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Key indicators

Single-crystal X-ray study
 $T = 110$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.041
 wR factor = 0.086
Data-to-parameter ratio = 35.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.***mer*-(2,2'-Bipyridine)trichloro(tetramethylene sulfoxide)ruthenium(III) dichloromethane solvate**

The title compound, *mer*-[RuCl₃(C₁₀H₈N₂)(C₄H₈OS)]·CH₂Cl₂ or *mer*-[RuCl₃(bpy)(TMSO)]·CH₂Cl₂, has the TMSO ligand coordinated through the S atom, with an Ru—S distance of 2.3042 (7) Å. The Ru—Cl distances are in the range 2.3324 (7)–2.3649 (8) Å, with the longest *trans* to N. The solvent molecule is ordered.

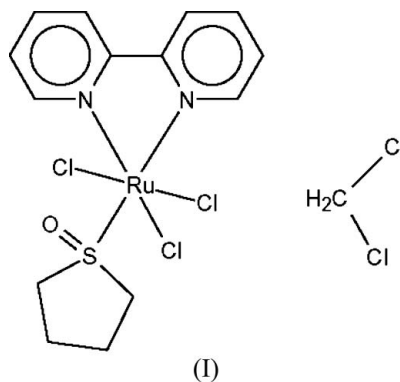
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Comment

Seeking to substitute one of the ligands in the title complex *mer*-[RuCl₃(bpy)(TMSO)] (bpy is 2,2'-bipyridine), (I), with *N,N*-diethyl-4-nitrosoaniline during the course of our general interest in Ru–halide TMSO complexes, the dichloromethane solvate of the starting material was recovered in crystalline form, as described in the *Experimental* section. Crystal structure analysis revealed that *N,N*-diethyl-4-nitrosoaniline did not substitute any of the ligands in the parent compound, presumably for steric reasons. Nevertheless, the results are of interest in view of the small amount of structural data available on Ru^{III}–TMSO complexes. We have also recently reported the structure of the unsolvated compound (Srivastava & Fronczek, 2005).



The TMSO ligand, which may be either S- or O-coordinated, coordinates *via* the S atom in this case, with the S=O bond *anti* to the central Ru—Cl bond, as was observed in the unsolvated compound (Srivastava & Fronczek, 2005). The bite angle of the bidentate bipyridine, N1—Ru—N2, is 78.49 (8)°, typical of bpy coordinated to second or third row transition metals (Constable, 1989; Constable *et al.*, 2003). This also agrees well with the bite angle of 78.16 (9)° in the unsolvated structure. The Ru—N bond lengths are slightly shorter than the typical Ru^{II}—N(bpy) distance of 2.099 Å (Garas *et al.*, 2000; Wang *et al.*, 1999), as expected from the smaller radius of Ru^{III}. The Ru—N1 distance, *trans* to Cl, is shorter than Ru—N2, *trans* to TMSO, by an amount of marginal statistical significance. However, the difference is probably real, as it was

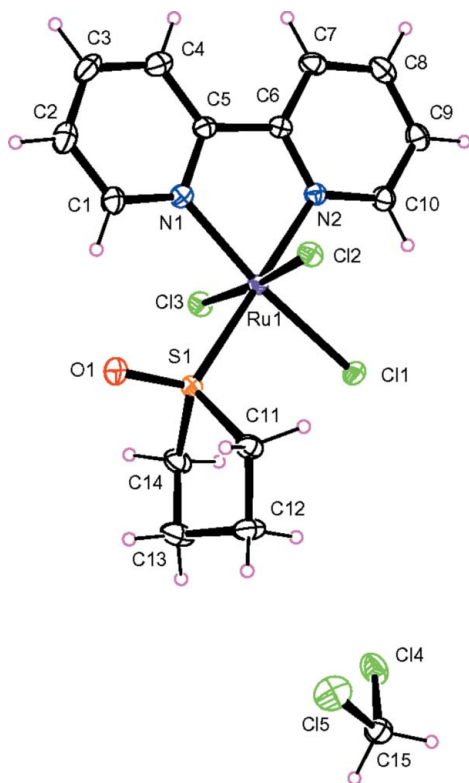


Figure 1
The structure of the asymmetric unit of (I), with displacement ellipsoids at the 50% probability level.

also observed in the unsolvated structure. The short Ru—N1 distance is accompanied by a Ru—Cl1 distance *trans* to it, which is somewhat longer than the two Ru—Cl distances *trans* to each other.

The Ru—S bond is significantly shorter than the average value of 2.3866 Å found for *mer*-[RuCl₃(TMSO)₃] (Yapp *et al.*, 1990) and other Ru^{III}—S compounds, *trans* to TMSO—S in the precursor (Alessio *et al.*, 1990, 1991; Calligaris *et al.*, 1993; Jaswal *et al.*, 1990). The Ru—S bond distances in the present compound do not show any substantial reduction with respect to *cis*-[RuCl₂(TMSO)₄] (Yapp *et al.*, 1990) and other Ru^{II}—S bond distances (Alessio *et al.*, 1990, 1991; Wang *et al.*, 1999). The Ru—S bond length depends upon the oxidation state and on the nature of coordinating *trans* ligands. The decrease in Ru—S bond length from Ru^{II} to Ru^{III} is partially due to the π -back-bonding contribution in the Ru—TMSO bond (Calligaris *et al.*, 1996; Srivastava *et al.*, 2004; Iengo *et al.*, 1999). The S—O bond length, however, is shorter than in crystalline TMSO (1.527 Å; Dodge *et al.*, 1972; Jaswal *et al.*, 1990), suggesting a significant S—O π -bonding, resulting from S-to-metal electron-density transfer (Davies, 1981; Jaswal *et al.*, 1990).

The solvent molecule is ordered, with statistically indistinguishable C—Cl distances.

Experimental

To a homogeneous solution of *mer*-[RuCl₃(bpy)(TMSO)] (0.23 mmol) in ClCH₂CH₂Cl (35 ml) was added solid *N,N*-diethyl-4-

nitrosoaniline (0.25 mmol), and the resulting solution was stirred at room temperature for 25 h. The solvent was removed under reduced pressure, triturated with diethyl ether (4 × 5 ml) and vacuum dried. On recrystallization from a CH₂Cl₂/hexane mixture (5:1 *v/v*), the title solvate was obtained as an orange crystalline solid.

Crystal data

[RuCl₃(C₁₀H₈N₂)(C₄H₈OS)]·
CH₂Cl₂
M_r = 552.69
Monoclinic, *P*2₁/*c*
a = 12.804 (2) Å
b = 10.139 (2) Å
c = 15.734 (2) Å
 β = 96.703 (11)°
V = 2028.6 (6) Å³
Z = 4

D_x = 1.810 Mg m⁻³
Mo *K*α radiation
Cell parameters from 8203
reflections
 θ = 2.5–33.7°
 μ = 1.54 mm⁻¹
T = 110 K
Prism, orange
0.15 × 0.12 × 0.10 mm

Data collection

Nonius KappaCCD diffractometer
(with an Oxford Cryosystems
Cryostream cooler)
 ω scans
Absorption correction: multi-scan
HKL SCALEPACK
(Otwinowski & Minor 1997)
T_{min} = 0.796, *T_{max}* = 0.857

56386 measured reflections
8048 independent reflections
5695 reflections with *I* > 2σ(*I*)
R_{int} = 0.044
 θ_{\max} = 33.7°
h = -19 → 19
k = -15 → 15
l = -24 → 23

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.041
wR (*F*²) = 0.086
S = 1.06
8048 reflections
226 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0297P)^2 + 1.6323P]$
where $P = (F_o^2 + 2F_c^2)/3$
(Δ/σ)_{max} = 0.001
 $\Delta\rho_{\max}$ = 2.23 e Å⁻³
 $\Delta\rho_{\min}$ = -1.01 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Ru1—N1	2.077 (2)	Ru1—Cl1	2.3649 (8)
Ru1—N2	2.086 (2)	S1—O1	1.4781 (19)
Ru1—S1	2.3042 (7)	Cl4—C15	1.769 (3)
Ru1—Cl3	2.3324 (7)	Cl5—C15	1.768 (3)
Ru1—Cl2	2.3414 (7)		
N1—Ru1—N2	78.49 (8)	Cl2—Ru1—Cl1	92.27 (2)
Cl3—Ru1—Cl1	93.49 (2)		
N1—Ru1—S1—O1	-3.66 (12)		

H atoms were placed in idealized positions, with C—H bond distances in the range 0.95–0.99 Å, and thereafter treated as riding. Displacement parameters were assigned as *U*_{iso}(H) = 1.2*U*_{eq} of the attached atom. The largest residual difference-map peak is 0.73 Å from the Ru atom and the deepest hole is 0.71 Å from the Ru atom.

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

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