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Key indicators

Single-crystal X-ray study T = 292 KMean $\sigma(C-C) = 0.006 \text{ Å}$ R factor = 0.051 wR factor = 0.141 Data-to-parameter ratio = 13.8

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

Triaqua(2,2'-bipyridine)maleatonickel(II) monohydrate

In the title complex, $[Ni(C_4H_2O_4)(C_{10}H_8N_2)_2(H_2O)_3]\cdot H_2O$, the Ni atom has a distorted NiO₄N₂ octahedral coordination, formed by one maleate dianion, one bipyridine and three water molecules. The crystal structure is stabilized by π - π stacking and hydrogen-bonding interactions.

Comment

Metal complexes with the maleate ligand have been studied extensively because of their wide applications (Marioka *et al.*, 1993; Ma *et al.*, 2003). Here we report the crystal structure of the title Ni^{II} complex, (I).



The molecular structure of (I) is shown in Fig. 1. The Ni atom has a distorted octahedral coordination, formed by one bipyridine ligand, three water molecules and one maleate dianion. The Ni-N bond distances and Ni-O bond distances (Table 1) are normal.

An extensive hydrogen-bonding network occurs in the crystal structure of (I) (Table 2). The centroid-to-centroid separation of 3.998 (2) Å suggests weak π - π stacking between neighbouring pyridine rings. These molecular interactions help to stabilize the crystal structure.

Experimental

 $Ni(NO_3)_2\cdot 6H_2O~(0.290~g,~1~mmol)$ was added slowly to an aqueous solution (15 ml) of maleic acid (0.116 g,~1~mmol). The mixture was stirred for 1 h at 353 K, then an ethanol solution (5 ml) of 2,2'-



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The molecular structure of (I), with 50% probability displacement ellipsoids (arbitrary spheres for H atoms).

bipyridine (0.156 g, 1 mmol) was added slowly with stirring. The solution was adjusted to pH = 7 with NaOH solution and stirred for 1.5 h at 353 K. The solution was then cooled to room temperature and filtered. Single crystals of (I) were obtained from the filtrate after two weeks.

 $D_r = 1.591 \text{ Mg m}^{-3}$

Cell parameters from 3123

 $0.40 \times 0.10 \times 0.10$ mm

Mo $K\alpha$ radiation

reflections

 $\theta=2.6{-}27.8^\circ$ $\mu = 1.20 \text{ mm}^{-1}$

T = 292 (2) K

Needle, green

Crvstal data

[Ni(C4H2O4)(C10H8N2)2- $(H_2O)_3$]·H2O $M_r = 401.01$ Monoclinic, $P2_1/c$ a = 9.4941 (19) Å b = 7.6031 (15) Å c = 23.408 (5) Å $\beta = 97.73 \ (3)^{\circ}$ V = 1674.3 (6) Å³ Z = 4

Data collection

5458 independent reflections
2764 reflections with $I > 2\sigma(I)$
$R_{\rm int} = 0.040$
$\theta_{\rm max} = 26.5^{\circ}$
$h = -11 \rightarrow 7$
$k = -9 \rightarrow 9$
$l = -20 \rightarrow 29$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0815P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.051$	+ 0.6287P]
$wR(F^2) = 0.141$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} = 0.001$
3438 reflections	$\Delta \rho_{\rm max} = 0.97 \text{ e } \text{\AA}^{-3}$
250 parameters	$\Delta \rho_{\rm min} = -0.37 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Selected geometric parameters (Å, °).

Ni1-O1	1.940 (2)	Ni1-O7	2.169 (3)
Ni1-O5	2.067 (2)	Ni1-N1	1.963 (3)
Ni1-O6	2.089 (3)	Ni1-N2	2.146 (3)
O1-Ni1-N1	176.36 (10)	O5-Ni1-N2	90.84 (10)
O1-Ni1-O6	89.16 (9)	N1-Ni1-O7	89.11 (12)

Table 2

H	yd	lrogen-	bond g	eome	try	(A, '	°).
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O5−H5A···O8	0.81 (2)	1.86 (2)	2.671 (4)	173 (4)
$O5-H5B\cdots O4^{i}$	0.81 (3)	2.14 (4)	2.928 (4)	165 (3)
$O6-H6A\cdots O2$	0.82(2)	1.85 (2)	2.633 (4)	160 (3)
$O6-H6B\cdots O4^{ii}$	0.81(3)	2.04 (3)	2.804 (4)	157 (3)
$O7-H7A\cdots O4$	0.81(4)	1.89 (4)	2.691 (4)	174 (3)
$O7 - H7B \cdot \cdot \cdot O3^{ii}$	0.83 (4)	1.92 (4)	2.724 (4)	163 (4)
O8−H8A…O3	0.83 (3)	2.13 (3)	2.918 (4)	159 (3)
$O8-H8B\cdots O2^{iii}$	0.82 (2)	1.94 (2)	2.725 (3)	159 (5)

Symmetry codes: (i) -x + 1, $y - \frac{1}{2}$, $-z + \frac{3}{2}$; (ii) -x + 1, $y + \frac{1}{2}$, $-z + \frac{3}{2}$; (iii) x, y - 1, z.

Water H atoms were initially located in a difference Fourier map and then refined with O-H = restrained to 0.809 (10)-0.828 (10) Å and $U_{iso}(H) = 1.5U_{eq}(O)$. Other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with C-H = 0.93 Å, and $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

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