metal-organic compounds

Acta Crystallographica Section C Crystal Structure Communications ISSN 0108-2701

Sulfido-bridged rhenium-carbonyl complexes with planar and folded Re₂S₂ cores

Ulrich Flörke,* Hans Egold and Detlev Schwarze

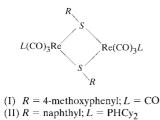
Fachbereich Chemie und Chemietechnik, Universität-GH Paderborn, Warburgerstr. 100, D-33098 Paderborn, Germany Correspondence e-mail: uf@chemie.uni-paderborn.de

Received 11 November 1999 Accepted 23 November 1999

The two sulfido-bridged dirhenium complexes bis(μ -4-methoxyphenylsulfido-S)bis(tetracarbonylrhenium), [Re₂- $(C_7H_7OS)_2(CO)_8$], and bis(μ -naphthylsulfido-S)bis[tricarbonyl(dicyclohexylphosphane)rhenium], [Re₂($C_{12}H_{23}P$)₂- $(C_{10}H_7S)_2(CO)_6$], show different geometries of the common Re₂S₂ core. The 4-methoxyphenyl derivative has crystallographic $\overline{1}$ symmetry and the naphthyl derivative has C^2 symmetry. This results from intramolecular repulsion due to different substitution patterns at the Re and S atoms.

Comment

We reported recently new synthetic routes to hydrido sulfidobridged dirhenium complexes (Egold *et al.*, 1999). In this context, the new doubly sulfido-bridged complexes $\text{Re}_2(\mu$ - $SR)_2(\text{CO})_8$, (I) (R = 4-methoxyphenyl), and $\text{Re}_2(\mu$ - $SR')_2(\text{CO})_6(\text{PHCy}_2)_2$, (II) (R' = naphthyl; Cy = cyclohexyl), were also prepared. These two compounds differ markedly in their molecular geometries.



In (I) (Fig. 1), each Re atom is octahedrally coordinated by two axially and two equatorially positioned carbonyl groups and two bridging S atoms. The CO groups show an ecliptic arrangement with torsion angles $C4-Re\cdots Re^{i}-C3^{i}$ and $C2-Re\cdots Re^{i}-C1^{i}$ of 0.4 (3) and 1.3 (3)°, respectively [symmetry code: (i) -1 - x, 1 - y, -z). The geometric centre of the molecule lies on a crystallographic inversion centre. Consequently, the Re₂S₂ core is planar and the methoxyphenyl groups attached to the μ -S atoms adopt *anti* positions. The two Re-S bond lengths of 2.524 (1) and 2.518 (1) Å differ only slightly, but they are significantly elongated

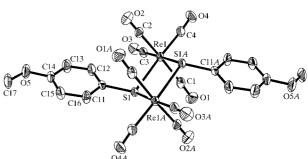


Figure 1

The molecular structure of (I) showing 50% probability displacement ellipsoids. H atoms have been omitted for clarity.

compared with those of 2.471 (3) or 2.478 (3) Å, respectively, found in the hydrido sulfido-bridged complexes $\text{Re}_2(\mu-\text{H})(\mu-SR)(\text{CO})_8$ (R = phenyl or naphthyl; Egold *et al.*, 1999). The related acetonitrile-substituted complex $\text{Re}_2(\mu-\text{S}^t\text{Bu})_2$ -($\text{CO})_6(\text{CH}_3\text{CN})_2$ (Eremenko *et al.*, 1992) exhibits Re-S bond lengths of 2.522 (2) and 2.534 (2) Å. The enclosed ring angles of (I) are Re-S-Re 98.47 (4)° and S-Re-S 81.53 (4)°, and the non-bonding $\text{Re}\cdots\text{Re}$ separation is 3.819 (1) Å. The basic structural elements of (I), the planar $M_2\text{S}_2$ ring and the *anti* position of the *R* residues at the sulfido bridges, are also valid for the above-mentioned rhenium acetonitrile complex, for the complex $\text{Re}_2(\mu-\text{S}R)_2(\text{CO})_8$ [$R = \text{CS}_2\text{Re}(\text{CO})_4$] (Thiele *et al.*, 1974) as well as for the two manganese compounds $\text{Mn}_2(\mu-\text{S}-\text{phenyl})_2(\text{CO})_8$ (Chen *et al.*, 1996).

In contrast to this, the core structure of (II) (Fig. 2) is entirely different showing a folded Re₂S₂ ring with an Re₂S/-SRe₂ dihedral angle of 137.8 (1)° and naphthyl residues at both bridging S atoms in the *syn* position. The coordination of Re by the two μ -S atoms, three CO groups and one axially attached PHCy₂ ligand is slightly distorted octahedral. These latter terminal ligands show a largely ecliptic arrangement with torsion angles of 6.4 (7)° for C3-Re···Reⁱⁱ-C1ⁱⁱ, 9.2 (1)° for P1-Re···Reⁱⁱ-P1ⁱⁱ and 10.3 (7)° for C2-Re···Reⁱⁱ-C2ⁱⁱ [symmetry code: (ii) -x, y, $\frac{1}{2} - z$). The phos-

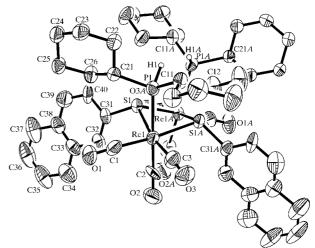


Figure 2

The molecular structure of (II) showing 50% probability displacement ellipsoids. Cyclohexyl- and naphthyl-H atoms have been omitted for clarity.

phido ligands at the Re atoms are in the syn position too, and a crystallographic twofold axis runs through the midpoint of the $Re \cdot \cdot \cdot Re$ vector. This geometry results in the axial PHCy₂ and CO ligands having intramolecular non-bonding distances of $PH \cdots HP$ 2.32 (15) Å [P-H is 1.44 (8) Å] and $O \cdots O$ 2.994 (15) Å. For a corresponding planar molecule like (I), the carbonyl $O \cdots O$ distances are about 3.2 Å, but the calculated PH···HP separation of 0.9 Å and cyclohexyl CH···HC separation of 1.8 Å would clearly be too short and the resulting repulsive forces give rise to the observed ring distortion. Similar ring folding in order to reduce intramolecular repulsion is also known for μ -P-bridged complexes (Flörke & Haupt, 1994). The Re–S bond lengths of 2.523 (2) and 2.526 (2) Å compare well with those of (I), but the Re-S-Re and S-Re-S angles of 98.93 (8) and 74.64 (8)°, respectively, reflect the folding of the ring. The non-bonding Re···Re distance is 3.838 (1) Å.

Experimental

For the preparation of (I), $\text{Re}_2(\text{CO})_{10}$ (200 mg, 0.307 mmol) and 4methoxy-thiophenol (75 µl, 0.612 mmol) were heated at 443 K in a sealed glass tube in the presence of xylene (1.5 ml) for 10 h. Subsequent thin-layer chromatography (TLC) separation (eluant: dichloromethane/*n*-hexane 1/5) gave (I) in 10% yield. Single crystals were grown from chloroform/*n*-pentane. For the preparation of (II), trimethylamine *N*-oxide (17 mg, 0.226 mmol) were added to a solution of $\text{Re}_2[\mu\text{-S}(2\text{-naphthyl})]_2(\text{CO})_8$ (100 mg, 0.109 mmol) and PHCy₂ (44 µl, 0.218 mmol) in tetrahydrofuran. The Re adduct was prepared according to Treichel & Tegen (1988). After stirring for 2 h, (II) was isolated by TLC separation (eluant: dichloromethane/*n*hexane 1/2) in 60% yield. Single crystals were grown from dichloromethane/*n*-pentane.

Compound (I)

Crystal data

2	
$[\operatorname{Re}_2(\operatorname{C_7H_7OS})_2(\operatorname{CO})_8]$	$D_x = 2.361 \text{ Mg m}^{-3}$
$M_r = 874.85$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 30
a = 9.731 (2) Å	reflections
b = 9.638(1) Å	$\theta = 7.323 - 22.046^{\circ}$
c = 13.772(1) Å	$\mu = 10.051 \text{ mm}^{-1}$
$\beta = 107.66 (1)^{\circ}$	T = 203 (2) K
V = 1230.8 (3) Å ³	Block, pale yellow
<i>Z</i> = 2	$0.44 \times 0.24 \times 0.18 \text{ mm}$
Data collection	

Siemens P4 diffractometer $2\theta/\omega$ scans Absorption correction: ψ scan (North et al., 1968) $T_{min} = 0.080, T_{max} = 0.164$ 3716 measured reflections 2832 independent reflections 2448 reflections with $I > 2\sigma(I)$ $\mu = 10.051 \text{ mm}^{-1}$ T = 203 (2) KBlock, pale yellow $0.44 \times 0.24 \times 0.18 \text{ mm}$ $R_{\text{int}} = 0.019$ $\theta_{\text{max}} = 27.50^{\circ}$ $h = -12 \rightarrow 1$ $k = -1 \rightarrow 12$ $l = -17 \rightarrow 17$ 3 standard reflections every 397 reflections

intensity decay: < 1%

Table 1

Selected geometric parameters (Å, °) for (I).

Re1-S1 ⁱ	2.5179 (12)	S1-C11	1.796 (4)		
Re1-S1	2.5243 (12)				
S1 ⁱ -Re1-S1	81.53 (4)	C11-S1-Re1	114.20 (16)		
$C11-S1-Re1^i$	111.92 (16)	Re1 ⁱ -S1-Re1	98.47 (4)		
Symmetry code: (i) -1	1 - r = 1 - n = 7				

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

+ 1.1289*P*]

 $w = 1/[\sigma^2(F_o^2) + (0.0352P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

Refinement

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Refinement on F^2

R[F^2 > 2\sigma(F^2)] = 0.026

wR(F^2) = 0.063

S = 1.049

2832 reflections

165 parameters

H-atom parameters constrained
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Compound (II)

Crystal data

$$\begin{split} & [\text{Re}_2(\text{C}_{12}\text{H}_{23}\text{P})_2(\text{C}_{10}\text{H}_7\text{S})_2(\text{CO})_6] \\ & M_r = 1255.44 \\ & \text{Monoclinic, } C2/c \\ & a = 24.561 (3) \text{ Å} \\ & b = 9.378 (3) \text{ Å} \\ & c = 23.722 (3) \text{ Å} \\ & \beta = 112.63 (1)^\circ \\ & V = 5043.3 (18) \text{ Å}^3 \\ & Z = 4 \end{split}$$

Data collection

Siemens P4 diffractometer ω scans Absorption correction: ψ scan (North et al., 1968) $T_{min} = 0.482, T_{max} = 0.671$ 6948 measured reflections 5783 independent reflections 2766 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.047$ $wR(F^2) = 0.117$ S = 1.0215783 reflections 284 parameters H atoms treated by a mixture of independent and constrained refinement

Table 2

Selected geometric parameters (Å, $^{\circ}$) for (II).

Re1-P1	2.484 (3)	Re1-S1	2.526 (2)
Re1-S1 ⁱ	2.523 (2)	S1-C31	1.780 (10)
$\begin{array}{c} S1^{i} - Re1 - S1 \\ C31 - S1 - Re1^{i} \end{array}$	74.64 (8)	C31-S1-Re1	115.7 (3)
	110.4 (3)	Re1 ⁱ -S1-Re1	98.93 (8)

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

In (II), H1 on P1 was refined isotropically $[P1-H1 \ 1.44 \ (8) \ \text{Å}]$, while all other H atoms were constrained and treated as riding atoms.

For both compounds, data collection: *XSCANS* (Siemens, 1995b); cell refinement: *XSCANS*; data reduction: *XSCANS*; program(s) used to solve structure: *SHELXTL* (Siemens, 1995a); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1446). Services for accessing these data are described at the back of the journal.

 $\begin{array}{l} (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.991 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.995 \ {\rm e} \ {\rm \AA}^{-3} \\ {\rm Extinction \ correction: \ SHELXTL} \\ ({\rm Siemens, \ } 1995a) \\ {\rm Extinction \ coefficient: \ } 0.00201 \ (18) \end{array}$

 $D_x = 1.653 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 35 reflections $\theta = 8.705 - 15.577^{\circ}$ $\mu = 4.988 \text{ mm}^{-1}$ T = 293 (2) KPrism, colourless $0.41 \times 0.10 \times 0.08 \text{ mm}$

 $\begin{aligned} R_{\rm int} &= 0.037\\ \theta_{\rm max} &= 27.50^\circ\\ h &= -31 \rightarrow 1\\ k &= -1 \rightarrow 12\\ l &= -28 \rightarrow 30\\ 3 \text{ standard reflections}\\ \text{every } 397 \text{ reflections}\\ \text{intensity decay: } 3\% \end{aligned}$

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\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0328P)^2 \\ &+ 11.0195P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ \Delta\rho_{\rm max} = 0.991 \ {\rm e}\ {\rm A}^{-3} \\ \Delta\rho_{\rm min} = -0.501 \ {\rm e}\ {\rm A}^{-3} \\ {\rm Extinction\ correction:\ SHELXTL} \\ &({\rm Siemens,\ }1995a) \\ {\rm Extinction\ coefficient:\ }0.00016\ (3) \end{split}
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