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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.006 \text{ Å}$ R factor = 0.044 wR factor = 0.131 Data-to-parameter ratio = 14.8

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

Hexakis(1*H*-imidazole- κN^3)cobalt(II) dichloride tetrahydrate

In the title compound, $[Co(C_3H_4N_2)_6]Cl_2\cdot 4H_2O$, the Co^{II} atom is situated on an inversion center. It is coordinated by six N atoms of six imidazole ligands and has a slightly distorted octahedral geometry. In the crystal structure, the Cl atoms, the water molecules of crystallization and the imidazole NH groups contribute to the formation of a three-dimensional hydrogen-bonded network.

Comment

Metal complexes with imidazole ligands have been of interest for decades due to the importance of the imidazole group in biological systems. We report here the crystal structure of a new complex, hexakis(1*H*-imidazole- κN^3)cobalt(II) dichloride tetrahydrate, (I).



The title compound, (I), consists of a centrosymmetric hexakis(imidazole)cobalt(II) dication, two chloride counterions, and four uncoordinated water molecules in the chemical structural unit, as shown in Fig. 1. Atom Co1 has an octahedral geometry and is coordinated by six N atoms from six imidazole



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Figure 2

The crystal packing of (I), showing the hydrogen-bonding interactions as dashed lines.

molecules. The three *trans* angles of the Co1 octahedron are 180° by symmetry, and the other angles vary from 89.39 (9) to 90.61 (9)°. The average Co-N bond length of 2.166 (2) Å is in the normal range for cobalt(II) complexes with imine N atoms. All the imidazole rings are planar, with a mean deviation of 0.0013 (2) Å. Opposing imidazole rings are parallel to one another, as required by the centrosymmetry. Neighboring imidazole rings are nearly perpendicular to one another, with dihedral angles of 84.4 (4), 85.2 (4) and 90.7 (4)°, thereby minimizing the steric effects between these rings.

In the crystal structure, the water molecules, involving atoms O1 and O2, the chloride anion Cl1, and all of the imidazole NH groups contribute to the formation of hydrogen bonds, as shown in Fig. 2. A number of intermolecular hydrogen-bond contacts involving NH groups and Cl⁻ anions link the cations into layers in the *ab* plane (see Table 1). These, in turn, are linked *via* the water molecules and Cl⁻ anions to form a three-dimensional network.

Experimental

 $CoCl_2 \cdot 6H_2O$ (1.0 mmol, 238 mg) and imidazole (6.0 mmol, 408 mg) were dissoved in water (20 ml). The mixture was stirred for a few minutes to obtain a clear pink solution. After allowing the resulting solution to stand in air for 20 d, dark-purple crystals were formed. The crystals were isolated, washed with water three times and dried in a vacuum desiccator using CaCl₂ (yield 39%). Elemental analysis found: C 35.50, H 5.35, N 27.69%; calculated for C₁₈H₃₂Cl₂CoN₁₂O₄: C 35.42, H 5.28, N 27.54%.

Crystal data

$[Co(C_3H_4N_2)_6]Cl_2\cdot 4H_2O$	Z = 1
$M_r = 610.39$	$D_x = 1.412 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 8.783 (4) Å	Cell parameters from 2655
b = 9.064 (4) Å	reflections
c = 10.575 (5) Å	$\theta = 2.6 - 26.3^{\circ}$
$\alpha = 75.156 \ (5)^{\circ}$	$\mu = 0.83 \text{ mm}^{-1}$
$\beta = 83.105 \ (6)^{\circ}$	T = 298 (2) K
$\gamma = 61.848 \ (5)^{\circ}$	Prism, dark purple
$V = 717.6 (5) \text{ Å}^3$	$0.42\times0.35\times0.21~\text{mm}$

Data collection

Bruker SMART CCD area-detector	2494 independent reflections
diffractometer	2186 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.021$
Absorption correction: multi-scan	$\theta_{\rm max} = 25.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -10 \rightarrow 9$
$T_{\min} = 0.715, \ T_{\max} = 0.843$	$k = -10 \rightarrow 6$
3747 measured reflections	$l = -12 \rightarrow 12$
Refinement	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0758P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.044$	+ 0.1518P]
$wR(F^2) = 0.131$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.23	$(\Delta/\sigma)_{\rm max} = 0.001$
2494 reflections	$\Delta \rho_{\rm max} = 0.51 \ {\rm e} \ {\rm \AA}^{-3}$
169 parameters	$\Delta \rho_{\rm min} = -0.62 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N2-H2B\cdots Cl1^{i}$	0.90	2.46	3.322 (3)	161
$N4-H4B\cdots Cl1^{ii}$	0.90	2.33	3.217 (3)	169
$N6-H6B\cdots Cl1^{iii}$	0.90	2.54	3.380 (3)	155
$O1 - H1B \cdot \cdot \cdot O2^{iii}$	0.85	2.42	2.720 (4)	101
$O1-H1C\cdots O1^{iv}$	0.85	2.23	2.815 (6)	126
$O2-H2C\cdots O1^{iii}$	0.85	2.19	2.720 (4)	121
$O2-H2D\cdots$ Cl1	0.85	2.44	3.178 (3)	145

Symmetry codes: (i) -x, 1-y, 1-z; (ii) x - 1, y, z - 1; (iii) 1 - x, 1 - y, 1 - z; (iv) -x, 2 - y, 1 - z.

All the H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms; N-H = 0.90 Å, C-H = 0.96 Å and O-H = 0.85 Å, with $U_{iso}(H)$ values fixed at 0.08 Å².

Data collection: *SMART* (Siemens, 1996); cell refinement: *SMART*; data reduction: *SAINT* (Siemens, 1996); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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