# metal-organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 290 KMean  $\sigma$ (C–C) = 0.008 Å R factor = 0.051 wR factor = 0.079 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.

# Bis(2-aminoacetato- $\kappa^2 N$ ,O)aquanickel(II)

The title compound,  $[Ni(C_2H_5NO_2)_2(H_2O)]$ , is a mononuclear nickel(II) complex. The Ni<sup>II</sup> atom is five-coordinated by two N atoms and two O atoms from two glycinate ligands in basal plane positions, and by one O atom from a coordinated water molecule in the apical position, forming a slightly distorted square-pyramidal geometry. In the crystal structure, molecules are linked through intermolecular  $N-H\cdots O$  and  $O-H\cdots O$  hydrogen bonds, forming a three-dimensional network.

### Comment

Transition metal compounds containing amino acid ligands are of great interest due to their biological activity and are present in the active sites of several important classes of metalloproteins. As an extension of the work on the structural characterization of amino acid complexes (Sun & You, 2004), a mononuclear nickel(II) compound, (I), is reported here.



The molecular structure of (I), with the atomic labeling scheme, is illustrated in Fig. 1, and selected bond distances, bond angles and torsion angles are given in Table 1. In (I), the Ni atom is in a square-pyramidal geometry, coordinated by two N atoms and two O atoms from two 2-aminoacetate acid ligands in the basal position, and by one O atom from a coordinated water molecule in the apical position. The slightly distorted square-pyramidal geometry is revealed by the bond angles between the apical and basal donor atoms (Table 1). In the basal positions, the four coordinating atoms around Ni are approximately coplanar, giving a square-planar coordination with an average deviation of 0.032(4) Å; the Ni atom lies 0.052 (2) Å above this plane; the two *trans* angles O1-Ni1-N2 and O3-Ni-N1 deviate from the ideal angle of 180° by 5.2 (2) and 2.3 (2) $^{\circ}$ , respectively; the other angles around the central Ni atom deviate from the ideal angle of 90°, ranging from 84.6 (2) to 95.7 (2) $^{\circ}$ ; this indicates a slightly distorted

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Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

square-pyramidal coordination of the Ni atom. The same character is also revealed by the torsion angles in the structure (Table 1).

In the crystal structure, the molecules are linked via three intermolecular N-H···O hydrogen bonds and two O- $H \cdots O$  hydrogen bonds, forming a three-dimensional network (details are given in Table 2 and Fig. 2).

## **Experimental**

All chemicals were obtained from a commercial source and used without further purification. 2-Aminoacetic acid (0.2 mmol, 15.0 mg) and Ni(CH<sub>3</sub>COO)<sub>2</sub>·2H<sub>2</sub>O (0.1 mmol, 21.3 mg) were dissolved in 20% EtOH aqueous solution (15 ml). The mixture was stirred for 30 min at room temperature to give a clear green solution. After allowing the solution to stand in air for 15 d, covered by a piece of filter paper, blue needle-shaped crystals of (I) formed at the bottom of the vessel on slow evaporation of the solvent.

#### Crystal data

$[Ni(C_2H_5NO_2)_2(H_2O)]$	Mo $K\alpha$ radiation		
$M_r = 224.85$	Cell parameters from 652		
Orthorhombic, $P2_12_12_1$	reflections		
a = 5.2174 (15)  Å	$\theta = 2.4  16.6^{\circ}$		
b = 10.823 (3) Å	$\mu = 2.53 \text{ mm}^{-1}$		
c = 13.517 (4) Å	T = 290 (2) K		
V = 763.3 (4) Å <sup>3</sup>	Needle, blue		
Z = 4	$0.13 \times 0.01 \times 0.01 \text{ mm}$		
$D_x = 1.957 \text{ Mg m}^{-3}$			

#### Data collection

Bruker APEX area-detector diffractometer  $\varphi$  and  $\varphi$  scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  $T_{\min} = 0.935, \ T_{\max} = 0.975$ 8284 measured reflections

### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.051$  $wR(F^2) = 0.079$ S = 1.011585 reflections 115 parameters H atoms treated by a mixture of independent and constrained refinement

Mo $K\alpha$ radiation
Cell parameters from 652
reflections
$\theta = 2.4 - 16.6^{\circ}$
$\mu = 2.53 \text{ mm}^{-1}$
T = 290 (2)  K
Needle, blue
$0.13 \times 0.01 \times 0.01 \text{ mm}$

1585 independent reflections 1245 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.105$  $\theta_{\rm max} = 26.5^{\circ}$  $h = -6 \rightarrow 6$  $k = -13 \rightarrow 13$  $l = -16 \rightarrow 16$ 

 $w = 1/[\sigma^2(F_0^2) + (0.0242P)^2]$ where  $P = (F_0^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.51 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\rm min} = -0.39 \text{ e} \text{ Å}^{-3}$ Absolute structure: Flack (1983), 634 Freidel pairs Flack parameter: 0.01 (4)



### Figure 2

The crystal packing of (I), viewed along the *a* axis. Hydrogen bonds are shown as dashed lines.

## Table 1

Selected geometric parameters (Å, °).

Ni1-N1	1.996 (4)	Ni1-O3	1.960 (3)
Ni1-N2	1.982 (4)	Ni1-O5	2.385 (4)
Ni1-O1	1.955 (4)		
O1-Ni1-N1	84.6 (2)	N2-Ni1-O5	95.7 (2)
O3-Ni1-N2	84.6 (2)	O1-Ni1-O5	89.2 (2)
O1-Ni1-O3	93.4 (2)	O3-Ni1-O5	91.3 (2)
N2-Ni1-N1	97.3 (2)	O1-Ni1-N2	174.8 (2)
N1-Ni1-O5	89.7 (2)	O3-Ni1-N1	177.7 (2)
01 N'1 01 C1	179.0 (4)	NO N'1 N1 CO	170.5 (4)
03-Nil-OI-CI	-1/8.9(4)	N2-Nil-NI-C2	1/9.5 (4)
N1 - Ni1 - O1 - C1	0.0 (4)	$05 - N_1 - N_1 - C_2$	-84.8 (4)
O5-Ni1-O1-C1	89.8 (4)	O3-Ni1-N2-C4	-4.9 (4)
O1-Ni1-O3-C3	177.6 (4)	N1-Ni1-N2-C4	176.3 (4)
N2-Ni1-O3-C3	2.5 (4)	O5-Ni1-N2-C4	85.9 (4)
O5-Ni1-O3-C3	-93.1(4)	O1-C1-C2-N1	8.0 (7)
O1-Ni1-N1-C2	4.4 (4)	O3-C3-C4-N2	-4.7 (7)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H1B \cdots O1^{i}$	0.90	2.22	3.110 (6)	169
$N2-H2A\cdots O3^{i}$	0.90	2.15	3.024 (6)	164
$N2-H2B\cdots O2^{n}$	0.90	2.09	2.980 (6)	171
$O5-H5A\cdots O4^{m}$	0.821 (10)	1.952 (12)	2.769 (6)	173 (5)
$O5-H5B\cdots O2^{W}$	0.823 (10)	1.986 (13)	2.808 (6)	177 (6)

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

The H atoms of the water ligand were located in a difference Fourier map and refined isotropically, with  $U_{iso}(H)$  values fixed at 0.08  $Å^2$ , and with O-H and H···H distances restrained to 0.84 (1) and 1.37 (2) Å, respectively. The other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H = 0.97 Å and N-H = 0.90 Å, and with  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C},{\rm N}).$ 

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Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); 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, 2002); software used to prepare material for publication: *SHELXTL*.

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