

Shu-Zhong Zhan,^{a*} Ying-Jun Zhang^a and Jian-Ge Wang^b^aDepartment of Chemistry, South China University of Technology, Guangzhou 510640, People's Republic of China, and ^bDepartment of Chemistry, Luoyang Normal University, Luoyang 471022, People's Republic of China

Correspondence e-mail: shzhzhan@scut.edu.cn

Key indicators

Single-crystal X-ray study

T = 291 K

Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$

R factor = 0.030

wR factor = 0.090

Data-to-parameter ratio = 16.7

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Bis{2-[(2-oxidoethyl)iminomethyl]phenolato- $\kappa^3\text{O},\text{N},\text{O}'$ }manganese(IV) hemihydrate

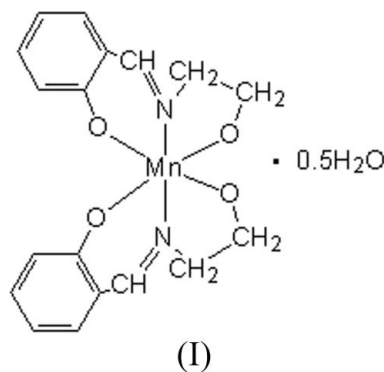
The crystal structure of the title complex, $[\text{Mn}(\text{C}_9\text{H}_9\text{NO}_2)_2] \cdot 0.5\text{H}_2\text{O}$, contains two Mn^{IV} units related by a twofold rotation axis and linked by intermolecular hydrogen bonds to a water molecule lying on this twofold axis.

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Comment

Manganese(II) centres coordinated by an *O,N,N*-tridentate Schiff base can form a variety of compounds which have been found to function as molecular nanomagnets (Aliaga *et al.*, 2001; Soler *et al.*, 2001; Yang *et al.*, 2003). Such molecules have been called single-molecule magnets (SMMs). Interest in SMMs has grown considerably since it was found that a reversal of the direction of magnetization can occur *via* quantum mechanical tunnelling. Recently, an Fe_4 SMM was synthesized by treating $\text{FeCl}_2 \cdot 4\text{H}_2\text{O}$ with a Schiff base ligand generated from the condensation of *o*-vanillin with 3-amino-propanol (Madhu *et al.*, 2005). We present here the structure of a new Mn^{IV} complex, (I) (Fig. 1).



The Mn^{IV} atom is coordinated by two N and four O atoms of the Schiff base ligands. The Mn–O bond lengths range from 1.8569 (12) to 1.9166 (11) \AA and the Mn–N distances are 1.9682 (12) and 1.9771 (13) \AA . The structure contains two Mn^{IV} units related by a twofold rotation axis and linked by intermolecular hydrogen bonds to a water molecule lying on this twofold axis.

Experimental

A solution of salicylaldehyde (10 mmol) in methanol (10 ml) was added to a solution of ethanolamine (10 mmol) in methanol (20 ml). The mixture was refluxed for 1 h before the addition of $\text{MnCl}_2 \cdot \text{H}_2\text{O}$ (10 mmol) and $\text{K}_2\text{S}_2\text{O}_8$ (20 mmol) dissolved in a minimum of water. After filtration, the solution was left to stand for several days, leading to the formation of yellow crystals (yield 65%). Crystals of (I)

suitable for single-crystal X-ray diffraction were selected directly from the sample as prepared.

Crystal data

[Mn(C₉H₉NO₂)₂].0.5H₂O
M_r = 390.29
 Orthorhombic, *Pbcn*
a = 27.6765 (16) Å
b = 10.6137 (6) Å
c = 11.6820 (7) Å
V = 3431.6 (3) Å³

Z = 8
D_x = 1.511 Mg m⁻³
 Mo *K*α radiation
 μ = 0.80 mm⁻¹
T = 291 (2) K
 Block, yellow
 0.39 × 0.28 × 0.21 mm

Data collection

Bruker SMART APEXII CCD diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.745, *T_{max}* = 0.850

19518 measured reflections
 3929 independent reflections
 3255 reflections with *I* > 2σ(*I*)
R_{int} = 0.018
 θ_{max} = 27.5°

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.030
wR (*F*²) = 0.090
S = 1.02
 3929 reflections
 235 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0512P)^2 + 0.8471P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} = 0.001
 Δρ_{max} = 0.25 e Å⁻³
 Δρ_{min} = -0.43 e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O5—H1W...O3	0.884 (9)	1.897 (10)	2.7666 (16)	168 (2)

H atoms were placed in calculated positions (N—H = 0.89 Å and C—H = 0.97 Å) and refined as riding, with *U*_{iso}(H) = 1.2*U*_{eq}(C,N). the water H atom was restrained, with H—O = 0.83 Å.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Siemens, 1995); software used to prepare material for publication: *SHELXTL*.

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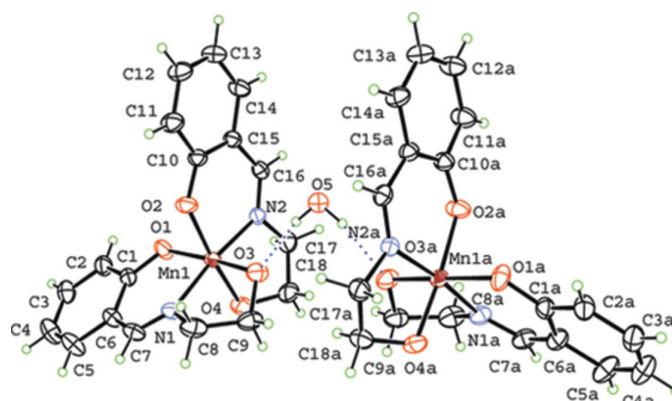


Figure 1

Two complex molecules linked by hydrogen bonds (dashed lines) to a water molecule. Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code: (a) 1 - *x*, *y*, ½ - *z*.]

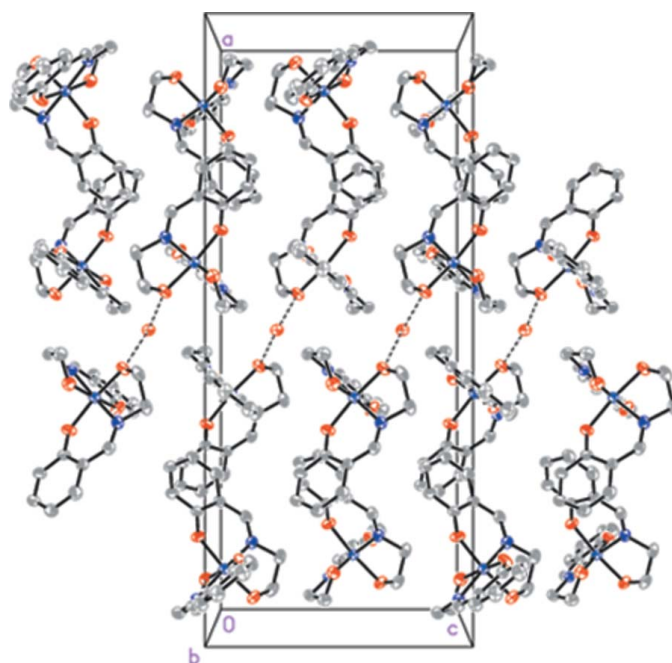


Figure 2

View along the *b* axis of the packing of the title compound. Dashed lines indicate hydrogen bonds.

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