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Key indicators

Single-crystal X-ray study  
 T = 295 K  
 Mean  $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$   
 R factor = 0.030  
 wR factor = 0.084  
 Data-to-parameter ratio = 13.2

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

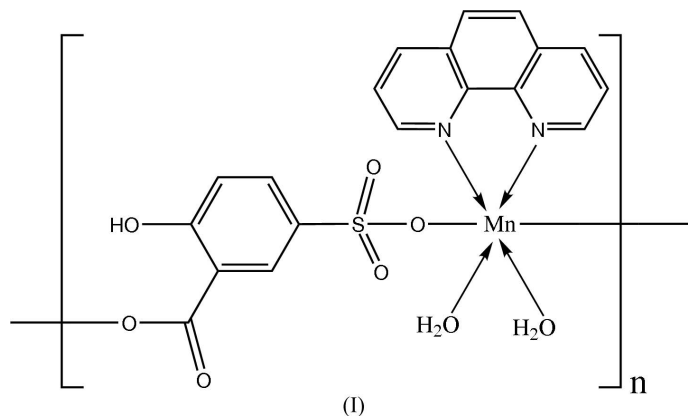
**catena-Poly[[*cis*-diaqua(1,10-phenanthroline- $\kappa^2N,N'$ )-  
 manganese(II)]- $\mu_2$ -5-sulfonatosalicylato- $\kappa^2O:O'$ ]**

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The title complex,  $[\text{Mn}(\text{C}_7\text{H}_4\text{O}_6\text{S})(\text{C}_{12}\text{H}_8\text{N}_2)(\text{H}_2\text{O})_2]_n$ , comprises a one-dimensional chain in which each Mn atom displays a distorted octahedral geometry. Hydrogen-bonding interactions between chains generate a two-dimensional architecture.

Comment

Recently, much attention has been paid to metal complexes of 5-sulfosalicylic acid ( $\text{H}_3\text{ssal}$ ) because of the structural diversity and biological interest of this ligand (Fan, Cai *et al.*, 2005; Gao *et al.*, 2005; Fan, Zhu *et al.*, 2005a). In our previous work on the manganese(II) 1,10-phenanthroline (phen) complex  $[\text{Mn}(\text{phen})_2(\text{H}_2\text{O})_2](\text{Hssal}) \cdot 4\text{H}_2\text{O}$ , (II), the doubly deprotonated 5-sulfosalicylate does not coordinate to the metal atom (Fan, Zhu *et al.*, 2005b), partly because the phen ligand can strongly chelate to the metal atom. Therefore, a new synthetic procedure, a two-step reaction method, was introduced to the  $\text{Mn}^{2+}/\text{H}_3\text{ssal}/\text{phen}$  system and, as expected, the  $\text{Hssal}^{2-}$  coordinated complex, (I), was synthesized.



In the title complex, the  $\text{Mn}^{\text{II}}$  atom adopts a distorted octahedral geometry defined by two N donors from one phen ligand, two O atoms from two *cis* water molecules and two O atoms, one from a sulfonyl and one from a carboxyl group of two different  $\text{Hssal}^{2-}$  ligands (Fig. 1 and Table 1). The  $\text{Mn}-\text{N}$  and  $\text{Mn}-\text{O}_w$  distances are similar to those of (II). The 5-sulfonatosalicylate ligand acts as a linker using one sulfonyl and one carboxyl O atom to extend the structure into a one-dimensional chain (Fig. 2). Hydroxy and uncoordinated carboxyl O atoms form intrachain hydrogen bonds with a carboxyl group and a water molecule, respectively. Between chains, water molecules and sulfonyl O atoms form intermolecular hydrogen bonds, generating a two-dimensional hydrogen-bonding network (Fig. 3 and Table 2).

Comparison of (II),  $[\text{Mn}(\text{phen})_2(\text{H}_2\text{O})_2]^{2+}$ , with (I),  $[\text{Mn}(\text{phen})(\text{H}_2\text{O})_2(\text{Hssal})]$ , shows that one phen ligand in II is replaced by an  $\text{Hssal}^{2-}$  ligand in (I). Two such diverse structures were constructed by different synthetic procedures. For (II), a one-pot synthesis was used, while for (I), a two-step reaction method was used, *i.e.* phen was added after the metal salt had completely reacted with 5-sulfosalicylic acid in the presence of a weak base. Therefore, multi-step reactions are very useful in the construction of different architectures.

## Experimental

A mixture of  $\text{MnCl}_2 \cdot 4\text{H}_2\text{O}$  (0.199 g, 1.0 mmol), 5-sulfosalicylic acid dihydrate (0.050 g, 0.20 mmol), and 4% pyridine (2 ml) in water (20 ml) was stirred for 17 h. 1,10-Phenanthroline (0.040 g, 0.20 mmol) was then added. The resulting solution was put aside and the solvent was allowed to evaporate at room temperature. After three weeks, pale-yellow block-shaped crystals were obtained.

### Crystal data

$[\text{Mn}(\text{C}_7\text{H}_4\text{O}_6\text{S})(\text{C}_{12}\text{H}_8\text{N}_2)(\text{H}_2\text{O})_2]$   
 $M_r = 487.34$   
 Monoclinic,  $P2_1/c$   
 $a = 14.4166$  (9) Å  
 $b = 7.7234$  (5) Å  
 $c = 18.6868$  (11) Å  
 $\beta = 107.421$  (1)°  
 $V = 1985.2$  (2) Å<sup>3</sup>  
 $Z = 4$

$D_x = 1.631$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 5871 reflections  
 $\theta = 2.3$ – $28.2$ °  
 $\mu = 0.82$  mm<sup>-1</sup>  
 $T = 295$  (2) K  
 Block, pale yellow  
 $0.43 \times 0.25 \times 0.16$  mm

### Data collection

Bruker SMART APEX area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2002)  
 $T_{\min} = 0.719$ ,  $T_{\max} = 0.880$   
 10821 measured reflections

3888 independent reflections  
 3537 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.019$   
 $\theta_{\text{max}} = 26.0$ °  
 $h = -14 \rightarrow 17$   
 $k = -9 \rightarrow 9$   
 $l = -23 \rightarrow 13$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.030$   
 $wR(F^2) = 0.084$   
 $S = 1.07$   
 3888 reflections  
 295 parameters  
 H-atom parameters constrained

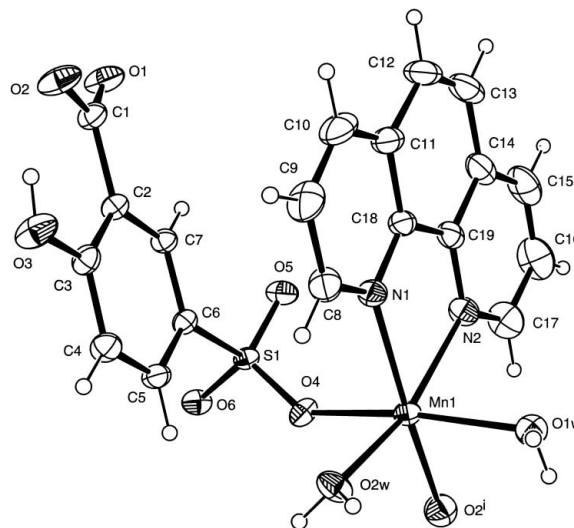
$w = 1/[\sigma^2(F_o^2) + (0.0466P)^2 + 0.5664P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.24$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.36$  e Å<sup>-3</sup>

**Table 1**

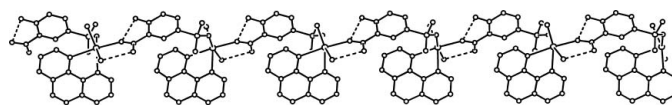
Selected geometric parameters (Å, °).

|                          |             |             |             |
|--------------------------|-------------|-------------|-------------|
| Mn1–O1w                  | 2.2316 (14) | Mn1–N2      | 2.2447 (15) |
| Mn1–O2w                  | 2.1255 (14) | S1–O4       | 1.4595 (12) |
| Mn1–O2 <sup>i</sup>      | 2.1174 (13) | S1–O5       | 1.4479 (13) |
| Mn1–O4                   | 2.1835 (12) | S1–O6       | 1.4627 (12) |
| Mn1–N1                   | 2.2659 (15) |             |             |
| O2 <sup>i</sup> –Mn1–O1w | 88.61 (5)   | O2w–Mn1–O1w | 94.12 (5)   |
| O2 <sup>i</sup> –Mn1–O2w | 97.59 (6)   | O2w–Mn1–N1  | 88.91 (6)   |
| O2 <sup>i</sup> –Mn1–O4  | 81.49 (5)   | O2w–Mn1–N2  | 162.63 (6)  |
| O2 <sup>i</sup> –Mn1–N1  | 173.00 (6)  | N2–Mn1–N1   | 73.76 (6)   |
| O2 <sup>i</sup> –Mn1–N2  | 99.68 (6)   | O5–S1–O4    | 113.94 (8)  |
| O4–Mn1–O1w               | 169.86 (5)  | O5–S1–O6    | 112.67 (8)  |
| O4–Mn1–N1                | 96.03 (5)   | O4–S1–O6    | 109.13 (7)  |
| O4–Mn1–N2                | 91.36 (5)   | O5–S1–C6    | 107.69 (8)  |
| O1w–Mn1–N1               | 93.55 (5)   | O4–S1–C6    | 105.45 (7)  |
| O1w–Mn1–N2               | 88.13 (6)   | O6–S1–C6    | 107.50 (8)  |
| O2w–Mn1–O4               | 89.39 (5)   |             |             |

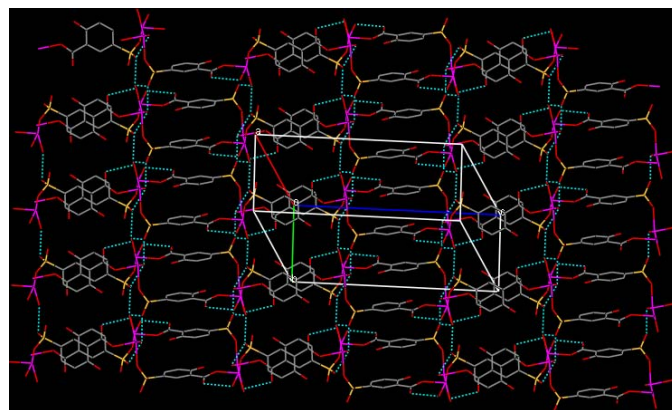
Symmetry code: (i)  $x, -y + \frac{1}{2}, z - \frac{1}{2}$ .



**Figure 1**  
 An ORTEP-3 (Farrugia, 1997) view of a segment of (I). Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code: (i)  $x, \frac{1}{2} - y, z - \frac{1}{2}$ ]



**Figure 2**  
 A view of the one-dimensional chain in (I). Hydrogen bonds are shown as dashed lines and H atoms have been omitted for clarity.



**Figure 3**  
 A view of the two-dimensional hydrogen-bonding (dashed lines) network for (I). The phen ligand and H atoms have been omitted for clarity.

**Table 2**

Hydrogen-bond geometry (Å, °).

| $D-H \cdots A$                     | $D-H$    | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|------------------------------------|----------|--------------|--------------|----------------|
| O2w–H1B $\cdots$ O6 <sup>ii</sup>  | 0.84 (1) | 1.89 (1)     | 2.7214 (18)  | 169 (2)        |
| O1w–H1A $\cdots$ O5 <sup>iii</sup> | 0.84 (1) | 2.01 (1)     | 2.8233 (18)  | 163 (2)        |
| O1w–H2A $\cdots$ O1 <sup>i</sup>   | 0.85 (1) | 1.90 (1)     | 2.7076 (19)  | 158 (2)        |
| O2w–H2B $\cdots$ O6 <sup>iii</sup> | 0.84 (1) | 1.88 (1)     | 2.7127 (17)  | 173 (2)        |
| O3–H3A $\cdots$ O2                 | 0.84 (1) | 1.80 (1)     | 2.578 (2)    | 152 (2)        |

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

The aromatic H atoms were positioned geometrically, and were included in the refinement in the riding-model approximation [C—

$H = 0.93 \text{ \AA}$  and  $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$ . The water and hydroxy H atoms were located in difference Fourier maps and were refined with a distance restraint of  $O-H = 0.85(1) \text{ \AA}$  and fixed isotropic displacement parameters of  $U_{\text{iso}}(H) = 0.05 \text{ \AA}^2$ .

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINTE* (Bruker, 2002); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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