

cis-Bis(2,2-bipyridine)dichloronickel(II) methanol solvate

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Key indicators

Single-crystal X-ray study

 $T = 293\text{ K}$ Mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$ R factor = 0.047 wR factor = 0.101

Data-to-parameter ratio = 14.7

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

The title complex, $[\text{NiCl}_2(\text{bipy})_2]\cdot\text{CH}_3\text{OH}$, where bipy is 2,2-bipyridine ($\text{C}_{10}\text{H}_8\text{N}_2$), consists of an Ni atom coordinated to two *cis* chlorides and two bidentate bipyridines. One molecule of methanol is hydrogen bonded to one of the two inequivalent chlorides.

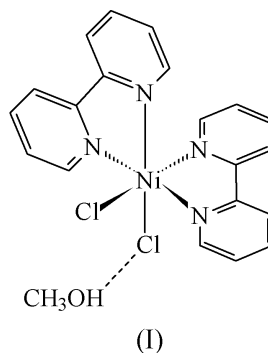
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Comment

In the course of our studies, the synthesis of $[\text{Ni}(\text{bipy})_2\text{Cl}_2]\cdot\text{CH}_3\text{OH}$, (I), was carried out using a method previously reported for the synthesis of $\text{Ni}(\text{pyridine})_4\text{Cl}_2$ (Long & Clarke, 1978). The molecule adopts a pseudo-octahedral geometry with N—Ni—N bite angles of 77.75 (14) and 78.00 (14) $^\circ$, and *cis* angles between 87.84 (10) and 97.58 (13) $^\circ$; *trans* angles are close to linearity. The structure of this complex (Fig. 1) is similar to the two structures of $\text{Ni}(\text{bipy})_2\text{Cl}_2$ previously reported.



If CH_3CN is used for recrystallization (Ferbinteanu *et al.*, 1998), two water molecules make strong hydrogen bonds with the chlorides. The structure of the dimethylformamide (DMF) solvate was also published the same year (Hipler *et al.*, 1998). In both published structures, a C_2 axis is observed on the nickel making both chlorides and bipy ligands equivalent by symmetry. Our studies revealed that the Ni—Cl bond lengths are quite different (more than 20σ) from one another. We believe that this is due to a methanol being hydrogen bonded only to Cl1 [3.276 (3) *versus* 3.467 (3) Å]. Also unexpected is that one of the bipy shows both of its Ni—N lengths [2.073 (3) and 2.077 (3) Å] shorter than the other bipy [2.084 (3) and 2.100 (4) Å].

Experimental

A solution of $\text{NiCl}_2\cdot 6\text{H}_2\text{O}$ (2 g) in methanol (25 ml) was added to a solution of 2,2-bipyridine (2.75 g) in methanol. This mixture was agitated for 1 h and concentrated under vacuum to a volume of 10 ml.

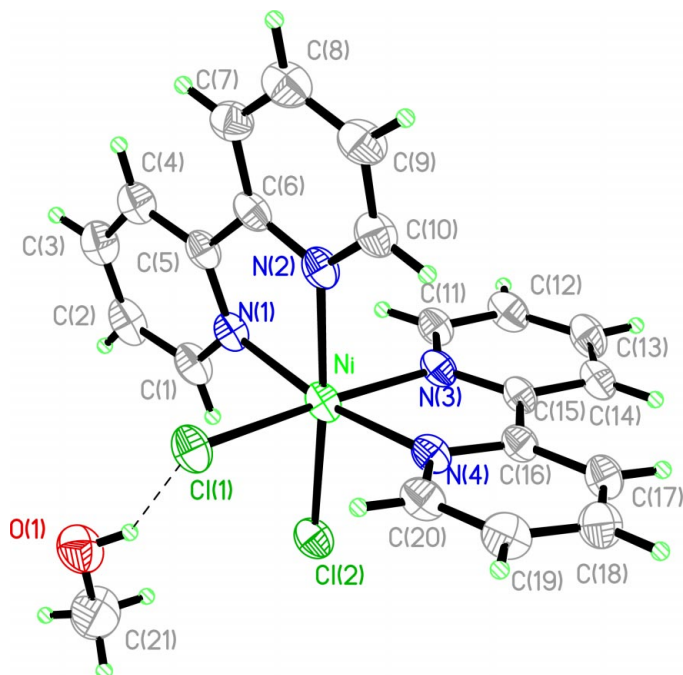


Figure 1

The structure of the title complex showing 30% probability displacement ellipsoids and the atom-numbering scheme.

Slow evaporation of the solvent in air allowed the formation of green crystals.

Crystal data

$[\text{NiCl}_2(\text{C}_{10}\text{H}_8\text{N}_2)_2] \cdot \text{CH}_4\text{O}$

$M_r = 474.02$

Orthorhombic, $Pbca$

$a = 10.455 (3) \text{ \AA}$

$b = 13.830 (4) \text{ \AA}$

$c = 28.548 (10) \text{ \AA}$

$V = 4128 (2) \text{ \AA}^3$

$Z = 8$

$D_x = 1.526 \text{ Mg m}^{-3}$

Cu $K\alpha$ radiation

Cell parameters from 25 reflections

$\theta = 20.0\text{--}23.0^\circ$

$\mu = 3.90 \text{ mm}^{-1}$

$T = 293 (2) \text{ K}$

Block, dark green

$0.23 \times 0.23 \times 0.16 \text{ mm}$

Data collection

Nonius CAD-4 diffractometer

$\omega/2\theta$ scans

Absorption correction: by integration

(*ABSORP* in *NRCVAX*;

Gabe *et al.*, 1989)

$T_{\min} = 0.432$, $T_{\max} = 0.593$

22 288 measured reflections

3908 independent reflections

2054 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.073$

$\theta_{\text{max}} = 69.9^\circ$

$h = 0 \rightarrow 12$

$k = 0 \rightarrow 16$

$l = 0 \rightarrow 34$

5 standard reflections

frequency: 60 min

intensity decay: none

Refinement

Refinement on F^2

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.101$

$S = 1.01$

3908 reflections

265 parameters

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.04P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} = 0.001$

$\Delta\rho_{\text{max}} = 0.31 \text{ e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.65 \text{ e \AA}^{-3}$

Extinction correction: *SHELXL96*

(Sheldrick, 1996)

Extinction coefficient: 0.00066 (5)

Table 1

Selected geometric parameters (\AA , $^\circ$).

Ni—N4	2.073 (3)	Ni—N2	2.100 (4)
Ni—N3	2.077 (3)	Ni—Cl1	2.3940 (13)
Ni—N1	2.084 (3)	Ni—Cl2	2.4213 (13)
N4—Ni—N3	78.00 (14)	N1—Ni—Cl1	88.60 (10)
N4—Ni—N1	170.07 (14)	N2—Ni—Cl1	87.84 (10)
N3—Ni—N1	97.58 (13)	N4—Ni—Cl2	94.06 (10)
N4—Ni—N2	93.37 (14)	N3—Ni—Cl2	88.70 (10)
N3—Ni—N2	91.44 (13)	N1—Ni—Cl2	94.73 (11)
N1—Ni—N2	77.75 (14)	N2—Ni—Cl2	172.43 (11)
N4—Ni—Cl1	95.58 (10)	Cl1—Ni—Cl2	92.87 (5)
N3—Ni—Cl1	173.48 (10)		

Table 2

Hydrogen-bonding geometry (\AA , $^\circ$).

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
O1—H1A \cdots Cl1	0.82	2.63	3.276 (3)	137
O1—H1A \cdots Cl2	0.82	2.78	3.467 (3)	143

H atoms were constrained to the parent site using a riding model ($C\cdots H$ 0.93–0.96 and $O\cdots H$ 0.82 \AA). The isotropic displacement parameters, U_{iso} , were adjusted to a value 50% higher than that of the parent site (methyl and OH) and 20% higher (others).

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *NRC-2* and *NRC-2A* (Ahmed *et al.*, 1973); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL96* (Sheldrick, 1996); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXL96*.

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