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Bis[2-methoxy-6-[tris(hydroxymethyl)-methyliminomethyl]phenolato- κ^3O,N,O']manganese(II) dimethanol solvate hemihydrate

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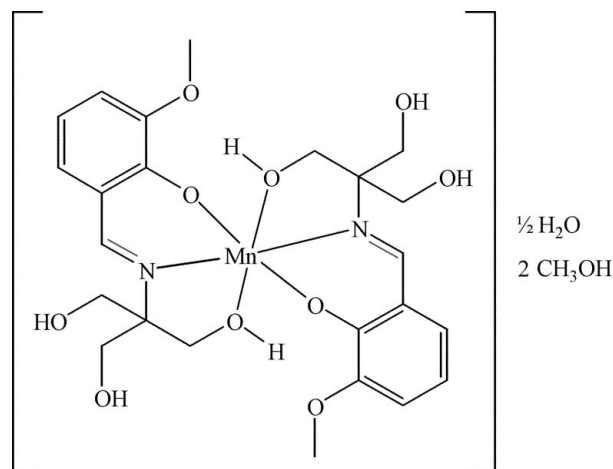
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(C-C) = 0.005$ Å; disorder in main residue; R factor = 0.051; wR factor = 0.148; data-to-parameter ratio = 12.7.

In the title complex, $[Mn(C_{12}H_{16}NO_5)_2] \cdot 2CH_3OH \cdot 0.5H_2O$, the Mn^{II} atom has a distorted octahedral coordination geometry in which two N atoms from two 6-methoxy-2-[tris(hydroxymethyl)methyliminomethyl]phenolate ligands adopt a *trans* arrangement. The $Mn-O(H)$ bonds (mean length 2.134 Å) are significantly longer than the $Mn-O$ and $Mn-N$ bonds (mean length 2.011 and 2.027 Å, respectively), and the dihedral angle between the mean planes through the aromatic rings of the two ligands is 76.8 (1)°. A complex network of $O-H \cdots O$ hydrogen bonds is formed between the complexes and the uncoordinated methanol and water molecules. The C and O atoms of one C-OH group are disordered with equal occupancies.

Related literature

For Schiff-base complexes of transition metals, see: Ward (2007). For details of the synthesis and a related structure, see: Wang *et al.* (2007).



Experimental

Crystal data

 $[Mn(C_{12}H_{16}NO_5)_2] \cdot 2CH_3OH \cdot 0.5H_2O$
 $M_r = 636.55$

 Monoclinic, $P2_1/n$
 $a = 8.141$ (2) Å

 $b = 18.130$ (5) Å

 $c = 20.211$ (6) Å

 $\beta = 93.590$ (4)°

 $V = 2977.2$ (14) Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 0.51$ mm⁻¹
 $T = 293$ K

 $0.12 \times 0.10 \times 0.09$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer

Absorption correction: multi-scan

 (*SADABS*; Sheldrick, 2003)

 $T_{min} = 0.941$, $T_{max} = 0.955$

14576 measured reflections

5287 independent reflections

 4036 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.031$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.148$
 $S = 1.05$

5287 reflections

416 parameters

16 restraints

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{max} = 0.93$ e Å⁻³
 $\Delta\rho_{min} = -0.65$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|--------------------------------------|----------|--------------|--------------|----------------|
| O3—H3 \cdots O8 | 0.86 (1) | 1.885 (10) | 2.706 (5) | 160 (4) |
| O4—H4 \cdots O9 | 0.86 (1) | 1.802 (12) | 2.664 (5) | 174 (5) |
| O5—H5 \cdots O2 ⁱ | 0.85 (1) | 1.886 (13) | 2.736 (4) | 173 (6) |
| O6—H6 \cdots O3 | 0.86 (1) | 1.886 (9) | 2.737 (7) | 172 (6) |
| O9—H9 \cdots O11 | 0.85 (1) | 1.888 (10) | 2.712 (5) | 162 (4) |
| O10—H10 \cdots O1W | 0.86 (1) | 1.90 (4) | 2.585 (6) | 136 (5) |
| O11—H11 \cdots O6 ⁱⁱ | 0.85 (1) | 1.897 (18) | 2.739 (4) | 169 (7) |
| O12—H12 \cdots O2 | 0.85 | 2.12 | 2.967 (7) | 180 |
| O12A—H12A \cdots O5 ⁱⁱⁱ | 0.85 | 2.10 | 2.946 (7) | 180 |
| O1W—H1W \cdots O5 | 0.85 | 1.81 | 2.657 (8) | 180 |
| O1W—H2W \cdots O12A ^{iv} | 0.85 | 2.14 | 2.990 (9) | 180 |

Symmetry codes: (i) $x - 1, y, z$; (ii) $x + \frac{1}{2}, -y + \frac{3}{2}, z + \frac{1}{2}$; (iii) $x + 1, y, z$; (iv) $-x + 1, -y + 1, -z + 1$.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to

solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); 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: BI2342).

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Wang, Q., Li, X., Wang, X. & Zhang, Y. (2007). *Acta Cryst.* **E63**, m2537.
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supporting information

Acta Cryst. (2009). E65, m293–m294 [doi:10.1107/S1600536809005364]

Bis{2-methoxy-6-[tris(hydroxymethyl)methyliminomethyl]phenolato- κ^3O,N,O }manganese(II) dimethanol solvate hemihydrate

Xiutang Zhang, Peihai Wei, Jianmin Dou, Bin Li and Bo Hu

S1. Comment

Transition metal Schiff-base complexes have been intensively studied owing to their interesting physical and chemical properties, including magnetic, optics and catalysis (Ward *et al.*). Herein, we report a crystal structure of an Mn^{II} complex incorporating the Schiff-base ligand, (*E*)-2-(2-hydroxy-3-methoxybenzylideneamino)-2-(hydroxymethyl)propane-1,3-diol.

The asymmetric unit (Fig. 1) comprises one [MnL₂] complex, two uncoordinated methanol molecules and one lattice water molecule. The coordination geometry of Mn^{II} is distorted octahedral, with the N atoms of the two ligands *trans* to each other. The Mn—O and Mn—N bond distances are in the normal range compared to similar reported complexes (for example, Wang *et al.*, 2007). A complex network of O—H \cdots O hydrogen bonds is formed between the complexes and the lattice methanol and water molecules.

S2. Experimental

The Schiff-base ligand (HL) was synthesized according to the reported literature procedure (Wang *et al.*). The title complex was then prepared by refluxing HL (0.050 g, 0.2 mmol) and MnSO₄·H₂O in the mixed solvent system CH₃OH:H₂O (4:1) until all solid was dissolved. The solution was then cooled to room temperature and filtered. Crystals for diffraction analysis were obtained by slow evaporation of the filtrate. Elemental analysis calculated: C 48.05, H 6.47, N 4.31%; found: C 49.89, H 6.39, N 4.28%.

S3. Refinement

H atoms bound to C atoms were refined using a riding model with C—H = 0.93 Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic H atoms, C—H = 0.96 Å, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and C—H = 0.97 Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for methylene H atoms. H atoms bound to most of the O atoms were located in difference Fourier maps and refined with O—H restrained to be 0.85 (1) Å and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. The exceptions were for the disordered C—OH groups and the lattice water molecules: in these cases, the H atoms were placed so as to form reasonable H-bonds with O—H = 0.85 Å and refined as riding with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. The C—O bonds of the disordered C—OH groups and the lattice methanol molecules were restrained to a common refined value with an uncertainty of 0.02 %Å.

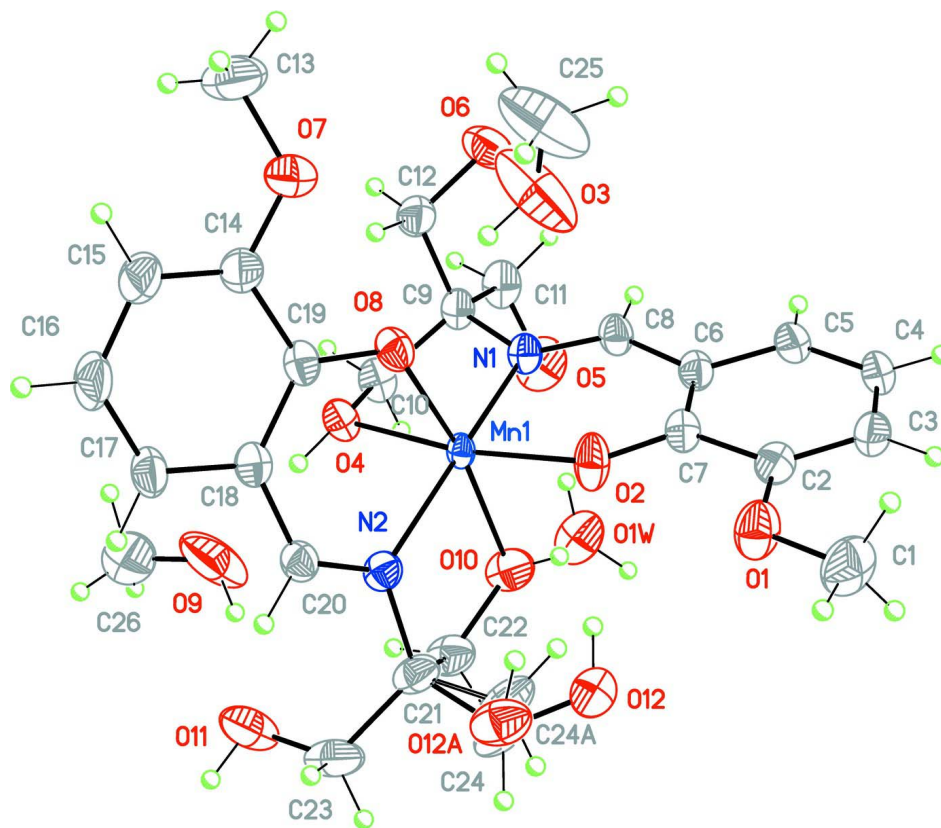


Figure 1

The molecular structure with displacement ellipsoids at 30% probability for non-H atoms.

Bis[2-methoxy-6-[tris(hydroxymethyl)methyliminomethyl]phenolato- κ^3O,N,O']manganese(II) dimethanol solvate hemihydrate

Crystal data

$[\text{Mn}(\text{C}_{12}\text{H}_{16}\text{NO}_5)_2] \cdot 2\text{CH}_4\text{O} \cdot 0.5\text{H}_2\text{O}$

$M_r = 636.55$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1/n$

$a = 8.141\ (2)\ \text{\AA}$

$b = 18.130\ (5)\ \text{\AA}$

$c = 20.211\ (6)\ \text{\AA}$

$\beta = 93.590\ (4)^\circ$

$V = 2977.2\ (14)\ \text{\AA}^3$

$Z = 4$

$F(000) = 1344$

$D_x = 1.420\ \text{Mg m}^{-3}$

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

Cell parameters from 5287 reflections

$\theta = 2.0\text{--}25.3^\circ$

$\mu = 0.51\ \text{mm}^{-1}$

$T = 293\ \text{K}$

Block, pink

$0.12 \times 0.10 \times 0.09\ \text{mm}$

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, 2003)

$T_{\min} = 0.941$, $T_{\max} = 0.955$

14576 measured reflections

5287 independent reflections

4036 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.031$

$\theta_{\max} = 25.3^\circ$, $\theta_{\min} = 2.0^\circ$

$h = -9 \rightarrow 9$

$k = -21 \rightarrow 17$

$l = -22 \rightarrow 24$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.148$
 $S = 1.05$
 5287 reflections
 416 parameters
 16 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0796P)^2 + 1.8727P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.93 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.65 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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) |
|------|-------------|--------------|--------------|----------------------------------|-----------|
| C1 | 0.8762 (6) | 0.5053 (3) | 0.3266 (3) | 0.0813 (14) | |
| H1A | 0.9812 | 0.5090 | 0.3509 | 0.122* | |
| H1B | 0.8876 | 0.5190 | 0.2813 | 0.122* | |
| H1C | 0.8368 | 0.4555 | 0.3285 | 0.122* | |
| C2 | 0.6105 (4) | 0.56122 (19) | 0.32411 (18) | 0.0476 (8) | |
| C3 | 0.5514 (5) | 0.5215 (2) | 0.26979 (19) | 0.0574 (10) | |
| H3A | 0.6184 | 0.4870 | 0.2506 | 0.069* | |
| C4 | 0.3910 (6) | 0.5330 (2) | 0.24340 (18) | 0.0604 (10) | |
| H4A | 0.3507 | 0.5059 | 0.2068 | 0.072* | |
| C5 | 0.2934 (5) | 0.5839 (2) | 0.27131 (17) | 0.0521 (9) | |
| H5A | 0.1873 | 0.5919 | 0.2530 | 0.062* | |
| C6 | 0.3507 (4) | 0.62479 (18) | 0.32757 (16) | 0.0440 (8) | |
| C7 | 0.5118 (4) | 0.61434 (18) | 0.35482 (16) | 0.0430 (8) | |
| C8 | 0.2333 (4) | 0.67685 (19) | 0.35280 (17) | 0.0450 (8) | |
| H8A | 0.1325 | 0.6813 | 0.3288 | 0.054* | |
| C9 | 0.1240 (4) | 0.76952 (19) | 0.42168 (17) | 0.0447 (8) | |
| C10 | 0.1292 (4) | 0.7749 (2) | 0.49750 (18) | 0.0524 (9) | |
| H10A | 0.0662 | 0.8172 | 0.5106 | 0.063* | |
| H10B | 0.0813 | 0.7309 | 0.5157 | 0.063* | |
| C11 | -0.0517 (4) | 0.7489 (2) | 0.3971 (2) | 0.0596 (10) | |
| H11A | -0.1283 | 0.7852 | 0.4124 | 0.071* | |
| H11B | -0.0608 | 0.7486 | 0.3490 | 0.071* | |
| C12 | 0.1665 (5) | 0.8444 (2) | 0.39232 (19) | 0.0563 (9) | |
| H12B | 0.0816 | 0.8796 | 0.4021 | 0.068* | |

| | | | | | |
|------|-------------|--------------|--------------|--------------|------|
| H12C | 0.2694 | 0.8617 | 0.4137 | 0.068* | |
| C13 | 0.5914 (9) | 1.0165 (3) | 0.3854 (3) | 0.1014 (18) | |
| H13A | 0.5234 | 1.0219 | 0.3452 | 0.152* | |
| H13B | 0.7017 | 1.0315 | 0.3780 | 0.152* | |
| H13C | 0.5488 | 1.0467 | 0.4194 | 0.152* | |
| C14 | 0.6726 (6) | 0.9237 (2) | 0.46473 (19) | 0.0594 (10) | |
| C15 | 0.7689 (7) | 0.9712 (2) | 0.5036 (2) | 0.0746 (13) | |
| H15A | 0.7847 | 1.0194 | 0.4894 | 0.090* | |
| C16 | 0.8427 (6) | 0.9479 (2) | 0.5637 (2) | 0.0721 (12) | |
| H16A | 0.9086 | 0.9800 | 0.5895 | 0.087* | |
| C17 | 0.8178 (5) | 0.8778 (2) | 0.58459 (19) | 0.0547 (9) | |
| H17A | 0.8657 | 0.8629 | 0.6254 | 0.066* | |
| C18 | 0.7211 (4) | 0.82661 (19) | 0.54598 (17) | 0.0445 (8) | |
| C19 | 0.6493 (4) | 0.84878 (19) | 0.48343 (16) | 0.0428 (8) | |
| C20 | 0.7043 (4) | 0.75505 (19) | 0.57352 (17) | 0.0441 (8) | |
| H20A | 0.7661 | 0.7456 | 0.6129 | 0.053* | |
| C21 | 0.6069 (5) | 0.63052 (19) | 0.5864 (2) | 0.0567 (10) | |
| C22 | 0.4262 (5) | 0.6059 (2) | 0.5794 (2) | 0.0630 (11) | |
| H22A | 0.4184 | 0.5544 | 0.5918 | 0.076* | |
| H22B | 0.3624 | 0.6346 | 0.6090 | 0.076* | |
| C23 | 0.6610 (6) | 0.6332 (3) | 0.6597 (2) | 0.0782 (14) | |
| H23A | 0.6497 | 0.5844 | 0.6786 | 0.094* | |
| H23B | 0.7764 | 0.6468 | 0.6646 | 0.094* | |
| C24 | 0.7115 (19) | 0.5685 (9) | 0.5636 (8) | 0.076 (4) | 0.50 |
| H24A | 0.8226 | 0.5871 | 0.5609 | 0.092* | 0.50 |
| H24B | 0.7156 | 0.5308 | 0.5977 | 0.092* | 0.50 |
| O12 | 0.6698 (8) | 0.5379 (3) | 0.5091 (3) | 0.0742 (17) | 0.50 |
| H12 | 0.6435 | 0.5701 | 0.4797 | 0.111* | 0.50 |
| C24A | 0.7240 (16) | 0.5822 (9) | 0.5488 (10) | 0.076 (4) | 0.50 |
| H24C | 0.7127 | 0.5329 | 0.5666 | 0.092* | 0.50 |
| H24D | 0.6770 | 0.5803 | 0.5035 | 0.092* | 0.50 |
| O12A | 0.8824 (6) | 0.5918 (3) | 0.5439 (3) | 0.0722 (16) | 0.50 |
| H12A | 0.8899 | 0.6168 | 0.5087 | 0.108* | 0.50 |
| C25 | 0.5935 (11) | 0.8441 (6) | 0.2719 (4) | 0.169 (4) | |
| H25A | 0.5563 | 0.8942 | 0.2672 | 0.254* | |
| H25B | 0.5871 | 0.8205 | 0.2293 | 0.254* | |
| H25C | 0.7054 | 0.8436 | 0.2900 | 0.254* | |
| C26 | 0.2701 (11) | 0.8216 (4) | 0.7004 (3) | 0.131 (3) | |
| H26A | 0.3613 | 0.8342 | 0.7307 | 0.197* | |
| H26B | 0.1866 | 0.7976 | 0.7241 | 0.197* | |
| H26C | 0.2256 | 0.8656 | 0.6799 | 0.197* | |
| Mn1 | 0.45435 (5) | 0.71273 (2) | 0.46924 (2) | 0.02874 (16) | |
| N1 | 0.2550 (3) | 0.71681 (14) | 0.40435 (14) | 0.0407 (6) | |
| N2 | 0.6132 (3) | 0.70187 (14) | 0.55018 (14) | 0.0433 (7) | |
| O1 | 0.7639 (3) | 0.55283 (16) | 0.35476 (15) | 0.0694 (8) | |
| O2 | 0.5780 (3) | 0.65014 (15) | 0.40653 (13) | 0.0586 (7) | |
| O3 | 0.5054 (6) | 0.8104 (4) | 0.3093 (2) | 0.159 (2) | |
| H3 | 0.545 (8) | 0.814 (5) | 0.3492 (7) | 0.239* | |

| | | | | | |
|-----|-------------|--------------|--------------|-------------|------|
| O4 | 0.2975 (3) | 0.78214 (14) | 0.52223 (12) | 0.0496 (6) | |
| H4 | 0.297 (6) | 0.782 (2) | 0.5650 (6) | 0.074* | |
| O5 | -0.0915 (3) | 0.67810 (19) | 0.42156 (19) | 0.0826 (10) | |
| H5 | -0.193 (2) | 0.668 (3) | 0.414 (3) | 0.124* | |
| O6 | 0.1812 (4) | 0.84266 (19) | 0.32360 (14) | 0.0779 (9) | |
| H6 | 0.279 (2) | 0.829 (3) | 0.316 (3) | 0.117* | |
| O7 | 0.5916 (5) | 0.94260 (17) | 0.40552 (15) | 0.0847 (10) | |
| O8 | 0.5665 (3) | 0.80546 (13) | 0.44233 (11) | 0.0475 (6) | |
| O9 | 0.3196 (7) | 0.7771 (3) | 0.65421 (19) | 0.145 (2) | |
| H9 | 0.383 (6) | 0.7408 (19) | 0.664 (5) | 0.218* | |
| O10 | 0.3606 (3) | 0.61528 (15) | 0.51303 (15) | 0.0656 (8) | |
| H10 | 0.294 (5) | 0.590 (2) | 0.488 (2) | 0.098* | |
| O11 | 0.5671 (5) | 0.6847 (2) | 0.69558 (16) | 0.0911 (10) | |
| H11 | 0.591 (8) | 0.679 (4) | 0.7369 (9) | 0.137* | |
| O1W | 0.0648 (7) | 0.5679 (4) | 0.4852 (4) | 0.0806 (18) | 0.50 |
| H1W | 0.0147 | 0.6032 | 0.4650 | 0.121* | 0.50 |
| H2W | 0.0795 | 0.5224 | 0.4770 | 0.121* | 0.50 |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C1 | 0.068 (3) | 0.080 (3) | 0.097 (4) | 0.026 (2) | 0.023 (3) | -0.006 (3) |
| C2 | 0.048 (2) | 0.0451 (19) | 0.050 (2) | -0.0014 (16) | 0.0098 (15) | -0.0001 (16) |
| C3 | 0.078 (3) | 0.046 (2) | 0.049 (2) | 0.0068 (19) | 0.0133 (19) | -0.0072 (16) |
| C4 | 0.090 (3) | 0.052 (2) | 0.038 (2) | 0.002 (2) | -0.0006 (19) | -0.0120 (16) |
| C5 | 0.064 (2) | 0.051 (2) | 0.0411 (19) | -0.0031 (17) | -0.0026 (16) | -0.0061 (16) |
| C6 | 0.0481 (19) | 0.0430 (18) | 0.0411 (18) | -0.0053 (15) | 0.0042 (14) | -0.0057 (14) |
| C7 | 0.0434 (18) | 0.0434 (18) | 0.0429 (18) | -0.0056 (14) | 0.0078 (14) | -0.0078 (14) |
| C8 | 0.0393 (18) | 0.050 (2) | 0.0450 (19) | -0.0023 (15) | -0.0040 (14) | -0.0103 (16) |
| C9 | 0.0371 (17) | 0.0474 (19) | 0.049 (2) | 0.0034 (14) | -0.0014 (14) | -0.0109 (15) |
| C10 | 0.0397 (18) | 0.065 (2) | 0.053 (2) | 0.0014 (16) | 0.0040 (15) | -0.0099 (17) |
| C11 | 0.0356 (19) | 0.071 (3) | 0.072 (3) | 0.0055 (18) | -0.0029 (17) | -0.018 (2) |
| C12 | 0.062 (2) | 0.049 (2) | 0.056 (2) | 0.0033 (18) | -0.0075 (18) | -0.0038 (17) |
| C13 | 0.143 (5) | 0.072 (3) | 0.091 (4) | 0.009 (3) | 0.019 (4) | 0.032 (3) |
| C14 | 0.082 (3) | 0.049 (2) | 0.048 (2) | -0.010 (2) | 0.0109 (19) | 0.0011 (17) |
| C15 | 0.115 (4) | 0.047 (2) | 0.063 (3) | -0.026 (2) | 0.009 (3) | -0.0032 (19) |
| C16 | 0.096 (3) | 0.055 (2) | 0.065 (3) | -0.032 (2) | 0.002 (2) | -0.018 (2) |
| C17 | 0.063 (2) | 0.053 (2) | 0.048 (2) | -0.0133 (18) | -0.0015 (17) | -0.0143 (17) |
| C18 | 0.0418 (18) | 0.047 (2) | 0.0447 (19) | -0.0047 (15) | 0.0031 (14) | -0.0075 (15) |
| C19 | 0.0405 (17) | 0.0482 (19) | 0.0404 (18) | -0.0071 (15) | 0.0072 (14) | -0.0087 (15) |
| C20 | 0.0417 (18) | 0.0465 (19) | 0.0434 (18) | -0.0017 (15) | -0.0040 (14) | -0.0038 (15) |
| C21 | 0.053 (2) | 0.0377 (19) | 0.078 (3) | -0.0042 (16) | -0.0116 (19) | 0.0085 (18) |
| C22 | 0.058 (2) | 0.053 (2) | 0.078 (3) | -0.0117 (19) | -0.007 (2) | 0.013 (2) |
| C23 | 0.075 (3) | 0.065 (3) | 0.091 (3) | -0.010 (2) | -0.026 (3) | 0.034 (3) |
| C24 | 0.055 (3) | 0.036 (5) | 0.140 (8) | -0.009 (3) | 0.016 (4) | 0.009 (6) |
| O12 | 0.086 (4) | 0.059 (4) | 0.077 (4) | 0.007 (3) | 0.000 (3) | -0.006 (3) |
| C24A | 0.055 (3) | 0.036 (5) | 0.140 (8) | -0.009 (3) | 0.016 (4) | 0.009 (6) |
| O12A | 0.053 (3) | 0.066 (4) | 0.096 (5) | 0.003 (3) | -0.010 (3) | 0.018 (3) |

| | | | | | | |
|-----|-------------|-------------|-------------|---------------|---------------|---------------|
| C25 | 0.160 (8) | 0.275 (12) | 0.074 (4) | -0.007 (8) | 0.019 (5) | 0.042 (6) |
| C26 | 0.208 (9) | 0.096 (5) | 0.090 (4) | 0.016 (5) | 0.006 (5) | 0.001 (4) |
| Mn1 | 0.0227 (2) | 0.0306 (3) | 0.0325 (3) | -0.00272 (17) | -0.00254 (16) | -0.00772 (18) |
| N1 | 0.0329 (14) | 0.0443 (15) | 0.0449 (16) | -0.0037 (11) | 0.0017 (11) | -0.0078 (12) |
| N2 | 0.0392 (15) | 0.0403 (15) | 0.0499 (16) | -0.0038 (12) | -0.0022 (12) | -0.0013 (12) |
| O1 | 0.0516 (16) | 0.0741 (19) | 0.083 (2) | 0.0111 (14) | 0.0072 (14) | -0.0217 (15) |
| O2 | 0.0378 (13) | 0.0666 (16) | 0.0705 (17) | -0.0002 (12) | -0.0023 (12) | -0.0337 (14) |
| O3 | 0.114 (3) | 0.299 (7) | 0.064 (3) | -0.080 (4) | -0.003 (2) | 0.000 (4) |
| O4 | 0.0462 (13) | 0.0615 (15) | 0.0408 (13) | 0.0030 (11) | -0.0002 (11) | -0.0082 (12) |
| O5 | 0.0400 (15) | 0.088 (2) | 0.119 (3) | -0.0172 (16) | -0.0023 (16) | -0.007 (2) |
| O6 | 0.089 (2) | 0.092 (2) | 0.0502 (17) | -0.0103 (19) | -0.0159 (16) | 0.0114 (15) |
| O7 | 0.128 (3) | 0.0641 (19) | 0.0599 (19) | -0.0113 (19) | -0.0078 (18) | 0.0153 (15) |
| O8 | 0.0468 (13) | 0.0554 (14) | 0.0400 (13) | -0.0114 (11) | 0.0000 (10) | -0.0050 (10) |
| O9 | 0.159 (4) | 0.228 (6) | 0.047 (2) | 0.108 (4) | -0.002 (2) | -0.008 (3) |
| O10 | 0.0547 (16) | 0.0536 (17) | 0.085 (2) | -0.0118 (13) | -0.0213 (14) | 0.0063 (14) |
| O11 | 0.098 (3) | 0.118 (3) | 0.0554 (19) | 0.015 (2) | -0.0102 (18) | 0.020 (2) |
| O1W | 0.050 (3) | 0.067 (4) | 0.123 (5) | -0.010 (3) | 0.000 (3) | -0.010 (4) |

Geometric parameters (Å, °)

| | | | |
|----------|-----------|-----------|------------|
| C1—O1 | 1.402 (5) | C19—O8 | 1.300 (4) |
| C1—H1A | 0.960 | C20—N2 | 1.288 (4) |
| C1—H1B | 0.960 | C20—H20A | 0.930 |
| C1—H1C | 0.960 | C21—N2 | 1.488 (4) |
| C2—O1 | 1.367 (5) | C21—C24 | 1.500 (10) |
| C2—C3 | 1.374 (5) | C21—C23 | 1.521 (6) |
| C2—C7 | 1.422 (5) | C21—C24A | 1.531 (9) |
| C3—C4 | 1.395 (6) | C21—C22 | 1.535 (5) |
| C3—H3A | 0.930 | C22—O10 | 1.422 (5) |
| C4—C5 | 1.363 (5) | C22—H22A | 0.970 |
| C4—H4A | 0.930 | C22—H22B | 0.970 |
| C5—C6 | 1.412 (5) | C23—O11 | 1.432 (7) |
| C5—H5A | 0.930 | C23—H23A | 0.970 |
| C6—C7 | 1.403 (5) | C23—H23B | 0.970 |
| C6—C8 | 1.458 (5) | C24—O12 | 1.261 (13) |
| C7—O2 | 1.316 (4) | C24—H24A | 0.970 |
| C8—N1 | 1.272 (4) | C24—H24B | 0.970 |
| C8—H8A | 0.930 | O12—H12 | 0.850 |
| C9—N1 | 1.490 (4) | C24A—O12A | 1.311 (12) |
| C9—C12 | 1.530 (5) | C24A—H24C | 0.970 |
| C9—C11 | 1.531 (5) | C24A—H24D | 0.970 |
| C9—C10 | 1.533 (5) | O12A—H12A | 0.850 |
| C10—O4 | 1.435 (4) | C25—O3 | 1.234 (8) |
| C10—H10A | 0.970 | C25—H25A | 0.960 |
| C10—H10B | 0.970 | C25—H25B | 0.960 |
| C11—O5 | 1.421 (5) | C25—H25C | 0.960 |
| C11—H11A | 0.970 | C26—O9 | 1.316 (7) |
| C11—H11B | 0.970 | C26—H26A | 0.960 |

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|------------|-----------|----------------|------------|
| C12—O6 | 1.402 (5) | C26—H26B | 0.960 |
| C12—H12B | 0.970 | C26—H26C | 0.960 |
| C12—H12C | 0.970 | Mn1—O8 | 2.005 (2) |
| C13—O7 | 1.400 (5) | Mn1—O2 | 2.017 (2) |
| C13—H13A | 0.960 | Mn1—N1 | 2.023 (3) |
| C13—H13B | 0.960 | Mn1—N2 | 2.030 (3) |
| C13—H13C | 0.960 | Mn1—O4 | 2.129 (2) |
| C14—O7 | 1.373 (5) | Mn1—O10 | 2.138 (3) |
| C14—C15 | 1.377 (6) | O3—H3 | 0.86 (1) |
| C14—C19 | 1.426 (5) | O4—H4 | 0.87 (1) |
| C15—C16 | 1.388 (6) | O5—H5 | 0.85 (1) |
| C15—H15A | 0.930 | O6—H6 | 0.86 (1) |
| C16—C17 | 1.358 (6) | O9—H9 | 0.85 (1) |
| C16—H16A | 0.930 | O10—H10 | 0.85 (1) |
| C17—C18 | 1.419 (5) | O11—H11 | 0.85 (1) |
| C17—H17A | 0.930 | O1W—H1W | 0.850 |
| C18—C19 | 1.417 (5) | O1W—H2W | 0.850 |
| C18—C20 | 1.422 (5) | | |
| O1—C1—H1A | 109.5 | N2—C21—C24A | 102.2 (8) |
| O1—C1—H1B | 109.5 | C23—C21—C24A | 110.5 (9) |
| H1A—C1—H1B | 109.5 | N2—C21—C22 | 105.7 (3) |
| O1—C1—H1C | 109.5 | C24—C21—C22 | 108.2 (7) |
| H1A—C1—H1C | 109.5 | C23—C21—C22 | 108.5 (4) |
| H1B—C1—H1C | 109.5 | C24A—C21—C22 | 114.2 (7) |
| O1—C2—C3 | 124.6 (3) | O10—C22—C21 | 110.7 (3) |
| O1—C2—C7 | 113.8 (3) | O10—C22—H22A | 109.5 |
| C3—C2—C7 | 121.6 (3) | C21—C22—H22A | 109.5 |
| C2—C3—C4 | 120.0 (3) | O10—C22—H22B | 109.5 |
| C2—C3—H3A | 120.0 | C21—C22—H22B | 109.5 |
| C4—C3—H3A | 120.0 | H22A—C22—H22B | 108.1 |
| C5—C4—C3 | 119.9 (3) | O11—C23—C21 | 112.5 (3) |
| C5—C4—H4A | 120.0 | O11—C23—H23A | 109.1 |
| C3—C4—H4A | 120.0 | C21—C23—H23A | 109.1 |
| C4—C5—C6 | 121.1 (4) | O11—C23—H23B | 109.1 |
| C4—C5—H5A | 119.4 | C21—C23—H23B | 109.1 |
| C6—C5—H5A | 119.4 | H23A—C23—H23B | 107.8 |
| C7—C6—C5 | 119.9 (3) | O12—C24—C21 | 118.1 (12) |
| C7—C6—C8 | 124.6 (3) | O12—C24—H24A | 107.8 |
| C5—C6—C8 | 115.5 (3) | C21—C24—H24A | 107.8 |
| O2—C7—C6 | 124.8 (3) | O12—C24—H24B | 107.8 |
| O2—C7—C2 | 117.8 (3) | C21—C24—H24B | 107.8 |
| C6—C7—C2 | 117.4 (3) | H24A—C24—H24B | 107.1 |
| N1—C8—C6 | 126.5 (3) | C24—O12—H12 | 110.5 |
| N1—C8—H8A | 116.8 | O12A—C24A—C21 | 127.4 (11) |
| C6—C8—H8A | 116.8 | O12A—C24A—H24C | 105.5 |
| N1—C9—C12 | 107.2 (3) | C21—C24A—H24C | 105.5 |
| N1—C9—C11 | 115.8 (3) | O12A—C24A—H24D | 105.5 |

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| C12—C9—C11 | 108.7 (3) | C21—C24A—H24D | 105.5 |
| N1—C9—C10 | 107.5 (3) | H24C—C24A—H24D | 106.0 |
| C12—C9—C10 | 109.8 (3) | C24A—O12A—H12A | 104.8 |
| C11—C9—C10 | 107.8 (3) | O3—C25—H25A | 109.7 |
| O4—C10—C9 | 108.7 (3) | O3—C25—H25B | 109.1 |
| O4—C10—H10A | 110.0 | H25A—C25—H25B | 109.5 |
| C9—C10—H10A | 110.0 | O3—C25—H25C | 109.7 |
| O4—C10—H10B | 110.0 | H25A—C25—H25C | 109.5 |
| C9—C10—H10B | 110.0 | H25B—C25—H25C | 109.5 |
| H10A—C10—H10B | 108.3 | O9—C26—H26A | 109.8 |
| O5—C11—C9 | 109.7 (3) | O9—C26—H26B | 109.5 |
| O5—C11—H11A | 109.7 | H26A—C26—H26B | 109.5 |
| C9—C11—H11A | 109.7 | O9—C26—H26C | 109.1 |
| O5—C11—H11B | 109.7 | H26A—C26—H26C | 109.5 |
| C9—C11—H11B | 109.7 | H26B—C26—H26C | 109.5 |
| H11A—C11—H11B | 108.2 | O8—Mn1—O2 | 92.73 (11) |
| O6—C12—C9 | 113.5 (3) | O8—Mn1—N1 | 98.86 (10) |
| O6—C12—H12B | 108.9 | O2—Mn1—N1 | 91.42 (10) |
| C9—C12—H12B | 108.9 | O8—Mn1—N2 | 91.25 (10) |
| O6—C12—H12C | 108.9 | O2—Mn1—N2 | 97.70 (11) |
| C9—C12—H12C | 108.9 | N1—Mn1—N2 | 166.06 (11) |
| H12B—C12—H12C | 107.7 | O8—Mn1—O4 | 86.48 (10) |
| O7—C13—H13A | 109.5 | O2—Mn1—O4 | 170.91 (10) |
| O7—C13—H13B | 109.5 | N1—Mn1—O4 | 79.77 (10) |
| H13A—C13—H13B | 109.5 | N2—Mn1—O4 | 91.37 (11) |
| O7—C13—H13C | 109.5 | O8—Mn1—O10 | 170.32 (10) |
| H13A—C13—H13C | 109.5 | O2—Mn1—O10 | 90.02 (12) |
| H13B—C13—H13C | 109.5 | N1—Mn1—O10 | 90.34 (10) |
| O7—C14—C15 | 124.5 (4) | N2—Mn1—O10 | 79.18 (10) |
| O7—C14—C19 | 113.8 (3) | O4—Mn1—O10 | 92.25 (11) |
| C15—C14—C19 | 121.7 (4) | C8—N1—C9 | 119.5 (3) |
| C14—C15—C16 | 120.6 (4) | C8—N1—Mn1 | 125.1 (2) |
| C14—C15—H15A | 119.7 | C9—N1—Mn1 | 115.3 (2) |
| C16—C15—H15A | 119.7 | C20—N2—C21 | 120.5 (3) |
| C17—C16—C15 | 119.5 (4) | C20—N2—Mn1 | 123.1 (2) |
| C17—C16—H16A | 120.2 | C21—N2—Mn1 | 116.1 (2) |
| C15—C16—H16A | 120.2 | C2—O1—C1 | 119.0 (3) |
| C16—C17—C18 | 121.9 (4) | C7—O2—Mn1 | 125.5 (2) |
| C16—C17—H17A | 119.0 | C25—O3—H3 | 110 (3) |
| C18—C17—H17A | 119.0 | C10—O4—Mn1 | 111.1 (2) |
| C19—C18—C17 | 119.3 (3) | C10—O4—H4 | 107 (3) |
| C19—C18—C20 | 124.3 (3) | Mn1—O4—H4 | 123 (3) |
| C17—C18—C20 | 116.4 (3) | C11—O5—H5 | 112 (4) |
| O8—C19—C18 | 124.5 (3) | C12—O6—H6 | 109 (4) |
| O8—C19—C14 | 118.6 (3) | C14—O7—C13 | 118.9 (4) |
| C18—C19—C14 | 116.9 (3) | C19—O8—Mn1 | 124.0 (2) |
| N2—C20—C18 | 127.4 (3) | C26—O9—H9 | 121 (7) |
| N2—C20—H20A | 116.3 | C22—O10—Mn1 | 111.5 (2) |

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| C18—C20—H20A | 116.3 | C22—O10—H10 | 133 (4) |
| N2—C21—C24 | 117.6 (8) | Mn1—O10—H10 | 115 (4) |
| N2—C21—C23 | 115.8 (3) | C23—O11—H11 | 109 (5) |
| C24—C21—C23 | 100.8 (7) | H1W—O1W—H2W | 135.0 |
| O1—C2—C3—C4 | 178.2 (4) | C12—C9—N1—Mn1 | 87.7 (3) |
| C7—C2—C3—C4 | 0.1 (6) | C11—C9—N1—Mn1 | -150.8 (3) |
| C2—C3—C4—C5 | 0.4 (6) | C10—C9—N1—Mn1 | -30.2 (3) |
| C3—C4—C5—C6 | -1.1 (6) | O8—Mn1—N1—C8 | 105.0 (3) |
| C4—C5—C6—C7 | 1.3 (6) | O2—Mn1—N1—C8 | 12.0 (3) |
| C4—C5—C6—C8 | -179.4 (3) | N2—Mn1—N1—C8 | -119.0 (5) |
| C5—C6—C7—O2 | 179.1 (3) | O4—Mn1—N1—C8 | -170.3 (3) |
| C8—C6—C7—O2 | -0.1 (6) | O10—Mn1—N1—C8 | -78.0 (3) |
| C5—C6—C7—C2 | -0.8 (5) | O8—Mn1—N1—C9 | -77.4 (2) |
| C8—C6—C7—C2 | 180.0 (3) | O2—Mn1—N1—C9 | -170.4 (2) |
| O1—C2—C7—O2 | 1.9 (5) | N2—Mn1—N1—C9 | 58.6 (5) |
| C3—C2—C7—O2 | -179.8 (3) | O4—Mn1—N1—C9 | 7.4 (2) |
| O1—C2—C7—C6 | -178.2 (3) | O10—Mn1—N1—C9 | 99.6 (2) |
| C3—C2—C7—C6 | 0.1 (5) | C18—C20—N2—C21 | -178.5 (3) |
| C7—C6—C8—N1 | -4.2 (6) | C18—C20—N2—Mn1 | -5.1 (5) |
| C5—C6—C8—N1 | 176.6 (3) | C24—C21—N2—C20 | -98.6 (8) |
| N1—C9—C10—O4 | 45.2 (4) | C23—C21—N2—C20 | 20.6 (5) |
| C12—C9—C10—O4 | -71.1 (4) | C24A—C21—N2—C20 | -99.6 (8) |
| C11—C9—C10—O4 | 170.7 (3) | C22—C21—N2—C20 | 140.6 (3) |
| N1—C9—C11—O5 | 56.9 (4) | C24—C21—N2—Mn1 | 87.6 (8) |
| C12—C9—C11—O5 | 177.6 (3) | C23—C21—N2—Mn1 | -153.2 (3) |
| C10—C9—C11—O5 | -63.4 (4) | C24A—C21—N2—Mn1 | 86.6 (7) |
| N1—C9—C12—O6 | 57.4 (4) | C22—C21—N2—Mn1 | -33.2 (4) |
| C11—C9—C12—O6 | -68.5 (4) | O8—Mn1—N2—C20 | 16.8 (3) |
| C10—C9—C12—O6 | 173.8 (3) | O2—Mn1—N2—C20 | 109.7 (3) |
| O7—C14—C15—C16 | 178.2 (5) | N1—Mn1—N2—C20 | -119.9 (4) |
| C19—C14—C15—C16 | -2.1 (7) | O4—Mn1—N2—C20 | -69.7 (3) |
| C14—C15—C16—C17 | -0.7 (8) | O10—Mn1—N2—C20 | -161.8 (3) |
| C15—C16—C17—C18 | 1.4 (7) | O8—Mn1—N2—C21 | -169.6 (3) |
| C16—C17—C18—C19 | 0.6 (6) | O2—Mn1—N2—C21 | -76.7 (3) |
| C16—C17—C18—C20 | -179.8 (4) | N1—Mn1—N2—C21 | 53.8 (6) |
| C17—C18—C19—O8 | 175.9 (3) | O4—Mn1—N2—C21 | 103.9 (3) |
| C20—C18—C19—O8 | -3.6 (5) | O10—Mn1—N2—C21 | 11.9 (3) |
| C17—C18—C19—C14 | -3.2 (5) | C3—C2—O1—C1 | 5.7 (6) |
| C20—C18—C19—C14 | 177.3 (3) | C7—C2—O1—C1 | -176.1 (4) |
| O7—C14—C19—O8 | 4.6 (5) | C6—C7—O2—Mn1 | 12.7 (5) |
| C15—C14—C19—O8 | -175.2 (4) | C2—C7—O2—Mn1 | -167.4 (2) |
| O7—C14—C19—C18 | -176.3 (3) | O8—Mn1—O2—C7 | -114.8 (3) |
| C15—C14—C19—C18 | 4.0 (6) | N1—Mn1—O2—C7 | -15.9 (3) |
| C19—C18—C20—N2 | -6.2 (6) | N2—Mn1—O2—C7 | 153.6 (3) |
| C17—C18—C20—N2 | 174.2 (3) | O10—Mn1—O2—C7 | 74.5 (3) |
| N2—C21—C22—O10 | 44.7 (4) | C9—C10—O4—Mn1 | -40.2 (3) |
| C24—C21—C22—O10 | -82.1 (8) | O8—Mn1—O4—C10 | 118.6 (2) |

| | | | |
|-------------------|------------|-----------------|------------|
| C23—C21—C22—O10 | 169.4 (3) | N1—Mn1—O4—C10 | 19.0 (2) |
| C24A—C21—C22—O10 | -66.9 (10) | N2—Mn1—O4—C10 | -150.2 (2) |
| N2—C21—C23—O11 | 57.4 (5) | O10—Mn1—O4—C10 | -71.0 (2) |
| C24—C21—C23—O11 | -174.6 (8) | C15—C14—O7—C13 | -6.8 (7) |
| C24A—C21—C23—O11 | 173.0 (7) | C19—C14—O7—C13 | 173.5 (4) |
| C22—C21—C23—O11 | -61.1 (4) | C18—C19—O8—Mn1 | 23.1 (5) |
| N2—C21—C24—O12 | -71.5 (16) | C14—C19—O8—Mn1 | -157.7 (3) |
| C23—C21—C24—O12 | 161.8 (13) | O2—Mn1—O8—C19 | -123.4 (3) |
| C22—C21—C24—O12 | 48.1 (16) | N1—Mn1—O8—C19 | 144.7 (3) |
| N2—C21—C24A—O12A | 64 (2) | N2—Mn1—O8—C19 | -25.6 (3) |
| C23—C21—C24A—O12A | -60 (2) | O4—Mn1—O8—C19 | 65.7 (3) |
| C22—C21—C24A—O12A | 177.2 (15) | C21—C22—O10—Mn1 | -36.8 (4) |
| C6—C8—N1—C9 | 177.6 (3) | O2—Mn1—O10—C22 | 112.2 (3) |
| C6—C8—N1—Mn1 | -4.9 (5) | N1—Mn1—O10—C22 | -156.3 (3) |
| C12—C9—N1—C8 | -94.5 (4) | N2—Mn1—O10—C22 | 14.4 (3) |
| C11—C9—N1—C8 | 26.9 (5) | O4—Mn1—O10—C22 | -76.6 (3) |
| C10—C9—N1—C8 | 147.5 (3) | | |

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

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|--|-------------|---------------|-----------------------|-------------------------|
| O3—H3...O8 | 0.86 (1) | 1.89 (1) | 2.706 (5) | 160 (4) |
| O4—H4...O9 | 0.86 (1) | 1.80 (1) | 2.664 (5) | 174 (5) |
| O5—H5...O2 ⁱ | 0.85 (1) | 1.89 (1) | 2.736 (4) | 173 (6) |
| O6—H6...O3 | 0.86 (1) | 1.89 (1) | 2.737 (7) | 172 (6) |
| O9—H9...O11 | 0.85 (1) | 1.89 (1) | 2.712 (5) | 162 (4) |
| O10—H10...O1 ^W | 0.86 (1) | 1.90 (4) | 2.585 (6) | 136 (5) |
| O11—H11...O6 ⁱⁱ | 0.85 (1) | 1.90 (2) | 2.739 (4) | 169 (7) |
| O12—H12...O2 | 0.85 | 2.12 | 2.967 (7) | 180 |
| O12A—H12A...O5 ⁱⁱⁱ | 0.85 | 2.10 | 2.946 (7) | 180 |
| O1 ^W —H1 ^W ...O5 | 0.85 | 1.81 | 2.657 (8) | 180 |
| O1 ^W —H2 ^W ...O12A ^{iv} | 0.85 | 2.14 | 2.990 (9) | 180 |

Symmetry codes: (i) $x-1, y, z$; (ii) $x+1/2, -y+3/2, z+1/2$; (iii) $x+1, y, z$; (iv) $-x+1, -y+1, -z+1$.