metal-organic papers

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Key indicators

Single-crystal X-ray study T = 296 KMean σ (C–C) = 0.003 Å R factor = 0.028 wR factor = 0.054 Data-to-parameter ratio = 16.6

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

cis-Tetraaqua(η^2 -pyridine-2,5-dicarboxylato- $\kappa^2 O, N$)nickel(II) dihydrate

The title compound, $[Ni(C_7H_3NO_4)(H_2O)_4]\cdot 2H_2O$, was obtained by the reaction of nickel nitrate with 2,5-pyridinedicarboxylic acid (molar ratio 1:1) in water. The compound contains a six-coordinate Ni^{II} ion, which is bonded to the N atom and an O–C group of the carboxylate group in the 2-position of the 2,5-pyridinedicarboxylate ligand, and four water O atoms. Intermolecular hydrogen-bonding interactions are present, linking the nickel complex and water molecules in the crystal structure.

Comment

The complexation of metal ions using the deprotonated conjugate base of 2,5-pyridinedicarboxylic acid as a ligand has been reported in the literature (Liang, Hong, Su et al., 2001; Liang, Hong & Cao, 2001; Plater et al., 1998). Presumably as a result of the presence of one pyridine N atom and two carboxylate groups in this ligand and/or of the hydrothermal synthetic conditions, it was found that the ligand coordinates to two or more metal ions in a bridging and chelating mode. In order to compare the results and to understand the possible experimental conditions under which a discrete molecule or a coordination polymer can be obtained, we carried out experiments under ambient conditions. We report here the synthesis and crystal structure of a discrete nickel complex, (I). It appears that the experimental conditions, viz. ambient or hydrothermal, are more critical than the multiple functionality of the organic ligands used to obtain discrete or polymeric products.



In the title compound, (I), atom Ni1 is octahedrally coordinated by N1 and O5 of the carboxylate group in the 2-position of the 2,5-pyridinedicaboxylate ligand, and four water O atoms, O1-O4 (Fig. 1). For the four Ni-O bond lengths involving the coordinated water O atoms, we might expect to observe two longer bond lengths for the water O atoms *trans* to N1 and O5 of the 2,5-pyridinedicarboxylate ligand, based on the *trans* influence; the opposite is, in fact, observed [Ni1-O4 = 2.080 (2) Å and Ni1-O2 = 2.022 (2) Å *versus* Ni1-O1 = 2.097 (2) Å and Ni1-O3 = 2.071 (2) Å; Table 1]. The contradiction is apparently caused by many

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 $D_r = 1.832 \text{ Mg m}^{-3}$

Cell parameters from 1951

 $0.25 \times 0.10 \times 0.10$ mm

2857 independent reflections

1894 reflections with $I > 2\sigma(I)$

H-atom parameters not refined

 $w = 1/[\sigma^2(F_o^2) + (0.0207P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

Mo $K\alpha$ radiation

reflections

 $\mu = 1.66 \text{ mm}^{-1}$

T = 296 (2) K

Tablet, blue

 $R_{\rm int} = 0.036$

 $\theta_{\rm max} = 28.3^{\circ}$ $h = -9 \rightarrow 9$

 $\begin{array}{l} k = -10 \rightarrow 11 \\ l = -15 \rightarrow 15 \end{array}$

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

 $\Delta \rho_{\rm max} = 0.33 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.35 \text{ e} \text{ Å}^{-3}$

 $\theta = 2.5 - 25^{\circ}$

Z = 2



Figure 1

The molecular structure of (I), showing the atom-numbering scheme and displacement ellipsoids at the 50% probability level.



Figure 2

The crystal packing diagram of (I), with hydrogen bonds shown as dashed lines, viewed along the b axis.

intermolecular hydrogen-bonding interactions between H atoms of the coordinated water molecules and O atoms of either the crystallization water molecules O9 and O10, or atoms O5–O8 of the 2,5-pyridinedicarboxylate ligand (Fig. 2). The hydrogen-bonding interactions are normal (Steed & Atwood, 2000), based on $H \cdots A$ distances of 1.79–2.08 Å, $D \cdots A$ distances of 2.663 (3)–2.933 (3) Å and bond angles of 147–175° (Table 2).

Experimental

2,5–Pyridinedicarboxylic acid (0.085 g, 0.5 mmol) and water (3 ml) were combined and the pH value adjusted to \sim 7.0 with drops of an aqueous NaOH solution. The resulting solution was layered on top of another aqueous solution (3 ml) of nickel(II) nitrate hexahydrate (0.151 g, 0.5 mmol). After one week, blue single crystals of (I) were collected and dried in air (yield 50%).

Crystal data

$$\begin{split} & [\text{Ni}(\text{C}_{7}\text{H}_{3}\text{NO}_{4})(\text{H}_{2}\text{O})_{4}]\cdot2\text{H}_{2}\text{O} \\ & M_{r} = 331.91 \\ & \text{Triclinic, } P\overline{1} \\ & a = 7.0177 \text{ (5) } \text{\AA} \\ & b = 8.3138 \text{ (6) } \text{\AA} \\ & c = 11.3906 \text{ (8) } \text{\AA} \\ & \alpha = 107.128 \text{ (1)}^{\circ} \\ & \beta = 99.039 \text{ (1)}^{\circ} \\ & \gamma = 102.748 \text{ (1)}^{\circ} \\ & V = 601.60 \text{ (7) } \text{\AA}^{3} \end{split}$$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1995) $T_{\rm min} = 0.733, T_{\rm max} = 0.847$ 6432 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.054$ S = 0.822857 reflections 172 parameters

Table 1

Selected geometric parameters (Å, °).

Ni1-O1	2.0970 (16)	Ni1-N1	2.0637 (16)
Ni1-O2	2.0217 (15)	O5-C6	1.258 (2)
Ni1-O3	2.0709 (15)	O6-C6	1.235 (2)
Ni1-O5	2.0324 (14)	O7-C7	1.253 (2)
Ni1-O4	2.0801 (14)	O8-C7	1.250 (3)
O1-Ni1-O3	174.32 (6)	O4-Ni1-N1	169.62 (6)
O2-Ni1-O5	175.81 (7)		

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

$D \cdot \cdot \cdot A$	
	$D = \Pi \cdots A$
2.933 (2)	160
2.776 (2)	175
2.680 (2)	171
2.712 (2)	168
2.674 (2)	158
2.663 (2)	157
2.848 (2)	153
2.727 (2)	168
2.758 (2)	147
2.722 (2)	157
2.779 (2)	164
2.853 (2)	158
	2.933 (2) 2.776 (2) 2.680 (2) 2.712 (2) 2.674 (2) 2.663 (2) 2.727 (2) 2.758 (2) 2.772 (2) 2.779 (2) 2.853 (2)

Symmetry codes: (i) -x, -y, -z; (ii) -x, -y, 1-z; (iii) 1-x, -y, 1-z; (iv) -x, -1-y, -z; (v) 1-x, 1-y, 1-z; (vi) 1+x, 1+y, 1+z; (vii) 1-x, -y, -z.

The H atoms were located in a difference Fourier synthesis but were not refined; their $U_{\rm iso}$ values were set to 0.05 Å².

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1997); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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