

supporting information

Acta Cryst. (2009). E65, m1518 [doi:10.1107/S1600536809043359]

Bis(dicyanamido- κN^1)bis[2-(2-hydroxyethyl)pyridine- $\kappa^2 N,O$]nickel(II)

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S1. Comment

In recent years there has been considerable interest in metal complexes supported by hydroxyethyl-pyridine, the ligand due to its versatile coordination activities and bridging function. (Sanudo *et al.*, 2003; Boskovic *et al.*, 2002). As an extension of this work, we have synthesized the title compound, (I), and report herein its crystal structure.

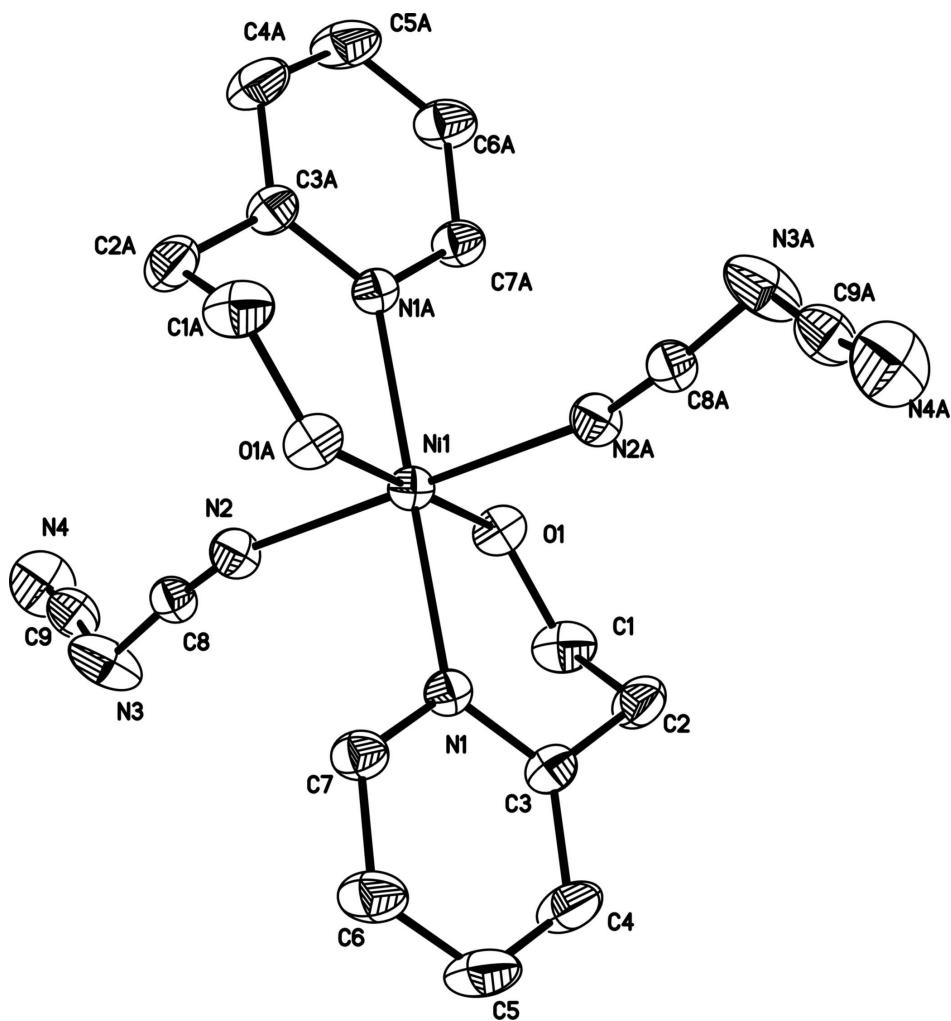
The complex (Fig. 1) consists of two L2- ($L = (\text{hydroxyethyl})(\text{pyridine})$) ligands, one Ni^{II} ion and two dicyanamiden ligands. The coordination geometry around the Ni center is octahedral with a NiN₄O₂ ligand set (Table 1). Two atoms N1 of hydroxyethylpyridine ligand occupy the axial sites. In the crystal structure, intermolecular O—H···N hydrogen bonds link molecules to form a one-dimensional chain along to the *c* axis (Table 2).

S2. Experimental

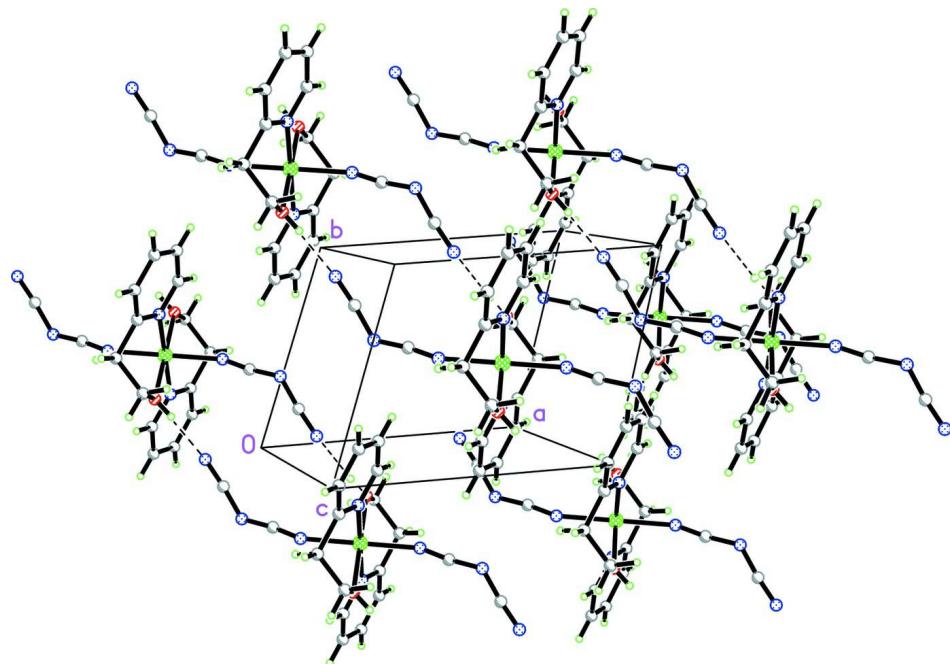
2-Hydroxyethylpyridine (0.123 g, 1 mmol) was deprotonated by Et₄NOH (25%) in the presence of nickel nitrate hexahydrate (0.5 mmol, 0.127 g) in a mixture of methanol and acetonitrile (V/V = 1:1) after the solution was stirred at room temperature for 0.5 h. Sodium dicyanamiden (5 mmol 0.486 g) was added to the above solution and then further stirred for 1 h. The resulting clear solution was filtered and left to stand at room temperature. Green blocks of (I) were obtained by slow evaporation of the solvents within 2 weeks. MP = 518–520 K (decomp).

S3. Refinement

All H atoms were placed geometrically and treated as riding on their parent atoms with C—H = 0.93–0.97 Å [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$] and O—H = 0.82 Å [$U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$].

**Figure 1**

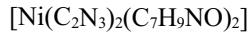
The structure of the title complex, showing 30% probability displacement ellipsoids. Atoms labelled with the suffix A are generated by the symmetry operation $(-x + 1, -y, -z + 1)$. H atoms have been omitted for clarity.

**Figure 2**

The crystal packing of (I), viewed approximately along the c axis.

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Crystal data



$$M_r = 437.11$$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$$a = 8.1498 (1) \text{ \AA}$$

$$b = 8.76020 (11) \text{ \AA}$$

$$c = 8.9201 (12) \text{ \AA}$$

$$\alpha = 100.841 (1)^\circ$$

$$\beta = 110.588 (2)^\circ$$

$$\gamma = 115.359 (2)^\circ$$

$$V = 493.66 (7) \text{ \AA}^3$$

$$Z = 1$$

$$F(000) = 226$$

$$D_x = 1.47 \text{ Mg m}^{-3}$$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1464 reflections

$$\theta = 2.7\text{--}26.3^\circ$$

$$\mu = 1.02 \text{ mm}^{-1}$$

$$T = 298 \text{ K}$$

Block, green

$$0.28 \times 0.20 \times 0.15 \text{ mm}$$

Data collection

Bruker SMART CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2003)

$$T_{\min} = 0.764, T_{\max} = 0.863$$

2566 measured reflections

1718 independent reflections

1579 reflections with $I > 2\sigma(I)$

$$R_{\text{int}} = 0.016$$

$$\theta_{\max} = 25.0^\circ, \theta_{\min} = 2.7^\circ$$

$$h = -9 \rightarrow 9$$

$$k = -7 \rightarrow 10$$

$$l = -10 \rightarrow 8$$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.034$
 $wR(F^2) = 0.095$
 $S = 1.00$
 1718 reflections
 133 parameters
 0 restraints
 Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
 Hydrogen site location: inferred from neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0648P)^2 + 0.0746P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.53 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.22 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Ni1	0.5000	0.0000	0.5000	0.03081 (18)
N1	0.7472 (3)	0.2759 (3)	0.6426 (3)	0.0341 (5)
N2	0.4639 (3)	0.0453 (3)	0.2748 (3)	0.0415 (5)
N3	0.3758 (4)	0.1688 (4)	0.0579 (4)	0.0700 (9)
N4	0.0438 (5)	0.0438 (4)	-0.1949 (4)	0.0731 (9)
O1	0.2970 (3)	0.0824 (2)	0.5031 (2)	0.0411 (4)
H1	0.1971	0.0414	0.4067	0.062*
C1	0.3611 (5)	0.2644 (4)	0.6012 (4)	0.0548 (8)
H1A	0.2523	0.2612	0.6245	0.066*
H1B	0.3849	0.3390	0.5343	0.066*
C2	0.5577 (5)	0.3491 (4)	0.7713 (4)	0.0537 (8)
H2A	0.5771	0.4562	0.8495	0.064*
H2B	0.5422	0.2608	0.8249	0.064*
C3	0.7468 (4)	0.4061 (4)	0.7512 (3)	0.0418 (6)
C4	0.9168 (5)	0.5852 (4)	0.8400 (4)	0.0614 (9)
H4	0.9143	0.6738	0.9136	0.074*
C5	1.0883 (5)	0.6328 (4)	0.8202 (5)	0.0653 (9)
H5	1.2039	0.7528	0.8817	0.078*
C6	1.0887 (5)	0.5030 (4)	0.7096 (4)	0.0559 (8)
H6	1.2034	0.5326	0.6932	0.067*
C7	0.9155 (4)	0.3274 (4)	0.6227 (4)	0.0428 (6)
H7	0.9151	0.2392	0.5456	0.051*
C8	0.4109 (4)	0.0950 (4)	0.1684 (3)	0.0375 (6)
C9	0.1941 (5)	0.0937 (4)	-0.0759 (4)	0.0489 (7)

O1—Ni1—N1 ⁱ	90.93 (7)	C5—C4—C3	120.3 (3)
O1 ⁱ —Ni1—N1 ⁱ	89.07 (7)	C5—C4—H4	119.8
N1—Ni1—N1 ⁱ	180.0	C3—C4—H4	119.8
C7—N1—C3	117.8 (2)	C6—C5—C4	119.4 (3)
C7—N1—Ni1	117.74 (17)	C6—C5—H5	120.3
C3—N1—Ni1	124.45 (18)	C4—C5—H5	120.3
C8—N2—Ni1	156.7 (2)	C5—C6—C7	118.3 (3)
C8—N3—C9	122.3 (3)	C5—C6—H6	120.8
C1—O1—Ni1	124.15 (16)	C7—C6—H6	120.8
C1—O1—H1	109.5	N1—C7—C6	123.6 (3)
Ni1—O1—H1	113.9	N1—C7—H7	118.2
O1—C1—C2	110.2 (2)	C6—C7—H7	118.2
O1—C1—H1A	109.6	N2—C8—N3	172.5 (3)
C2—C1—H1A	109.6	N4—C9—N3	173.2 (3)
N2 ⁱ —Ni1—N1—C7	115.4 (2)	O1—C1—C2—C3	74.7 (3)
N2—Ni1—N1—C7	−64.6 (2)	C7—N1—C3—C4	−1.0 (4)
O1—Ni1—N1—C7	−152.00 (19)	Ni1—N1—C3—C4	−179.8 (2)
O1 ⁱ —Ni1—N1—C7	28.00 (19)	C7—N1—C3—C2	179.1 (3)
N2 ⁱ —Ni1—N1—C3	−65.8 (2)	Ni1—N1—C3—C2	0.3 (3)
N2—Ni1—N1—C3	114.2 (2)	C1—C2—C3—N1	−56.9 (4)
O1—Ni1—N1—C3	26.8 (2)	C1—C2—C3—C4	123.2 (3)
O1 ⁱ —Ni1—N1—C3	−153.2 (2)	N1—C3—C4—C5	−0.5 (5)
O1—Ni1—N2—C8	9.2 (5)	C2—C3—C4—C5	179.4 (3)
O1 ⁱ —Ni1—N2—C8	−170.8 (5)	C3—C4—C5—C6	1.3 (6)
N1—Ni1—N2—C8	−80.0 (5)	C4—C5—C6—C7	−0.6 (5)
N1 ⁱ —Ni1—N2—C8	100.0 (5)	C3—N1—C7—C6	1.8 (4)
N2 ⁱ —Ni1—O1—C1	84.2 (2)	Ni1—N1—C7—C6	−179.3 (2)
N2—Ni1—O1—C1	−95.8 (2)	C5—C6—C7—N1	−1.0 (5)
N1—Ni1—O1—C1	−7.7 (2)	Ni1—N2—C8—N3	104 (2)
N1 ⁱ —Ni1—O1—C1	172.3 (2)	C9—N3—C8—N2	177 (2)
Ni1—O1—C1—C2	−34.9 (3)	C8—N3—C9—N4	172 (3)

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

Hydrogen-bond geometry (\AA , °)

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N4 ⁱⁱ	0.82	1.89	2.711 (3)	175

Symmetry code: (ii) $-x, -y, -z$.