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Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.006$ Å
 R factor = 0.042
 wR factor = 0.103
Data-to-parameter ratio = 17.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Aqua[*N*-(2-oxidophenylmethylimino)-3-aza-
pentane-1,5-diamine]cobalt(II) chloride

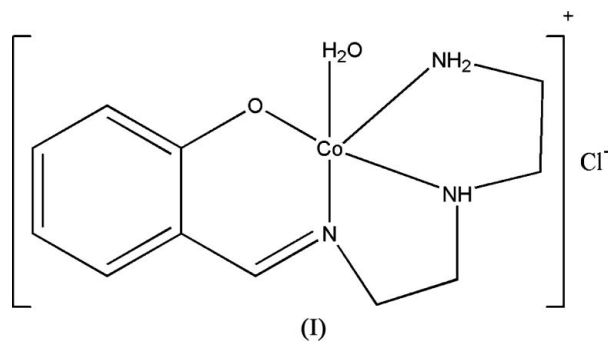
The title compound, $[\text{Co}(\text{C}_{11}\text{H}_{16}\text{N}_3\text{O})(\text{H}_2\text{O})]\text{Cl}$, consists of a $[\text{Co}L]^+$ cation and a chloride anion, where L is the *N*-(2-oxidophenylmethylimino)-3-azapentane-1,5-diamine anion. In the complex cation, the Co atom is coordinated by three N and two O atoms from the ligand and an apical water molecule in a distorted square-pyramidal geometry. The crystal structure is stabilized by $\text{N}-\text{H}\cdots\text{O}$, $\text{N}-\text{H}\cdots\text{Cl}$, $\text{O}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{Cl}$ hydrogen bonds.

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Comment

Schiff base complexes are of great interest in coordination chemistry (Goswami & Eichhorn, 1999; Dominguez-Vera *et al.*, 1998; Bernardo *et al.*, 1996). In recent years, a few Schiff base complexes with the *N*-(2-hydroxyphenylmethylimino)-3-azapentane-1,5-diamine anion ($\text{C}_{11}\text{H}_{16}\text{N}_3\text{O}^-$) have been reported. Most of these compounds consist of a univalent mononuclear complex cation and a simple anion, such as Cl^- , Br^- , ClO_4^- , SCN^- , PF_6^- and $[\text{B}(\text{Ph})_4]^-$ (Kratochvíl *et al.*, 1989, 1991; Liu *et al.*, 2004; Loub *et al.*, 1989.; Podlahová *et al.*, 1988; Cusmano Priolo *et al.*, 1983; Rotondo *et al.*, 1983; Zhu *et al.*, 2004). To date, there is only one dinuclear compound, $(\text{C}_{26}\text{H}_{44}\text{Cd}_2\text{N}_6\text{O}_4^{2+}\cdot 2\text{ClO}_4^-)$, in which the two Cd atoms are connected by two O atoms of the hydroxyl group (Loub *et al.*, 1990). We report here the structure of another mononuclear Co compound, (I), which is isostructural with the copper analog $[\text{Cu}(\text{C}_{11}\text{H}_{16}\text{N}_3\text{O})(\text{H}_2\text{O})]\text{Cl}$ (Zhu *et al.*, 2004).



As shown in Fig. 1, the Co^{II} atom in (I) is in a distorted square-pyramidal geometry, being coordinated by three N atoms and one O atom from the ligand and one O atom from a water molecule (Table 1). The four coordinating atoms from the ligand constitute the basal plane, with a mean deviation from the plane of 0.079 Å, and the O atom from the water molecule occupies the apical position. The Co^{II} atom is displaced by 0.1779 (16) Å above the basal plane towards the water molecule. The three $\text{Co}-\text{N}$ bonds represent the inter-

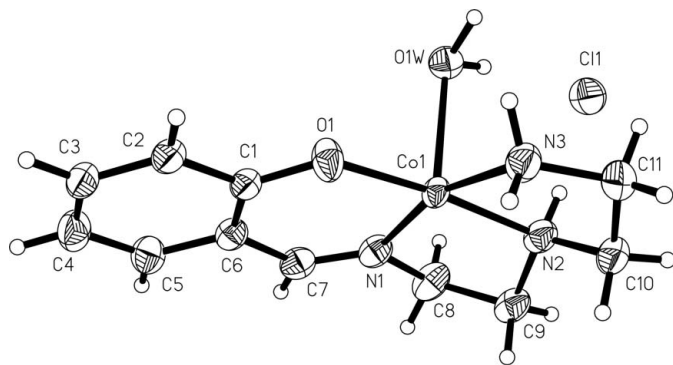


Figure 1
The structure of (I), showing 35% probability displacement ellipsoids (arbitrary spheres for the H atoms).

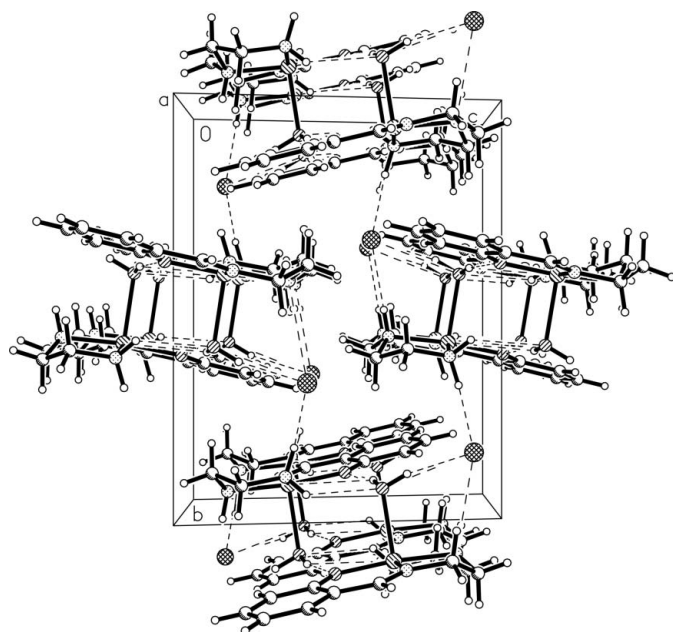


Figure 2
The packing of (I), showing hydrogen-bonded (dashed lines) layers in the *bc* plane.

action of the cobalt(II) centre with the primary amine, secondary amine and imine; they are in the normal bond-length ranges. The dihedral angle between the N1/N2/N3/O1 mean plane and the C1–C6 benzene ring mean plane is 11.90 (8)°.

Except for the imine N atom, all the other coordinating atoms and the uncoordinated chloride anion contribute to the formation of a hydrogen-bond network. The hydrogen bonds (Table 2) join two neighbouring cations to form a dimer. Further hydrogen bonds between the chloride ion and atoms O1W, N2, and N3 connect the dimers, forming a two-dimensional network in the *bc* plane, as shown in Fig. 2.

Experimental

A mixture of diethylenetriamine (1 mmol, 103 mg), salicylaldehyde (1 mmol, 122 mg) and $\text{CoCl}_2 \cdot 6\text{H}_2\text{O}$ (1 mmol, 238 mg) was stirred in

ethanol (10 ml) for 1 h. The precipitate was recrystallized from acetonitrile, and red blocks of (I) were filtered off and washed with MeCN and EtOH. Analysis found: C 41.40, H 5.72, N 13.23%; calculated for $\text{C}_{11}\text{H}_{18}\text{ClCoN}_3\text{O}_2$: C 41.46, H 5.69, N 13.19%.

Crystal data

$[\text{Co}(\text{C}_{11}\text{H}_{16}\text{N}_3\text{O})(\text{H}_2\text{O})]\text{Cl}$
 $M_r = 318.66$
 Monoclinic, $P2_1/c$
 $a = 9.047$ (4) Å
 $b = 13.633$ (5) Å
 $c = 10.754$ (5) Å
 $\beta = 101.87$ (1)°
 $V = 1298.0$ (10) Å³

$Z = 4$
 $D_x = 1.631$ Mg m⁻³
 Mo $K\alpha$ radiation
 $\mu = 1.53$ mm⁻¹
 $T = 293$ (2) K
 Block, red
 0.28 × 0.25 × 0.20 mm

Data collection

Bruker SMART CCD diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2001)
 $T_{\min} = 0.675$, $T_{\max} = 0.750$

2993 measured reflections
 2820 independent reflections
 2276 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$
 $\theta_{\max} = 27.0^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.103$
 $S = 1.08$
 2820 reflections
 164 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0306P)^2 + 2.5584P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.46$ e Å⁻³
 $\Delta\rho_{\min} = -0.40$ e Å⁻³
 Extinction correction: SHELXL97
 Extinction coefficient: 0.0105 (12)

Table 1

Selected bond lengths (Å).

Co1–O1	1.900 (3)	Co1–N3	2.007 (3)
Co1–N1	1.923 (3)	Co1–O1W	2.300 (3)
Co1–N2	1.994 (3)		

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
N3–H3NB...O1W ⁱ	0.92	2.42	3.096 (4)	130
N3–H3NA...Cl1 ⁱⁱ	0.98	2.38	3.330 (4)	162
N2–H2N...Cl1	0.90	2.35	3.241 (3)	170
O1W–H1WB...Cl1	0.91	2.23	3.117 (3)	165
O1W–H1WA...O1 ⁱ	1.01	1.69	2.665 (4)	163

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

All the H atoms, except for H1WA and H1WB attached to O1W, H3NA and H3NB attached to N3, were placed in idealized positions (C–H = 0.96 Å, N–H = 0.90–0.98 Å and O–H = 0.91 and 1.01 Å) and refined as riding with $U_{\text{iso}}(\text{H})$ values fixed at 0.08 Å² (C-bound H atoms) and 0.05 Å² (N- and O-bound H atoms). Atoms H1WA, H1WB, H3NA and H3NB were located from a difference Fourier map and it was allowed to attached atom during the final cycles of refinement.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:

SHELXTL (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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