cl_2(\mu-Cl)(\mu-OH)]$ (tacn = 1,4,7-triazacyclononane) cation, but were unable to locate the H atom of the hydroxy group; thus the assignment of the OH⁻ rather than an O^{2-} bridge was based solely on charge. In a study involving the reactivity of rhenium with pyrazolato ligands, Ardizzoia et al. (1998) prepared (Et₃NH)[Re₂(CO)₆- $(pz)_2OH]^-$ by hydrolysis of a bromide-bridged species. The crystal structure reported for the bromide complex shows two additional bridging pyrazoles, with an Re...Re separation of 3.825 (1) Å. Unfortunately, the structure determination of the hydroxo-bridged compound was an ancillary study for which only unit-cell parameters are available. A third structurally characterized binuclear rhenium complex with N donor atoms is $\operatorname{Re}_2(\operatorname{CO})_3(\operatorname{NO})_2(\operatorname{SMe})(\mu$ -'BuS)(μ -OH) (Nefedov *et al.*, 1998), which has two nitrosyl groups trans to the hydroxo bridge and two bulky tert-butylthiolato groups completing the triply-bridged core. Cotton et al. (1996) reported the crystal structure of the tetranuclear compound [Re₄(C₆H₅NCO- $CH_3)_6(Cl)(\mu-O)(\mu-OH)(MeOH)_3][ReO_4]_2$, formed by bridging two dirhenium units with one oxo and one hydroxo ligand.

Compound (I) crystallizes with one hydroxo-bridged $[\text{Re}_2(\text{CO})_6(4,4'-\text{dmbpy})_2\text{OH}]^+$ cation (dmbpy = dimethyl-2,2'bipyridine), one perrhenate anion and one-half disordered acetone solvent molecule per asymmetric unit; an *ORTEP*-3 diagram (Farrugia, 1997) of (I) illustrates the structures of the Received 12 August 2003 Accepted 10 September 2003 Online 18 September 2003

metal-organic papers

independent entities in Fig. 1. Each rhenium atom displays distorted octahedral geometry, consistent with rhenium dimethylpyridine compounds that do not contain a hydroxo bridge (Gibson *et al.*, 2003).



The rhenium atoms are bridged by a single hydroxo ligand, unsupported by a metal-metal bond, as indicated by the large Re...Re separation [4.0185 (3) Å]. In comparison, the multiply bridged complexes described above show strong Re-Re bonds ranging from 2.213 (2) Å to 2.528 (1) for the tetranuclear acetanilide and triazacyclonane complexes, respectively. The position of the H atom of the hydoxo bridge in (I) was determined from an electron density maps and its presence is corroborated by two long Re-O bonds [2.127 (3) and 2.155 (3) Å]. In [Re₄(C₆H₅NCOCH₃)₆(Cl)(μ -O)(μ -OH)(MeOH)₃], both Re-O bonds in the oxo bridge are strong [1.88 (2) and 1.94 (2) Å], while its hydroxo bridge has one moderately strong 2.06 (3) and one weak 2.29 (2) Å Re-O bond.

The packing arrangement about the *ac* crystal face for (I) (Fig. 2) shows that the dirhenium cations pack in a threedimensional network arising from intermolecular π - π -stacking between dimethylbipyridine ligands of adjacent cations. One six-membered ring (N2 and C12–C16) of the first dmpy stacks together with a second six-membered ring (N2' and C12'–C16') of a neighboring ligand, forming a π - π interaction (3.959 Å centroid to centroid). Also seen in this illustration is the hydrogen-bonding interaction between the H



Figure 1

ORTEP-3 (Farrugia, 1997) plot, showing 40% displacement ellipsoids. Aromatic and methyl H atoms are omitted for clarity. The hydoxyl H atom is shown as a small sphere of arbitrary radius.



Figure 2

Packing diagram for (I), displaying intermolecular π stacking interactions between neighboring dimethylpyridyl rings, the hydrogen-bonding interaction between the hydroxo-bridged H atom and the perrhenate anion, and disorder of the solvent molecule about an inversion center.

atom of the hydroxo bridge and the perrhenate anion. The O7…O11ⁱ [2.761 (1) Å] and H7o…O2ⁱ [2.015 (1) Å] separations, as well as the O7—H7o…O11ⁱ [169.4 (2)°] angle [symmetry code: (i) x, y, z - 1] indicate strong hydrogen bonding.

Experimental

A mixture of *fac*-Re(dmbpy)(CO)₃OTf (Gibson & Yin, 1998) and Na₂CO₃ (1:2) in acetone was stirred for 1 d at 295 K. The reaction mixture was filtered, the filtrate was concentrated and a solution of NH₄ReO₄ in water was added. After stirring for 19 h, the crude product precipitated from the solution. Repeated recrystallizations from acetone/ether at 273 K gave the pure compound *fac*,*fac*-Re(dmbpy)(CO)₃(μ -OH)Re(dmbpy)(CO)₃ReO₄ as plate-shaped crystals.

Crystal data

 $T_{\min} = 0.180, \ T_{\max} = 0.900$

29838 measured reflections

[Re2(OH)(C12H12N2)2(CO)6]- $D_x = 2.320 \text{ Mg m}^{-3}$ [ReO₄] 0.5C₃H₆O Mo $K\alpha$ radiation $M_r = 1205.16$ Cell parameters from 8716 Monoclinic, $P2_1/c$ reflections a = 10.6841 (7) Å $\theta = 2.4 - 28.1^{\circ}$ b = 31.649 (2) Å $\mu = 10.56 \text{ mm}^{-1}$ c = 10.5758 (7) Å T = 110 (2) K $\beta = 105.222 (1)^{\circ}$ Rectangular plate, yellow $V = 3450.7 (4) \text{ Å}^3$ $0.29 \times 0.14 \times 0.01 \text{ mm}$ Z = 4Data collection Bruker SMART APEX CCD 7989 independent reflections diffractometer 7184 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.042$ ω scans Absorption correction: multi-scan $\theta_{\rm max} = 28.3^{\circ}$ (SADABS; Sheldrick, 2001) $h=-13\rightarrow14$

 $k = -41 \rightarrow 42$

 $l = -14 \rightarrow 14$

$w = 1/[\sigma^2(F_o^2) + (0.039P)^2]$
+ 1.9625P]
where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} = 0.003$
$\Delta \rho_{\rm max} = 3.70 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -1.94 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Re1-C1	1.936 (4)	Re1-N2	2.186 (4)
Re1-C2	1.905 (5)	Re1-O7	2.155 (3)
Re1-C3	1.920 (5)	Re2-O7	2.127 (3)
Re1-N1	2.169 (4)		
C2-Re1-C3	89.35 (19)	C1-Re1-N1	171.52 (16)
C2-Re1-C1	89.70 (19)	O7-Re1-N1	83.08 (12)
C3-Re1-C1	88.81 (18)	C2-Re1-N2	171.33 (16)
C2-Re1-O7	94.05 (16)	C3-Re1-N2	96.26 (16)
C3-Re1-O7	175.33 (15)	C1-Re1-N2	97.00 (16)
C1-Re1-O7	94.41 (15)	O7-Re1-N2	80.00 (12)
C2-Re1-N1	98.54 (16)	N1-Re1-N2	74.60 (13)
C3-Re1-N1	93.26 (16)	Re2-O7-Re1	139.63 (17)

Half-occupancy atoms C60–C62 and O60 in the disordered acetone solvent were refined isotropically. A second 50% acetone is generated by the symmetry operator -x, -y, 2 - z, resulting in one full-occupancy disordered solvent located at each corner of the unit cell. Positions for H atoms in the solvent could not be accurately determined and were not included in the model. Aromatic H-atom positions were calculated, and included as fixed contributions, with $U_{iso}(H) = 1.2U_{eq}(C)$. Methyl H atoms of the cation were placed in calculated positions and allowed to ride (the torsion angle which defines its orientation was allowed to refine) on the attached C atom,

and these atoms were assigned $U_{\rm iso}({\rm H}) = 1.5 U_{\rm eq}({\rm C})$. The H atom of the hydroxo bridge was located in a difference map and refined isotropically. The highest peak and deepest hole are located at 0.77 and 0.66 Å from Re3, respectively.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*86 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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