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Key indicators

Single-crystal X-ray study T = 299 KMean σ (C–C) = 0.003 Å R factor = 0.023 wR factor = 0.057 Data-to-parameter ratio = 14.8

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Bis(piperazinium) di-µ-chloro-bis[diaquadichloromanganate(II)] dichloride

The title compound, $(C_4H_{12}N_2)_2[Mn_2Cl_6(H_2O)_4]Cl_2$, consists of two piperazinium cations, two chloride anions and a dimanganese complex anion, which lies on a center of symmetry. Each Mn^{II} atom is located in a distorted octahedral environment. The cations and the anions are held together by $O-H\cdots Cl$ and $N-H\cdots Cl$ hydrogen bonds. The structure features the formation of inorganic and organic ionic layers, which interpenetrate each other and which are linked by hydrogen bonds. Received 19 July 2005 Accepted 1 August 2005 Online 6 August 2005

Comment

The chloride anion, as a bridging ligand, exhibits a good binding ability in the formation of coordination complexes (Caneschi *et al.*, 1990; Fryzuk *et al.*, 1994; Rancurel *et al.*, 2000; Wang *et al.*, 2003). Piperazine is a strong basic amine which readily forms dications, where all four N-H bonds are generally active in hydrogen-bond formation (Coupar *et al.*, 1996; Ferguson *et al.*, 1998). Moreover, dinuclear manganese complexes are active in biological redox reactions involving water, peroxide and dioxygen (Barynin *et al.*, 1997; Zhang *et al.*, 2001). Recently, we have synthesized the title compound, (I), a dinuclear manganese(II) complex.



The title compound consists of two diprotonated piperazinium ions, two chloride anions and a dimanganese(II) anion (Fig. 1). The Mn^{II} complex is built up from two inver-



Figure 1

The formula unit with atom labels, showing 40% probability displacement ellipsoids. Unlabeled atoms are related by the symmetry code (-x, -y, 1-z). Hydrogen bonds are illustrated as dashed lines.

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Figure 2



sion-related Mn^{II} atoms bridged by two Cl⁻ anions, where the Mn-Cl-Mn bond angle is 94.41 (2) $^{\circ}$, smaller than the value of 96.82 $(4)^{\circ}$ in the pentacoordinated dimanganese compound $[MnCl_2(HDPhF)_2]_2$ (HDPhF is diphenylformamidine) (Arnold et al., 1997). The Mn^{II} atoms in (I) are separated by a distance of 3.843 (7) Å, which is quite large and seems to rule out any strong direct metal-metal interaction. Each Mn^{II} atom is located in a distorted octahedral environment; the equatorial plane is formed by coordinated atoms O1, O2, Cl1 and Cl1(-x, -y, 1-z). The O2 water molecule is firmly bonded to the Mn^{II} atom, with a bond length of 2.1776 (18) Å, which is one of the shortest such lengths observed (Orpen et al., 1989; Garoufis et al., 1999). The bridging Mn-Cl bonds, viz. Mn-Cl1 and Mn-Cl1(-x, -y, 1-z), are obviously elongated, while the terminal Mn-Cl2 and Mn-Cl3 bonds are normal for hexacoordinated Mn^{II} complexes (Orpen et al., 1989).

In the structure of (I), a piperazinium cation is linked to an uncoordinated chloride ion by an N2-H2NB···Cl4 hydrogen bond, and the chloride ion is, in turn, linked to a coordinated water molecule by an O1-H1OA···Cl4 hydrogen bond (Fig. 1 and Table 2). In addition, the cation is linked to a coordinated Cl atom by an N1-H1NA···Cl2 hydrogen bond. The hydrogen bonds mentioned above result in a supramolecular ring, which may be described by the graph set $R_2^3(11)$ (Etter, 1990; Grell *et al.*, 2000). As shown in Fig. 2, the structure features inorganic and organic ionic layers parallel to the (100) plane, which interpenetrate and are linked by hydrogen bonds (Table 2).

Experimental

Piperazine, MnCl₂ and HCl in a molar ratio of 1:2:4 were mixed, dissolved in water and heated to a temperature where a clear solution resulted. Single crystals of (I) formed when the solution was left standing overnight at 293 K. IR (KBr, cm⁻¹): 3441 (*b*), 3008 (*w*), 2995 (*w*), 2748 (*m*), 2459 (*w*), 1629 (*s*), 1464 (*m*), 1354 (*m*), 1234 (*m*), 1091 (*w*), 812 (*m*), 791 (*m*), 718 (*m*), 599 (*m*), 570 (*w*), 446 (*w*).

 $\begin{array}{l} ({\rm C}_4{\rm H}_{12}{\rm N}_{2})_2[{\rm Mn}_2{\rm Cl}_6({\rm H}_2{\rm O})_4]{\rm Cl}_2\\ M_r = 641.86\\ {\rm Monoclinic}, P_{2_1}/c\\ a = 9.865 \ (2) \ {\rm \AA}\\ b = 9.429 \ (1) \ {\rm \AA}\\ c = 13.506 \ (2) \ {\rm \AA}\\ \beta = 101.74 \ (1)^\circ\\ V = 1230.0 \ (3) \ {\rm \AA}^3\\ Z = 2 \end{array}$

Data collection

Siemens P4 diffractometer ω scans Absorption correction: ψ scan (North et al., 1968) $T_{min} = 0.356, T_{max} = 0.608$ 2623 measured reflections 2235 independent reflections 1931 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_0^2) + (0.0296P)^2]$ where $P = (F_0^2 + 2F_c^2)/3$ $R[F^2 > 2\sigma(F^2)] = 0.023$ $wR(F^2) = 0.057$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.26 \text{ e} \text{ Å}^{-3}$ S = 1.09 $\Delta \rho_{\rm min} = -0.23 \ {\rm e} \ {\rm \AA}^{-3}$ 2235 reflections 151 parameters Extinction correction: SHELXL97 H atoms treated by a mixture of Extinction coefficient: 0.0331 (10) independent and constrained refinement

 $D_x = 1.733 \text{ Mg m}^{-3}$ Mo *K* α radiation

reflections

 $\theta = 5.2 - 15.1^{\circ}$ $\mu = 1.92 \text{ mm}^{-1}$

T = 299 (2) K

Block, pink

 $R_{\rm int} = 0.012$

 $\theta_{\rm max} = 25.3^{\circ}$

 $h = 0 \rightarrow 11$

 $k = 0 \rightarrow 11$

 $l = -16 \rightarrow 15$

3 standard reflections

every 97 reflections

intensity decay: 1.2%

Cell parameters from 36

 $0.56 \times 0.56 \times 0.26$ mm

Table 1

Selected geometric parameters (Å, °).

Mn-O2	2.1776 (18)	N1-C4	1.488 (3)
Mn-O1	2.2308 (18)	N1-C1	1.489 (3)
Mn-Cl2	2.4540 (7)	N2-C2	1.488 (3)
Mn-Cl3	2.5038 (7)	N2-C3	1.492 (3)
Mn-Cl1	2.5939 (7)	C1-C2	1.509 (3)
Mn-Cl1 ⁱ	2.6429 (6)	C3-C4	1.507 (3)
O2-Mn-O1	91.14 (9)	O2-Mn-Cl1	93.56 (7)
O2-Mn-Cl2	86.86 (5)	Cl2-Mn-Cl1	90.36 (2)
O1-Mn-Cl2	91.21 (5)	Cl3-Mn-Cl1	92.32 (2)
O2-Mn-Cl3	86.35 (5)	Mn-Cl1-Mn ⁱ	94.41 (2)
O1-Mn-Cl3	86.66 (5)		

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

Table 2				
Hydrogen-bond	geometry	(Å,	°).	

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O1−H1OA…Cl4	0.81 (1)	2.40(1)	3.197 (2)	167 (4)
$O1 - H1OB \cdot \cdot \cdot Cl1^{ii}$	0.81 (1)	2.77 (2)	3.5411 (19)	160 (3)
O2−H2OB···Cl3 ⁱⁱⁱ	0.81(1)	2.45 (1)	3.2433 (19)	167 (3)
O2−H2OA···Cl4 ^{iv}	0.81(1)	2.37 (1)	3.174 (2)	172 (3)
$N1 - H1NA \cdots Cl2$	0.89 (1)	2.50 (2)	3.168 (2)	132 (2)
$N1 - H1NB \cdot \cdot \cdot Cl3^{v}$	0.89(1)	2.39 (1)	3.223 (2)	156 (2)
$N2 - H2NB \cdot \cdot \cdot Cl4$	0.91 (1)	2.25 (1)	3.148 (2)	169 (2)
$N2-H2NA\cdots Cl1^{vi}$	0.90 (1)	2.49 (2)	3.247 (2)	141 (2)

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

H atoms attached to O and N atoms were located in a difference Fourier map and refined isotropically. Other H atoms were placed in calculated positions and allowed to ride on their parent atoms at a distance of 0.97 Å, with isotropic displacement parameters 1.2 times $U_{\rm eq}$ of the parent atoms.

Data collection: XSCANS (Siemens, 1994); cell refinement: XSCANS; data reduction: SHELXTL (Siemens, 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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