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Key indicators

Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
Disorder in solvent or counterion
 R factor = 0.041
 wR factor = 0.119
Data-to-parameter ratio = 13.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

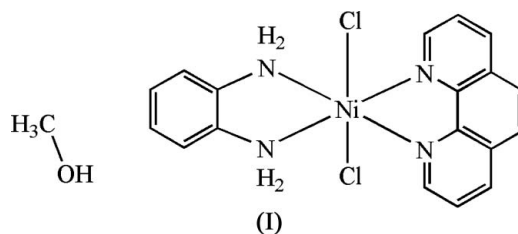
Dichloro(1,10-phenanthroline)(benzene-1,2-diamine)nickel(II) methanol solvate

The complex molecule of the title compound, $[\text{NiCl}_2(\text{C}_6\text{H}_8\text{N}_2)(\text{C}_{12}\text{H}_8\text{N}_2)] \cdot \text{CH}_4\text{O}$, has twofold rotation symmetry. The complex has a distorted octahedral NiCl_2N_4 coordination. Hydrogen-bonding and π - π stacking interactions help to stabilize the crystal structure.

Received 24 April 2006
Accepted 6 May 2006

Comment

The use of diamine complexes as hydrogenation catalysts or precatalysts has been reported previously (Noyori *et al.*, 2001; Haack *et al.*, 1997; Ohkuma *et al.*, 2002; Hedberg *et al.*, 2005; Ito *et al.*, 2001). In order to investigate the application of diamine complexes of Ni^{II} as hydrogenation catalysts, we have prepared the title complex, (I), and report here its structure.



The molecular structure of (I) is shown in Fig. 1. The molecule has a twofold rotation symmetry; the Ni^{II} ion and midpoints of the C3—C3A and C9—C9A bonds [symmetry code: (A) $2 - x, y, \frac{1}{2} - z$] are located on a twofold axis. The Ni^{II} ion is coordinated by two Cl^- anions and four N atoms in a distorted octahedral geometry (Table 1). The parallel phenanthroline rings of neighboring complexes are partially overlapped with a face-to-face separation of 3.463 (3) Å, indicating the existence of π - π stacking. Hydrogen-bonding and π - π stacking interactions (Table 2) help to stabilize the crystal structure (Fig. 2).

Experimental

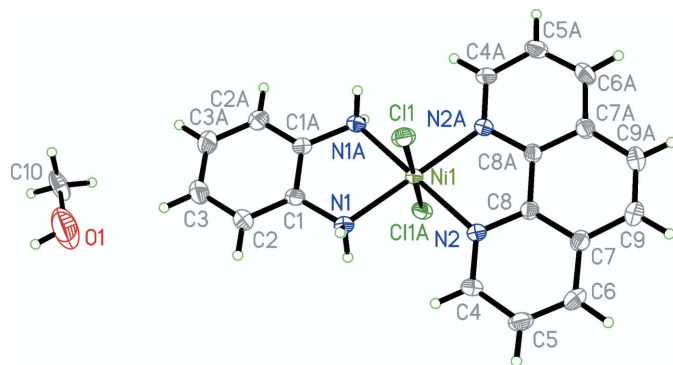
$\text{NiCl}_2 \cdot 6\text{H}_2\text{O}$ (1 mmol), 1,10-phenanthroline (1 mmol) and 1,2-phenylenediamine (1 mmol) were dissolved in methanol (12 ml). The solution was transferred into a 23 ml Teflon-lined autoclave and heated at 433 K for 80 h. The autoclave was then cooled over a period of 30 h to give purple single crystals of (I).

Crystal data

 $[\text{NiCl}_2(\text{C}_6\text{H}_8\text{N}_2)(\text{C}_{12}\text{H}_8\text{N}_2)] \cdot \text{CH}_4\text{O}$ $M_r = 450.00$ Monoclinic, $C2/c$ $a = 13.898$ (4) Å $b = 18.246$ (5) Å $c = 10.015$ (3) Å $\beta = 126.313$ (3)° $V = 2046.4$ (10) Å³ $Z = 4$ $D_x = 1.461$ Mg m⁻³Mo $K\alpha$ radiation $\mu = 1.23$ mm⁻¹ $T = 298$ (2) K

Block, purple

 $0.26 \times 0.22 \times 0.17$ mm


Figure 1

The molecular structure of (I) with 30% probability displacement ellipsoids (arbitrary spheres for H atoms) [symmetry code: (A) $2 - x, y, \frac{1}{2} - z$].

Data collection

Bruker SMART CCD area-detector diffractometer	5191 measured reflections
φ and ω scans	1794 independent reflections
Absorption correction: multi-scan (SADABS; Bruker, 1998)	1204 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.741, T_{\max} = 0.819$	$R_{\text{int}} = 0.034$
	$\theta_{\text{max}} = 25.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0592P)^2 + 1.4849P]$
$R[F^2 > 2\sigma(F^2)] = 0.041$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.119$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.07$	$\Delta\rho_{\text{max}} = 0.53 \text{ e } \text{\AA}^{-3}$
1794 reflections	$\Delta\rho_{\text{min}} = -0.28 \text{ e } \text{\AA}^{-3}$
132 parameters	
H-atom parameters constrained	

Table 1

Selected bond lengths (\AA).

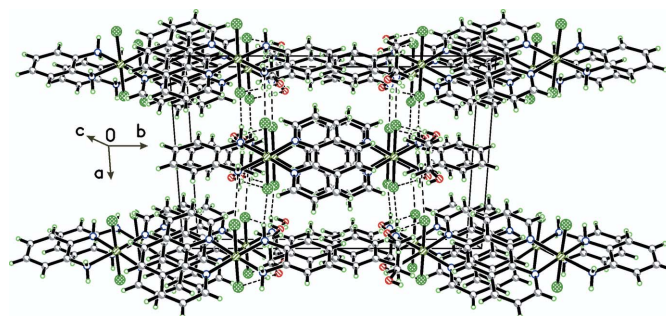
Ni1—N1	2.080 (3)	Ni1—Cl1	2.4671 (12)
Ni1—N2	2.077 (3)		

Table 2

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N1—H1A \cdots Cl1 ⁱ	0.90	2.49	3.267 (3)	144
O1—H1 \cdots Cl1 ⁱⁱ	0.82	2.36	3.168 (10)	171

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


Figure 2

A packing diagram, with dashed lines indicating hydrogen bonds.

The methanol solvent molecule is disordered over a twofold axis and its occupancy was fixed at 0.5 for the unique location. H atoms of the methanol molecule were located in a difference Fourier map and refined as riding with $O-H = 0.82, C-H = 0.96 \text{ \AA}$ and with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(O,C)$. Other H atoms were placed in calculated positions, with $N-H = 0.90 \text{ \AA}$ and $C-H = 0.93 \text{ \AA}$, and refined as riding, with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(N,C)$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

We thank the Scientific Research Foundation of Guangxi Normal University and the Science Foundation of Guangxi (Guikeqiang 0542021), China.

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