

Poly[diaqua-di- μ_3 -malonato- μ -pyrazine-dimanganese(II)]

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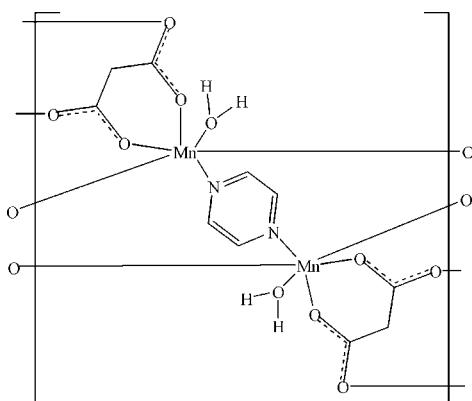
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; R factor = 0.045; wR factor = 0.117; data-to-parameter ratio = 10.9.

The title compound, $[\text{Mn}_2(\text{C}_3\text{H}_2\text{O}_4)_2(\text{C}_4\text{H}_4\text{N}_2)(\text{H}_2\text{O})_2]_n$, is isostructural with its Co^{II} , Ni^{II} , Zn^{II} and Cd^{II} analogues, and the complex resides on a crystallographic centre of inversion (at the pyrazine ring centroid). The Mn^{II} atoms are linked via coordinated malonates, forming a two-dimensional network with cavities. These sheets are further connected into a three-dimensional network by bridging pyrazine ligands which have inversion symmetry. The coordination geometry around the Mn^{II} atom is a tetragonally elongated octahedron, with pyrazine N and aqua O atoms at the axial positions.

Related literature

For studies and reviews of inorganic–organic hybrid materials, see: Chung *et al.* (1971); Okabe & Oya (2000); Serre *et al.* (2005); Pocker & Fong (1980); Scapin *et al.* (1997); Kim *et al.* (2001). For the isostructural analogues, see: Co^{II} : Delgado *et al.* (2003); Ni^{II} : Liu *et al.* (2005); Zn^{II} : Zhang *et al.* (2003); Delgado *et al.* (2003); Cd^{II} : Mao *et al.* (2004).



Experimental

Crystal data

$[\text{Mn}_2(\text{C}_3\text{H}_2\text{O}_4)_2(\text{C}_4\text{H}_4\text{N}_2)(\text{H}_2\text{O})_2]$	$V = 748.45$ (18) Å ³
$M_r = 430.10$	$Z = 2$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 7.0127$ (10) Å	$\mu = 1.74$ mm ⁻¹
$b = 14.490$ (2) Å	$T = 293$ (2) K
$c = 7.3711$ (10) Å	$0.36 \times 0.28 \times 0.24$ mm
$\beta = 92.182$ (1)°	

Data collection

Bruker APEX II CCD area-detector diffractometer	2397 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2001)	1314 independent reflections
$T_{\min} = 0.573$, $T_{\max} = 0.680$	1103 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.023$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.117$	$\Delta\rho_{\text{max}} = 0.72$ e Å ⁻³
$S = 1.00$	$\Delta\rho_{\text{min}} = -0.40$ e Å ⁻³
1314 reflections	
116 parameters	
3 restraints	

Table 1
Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
O1—H2W···O2 ⁱ	0.83 (4)	1.97 (4)	2.705 (5)	148 (7)
O1—H1W···O3 ⁱⁱ	0.82 (4)	1.90 (4)	2.644 (5)	149 (7)

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

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GG2036).

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supplementary materials

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Poly[diaqua-di- μ_3 -malonato- μ -pyrazine-dimanganese(II)]

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Comment

In recent years, dicarboxylic acids have been widely used as polydentate ligands, which undergo various metal chelation reactions to form transition or rare earth metal complexes with interesting properties in materials science and in biological systems (Church *et al.*, 1971; Okabe & Oya, 2000; Serre *et al.*, 2005; Pocker & Fong, 1980; Scapin *et al.*, 1997). For example, Kim *et al.* (2001) focused on the syntheses of transition metal complexes containing benzene dicarboxylate and rigid aromatic pyridine ligands in order to study their electronic conductivity and magnetic properties. The importance of transition metal dicarboxylate complexes in materials science and biological systems prompted us to pursue synthetic strategies for these compounds, using malonate as a polydentate ligand and pyrazine as a rigid aromatic ligand. In this paper, we report the synthesis and X-ray crystal structure analysis of the title compound, $[\text{Mn}_2(\text{C}_3\text{H}_2\text{O}_4)_2(\text{C}_4\text{H}_4\text{N}_2)(\text{H}_2\text{O})_2]_n$.

The Mn^{II} atom has sixfold coordination, chelated by two O atoms from one malonate ligand to form a six-membered boat-type ring, and by two O atoms from two neighbouring malonates, one aqua molecule and one N atom from the bridging pyrazine ligand (Fig. 1). The $\text{Mn}—\text{O}(\text{carboxylate})$ and $\text{Mn}—\text{N}$ bond lengths are in the range 2.060 (4) to 2.220 (4) and 2.269 (4) Å, respectively.

The packing diagram is shown in Fig. 2. If the pyrazine bridges are neglected, a two-dimensional network is formed by the $[\text{Mn}(\text{malonate})(\text{H}_2\text{O})]$ moieties parallel to the (010) plane. There are hydrogen bonds (Table 2) between the aqua and malonate ligands in this network.

Experimental

A mixture of manganese(II) sulfate (0.5 mmol), malonate acid (0.5 mmol), sodium hydroxide (1 mmol), pyrazine (1 mmol) and H_2O (8 ml) in a 25 ml Teflon-lined stainless steel autoclave was heated at 443 K for two days, and then cooled to room temperature. block crystals of complex (I) were obtained with a yield of 22%. Anal. Calc. for $\text{C}_5\text{H}_6\text{NMnO}_5$: C 27.91, H 2.79, N 6.51%; Found: C 27.88, H 2.75, N 6.47%.

Refinement

All H atoms on C atoms were generated geometrically and treated as riding atoms with $\text{C}—\text{H}= 0.93\text{\AA}$ and $U_{\text{iso}}(\text{H})= 1.2$ times $U_{\text{eq}}(\text{C})$. The H atoms of the water molecule were located from difference density maps and were refined with distance restraints of $d(\text{H}—\text{H}) = 1.38$ (2) Å, $d(\text{O}—\text{H}) = 0.82$ (1) Å.

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Figures

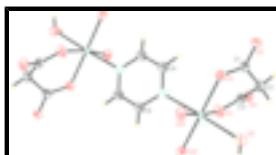


Fig. 1. Atom labeling scheme for the title complex (I), showing 30% probability displacement ellipsoids for non-H atoms. Atoms labeled with i are at the symmetry position ($-x + 1, -y, -z + 1$).

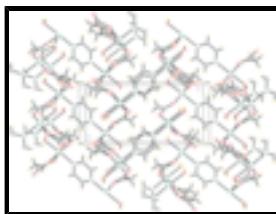


Fig. 2. Packing diagram for the title complex along the c axis.

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Crystal data

[Mn ₂ (C ₃ H ₂ O ₄) ₂ (C ₄ H ₄ N ₂)(H ₂ O) ₂]	$F_{000} = 432$
$M_r = 430.10$	$D_x = 1.908 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
Hall symbol: -P 2yn	$\lambda = 0.71073 \text{ \AA}$
$a = 7.0127 (10) \text{ \AA}$	Cell parameters from 1314 reflections
$b = 14.490 (2) \text{ \AA}$	$\theta = 2.8\text{--}25.1^\circ$
$c = 7.3711 (10) \text{ \AA}$	$\mu = 1.74 \text{ mm}^{-1}$
$\beta = 92.182 (1)^\circ$	$T = 293 (2) \text{ K}$
$V = 748.45 (18) \text{ \AA}^3$	Block, colorless
$Z = 2$	$0.36 \times 0.28 \times 0.24 \text{ mm}$

Data collection

Bruker APEX II CCD area-detector diffractometer	1314 independent reflections
Radiation source: fine-focus sealed tube	1103 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.023$
$T = 293(2) \text{ K}$	$\theta_{\max} = 25.1^\circ$
φ and ω scans	$\theta_{\min} = 2.8^\circ$
Absorption correction: multi-scan (SADABS; Bruker, 2001)	$h = -8 \rightarrow 3$
$T_{\min} = 0.573, T_{\max} = 0.680$	$k = -13 \rightarrow 16$
2397 measured reflections	$l = -8 \rightarrow 8$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites

$R[F^2 > 2\sigma(F^2)] = 0.045$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.117$	$w = 1/\sigma^2(F_o^2) + (0.0636P)^2 + 2.2589P$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.00$	$(\Delta/\sigma)_{\max} = 0.013$
1314 reflections	$\Delta\rho_{\max} = 0.72 \text{ e \AA}^{-3}$
116 parameters	$\Delta\rho_{\min} = -0.40 \text{ e \AA}^{-3}$
3 restraints	Extinction correction: SHELXL97 (Sheldrick, 1997), $F_c^* = k F_c [1 + 0.001 x F_c^2 \lambda^3 / \sin(2\theta)]^{1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.051 (5)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C3	0.5846 (7)	0.3318 (3)	0.7561 (7)	0.0344 (11)
C4	0.6125 (8)	0.3910 (4)	0.5887 (7)	0.0432 (12)
H4A	0.7407	0.4162	0.5951	0.052*
H4B	0.5240	0.4424	0.5904	0.052*
C5	0.5834 (7)	0.3404 (3)	0.4123 (7)	0.0348 (11)
C2	0.4526 (8)	0.0744 (4)	0.5906 (8)	0.0507 (14)
H2	0.4135	0.1258	0.6549	0.061*
C1	0.6785 (7)	-0.0046 (4)	0.4448 (8)	0.0468 (13)
H1	0.8016	-0.0106	0.4036	0.056*
Mn1	0.83760 (8)	0.19134 (4)	0.58103 (7)	0.0172 (3)
N1	0.6323 (6)	0.0717 (3)	0.5372 (6)	0.0389 (10)
O2	0.6638 (5)	0.2528 (3)	0.7693 (5)	0.0433 (9)
O5	0.4862 (5)	0.3675 (2)	0.8747 (5)	0.0435 (9)
O3	0.6698 (5)	0.2644 (3)	0.3903 (5)	0.0464 (9)
O4	0.4753 (5)	0.3775 (2)	0.2943 (5)	0.0427 (9)
O1	1.0317 (6)	0.3107 (3)	0.5859 (5)	0.0473 (9)
H2W	1.087 (8)	0.313 (5)	0.489 (4)	0.080*
H1W	1.104 (7)	0.303 (5)	0.675 (5)	0.080*

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Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C3	0.038 (2)	0.031 (3)	0.034 (2)	0.000 (2)	0.0003 (19)	-0.005 (2)
C4	0.060 (3)	0.035 (3)	0.034 (3)	0.005 (2)	0.003 (2)	0.000 (2)
C5	0.039 (2)	0.029 (3)	0.036 (3)	0.003 (2)	0.005 (2)	0.004 (2)
C2	0.048 (3)	0.042 (3)	0.063 (4)	-0.006 (3)	0.018 (3)	-0.014 (3)
C1	0.041 (3)	0.037 (3)	0.063 (4)	-0.002 (2)	0.012 (2)	-0.009 (3)
Mn1	0.0218 (4)	0.0155 (4)	0.0144 (4)	0.0007 (2)	0.0018 (2)	-0.0006 (2)
N1	0.039 (2)	0.036 (2)	0.042 (2)	-0.0017 (18)	0.0021 (17)	-0.0012 (19)
O2	0.056 (2)	0.039 (2)	0.0350 (19)	0.0076 (17)	0.0117 (16)	0.0031 (15)
O5	0.049 (2)	0.040 (2)	0.043 (2)	0.0044 (16)	0.0172 (16)	-0.0004 (17)
O3	0.058 (2)	0.045 (2)	0.0355 (19)	0.0118 (18)	-0.0044 (16)	-0.0029 (16)
O4	0.053 (2)	0.0347 (19)	0.0399 (19)	0.0032 (17)	-0.0064 (16)	0.0032 (16)
O1	0.051 (2)	0.055 (2)	0.037 (2)	-0.0021 (18)	0.0019 (16)	0.0004 (18)

Geometric parameters (\AA , $^\circ$)

C3—O5	1.246 (6)	C1—C2 ⁱ	1.386 (8)
C3—O2	1.275 (6)	C1—H1	0.9300
C3—C4	1.521 (7)	Mn1—O5 ⁱⁱ	2.060 (3)
C4—C5	1.500 (7)	Mn1—O4 ⁱⁱⁱ	2.070 (4)
C4—H4A	0.9700	Mn1—O3	2.087 (4)
C4—H4B	0.9700	Mn1—O2	2.082 (3)
C5—O4	1.254 (6)	Mn1—O1	2.200 (4)
C5—O3	1.270 (7)	Mn1—N1	2.269 (4)
C2—N1	1.335 (7)	O5—Mn1 ^{iv}	2.060 (3)
C2—C1 ⁱ	1.386 (8)	O4—Mn1 ^v	2.070 (4)
C2—H2	0.9300	O1—H2W	0.83 (4)
C1—N1	1.344 (7)	O1—H1W	0.82 (4)
O5—C3—O2	124.7 (5)	O4 ⁱⁱⁱ —Mn1—O2	88.18 (14)
O5—C3—C4	115.3 (4)	O3—Mn1—O2	84.41 (14)
O2—C3—C4	119.9 (4)	O5 ⁱⁱ —Mn1—O1	90.42 (14)
C3—C4—C5	114.2 (4)	O4 ⁱⁱⁱ —Mn1—O1	95.52 (14)
C3—C4—H4A	108.7	O3—Mn1—O1	86.84 (16)
C5—C4—H4A	108.7	O2—Mn1—O1	91.75 (15)
C3—C4—H4B	108.7	O5 ⁱⁱ —Mn1—N1	85.11 (15)
C5—C4—H4B	108.7	O4 ⁱⁱⁱ —Mn1—N1	90.85 (15)
H4A—C4—H4B	107.6	O3—Mn1—N1	87.32 (16)
O4—C5—O3	124.2 (5)	O2—Mn1—N1	92.14 (15)
O4—C5—C4	116.8 (4)	O1—Mn1—N1	172.65 (15)
O3—C5—C4	119.0 (4)	C2—N1—C1	115.0 (4)
N1—C2—C1 ⁱ	123.3 (5)	C2—N1—Mn1	122.4 (4)
N1—C2—H2	118.3	C1—N1—Mn1	122.4 (3)
C1 ⁱ —C2—H2	118.3	C3—O2—Mn1	126.7 (3)

N1—C1—C2 ⁱ	121.7 (5)	C3—O5—Mn1 ^{iv}	131.0 (3)
N1—C1—H1	119.2	C5—O3—Mn1	127.8 (3)
C2 ⁱ —C1—H1	119.2	C5—O4—Mn1 ^v	124.9 (3)
O5 ⁱⁱ —Mn1—O4 ⁱⁱⁱ	97.27 (15)	Mn1—O1—H2W	109 (5)
O5 ⁱⁱ —Mn1—O3	90.03 (15)	Mn1—O1—H1W	105 (5)
O4 ⁱⁱⁱ —Mn1—O3	172.30 (14)	H2W—O1—H1W	114 (3)
O5 ⁱⁱ —Mn1—O2	173.92 (15)		

Symmetry codes: (i) $-x+1, -y, -z+1$; (ii) $x+1/2, -y+1/2, z-1/2$; (iii) $x+1/2, -y+1/2, z+1/2$; (iv) $x-1/2, -y+1/2, z+1/2$; (v) $x-1/2, -y+1/2, z-1/2$.

Hydrogen-bond geometry (\AA , $^\circ$)

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
O1—H2W \cdots O2 ⁱⁱ	0.83 (4)	1.97 (4)	2.705 (5)	148 (7)
O1—H1W \cdots O3 ⁱⁱⁱ	0.82 (4)	1.90 (4)	2.644 (5)	149 (7)

Symmetry codes: (ii) $x+1/2, -y+1/2, z-1/2$; (iii) $x+1/2, -y+1/2, z+1/2$.

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Fig. 1

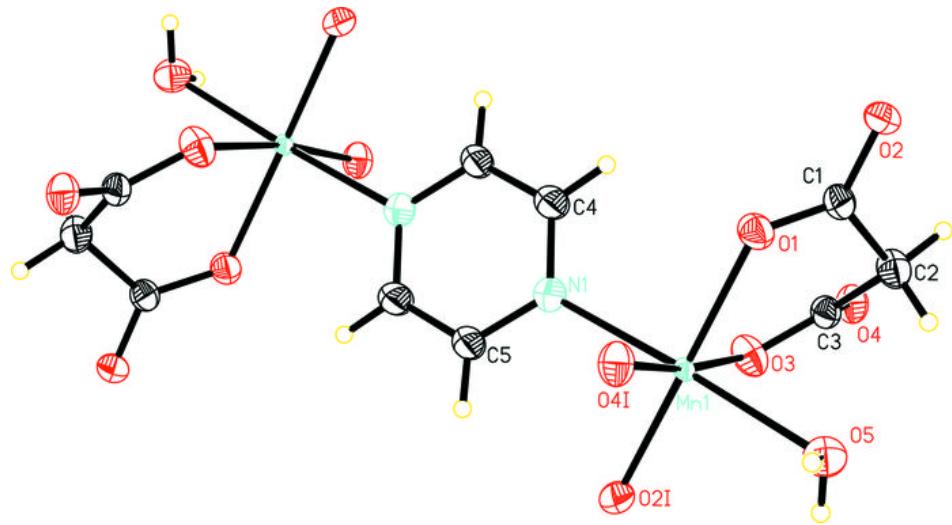


Fig. 2

