metal-organic papers

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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.007 Å Disorder in solvent or counterion R factor = 0.064 wR factor = 0.180 Data-to-parameter ratio = 15.0

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

Bis{2-[2-(diethylammonio)ethyliminomethyl]-4-nitrophenolato}dimethanolnickel(II) dinitrate

In the title mononuclear nickel(II) compound, [Ni(C₁₃H₁₉- $N_3O_3)_2(CH_4O)_2](NO_3)_2$, the Ni^{II} atom is coordinated by two N and four O atoms, giving an octahedral geometry. The cationic complex lies on a centre of symmetry. In the crystal structure, the molecules are linked through intermolecular $O-H\cdots O$, $N-H\cdots N$ and $N-H\cdots O$ hydrogen bonds, forming chains running along the b axis.

Comment

Nickel(II) complexes 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 complexes have been found to have pharmacological and catalytic properties (Brückner et al., 2000; Harrop et al., 2003). As part of our investigations of non-covalent interactions in metal complexes (Chen, 2005), the new title Ni^{II} complex, (I), has been prepared and its crystal structure is presented here.



The structure of (I) is shown in Fig. 1. The complex consists of an $[Ni(C_{13}H_{19}N_3O_3)_2(CH_3OH)_2]^{2+}$ cation, which possesses a crystallographically imposed centre of symmetry, and two disordered nitrate anions. The coordination sites are occupied by four donor atoms from two Schiff base ligands and two O atoms from two methanol molecules, giving a slightly distorted octahedral geometry. All the bond lengths (Table 1) around the metal centre are comparable with those of similar compounds (Gomes et al., 2000; Arici et al., 2005; Jircitano et al., 1990).

In the crystal structure, the molecules are linked through intermolecular O-H···O, N-H···N and N-H···O hydrogen bonds (Table 2), forming chains running along the b axis (Fig. 2).

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H atoms treated by a mixture of

 $w = 1/[\sigma^2(F_0^2) + (0.1069P)^2]$ where $P = (F_0^2 + 2F_c^2)/3$

independent and constrained

refinement

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 1.18 \text{ e } \text{\AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.59 \ {\rm e} \ {\rm \AA}^{-3}$



Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level. Only the major components of the disordered nitrate anions are shown. Unlabelled atoms are at the symmetry position (-x, 2 - y, 2 - z).



Figure 2

The molecular packing of (I). Intermolecular hydrogen bonds are shown as dashed lines. H atoms and anions have been omitted.

Experimental

All chemicals were of AR grade. 5-Nitro-2-hydroxybenzaldehyde (33.5 mg, 0.2 mmol), N,N-diethylethane-1,2-diamine (23.2 mg, 0.2 mmol) and Ni(NO₃)₂·6H₂O (29.1 mg, 0.1 mmol) were refluxed in methanol (30 ml) for 30 min at 338 K. The mixture was then cooled to room temperature and filtered. After keeping the filtrate in air for 15 d, green block crystals of (I) suitable for X-ray analysis were obtained.

Crystal data

[Ni(C ₁₃ H ₁₉ N ₃ O ₃) ₂ (CH ₄ O) ₂](NO ₃) ₂	Z = 1
$M_r = 777.44$	$D_x = 1.434 \text{ Mg m}^{-3}$
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation
a = 8.351 (1) Å	Cell parameters from 2165
b = 11.000 (1) Å	reflections
c = 11.097 (1) Å	$\theta = 2.5-24.3^{\circ}$
$\alpha = 69.092 \ (2)^{\circ}$	$\mu = 0.62 \text{ mm}^{-1}$
$\beta = 70.963 \ (2)^{\circ}$	T = 298 (2) K
$\gamma = 84.048 \ (2)^{\circ}$	Block, green
$V = 900.09 (16) \text{ Å}^3$	$0.12 \times 0.11 \times 0.10 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-	3889 independent reflections
detector diffractometer	2804 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.033$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(SADABS; Bruker, 2000)	$h = -10 \rightarrow 10$
$T_{\min} = 0.930, \ T_{\max} = 0.941$	$k = -13 \rightarrow 14$
7414 measured reflections	$l = -14 \rightarrow 14$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.064$ wR(F²) = 0.180 S=1.003889 reflections 260 parameters

Table 1

Selected geometric parameters (Å, °).

Ni1-O1	1.999 (3)	Ni1-N1	2.066 (3)
Ni1-O1 ⁱ	1.999 (3)	Ni1-O2	2.120 (3)
Ni1-N1 ⁱ	2.066 (3)	Ni1-O2 ⁱ	2.120 (3)
$01 - Ni1 - 01^i$	180	Ω^{1i} -Ni1- Ω^{2}	91.87 (13)
$O1-Ni1-N1^{i}$	90.79 (11)	N1 - Ni1 - O2	89.96 (12)
O1-Ni1-N1	89.21 (11)	N1-Ni1-O2 ⁱ	90.04 (12)
N1 ⁱ -Ni1-N1	180	O2-Ni1-O2 ⁱ	180
O1-Ni1-O2	88.13 (13)		

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

Table 2 Hydrogen-bond geometry (Å, $^\circ).$

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O2−H2···N3 ⁱⁱ	0.85 (4)	2.58 (4)	3.416 (5)	170 (5)
O2−H2···O4 ⁱⁱ	0.85 (4)	2.31 (3)	3.039 (4)	145 (4)
O2−H2···O3 ⁱⁱ	0.85 (4)	2.26 (3)	3.038 (5)	154 (5)
$N2-H2A\cdots N4^{iii}$	0.90(3)	2.625 (15)	3.511 (5)	169 (4)
$N2-H2A\cdots O7'^{iii}$	0.90 (3)	2.55 (3)	3.317 (11)	143 (4)
$N2-H2A\cdots O6^{iii}$	0.90(3)	2.53 (3)	3.321 (11)	147 (4)
$N2-H2A\cdots O7^{iii}$	0.90(3)	1.96 (2)	2.803 (10)	157 (4)
$N2-H2A\cdots O6'^{iii}$	0.90 (3)	1.875 (19)	2.749 (9)	164 (5)

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

Atoms H2 and H2A were located in a difference Fourier map and refined isotropically, with the O-H2 distance restrained to 0.84 (1) Å and the N-H2A distance restrained to 0.90 (1) Å. All other H atoms were placed in idealized positions and constrained to ride on their parent atoms, with C-H distances in the range 0.93-0.97 Å, and with $U_{iso}(H) = 1.2$ or $1.5U_{eq}(C)$. The O atoms of the nitrate anions were disordered over two distinct sites with occupancies of 0.536 (2) and 0.464 (2). The N–O and O···O distances in both disordered components were restrained to be equal. An unassigned maximum residual density was observed 1.15 Å from atom C12.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; 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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