Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Deguang Huang, Xiaofeng Zhang, Hongping Zhu, Changneng Chen and Qiutian Liu*

State Key Laboratory of Structural Chemistry, Fujian Institute of Research on the Structure of Matter, Fuzhou, Fujian 350002, People's Republic of China

Correspondence e-mail: lqt@ms.fjirsm.ac.cn

Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.005 \text{ Å}$ R factor = 0.037 wR factor = 0.130 Data-to-parameter ratio = 11.8

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

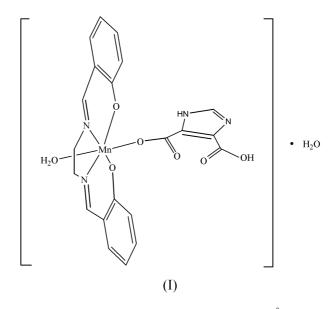
Aqua(N,N'-disalicylideneethylenediamino-N,N',O,O')(4,5-imidazoledicarboxylato-O)-manganese(III) monohydrate

In the title compound, $[Mn^{III}(salen)(OCOC_3H_2N_2COOH)-(H_2O)]\cdotH_2O$, $(H_2salen = N,N'$ -disalicylideneethylenediamine, $C_{16}H_{14}N_2O_2$), two N and two O atoms from salen chelate to the Mn^{III} ion, forming a distorted octahedral coordination sphere together with two other axial O atoms from a carboxyl group and a water molecule. Extended intermolecular hydrogen bonds are observed to link the complex molecules, forming a one-dimensional chain structure.

Received 13 August 2001 Accepted 10 September 2001 Online 20 September 2001

Comment

The imidazole N atom of histidine has been described as a functional group (Wieghardt, 1989) in the oxygen-evolving complex (EOC) of photosystem II (PS II). 4,5-Imidazoledicarboxylate was suggested as a bridging ligand (Rypniewski *et al.*, 1995; Tainer *et al.*, 1982) and a dinuclear Mn^{IV} cluster in 4,5-imidazoledicarboxylate-bridged dimanganese(IV) has been reported (Caudle *et al.*, 1997). It is interesting to study the differing coordination behavior of 4,5-dicarboxyimidazole. We present here a new complex, [Mn(salen)(OCO- $C_3H_2N_2COOH$)(H_2O)]· H_2O , (I), in which 4,5-imidazoledicarboxylate coordinates to Mn^{III} as a terminal ligand. The structure of (I) is shown in Fig. 1. The Mn atom is chelated by two N [Mn–N1 1.991 (2) Å and Mn–N2 1.989 (2) Å] and two O atoms [Mn–O1 1.8793 (18) Å and Mn–O2 1.8944 (18) Å] from the salen ligand to form an equatorial



plane with a maximum deviation of 0.0052 (10) Å from the least-squares plane, while the opposite axial positions are occupied by another two O atoms from the 4,5-dicarboxyl-atoimidazole [Mn-O4 2.1815 (19) Å] and the ligated aqua

 \odot 2001 International Union of Crystallography Printed in Great Britain – all rights reserved

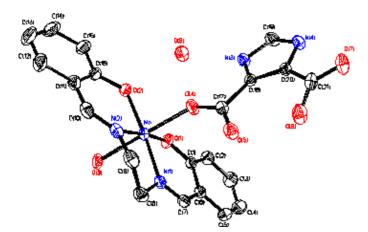


Figure 1

Diagram (*SHELXTL*; Bruker, 1998) of (I) showing the atom-numbering scheme and displacement ellipsoids at the 30% probability level. H atoms have been omitted for clarity.

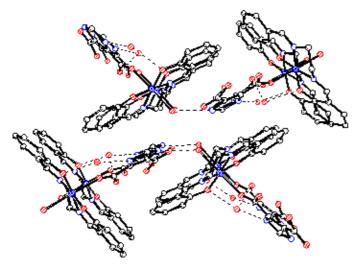


Figure 2 A packing diagram of the title compound.

[Mn–O3 2.249 (2) Å]. The Mn site is surrounded by a distorted octahedral coordination sphere, with the *trans* angles ranging from 172.83 (9) to 178.04 (9)° and the other angles ranging from N2–Mn–N1 of 82.50 (9)° to O2–Mn–O1 of 94.69 (8)°. All the bond lengths and angles are comparable to those of other six-coordinate Mn^{III} Schiff base complexes (Shyu *et al.*, 1999; Watkinson *et al.*, 1999; Oshio *et al.*, 1995), while the Mn–O bonds of (I) are longer than those reported by Caudle *et al.* (1997), where Mn possesses a 4+ oxidation state, higher than that in (I).

A packing diagram of the title compound is presented in Fig. 2 and shows the extensive intermolecular hydrogenbonding interactions, which contain $N_{imidazo} \cdots H - O_{aqua}$ [2.992 (3) Å], $O_{carboxy} \cdots H - O_{aqua}$ [2.880 (3) Å] and $N_{imidazo} \cdots H - O_{aqua}$ [2.861 (3) Å]. These hydrogen bonds produce a one-dimensional polymeric structure.

Experimental

To a solution of 0.268 g (1 mmol) of H₂salen in 40 ml of CH₂Cl₂/MeCN (ν/ν , 1:1), 0.156 g (1 mmol) of 4,5-dicarboxyimidazole was added and the resulting solution stirred. The reaction solution was refluxed for 18 h, accompanied by the addition of 0.245 g (1 mmol) of Mn(CH₃COO)₂.4H₂O. After filtration, the filtrate was allowed to stand for several days at room temperature to deposit dark-red crystals of (I), which were recrystallized from ethanol solution to obtain the crystals suitable for X-ray diffraction.

Crystal data

$$\begin{split} & [\mathrm{Mn}(\mathrm{C}_{5}\mathrm{H}_{3}\mathrm{N}_{2}\mathrm{O}_{4})(\mathrm{C}_{16}\mathrm{H}_{14}\mathrm{N}_{2}\mathrm{O}_{2}) - \\ & (\mathrm{H}_{2}\mathrm{O})] \cdot \mathrm{H}_{2}\mathrm{O} \\ & M_{r} = 512.36 \end{split}$$

Monoclinic, $P2_1/n$ a = 7.009 (3) Å b = 19.8634 (4) Å c = 16.2811 (3) Å $\beta = 101.422$ (1)° V = 2221.93 (6) Å³ Z = 4

Data collection

Siemens SMART CCD diffractometer ω scans Absorption correction: empirical (*SADABS*; Sheldrick, 1996) $T_{min} = 0.611, T_{max} = 0.802$ 7277 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.130$ S = 0.973856 reflections 327 parameters $D_x = 1.532 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 4856 reflections $\theta = 1.6-25.0^{\circ}$ $\mu = 0.65 \text{ mm}^{-1}$ T = 293 (2) KBlock, dark-red $0.76 \times 0.54 \times 0.34 \text{ mm}$

3856 independent reflections 3101 reflections with $I > 2\sigma(I)$ $R_{int} = 0.021$ $\theta_{max} = 25.0^{\circ}$ $h = -8 \rightarrow 6$ $k = -23 \rightarrow 17$ $l = -19 \rightarrow 16$

H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.33 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.45 \text{ e} \text{ Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Mn-O1	1.8793 (18)	Mn-N1	1.991 (2)
Mn-O2	1.8944 (18)	Mn-O4	2.1815 (19)
Mn-N2	1.989 (2)	Mn-O3	2.249 (2)
O1-Mn-O2	94.69 (8)	N2-Mn-O4	92.95 (9)
O1-Mn-N2	174.55 (9)	N1-Mn-O4	91.45 (8)
O2-Mn-N2	90.42 (9)	O1-Mn-O3	89.20 (10)
O1-Mn-N1	92.35 (9)	O2-Mn-O3	90.78 (9)
O2-Mn-N1	172.83 (9)	N2-Mn-O3	88.84 (10)
N2-Mn-N1	82.50 (9)	N1-Mn-O3	87.97 (9)
O1-Mn-O4	88.95 (9)	O4-Mn-O3	178.04 (9)
O2-Mn-O4	90.02 (8)		

H atoms were located and included in the structure-factor calculations but their positions were not refined, except for those of the aqua molecule coordinated to Mn.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SMART* and *SAINT* (Siemens, 1994); data reduction: *SHELXTL XPREP* (Siemens, 1994); program(s) used to solve structure: *SHELXTL* (Siemens, 1994); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

This work was supported by The State Key Basic Research and Development Plan (G1998010100) and NNSFC (No. 29733090 and 29973047).

References

- Bruker (1998). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.
- Caudle, M. T., Kampf, J. W., Kirk, M. L., Rasmussen, P. G. & Pecoraro, V. L. (1997). J. Am. Chem. Soc. 119, 9297–9298.
- Oshio, H., Ino, E., Ito, T. & Maeda, Y. (1995). Bull. Chem. Soc. Jpn. 68, 889– 897.

- Rypniewski, W. R., Mangani, S., Bruni, S., Orioli, P. L., Casati, M. & Wilson, K. (1995). J. Mol. Biol. 251, 282–296.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
- Shyu, H. L., Wei, H. H. & Wang, Y. (1999). Inorg. Chim. Acta, 290, 8-13.
- Siemens (1994). SAINT and SHELXTL (Version 5.0). Siemens Analytical Xray Instruments Inc., Madison, Wisconsin, USA.
- Siemens (1996). SMART. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.
- Tainer, J. A., Getzoff, E. D., Beem, K. M., Richardson, J. S. & Richardson, D. C. (1982). J. Mol. Biol. 160, 181–217.
- Watkinson, M., Fondo, M., Bermejo, M. R., Sousa, A., McAuliffe, C. A., Pritchard, R. G., Jaiboon, N., Aurangzeb, N. & Naeem, M. (1999). J. Chem. Soc. Dalton Trans. pp. 31–41.
- Wieghardt, K. (1989). Angew. Chem. Int. Ed. Engl. 28, 1153-1172.