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Key indicators

Single-crystal X-ray study
T = 298 K
Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$
R factor = 0.029
wR factor = 0.078
Data-to-parameter ratio = 11.7

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

μ -Benzene-1,2,4,5-tetracarboxylato- bis[triaqua(2,2'-bipyridine)nickel(II)] dihydrate

In the title compound, $[\text{Ni}_2(\text{C}_{10}\text{H}_2\text{O}_8)(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_6] \cdot 2\text{H}_2\text{O}$, the benzenetetracarboxylate anion, which lies on a center of symmetry, is monodentate to two bipyridine-chelated water-coordinated Ni atoms. The geometry of the Ni atom is that of an octahedron. The coordinated and uncoordinated water molecules are linked by hydrogen bonds into a network structure.

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Comment

The benzene-1,2,4,5-tetracarboxylate anion binds to metal atoms in a variety of bonding modes (Cao *et al.*, 2002), and a large number of coordination polymers have been structurally characterized (Hu *et al.*, 2003; Xiao & Zhu, 2003). These studies have been extended to the present binuclear nickel(II) complex, (I), in which the anion is present in a μ_2 -bridging mode (Fig. 1). The anion lies on a special position of $\bar{1}$ site symmetry and binds to each Ni atom through only one carboxylate O atom. The geometry around the Ni atoms is that of an octahedron, being chelated by the heterocycle and also bonded to three water molecules. The coordinated and uncoordinated water molecules interact through hydrogen bonds (Table 2), generating a three-dimensional network.

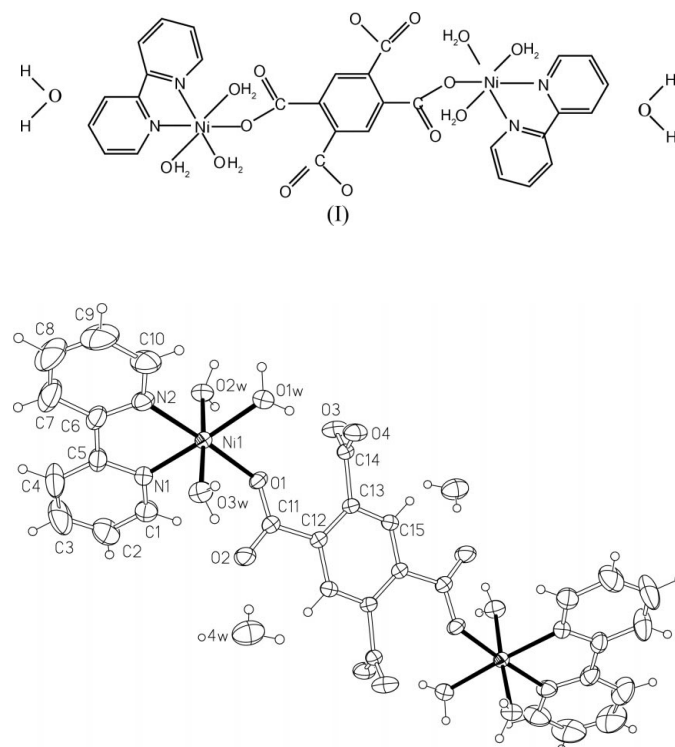


Figure 1
View of the structure of (I). Displacement ellipsoids are drawn at the 50% probability level.

Experimental

A DMF (10 ml) solution of $\text{Ni}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (0.15 g, 0.5 mmol) and benzene-1,2,4,5-tetracarboxylic acid (0.13 g, 0.5 mmol) was mixed with a DMF solution (10 ml) of 2,2'-bipyridine (0.08 g, 0.5 mmol). After one month, green crystals separated from the solution.

Crystal data

$[\text{Ni}_2(\text{C}_{10}\text{H}_2\text{O}_8)(\text{C}_{10}\text{H}_8\text{N}_2)_2 \cdot (\text{H}_2\text{O})_6] \cdot 2\text{H}_2\text{O}$	$Z = 1$
$M_r = 824.03$	$D_x = 1.617 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 7.5839$ (4) Å	Cell parameters from 4405 reflections
$b = 9.7257$ (6) Å	$\theta = 2.5\text{--}25.2^\circ$
$c = 11.9698$ (7) Å	$\mu = 1.19 \text{ mm}^{-1}$
$\alpha = 76.927$ (1)°	$T = 298$ (2) K
$\beta = 87.388$ (1)°	Prism, green
$\gamma = 79.807$ (1)°	$0.35 \times 0.32 \times 0.17 \text{ mm}$
$V = 846.41$ (8) Å ³	

Data collection

Bruker APEX area-detector diffractometer	3036 independent reflections
φ and ω scans	2903 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 2002)	$R_{\text{int}} = 0.013$
$T_{\text{min}} = 0.680$, $T_{\text{max}} = 0.823$	$\theta_{\text{max}} = 25.2^\circ$
6271 measured reflections	$h = -9 \rightarrow 9$
	$k = -11 \rightarrow 11$
	$l = -14 \rightarrow 14$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0459P)^2 + 0.4734P]$
$R[F^2 > 2\sigma(F^2)] = 0.029$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.078$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.02$	$\Delta\rho_{\text{max}} = 0.27 \text{ e \AA}^{-3}$
3036 reflections	$\Delta\rho_{\text{min}} = -0.31 \text{ e \AA}^{-3}$
259 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1

Selected geometric parameters (Å, °).

Ni1—N1	2.061 (2)	Ni1—O1w	2.075 (2)
Ni1—N2	2.065 (2)	Ni1—O2w	2.062 (2)
Ni1—O1	2.052 (1)	Ni1—O3w	2.082 (2)
N1—Ni1—N2	79.65 (7)	N2—Ni1—O3w	86.85 (7)
N1—Ni1—O1	97.78 (6)	O1—Ni1—O1w	85.00 (6)
N1—Ni1—O1w	176.98 (6)	O1—Ni1—O2w	89.71 (6)
N1—Ni1—O2w	88.26 (6)	O1—Ni1—O3w	91.54 (6)
N1—Ni1—O3w	89.09 (6)	O1w—Ni1—O2w	90.60 (6)
N2—Ni1—O1	176.99 (6)	O1w—Ni1—O3w	92.01 (6)
N2—Ni1—O1w	97.59 (7)	O2w—Ni1—O3w	177.20 (6)
N2—Ni1—O2w	91.80 (7)		

Table 2

Hydrogen-bonding geometry (Å, °).

$D\text{—}H \cdots A$	$D\text{—}H$	$H \cdots A$	$D \cdots A$	$D\text{—}H \cdots A$
O1w—H1w1 \cdots O4	0.84 (1)	2.15 (1)	2.989 (2)	176 (2)
O1w—H1w2 \cdots O3 ⁱ	0.85 (1)	1.87 (1)	2.714 (2)	174 (2)
O2w—H2w2 \cdots O4 ⁱ	0.85 (1)	1.90 (1)	2.737 (2)	172 (2)
O2w—H2w1 \cdots O4w ⁱⁱ	0.83 (2)	1.85 (2)	2.674 (2)	178 (2)
O3w—H3w1 \cdots O2	0.85 (1)	1.90 (1)	2.701 (2)	156 (2)
O3w—H3w2 \cdots O4 ⁱⁱⁱ	0.84 (1)	1.93 (1)	2.746 (2)	163 (2)
O4w—H4w1 \cdots O2	0.85 (1)	1.83 (1)	2.676 (2)	173 (3)
O4w—H4w2 \cdots O3 ^{iv}	0.85 (1)	1.96 (1)	2.781 (3)	161 (3)

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

The aromatic H atoms were positioned geometrically and were included in the refinement in the riding model approximation ($C\text{—}H = 0.93$ Å). The water H atoms were located and refined with distance restraints [$O\text{—}H = 0.85$ (1) Å and $H \cdots H = 1.39$ (1) Å]. The displacement parameters for all H atoms were $1.2U_{\text{eq}}(C, O)$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINTE* (Bruker, 2002); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2002); software used to prepare material for publication: *SHELXL97*.

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