metal-organic papers

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.030 wR factor = 0.064 Data-to-parameter ratio = 12.2

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

Bis{*N*-(2-hydroxybenzoyl)-*N*'-[1-(2-pyridyl)ethylidene]hydrazinato}nickel(II)

The title complex, $[Ni(C_{14}H_{12}N_3O_2)_2]$, consists of two deprotonated 2-acetylpyridine salicylhydrazone ligands coordinated to the nickel as N,N',O-tridentate ligands, forming a neutral complex. The Ni atom lies on a twofold rotation axis. The structure is stabilized by intramolecular $O-H\cdots N$ hydrogen bonding and $\pi-\pi$ stacking interactions between the pyridyl and benzene rings, leading to a two-dimensional network in the crystal structure.

Comment

Diazine Schiff base ligands with two coordination compartments linked by a single N-N bridge are employed for their ease of synthesis and high yield in a single-step reaction from commercial and inexpensive reagents (Grove et al., 2004; Bai et al., 2006). The most notable feature of this type of polydentate ligand is that the single N-N bridge of the diazine fragment can rotate freely (Xu et al., 2000; Zhao et al., 2004) so that it usually leads to a situation where a variety of different structural types, such as chains, metallomacrocycles, and metallocrowns are observed when the ligands combine with metals (Bai et al., 2005; Song et al., 2004). On the other hand, the self-assembly of transition metal with diazine ligands introduces many special functional properties, such as magnetism and redox activity, into the structure apart from their particular structural features (Leininger et al., 2000). In our previous work, a mononuclear copper(II) complex generated from the ligand 2-acetylpyridine salicylhydrazone (H₂L) was reported (Dang et al., 2006). Here, a new mononuclear complex of nickel(II) was constructed via selfassembly using the same ligand.



Reaction of NiCl₂· $6H_2O$ with the pentadentate Schiff base ligand 2-acetylpyridine salicylhydrazone (H₂L) leads to the formation of the title nickel(II) complex, (I), which consists of

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2171 independent reflections 1934 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.041$

 $\theta_{\rm max} = 25.0^{\circ}$



Figure 1

View of the molecular structure of the title complex, showing the atomlabelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms other than the hydroxyl proton have been omitted for clarity. Hydrogen bonds are indicated by dashed lines. [Symmetry code (i) -x + 2, -y + 1, z.]



Figure 2

Intermolecular π - π stacking (dashed lines) in the title complex, viewed along the *a* axis. H atoms have been omitted for clarity.

two HL⁻ ligands coordinated to the nickel as N,N',O-tridentate ligands, forming a neutral complex (Fig. 1). The structure is similar to that found for the related copper(II) complex (Dang et al., 2006). The Ni atom lies on a twofold rotation axis and is coordinated in a distorted octahedral configuration by two carbonyl O atoms, two imine N atoms and two pyridine N atoms. Just as has been found in the copper(II) complex, the ligands bond to the nickel in the mer configuration with pairs of carbonyl O atoms and pyridyl N atoms each bearing a cis relationship, whereas the imino N atoms are trans to each other. Intramolecular hydrogen bonds between imino N atoms and hydroxy groups stabilize the molecule. In the crystal structure, molecules are connected together by $\pi - \pi$ interactions between the benzene and pyridine rings, generating a two-dimensional network (Fig. 2). The dihedral angle and the centroid-entroid distance between the interacting pair are $10.8 (1)^{\circ}$ and 3.859 (1) Å, respectively.

Experimental

All chemicals were of reagent grade quality obtained from commercial sources and used without further purification. The ligand H_2L was synthesized according to the literature methods (Dang *et al.*, 2006). A solution of H_2L (0.025 g, 0.1 mmol) and Et_3N (0.011 g, 0.1 mmol) in methanol (6 ml) was layered onto a solution of DMF (6 ml) containing NiCl₂·6H₂O (0.024 g, 0.1 mmol) at the bottom. The solutions were allowed to stand for few weeks at room temperature in darkness, giving X-ray quality crystals (yield, 85%). Analysis calculated for C₂₈H₂₄N₆NiO₄: C 59.3, H 4.3, N 14.8%; found: C 59.3, H 4.4, N 14.2%.

Crystal data

$[Ni(C_{14}H_{12}N_3O_2)_2]$	Z = 4
$M_r = 567.24$	$D_x = 1.531 \text{ Mg m}^{-3}$
Orthorhombic, Aba2	Mo $K\alpha$ radiation
a = 12.495 (3) Å	$\mu = 0.84 \text{ mm}^{-1}$
b = 18.672 (4) Å	T = 293 (2) K
c = 10.549 (2) Å	Block, green
$V = 2461.2 (9) \text{ Å}^3$	$0.20 \times 0.18 \times 0.15 \text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer $\omega | \omega$ scans Absorption correction: none 5887 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.028P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_0^2 + 2F_c^2)/3$
$vR(F^2) = 0.064$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 1.01	$\Delta \rho_{\rm max} = 0.31 \ {\rm e} \ {\rm \AA}^{-3}$
2171 reflections	$\Delta \rho_{\rm min} = -0.16 \text{ e} \text{ Å}^{-3}$
78 parameters	Absolute structure: Flack (1983)
H-atom parameters constrained	1011 Friedel pairs
	Flack parameter: -0.033 (16)

Table 1

Selected geometric parameters (Å, °).

Ni1-N2 Ni1-N1	1.9830 (19) 2.081 (2)	Ni1-01	2.0977 (19)
$\begin{array}{l} N2 - Ni1 - N2^{i} \\ N2 - Ni1 - N1^{i} \\ N2 - Ni1 - N1 \\ N1^{i} - Ni1 - N1 \\ N2 - Ni1 - O1 \end{array}$	173.24 (16) 107.31 (9) 77.65 (10) 89.53 (12) 77.08 (8)	N1-Ni1-O1 $N2-Ni1-O1^{i}$ $N1-Ni1-O1^{i}$ $O1-Ni1-O1^{i}$	154.31 (7) 98.26 (8) 93.59 (8) 94.52 (11)

Symmetry code: (i) -x + 2, -y + 1, z.

Table 2 Hydrogen-bond geometry (Å, °).

	D II	11 4	D 4	
$D - H \cdots A$	D-H	H···A	$D \cdots A$	$D - H \cdots A$
$O2-H2B\cdots N3$	0.82	1.82	2.541 (3)	146

H atoms bonded to C and O atoms were constrained to ride on their parent atoms, with distances of 0.96 (CH₃), 0.93 (aromatic) and 0.82 Å (OH), and $U_{iso}(H) = 1.5U_{eq}(C)$ for CH₃, $1.2U_{eq}(C)$ for aromatic and $1.2U_{eq}(O)$ for OH.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve structure: *SHELXTL* (Bruker, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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