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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.004 Å R factor = 0.036 wR factor = 0.111 Data-to-parameter ratio = 18.0

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title compound, $[Ni(C_{14}H_{10}N_3O_4)_2(C_5H_5N)_2]$, contains a Ni atom (site symmetry 2) six-coordinated by two *N*,*O*-bidentate Schiff base ligands and two pyridine N atoms, giving a distorted *trans*-NiN₄O₂ octahedral coordination environment for the metal atom.

Bis[(E)-N'-(2-nitrobenzylidene)-2-oxido-

benzohydrazide]bis(pyridine-*kN*)nickel(II)

Comment

Recently, we have reported some Schiff base complexes (Qiu *et al.*, 2004; Zhu *et al.*, 2003). As an extension of this work the synthesis and structure of the title nickel(II) complex, (I) (Fig. 1), is reported here. The Ni atom (site symmetry 2) in (I) is six-coordinated by two N atoms and two O atoms from two Schiff base ligands and two N atoms from two pyridine molecules. The Schiff base acts as a bidentate ligand and ligates to the Ni atom through an O atom and an N atom. The three *trans* angles at Ni are close to 180° . The *cis* bond angles range from 77.61 (5) to 101.97 (5)°, which indicates a slightly distorted octahedral geometry about Ni.



The Ni1-O2 bond length in (I) is comparable to the value of 2.038 (2) Å observed by us in a similar nickel(II) compound (You *et al.*, 2004). The Ni-N bond distances are a little longer than the value of 2.068 (3) Å observed in the nickel(II) complex cited above. The conformation of the five-membered chelate ring containing atoms Ni1, O2, C7, N1 and N2 is close to planar.

Experimental

A methanol (10 ml) and pyridine (4 ml) mixture of Ni(CH₃COO)₂·-6H₂O (0.5 mmol, 142.5 mg) was added to a methanol (10 ml) solution of HL (1.0 mmol, 285.0 mg), where HL is (E)-N'-(2-nitrobenzylidene)-2-hydroxybenzohydrazide. The reaction mixture was stirred for 20 min to give a green solution. After standing in air for 8 d, green blocks of (I) formed at the bottom of the vessel on slow evaporation of the solvent. The crystals were isolated, washed three times with

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Figure 1

The structure of (I), showing 30% probability displacement ellipsoids (arbitrary spheres for the H atoms). Atoms with the suffix A are generated by the symmetry operation $(\frac{3}{2} - x, 2 - y, z)$.

methanol and dried in a vacuum desiccator using anhydrous $CaCl_2$ (yield 63%).

Crystal data

 $\begin{bmatrix} \text{Ni}(\text{C}_{14}\text{H}_{10}\text{N}_{3}\text{O}_{4})_{2}(\text{C}_{5}\text{H}_{5}\text{N})_{2} \end{bmatrix} \qquad Z = 8 \\ M_{r} = 785.41 \qquad D_{x} = 1.406 \text{ Mg m}^{-3} \\ \text{Orthorhombic, } Ccca \qquad \text{Mo } K\alpha \text{ radiation} \\ a = 14.9148 (2) \text{ Å} \qquad \mu = 0.59 \text{ mm}^{-1} \\ b = 22.1842 (4) \text{ Å} \qquad T = 298 (2) \text{ K} \\ c = 22.4259 (4) \text{ Å} \qquad \text{Block, green} \\ V = 7420.1 (2) \text{ Å}^{3} \qquad 0.46 \times 0.24 \times 0.13 \text{ mm} \\ \end{bmatrix}$

Data collection

Bruker SMART APEX areadetector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{min} = 0.841, T_{max} = 0.925$ 22385 measured reflections 4484 independent reflections 3308 reflections with $I > 2\sigma(I)$

 $\begin{aligned} R_{\rm int} &= 0.019\\ \theta_{\rm max} &= 28.3^\circ \end{aligned}$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0557P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 3.9687P]
$wR(F^2) = 0.111$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} = 0.001$
4484 reflections	$\Delta \rho_{\rm max} = 0.37 \ {\rm e} \ {\rm A}^{-3}$
249 parameters	$\Delta \rho_{\rm min} = -0.43 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected bond lengths (Å).

Ni1-O2	2.0064 (11)	Ni1-N2	2.1416 (14)
Ni1-N4	2.1184 (15)		

All H atoms were placed in geometrically idealized positions (C– H = 0.93 and N–H = 0.86 Å) and refined as riding with $U_{iso}(H) = 1.2U_{eq}(carrier)$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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