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#### **Key indicators**

Single-crystal X-ray study T = 295 KMean  $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$  R factor = 0.038 wR factor = 0.107 Data-to-parameter ratio = 16.2

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

# Bis(2-methyl-1*H*-indole-3-carbaldehyde 4-hydroxybenzoylhydrazonato- $\kappa^2 N$ ,*O*)nickel(II) *N*,*N*-dimethylformamide disolvate dihydrate

The title compound,  $[Ni(C_{17}H_{14}N_3O_2)_2]\cdot 2C_3H_7NO\cdot 2H_2O$ , contains Ni<sup>II</sup> cations (site symmetry  $\overline{1}$ ) chelated by two deprotonated Schiff bases in a *trans*-NiN<sub>2</sub>O<sub>2</sub> square-planar geometry. The complex molecule and the dimethylformamide (DMF) and water molecules interact by way of  $O-H\cdots O$  and  $N-H\cdots O$  hydrogen bonds, resulting in a layered structure.

### Comment

This report continues on from our previous study on nickel derivatives of indole-3-carbaldehyde aroylhydrazone (Ali *et al.*, 2005). The title compound, (I), has the Ni atom (site symmetry  $\overline{1}$ ) in a square-planar geometry (Table 1 and Fig. 1). The dimethylformamide solvent used in recrystallization is incorporated into the crystal structure; however, it does not participate in bonding with the metal but interacts instead with the water molecule. Together, the constituent molecules interact by way of  $O-H\cdots O$  and  $N-H\cdots O$  hydrogen bonds (Table 2), resulting in a sheet structure.



## Experimental

The Schiff base was synthesized from 2-methylindole-3-carbaldehyde (0.44 g, 2.7 mmol) and 4-hydroxybenzhydrazide (0.42 g, 2.7 mmol) in ethanol (100 ml). The reactants were heated under reflux for 2 h. The solvent was removed to give the organic compound. This compound (0.18 g, 0.6 mmol) and nickel(II) acetate dihydrate (0.08 g, 0.3 mmol) were heated in ethanol (100 ml) for 5 h. The complex was obtained as a red powder which was recrystallized from dimethylformamide, yielding crystals of (I).

#### Crystal data

 $[Ni(C_{17}H_{14}N_3O_2)_2] \cdot 2C_3H_7NO \cdot 2H_2O$ V = 1034.5 (2) Å<sup>3</sup>  $M_r = 825.56$ Z = 1Triclinic,  $P\overline{1}$  $D_{\rm x} = 1.325 {\rm Mg m}^{-3}$ a = 7.2441 (6) Å Mo  $K\alpha$  radiation b = 12.326 (1) Å  $\mu = 0.53 \text{ mm}^{-1}$ T = 295 (2) K c = 12.395(1) Å  $\alpha = 76.336 \ (1)^{\circ}$ Irregular block, red  $\beta = 74.120(1)^{\circ}$  $0.45 \times 0.35 \times 0.15$  mm  $\gamma = 86.253 (1)^{\circ}$ 

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# metal-organic papers

### Data collection

Bruker APEX-II CCD diffractometer  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\min} = 0.801, T_{\max} = 0.925$ 

### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.038$   $wR(F^2) = 0.107$  S = 1.024495 reflections 278 parameters H atoms treated by a mixture of independent and constrained refinement 9651 measured reflections 4495 independent reflections 3822 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.017$  $\theta_{\text{max}} = 27.1^{\circ}$ 

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0595P)^2 \\ &+ 0.185P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\rm max} = 0.001 \\ \Delta\rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.20 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

## Table 1

Selected geometric parameters (Å, °).

O1-Ni1-N2	83.52 (6)	O1-Ni1-N2 <sup>i</sup>	96.48 (6)
Ni1-O1	1.833 (1)	Ni1-N2	1.862 (1)

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

Table 2

Hydrogen-bond	geometry	(A, '	°).
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O2−H2o···O1w <sup>ii</sup>	0.85(1)	1.85 (1)	2.688 (2)	171 (3)
$O1w-H1w1\cdots O1$	0.84(1)	2.02(1)	2.852 (2)	173 (3)
$O1w - H1w2 \cdots O3$	0.84 (1)	1.94 (1)	2.754 (2)	163 (3)
$N3-H3n\cdots O3^{iii}$	0.86 (1)	2.00 (1)	2.859 (2)	173 (2)

Symmetry codes: (ii) -x + 2, -y + 1, -z; (iii) x, y + 1, z.

The carbon-bound H atoms were placed at calculated positions (C-H = 0.93-0.96 Å) and refined as riding, with  $U_{iso}(H) = 1.2$  or 1.5 times  $U_{eq}(C)$ . The methyl group was rotated to best fit the electron density. The water, hydroxy and amino H atoms were located in a difference Fourier map and were refined with distance restraints [O-H = N-H = 0.85 (1) Å]; the  $U_{iso}$  values were freely refined.



### Figure 1

The molecular structure of (I) with displacement ellipsoids drawn at the 50% probability level (arbitrary spheres for the H atoms). The dashed lines represent hydrogen bonds. [Symmetry code: (i) 1 - x, 1 - y, 1 - z.]

Data collection: *APEXII* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97*.

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