

Polymeric tetraqua(1,2,4,5-benzenetetracarboxylato)(pyrazine)nickel(II) dihydrate

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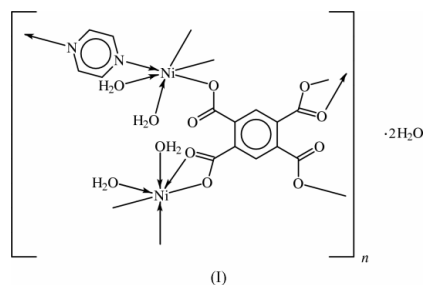
Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$
 R factor = 0.065
 wR factor = 0.142
Data-to-parameter ratio = 13.2For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The Ni atom in the title compound, $\{[\text{Ni}_2(\text{C}_{10}\text{H}_2\text{O}_8)(\text{C}_4\text{H}_4\text{N}_2)(\text{H}_2\text{O})_4] \cdot 2\text{H}_2\text{O}\}_n$, exists in an octahedral NO_5Ni coordination environment that is defined by the two O atoms of a chelating carboxyl group, the O atom of a monodentate carboxyl group belonging to another carboxylato tetraanion, two water molecules and the N atom of the *N*-heterocycle. The tetracarboxylato anion and the pyrazine molecule are each located on a center of inversion. The bonding pattern leads to the formation of a layer structure; the layers are linked into a three-dimensional network by extensive hydrogen bonds involving the water molecules and the O atoms of the anionic unit.

Comment

A number of adducts of nickel benzenepolycarboxylates with *N*-heterocycles have been reported (Li *et al.*, 2003; Yuan *et al.*, 2001). To this number has been added the 4,4'-bipyridine adduct of nickel terephthalate (Yang *et al.*, 2003a) and of nickel phthalate (Yang *et al.*, 2003b). The present pyrazine adduct, (I), of dinickel 1,2,4,5-benzenetetracarboxylate exists as the dihydrate (Fig. 1).



The Ni atom in (I) exists in an octahedral NO_5Ni environment that is defined by the two O atoms of a chelating $-\text{CO}_2$ group, the O atom of a monodentate CO_2 group of another tetraanion, two water molecules and the N atom of pyrazine (Table 1). The tetracarboxylate anion and the pyrazine molecule are each located on a center of inversion. The bonding pattern leads to the formation of layers; the layers are linked into a three-dimensional network by hydrogen bonds involving the water molecules and the O atoms of the anionic unit (Table 2).

Experimental

1,2,4,5-Benzenetetracarboxylic acid anhydride (pyromellitic anhydride, 0.22 g, 1 mmol) was dissolved in water (15 ml) containing sodium hydroxide (0.16 g, 4 mmol). Nickel dinitrate hexahydrate

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(0.58 g, 2 mmol) and pyrazine (0.16 g, 2 mmol) dissolved in water (3 ml) were then added. The mixture was placed in a 20 ml Teflon-lined stainless-steel bomb. The bomb was heated at 453 K for 100 h. Crystals separated from the solution when the bomb was cooled down at a rate of 5 K h⁻¹.

Crystal data

[Ni₂(C₁₀H₂O₈)(C₄H₄N₂)-
(H₂O)₄]-2H₂O
M_r = 555.72
Triclinic, P1
a = 7.2005 (5) Å
b = 8.0322 (5) Å
c = 9.3735 (6) Å
α = 96.175 (1)°
β = 101.872 (1)°
γ = 116.245 (1)°
V = 463.50 (5) Å³

Z = 1
D_x = 1.991 Mg m⁻³
Mo Kα radiation
Cell parameters from 2804
reflections
θ = 2.3–28.3°
μ = 2.12 mm⁻¹
T = 298 (2) K
Irregular block, green
0.14 × 0.13 × 0.04 mm

Data collection

Bruker SMART APEX area-
detector diffractometer
φ and ω scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
T_{min} = 0.616, T_{max} = 0.919
5415 measured reflections

2152 independent reflections
2055 reflections with I > 2σ(I)
R_{int} = 0.041
θ_{max} = 28.3°
h = -9 → 9
k = -10 → 10
l = -12 → 12

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.065
wR(F²) = 0.142
S = 1.25
2152 reflections
163 parameters
H atoms treated by a mixture of
independent and constrained
refinement

$w = 1/[\sigma^2(F_o^2) + (0.0629P)^2 + 0.259P]$
where $P = (F_o^2 + 2F_c^2)/3$
(Δ/σ)_{max} = 0.001
Δρ_{max} = 0.95 e Å⁻³
Δρ_{min} = -0.65 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Ni1—O1	2.126 (3)	Ni1—O1w	2.049 (3)
Ni1—O2	2.123 (3)	Ni1—O2w	2.025 (4)
Ni1—O3 ⁱ	2.014 (3)	Ni1—N1	2.068 (4)
O1—Ni1—O2	61.7 (1)	O2—Ni1—N1	85.8 (1)
O1—Ni1—O3 ⁱ	84.9 (1)	O3 ⁱ —Ni1—O1w	93.2 (1)
O1—Ni1—O1w	103.2 (1)	O3 ⁱ —Ni1—O2w	89.8 (1)
O1—Ni1—O2w	162.8 (1)	O3 ⁱ —Ni1—N1	173.6 (1)
O1—Ni1—N1	91.5 (1)	O1w—Ni1—O2w	93.4 (1)
O2—Ni1—O3 ⁱ	87.8 (1)	O1w—Ni1—N1	92.8 (1)
O2—Ni1—O1w	164.8 (1)	O2w—Ni1—N1	92.2 (2)
O2—Ni1—O2w	101.8 (1)		

Symmetry code: (i) 1 - x, 1 - y, 2 - z.

Table 2

Hydrogen-bonding geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O1w—H1w2...O1 ⁱⁱ	0.85 (1)	1.87 (3)	2.660 (5)	155 (5)
O1w—H1w1...O4 ⁱ	0.84 (1)	1.82 (2)	2.621 (5)	157 (5)
O2w—H2w2...O1w ⁱⁱⁱ	0.85 (1)	1.96 (1)	2.807 (5)	176 (6)
O2w—H2w1...O3w ^{iv}	0.85 (1)	1.90 (1)	2.744 (5)	176 (6)
O3w—H3w1...O2	0.85 (1)	2.26 (3)	3.047 (5)	155 (6)
O3w—H3w2...O3 ^v	0.85 (1)	2.02 (1)	2.866 (5)	174 (7)

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

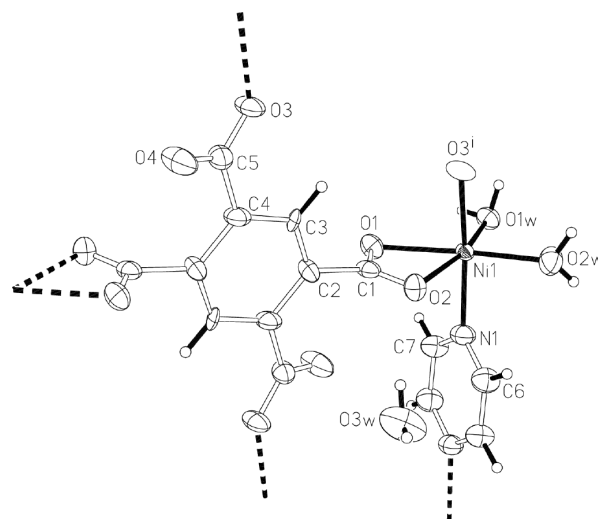


Figure 1

ORTEPII (Johnson, 1976) plot of a segment of the title structure, with displacement ellipsoids at the 75% probability level. [Symmetry code: (i) 1 - x, 1 - y, 2 - z.]

The use of too small a value for the θ-dependent absorption correction in SADABS (Sheldrick, 1996) led to non-positive definite displacement parameters for several atoms, and a compromise value of 0.5 was used, for which parameters of only one atom (C3) were non-positive definite. The displacement parameter of this atom was restrained to be close to isotropic. The aromatic H atoms were positioned geometrically (C—H = 0.93 Å) and were included in the refinement in the riding-model approximation, with U_{iso}(H) = 1.2U_{eq}(C). The water H atoms were located and refined with restraints of O—H = 0.85 (1) Å, H...H = 1.39 (1) Å and U_{iso}(H) = 1.2U_{eq}(O).

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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