# metal-organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma$ (C–C) = 0.008 Å R factor = 0.056 wR factor = 0.159 Data-to-parameter ratio = 14.0

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

# Bis{*N*-[2-(2-hydroxyethylamino)ethyl]salicylideneimine}chromium(III) chloride

The title compound,  $[Cr(C_{11}H_{15}N_2O_2)_2]Cl$ , is a mononuclear chromium(III) complex with a distorted octahedral geometry defined by an  $N_4O_2$  donor set.

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### Comment

Schiff bases obtained by condensation of *N*-2-hydroxyethylaminoethylamine and salicylaldehyde or analogues (*L*) have strong coordination abilities to transition metal ions. Recently, many such metal complexes have been structurally characterised: zinc(II) (Usman *et al.*, 2003; You *et al.*, 2004), cobalt(III) (Zhu *et al.*, 2003), silver(I) (Zhu *et al.*, 2000; Zou *et al.*, 2004), copper(II) (Qu *et al.*, 2004), nickel(II) (Ma *et al.*, 2005) and cadmium(II) (Yang *et al.*, 2004). However, no reports on related chromium(III) complexes have appeared. Here, the crystal structure of the title complex, (I), is described.



The mononuclear chromium(III) complex (Fig. 1) has a structure similar to that found for the cobalt(III) analogue (Zhu *et al.*, 2003), and comprises a complex cation and  $Cl^-$  anion. In the cation, the central atom is six-coordinated by four N and two O atoms derived from the two Schiff base ligands, forming a slightly distorted octahedron. In the crystal structure, there is a close association between the complex cation and  $Cl^-$  anion (Table 1). The  $Cl^-$  anion is stabilized by hydrogen bonding in a hydrophilic pocket located to one side of the cation and defined by two amine NH and two hydroxyl groups.

#### Experimental

© 2007 International Union of Crystallography All rights reserved Using a similar procedure to that in the literature (Zhu *et al.*, 2003), complex (I) was prepared as follows. Equimolar salicylaldehyde and



#### Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

2-hydroxylaminoethylamine were dissolved in anhydrous ethanol. The mixture was stirred to give a clear colorless solution of N-[2-(2hydroxyethylamino)ethyl]salicylideneimine. To this solution was added equimolar CrCl<sub>3</sub>·6H<sub>2</sub>O in anhydrous ethanol. After standing, dark-blue crystals of (I) were formed. They were isolated, washed with ethanol three times and dried in a vacuum desiccator using silica gel (yield 48%; m.p. 584 K).

#### Crystal data

[Cr(C<sub>11</sub>H<sub>15</sub>N<sub>2</sub>O<sub>2</sub>)<sub>2</sub>]Cl  $M_r = 501.95$ Monoclinic,  $P2_1/n$ a = 9.818 (7) Å b = 24.895 (18) Å c = 10.431 (7) Å  $\beta = 115.724 \ (11)^{\circ}$ 

#### Data collection

Bruker SMART CCD diffractometer Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  $T_{\min} = 0.795, T_{\max} = 0.920$ 

V = 2297 (3) Å<sup>3</sup> Z = 4Mo  $K\alpha$  radiation  $\mu = 0.65 \text{ mm}^{-1}$ T = 298 (2) K  $0.37 \times 0.25 \times 0.13 \text{ mm}$ 

12142 measured reflections 4052 independent reflections 2404 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.069$ 

$R[F^2 > 2\sigma(F^2)] = 0.056$	289 parameters
$wR(F^2) = 0.159$	H-atom parameters constrained
S = 0.93	$\Delta \rho_{\rm max} = 0.69 \text{ e} \text{ \AA}^{-3}$
4052 reflections	$\Delta \rho_{\rm min} = -0.38 \text{ e } \text{\AA}^{-3}$

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N2-H2···Cl1	0.91	2.55	3.457 (5)	172
N4-H4···Cl1	0.91	2.26	3.152 (4)	168
$O2-H2A\cdots Cl1$	0.82	2.74	3.498 (5)	154
$O4-H4A\cdots Cl1$	0.82	2.35	3.157 (5)	170

C-bound H atoms were included in the riding-model approximation with C-H = 0.93–0.97Å, and with  $U_{iso}(H) = 1.2U_{eq}(C)$ . The H atoms attached to N and O atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with N-H and C-H distances of 0.90 and 0.96 Å, respectively, and  $U_{iso}(H) = 1.2U_{eq}(N)$  and  $U_{iso}(H) = 1.5U_{eq}(O)$ .

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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