

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Dichlorido(2,9-dimethyl-1,10-phenanthroline)manganese(II) hemihydrate

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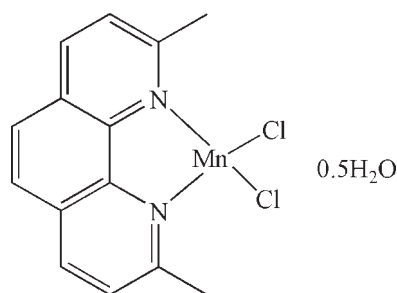
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Received 12 November 2009; accepted 18 November 2009

 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; disorder in solvent or counterion; R factor = 0.044; wR factor = 0.134; data-to-parameter ratio = 12.6.

In the title compound, $[\text{MnCl}_2(\text{C}_{14}\text{H}_{12}\text{N}_2)] \cdot 0.5\text{H}_2\text{O}$, all of the non-H atoms apart from the Cl atom lie on a mirror plane and the methyl H atoms are disordered over two sites of equal occupancy about the mirror plane. The Mn^{II} ion is coordinated in a distorted tetrahedral environment by two N atoms of the phenanthroline ligand and two chloride ions. A half-occupancy solvent water molecule lies on a mirror plane and close to an inversion center.

Related literature

 For related crystal structures, see: McCann *et al.* (1998); Pan & Xu (2005); Xu *et al.* (2009).


Experimental

Crystal data

 $[\text{MnCl}_2(\text{C}_{14}\text{H}_{12}\text{N}_2)] \cdot 0.5\text{H}_2\text{O}$
 $M_r = 343.10$

 Monoclinic, $C2/m$
 $a = 18.763$ (4) Å

 $b = 7.7343$ (15) Å

 $c = 11.362$ (2) Å

 $\beta = 101.532$ (3)°

 $V = 1615.5$ (5) Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 1.14$ mm⁻¹
 $T = 293$ K

 $0.31 \times 0.23 \times 0.19$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer

Absorption correction: multi-scan (SADABS; Sheldrick, 2008a)

 $T_{\text{min}} = 0.719$, $T_{\text{max}} = 0.813$

4794 measured reflections

1511 independent reflections

 1199 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.134$
 $S = 1.01$

1511 reflections

120 parameters

 $\Delta\rho_{\text{max}} = 0.59$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.31$ e Å⁻³

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT-Plus (Bruker, 2001); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008b); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008b); molecular graphics: SHELXTL (Sheldrick, 2008b); software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2953).

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supporting information

Acta Cryst. (2009). E65, m1659 [doi:10.1107/S1600536809049149]

Dichlorido(2,9-dimethyl-1,10-phenanthroline)manganese(II) hemihydrate**Xinfang Wang****S1. Comment**

1,10-phenanthroline is a good bidentate chelating ligand and here, we present the crystal structure of the title complex based on 2,9-dimethyl-1,10-phenanthroline.

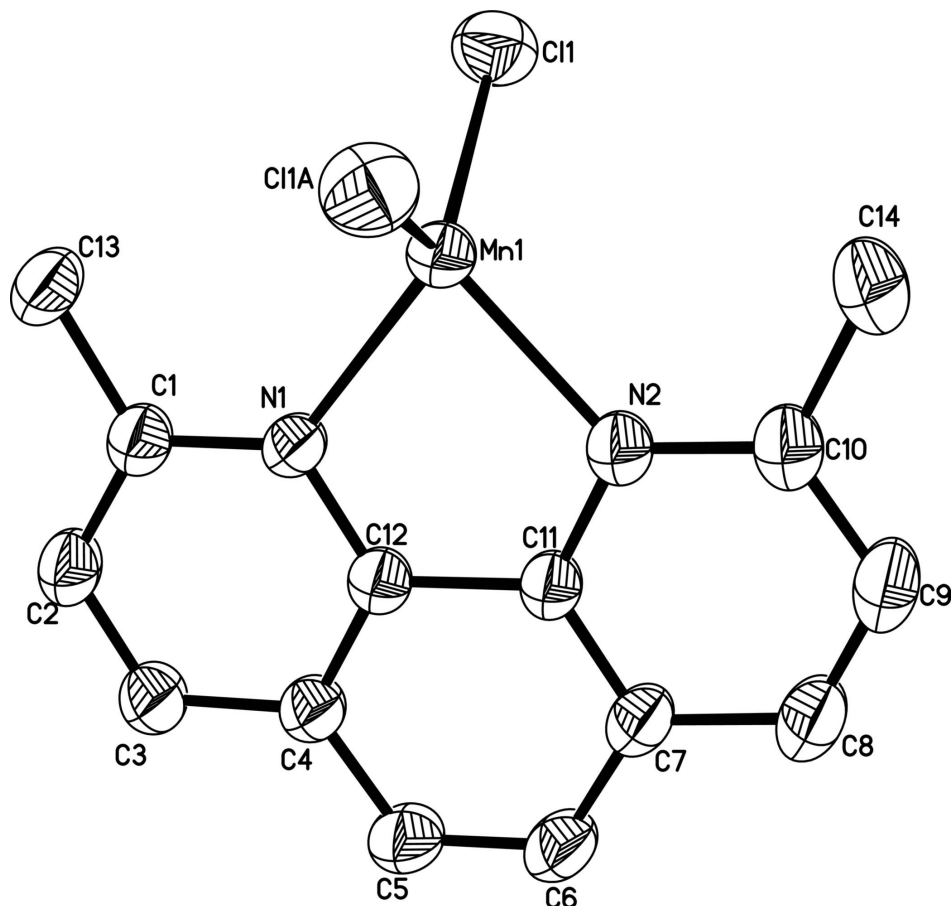
The crystal structure of the title compound is shown in Fig. 1. The coordination environment of the Mn^{II} ion is distorted tetrahedral, in which two sites are occupied by the two N atoms of the chelating 2,9-dimethyl-1,10-phenanthroline ligand and the other two from two chloride ions. For Mn—N and Mn—Cl bond lengths in other manganese biphenanthroline complexes, see e.g. McCann, *et al.* (1998); Pan & Xu (2005); Xu *et al.* (2009). The location of the water H atoms is such that they are disordered over several sites imposed by the crystal symmetry and hence any potential hydrogen bonding is not discussed.

S2. Experimental

A mixture of 2,9-dimethyl-1,10-phenanthroline, MnCl₂·4H₂O (1:2, molar ratio) and water (20 ml) was sealed in a Teflon-lined autoclave (25 ml) and heated 393 K for two days. Upon cooling slowly and opening the bomb, yellow crystals suitable for X-ray diffraction were obtained with a yield about 40% (based on phenanthroline).

S3. Refinement

All H atoms bonded to C atoms were included using the HFIX commands in SHELXL-97 (Sheldrick, 2008b/i>) with C—H distances of 0.93 and 0.96 Å, and were allowed for as riding atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{C}_{\text{methyl}})$. The H atoms of the disordered water molecule were found in a difference Fourier map and were refined as riding with O-H fixed at 0.85 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$.

**Figure 1**

The molecular structure of the title compound with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. The disordered water molecule and all the H-atoms are omitted for clarity.

Dichlorido(2,9-dimethyl-1,10-phenanthroline)manganese(II) hemihydrate

Crystal data

$[\text{MnCl}_2(\text{C}_{14}\text{H}_{12}\text{N}_2)] \cdot 0.5\text{H}_2\text{O}$

$M_r = 343.10$

Monoclinic, $C2/m$

Hall symbol: $-C 2y$

$a = 18.763 (4) \text{ \AA}$

$b = 7.7343 (15) \text{ \AA}$

$c = 11.362 (2) \text{ \AA}$

$\beta = 101.532 (3)^\circ$

$V = 1615.5 (5) \text{ \AA}^3$

$Z = 4$

$F(000) = 696$

$D_x = 1.411 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1089 reflections

$\theta = 2.9\text{--}26.8^\circ$

$\mu = 1.14 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Block, yellow

$0.31 \times 0.23 \times 0.19 \text{ mm}$

Data collection

Bruker APEXII CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2008*a*)

$T_{\min} = 0.719$, $T_{\max} = 0.813$

4794 measured reflections

1511 independent reflections

1199 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.020$
 $\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 2.2^\circ$
 $h = -22 \rightarrow 22$

$k = -8 \rightarrow 9$
 $l = -13 \rightarrow 13$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.134$
 $S = 1.01$
 1511 reflections
 120 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 $w = 1/[\sigma^2(F_o^2) + (0.1015P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.59 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.31 \text{ e } \text{\AA}^{-3}$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Mn1	0.35177 (3)	0.0000	0.27709 (5)	0.0558 (3)	
Cl1	0.40493 (4)	0.25282 (10)	0.24304 (8)	0.0815 (3)	
O1	0.5351 (5)	1.0000	0.0528 (9)	0.155 (4)	0.50
H2A	0.5257	0.9426	0.1128	0.186*	0.25
H1A	0.5745	0.9558	0.0399	0.186*	0.25
N1	0.30817 (15)	0.0000	0.4374 (3)	0.0555 (8)	
N2	0.24030 (17)	0.0000	0.2034 (3)	0.0601 (8)	
C1	0.3433 (2)	0.0000	0.5521 (4)	0.0625 (10)	
C2	0.3066 (2)	0.0000	0.6486 (4)	0.0715 (12)	
H2	0.3329	0.0000	0.7272	0.086*	
C3	0.2337 (2)	0.0000	0.6273 (4)	0.0715 (12)	
H3	0.2089	0.0000	0.6905	0.086*	
C4	0.1951 (2)	0.0000	0.5056 (3)	0.0604 (10)	
C5	0.1191 (2)	0.0000	0.4745 (4)	0.0781 (13)	
H5	0.0919	0.0000	0.5346	0.094*	
C6	0.0848 (2)	0.0000	0.3584 (4)	0.0768 (13)	
H6	0.0342	0.0000	0.3399	0.092*	
C7	0.1240 (2)	0.0000	0.2635 (4)	0.0655 (11)	
C8	0.0910 (3)	0.0000	0.1374 (4)	0.0840 (15)	
H8	0.0406	0.0000	0.1136	0.101*	
C9	0.1330 (3)	0.0000	0.0535 (4)	0.0875 (15)	
H9	0.1113	0.0000	-0.0276	0.105*	

C10	0.2073 (3)	0.0000	0.0872 (4)	0.0703 (11)	
C11	0.1985 (2)	0.0000	0.2896 (3)	0.0562 (9)	
C12	0.2350 (2)	0.0000	0.4153 (3)	0.0532 (9)	
C13	0.4239 (2)	0.0000	0.5734 (4)	0.0819 (14)	
H13A	0.4397	-0.0724	0.5150	0.123*	0.50
H13B	0.4431	-0.0434	0.6525	0.123*	0.50
H13C	0.4410	0.1158	0.5666	0.123*	0.50
C14	0.2542 (3)	0.0000	-0.0069 (4)	0.0933 (17)	
H14A	0.2713	0.1152	-0.0163	0.140*	0.50
H14B	0.2260	-0.0395	-0.0820	0.140*	0.50
H14C	0.2950	-0.0756	0.0181	0.140*	0.50

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Mn1	0.0481 (4)	0.0703 (5)	0.0501 (4)	0.000	0.0127 (3)	0.000
Cl1	0.0781 (6)	0.0818 (7)	0.0876 (6)	-0.0094 (4)	0.0235 (4)	0.0054 (4)
O1	0.108 (7)	0.226 (12)	0.134 (9)	0.000	0.031 (6)	0.000
N1	0.0413 (15)	0.074 (2)	0.0503 (17)	0.000	0.0063 (13)	0.000
N2	0.0572 (17)	0.073 (2)	0.0483 (17)	0.000	0.0069 (14)	0.000
C1	0.056 (2)	0.076 (3)	0.053 (2)	0.000	0.0032 (17)	0.000
C2	0.061 (2)	0.100 (4)	0.049 (2)	0.000	-0.0002 (18)	0.000
C3	0.064 (2)	0.101 (4)	0.050 (2)	0.000	0.0127 (19)	0.000
C4	0.053 (2)	0.077 (3)	0.052 (2)	0.000	0.0100 (17)	0.000
C5	0.055 (2)	0.112 (4)	0.071 (3)	0.000	0.020 (2)	0.000
C6	0.045 (2)	0.110 (4)	0.073 (3)	0.000	0.006 (2)	0.000
C7	0.047 (2)	0.080 (3)	0.064 (3)	0.000	-0.0017 (18)	0.000
C8	0.062 (3)	0.111 (4)	0.070 (3)	0.000	-0.008 (2)	0.000
C9	0.078 (3)	0.120 (4)	0.056 (3)	0.000	-0.010 (2)	0.000
C10	0.078 (3)	0.076 (3)	0.054 (2)	0.000	0.006 (2)	0.000
C11	0.051 (2)	0.070 (3)	0.0456 (19)	0.000	0.0040 (16)	0.000
C12	0.053 (2)	0.054 (2)	0.052 (2)	0.000	0.0073 (16)	0.000
C13	0.049 (2)	0.127 (4)	0.064 (3)	0.000	0.000 (2)	0.000
C14	0.095 (3)	0.133 (5)	0.052 (2)	0.000	0.014 (2)	0.000

Geometric parameters (Å, °)

Mn1—N2	2.092 (3)	C5—C6	1.347 (6)
Mn1—N1	2.140 (3)	C5—H5	0.9300
Mn1—Cl1	2.2633 (9)	C6—C7	1.421 (6)
Mn1—Cl1 ⁱ	2.2633 (9)	C6—H6	0.9300
O1—O1 ⁱⁱ	1.59 (2)	C7—C11	1.371 (5)
O1—H2A	0.8610	C7—C8	1.443 (6)
O1—H1A	0.8530	C8—C9	1.353 (7)
N1—C1	1.338 (5)	C8—H8	0.9300
N1—C12	1.345 (5)	C9—C10	1.370 (7)
N2—C10	1.342 (5)	C9—H9	0.9300
N2—C11	1.371 (5)	C10—C14	1.514 (7)

C1—C2	1.407 (6)	C11—C12	1.454 (5)
C1—C13	1.482 (6)	C13—H13A	0.9600
C2—C3	1.341 (6)	C13—H13B	0.9600
C2—H2	0.9300	C13—H13C	0.9600
C3—C4	1.428 (6)	C14—H14A	0.9600
C3—H3	0.9300	C14—H14B	0.9600
C4—C12	1.387 (5)	C14—H14C	0.9600
C4—C5	1.399 (6)		
N2—Mn1—N1	79.57 (12)	C7—C6—H6	119.2
N2—Mn1—C11	111.79 (4)	C11—C7—C6	119.7 (4)
N1—Mn1—C11	113.70 (4)	C11—C7—C8	115.5 (4)
N2—Mn1—C11 ⁱ	111.79 (4)	C6—C7—C8	124.8 (4)
N1—Mn1—C11 ⁱ	113.70 (4)	C9—C8—C7	120.4 (4)
C11—Mn1—C11 ⁱ	119.53 (5)	C9—C8—H8	119.8
O1 ⁱⁱ —O1—H2A	109.0	C7—C8—H8	119.8
O1 ⁱⁱ —O1—H1A	119.0	C8—C9—C10	120.4 (4)
H2A—O1—H1A	104.5	C8—C9—H9	119.8
C1—N1—C12	117.9 (3)	C10—C9—H9	119.8
C1—N1—Mn1	129.1 (3)	N2—C10—C9	121.2 (4)
C12—N1—Mn1	113.0 (2)	N2—C10—C14	118.4 (4)
C10—N2—C11	119.1 (3)	C9—C10—C14	120.3 (4)
C10—N2—Mn1	128.4 (3)	C7—C11—N2	123.3 (3)
C11—N2—Mn1	112.5 (2)	C7—C11—C12	118.2 (3)
N1—C1—C2	122.5 (3)	N2—C11—C12	118.5 (3)
N1—C1—C13	116.6 (4)	N1—C12—C4	123.0 (3)
C2—C1—C13	121.0 (4)	N1—C12—C11	116.5 (3)
C3—C2—C1	120.0 (4)	C4—C12—C11	120.5 (3)
C3—C2—H2	120.0	C1—C13—H13A	109.5
C1—C2—H2	120.0	C1—C13—H13B	109.5
C2—C3—C4	118.5 (4)	H13A—C13—H13B	109.5
C2—C3—H3	120.8	C1—C13—H13C	109.5
C4—C3—H3	120.8	H13A—C13—H13C	109.5
C12—C4—C5	119.2 (4)	H13B—C13—H13C	109.5
C12—C4—C3	118.2 (4)	C10—C14—H14A	109.5
C5—C4—C3	122.6 (4)	C10—C14—H14B	109.5
C6—C5—C4	120.7 (4)	H14A—C14—H14B	109.5
C6—C5—H5	119.7	C10—C14—H14C	109.5
C4—C5—H5	119.7	H14A—C14—H14C	109.5
C5—C6—C7	121.7 (4)	H14B—C14—H14C	109.5
C5—C6—H6	119.2		

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