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Key indicators

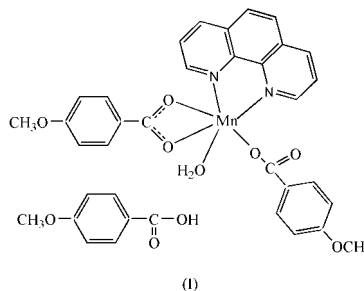
Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$
 R factor = 0.051
 wR factor = 0.127
Data-to-parameter ratio = 12.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Aquabis(*p*-methoxybenzoato)(1,10-phenanthroline)manganese(II) *p*-methoxybenzoic acid

The title compound, $[\text{Mn}(\text{C}_8\text{H}_7\text{O}_3)_2(\text{C}_{12}\text{H}_8\text{N}_2)(\text{H}_2\text{O})] \cdot \text{C}_8\text{H}_8\text{O}_3$, has been synthesized by the reaction of $\text{MnCl}_2 \cdot 6\text{H}_2\text{O}$ with *p*-anisic acid and 1,10-phenanthroline in an alcohol–water solution. The Mn atom is six-coordinated by two N atoms and four O atoms, forming a distorted octahedral geometry. The Mn–O and Mn–N distances range from 2.113 (3) to 2.410 (3) Å.

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Comment

In order to study the role of manganese ions and the structures of the active sites in manganese enzymes, small-molecule complexes of Mn are often used as model systems (Shi *et al.*, 2000). In recent years, Mn–phen (phen is 1,10-phenanthroline) complexes have been studied extensively as model compounds for simulating the oxygen-evolving center (OEC) of photosystem II (PSII) (Manchanda *et al.*, 1994). In this paper, we report the synthesis and structure of aquabis(*p*-methoxybenzoato)(1,10-phenanthroline)manganese(II) *p*-methoxybenzoic acid, (I) (Fig. 1).



The Mn atom in (I) is located in a very distorted octahedral environment containing two N atoms from a chelating phen ligand, three O atoms from monodentate and bidentate carboxylate groups and one O atom from a water molecule. The Mn–O bond lengths (Table 1) in (I) are in the range 2.118 (2)–2.324 (2) Å, and the Mn–N bond lengths are 2.254 (2) and 2.246 (2) Å. The phenanthroline ligand exhibits the usual short N···N bite distance [N1···N2 = 2.696 (2) Å] and acute N–Mn–N angle [N1–Mn1–N2 = 73.61 (8)°]; these are very close to those found in $[\text{Mn}(\text{phen})_2\text{Cl}_2]$ (McCann *et al.*, 1998), $[\text{Mn}(\text{phen})_2(\text{NCS})_2]$ (Holleman *et al.*, 1994) and $[\text{Mn}(\text{phen})_2(\text{N}_3)_2]$ (Shen *et al.*, 1999). The dihedral angles between the chelating phen plane and chelating carboxylate planes are 106.1 (3) and 73.9 (2)°. There exist strong O–H···O hydrogen bonds between the carboxylic acid group of *p*-methoxybenzoic acid and the water molecule (Table 2). These hydrogen bonds produce dimeric clusters (Fig. 2). The dimers interact by way of π – π stacking interactions in the sequence benz–phen–phen–benz (Fig. 3), with the neighbouring phen planes separated by 3.447 Å.

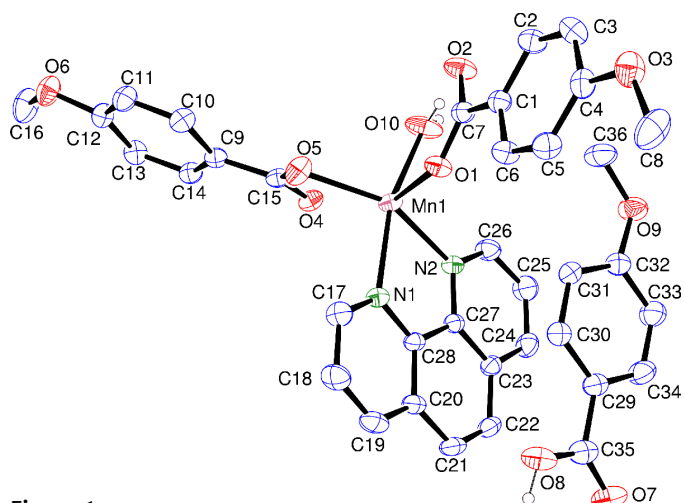


Figure 1
View of (I), showing 30% probability displacement ellipsoids. H atoms attached to C have been omitted for clarity.

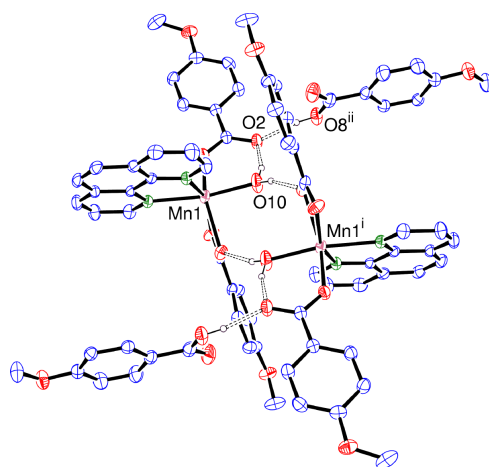


Figure 2
Hydrogen bonding (dashed lines) in the dimeric cluster of (I) (30% probability displacement ellipsoids). [Symmetry codes: (i) $2 - x, 1 - y, z$; (ii) $1 + x, y, z$.]

Experimental

The reaction of *p*-anisic acid (0.456 g, 3 mmol) and NaOH (0.12 g, 3 mmol) in a 1:1 molar ratio in a water–alcohol (3:1) solution (40 ml) at room temperature for 30 min produced a colorless solution, to which $\text{MnCl}_2 \cdot 6\text{H}_2\text{O}$ (0.198 g, 1 mmol) and 1,10-phenanthroline (0.198 g, 1 mmol) were added, yielding a yellow solution. The reaction solution was kept at room temperature after stirring for an hour. Yellow crystals were obtained after a few days; they were washed with water and ethanol several times and then dried in air.

Crystal data

$[\text{Mn}(\text{C}_8\text{H}_7\text{O}_3)_2(\text{C}_{12}\text{H}_8\text{N}_2) \cdot (\text{H}_2\text{O})] \cdot \text{C}_8\text{H}_8\text{O}_3$	$Z = 2$
$M_r = 707.58$	$D_x = 1.431 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 10.7353(6) \text{ \AA}$	Cell parameters from 5737 reflections
$b = 11.3832(2) \text{ \AA}$	$\theta = 2.0\text{--}25.0^\circ$
$c = 16.0107(1) \text{ \AA}$	$\mu = 0.47 \text{ mm}^{-1}$
$\alpha = 76.41(3)^\circ$	$T = 293(2) \text{ K}$
$\beta = 71.56(2)^\circ$	Block, yellow
$\gamma = 62.855(18)^\circ$	$0.55 \times 0.40 \times 0.30 \text{ mm}$
$V = 1642.1(3) \text{ \AA}^3$	

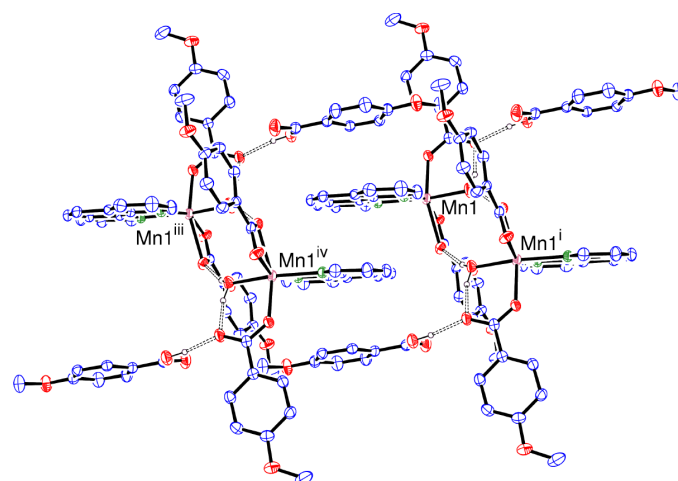


Figure 3
Diagram showing the π - π stacking interactions in (I) (30% probability displacement ellipsoids). [Symmetry codes: (i) $2 - x, 1 - y, z$; (iii) $-x, 1 - y, z$; (iv) $1 - x, 1 - y, z$.]

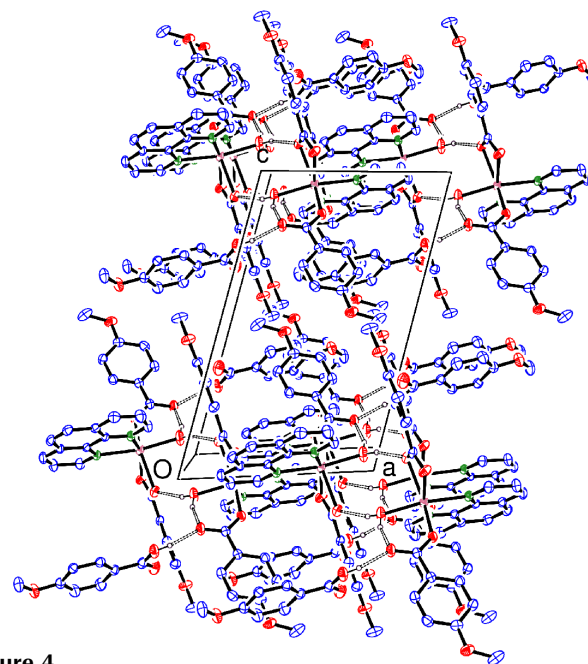


Figure 4
Packing diagram of (I), viewed down $[010]$, showing hydrogen bonds as dashed lines (30% probability displacement ellipsoids).

Data collection

Mercury diffractometer	4803 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.024$
Absorption correction: multi-scan	$\theta_{\text{max}} = 25.0^\circ$
(CrystalClear; Rigaku, 2000)	$h = -12 \rightarrow 12$
$T_{\text{min}} = 0.784, T_{\text{max}} = 0.873$	$k = -13 \rightarrow 13$
10 120 measured reflections	$l = -19 \rightarrow 17$
5737 independent reflections	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.062P)^2 + 0.3137P]$
$R[F^2 > 2\sigma(F^2)] = 0.051$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.127$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.10$	$\Delta\rho_{\text{max}} = 0.26 \text{ e \AA}^{-3}$
5737 reflections	$\Delta\rho_{\text{min}} = -0.24 \text{ e \AA}^{-3}$
454 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1
Selected geometric parameters (Å).

Mn1—O1	2.118 (2)	Mn1—N2	2.246 (2)
Mn1—O10	2.156 (2)	Mn1—N1	2.254 (2)
Mn1—O5	2.181 (2)	Mn1—O4	2.324 (2)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O10—H10B \cdots O2	0.84 (4)	1.85 (2)	2.649 (3)	159 (4)
O10—H10C \cdots O4 ⁱ	0.84 (4)	1.91 (2)	2.726 (3)	165 (4)
O8—H8D \cdots O2 ^v	0.85 (4)	1.77 (2)	2.615 (3)	171 (4)

Symmetry codes: (i) $2-x, 1-y, -z$; (v) $x-1, y, z$.

Hydroxyl H atoms were located in difference maps and freely refined, while H atoms attached to C were placed in idealized positions ($C-H = 0.93-0.96$ Å) and refined as riding, with $U_{iso} = 1.2$ or $1.5U_{eq}$ of the parent atom.

Data collection: *Crystal Clear* (Rigaku, 2000); cell refinement: *Crystal Clear*; data reduction: *Crystal Clear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXL97*; software used to prepare material for publication: *SHELXL97*.

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