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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma$ (C–C) = 0.007 Å R factor = 0.054 wR factor = 0.131 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.

# catena-Poly[[aquamanganese(II)]-di-μ-4-pyridylthioacetato-κ<sup>6</sup>O,O':N;N:O,O']

The Mn atom and the coordinated water molecule in the title compound,  $[Mn(C_7H_6NO_2S)_2(H_2O)]_n$ , lie on a twofold axis; the Mn atom is chelated by the carboxyl –CO<sub>2</sub> units and is also coordinated by the pyridyl N atoms of two adjacent anionic groups in a seven-coordinate *trans*-pentagonal bipyramidal geometry. The polymeric chain runs in a zigzag manner along the *c* axis, and neighboring chains are linked into a hydrogenbonded layer structure.

#### Comment

The reaction of a divalent transition metal ion with the anion of 4-pyridylthioacetic acid affords different products depending on the reaction conditions and the nature of the metal ion. Under hydrothermal conditions, the reaction with  $Zn^{2+}$  affords polymeric  $[Zn(C_7H_6NO_2S)_2]_n$  in which the metal atom is four-coordinate in a tetrahedral environment, the metal atom being linked to two N and two O atoms (Zhang et al., 2003). On the other hand, the Ni<sup>II</sup> derivative exists as a zwitterionic compound,  $[Ni(C_7H_6NO_2S)_2(H_2O)_4]$ , in which the metal atom is coordinated by four water molecules and the pyridyl N atom of the anionic group; the metal atom exists in an all-trans octahedral geometry (Zhang et al., 2004). In the present study, the Mn<sup>II</sup> derivative is coordinated by only one water molecule, but the coordination number is seven as the two carboxyl  $-CO_2$  units behave in a chelating mode; the donor set is completed by the pyridyl N atoms of two adjacent anions (Fig. 1) that occupy the apical sites of the pentagonal bipyramidal polyhedron. Bridging gives rise to the formation of a linear zigzag chain that runs along the c axis (Fig. 2); adjacent chains are linked by a hydrogen bond  $[O1w \cdots O1^{i} =$ 2.781 (4) Å; symmetry code: (i) -x, y - 1,  $\frac{1}{2} - z$ ] into layers.



Seven-coordinate Mn complexes are relatively less common than six-coordinate Mn complexes; in fact, there appears to be only one example of an  $MnN_2O_5$  fragment in which four of the O atoms belong to a pair of carboxyl  $-CO_2$  units. The dinuc-

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## Figure 1

*ORTEPII* (Johnson, 1976) plot of a fragment of the  $[Mn(C_7H_6NO_2S)_2(H_2O)]_n$  chain, with displacement ellipsoids drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.



Figure 2

ORTEPII (Johnson, 1976) plot of the pyridyl-bridged zigzag  $[Mn(C_7H_6NO_2S)_2(H_2O)]_n$  chain. The chains are linked by hydrogen bonds into layers.

lear 2,2'-bipyridine-chelated bicyclo[2.2.1]hept-5-ene-2-*exo*,3*exo*-dicarboxylate also features a coordinated water molecule  $[Mn-O_{water} = 2.208 (1) \text{ Å}]$ ; the N atoms of the chelating ligand occupy adjacent sites of the pentagonal plane (Baumeister & Hartung, 1997).

# **Experimental**

A mixture of manganese acetate tetrahydrate (0.25 g, 1 mmol), 4pyridylthioacetic acid (0.20 g, 1.2 mmol) and imidazole (0.03, 0.5 mmol) in water (7 ml) was treated with several drops of 2 N sodium hydroxide to a pH of approximately 6. The solution was placed in a 15 ml Teflon-lined stainless-steel bomb, which was heated at 433 K for 96 h. Colorless crystals of the title compound were obtained in about 40% yield; the imidazole component was not incorporated into the molecule.

## Crystal data

$[Mn(C_7H_6NO_2S)_2(H_2O)]$	$D_x = 1.693 \text{ Mg m}^{-3}$
$M_r = 409.33$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 950
a = 21.595 (3) Å	reflections
b = 6.4904 (8)  Å	$\theta = 2.5-24.7^{\circ}$
c = 15.611 (2)  Å	$\mu = 1.11 \text{ mm}^{-1}$
$\beta = 132.774 \ (1)^{\circ}$	T = 298 (2)  K
$V = 1606.1 (3) \text{ Å}^3$	Block, colorless
Z = 4	$0.09\times0.08\times0.06~\text{mm}$

#### Data collection

Bruker SMART APEX area- detector diffractometer $\varphi$ and $\omega$ scans Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 1996) $T_{\min} = 0.525, T_{\max} = 0.936$	1692 independent reflections 1419 reflections with $I > 2\sigma(I)$ $R_{int} = 0.034$ $\theta_{max} = 27.0^{\circ}$ $h = -27 \rightarrow 27$ $k = -8 \rightarrow 5$
4507 measured reflections	$l = -19 \rightarrow 19$
Refinement	
Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.054$ $wR(F^2) = 0.131$ S = 1.12 1692 reflections 111 parameters H-atom parameters constrained	$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0629P)^2 \\ &+ 0.9714P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.60 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.38 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$

# Table 1

Selected geometric parameters (Å, °).

Mn1-O1	2.378 (3)	Mn1-N1 <sup>i</sup>	2.276 (3)
Mn1-O2	2.280 (2)	Mn1-N1 <sup>ii</sup>	2.276 (3)
Mn1 - O1w	2.218 (4)		
O1-Mn1-O1 <sup>iii</sup>	74.5 (1)	O2-Mn1-O2 <sup>iii</sup>	173.3 (1)
O1-Mn1-O2	56.1 (1)	O2-Mn1-O1w	86.7 (1)
O1-Mn1-O2 <sup>iii</sup>	130.6 (1)	O2-Mn1-N1 <sup>i</sup>	90.1 (1)
O1 - Mn1 - O1w	142.8 (1)	O2-Mn1-N1 <sup>ii</sup>	90.2 (1)
O1-Mn1-N1 <sup>i</sup>	88.0 (1)	$O1w-Mn1-N1^{i}$	92.6 (1)
O1-Mn1-N1 <sup>ii</sup>	87.9 (1)	N1 <sup>i</sup> -Mn1-N1 <sup>ii</sup>	174.9 (2)

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

A dimensionless value  $\mu \times 2r = 0.10$  was used in the absorption correction. The H atoms were placed at calculated positions in the riding-model approximation, with C-H = 0.98 Å for aliphatic H atoms, C-H = 0.93 Å for aromatic H atoms and O-H = 0.82 Å; the displacement parameters were tied to those  $(U_{eq})$  of the parent atoms by a factor of 1.2. The H atom belonging to the O1w water molecule, which lies on the twofold axis, was generated by the HFIX 147 instruction in *SHELXL97* (Sheldrick, 1997).

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL*97.

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