metal-organic papers

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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$ Disorder in main residue R factor = 0.033 wR factor = 0.101 Data-to-parameter ratio = 14.4

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

Diaquabis(4-formylbenzoato- κ O)bis(1*H*-imidazole- κ N³)manganese(II)

The Mn atom in the title compound, $[Mn(C_8H_5O_3)_2 (C_3H_4N_2)_2(H_2O)_2]$, is datively linked to the imidazole heterocycles and water molecules, and covalently linked to the substituted benzoate groups in an all-*trans* octahedral geometry; the Mn atom lies on a center of inversion.

Comment

Benzoic acid and imidazole are used in the synthesis of a large number of transition metal complexes; the crystal structures of some of these have been reported (Cambridge Structural Database, Version 5.25; Allen, 2002). An example is diaquabis(4-nitrobenzoato)bis(imidazole)manganese, which exists as a centrosymmetric six-coordinate molecule that is hydrogenbonded into a three-dimensional network motif (Xu & Xu, 2004). However, among the substituted benzoic acids, there is not much information about metal derivatives of 4-formylbenzoic acid in the literature (American Chemical Society, 2006). Although the crystal structure of the acid is known (Haisa *et al.*, 1976), no crystal structure of its metal derivatives has been reported.



The centrosymmetric title complex, (I), has the metal atom in an all-*trans* octahedral geometry; the metal atom is linked covalently to the 4-formylbenzoate group, and datively to the water and imidazole donor ligands (Fig. 1). The coordinated water molecules engage in hydrogen-bonding interactions (Table 2), giving rise to a three-dimensional network motif. The interaction involving the formyl unit is weak, however, and may account for the unit adopting two conformations.

Experimental

© 2006 International Union of Crystallography All rights reserved Manganese diacetate tetrahydrate (0.122 g, 0.5 mmol) was added to an aqueous solution (10 ml) of 4-formylbenzoic acid (0.15 g, 1 mmol) and imidazole (0.068 g, 1 mmol). The pH value of the mixture was Received 3 August 2006 Accepted 4 August 2006 about 5. The filtered solution was allowed to evaporate at room temperature; colorless prismatic crystals separated after several days. Elemental analysis calculated for $C_{22}H_{22}N_4O_8Mn$: C 50.29, H 4.22, N 10.67%; found: C 50.44, H 3.88, N 10.69%.

V = 574.3 (4) Å³

 $D_x = 1.519 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation $\mu = 0.63 \text{ mm}^{-1}$

T = 295 (2) K

 $R_{\rm int} = 0.017$

 $\theta_{\rm max} = 27.5^{\circ}$

Prism, colorless

0.36 \times 0.28 \times 0.15 mm

5688 measured reflections

2616 independent reflections

2474 reflections with $I > 2\sigma(I)$

Z = 1

Crystal data

$$\begin{split} & \left[\mathrm{Mn}(\mathrm{C_8H_5O_3})_2(\mathrm{C_3H_4N_2})_2(\mathrm{H_2O})_2 \right] \\ & M_r = 525.38 \\ & \mathrm{Triclinic}, \ P\overline{1} \\ & a = 5.644 \ (2) \ \text{\AA} \\ & b = 8.141 \ (4) \ \text{\AA} \\ & c = 12.940 \ (5) \ \text{\AA} \\ & \alpha = 75.626 \ (1)^{\circ} \\ & \beta = 86.664 \ (1)^{\circ} \\ & \gamma = 86.384 \ (1)^{\circ} \end{split}$$

Data collection

Rigaku RAXIS-RAPID IP diffractometer ω scans Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995) $T_{min} = 0.662, T_{max} = 0.911$

Refinement

 Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.061P)^2$
 $R[F^2 > 2\sigma(F^2)] = 0.033$ $w = 1/[\sigma^2(F_o^2) + (0.061P)^2$
 $wR(F^2) = 0.101$ where $P = (F_o^2 + 2F_c^2)/3$

 S = 1.15 $(\Delta/\sigma)_{max} = 0.001$

 2616 reflections
 $\Delta\rho_{max} = 0.36$ e Å⁻³

 182 parameters
 $\Delta\rho_{min} = -0.32$ e Å⁻³

 H atoms treated by a mixture of independent and constrained refinement
 A^{-3}

Table 1

Selected geometric parameters (Å, $^{\circ}$).

Mn1-O1 Mn1-O1w	2.221 (1) 2.218 (1)	Mn1-N1	2.230 (2)
$\begin{array}{c} O1 - Mn1 - O1^{i} \\ O1 - Mn1 - O1w \\ O1 - Mn1 - O1w^{i} \\ O1 - Mn1 - N1 \\ O1 - Mn1 - N1^{i} \end{array}$	180 88.87 (5) 91.13 (5) 93.16 (5) 86.84 (5)	$01w - Mn1 - 01w^{i}$ 01w - Mn1 - N1 $01w - Mn1 - N1^{i}$ $N1 - Mn1 - N1^{i}$	180 90.28 (5) 89.72 (5) 180

Symmetry code: (i) -x + 1, -y + 1, -z + 1.

Table 2

H	lyd	lrogen-	bond	geomet	ry	(A,	°)).
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1w - H1w1 \cdots O1^{ii}$	0.85 (1)	1.97 (1)	2.804 (2)	168 (2)
$O1w - H1w2 \cdots O2^{i}$	0.85(1)	1.84 (1)	2.667 (2)	168 (3)
$N2-H2n \cdot \cdot \cdot O2^{iii}$	0.85 (1)	2.37 (2)	3.089 (3)	142 (3)
$N2-H2n\cdots O3^{iv}$	0.85 (1)	2.43 (2)	3.044 (3)	129 (3)



Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are drawn as spheres of arbitrary radii. The minor disorder component is not shown. [Symmetry code: (i) 1 - x, 1 - y, 1 - z.]

The formyl group is disordered over two positions; the occupancies refined to 0.639 (5):0.361 (5). The group was refined as two CH(==O) groups sharing a common C atom; this was necessary for the generation of the riding H atoms. The carbon-bound H atoms were positioned geometrically (C-H = 0.93 Å) and were included in the refinement in the riding model approximation, with $U_{\rm iso}(H) = 1.2U_{\rm eq}(C)$. The amino and water H atoms were located in a difference Fourier map, and were refined with a distance restraint of N-H = O-H 0.85 (1) Å.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97*.

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