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#### **Key indicators**

Single-crystal X-ray study T = 295 KMean  $\sigma(\text{C-C}) = 0.004 \text{ Å}$  R factor = 0.042 wR factor = 0.111 Data-to-parameter ratio = 15.2

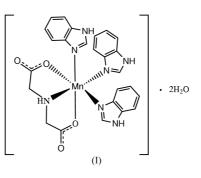
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# Tris(1*H*-benzimidazole- $\kappa N^3$ )(iminodiacetato- $\kappa^2 O, O'$ )manganese(II) dihydrate

In the title compound,  $[Mn(C_4H_5NO_4)(C_7H_6N_2)_3]\cdot 2H_2O$ , the  $Mn^{II}$  atom is surrounded by one iminodiacetate dianion and three benzimidazole molecules in a distorted octahedral geometry. The tridentate iminodiacetate dianion chelates the  $Mn^{II}$  atom in the facial mode. Extensive intermolecular,  $N - H \cdots O$  and  $O - H \cdots O$  hydrogen bonding and  $C - H \cdots \pi$ (arene) interactions are present in the crystal structure.

#### Comment

As part of investigations of the non-covalent interactions between aromatic rings in metal complexes, the title benzimidazole (BZIM) complex of Mn<sup>II</sup>, (I), has been prepared and its crystal structure is presented here.



The molecular structure of (I) is illustrated in Fig. 1. The  $Mn^{II}$  atom is surrounded by one iminodiacetate dianion (IDA) and three benzimidazole molecules (BZIM) in a distorted octahedral coordination geometry, similar to that found in the Ni<sup>II</sup> analogue (Su *et al.*, 2004). The IDA chelates, in a tridentate fashion, the Mn<sup>II</sup> atom in a facial configuration. Of the two five-membered chelating rings, the O1-containing ring is nearly planar, the maximum deviation being 0.061 (2) Å (for atom C2), whereas the O3-containing ring displays an envelope conformation, the Mn atom lying in the flap position and out of the mean plane formed by the other four atoms by 0.867 (4) Å. Such a facial configuration of IDA agrees with the situation found in other reported IDA complexes (Yasui *et al.*, 1987; Castineiras *et al.*, 1993).

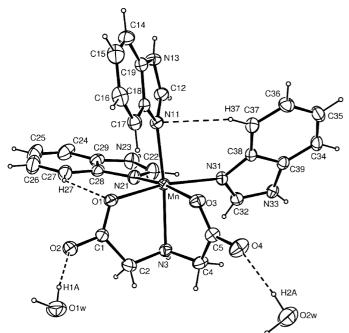
The carboxyl groups of IDA coordinate, in a monodentate fashion, to the Mn<sup>II</sup> atom, whereas uncoordinated carboxy O atoms (O2 and O4) are hydrogen bonded to water molecules and to BZIM of neighboring complex molecules, as shown in Fig. 2 and Table 1.

A *PLATON* analysis (Spek, 1980) gave the following  $H \cdots Cg$  distances and  $C-H \cdots Cg$  angles (*Cg* is the ring centroid), suggesting weak  $C-H \cdots \pi$ (arene) interactions in the crystal structure, as shown in Fig. 3 (Nishio *et al.*, 1998):

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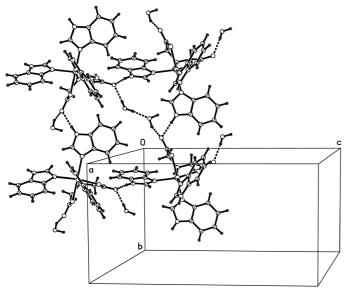
Acta Cryst. (2004). E60, m1063–m1065 DOI: 10.1107/S1600536804015910 Su, Yin and Xu • [Mn(C<sub>4</sub>H<sub>5</sub>NO<sub>4</sub>)(C<sub>7</sub>H<sub>6</sub>N<sub>2</sub>)<sub>3</sub>]·2H<sub>2</sub>O **m1063** 

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The molecular structure of (I), shown with 30% probability displacement ellipsoids. Dashed lines indicate hydrogen bonding, including possible weak intramolecular  $C-H\cdots O$  and  $C-H\cdots N$  interactions.



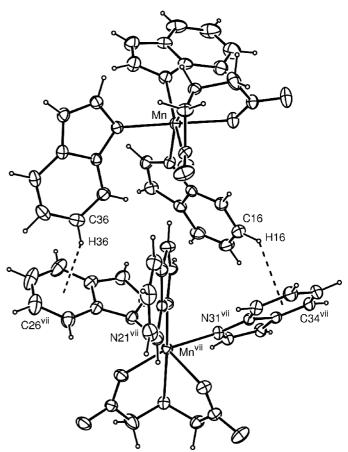
### Figure 2

A partial packing diagram of (I), showing the hydrogen bonding (dashed lines).

H16···*Cg*(C34<sup>vii</sup> ring) = 2.78 Å, C16–H16···*Cg*(C34<sup>vii</sup> ring) = 139°; H36··*Cg*(C26<sup>vii</sup> ring) = 2.79 Å, C36–H36··*Cg*(C26<sup>vii</sup> ring) = 148° [symmetry code: (vii)  $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$ ].

## Experimental

All reagents were commercially available and of analytical grade. Mn(CH<sub>3</sub>COO)<sub>2</sub>·4H<sub>2</sub>O (0.24 g, 1 mmol), H<sub>2</sub>IDA (0.13 g, 1 mmol), BZIM (0.24 g, 2 mmol) and Na<sub>2</sub>CO<sub>3</sub> (0.11 g, 1 mmol) were dissolved in a water (6 ml)–ethanol(2 ml) mixture. The solution was stirred at room temperature until the bubbles of CO<sub>2</sub> disappeared. The solution was then refluxed for 4 h and a small amount of precipitate





A diagram showing the intermolecular  $C-H\cdots\pi$  interactions (dashed lines). [Symmetry code: (vii) 1 - x,  $y - \frac{1}{2}, \frac{1}{2} - z$ .]

appeared. The solution was filtered into hot water (10 ml), then kept at room temperature. Colorless single crystals were obtained from the filtrate after 2 d.

#### Crystal data

 $[Mn(C_4H_5NO_4)(C_7H_6N_2)_3]\cdot 2H_2O$  $D_x = 1.426 \text{ Mg m}^{-3}$  $M_r = 576.48$ Mo  $K\alpha$  radiation Monoclinic,  $P2_1/c$ Cell parameters from 16 558  $a = 13.5852 (7) \text{\AA}$ reflections b = 10.4183 (4) Å $\theta = 2.0-25.0^{\circ}$  $\mu = 0.55 \text{ mm}^{-1}$ c = 19.5622 (7) Å $\beta = 104.065 (2)^{\circ}$ T = 295 (2) K $V = 2685.7 (2) \text{ Å}^3$ Prism, colorless  $0.27 \times 0.21 \times 0.11 \ \mathrm{mm}$ Z = 4

# Data collection

Rigaku R-AXIS RAPID diffractometer	5333 independent reflections 4315 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\rm int} = 0.035$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.1^{\circ}$
(ABSCOR; Higashi, 1995)	$h = -16 \rightarrow 16$
$T_{\min} = 0.86, T_{\max} = 0.94$	$k = -12 \rightarrow 12$
21 886 measured reflections	$l = -24 \rightarrow 24$

## Refinement

Table 1	
Hydrogen-bonding geometry (Å, °).	

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
O1W−H1A···O2	0.84	2.00	2.822 (3)	164
$O1W - H1B \cdot \cdot \cdot O2W^{i}$	0.95	1.89	2.814 (3)	163
$O2W-H2A\cdots O4$	0.96	1.90	2.797 (3)	155
$O2W - H2B \cdot \cdot \cdot O2^{ii}$	0.95	2.10	3.001 (3)	159
$N3-H3\cdots O1W^{iii}$	0.91	2.14	3.025 (3)	164
$N13-H13\cdots O3^{iv}$	0.86	2.19	3.037 (3)	167
$N23-H23\cdots O4^{v}$	0.86	1.93	2.770 (3)	165
$N33{-}H33{\cdots}O2^{vi}$	0.86	1.98	2.819 (3)	166

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

H atoms of water molecules were located in a difference Fourier map and included in the structure-factor calculations with fixed positional and isotropic displacement parameters of  $U_{\rm iso}({\rm H}) = 0.08 \text{ Å}^2$ . Other H atoms were placed in calculated positions, with C–H = 0.93 (aromatic) or 0.97 Å (methylene) and N–H = 0.86 (imidazole) or 0.91 Å (imino), and were included in the final cycles of refinement in riding mode, with  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$ (carrier atoms).

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/ MSC, 2002); program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *XP* (Siemens, 1994); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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