metal-organic compounds

Acta Crystallographica Section C

Crystal Structure Communications

ISSN 0108-2701

Bis(μ -6-hydroxypicolinato)- μ -oxobis[dipyridinemanganese(III)] monohydrate

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Received 6 May 2004 Accepted 24 May 2004 Online 22 June 2004

The title compound, $[Mn_2(\mu\text{-}O)(C_6H_3NO_3)_2(C_5H_5N)_4]\cdot H_2O,$ was isolated from the reaction of 2,6-pyridinedicarboxylic acid with $[Mn_{12}O_{12}(CH_3COO)_{16}(H_2O)_4]$ in pyridine. The dimanganese complex has twofold symmetry; the Mn^{III} atoms are bridged by one oxo and two amidate ligands and show compressed octahedral Jahn–Teller distortion. The molecular packing comprises a three-dimensional structure constructed by means of extensive intermolecular interactions, including three kinds of hydrogen bonds and $\pi\text{-}\pi$ interactions.

Comment

The discovery of the role of manganese at the active sites in various enzymatic systems has given considerable impetus to research in the field of polynuclear complexes of manganese. The manganese catalase in *Lactobacillus platarum* has been shown to contain a dimanganese unit with $[Mn_2^{III}(\mu-oxo)(\mu-carboxylato)_2]$ cores (Fronko *et al.*, 1988). Several Mn complexes containing the $[Mn_2(\mu-O)]^{4+}$ core have been

reported so far (Sheats et al., 1987; Wieghardt et al., 1988; Wu et al., 1990; Hotzelmann et al., 1992; Vincent et al., 1993; Gultneh et al., 1995; Canada-Vilalta et al., 2004). The two Mn ions in those complexes are bridged by oxo and dicarboxylate ligands, and the six-coordination around each Mn ion is completed by ligands such as water, CH₃CN and bipyridine (py). We report here the crystal structure of the title

compound, $[Mn_2(\mu-O)(hpc)_2(py)_4] \cdot H_2O$, (I), containing an $[Mn_2^{III}(\mu-O)]^{4+}$ core bridged by a 6-hydroxypicolinate ion (hpc^{2-}) .

Complex (I) was obtained during the course of our attempt to synthesize new multinuclear manganese complexes using 2,6-pyridinedicarboxylic acid (pdcH₂) and [Mn₁₂O₁₂(CH₃-COO)₁₆(H₂O)₄] (abbreviated to Mn₁₂-ac) in pyridine (see equation 1). In this reaction, the pdcH₂ ligand could not replace the acetate ions of Mn₁₂-ac but was oxidized to hpc²- (see equation 2). A possible mechanism may be oxidation of pdcH₂ followed by removal of carbon dioxide.

$$Mn_{12}-ac + pdcH_2 \xrightarrow{py} [Mn_2O(hpc)_2(py)_4]\cdot H_2O$$
 (1)

$$\frac{Mn_{12}\text{-ac}}{\text{pyridinc}} - OOO^{-}$$
(11)

Complex (I) consists of two hpc²⁻ ions, four pyridine rings and two Mn^{III} ions bridged by an oxo ion (Fig. 1 and Table 1). Each Mn ion is coordinated by two hpc²⁻ ions in bridging and chelating fashions, so that the molecule has a twofold axis perpendicular to the line connecting the two Mn^{III} ions and passing through the μ -oxo ion. The N1/C1/O1 amidate group of each hpc²⁻ ion bridges the two Mn^{III} ions, while the N1/C5/C6/O2 carboxypyridine group chelates to an Mn^{III} ion to form a five-membered ring. Coordination of two pyridine rings completes a distorted octahedral six-coordination around each Mn ion.

The Mn ions exhibit a marked Jahn–Teller (J–T) distortion, as expected for a high-spin d^4 Mn^{III} ion (Cotton *et al.*, 1999). The Mn1–O2 and Mn1–O4 bond distances [1.939 (2) and 1.789 (1) Å, respectively] are the shortest among the six Mn–N and Mn–O bonds, and the other four are all longer than 2.00 Å, indicating a compression type of J–T distortion.

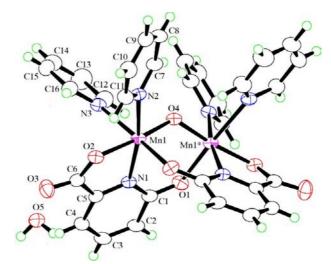


Figure 1 The molecular structure of (I), with the atom-numbering scheme, showing 50% probability displacement ellipsoids. The atom labelled with an asterisk (*) is at the symmetry position $(1 - x, y, \frac{1}{2} - z)$.

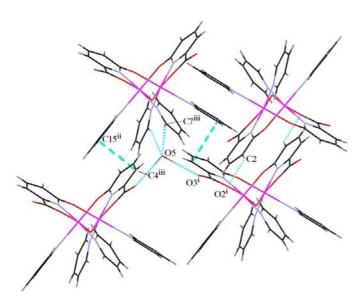


Figure 2 The molecular packing, showing intermolecular π - π interactions (dashed lines) and hydrogen bonds (dotted lines). [Symmetry codes: (i) x, -y, $z + \frac{1}{2}$; (ii) $x + \frac{1}{2}$, $\frac{1}{2} - y$, $z + \frac{1}{2}$; (iii) 2 - x, -y, 1 - z.]

Among nine complexes with a similar $[Mn_2(\mu-O)]^{4+}$ core structure, three show compressed J-T distortion; two of them (Wieghardt et al., 1985; Sheats et al., 1987) have tridentate capping ligands, such as Me₃TACN (Me₃TACN is N,N',N"trimethyl-1,4,7-triazacyclononane) or $HB(pz)_3$ [$HB(pz)_3$ is hydrotris(1-pyrazolyl)borate], and the other example (Vincent et al., 1993) has a strong coordination with an N₃⁻ ligand. In (I), the effect of the tridentate hpc²⁻ ion together with the strong coordination of the μ -oxo ligand probably induces the compressed J-T distortion. The Mn1-N1(hpc²⁻) distance [2.094 (2) Å] is shorter than the Mn1-N2/N3(py) bonds [2.169 (2) and 2.325 (2) Å], which is also attributable to the effect of the tridentate nature of the hpc²⁻ ion. The $Mn1 \cdots Mn1^{iv}$ [symmetry code: (iv) 1 - x, y, $\frac{1}{2} - z$] distance is 3.172 (1) Å, which is longer than the distances reported so far (3.14-3.16 Å) for complexes with the similar $[\text{Mn}_2(\mu\text{-O})]^{4+}$ core structure. This difference is due to the larger Mn-O- Mn^{iv} angle [125.0 (1)°] in (I).

The molecular packing is shown in Fig. 2. Water atom O5 lies on a twofold axis. Each complex molecule is connected to neighboring molecules via π – π interactions and hydrogen bonds (Table 2), resulting in a three-dimensional structure. The shortest $C \cdots C$ distance between the pyridine and hpc²-groups of neighboring complex molecules is 3.292 (4) Å for $C4 \cdots C15(\frac{3}{2}-x, y-\frac{1}{2},\frac{1}{2}-z)$, indicating a strong π – π interaction. The dihedral angle between these two planes is 17.28 (10)°. The absence of any counter-ions in (I) enables such intermolecular interactions.

Experimental

 $\rm Mn_{12}$ -ac was prepared according to a literature method (Lis, 1980). To a solution of $\rm Mn_{12}$ -ac (0.05 mmol) in pyridine (10 ml) was added pdcH₂ (0.4 mmol). After stirring for 30 min, the mixture was filtered and the dark-brown filtrate was left to stand, allowing slow

evaporation of the solvent at room temperature. Brown block-shaped crystals of (I) suitable for X-ray analysis were obtained after four weeks. IR (KBr pellet, cm⁻¹): 3198 (*b*), 3087 (*m*), 2923 (*m*), 1654 (*m*), 1632 (*m*), 1592 (*s*), 1570 (*m*), 1435 (*m*), 1396 (*m*), 1371 (*m*), 1278 (*w*), 1193 (*w*), 1015 (*w*), 910 (*w*), 785 (*w*), 720 (*m*), 699 (*w*), 661 (*w*).

Crystal data

$[Mn_2O(C_6H_3NO_3)_2(C_5H_5N)_4]\cdot H_2O$	Mo $K\alpha$ radiation
$M_r = 734.48$	Cell parameters from 4653
Monoclinic, $C2/c$	reflections
a = 13.606 (8) Å	$\theta = 3.1 - 27.5^{\circ}$
b = 16.379 (9) Å	$\mu = 0.88 \text{ mm}^{-1}$
c = 14.289 (9) Å	T = 150.2 K
$\beta = 104.118 (7)^{\circ}$	Block, brown
$V = 3088 (3) \text{ Å}^3$	$0.20 \times 0.15 \times 0.15 \text{ mm}$
Z = 4	
$D_x = 1.580 \text{ Mg m}^{-3}$	

Data collection

Rigaku/MSC Mercury CCD	3512 independent reflections
diffractometer	3144 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.034$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(Jacobson, 1998)	$h = -13 \rightarrow 17$
$T_{\min} = 0.764, T_{\max} = 0.876$	$k = -18 \rightarrow 21$
11 903 measured reflections	$l = -18 \rightarrow 18$

Refinement

J	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0468P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.050$	+ 5.1887 <i>P</i>]
$wR(F^2) = 0.109$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.12	$(\Delta/\sigma)_{\rm max} < 0.001$
3144 reflections	$\Delta \rho_{\text{max}} = 0.40 \text{ e Å}^{-3}$
260 parameters	$\Delta \rho_{\min} = -0.46 \text{ e Å}^{-3}$
Only coordinates of H atoms	
refined	

Table 1 Selected geometric parameters (Å, °).

$Mn1-Mn1^{iv}$	3.172 (1)	Mn1-N1	2.094(2)
$Mn1-O1^{iv}$	2.083 (2)	Mn1-N2	2.169(2)
Mn1-O2	1.939(2)	Mn1-N3	2.325 (2)
Mn1-O4	1.789 (1)		` ,
$O1^{iv}$ -Mn1-O2	90.95 (8)	O2-Mn1-N3	84.24 (8)
$O1^{iv}-Mn1-O4$	96.18 (8)	O4-Mn1-N1	97.59 (7)
$O1^{iv}-Mn1-N1$	93.10 (7)	O4-Mn1-N2	91.55 (8)
$O1^{iv}-Mn1-N2$	90.41 (7)	O4-Mn1-N3	88.71 (8)
$O1^{iv}-Mn1-N3$	173.97 (8)	N1-Mn1-N2	169.78 (9)
O2-Mn1-O4	172.80 (6)	N1-Mn1-N3	89.75 (7)
O2-Mn1-N1	80.98 (8)	N2-Mn1-N3	85.91 (7)
O2-Mn1-N2	89.37 (8)	$Mn1-O4-Mn1^{iv}$	125.0 (1)

Symmetry code: (iv) 1 - x, y, $\frac{1}{2} - z$.

Table 2 Hydrogen-bonding geometry (Å, °).

D $ H$ $\cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$\begin{array}{l} O5 - H14 \cdots O3^{iii} \\ C2 - H1 \cdots O2^{i} \\ C7 - H4 \cdots O5^{v} \end{array}$	0.87 (2)	1.88 (2)	2.748 (3)	178 (1)
	0.96 (3)	2.63 (2)	3.340 (3)	131 (2)
	0.92 (3)	2.38 (2)	3.137 (3)	140 (2)

Symmetry codes: (i) $x, -y, z + \frac{1}{5}$; (iii) 2 - x, -y, 1 - z; (v) $\frac{3}{2} - x, \frac{1}{2} - y, 1 - z$.

H atoms were located from difference density maps and their coordinates were refined, with $U_{\rm iso}({\rm H})=0.0146~{\rm Å}^2$. The C-H and O-H bond lengths are 0.80 (3)–1.00 (3) and 0.87 (2) Å, respectively.

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Data collection: CrystalClear (Rigaku, 2001); cell refinement: CrystalClear; data reduction: TEXSAN (Molecular Structure Corporation & Rigaku, 2000); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976) and MERCURY (Version 1.2.1; Bruno et al., 2002); software used to prepare material for publication: TEXSAN.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: OB1180). Services for accessing these data are described at the back of the journal.

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