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Key indicators

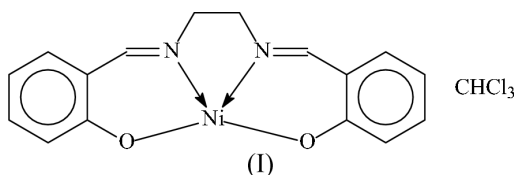
Single-crystal X-ray study
 $T = 295$ K
Mean $\sigma(\text{C}-\text{C}) = 0.006$ Å
 R factor = 0.046
 wR factor = 0.147
Data-to-parameter ratio = 18.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**[*N,N'*-Ethylenebis(salicylideneiminato)]nickel(II)
chloroform solvate**In the crystal structure of the title compound, $[\text{Cu}(\text{C}_8\text{H}_7\text{NO})_2]\cdot\text{CHCl}_3$, the nickel complex and the solvent molecule both have crystallographic mirror symmetry. The Ni atom exists in square-planar geometry.

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Comment

The crystal structure of [*N,N'*-ethylenebis(salicylideneiminato)]nickel(II) was first reported in 1970 (Shkol'nikova *et al.*, 1970) and redetermined to improved precision some 13 years later (Manfredotti & Guastini, 1983). It has since been re-determined twice more (DiMauro & Kozlowski, 2002; Kondo *et al.*, 2003). The same complex has now been isolated as the chloroform solvate, (I) (Fig. 1), and is described here.

In the title solvate, the square-planar geometry of the Ni atom is almost the same as in the unsolvated structure. Atoms Ni1, C9, and Cl1 occupy special positions on a mirror plane.

The CHCl_3 molecule interacts with the nickel complex by way of a bifurcated $\text{C}-\text{H}\cdots(\text{O},\text{O}')$ bond (Fig. 1, Table 2).

Experimental

Nickel nitrate hexahydrate (0.58 g, 2 mmol) and an excess of triethylamine (1 ml) were added to *N,N'*-ethylenebis(salicylideneimine) (0.54 g, 2 mmol) dissolved in a small volume of ethanol. The mixture was heated for 1 h. After removal of the solvent, a red solid was collected, and this was purified by recrystallization from chloroform. Red prismatic crystals of (I) were obtained. CHN elemental analysis, calculated for $\text{C}_{17}\text{H}_{15}\text{N}_2\text{O}_2\text{Cl}_3\text{Ni}$: C 45.95, H 3.40, N 6.30%; found: C 45.91, H 3.43, N 6.28%.

Crystal data

$[\text{Cu}(\text{C}_8\text{H}_7\text{NO})_2]\cdot\text{CHCl}_3$
 $M_r = 444.37$
 Orthorhombic, $Pnmm$
 $a = 6.997$ (1) Å
 $b = 14.221$ (3) Å
 $c = 18.355$ (4) Å
 $V = 1826.4$ (6) Å³
 $Z = 4$
 $D_x = 1.616$ Mg m⁻³

Mo $K\alpha$ radiation
 Cell parameters from 14 841
 reflections
 $\theta = 3.1\text{--}27.5^\circ$
 $\mu = 1.51$ mm⁻¹
 $T = 295$ (2) K
 Prism, red
 $0.35 \times 0.26 \times 0.18$ mm

Data collection

Rigaku R-AXIS RAPID
diffractometer
 ω scans
Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.507$, $T_{\max} = 0.764$
16 264 measured reflections

2150 independent reflections
1766 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -9 \rightarrow 8$
 $k = -18 \rightarrow 18$
 $l = -22 \rightarrow 23$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.147$
 $S = 1.07$
2150 reflections
118 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0952P)^2 + 0.4858P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.73 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.56 \text{ e } \text{\AA}^{-3}$
Extinction correction: none

Table 1

Selected geometric parameters (\AA , $^\circ$).

| | | | |
|------------------------|-----------|------------------------|-----------|
| Ni1—O1 | 1.844 (2) | Ni1—N1 | 1.843 (3) |
| O1—Ni1—O1 ⁱ | 84.6 (1) | O1—Ni1—N1 ⁱ | 178.9 (1) |
| O1—Ni1—N1 | 94.8 (1) | N1—Ni1—N1 ⁱ | 85.9 (2) |

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

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|--------------------------------|-------|-------------|-------------|---------------|
| C9—H9 \cdots O1 | 0.98 | 2.43 | 3.292 (5) | 147 |
| C9—H9 \cdots O1 ⁱ | 0.98 | 2.43 | 3.292 (5) | 147 |

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

H atoms were placed in calculated positions ($C-H = 0.93-0.98 \text{ \AA}$) and refined as riding, with the constraint $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ applied in all cases.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

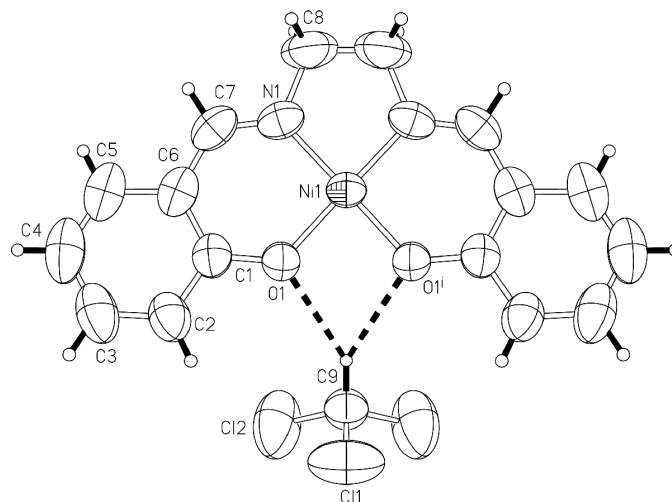


Figure 1

A view of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. C—H \cdots O interactions are shown as dashed lines. [Symmetry code (i): $x, y, 1 - z$.]

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