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Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$

R factor = 0.063

w R factor = 0.148

Data-to-parameter ratio = 16.1

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

trans-Bis{2-[3-(cyclohexylamino)propylimino-methyl]phenolato}nickel(II) dinitrate

The title compound, $[\text{Ni}(\text{C}_{16}\text{H}_{24}\text{N}_2\text{O})_2](\text{NO}_3)_2$, is a mononuclear nickel(II) compound. The Ni^{II} atom is coordinated by two N atoms and two O atoms from the Schiff base ligands. The four atoms around the metal are coplanar, constituting a slightly distorted square-planar geometry. The complex is located on an inversion center. All of the nitrate O atoms and all amine N atoms in the Schiff base ligands contribute to hydrogen bonds, leading to the formation of a three-dimensional network.

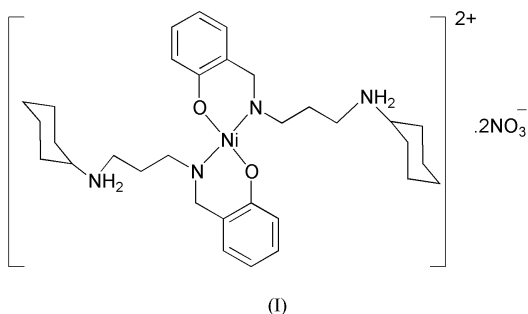
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Comment

Recently, we have reported a few Schiff base complexes (You, Lin *et al.*, 2003; You, Qu *et al.*, 2003; You, Xiong *et al.*, 2004; You, Zhu & Liu, 2004). As an extension of our work on the structural characterization of Schiff base complexes, a mononuclear nickel(II) complex, (I), is reported here.



The structure of the title compound, (I) (Fig. 1), consists of a mononuclear $[\text{Ni}(\text{C}_{16}\text{H}_{24}\text{N}_2\text{O})_2]^{2+}$ cation and two nitrate anions. The Ni atom, on an inversion center, is in a square-planar geometry and is four-coordinated by two N atoms and two O atoms from the Schiff base ligands. The four coordi-

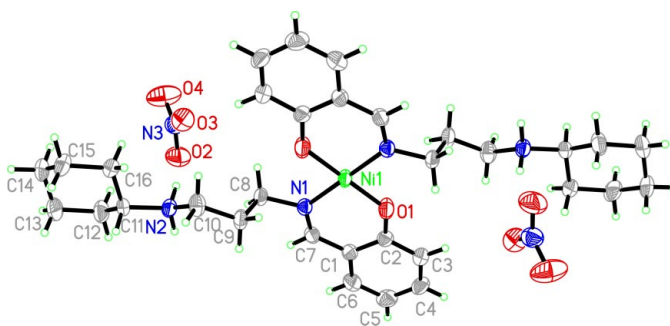


Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.

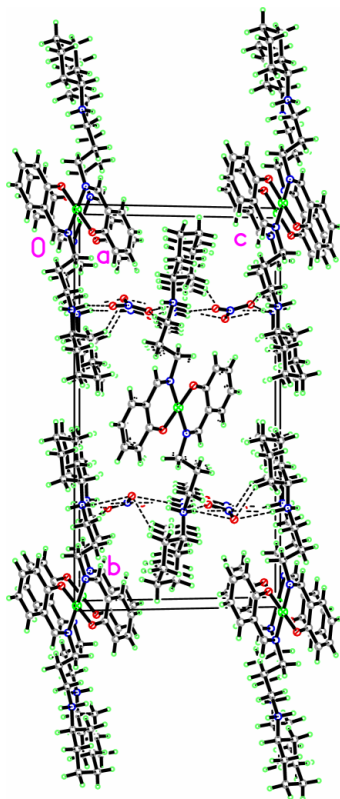


Figure 2
The crystal packing of (I), viewed along the *a* axis. Hydrogen bonds are shown as dashed lines.

nating atoms around the central metal are coplanar, forming a square-planar geometry around the Ni atom. The two *trans* angles at the nickel(II) center are 180° , by symmetry (Table 1), and all other angles are close to 90° , *viz.* $87.15(12)$ and $92.85(12)^\circ$, thus indicating a slightly distorted square-planar geometry. The Ni1—O1 bond length [$1.827(3)$ Å] is comparable to that observed in another Schiff base complex [$1.833(3)$ Å; Zhu *et al.*, 2004]. The Ni1—N1 bond distance [$1.919(3)$ Å] is a little longer than the value [$1.876(3)$ Å] observed in the same previously reported complex.

In the crystal structure, all of the nitrate O atoms and all the amine N atoms in the Schiff base ligands contribute to the formation of hydrogen bonds, leading to the formation of a three-dimensional network (Fig. 2 and Table 2). As expected, the cyclohexyl groups in the complex adopt chair conformations to minimize steric effects.

Experimental

N-Cyclohexyl-1,3-diaminopropane and salicylaldehyde were available commercially and were used without further purification. *N*-Cyclohexyl-1,3-diaminopropane (0.2 mmol, 31.2 mg) and salicylaldehyde (0.2 mmol, 24.4 mg) were dissolved in methanol (10 ml). The mixture was stirred for 1 h to obtain a clear orange solution of *L* (0.2 mmol), where *L* is 2-[(3-cyclohexylaminopropylimino)methyl]-phenol. To the solution of *L* was added a solution of $\text{Ni}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (0.1 mmol, 29.1 mg) in methanol (10 ml), with stirring. After keeping the resulting solution in air for 9 d, green block-shaped crystals were formed at the bottom of the vessel on slow evaporation of the

solvents. The crystals were isolated, washed three times with methanol and dried in a vacuum desiccator using anhydrous CaCl_2 (yield 70.1%). Analysis found: C 54.6, H 6.9, N 11.9%; calculated for $\text{C}_{32}\text{H}_{48}\text{N}_6\text{NiO}_8$: C 54.8, H 7.1, N 11.8%.

Crystal data

$[\text{Ni}(\text{C}_{16}\text{H}_{24}\text{N}_2\text{O}_2)](\text{NO}_3)_2$
 $M_r = 703.47$
 Monoclinic, $P2_1/c$
 $a = 6.444(2)$ Å
 $b = 22.946(5)$ Å
 $c = 11.961(2)$ Å
 $\beta = 102.02(3)^\circ$
 $V = 1729.9(6)$ Å³
 $Z = 2$

$D_x = 1.351$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 1144 reflections
 $\theta = 2.5\text{--}19.9^\circ$
 $\mu = 0.62$ mm⁻¹
 $T = 293(2)$ K
 Block, green
 $0.18 \times 0.15 \times 0.13$ mm

Data collection

Siemens SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Blessing, 1995)
 $T_{\min} = 0.897$, $T_{\max} = 0.924$
 7948 measured reflections

3573 independent reflections
 1985 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.063$
 $\theta_{\text{max}} = 26.5^\circ$
 $h = -8 \rightarrow 4$
 $k = -28 \rightarrow 27$
 $l = -14 \rightarrow 15$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.063$
 $wR(F^2) = 0.148$
 $S = 0.95$
 3573 reflections
 222 parameters

H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0601P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.48$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.24$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Ni1—O1	1.827 (3)	Ni1—N1	1.919 (3)
O1 ⁱ —Ni1—O1	180	O1—Ni1—N1	92.85 (12)
O1 ⁱ —Ni1—N1	87.15 (12)	N1—Ni1—N1 ⁱ	180

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

Table 2

Hydrogen-bonding 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>
N2—H2B...N3 ⁱⁱ	0.932 (10)	2.542 (12)	3.458 (6)	168 (3)
N2—H2B...O3 ⁱⁱ	0.932 (10)	2.36 (2)	3.125 (5)	139 (3)
N2—H2B...O4 ⁱⁱ	0.932 (10)	1.980 (15)	2.887 (5)	164 (3)
N2—H2A...N3	0.926 (10)	2.558 (15)	3.462 (6)	165 (3)
N2—H2A...O2	0.926 (10)	2.26 (2)	3.105 (5)	150 (3)
N2—H2A...O3	0.926 (10)	2.234 (17)	3.076 (5)	151 (3)

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

All H atoms, except H2A and H2B, were placed in idealized positions and constrained to ride on their parent atoms, with C—H distances of 0.93–0.98 Å, and with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$. Atoms H2A and H2B were located in a difference Fourier map and were refined isotropically.

Data collection: SMART (Siemens, 1996); cell refinement: SMART; data reduction: SAINT (Siemens, 1996); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics:

SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: *SHELXTL*.

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