

[(*R,R*)-2-Amino-1-(*p*-tolylsulfonylamido)cyclohexane- κ^2 N,*N'*]chloro(η^5 -pentamethylcyclopentadienyl)iridium(III) chloroform solvate

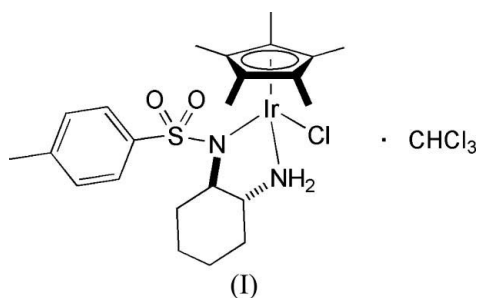
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The enantiomerically pure iridium title complex, $[\text{Ir}(\text{C}_{13}\text{H}_{19}\text{N}_2\text{O}_2\text{S})(\text{C}_{10}\text{H}_{15})\text{Cl}]\cdot\text{CHCl}_3$, has the Ir atom in the *S* configuration, with a distorted octahedral geometry.

Comment

Recently, we reported an asymmetric transfer hydrogenation reaction in aqueous solution using η^6 -arene ruthenium *N*-tosyl-*trans*-1,2-diaminocyclohexane complexes as catalysts (Canivet *et al.*, 2005). In order to extend our studies to iridium complexes, we synthesized the $[\text{Cp}^*\text{IrCl}\{(R,R)\text{-}L\}]$ complex, (I) [Cp^* = pentamethylcyclopentadienyl and *LH* = (*R,R*)-*N*-(*p*-toluenesulfonyl)-1,2-cyclohexanediamine], which crystallized from chloroform to give a polymorph of the known complex described by Murata *et al.* (1999). The bond distances and angles are similar to those found in the parent compound $[\text{Cp}^*\text{IrCl}\{(R,R)\text{-}L\}]$ (Murata *et al.*, 1999), the rhodium analogue $[\text{Cp}^*\text{RhCl}\{(R,R)\text{-}L\}]$ (Murata *et al.*, 1999) and the nitromethyl derivative $[\text{Cp}^*\text{Ir}(\text{nitromethyl})\{(R,R)\text{-}L\}]$ (Murata *et al.*, 2002).



The single-crystal X-ray structure analysis of (I) reveals a distorted octahedral geometry for the Ir atom with Cp*, amino, sulfonamido and chloro ligands. Upon formation of complex (I) the iridium metal center becomes chiral. The iridium possesses the *S* configuration as observed in the ruthenium analogue $[(\eta^6\text{-mesitylene})\text{RuCl}\{(R,R)\text{-}L'\}]$ [*L'H* = (*R,R*)-*N*-(*p*-toluenesulfonyl)-1,2-diphenylethylenediamine] (Uematsu *et al.*, 1996) and $[\text{Cp}^*\text{IrCl}\{(R,R)\text{-}L\}]$.

In contrast with the parent compound which was crystallized from water, (I) was obtained in the presence of chloroform, thus giving rise to a completely different crystal packing. Indeed, two intermolecular hydrogen-bonded systems are observed: an (amino) N—H...O—S (sulfonamido) hydrogen bonding between two symmetry-related molecules and a (chloro) Ir—Cl...H—C (chloroform) hydrogen bonding between complex (I) and the solvent molecule (see Table 1).

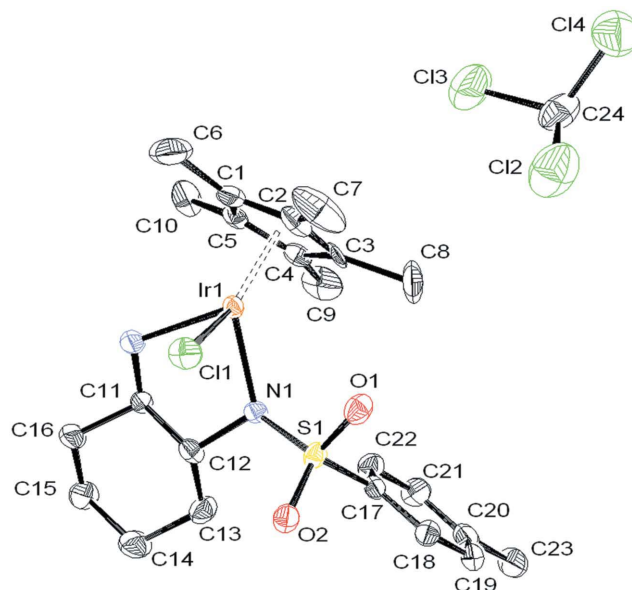


Figure 1

The asymmetric unit of (I), showing 50% probability displacement ellipsoids (H atoms omitted for clarity).

Experimental

$[\text{Cp}^*\text{IrCl}\{(R,R)\text{-}L\}]$ was synthesized according to the published method (Murata *et al.*, 1999). The complex was dissolved in chloroform at room temperature. Yellow crystals of (I) suitable for X-ray diffraction analysis were obtained after three days by slow evaporation of the solvent.

Crystal data

$[\text{Ir}(\text{C}_{13}\text{H}_{19}\text{N}_2\text{O}_2\text{S})(\text{C}_{10}\text{H}_{15})\text{Cl}]\cdot\text{CHCl}_3$
 $M_r = 749.60$
Orthorhombic, $P2_12_12_1$
 $a = 9.4012$ (5) Å
 $b = 12.8491$ (7) Å
 $c = 23.7478$ (17) Å
 $V = 2868.7$ (3) Å³

$Z = 4$
 $D_x = 1.736$ Mg m⁻³
Mo $K\alpha$ radiation
 $\mu = 5.12$ mm⁻¹
 $T = 173$ (2) K
Needle, yellow
 $0.45 \times 0.15 \times 0.08$ mm

Data collection

Stoe IPDS diffractometer
 φ scans
Absorption correction: multi-scan (*MULABS* in *PLATON*; Spek, 2003)
 $T_{\min} = 0.360$, $T_{\max} = 0.557$
(expected range = 0.428–0.664)

15012 measured reflections
5576 independent reflections
4830 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.044$
 $\theta_{\text{max}} = 25.9^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.028$
 $wR(F^2) = 0.059$
 $S = 0.94$
 5576 reflections
 313 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0306P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 1.53 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -1.82 \text{ e } \text{Å}^{-3}$
 Absolute structure: Flack (1983),
 2413 Friedel pairs
 Flack parameter: 0.024 (7)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C24—H24A \cdots C11	1.00	2.36	3.357 (8)	178
N2—H2A \cdots C11	0.92	2.60	2.996 (5)	106
N2—H2B \cdots O2 ⁱ	0.92	2.26	3.063 (7)	145

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

All H atoms were included in calculated positions (C—H = 0.95 Å for CH_{arom}, 1.00 Å for CH, 0.99 Å for CH₂ and 0.98 Å for CH₃; N—H = 0.92 Å) and treated as riding atoms with the constraint $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{carrier})$ applied. The highest residual electron density and deepest hole are both located at less than 1.1 Å from the Ir atom.

Data collection: *EXPOSE* in *IPDS Software* (Stoe & Cie, 2000); cell refinement: *CELL* in *IPDS Software*; data reduction: *INTEGRATE* in *IPDS Software*; 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: *SHELXL97*.

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