

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

## fac-Aquadichloridotris(tetramethylene sulfoxide- $\kappa$ S)ruthenium(II)

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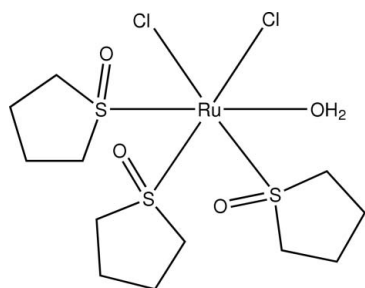
Received 5 December 2008; accepted 6 January 2009

Key indicators: single-crystal X-ray study;  $T = 90$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.037;  $wR$  factor = 0.076; data-to-parameter ratio = 30.1.

The title molecule,  $[\text{RuCl}_2(\text{C}_4\text{H}_8\text{OS})_3(\text{H}_2\text{O})]$ , is the isomer with the two chloride ligands *cis* and the three *S*-coordinated tetramethylene sulfoxide ligands facial relative to the Ru(II) center. The Ru—Cl distances are 2.4161 (7) and 2.4317 (7) Å, the Ru—O distance is 2.1540 (19) Å, and the Ru—S distances are in the range 2.2254 (8)–2.2657 (7) Å, with the shortest being that *trans* to the aqua ligand. The coordinated water molecule forms intermolecular hydrogen bonds with Cl and sulfoxide O atoms.

### Related literature

For background literature, see: Aldinucci *et al.* (2007). For related structures, see: Srivastava & Fronczek (2003); Srivastava *et al.* (2004); Allen (2002). For hydrogen-bonding patterns, see: Etter (1990).



### Experimental

#### Crystal data

$[\text{RuCl}_2(\text{C}_4\text{H}_8\text{OS})_3(\text{H}_2\text{O})]$   
 $M_r = 502.48$   
 Monoclinic,  $P2_1/n$   
 $a = 14.302$  (3) Å

$b = 7.7877$  (15) Å  
 $c = 17.248$  (3) Å  
 $\beta = 109.917$  (9)°  
 $V = 1806.2$  (6) Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 1.52$  mm<sup>-1</sup>

$T = 90.0$  (5) K  
 $0.22 \times 0.10 \times 0.05$  mm

#### Data collection

Nonius KappaCCD diffractometer (with an Oxford Cryosystems Cryostream cooler)  
 Absorption correction: multi-scan (SCALEPACK; Otwinowski & Minor, 1997)  
 $T_{\min} = 0.731$ ,  $T_{\max} = 0.928$   
 25916 measured reflections  
 5982 independent reflections  
 4610 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.045$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.076$   
 $S = 1.02$   
 5982 reflections  
 199 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.87$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -1.12$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O4—H41...O1 <sup>i</sup>	0.80	1.99	2.785 (3)	169
O4—H42...Cl2 <sup>ii</sup>	0.80	2.37	3.116 (2)	156

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

Data collection: COLLECT (Nonius, 2000); cell refinement: SCALEPACK (Otwinowski & Minor, 1997); data reduction: DENZO (Otwinowski & Minor, 1997) and SCALEPACK; program(s) used to solve structure: SIR97 (Altomare *et al.*, 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

Financial support provided by the Research Corporation (Cottrell College Science Award, #CC 6234 to RSS) and the BoR, Louisiana, is greatly appreciated. The purchase of the diffractometer was made possible by grant No. LEQSF (1999–2000)-ENH-TR-13, administered by the Louisiana Board of Regents.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2128).

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## supporting information

*Acta Cryst.* (2009). E65, m170 [doi:10.1107/S1600536809000439]

***fac*-Aquadichloridotris(tetramethylene sulfoxide- $\kappa$ S)ruthenium(II)****Radhey S. Srivastava, Carlos F. Gonzales and Frank R. Fronczek****S1. Comment**

During the course of our studies on ruthenium-DMSO/TMSO complexes, *mer*-RuCl<sub>3</sub>(TMSO)<sub>3</sub> was refluxed with methyl-*p*-tolylsulfide in absolute ethanol for 1 h. In view of anticancer properties of Ru-DMSO/TMSO complexes, we envision to interact *mer*-RuCl<sub>3</sub>(TMSO)<sub>3</sub>, (1) with sulfur donor ligands, because sulfur-containing ligands are able to bind the metal center strongly and prevent interactions with sulfur-containing enzymes (Aldinucci *et al.*, 2007). In fact, these reactions are believed to be responsible for the nephrotoxicity induced by the platinum (II)-based drugs. However, the compound (2) was hydrolyzed on long standing in solution, and finally the title compound, *fac*-[RuCl<sub>2</sub>(TMSO)<sub>3</sub>(H<sub>2</sub>O)] (3) was isolated. A plausible mechanism of the formation of (3) is shown in scheme 2.

There are three geometrical isomers of the title compound: *trans,mer*; *cis,mer*; and *fac*. The reported structure is found to be the latter, with chloro groups *cis* and TMSO groups facial, as shown in Fig. 1. More isomers are possible, considering that the TMSO ligands may be coordinated to Ru through either S or O in the same complex, (Srivastava & Fronczek, 2003; Srivastava *et al.*, 2004); however, all are S-coordinated here. Relevant bond distances are given in the supplementary Tables. Most noteworthy is that the Ru—S<sub>3</sub> distance, *trans* to water, is 0.03–0.04 Å shorter than the two Ru—S distances *trans* to Cl. While a search of the Cambridge Structural Database (version 5.29, Jan. 2008; Allen, 2002) for Ru complexes with S-bonded TMSO *trans* to water produced no hits, eleven examples of such DMSO complexes were found, refcodes AQAXIZ, AQAXOF, AQAXUL, BINBAC, CECSUZ, QUDRUC, TEXMOZ, TEXNEQ, TEXNIU, WOHNEM, AND WOHNIQ. Those have mean Ru—O distance 2.142 Å and mean Ru—S distance 2.256 Å. Our Ru—O distance, 2.1540 (19) Å, is near the high end of that sample, and our Ru—S distance, Ru1 S3 2.2254 (8) Å, is shorter than any in that sample.

The coordinated water molecule donates an intermolecular hydrogen bond to sulfoxide O and another to Cl, on two different molecules related by unit translation in the b direction. Thus, rings of graph set (Etter, 1990) symbol  $R^2_2(9)$  form chains along [010], propagated by the 2<sub>1</sub> axis, as shown in Fig. 2.

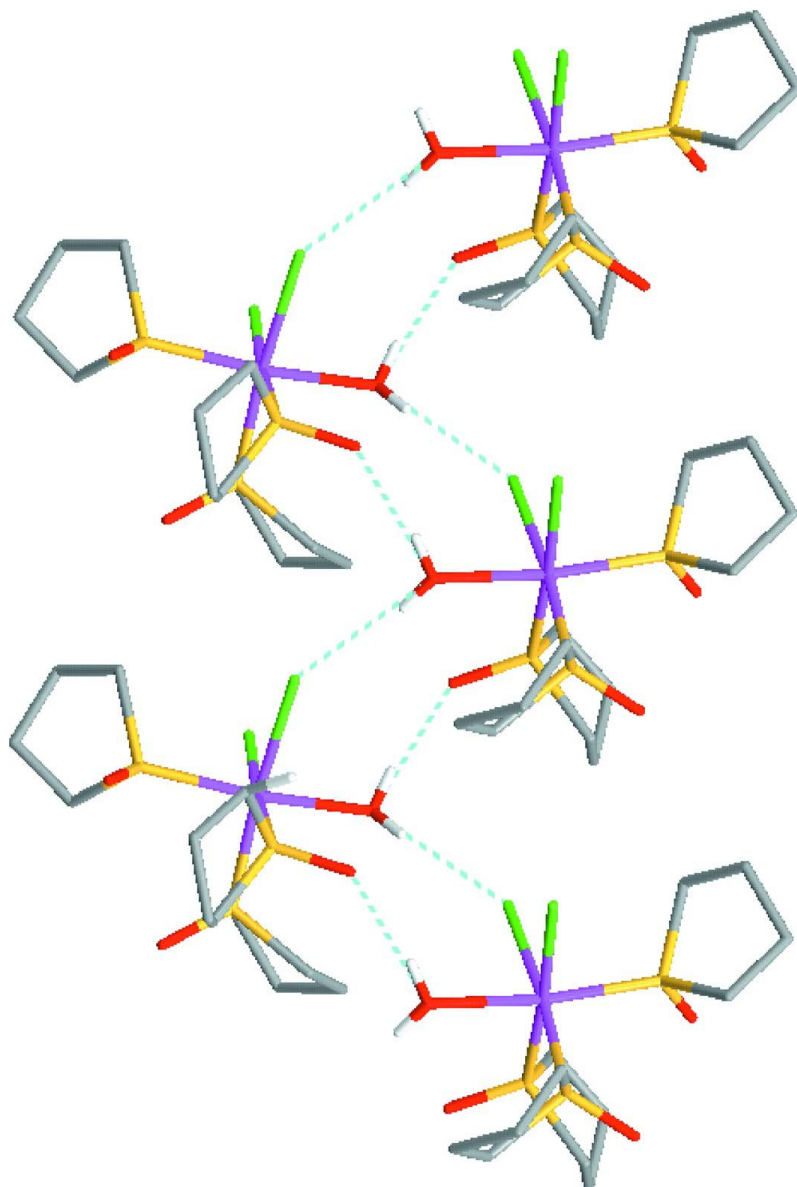
**S2. Experimental**

*mer*-RuCl<sub>3</sub>(TMSO)<sub>3</sub> (0.166 g, 0.233 mmol) was refluxed with methyl-*p*-tolylsulfide (73 μl, 0.533 mmol) in absolute ethanol (15 ml) for 2 h, followed by cooling to room temperature. Upon standing for eight months, colorless needles of the title compound formed.

**S3. Refinement**

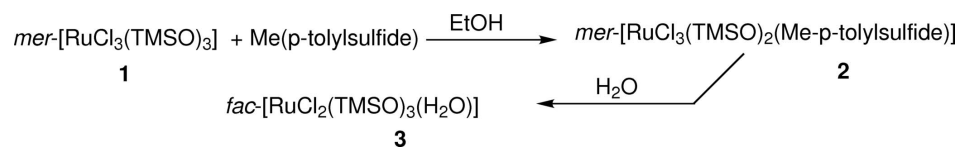
H atoms on C were placed in idealized positions with C—H distances 0.99 Å and thereafter treated as riding. Water H atoms were located in difference maps, idealized to have O—H distance 0.80 Å, and treated as riding.  $U_{iso}$  for H was assigned as 1.2 times  $U_{eq}$  of the attached atoms (1.5 for H<sub>2</sub>O). The largest negative feature in the final difference map was located 0.75 Å from the Ru position.





**Figure 2**

A portion of a hydrogen-bonded chain in the [010] direction.



**Figure 3**

The formation of the title compound.

**fac-Aquadichloridotris(tetramethylene sulfoxide- $\kappa$ S)ruthenium(II)***Crystal data*[RuCl<sub>2</sub>(C<sub>4</sub>H<sub>8</sub>OS)<sub>3</sub>(H<sub>2</sub>O)] $M_r = 502.48$ Monoclinic,  $P2_1/n$ 

Hall symbol: -P 2yn

 $a = 14.302$  (3) Å $b = 7.7877$  (15) Å $c = 17.248$  (3) Å $\beta = 109.917$  (9)° $V = 1806.2$  (6) Å<sup>3</sup> $Z = 4$  $F(000) = 1024$  $D_x = 1.848$  Mg m<sup>-3</sup>Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 6027 reflections

 $\theta = 2.5$ – $31.5$ ° $\mu = 1.52$  mm<sup>-1</sup> $T = 90$  K

Needle, colorless

 $0.22 \times 0.10 \times 0.05$  mm*Data collection*

Nonius KappaCCD

diffractometer (with an Oxford Cryosystems  
Cryostream cooler)

Radiation source: fine-focus sealed tube

Graphite monochromator

 $\omega$  and  $\varphi$  scans

Absorption correction: multi-scan

(SCALEPACK; Otwinowski &amp; Minor, 1997)

 $T_{\min} = 0.731$ ,  $T_{\max} = 0.928$ 

25916 measured reflections

5982 independent reflections

4610 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.045$  $\theta_{\max} = 31.5$ °,  $\theta_{\min} = 2.9$ ° $h = -20$ → $21$  $k = -11$ → $11$  $l = -25$ → $25$ *Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.037$  $wR(F^2) = 0.076$  $S = 1.02$ 

5982 reflections

199 parameters

0 restraints

Primary atom site location: structure-invariant  
direct methodsSecondary atom site location: difference Fourier  
mapHydrogen site location: inferred from  
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0256P)^2 + 2.7566P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\max} = 0.002$  $\Delta\rho_{\max} = 0.87$  e Å<sup>-3</sup> $\Delta\rho_{\min} = -1.12$  e Å<sup>-3</sup>*Special details*

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å<sup>2</sup>)*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
Ru1	0.589322 (15)	0.17421 (3)	0.298795 (12)	0.00590 (5)
Cl1	0.63294 (5)	-0.00620 (8)	0.41988 (4)	0.01063 (12)
Cl2	0.59746 (5)	-0.07251 (8)	0.21491 (4)	0.00956 (12)

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S1	0.55748 (5)	0.33080 (8)	0.18270 (4)	0.00718 (12)
S2	0.59211 (5)	0.41572 (8)	0.37280 (4)	0.00785 (12)
S3	0.42982 (5)	0.11754 (8)	0.27675 (4)	0.00811 (12)
O1	0.64973 (14)	0.3915 (2)	0.16851 (11)	0.0112 (4)
O2	0.49707 (14)	0.5110 (2)	0.35546 (12)	0.0125 (4)
O3	0.35604 (14)	0.1775 (3)	0.19762 (12)	0.0133 (4)
O4	0.74739 (13)	0.1973 (2)	0.32611 (11)	0.0094 (4)
H41	0.7697	0.1034	0.3254	0.014*
H42	0.7735	0.2581	0.3021	0.014*
C1	0.4799 (2)	0.2324 (3)	0.08568 (16)	0.0117 (5)
H1A	0.5196	0.2111	0.0495	0.014*
H1B	0.4527	0.1216	0.0965	0.014*
C2	0.3955 (2)	0.3583 (4)	0.04466 (17)	0.0142 (6)
H2A	0.3353	0.3260	0.0573	0.017*
H2B	0.3790	0.3561	-0.0159	0.017*
C3	0.4301 (2)	0.5380 (4)	0.07805 (16)	0.0129 (5)
H3A	0.4794	0.5831	0.0545	0.016*
H3B	0.3731	0.6183	0.0645	0.016*
C4	0.4770 (2)	0.5151 (3)	0.17114 (16)	0.0109 (5)
H4A	0.4256	0.4934	0.1964	0.013*
H4B	0.5157	0.6181	0.1970	0.013*
C5	0.6859 (2)	0.5663 (3)	0.36345 (16)	0.0108 (5)
H5A	0.7141	0.5249	0.3218	0.013*
H5B	0.6559	0.6808	0.3463	0.013*
C6	0.7667 (2)	0.5771 (4)	0.44766 (17)	0.0152 (6)
H6A	0.8007	0.6897	0.4548	0.018*
H6B	0.8167	0.4854	0.4538	0.018*
C7	0.7149 (2)	0.5548 (4)	0.51139 (17)	0.0145 (6)
H7A	0.7644	0.5366	0.5671	0.017*
H7B	0.6748	0.6576	0.5128	0.017*
C8	0.6484 (2)	0.3978 (3)	0.48391 (16)	0.0110 (5)
H8A	0.5967	0.3964	0.5101	0.013*
H8B	0.6881	0.2911	0.4990	0.013*
C9	0.3821 (2)	0.1780 (4)	0.35774 (17)	0.0132 (5)
H9A	0.4359	0.1791	0.4121	0.016*
H9B	0.3507	0.2928	0.3471	0.016*
C10	0.3059 (2)	0.0401 (4)	0.35450 (18)	0.0148 (6)
H10A	0.2462	0.0530	0.3046	0.018*
H10B	0.2856	0.0455	0.4039	0.018*
C11	0.3595 (2)	-0.1292 (4)	0.35214 (18)	0.0149 (6)
H11A	0.4113	-0.1516	0.4063	0.018*
H11B	0.3117	-0.2258	0.3390	0.018*
C12	0.4068 (2)	-0.1113 (3)	0.28551 (18)	0.0119 (5)
H12A	0.3614	-0.1562	0.2323	0.014*
H12B	0.4699	-0.1764	0.3010	0.014*

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Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Ru1	0.00693 (9)	0.00402 (9)	0.00659 (9)	0.00017 (7)	0.00210 (7)	0.00032 (7)
Cl1	0.0146 (3)	0.0069 (3)	0.0095 (3)	0.0008 (2)	0.0030 (2)	0.0025 (2)
Cl2	0.0109 (3)	0.0061 (3)	0.0119 (3)	0.0004 (2)	0.0042 (2)	-0.0016 (2)
S1	0.0081 (3)	0.0058 (3)	0.0071 (3)	-0.0008 (2)	0.0020 (2)	0.0000 (2)
S2	0.0099 (3)	0.0056 (3)	0.0077 (3)	0.0003 (2)	0.0026 (2)	0.0005 (2)
S3	0.0086 (3)	0.0069 (3)	0.0087 (3)	0.0000 (2)	0.0028 (2)	0.0008 (2)
O1	0.0116 (9)	0.0101 (9)	0.0133 (9)	-0.0033 (7)	0.0060 (8)	0.0002 (7)
O2	0.0117 (9)	0.0097 (9)	0.0171 (10)	0.0036 (7)	0.0060 (8)	0.0003 (8)
O3	0.0093 (9)	0.0158 (9)	0.0127 (9)	-0.0009 (8)	0.0009 (7)	0.0055 (8)
O4	0.0108 (9)	0.0049 (8)	0.0132 (9)	-0.0012 (7)	0.0049 (7)	-0.0001 (7)
C1	0.0127 (13)	0.0110 (12)	0.0102 (12)	-0.0029 (10)	0.0024 (10)	-0.0015 (10)
C2	0.0131 (13)	0.0156 (14)	0.0110 (13)	-0.0005 (10)	0.0005 (10)	0.0025 (10)
C3	0.0146 (13)	0.0125 (12)	0.0113 (13)	0.0028 (11)	0.0038 (10)	0.0071 (10)
C4	0.0115 (12)	0.0086 (12)	0.0121 (13)	0.0040 (10)	0.0034 (10)	0.0024 (10)
C5	0.0143 (13)	0.0061 (11)	0.0123 (12)	-0.0014 (10)	0.0052 (10)	0.0009 (10)
C6	0.0160 (14)	0.0133 (13)	0.0135 (13)	-0.0039 (11)	0.0014 (11)	-0.0030 (11)
C7	0.0197 (15)	0.0130 (13)	0.0083 (12)	-0.0027 (11)	0.0014 (11)	-0.0033 (10)
C8	0.0161 (13)	0.0092 (12)	0.0080 (12)	0.0007 (10)	0.0044 (10)	0.0006 (10)
C9	0.0135 (13)	0.0131 (12)	0.0159 (13)	-0.0003 (11)	0.0086 (11)	-0.0031 (11)
C10	0.0138 (13)	0.0178 (14)	0.0156 (14)	-0.0047 (11)	0.0087 (11)	0.0012 (11)
C11	0.0180 (14)	0.0121 (12)	0.0156 (14)	-0.0055 (11)	0.0068 (11)	0.0024 (11)
C12	0.0107 (12)	0.0063 (11)	0.0199 (14)	-0.0030 (10)	0.0067 (11)	0.0009 (10)

Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

Ru1—O4	2.1540 (19)	C3—H3B	0.9900
Ru1—S3	2.2254 (8)	C4—H4A	0.9900
Ru1—S1	2.2546 (7)	C4—H4B	0.9900
Ru1—S2	2.2657 (7)	C5—C6	1.519 (4)
Ru1—Cl1	2.4161 (7)	C5—H5A	0.9900
Ru1—Cl2	2.4317 (7)	C5—H5B	0.9900
S1—O1	1.498 (2)	C6—C7	1.531 (4)
S1—C4	1.807 (3)	C6—H6A	0.9900
S1—C1	1.831 (3)	C6—H6B	0.9900
S2—O2	1.487 (2)	C7—C8	1.521 (4)
S2—C8	1.814 (3)	C7—H7A	0.9900
S2—C5	1.830 (3)	C7—H7B	0.9900
S3—O3	1.487 (2)	C8—H8A	0.9900
S3—C9	1.813 (3)	C8—H8B	0.9900
S3—C12	1.828 (3)	C9—C10	1.517 (4)
O4—H41	0.8000	C9—H9A	0.9900
O4—H42	0.8000	C9—H9B	0.9900
C1—C2	1.529 (4)	C10—C11	1.532 (4)
C1—H1A	0.9900	C10—H10A	0.9900
C1—H1B	0.9900	C10—H10B	0.9900

C2—C3	1.530 (4)	C11—C12	1.525 (4)
C2—H2A	0.9900	C11—H11A	0.9900
C2—H2B	0.9900	C11—H11B	0.9900
C3—C4	1.525 (4)	C12—H12A	0.9900
C3—H3A	0.9900	C12—H12B	0.9900
O4—Ru1—S3	172.92 (5)	C3—C4—S1	104.15 (18)
O4—Ru1—S1	91.63 (5)	C3—C4—H4A	110.9
S3—Ru1—S1	94.04 (3)	S1—C4—H4A	110.9
O4—Ru1—S2	89.52 (5)	C3—C4—H4B	110.9
S3—Ru1—S2	94.66 (3)	S1—C4—H4B	110.9
S1—Ru1—S2	90.62 (3)	H4A—C4—H4B	108.9
O4—Ru1—C11	85.23 (5)	C6—C5—S2	107.04 (18)
S3—Ru1—C11	88.86 (3)	C6—C5—H5A	110.3
S1—Ru1—C11	175.45 (2)	S2—C5—H5A	110.3
S2—Ru1—C11	92.63 (3)	C6—C5—H5B	110.3
O4—Ru1—C12	86.37 (5)	S2—C5—H5B	110.3
S3—Ru1—C12	89.75 (2)	H5A—C5—H5B	108.6
S1—Ru1—C12	86.28 (3)	C5—C6—C7	106.5 (2)
S2—Ru1—C12	174.78 (2)	C5—C6—H6A	110.4
C11—Ru1—C12	90.24 (3)	C7—C6—H6A	110.4
O1—S1—C4	107.13 (12)	C5—C6—H6B	110.4
O1—S1—C1	106.01 (12)	C7—C6—H6B	110.4
C4—S1—C1	93.82 (12)	H6A—C6—H6B	108.6
O1—S1—Ru1	113.14 (8)	C8—C7—C6	105.8 (2)
C4—S1—Ru1	117.12 (9)	C8—C7—H7A	110.6
C1—S1—Ru1	117.49 (9)	C6—C7—H7A	110.6
O2—S2—C8	107.31 (12)	C8—C7—H7B	110.6
O2—S2—C5	108.06 (12)	C6—C7—H7B	110.6
C8—S2—C5	93.85 (12)	H7A—C7—H7B	108.7
O2—S2—Ru1	117.48 (8)	C7—C8—S2	105.84 (18)
C8—S2—Ru1	116.59 (9)	C7—C8—H8A	110.6
C5—S2—Ru1	110.80 (9)	S2—C8—H8A	110.6
O3—S3—C9	106.85 (13)	C7—C8—H8B	110.6
O3—S3—C12	106.88 (12)	S2—C8—H8B	110.6
C9—S3—C12	93.64 (13)	H8A—C8—H8B	108.7
O3—S3—Ru1	117.39 (8)	C10—C9—S3	104.01 (19)
C9—S3—Ru1	116.73 (10)	C10—C9—H9A	111.0
C12—S3—Ru1	112.49 (9)	S3—C9—H9A	111.0
Ru1—O4—H41	108.3	C10—C9—H9B	111.0
Ru1—O4—H42	125.1	S3—C9—H9B	111.0
H41—O4—H42	105.9	H9A—C9—H9B	109.0
C2—C1—S1	106.96 (18)	C9—C10—C11	104.5 (2)
C2—C1—H1A	110.3	C9—C10—H10A	110.9
S1—C1—H1A	110.3	C11—C10—H10A	110.9
C2—C1—H1B	110.3	C9—C10—H10B	110.9
S1—C1—H1B	110.3	C11—C10—H10B	110.9
H1A—C1—H1B	108.6	H10A—C10—H10B	108.9



C1—C2—C3	108.0 (2)	C12—C11—C10	107.2 (2)
C1—C2—H2A	110.1	C12—C11—H11A	110.3
C3—C2—H2A	110.1	C10—C11—H11A	110.3
C1—C2—H2B	110.1	C12—C11—H11B	110.3
C3—C2—H2B	110.1	C10—C11—H11B	110.3
H2A—C2—H2B	108.4	H11A—C11—H11B	108.5
C4—C3—C2	105.0 (2)	C11—C12—S3	106.86 (19)
C4—C3—H3A	110.7	C11—C12—H12A	110.4
C2—C3—H3A	110.7	S3—C12—H12A	110.4
C4—C3—H3B	110.7	C11—C12—H12B	110.4
C2—C3—H3B	110.7	S3—C12—H12B	110.4
H3A—C3—H3B	108.8	H12A—C12—H12B	108.6
O4—Ru1—S1—O1	-0.88 (10)	S1—Ru1—S3—C12	129.51 (10)
S3—Ru1—S1—O1	-176.63 (9)	S2—Ru1—S3—C12	-139.54 (10)
S2—Ru1—S1—O1	88.66 (9)	C11—Ru1—S3—C12	-46.99 (10)
C12—Ru1—S1—O1	-87.14 (9)	C12—Ru1—S3—C12	43.26 (10)
O4—Ru1—S1—C4	-126.24 (11)	O1—S1—C1—C2	-105.28 (19)
S3—Ru1—S1—C4	58.01 (11)	C4—S1—C1—C2	3.8 (2)
S2—Ru1—S1—C4	-36.70 (11)	Ru1—S1—C1—C2	127.11 (16)
C12—Ru1—S1—C4	147.50 (11)	S1—C1—C2—C3	23.5 (3)
O4—Ru1—S1—C1	123.22 (11)	C1—C2—C3—C4	-46.0 (3)
S3—Ru1—S1—C1	-52.53 (11)	C2—C3—C4—S1	47.2 (2)
S2—Ru1—S1—C1	-147.24 (10)	O1—S1—C4—C3	78.5 (2)
C12—Ru1—S1—C1	36.96 (10)	C1—S1—C4—C3	-29.5 (2)
O4—Ru1—S2—O2	158.55 (10)	Ru1—S1—C4—C3	-153.17 (15)
S3—Ru1—S2—O2	-27.18 (9)	O2—S2—C5—C6	117.47 (19)
S1—Ru1—S2—O2	66.92 (9)	C8—S2—C5—C6	7.9 (2)
C11—Ru1—S2—O2	-116.25 (9)	Ru1—S2—C5—C6	-112.51 (17)
O4—Ru1—S2—C8	-72.00 (12)	S2—C5—C6—C7	-33.1 (3)
S3—Ru1—S2—C8	102.28 (11)	C5—C6—C7—C8	48.6 (3)
S1—Ru1—S2—C8	-163.62 (11)	C6—C7—C8—S2	-41.6 (3)
C11—Ru1—S2—C8	13.20 (11)	O2—S2—C8—C7	-90.7 (2)
O4—Ru1—S2—C5	33.70 (11)	C5—S2—C8—C7	19.5 (2)
S3—Ru1—S2—C5	-152.03 (9)	Ru1—S2—C8—C7	135.11 (16)
S1—Ru1—S2—C5	-57.93 (10)	O3—S3—C9—C10	80.2 (2)
C11—Ru1—S2—C5	118.90 (10)	C12—S3—C9—C10	-28.6 (2)
S1—Ru1—S3—O3	4.88 (10)	Ru1—S3—C9—C10	-146.10 (16)
S2—Ru1—S3—O3	95.83 (10)	S3—C9—C10—C11	48.1 (2)
C11—Ru1—S3—O3	-171.62 (10)	C9—C10—C11—C12	-48.9 (3)
C12—Ru1—S3—O3	-81.37 (10)	C10—C11—C12—S3	26.8 (3)
S1—Ru1—S3—C9	-123.92 (11)	O3—S3—C12—C11	-107.7 (2)
S2—Ru1—S3—C9	-32.96 (11)	C9—S3—C12—C11	1.2 (2)
C11—Ru1—S3—C9	59.59 (11)	Ru1—S3—C12—C11	122.10 (17)
C12—Ru1—S3—C9	149.83 (11)		

*Hydrogen-bond geometry (Å, °)*

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
O4—H41···O1 <sup>i</sup>	0.80	1.99	2.785 (3)	169
O4—H42···Cl2 <sup>ii</sup>	0.80	2.37	3.116 (2)	156

Symmetry codes: (i)  $-x+3/2, y-1/2, -z+1/2$ ; (ii)  $-x+3/2, y+1/2, -z+1/2$ .