

Diaquaiminodiacetonickel(II)

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Key indicators

Single-crystal X-ray study

T = 298 K

Mean $\sigma(\text{C}-\text{C}) = 0.008 \text{ \AA}$

R factor = 0.054

wR factor = 0.105

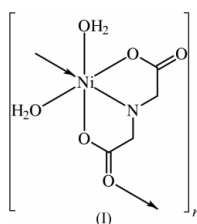
Data-to-parameter ratio = 13.0

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

The iminodiacetate dianion in diaquaiminodiacetonickel(II), $[\text{Ni}(\text{C}_4\text{H}_5\text{NO}_4)(\text{H}_2\text{O})_2]$, chelates to the Ni atom through two carboxyl O and one imino N atoms; the six-fold coordination environment comprises these three atoms, the water molecules and the carbonyl O atom of an adjacent dianionic group. The dative $\text{Ni} \leftarrow \text{O}$ linkage leads to the formation of a helical chain running along the *a* axis of the orthorhombic crystal; adjacent chains are held in a network motif by hydrogen bonds.

Comment

Unlike iminodiacetatecopper(II), a compound that forms a large number of coordination complexes, as well as complexes with metal salts (Román-Alpiste *et al.*, 1999), the nickel(II) analog is much less studied, and the crystal structure of iminodiacetonickel(II) has not been reported. The limited number of derivatives of this compound comprise the trihydrated 1:1 complex of dipotassium bis(iminodiacetato)nickelate (Agre *et al.*, 1984), caesium bis(iminodiacetato)nickelate tetrahydrate (Mammano *et al.*, 1977), lithium bis(iminodiacetato)nickelate tetrahydrate (Kramarenko *et al.*, 1974; Mammano *et al.*, 1977) and the only *N*-heterocycle adduct, tris(imidazole)iminodiacetonickel hydrate (Polyakova *et al.*, 2000).



Iminodiacetonickel crystallizes as a dihydrate, (I), in which the Ni atom exists in an octahedral environment; this comprises the *O,N,O'*-chelating iminodiacetate ligand, the two water molecules and the carbonyl O atom of an adjacent molecule. The dative $\text{Ni} \leftarrow \text{O}$ linkage [2.011 (4) Å] leads to the formation of a helical chain running along the *a* axis of the orthorhombic crystal (Fig. 2). The other carbonyl O atom does not participate in bonding to the Ni atom but is, instead, engaged in hydrogen bonding with a water molecule and the imino N atom of an adjacent molecule. The extensive hydrogen bonds consolidate the structure into a tightly held network.

Experimental

Nickel nitrate hexahydrate (0.29 g, 1 mmol) dissolved in water (2 ml) was reacted with iminodiacetic acid (0.13 g, 1 mmol) dissolved in

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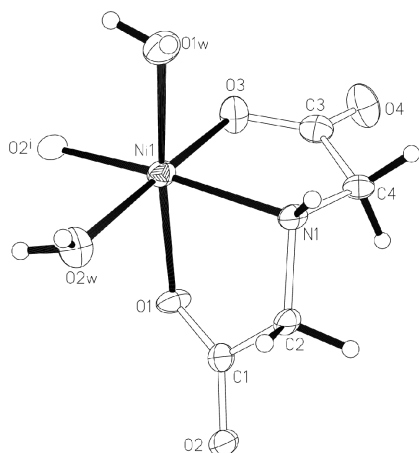


Figure 1
ORTEPII (Johnson, 1976) plot of the repeat unit of diaquaimino-diacetatonickel, with displacement ellipsoids drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.

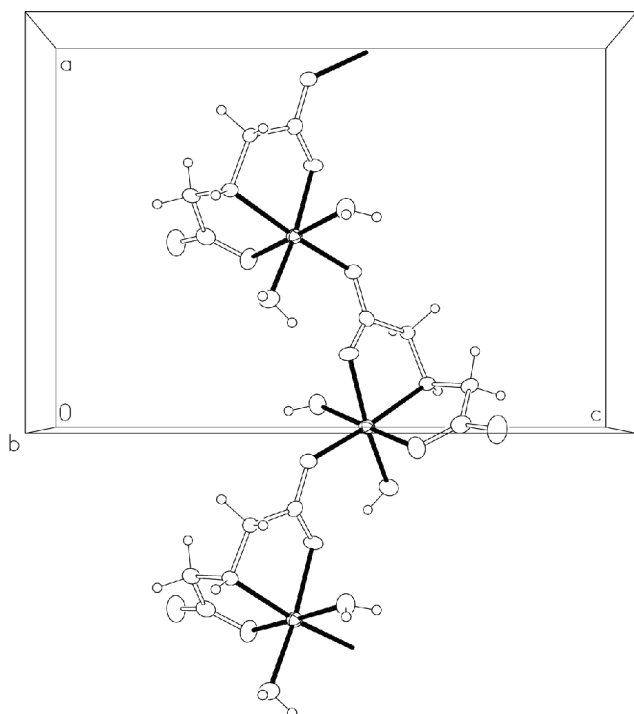


Figure 2
ORTEPII (Johnson, 1976) plot depicting the helical chain propagating along the *a* axis.

ethanol (8 ml). The pH of the solution was adjusted to 5 by the addition of aqueous sodium hydroxide. The solution was then sealed in a Teflon-lined Parr stainless-steel vessel, which was heated to 433 K for 2 h. The title compound separated as plates.

Crystal data

[Ni(C₄H₅NO₄)(H₂O)₂]
M_r = 225.83
 Orthorhombic, *P*2₁*ca*
a = 9.7610 (3) Å
b = 5.2219 (2) Å
c = 14.1713 (4) Å
V = 722.32 (4) Å³
Z = 4
D_x = 2.08 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 1988 reflections
 θ = 2.9–28.3°
 μ = 2.68 mm⁻¹
T = 298 (2) K
 Plate, green
 0.34 × 0.07 × 0.04 mm

Data collection

Bruker SMART area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 T_{\min} = 0.728, T_{\max} = 0.898
 4084 measured reflections

1613 independent reflections
 1585 reflections with $I > 2\sigma(I)$
 R_{int} = 0.035
 θ_{max} = 28.3°
 h = -12 → 12
 k = -6 → 2
 l = -16 → 18

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)]$ = 0.054
 $wR(F^2)$ = 0.105
 S = 1.25
 1613 reflections
 124 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0418P)^2 + 0.7011P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}}$ = 0.001
 $\Delta\rho_{\text{max}}$ = 0.89 e Å⁻³
 $\Delta\rho_{\text{min}}$ = -1.03 e Å⁻³
 Absolute structure: Flack (1983), 718 Friedel pairs
 Flack parameter = 0.12 (3)

Table 1

Selected geometric parameters (Å, °).

| | | | |
|--------------------------|-----------|--------------------------|-----------|
| Ni1—O1 | 2.123 (4) | Ni1—O1w | 2.087 (4) |
| Ni1—O2 ⁱ | 2.011 (4) | Ni1—O2w | 2.112 (4) |
| Ni1—O3 | 2.027 (4) | Ni1—N1 | 2.055 (5) |
| O1—Ni1—O2 ⁱ | 90.5 (2) | O2 ⁱ —Ni1—O2w | 91.7 (2) |
| O1—Ni1—O3 | 92.8 (2) | O3—Ni1—O1w | 92.7 (2) |
| O1—Ni1—O1w | 169.0 (2) | O3—Ni1—N1 | 83.6 (2) |
| O1—Ni1—O2w | 85.7 (2) | O3—Ni1—O2w | 178.5 (2) |
| O1—Ni1—N1 | 79.1 (2) | O1w—Ni1—O2w | 88.8 (2) |
| O2 ⁱ —Ni1—O3 | 88.2 (2) | O1w—Ni1—N1 | 92.0 (2) |
| O2 ⁱ —Ni1—N1 | 166.4 (2) | O2w—Ni1—N1 | 96.2 (2) |
| O2 ⁱ —Ni1—O1w | 99.2 (2) | | |

Symmetry code: (i) $x - \frac{1}{2}, -y, 1 - z$.

Table 2

Hydrogen-bonding geometry (Å, °).

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|------------------------------|-------------|---------------|-----------------------|-------------------------|
| O1w—H1w1...O1 ⁱ | 0.85 (5) | 2.05 (5) | 2.837 (6) | 154 (5) |
| O1w—H1w2...O3 ⁱⁱⁱ | 0.85 (2) | 2.10 (4) | 2.863 (6) | 150 (2) |
| O2w—H2w1...O2 ⁱⁱⁱ | 0.85 (2) | 2.19 (4) | 2.994 (6) | 159 (2) |
| O2w—H2w2...O4 ^{iv} | 0.85 (3) | 1.86 (3) | 2.673 (6) | 160 (3) |
| N1—H1n...O4 ⁱⁱ | 0.85 (5) | 2.11 (5) | 2.913 (7) | 157 (3) |

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

The H atoms of the imino N and water O atoms were located and refined, subject to O—H = N—H = 0.85 (1) Å; for the water H atoms, H...H = 1.39 (1) Å. The C-bound H atoms were generated geometrically and constrained with a riding model (C—H = 0.97 Å). The displacement parameters of all H atoms were set to 1.2 times U_{eq} of their parent atoms. The largest peak in the final difference map was about 1 Å from Ni1 and the deepest hole about 1 Å from C3. The structure was refined in the setting the data were connected on. The matrix (00 $\bar{1}$,010,100) transforms the cell into the standard *Pca*₂₁ setting.

Data collection: SMART (Bruker, 1997); cell refinement: SMART; data reduction: SAINT (Bruker, 1997); 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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