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## Structure Reports

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# Bis[(2-aminophenyl)methanol- $\kappa^2O,N$ ]-bis(nitrato- $\kappa O$ )manganese(II)

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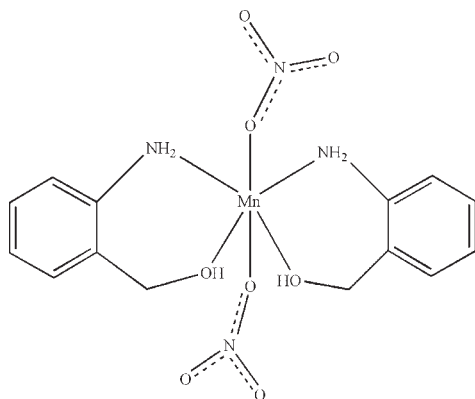
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Key indicators: single-crystal X-ray study;  $T = 120$  K; mean  $\sigma(C-C) = 0.002$  Å;  $R$  factor = 0.038;  $wR$  factor = 0.071; data-to-parameter ratio = 17.3.

In the title compound,  $[Mn(NO_3)_2(C_7H_9NO)_2]$ , the  $Mn^{II}$  atom (site symmetry 2) is coordinated by two  $N,O$ -bidentate (2-aminophenyl)methanol ligands and two monodentate nitrate anions in a distorted  $cis$ - $MnN_2O_4$  octahedral coordination geometry. In the crystal,  $N-H\cdots O$ ,  $O-H\cdots O$  and  $C-H\cdots O$  hydrogen bonds help to establish the packing.

## Related literature

For structures involving the same ligand with other metal ions, see: Bandoli *et al.* (2002); Lewirski *et al.* (1998).



## Experimental

### Crystal data

$[Mn(NO_3)_2(C_7H_9NO)_2]$   $b = 10.1929$  (12) Å  
 $M_r = 425.26$   $c = 7.3336$  (6) Å  
 Orthorhombic,  $Pbcn$   $V = 1747.2$  (3) Å<sup>3</sup>  
 $a = 23.374$  (2) Å  $Z = 4$

Mo  $K\alpha$  radiation  
 $\mu = 0.81$  mm<sup>-1</sup>

$T = 120$  K  
 $0.40 \times 0.10 \times 0.06$  mm

### Data collection

Bruker SMART CCD diffractometer  
 Absorption correction: multi-scan (SADABS; Bruker, 1998)  
 $T_{min} = 0.907$ ,  $T_{max} = 0.955$

6668 measured reflections  
 2335 independent reflections  
 1928 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.046$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$   
 $wR(F^2) = 0.071$   
 $S = 1.10$   
 2335 reflections  
 135 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{max} = 0.31$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.32$  e Å<sup>-3</sup>

**Table 1**

Selected bond lengths (Å).

Mn1—N1	2.2469 (15)	Mn1—O2	2.2203 (12)
Mn1—O1	2.2041 (13)		

**Table 2**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1C $\cdots$ O3 <sup>i</sup>	0.87 (2)	2.16 (2)	2.9775 (19)	156 (2)
N1—H1D $\cdots$ O4 <sup>ii</sup>	0.90 (3)	2.15 (3)	3.037 (2)	169 (2)
O1—H1E $\cdots$ O2 <sup>iii</sup>	0.82 (3)	1.88 (3)	2.6937 (17)	176 (3)
C1—H1B $\cdots$ O4 <sup>ii</sup>	0.97	2.60	3.479 (2)	151

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

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

We are grateful to the University of Urmieh for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5561).

## References

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## supporting information

*Acta Cryst.* (2010). E66, m1052 [https://doi.org/10.1107/S1600536810030114]

**Bis[(2-aminophenyl)methanol- $\kappa^2$ O,N]bis(nitrato- $\kappa$ O)manganese(II)****Majid Esmhosseini and Shahrzad Maleki****S1. Comment**

(2-Aminophenyl)methanol is a bidentate ligand. There are only two complexes with this ligand that have been prepared: those of Re (Bandoli *et al.*, 2002) and Al (Lewiriski *et al.* 1998). We report herein the synthesis and crystal structure of the title compound, (I).

The asymmetric unit of the title compound, Fig. 1, contains half molecule. The Mn<sup>II</sup> atom is six-coordinated in distorted hexagonal configurations by two N and two O atoms from two (2-aminophenyl)methanol ligand and two O atoms from two nitrate anions. The Mn—O and Mn—N bond lengths and angles are collected in Table 1.

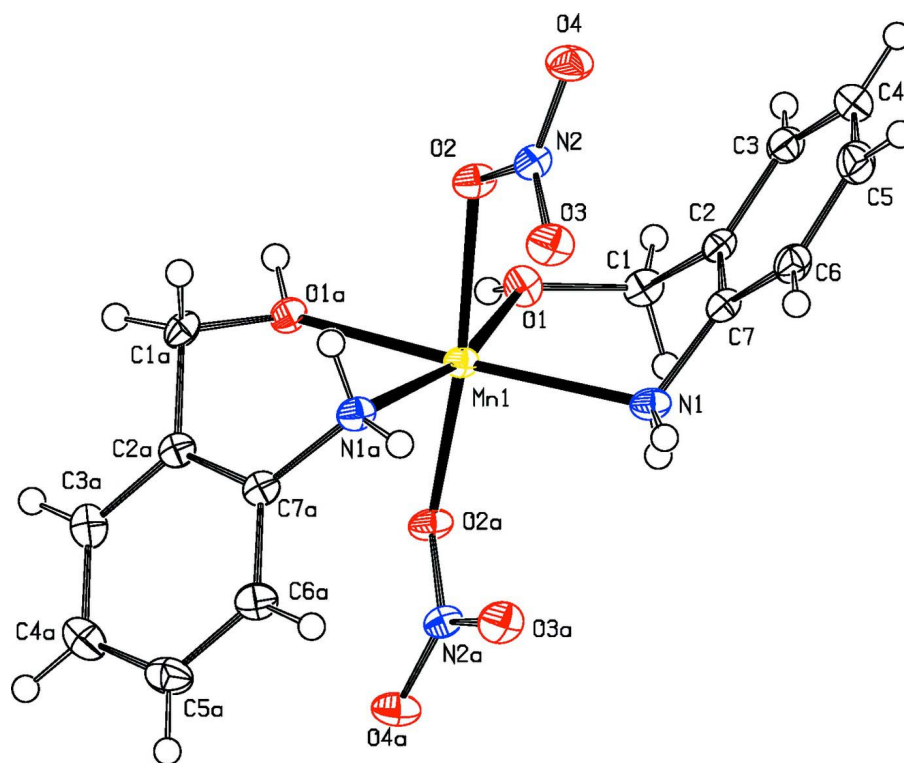
Intermolecular N—H $\cdots$ O, O—H $\cdots$ O and C—H $\cdots$ O hydrogen bonding may stabilize the structure, (Table 2, Fig. 2).

**S2. Experimental**

A solution of (2-aminophenyl)methanol (0.25 g, 2.00 mmol) in methanol (10 ml) was added to a solution of Mn(NO<sub>3</sub>)<sub>2</sub>·4H<sub>2</sub>O (0.25 g, 1.00 mmol) in methanol (10 ml) and the resulting colorless solution was stirred for 20 min at 313 K. This solution was left to evaporate slowly at room temperature. After one week, colorless needles of (I) were isolated (yield 0.33 g, 77.6%).

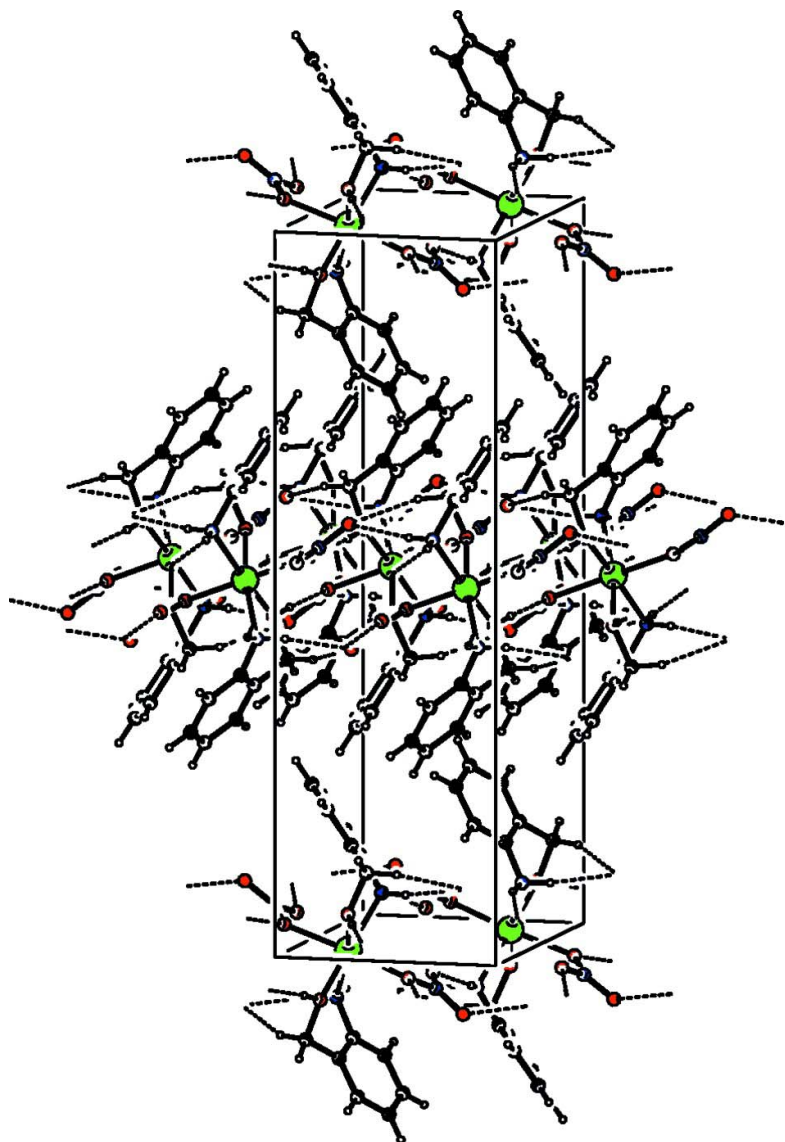
**S3. Refinement**

The N- and O-bound H atoms were located in a difference map and freely refined. All C-bound H atoms were positioned geometrically, with C—H = 0.93 Å and constrained to ride on their parent atoms, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .



**Figure 1**

The molecular structure of (I) with displacement ellipsoids drawn at the 50% probability level. Atoms with suffix a are generated by  $(1-x, y, 1/2-z)$ .



**Figure 2**

Unit-cell packing diagram for (I). Hydrogen bonds are shown as dashed lines.

**Bis[(2-aminophenyl)methanol- $\kappa^2O,N$ ]bis(nitrato- $\kappa O$ )manganese(II)**

*Crystal data*

$[\text{Mn}(\text{NO}_3)_2(\text{C}_7\text{H}_9\text{NO})_2]$

$M_r = 425.26$

Orthorhombic, *Pbcn*

Hall symbol:  $-P\ 2n\ 2ab$

$a = 23.374\ (2)\ \text{\AA}$

$b = 10.1929\ (12)\ \text{\AA}$

$c = 7.3336\ (6)\ \text{\AA}$

$V = 1747.2\ (3)\ \text{\AA}^3$

$Z = 4$

$F(000) = 876$

$D_x = 1.617\ \text{Mg m}^{-3}$

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

Cell parameters from 1154 reflections

$\theta = 2.2\text{--}29.2^\circ$

$\mu = 0.81\ \text{mm}^{-1}$

$T = 120\ \text{K}$

Block, colorless

$0.40 \times 0.10 \times 0.06\ \text{mm}$

*Data collection*

Bruker SMART CCD diffractometer	6668 measured reflections
Radiation source: fine-focus sealed tube	2335 independent reflections
Graphite monochromator	1928 reflections with $I > 2\sigma(I)$
phi and $\omega$ scans	$R_{\text{int}} = 0.046$
Absorption correction: multi-scan (SADABS; Bruker, 1998)	$\theta_{\text{max}} = 29.2^\circ$ , $\theta_{\text{min}} = 2.2^\circ$
$T_{\text{min}} = 0.907$ , $T_{\text{max}} = 0.955$	$h = -31 \rightarrow 23$
	$k = -13 \rightarrow 10$
	$l = -9 \rightarrow 9$

*Refinement*

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.038$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.071$	$w = 1/[\sigma^2(F_o^2) + (0.017P)^2 + 1.3224P]$
$S = 1.10$	where $P = (F_o^2 + 2F_c^2)/3$
2335 reflections	$(\Delta/\sigma)_{\text{max}} = 0.007$
135 parameters	$\Delta\rho_{\text{max}} = 0.31 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.32 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.61659 (7)	0.43167 (16)	0.1127 (2)	0.0177 (3)
H1A	0.6380	0.5132	0.1158	0.021*
H1B	0.6137	0.4033	-0.0132	0.021*
C2	0.64736 (7)	0.32879 (16)	0.2234 (2)	0.0160 (3)
C3	0.69730 (8)	0.36086 (18)	0.3180 (2)	0.0208 (3)
H3	0.7119	0.4455	0.3095	0.025*
C4	0.72554 (8)	0.26846 (19)	0.4247 (3)	0.0248 (4)
H4	0.7586	0.2911	0.4876	0.030*
C5	0.70376 (8)	0.14206 (19)	0.4363 (3)	0.0248 (4)
H5	0.7221	0.0802	0.5087	0.030*
C6	0.65484 (8)	0.10717 (17)	0.3408 (2)	0.0197 (3)
H6	0.6411	0.0217	0.3473	0.024*
C7	0.62624 (7)	0.20036 (16)	0.2348 (2)	0.0162 (3)
N1	0.57317 (7)	0.16868 (14)	0.1479 (2)	0.0168 (3)
H1D	0.5731 (10)	0.194 (2)	0.030 (4)	0.033 (6)*

H1C	0.5657 (10)	0.085 (2)	0.156 (3)	0.027 (6)*
N2	0.55303 (6)	0.21326 (14)	0.61414 (18)	0.0165 (3)
O1	0.55977 (5)	0.45233 (12)	0.18714 (17)	0.0181 (2)
H1E	0.5502 (11)	0.522 (3)	0.141 (4)	0.046 (8)*
O2	0.53386 (6)	0.31604 (10)	0.53077 (16)	0.0184 (3)
O3	0.53757 (6)	0.10431 (11)	0.55841 (18)	0.0241 (3)
O4	0.58609 (6)	0.22845 (13)	0.74423 (18)	0.0249 (3)
Mn1	0.5000	0.29131 (3)	0.2500	0.01412 (9)

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0222 (8)	0.0131 (7)	0.0179 (8)	−0.0029 (6)	0.0022 (7)	0.0018 (6)
C2	0.0186 (7)	0.0156 (7)	0.0137 (8)	0.0016 (6)	0.0017 (6)	−0.0019 (6)
C3	0.0197 (8)	0.0212 (8)	0.0216 (8)	−0.0015 (7)	0.0024 (7)	−0.0055 (7)
C4	0.0196 (8)	0.0306 (10)	0.0243 (8)	0.0028 (8)	−0.0040 (7)	−0.0056 (8)
C5	0.0244 (9)	0.0276 (10)	0.0224 (8)	0.0090 (8)	−0.0023 (7)	0.0006 (8)
C6	0.0237 (8)	0.0152 (7)	0.0203 (8)	0.0043 (7)	0.0000 (7)	−0.0005 (7)
C7	0.0191 (7)	0.0151 (7)	0.0143 (7)	0.0014 (6)	0.0016 (6)	−0.0017 (6)
N1	0.0230 (7)	0.0109 (6)	0.0165 (7)	−0.0018 (5)	−0.0026 (6)	−0.0005 (5)
N2	0.0213 (7)	0.0139 (6)	0.0144 (6)	0.0013 (6)	0.0004 (5)	0.0012 (5)
O1	0.0204 (6)	0.0102 (5)	0.0238 (6)	−0.0002 (5)	−0.0008 (5)	0.0045 (5)
O2	0.0288 (6)	0.0087 (5)	0.0177 (6)	0.0011 (5)	−0.0042 (5)	0.0014 (4)
O3	0.0316 (7)	0.0103 (5)	0.0305 (7)	−0.0017 (5)	−0.0047 (6)	0.0007 (5)
O4	0.0308 (6)	0.0276 (7)	0.0163 (6)	0.0021 (5)	−0.0072 (6)	0.0004 (6)
Mn1	0.01774 (16)	0.00979 (14)	0.01484 (16)	0.000	−0.00214 (14)	0.000

*Geometric parameters (Å, °)*

C1—O1	1.451 (2)	C7—N1	1.431 (2)
C1—C2	1.509 (2)	Mn1—N1	2.2469 (15)
C1—H1A	0.9700	N1—H1D	0.90 (3)
C1—H1B	0.9700	N1—H1C	0.87 (2)
C2—C3	1.397 (2)	N2—O3	1.2372 (19)
C2—C7	1.402 (2)	N2—O4	1.2375 (18)
C3—C4	1.391 (3)	N2—O2	1.2931 (18)
C3—H3	0.9300	Mn1—O1	2.2041 (13)
C4—C5	1.388 (3)	O1—H1E	0.82 (3)
C4—H4	0.9300	Mn1—O2	2.2203 (12)
C5—C6	1.387 (3)	Mn1—O1 <sup>i</sup>	2.2041 (13)
C5—H5	0.9300	Mn1—O2 <sup>i</sup>	2.2202 (12)
C6—C7	1.398 (2)	Mn1—N1 <sup>i</sup>	2.2469 (15)
C6—H6	0.9300		
O1—C1—C2	109.56 (13)	Mn1—N1—H1D	99.2 (16)
O1—C1—H1A	109.8	C7—N1—H1C	111.3 (15)
C2—C1—H1A	109.8	Mn1—N1—H1C	111.6 (15)
O1—C1—H1B	109.8	H1D—N1—H1C	110 (2)

C2—C1—H1B	109.8	O3—N2—O4	123.32 (14)
H1A—C1—H1B	108.2	O3—N2—O2	118.03 (13)
C3—C2—C7	118.92 (15)	O4—N2—O2	118.65 (14)
C3—C2—C1	120.20 (15)	C1—O1—Mn1	123.42 (10)
C7—C2—C1	120.88 (15)	C1—O1—H1E	102.7 (19)
C4—C3—C2	121.16 (17)	Mn1—O1—H1E	124.1 (19)
C4—C3—H3	119.4	N2—O2—Mn1	118.03 (10)
C2—C3—H3	119.4	O1 <sup>i</sup> —Mn1—O1	83.74 (7)
C5—C4—C3	119.26 (17)	O1 <sup>i</sup> —Mn1—O2 <sup>i</sup>	83.31 (5)
C5—C4—H4	120.4	O1—Mn1—O2 <sup>i</sup>	86.99 (5)
C3—C4—H4	120.4	O1 <sup>i</sup> —Mn1—O2	86.99 (5)
C6—C5—C4	120.63 (17)	O1—Mn1—O2	83.31 (5)
C6—C5—H5	119.7	O2 <sup>i</sup> —Mn1—O2	166.96 (6)
C4—C5—H5	119.7	O1 <sup>i</sup> —Mn1—N1 <sup>i</sup>	82.07 (5)
C5—C6—C7	120.05 (17)	O1—Mn1—N1 <sup>i</sup>	165.12 (5)
C5—C6—H6	120.0	O2 <sup>i</sup> —Mn1—N1 <sup>i</sup>	95.79 (5)
C7—C6—H6	120.0	O2—Mn1—N1 <sup>i</sup>	91.45 (5)
C6—C7—C2	119.96 (15)	O1 <sup>i</sup> —Mn1—N1	165.11 (5)
C6—C7—N1	120.58 (15)	O1—Mn1—N1	82.07 (5)
C2—C7—N1	119.31 (14)	O2 <sup>i</sup> —Mn1—N1	91.45 (5)
C7—N1—Mn1	112.68 (10)	O2—Mn1—N1	95.79 (5)
C7—N1—H1D	111.3 (16)	N1 <sup>i</sup> —Mn1—N1	112.40 (8)
O1—C1—C2—C3	-117.48 (17)	O4—N2—O2—Mn1	159.69 (11)
O1—C1—C2—C7	61.89 (19)	C1—O1—Mn1—O1 <sup>i</sup>	-175.17 (15)
C7—C2—C3—C4	-1.1 (3)	C1—O1—Mn1—O2 <sup>i</sup>	-91.58 (12)
C1—C2—C3—C4	178.31 (16)	C1—O1—Mn1—O2	97.15 (12)
C2—C3—C4—C5	0.4 (3)	C1—O1—Mn1—N1 <sup>i</sup>	167.15 (17)
C3—C4—C5—C6	0.9 (3)	C1—O1—Mn1—N1	0.31 (12)
C4—C5—C6—C7	-1.4 (3)	N2—O2—Mn1—O1 <sup>i</sup>	148.15 (12)
C5—C6—C7—C2	0.7 (2)	N2—O2—Mn1—O1	-127.80 (12)
C5—C6—C7—N1	-174.78 (16)	N2—O2—Mn1—O2 <sup>i</sup>	-169.96 (11)
C3—C2—C7—C6	0.5 (2)	N2—O2—Mn1—N1 <sup>i</sup>	66.17 (12)
C1—C2—C7—C6	-178.84 (15)	N2—O2—Mn1—N1	-46.52 (12)
C3—C2—C7—N1	176.07 (15)	C7—N1—Mn1—O1 <sup>i</sup>	69.1 (2)
C1—C2—C7—N1	-3.3 (2)	C7—N1—Mn1—O1	51.34 (11)
C6—C7—N1—Mn1	117.00 (14)	C7—N1—Mn1—O2 <sup>i</sup>	138.10 (11)
C2—C7—N1—Mn1	-58.50 (17)	C7—N1—Mn1—O2	-31.05 (12)
C2—C1—O1—Mn1	-49.35 (17)	C7—N1—Mn1—N1 <sup>i</sup>	-125.03 (12)
O3—N2—O2—Mn1	-19.78 (18)		

Symmetry code: (i)  $-x+1, y, -z+1/2$ .

#### Hydrogen-bond geometry ( $\text{\AA}$ , $^\circ$ )

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1C $\cdots$ O3 <sup>ii</sup>	0.87 (2)	2.16 (2)	2.9775 (19)	156 (2)
N1—H1D $\cdots$ O4 <sup>iii</sup>	0.90 (3)	2.15 (3)	3.037 (2)	169 (2)

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O1—H1E···O2 <sup>iv</sup>	0.82 (3)	1.88 (3)	2.6937 (17)	176 (3)
C1—H1B···O4 <sup>iii</sup>	0.97	2.60	3.479 (2)	151

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