Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### Key indicators

Single-crystal X-ray study  $T = 100 K$ Mean  $\sigma$ (C–C) = 0.003 Å  $R$  factor = 0.025  $wR$  factor =  $0.060$ Data-to-parameter ratio = 14.8

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# A bis(benzenedithiolate)-bridged dinuclear Rh<sup>III</sup> complex with capping triphenylphosphine ligands and an  $Rh<sub>2</sub>S<sub>2</sub>$  core

The title compound,  $bis(\mu$ -benzene-1,2-dithiolato)bis[(2mercaptobenzenethiolato)triphenylphosphine)rhodium(III)] dichloromethane tetrasolvate,  $[Rh_2(C_6H_4S_2)_2(C_6H_5S_2)_2$ - $(C_{18}H_{15})_2$ ] 4CH<sub>2</sub>Cl<sub>2</sub>, isolated from the reaction of Wilkinson's catalyst  $RhCl(PPh<sub>3</sub>)<sub>3</sub>$  and benzo[1,3,2]dithiaborole, is a neutral dinuclear species with two bridging benzenedithiolate ligands and two 2-mercaptobenzenethiolate ligands chelating each Rh<sup>III</sup> ion. This compound is derived from decomposition of the borole derivative after its oxidative addition to Rh<sup>I</sup>. The compound crystallizes as the dichloromethane tetrasolvate. The molecule is centrosymmetric. The Rh—S distances in the  $Rh<sub>2</sub>S<sub>2</sub>$  core are inequivalent, measuring 2.383 (1) and  $2.409(1)$  Å.

### Comment

Since Männig & Nöth (1985) discovered that  $RhCl(PPh<sub>3</sub>)<sub>3</sub>, (I)$ , catalyzes the addition of benzo[1,3,2]dioxaborole (HBCat) to alkenes, several investigations have been reported that clarify the mechanisms of transition metal-catalyzed hydroborations (Evans et al., 1992; Burgess et al., 1991). Examples of crystal structures with Rh—B bonds involving BCat are now known, including  $RhCl(BCat)<sub>2</sub>(PPh<sub>3</sub>)<sub>2</sub>$  (Baker et al., 1993),  $RhHCl(BCat)(P<sup>i</sup>Pr<sub>3</sub>)<sub>2</sub>$  (Westcott *et al.*, 1991) and  $RhCl<sub>2</sub>$ - $(BCat)(PMe<sub>3</sub>)<sub>3</sub>$  (Souza *et al.*, 2005). However, the use of benzo[1,3,2]dithiaborole, (II), in Rh-catalyzed hydroborations has not been reported, despite our recent microwave-assisted hydroborations with the dioxaborole analogue (Hadebe & Robinson, 2006a,b).



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Compound (II) oxidatively adds to Wilkinson's catalyst (I) to give (III). This key reaction step was evident from the Received 4 December 2006 Accepted 7 December 2006



#### Figure 1

The molecular structure of (IV), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 80% probability level. All H atoms other than those of the SH groups, and the four dichloromethane solvent molecules, have been omitted for clarity. Unlabelled atoms are related to labelled atoms by  $(1 - x, 1 - y, 1 - z)$ .



Figure 2

The unit-cell contents of (IV). Displacement ellipsoids for solvent non-H atoms are at the 50% probability level. All other atoms are shown only as the intersections and end-points of bonds represented as cylinders (including solvent H atoms). H atoms involved in intramolecular (red dashed lines) and intermolecular (blue lines) hydrogen bonds are shown and labelled. Unlabelled atoms are related to labelled atoms by  $(1 - x,$  $1 - y$ ,  $1 - z$ ).

dramatic colour change of the solution (brick red to dark green). Crystallization of (III) was, however, thwarted by its decomposition (mechanism unknown) to the dinuclear  $Rh<sup>III</sup>$ complex, (IV) (18 electrons per Rh centre), the structure of which we present here.

The centrosymmetric compound, (IV), belongs to an interesting class of cluster compounds containing an  $Rh_2S_2$ core bridged by two benzene-1,2-bis(thiolate) groups (Fig. 1). The Rh–S bond distances of (IV) within the  $Rh<sub>2</sub>S<sub>2</sub>$  core (Table 1) are significantly different, at 2.3829 (5) (Rh—S4) and 2.4093 (5)  $\AA$  (Rh - S4<sup>i</sup>) [symmetry code: (i)  $1 - x$ ,  $1 - y$ ,  $1 - z$ . Similar asymmetry for the in-plane Rh–S distances exists for the dinuclear complex  $[(Cp*)Rh-(1,2-C<sub>6</sub>H<sub>4</sub>S<sub>2</sub>)<sub>2</sub>$  $Rh(Cp^*)$ ] (Xi *et al.*, 1997). This most likely reflects the fact that the trans ligands differ for each bridging sulfide atom. Indeed, the shorter Rh—S distance is observed for the sulfide donor

*trans* to P (a good  $\pi$ -acceptor ligand), while the longer Rh–S distance occurs for the sulfide donor *trans* to a good  $\sigma$ -donor  $(i.e.$  the sulfide donor of the chelating mercaptobenzenethiolate ligand). The internuclear Rh. Rh distance is 3.6056 (5)  $\AA$ , slightly longer than the Rh $\cdots$ Rh distance of 3.531 (1) Å reported for  $[(Cp^*)Rh-(1,2-C<sub>6</sub>H<sub>4</sub>S<sub>2</sub>)<sub>2</sub>-Rh(Cp^*)]$ (Xi et al., 1997), a finding consistent with slight expansion of the Rh<sub>2</sub>S<sub>2</sub> core of (IV) driven by the greater  $\sigma$ -donor power of the *trans* ligands relative to  $Cp^*$  (where  $Cp^*$  = pentamethylcyclopentadienate).

The unit-cell contents of (IV) are shown in Fig. 2. Four dichloromethane solvent molecules are present. The most significant intermolecular interaction is the  $C-H \cdots S$ hydrogen bond that exists between one of the solvent molecules and atom S1 of the mercaptobenzenethiolate ligand. The geometries of this interaction and of the three intramolecular hydrogen bonds for (IV) are listed in Table 2.

# Experimental

A solution of benzo[1,3,2]dithiaborole in  $CH_2Cl_2$  (0.4 ml, 0.16 mmol) was injected into a flame-dried nitrogen-purged septum-capped test tube. To this solution was added  $RhCl(PPh<sub>3</sub>)<sub>3</sub>$  (4.72 mg, 5.1  $\mu$ mol), which had been dissolved in dichloromethane  $(3.5 \text{ ml})$  n a separate flame-dried nitrogen-flushed test tube. The resulting mixture was shaken vigorously and allowed to stand for ca 2 weeks. Evaporation of the solvent through the septum afforded a dark-green powder containing, as the minor component, brick-red crystals of (IV) (m.p. 418.5–422.6 K).

Crystal data



Data collection

Oxford Diffraction Xcalibur2 CCD area-detector diffractometer  $\omega$  scans

Absorption correction: analytical (CrysAlis RED; Oxford Diffraction, 2005; Clark & Reid,

# Refinement

Refinement on  $F^2$ 

- $R[F^2 > 2\sigma(F^2)] = 0.025$
- $wR(F^2) = 0.06$  $S = 1.08$
- 5650 reflections
- 383 parameters
- H atoms treated by a mixture of
- independent and constrained refinement

1995)  $T_{\text{min}} = 0.808$ ,  $T_{\text{max}} = 0.880$ 15977 measured reflections 5650 independent reflections 5476 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.021$  $\theta_{\text{max}} = 25.1^{\circ}$ 



Table 1 Selected geometric parameters  $(\AA, \degree)$ .

$P1 - Rh1$	2.3542(5)	$Rh1-S4$	2.3829(5)
$Rh1-S1$	2.3483(5)	$Rh1-S4$ <sup>i</sup>	2.4093(5)
$Rh1-S3$	2.3512(6)	$S2 - H100$	1.31(3)
$Rh1 - S2$	2.3614(6)		
$S1 - Rh1 - S3$	89.708 (19)	$S2 - Rh1 - S4$	88.307 (19)
$S1 - Rh1 - P1$	87.264 (19)	$S1 - Rh1 - S4$	172.623 (19)
$S3 - Rh1 - P1$	94.227 (19)	$S3 - Rh1 - S4$	86.131 (19)
$S1 - Rh1 - S2$	87.451 (19)	$P1 - Rh1 - S4$	99.105 (19)
$S3 - Rh1 - S2$	173.370 (19)	$S2 - Rh1 - S4$	96.034 (19)
$P1 - Rh1 - S2$	91.621 (19)	$S4 - Rh1 - S4$	82.405 (19)
$S1 - Rh1 - S4$	91.223 (19)	$Rh1 - S2 - H100$	102.1(12)
$S3 - Rh1 - S4$	85.765 (18)	$Rh1 - S4 - Rh1$	97.595 (19)
$P1 - Rh1 - S4$	178.486 (19)		
$C26 - C21 - S3 - Rh1$	17.52 (19)	$Rh1 - S4 - C26 - C21$	$-21.29(18)$
$C22 - C21 - S3 - Rh1$	$-164.84(16)$	$Rh1 - S4 - C26 - C25$	160.60(15)

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

Table 2 Hydrogen-bond geometry  $(\mathring{A}, \degree)$ .

$D - H \cdots A$	$D-H$	$H \cdots A$	$D\cdots A$	$D - H \cdots A$
$S2 - H100 \cdots S3^i$	1.31(3)	2.78(3)	3.5137(8)	113.8(18)
$C1S-H1S1\cdots S1$	0.99	2.86	3.660(3)	139
$C56 - H56 \cdots S1$	0.95	2.77	3.396(2)	125
$C56 - H56 \cdots S3$	0.95	2.70	3.395(2)	131

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

The H atom attached to atom S2 was located in a difference Fourier map and refined isotropically without constraints. The remaining H atoms were positioned geometrically and refined using a riding model, with C-H = 0.97 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$  for methylene H, and with C—H = 0.93 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$  for aromatic H.

Data collection: CrysAlis CCD (Oxford Diffraction, 2005); cell refinement: CrysAlis RED (Oxford Diffraction, 2005); data reduction: CrysAlis RED; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

Financial support from SASOL is gratefully acknowledged, as are Mr Arno de Klerk and Dr Hein Strauss for collaborative involvement in this ongoing project. We thank Mr Craig Grimmer for NMR assistance and the University of KwaZulu-Natal for financial support.

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