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#### **Key indicators**

Single-crystal X-ray study T = 292 KMean  $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$  R factor = 0.038 wR factor = 0.101 Data-to-parameter ratio = 16.4

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

# [Bis(1*H*-benzimidazol-2-ylmethyl)amine]dichloro(ethanol)manganese(II)

In the molecular structure of the title compound,  $[MnCl_2(C_{16}H_{15}N_5)(C_2H_6O)]$ , the Mn cation is in a distorted octahedral configuration. A three-dimensional framework structure is formed by means of  $N-H\cdots Cl$  and  $O-H\cdots Cl$ hydrogen bonds.

#### Comment

Histidine is an important ligand in iron, copper, zinc and manganese metalloproteins such as superoxide dismutases, lipoxygenases, tyrosinases, amine oxidases and hemocyanins (Que & Ho, 1996; Kaim & Rall, 1996). Bis(1*H*-benzimidazol-2-ylmethyl)amine (IDB) is a benzimidazole-rich ligand, which has the advantage that the basicity of the coordinating group approximates that of histidine (pK<sub>b</sub>: histidine = 7.96 and benzimidazole = 8.47; Main, 1992). We obtained the title compound, (I), in the process of synthesizing an IDB–metal complex in 95% ethanol with 1% dilute hydrochloric acid.



In (I), the Mn atom displays a distorted octahedral coordination geometry provided by the tridentate IDB ligand, two chloride anions and the O atom of an ethanol molecule (Fig. 1). The IDB ligand is *mer*-coordinated, with the central N3 atom in the axial position and atoms N2 and N4 of the benzimidazoyl groups in the equatorial positions. The remaining two equatorial positions are occupied by atoms Cl2 and O1. The bond distance between the Mn<sup>II</sup> ion and the axial N3 atom is about 0.21 Å longer than those between the Mn<sup>II</sup> ion and the equatorial N atoms of the benzimidazolyl groups (mean value = 2.210 Å; Table 1). The Mn–Cl bond length involving the equatorial atom Cl2 is about 0.24 Å longer than that involving the axial atom Cl1.

In the crystal structure, the molecules are linked into a three-dimensional framework by intermolecular N-H···Cl and O-H···Cl hydrogen bonds (Table 2 and Fig. 2). In addition to these interactions,  $\pi$ - $\pi$  stacking interactions between centrosymmetrically related imidazole rings at (x, y, z) and (-x, 1 - y, 1 - z) are observed, with an interplanar spacing of 3.379 Å and a centroid-centroid distance of 3.634 (1) Å.

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# Experimental

All reagents and solvents were used as obtained without further purification. Compound (I) was synthesized by refluxing stoichiometric quantities (1:1 molar ratio) of IDB (0.28 g, 1.0 mmol) and manganese(II) chloride tetrahydrate (0.20 g, 1 mmol) in 95% ethanol (30 ml) at 333 K for 4 h. The solution was cooled to room temperature, filtered and evaporated to obtain the product (yield 30%). Crystals of (I) were grown by slow evaporation of an ethanol solution.

Z = 4

 $D_x = 1.517 \text{ Mg m}^{-3}$ Mo *Ka* radiation  $\mu = 0.96 \text{ mm}^{-1}$ T = 292 (2) K

Block, colourless

 $\begin{aligned} R_{\rm int} &= 0.038\\ \theta_{\rm max} &= 27.0^\circ \end{aligned}$ 

 $0.30 \times 0.20 \times 0.10 \ \mathrm{mm}$ 

15190 measured reflections

4278 independent reflections

3433 reflections with  $I > 2\sigma(I)$ 

#### Crystal data

[MnCl <sub>2</sub> (C <sub>16</sub> H <sub>15</sub> N <sub>5</sub> )(C <sub>2</sub> H <sub>6</sub> O)]
$M_r = 449.24$
Monoclinic, $P2_1/c$
a = 7.2920 (5) Å
b = 13.8686 (10)  Å
c = 19.6429 (14)  Å
$\beta = 98.009 \ (1)^{\circ}$
$V = 1967.1 (2) \text{ Å}^3$

#### Data collection

Bruker SMART APEX CCD areadetector diffractometer  $\omega$  scans Absorption correction: multi-scan

(SADABS; Sheldrick, 2001) $T_{min} = 0.761, T_{max} = 0.910$ 

#### Refinement

 Refinement on  $F^2$   $w = 1/[\sigma^2(F_o^2) + (0.0569P)^2$ 
 $R[F^2 > 2\sigma(F^2)] = 0.038$   $w = 1/[\sigma^2(F_o^2) + (0.0569P)^2$ 
 $wR(F^2) = 0.101$  where  $P = (F_o^2 + 2F_c^2)/3$  

 S = 1.02  $(\Delta/\sigma)_{max} = 0.001$  

 4278 reflections
  $\Delta\rho_{max} = 0.32$  e Å<sup>-3</sup>

 261 parameters
  $\Delta\rho_{min} = -0.26$  e Å<sup>-3</sup>

 H atoms treated by a mixture of independent and constrained
  $\sigma_{min} = -0.26$  e Å<sup>-3</sup>

### Table 1

refinement

Selected	geometric	parameters	(Å,	°).
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Mn1-N4	2.2027 (17)	Mn1-Cl1	2.4057 (7)
Mn1-N2	2.2159 (17)	Mn1-N3	2.4210 (19)
Mn1-O1	2.2512 (17)	Mn1-Cl2	2.6444 (7)
N4-Mn1-N2	143.31 (7)	O1 - Mn1 - N3	79.89 (7)
N4-Mn1-O1	89.40 (6)	Cl1-Mn1-N3	176.75 (5)
N2-Mn1-O1	87.21 (6)	N4-Mn1-Cl2	91.04 (5)
N4-Mn1-Cl1	107.75 (5)	N2-Mn1-Cl2	84.62 (5)
N2-Mn1-Cl1	108.93 (5)	O1-Mn1-Cl2	166.90 (5)
O1-Mn1-Cl1	96.94 (5)	Cl1-Mn1-Cl2	95.38 (2)
N4-Mn1-N3	71.62 (6)	N3-Mn1-Cl2	87.83 (5)
N2-Mn1-N3	71.82 (6)		

Table 2

Hydrogen-bond geometry (Å, °).

$\overline{D-\mathrm{H}\cdots A}$	$D-{\rm H}$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
$\begin{array}{c} N5-H5A\cdots Cl2^{i}\\ O1-H1A\cdots Cl2^{ii}\\ N1-H1\cdots Cl2^{iii} \end{array}$	0.86 (1) 0.82 (3) 0.86 (1)	2.48 (2) 2.34 (2) 2.41 (2)	3.215 (2) 3.1265 (18 3.186 (2)	144 (2)           3)         161 (3)           150 (2)
Symmetry codes: -x + 1, -y + 1, -z + 1.	(i) -	$-x+1, y+\frac{1}{2}, -z+\frac{1}{2};$	(ii)	x - 1, y, z; (iii)



#### Figure 1

The molecular structure of (I), showing displacement ellipsoids drawn at the 30% probability level.



#### Figure 2

A partial packing diagram for (I), showing the intermolecular hydrogen bonds as dashed lines.

H atoms bonded to C atoms were placed in calculated positions, with C–H distances of 0.93–0.97 Å, and refined using a riding model, with  $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$ . H atoms bonded to N and O atoms were located in difference density maps and isotropically refined with soft restraints, N–H = 0.86 (1) Å and O–H = 0.82 (1) Å.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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