Diazido(2,2′-bipyridyl)dimethanol-nickel(II)

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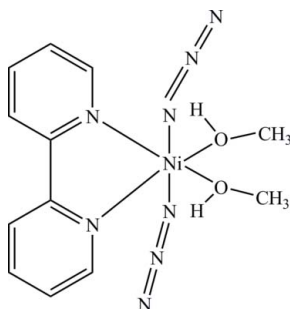Correspondence e-mail: weifanglhg@126.com

The title complex, $[Ni(N_3)_2(C_{10}H_8N_2)(CH_3OH)_2]$, lies on a twofold roation axis which runs through the Ni^{II} ion and the mid-point of the bipyridine ligand. The Ni^{II} ion is coordinated in a distorted octahedral environment by two azide ligands in a *trans* configuration. The methanol ligands are in a *cis* configuration and their hydroxy groups form intramolecular O—H···(N,N) hydrogen bonds with the azide ligands.

Related literature

For related structures, see: Urtiaga *et al.* (1995); Phatchimkun *et al.* (2009); Kou *et al.* (2008); Hou (2008).

Experimental

Crystal data

$[Ni(N_3)_2(C_{10}H_8N_2)(CH_4O)_2]$
M_r = 363.04
Monoclinic, $C2/c$
a = 16.8173 (15) Å
b = 13.4470 (12) Å
c = 7.1848 (6) Å
β = 111.238 (1)°

V = 1514.4 (2) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 1.31 mm^{-1}
T = 293 K
$0.32 \times 0.24 \times 0.19$ mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2008a)
T_{min} = 0.680, T_{max} = 0.790

3647 measured reflections
1333 independent reflections
1247 reflections with $I > 2\sigma(I)$
R_{int} = 0.012

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.058
$wR(F^2)$ = 0.210
S = 1.02
1333 reflections

106 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 1.08 e Å^{-3}
$\Delta\rho_{min}$ = −0.68 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1a···N3	0.93	2.25	2.721 (4)	111
O1—H1a···N4	0.93	2.48	3.227 (5)	137

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008b); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008b); molecular graphics: *SHELXTL* (Sheldrick, 2008b); software used to prepare material for publication: *SHELXTL*.

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supplementary materials

Diazido(2,2'-bipyridyl)dimethanolnickel(II)

H.-G. Li, S.-Y. Li and L.-J. Shao

Comment

As has been known for some time, 2,2'-bipyridine is a good bidentate chelating ligand and the azido group a good bridging ligand. Here, we present a new six-coordinate nickel(II) complex based on 2,2'-bipyridine azido ligands.

The molecular structure of the title compound is shown in Fig. 1. The coordination environment of the Ni^{II} ion distorted octahedral, in which two sites are occupied by the two N atoms of the chelating 2,2'-bipyridine ligand and the *trans* positions are occupied by two N atoms of two azido ligands. Two O atoms of two methanol ligands complete the coordination. The $Ni—N_{bipyridine}$ and $Ni—N_{azido}$ bond lengths are basically consistant with the corresponding distances found in other nickel bipyridine azido copmplexes (Urtiaga, *et al.*, 1995; Phatchimkun, *et al.*, 2009; Kou, *et al.*, 2008; Hou, 2008.).

Experimental

A mixture of 2,2'-bipyridine, $NiCl_2.6H_2O$ (1:1, molar ratio), excessive NaN_3 and methanol (20 ml) was sealed in a Teflon-lined autoclave (25 ml) and heated 353 K for 12 h. Upon cooling slowly and opening the bomb, green crystals suitable for X-ray diffraction were obtained with a yield about 55% (based on 2,2'-bipyridine).

Refinement

The H atoms bonded to C atoms were placed using the HFIX commands in *SHELXL-97* (Sheldrick, 2008*a*), with C—H distances of 0.93 and 0.96 Å, and were allowed for as riding atoms with $U_{iso}(H) = 1.2U_{eq}(C)$ and $1.5U_{eq}(C)$, respectively. The hydroxy H atom could not be found in a difference map but by including it using the HFIX 43 instruction in *SHELXL-97* there are likely acceptors for hydrogen bonds [see Table 1].
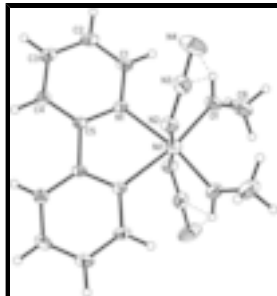
Figures

Fig. 1. The molecular structure of the title compound with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. Hydrogen bonds are shown as dashed lines. Unlabeled atoms are related by the symmetry operator -x+2, y, -z+1/2.

supplementary materials

Diazido(2,2'-bipyridyl)dimethanolnickel(II)

Crystal data

$[Ni(N_3)_2(C_{10}H_8N_2)(CH_4O)_2]$	$F(000) = 752$
$M_r = 363.04$	$D_x = 1.592$ Mg m^{-3}
Monoclinic, $C2/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: -C 2yc	Cell parameters from 1564 reflections
$a = 16.8173$ (15) Å	$\theta = 2.8–27.4°$
$b = 13.4470$ (12) Å	$\mu = 1.31$ mm^{-1}
$c = 7.1848$ (6) Å	$T = 293$ K
$\beta = 111.238$ (1)°	Block, green
$V = 1514.4$ (2) Å^3	$0.32 \times 0.24 \times 0.19$ mm
$Z = 4$	

Data collection

Bruker APEXII CCD diffractometer	1333 independent reflections
Radiation source: fine-focus sealed tube	1247 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.012$
φ and ω scans	$\theta_{max} = 25.0°$, $\theta_{min} = 2.0°$
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2008*a*)	$h = -15 \rightarrow 20$
$T_{min} = 0.680$, $T_{max} = 0.790$	$k = -15 \rightarrow 11$
3647 measured reflections	$l = -8 \rightarrow 8$

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.058$	Hydrogen site location: inferred from neighbouring sites
$wR(F^2) = 0.210$	H-atom parameters constrained
$S = 1.02$	$w = 1/[\sigma^2(F_o^2) + (0.1511P)^2 + 7.4217P]$ where $P = (F_o^2 + 2F_c^2)/3$
1333 reflections	$(\Delta/\sigma)_{max} = 0.011$
106 parameters	$\Delta\rho_{max} = 1.08$ e Å^{-3}
0 restraints	$\Delta\rho_{min} = -0.68$ e Å^{-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 ($Å^2$)

	x	y	z	$U_{iso}*/U_{eq}$
Ni1	1.0000	0.74532 (6)	0.2500	0.0357 (4)
O1	0.9127 (2)	0.6557 (3)	0.1994 (5)	0.0403 (8)
H1A	0.8679	0.6589	0.0766	0.048*
N1	0.9185 (2)	0.8653 (3)	0.2091 (5)	0.0267 (8)
N2	0.9797 (3)	0.7545 (3)	−0.0390 (6)	0.0331 (10)
N3	0.9118 (3)	0.7278 (3)	−0.1552 (6)	0.0393 (10)
N4	0.8473 (3)	0.7061 (5)	−0.2714 (8)	0.0713 (16)
C1	0.8326 (2)	0.8578 (3)	0.1560 (6)	0.0347 (10)
H1	0.8083	0.7953	0.1514	0.042*
C2	0.7806 (3)	0.9397 (4)	0.1091 (7)	0.0397 (11)
H2	0.7220	0.9327	0.0750	0.048*
C3	0.8157 (3)	1.0328 (4)	0.1126 (6)	0.0374 (10)
H3	0.7811	1.0889	0.0782	0.045*
C4	0.9035 (3)	1.0414 (3)	0.1684 (6)	0.0323 (9)
H4	0.9287	1.1033	0.1723	0.039*
C5	0.9526 (2)	0.9565 (3)	0.2179 (5)	0.0244 (8)
C6	0.9097 (4)	0.5858 (5)	0.3297 (8)	0.0599 (16)
H6A	0.9663	0.5619	0.4021	0.090*
H6B	0.8745	0.5317	0.2589	0.090*
H6C	0.8863	0.6138	0.4215	0.090*

Atomic displacement parameters ($Å^2$)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Ni1	0.0333 (6)	0.0382 (6)	0.0347 (6)	0.000	0.0114 (4)	0.000
O1	0.0367 (17)	0.0404 (19)	0.0380 (17)	−0.0107 (14)	0.0068 (13)	0.0056 (14)
N1	0.0201 (16)	0.0322 (18)	0.0275 (17)	−0.0002 (13)	0.0083 (13)	0.0000 (13)
N2	0.032 (2)	0.043 (2)	0.024 (2)	−0.0043 (14)	0.0095 (18)	0.0006 (13)
N3	0.048 (3)	0.041 (2)	0.031 (2)	−0.0041 (19)	0.016 (2)	0.0055 (17)
N4	0.057 (3)	0.100 (4)	0.042 (3)	−0.034 (3)	0.000 (2)	0.003 (3)
C1	0.0205 (19)	0.043 (3)	0.038 (2)	−0.0039 (17)	0.0078 (16)	0.0002 (18)
C2	0.024 (2)	0.054 (3)	0.041 (2)	0.0030 (19)	0.0108 (18)	−0.002 (2)
C3	0.029 (2)	0.043 (3)	0.037 (2)	0.0113 (18)	0.0087 (18)	0.0006 (18)
C4	0.031 (2)	0.033 (2)	0.032 (2)	0.0023 (17)	0.0103 (17)	−0.0023 (17)
C5	0.021 (2)	0.033 (2)	0.0198 (17)	0.0004 (15)	0.0069 (14)	−0.0010 (15)
C6	0.052 (3)	0.069 (4)	0.049 (3)	−0.024 (3)	0.008 (2)	0.008 (3)

Geometric parameters (Å, °)

Ni1—O1	1.831 (3)	C1—C2	1.370 (7)
Ni1—O1^i	1.831 (3)	C1—H1	0.9300
Ni1—N2	1.982 (4)	C2—C3	1.380 (7)
Ni1—N2^i	1.982 (4)	C2—H2	0.9300
Ni1—N1^i	2.068 (4)	C3—C4	1.387 (6)
Ni1—N1	2.068 (4)	C3—H3	0.9300
O1—C6	1.340 (6)	C4—C5	1.379 (6)
O1—H1A	0.9300	C4—H4	0.9300
N1—C5	1.346 (5)	C5—C5^i	1.490 (7)
N1—C1	1.358 (5)	C6—H6A	0.9600
N2—N3	1.199 (6)	C6—H6B	0.9600
N3—N4	1.142 (6)	C6—H6C	0.9600
O1—Ni1—O1^i	97.7 (2)	N4—N3—N2	176.8 (5)
O1—Ni1—N2	90.64 (15)	N1—C1—C2	121.9 (4)
O1^i—Ni1—N2	94.04 (16)	N1—C1—H1	119.1
O1—Ni1—N2^i	94.04 (16)	C2—C1—H1	119.1
O1^i—Ni1—N2^i	90.64 (15)	C1—C2—C3	119.6 (4)
N2—Ni1—N2^i	172.9 (2)	C1—C2—H2	120.2
O1—Ni1—N1^i	169.78 (14)	C3—C2—H2	120.2
O1^i—Ni1—N1^i	92.41 (16)	C2—C3—C4	119.0 (4)
N2—Ni1—N1^i	87.38 (14)	C2—C3—H3	120.5
N2^i—Ni1—N1^i	87.08 (15)	C4—C3—H3	120.5
O1—Ni1—N1	92.41 (16)	C5—C4—C3	118.8 (4)
O1^i—Ni1—N1	169.78 (14)	C5—C4—H4	120.6
N2—Ni1—N1	87.08 (15)	C3—C4—H4	120.6
N2^i—Ni1—N1	87.38 (14)	N1—C5—C4	122.5 (3)
N1^i—Ni1—N1	77.48 (19)	N1—C5—C5^i	113.9 (2)
C6—O1—Ni1	123.5 (3)	C4—C5—C5^i	123.6 (2)
C6—O1—H1A	118.2	O1—C6—H6A	109.5
Ni1—O1—H1A	118.2	O1—C6—H6B	109.5
C5—N1—C1	118.2 (3)	H6A—C6—H6B	109.5
C5—N1—Ni1	117.0 (2)	O1—C6—H6C	109.5
C1—N1—Ni1	124.4 (3)	H6A—C6—H6C	109.5
N3—N2—Ni1	118.1 (3)	H6B—C6—H6C	109.5

Symmetry codes: (i) −*x*+2, *y*, −*z*+1/2.

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O1—H1a···N3	0.93	2.25	2.721 (4)	111
O1—H1a···N4	0.93	2.48	3.227 (5)	137

Fig. 1