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Sheng-Gui Liu, Yi-Zhi Li,* Jing-Lin Zuo* and Xiao-Zeng You

Coordination Chemistry Institute, State Key Laboratory of Coordination Chemistry, Nanjing University, Nanjing 210093, People's Republic of China

Correspondence e-mail: llyyjz@nju.edu.cn

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.005 Å R factor = 0.052 wR factor = 0.132 Data-to-parameter ratio = 14.5

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

{*N*,*N*'-Bis[2-(methylsulfanyl)phenyl]pyridine-2,6-dicarboxamidato(2–)}nickel(II)

In the molecule of the title complex, $[Ni(C_{21}H_{17}N_3O_2S_2)]$, the Ni atom is coordinated by three N atoms and one S atom of the pyridine-2,6-dicarboxamide ligand. The geometry around the Ni^{II} atom is approximately square planar. The crystal structure is stabilized by weak π - π and C-H···O intermolecular interactions.

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Comment

Nickel complexes with *N*,*S*-donor chelate ligands have received much attention in recent years. As well as being employed as models of metalloenzymes such as [NiFe]hydrogenases (Marganian *et al.*, 1995), some of the complexes have been found to have pharmacological and catalytic properties (Brückner *et al.*, 2000; Harrop *et al.*, 2003). In order to develop further the coordination chemistry of nickel complexes with *N*,*S*-donor chelate ligands, we have synthesized a nickel complex based on the ligand *N*,*N'*-bis[2-(methylsulfanyl)phenyl]pyridine-2,6-dicarboxamide, L^{2-} , which contains deprotonated carboxamide nitrogen and sulfur as donor atoms.



The title complex, (I), was obtained from reaction of L^{2-} and NiCl₂ in DMF solution in high yield. The absence of ν (NH) (3170 cm⁻¹) in the IR spectrum confirms that the ligand is coordinated to the Ni atom in the deprotonated form. The C=O stretching vibration for the complex is at 1635 cm⁻¹, lower than that in the free ligand H₂L (1676 cm⁻¹). Such behavior is also observed in another deprotonated amide complex (Patra *et al.*, 2000). Compared to the UV spectrum of the free ligand, there is a new absorption band at 342 nm for the complex, which is caused by a ligand-to-metal charge transfer transition.

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Figure 1

The molecular structure of (I), with anisotropic displacement ellipsoids drawn at the 30% probability level.



Figure 2

The packing of the complex. The dashed lines show weak π - π and C- $H \cdots O$ interactions, as detailed in Table 2 and the main text.

The molecular structure of the title complex is shown in Fig. 1. It is a mononuclear complex. The geometry around the central Ni atom is approximately square planar (r.m.s. deviation = 0.022 Å). Three of the coordination sites are occupied by N atoms from two amidate moieties and one pyridine ring; the fourth donor is an S atom. The C1-C6 benzene ring, the central pyridine ring and the two carboxamide groups are coplanar with the nickel coordination plane, which makes a dihedral angle of 75.74 (8)° with the other benzene ring, C14-C19. The average Ni $-N_{amide}$ distance [1.8685 (2) Å] is appreciably longer than that for Ni-N_{py} [py is pyridine; 1.821 (2) Å], because of the steric requirement of the rigid ligand. This behavior has also been observed in an NiN₃O square-planar complex (Kawamoto et al., 1998). The Ni1-S1 bond distance is shorter than that observed in another complex [2.1988 (9) and 2.1740 (9) Å; Brückner et al., 2000]. The bond lengths for C7–O1 and C13–O2 are only slightly shorter than those reported for a C=O bond [1.234 (6) and 1.242 (6) Å; Redmore et al., 1997]. Subtle changes within a ligand framework may result in entirely different structures when coordinated to a metal center. An NiN₆ compressed octahedral coordination for 2,6-bis(N-phenylcarbamoyl)pyridine has been reported (Patra & Mukherjee, 1999).

The crystal packing of the title complex is shown in Fig. 2. Two adjacent molecules form a dimer in the head-to-head fashion by two weak interactions; one is an intermolecular π - π interaction between the pyridine ring and the C1–C6 benzene ring [the distance $Cg1\cdots Cg2^{i}$ is 3.691 (16) Å; symmetry code: (i) -x, 2 - y, -z; Cg1 is the center of the benzene ring and Cg2 is the center of the pyridine ring], and the other is an intermolecular weak interaction C15-H15...O1ⁱ (Table 2). In the dimer, the Ni1...Ni1ⁱ distance is 4.2943 (11) Å. In the crystal structure, these dimers form a two-dimensional layer through two further weak interactions (see Table 2).

Experimental

All the chemicals were of AR grade reagent. CH2Cl2 and DMF were dried over CaH₂ and distilled before use. Pyridine-2,6-dicarboxylic acid (3.34 g, 20 mmol) was refluxed in 10 ml thionyl chloride for 8 h. Excess thionyl chloride was removed under vacuum. After cooling to room temperature, 2-(methylthio)aniline (5.56 g, 40 mmol) and triethylamine (3 ml) in CH₂Cl₂ (40 ml) were added to the solution of the residue in dry CH₂Cl₂ (30 ml). After the mixture had been stirred for 2 h at ambient temperature, the precipitated white solid was collected by filtration, washed with water and dried in vacuo. The product was recrystallized from ethanol to give the ligand H_2L (yield: 7.40 g, 90.5%). Analysis calculated for C₂₁H₁₉N₃O₂S₂: C 61.54, H 4.64, N 10.26%; found: C 61.62, H 4.63, N 10.29%. IR (KBr pellets, cm⁻¹): 3450, 3170, 1676, 1655, 1585, 1533, 1458, 1257, 1073, 768.

The ligand H_2L (257 mg, 0.63 mmol) and NaH (30 mg, 1.25 mmol) in 10 ml N,N-dimethylformamide (DMF) were added to a solution of NiCl₂·6H₂O (74 mg, 0.315 mmol) in DMF (5 ml) with stirring. The resulting red solution was stirred for 15 min at room temperature. Exposure of the solution to the air for a week gave red single crystals suitable for structure determination (yield: 180 mg, 69.20%). Analysis calculated for C₂₁H₁₉N₃NiO₂S₂: C 54.05, H 3.65, N 9.00%; found: C 54.13, H 3.63, N 9.04%. IR: (KBr pellets, cm⁻¹): 3445, 1635, 1578, 1466, 1360, 750. UV (DMF): 342 nm.

Crystal data

$[Ni(C_{21}H_{17}N_3O_2S_2)]$	$D_x = 1.585 \text{ Mg m}^{-3}$
$M_r = 466.21$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 775
a = 11.430(2) Å	reflections
b = 11.406 (2) Å	$\theta = 2.5 - 25.0^{\circ}$
c = 15.087 (3) Å	$\mu = 1.23 \text{ mm}^{-1}$
$\beta = 96.765 \ (4)^{\circ}$	T = 293 (2) K
V = 1953.2 (6) Å ³	Prism, red
Z = 4	$0.32 \times 0.22 \times 0.20 \text{ mm}$
Data collection	
Bruker SMART APEX CCD area-	3834 independent reflections
detector diffractometer	3039 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.066$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.0^{\circ}$
(SADABS; Bruker, 2000)	$h = -8 \rightarrow 14$
$T_{\min} = 0.669, \ T_{\max} = 0.782$	$k = -14 \rightarrow 13$
10243 measured reflections	$l = -18 \rightarrow 18$

Refinement

Refinement on F^2	$w = 1/(\sigma^2(F_o^2) + (0.07P)^2)$
$R[F^2 > 2\sigma(F^2)] = 0.052$	+ 1.55P)
$wR(F^2) = 0.132$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.92	$(\Delta/\sigma)_{\rm max} < 0.001$
3834 reflections	$\Delta \rho_{\rm max} = 0.87 \ {\rm e} \ {\rm \AA}^{-3}$
264 parameters	$\Delta \rho_{\rm min} = -0.83 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

Ni1-N2	1.821 (3)	Ni1-N3	1.876 (3)
Ni1-N1	1.861 (3)	Ni1-S1	2.1535 (10)
N2-Ni1-N1	83.88 (13)	N2-Ni1-S1	173.58 (10)
N2-Ni1-N3	83.42 (13)	N1-Ni1-S1	89.88 (10)
N1-Ni1-N3	167.19 (13)	N3-Ni1-S1	102.85 (10)

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot \cdot \cdot A$
C15-H15···O1 ⁱ	0.93	2.36	3.272 (6)	168
$C11-H11\cdots O2^{ii}$	0.93	2.77	3.134 (5)	105
$\rm C18{-}H18{\cdot}{\cdot}\rm O2^{iii}$	0.93	2.32	3.235 (5)	167
Symmetry codes: (i)	x, 2 - y, -z; (ii	(1 - x, 2 - y, -	-z; (iii) $1 - x, y - z$	$\frac{1}{2}, \frac{1}{2} - z.$

H atoms were positioned geometrically and constrained as riding atoms, with C–H distances of 0.93–0.96 Å and $U_{\rm iso}({\rm H})$ set to 1.2–1.5 $U_{\rm eq}$ of the parent atom.

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* and *MERCURY* (Version 1.2.1; Bruno *et al.*, 2002); software used to prepare material for publication: *SHELXTL*.

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References

Brückner, C., Retting, S. J. & Dolphin, D. (2000). Inorg. Chem. 39, 6100-6106.

- Bruker (2000). *SMART* (Version 5.625), *SAINT* (Version 6.01), *SHELXTL* (Version 6.10) and *SADABS* (Version 2.03). Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruno, I. J., Cole, J. C., Edgington, P. R., Kessler, M., Macrae, C. F., McCabe, P., Pearson, J. & Taylor, R. (2002). Acta Cryst. B58, 389–397.
- Harrop, T. C., Olmstead, M. M. & Mascharak, P. K. (2003). *Chem. Commun.* pp. 410–411.
- Kawamoto, T., Hamme, B. S., Ostrander, R., Rheingold, A. L. & Borovik, A. S. (1998). Inorg. Chem. 37, 3424–3427.
- Marganian, C. A., Vazir, H, Baidya, N., Olmstead, M. M. & Mascharak, P. K. (1995). J. Am. Chem. Soc. 117, 1584–1594.
- Patra, A. K. & Mukherjee, R. (1999). Inorg. Chem. 38, 1388-1393.
- Patra, A. K., Ray, M. & Mukherjee, R. (2000). Inorg. Chem. 39, 652-657.
- Redmore, S. M., Rickard, C. E. F., Webb, S. J. & Wright, L. J. (1997). Inorg. Chem. 36, 4743–4748.