metal-organic compounds

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catena-Poly[[[(1,10-phenanthroline- $\kappa^2 N, N'$)manganese(II)]- μ -L-tartrato- $\kappa^4 O^1, O^2: O^3, O^4$] hexahydrate]

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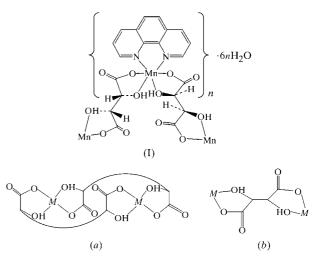
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The title compound, {[Mn(C₄H₄O₆)(C₁₂H₈N₂)]·6H₂O]_n, has a linear chain structure containing monomeric [Mn(C₄H₄O₆)-(C₁₂H₈N₂)] repeat units. Each manganese ion is six-coordinate, with the two phenanthroline N atoms [Mn-N = 2.229 (2) and 2.235 (2) Å] and four O atoms from two tartrate anions [Mn-O_{COO} = 2.1252 (19) and 2.1310 (19) Å, and Mn-O_{OH} = 2.2404 (19) and 2.2424 (19) Å] forming a seriously distorted octahedral coordination environment. Six water molecules exist outside every repeat unit as solvate molecules. Extensive hydrogen-bonding interactions and π - π stacking of the phenanthroline moieties exist between the chains.

Comment

The structures and properties of polymanganese complexes are currently of great interest on account of their promising applications in diverse areas of technology (Aubin et al., 1997; Manson et al., 1998) and as model complexes for the study of the photosynthetic oxygen-evolving complexes (OEC) of photosystem II in green plants (Wieghardt, 1989). L-Tartaric acid, in its role as a polydentate ligand, has been extensively studied in transition metal compounds. The coordination modes of L-tartrate are shown as (a) and (b) in the Scheme below, though mode (a) is seldom found. However, only two manganese-L-tartrate compounds have been reported (Ruiz-Pérez et al., 1996; Soylu, 1983) up to now. We report here the synthesis and crystal structure of the first mixed-ligand manganese-L-tartrate complex, viz. $\{[Mn(C_4H_4O_6)(C_{12}H_8N_2)]\}$ $6H_2O_n$ [C₄H₄O₆ is the dianion of L(+)-tartaric acid and $C_{12}H_8N_2$ is 1,10-phenanthroline], (I), which has a polymeric linear chain structure. A similar copper complex has been reported (McCann et al., 1997).

The crystal structure of (I) consists of an infinite $\{[Mn(C_4H_4O_6)(C_{12}H_8N_2)].6H_2O\}_n$ chain and the asymmetric unit is shown in Fig. 1. The Mn ion is located in a seriously distorted octahedral environment, with the three *trans* angles ranging from 155.82 (7) to 157.17 (9)°. The Mn ion is coordinated by two phenanthroline N atoms [Mn-N = 2.229 (2) and 2.235 (2) Å] and four O atoms [Mn-O = 2.1252 (19)-2.2424 (19) Å] from opposite ends of two tartrate ligands, which are symmetry related to form three chelating rings (symmetry code: x - 1, y, z). A tartrate anion links two neighboring Mn ions by a single ligand bridge, forming two chelating rings [form (*b*) in the *Scheme*]. Thus, Mn ions are bridged by tetradentate L-tartrate, forming a linear chain along the *a* axis of the cell. Six water molecules exist outside every repeat unit as solvate molecules.



The distortion of the coordination geometry is caused mainly by the three chelate five-membered rings. Bond lengths and angles for the L-tartrate ligand are usual and do not

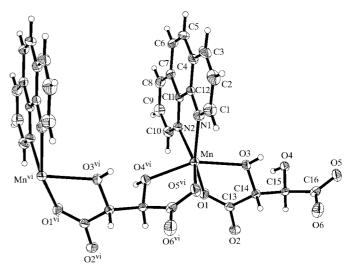
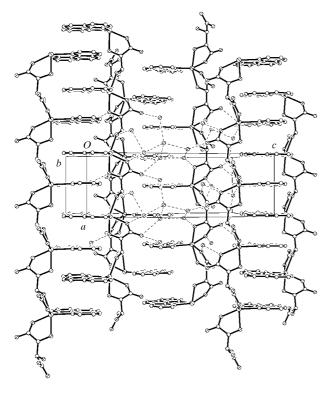
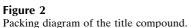


Figure 1

Part of the structure of (I), showing the bridging mode of the L-tartrate anion and one of the independent structural units with the atomnumbering scheme [symmetry code: (vi) x - 1, y, z].





deviate significantly from those in other reported L-tartrate coordination compounds (Ortega *et al.*, 1982; Ruiz-Pérez *et al.*, 1996). The average $Mn-O_{OH}$ distance [2.2414 (19) Å] is longer than that of $Mn-O_{COO}$ [2.1281 (19) Å], which indicates that the hydroxyl groups of the L-tartrate ligand are not deprotonated, analogous to the reaction of L-tartrate with other metal ions, such as Cu^{II} , Ni^{II} , Zn^{II} , Co^{II} and Cd^{II} . According to charge-balance requirements, the oxidation state of Mn is +2. In addition, the room-temperature magnetic moments of a powdered sample of (I) (5.89 BM) were in the range expected for a manganese(II) complex where there is no significant exchange interaction between metal centres.

Extensive hydrogen-bonding interactions are observed between the chains. The packing diagram of the complex along the *a* axis is presented in Fig. 2, showing three types of hydrogen-bonding interactions as follows: (i) O_{coord} to water [2.854 (3) and 2.889 (3) Å], (ii) $O_{uncoord}$ to water [2.709 (3)– 2.904 (3) Å] and (iii) water to water [2.775 (4)–2.832 (4) Å] (hydrogen-bond data are listed in Table 2). In addition, the phenanthroline groups of each polymeric chain are interwoven between the chains and form a π - π stacking along the *a* axis, with interplanar distances ranging from 3.22 to 3.35 Å.

Experimental

A mixture of Mn(CH₃COO)₂·4H₂O (0.25 g, 1.02 mmol), phen (2.0 g, 1 mmol), NaOCH₃ (0.11 g, 2 mmol) and L(+)-tartaric acid (0.15 g, 1 mmol) in MeOH/H₂O (16 ml, ν/ν , 1:1) was sealed in a 25 ml stainless-steel Teflon-lined reactor. The reaction system was heated at

Crystal data

$$\begin{split} & [\mathrm{Mn}(\mathrm{C_4H_4O_6})(\mathrm{C_{12}H_8N_2})]\cdot \mathrm{6H_2O} \\ & M_r = 491.31 \\ & \mathrm{Orthorhombic}, \ P2_12_12_1 \\ & a = 6.7093 \ (9) \ \mathrm{\mathring{A}} \\ & b = 15.336 \ (3) \ \mathrm{\mathring{A}} \\ & c = 20.210 \ (3) \ \mathrm{\mathring{A}} \\ & V = 2079.4 \ (6) \ \mathrm{\mathring{A}}^3 \\ & Z = 4 \\ & D_x = 1.569 \ \mathrm{Mg \ m^{-3}} \end{split}$$

Data collection

Siemens SMART CCD diffractometer φ and ω scans Absorption correction: empirical (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.777, T_{\max} = 0.869$ 6282 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.074$ S = 1.023372 reflections 337 parameters H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0341P)^2 + 1.1316P]$ where $P = (F_o^2 + 2F_c^2)/3$

Cell parameters from 140
reflections
$$\theta = 1.7-25.1^{\circ}$$

 $\mu = 0.70 \text{ mm}^{-1}$
 $T = 293 (2) \text{ K}$
Block, deep red
 $0.36 \times 0.24 \times 0.20 \text{ mm}$

Mo $K\alpha$ radiation

3372 independent reflections 3183 reflections with $I > 2\sigma(I)$ $R_{int} = 0.022$ $\theta_{max} = 25.1^{\circ}$ $h = -7 \rightarrow 7$ $k = -16 \rightarrow 18$ $l = -24 \rightarrow 10$

 $\begin{array}{l} (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.33 \ {\rm e} \ {\rm \AA}^{-3} \\ {\rm Extinction correction: $SHELXTL$} \\ ({\rm Bruker, 1997}) \\ {\rm Extinction coefficient: } 0.0119 \ (8) \\ {\rm Absolute structure: Flack (1983), } \\ 1242 \ {\rm Friedel pairs} \\ {\rm Flack parameter} = 0.01 \ (2) \end{array}$

Table 1

Selected geometric parameters (Å, °).

Mn-O5 ^{vi}	2.1252 (19)	Mn-N2	2.235 (2)
Mn-O1	2.1310 (19)	Mn-O3	2.2404 (19)
Mn-N1	2.229 (2)	$Mn-O4^{vi}$	2.2424 (19)
O5 ^{vi} -Mn-O1	103.65 (8)	N1-Mn-O3	90.36 (8)
O5 ^{vi} -Mn-N1	93.33 (8)	N2-Mn-O3	108.58 (8)
O1-Mn-N1	155.84 (9)	O5 ^{vi} -Mn-O4 ^{vi}	73.36 (7)
O5 ^{vi} -Mn-N2	157.17 (9)	O1-Mn-O4 ^{vi}	93.42 (7)
O1-Mn-N2	94.25 (8)	N1-Mn-O4 ^{vi}	108.02 (8)
N1-Mn-N2	74.56 (8)	N2-Mn-O4 ^{vi}	91.76 (8)
O5 ^{vi} -Mn-O3	90.44 (8)	O3-Mn-O4 ^{vi}	155.82 (7)
O1-Mn-O3	72.69 (7)		
O5 ^{vi} -Mn-N1-C1	19.4 (3)	O5 ^{vi} -Mn-N2-C11	61.1 (3)
$O4^{vi}$ -Mn-N1-C1	93.0 (3)	N1-Mn-N2-C11	1.1 (2)
O1-Mn-N1-C12	62.3 (3)	O5 ^{vi} -Mn-O3-C14	77.46 (18)
O3-Mn-N1-C12	106.9 (2)	O4 ^{vi} -Mn-O3-C14	30.5 (3)

Symmetry code: (vi) x - 1, y, z.

H atoms bonded to C atoms were placed at calculated idealized positions using a riding model (C–H = 0.93 Å for 1,10-phenanthroline and 0.98 Å for L-tartrate). H atoms bonded to O atoms were located in a difference Fourier synthesis and were subsequently refined with restrained O–H distances approximately equal to 0.85 Å.

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O8-H8B\cdots O11^{i}$	0.85 (3)	1.92 (6)	2.775 (4)	168 (6)
$O9-H9B\cdots O6^{ii}$	0.834 (19)	1.88 (2)	2.709 (3)	173 (4)
$O10-H10B\cdots O8^{iii}$	0.85 (2)	2.07 (3)	2.820 (4)	147 (4)
$O9-H9C\cdots O7^{iv}$	0.85 (2)	1.95 (2)	2.782 (4)	168 (5)
$O10-H10C\cdots O12^{v}$	0.837 (19)	2.03 (2)	2.832 (4)	160 (4)
$O11-H11C\cdots O8^{vi}$	0.86 (3)	1.92 (3)	2.775 (5)	169 (6)
$O7 - H7C \cdot \cdot \cdot O5^{ii}$	0.840 (19)	2.02 (2)	2.854 (3)	170 (5)
$O11 - H11B \cdots O7$	0.86 (4)	1.96 (5)	2.790 (4)	159 (4)
$O7 - H7B \cdots O1$	0.845 (19)	2.07 (2)	2.889 (3)	163 (4)
O12−H12B···O11	0.88 (5)	1.94 (5)	2.817 (4)	174 (5)
O8−H8C···O2	0.86 (6)	1.94 (6)	2.794 (4)	173 (6)
O12−H12C···O2	0.85(4)	2.06(2)	2.904 (3)	173 (4)

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

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1994); data reduction: *XPREP* in *SHELXTL* (Siemens, 1994); program(s) used to solve structure: *SHELXTL* (Bruker, 1997); program(s) used to refine structure: *SHELXTL* (Bruker, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL* (Bruker, 1997).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1417). Services for accessing these data are described at the back of the journal.

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