

Pentaqua[5,5'-(*m*-phenylene)ditetra-zolato- κN^2]manganese(II) dihydrate

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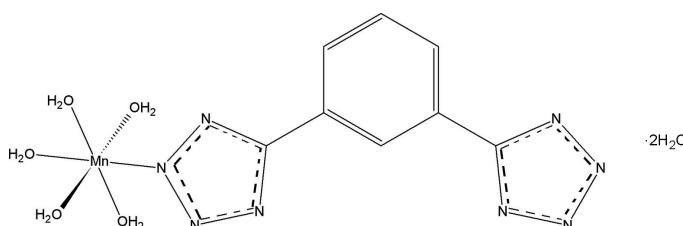
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(C-C) = 0.003$ Å; R factor = 0.038; wR factor = 0.095; data-to-parameter ratio = 13.6.

The title compound, $[Mn(C_8H_4N_8)_2(H_2O)_5] \cdot 2H_2O$, is the fourth transition metal complex containing the 5,5'-(*m*-phenylene)ditetrazolate ligand to be structurally characterized. The Mn^{II} cation has a distorted octahedral coordination geometry. The 5,5'-(*m*-phenylene)ditetrazolate ligand is planar. All H atoms bonded to O atoms participate in hydrogen bonds, which link the molecules into a framework structure.

Related literature

For similar complexes, see: Jiang *et al.* (2004); Hill *et al.* (1996).



Experimental

Crystal data

$[Mn(C_8H_4N_8)_2(H_2O)_5] \cdot 2H_2O$
 $M_r = 393.24$

Triclinic, $P\bar{1}$
 $a = 6.5932$ (1) Å
 $b = 10.0711$ (2) Å
 $c = 12.9857$ (3) Å
 $\alpha = 68.296$ (1) $^\circ$
 $\beta = 77.213$ (3) $^\circ$

$\gamma = 77.280$ (5) $^\circ$
 $V = 772.10$ (3) Å³
 $Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.91$ mm⁻¹
 $T = 296$ (2) K
 $0.26 \times 0.14 \times 0.08$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.798$, $T_{\max} = 0.931$

7710 measured reflections
3704 independent reflections
2846 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.095$
 $S = 1.02$
3704 reflections
273 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.36$ e Å⁻³
 $\Delta\rho_{\min} = -0.30$ e Å⁻³

Table 1
Hydrogen-bond geometry (Å, °).

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|-----------------------------|----------|--------------|--------------|----------------|
| O7—H7B···O6 | 0.57 (6) | 2.31 (6) | 2.852 (3) | 162 (8) |
| O5—H5A···O5 ⁱ | 0.57 (6) | 2.41 (5) | 2.910 (6) | 148 (9) |
| O6—H6B···O7 ⁱⁱ | 0.61 (5) | 2.21 (5) | 2.814 (3) | 171 (6) |
| O4—H4A···O1 ⁱⁱⁱ | 0.67 (4) | 2.38 (4) | 3.035 (3) | 167 (5) |
| O3—H3A···O7 ⁱ | 0.84 (4) | 1.91 (4) | 2.747 (3) | 171 (3) |
| O3—H3B···O6 ^{iv} | 0.82 (3) | 1.98 (3) | 2.794 (3) | 176 (3) |
| O2—H2B···N1 ^v | 0.75 (3) | 2.06 (3) | 2.800 (3) | 173 (3) |
| O1—H1B···N6 ^{vi} | 0.85 (4) | 1.89 (4) | 2.730 (3) | 176 (3) |
| O5—H5B···N8 ^{vii} | 0.75 (4) | 2.07 (4) | 2.810 (3) | 168 (4) |
| O7—H7A···N5 ⁱⁱ | 0.73 (4) | 2.10 (4) | 2.828 (3) | 173 (4) |
| O4—H4B···N3 ^{iv} | 0.88 (4) | 1.80 (4) | 2.681 (3) | 175 (3) |
| O6—H6A···N4 | 0.82 (3) | 2.07 (4) | 2.886 (3) | 176 (3) |
| O1—H1A···N7 ^{viii} | 0.78 (3) | 1.99 (3) | 2.771 (3) | 175 (3) |

Symmetry codes: (i) $-x + 1, -y + 1, -z + 1$; (ii) $-x + 1, -y + 1, -z$; (iii) $x - 1, y, z$; (iv) $-x, -y + 1, -z + 1$; (v) $-x, -y + 2, -z + 1$; (vi) $x, y, z + 1$; (vii) $x, y - 1, z + 1$; (viii) $-x + 1, -y + 2, -z$.

Data collection: *SMART* (Bruker, 2007); cell refinement: *SMART*; data reduction: *SAINT-Plus* (Bruker, 2007); 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: EZ2139).

References

- Bruker (2007). *SMART* and *SAINT-Plus*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Hill, M., Mahon, M. F., McGinley, J. & Molloy, K. C. (1996). *J. Chem. Soc. Dalton Trans.*, pp. 835–845.
- Jiang, C., Yu, Z., Jiao, C., Wang, S., Li, J., Wang, Z. & Cui, Y. (2004). *Eur. J. Inorg. Chem.*, pp. 4669–4674.
- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.

supporting information

Acta Cryst. (2008). E64, m1255 [doi:10.1107/S1600536808028316]

Pentaqua[5,5'-(*m*-phenylene)ditetrazolato- κN^2]manganese(II) dihydrate

Yuanqi Lü

S1. Comment

The 1,3-di(2*H*-tetrazol-5-yl)benzene (DHTB) ligand has hitherto been reported in the twice deprotonated form in the crystal structures of its complexes with zinc, cadmium and tin (Jiang *et al.*, 2004; Hill *et al.*, 1996), where it acts as a bridging ligand. This paper provides the first structural characterization of a DHTB complex with the transition metal Mn(II); the ligand in this complex is also twice deprotonated, but coordinated to the Mn atom as a terminal ligand.

The molecule of Mn(DHTB)(H₂O)₅ occupies a general position in the unit cell; the Mn atom has a non-distorted octahedral coordination as indicated by bond lengths and angles (Fig. 1). The DHTB ligand has an essentially planar conformation, with the maximum deviation from the mean plane being 0.054 (2) Å by atom C7. The geometry of the ligand is similar to that observed in Jiang *et al.* (2004) and Hill *et al.* (1996).

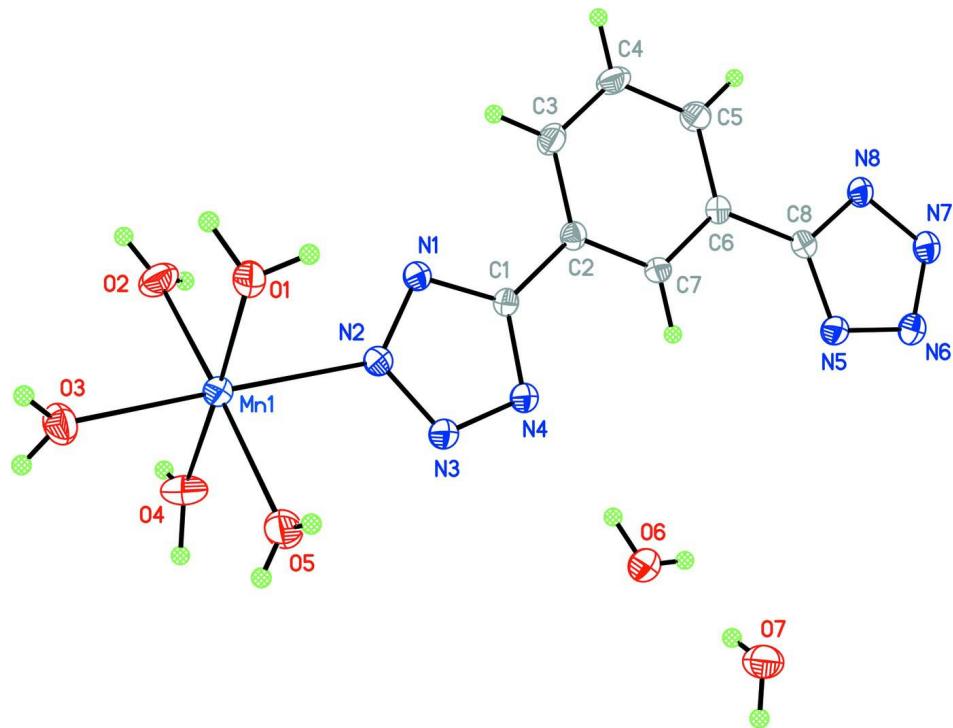
Strong π – π interactions between the aromatic rings are indicated by the short distance of 3.324 (3) Å between C1 and C8ⁱ [Symmetry code: (i) 1-x, 2-y, -z]. All hydrogen atoms that are bonded to oxygen atoms participate in H-bonding (Table 1); the extensive H-bond system and the strong π – π interactions link molecules of the complex and non-coordinated water molecules into a three-dimensional infinite network (Fig. 2).

S2. Experimental

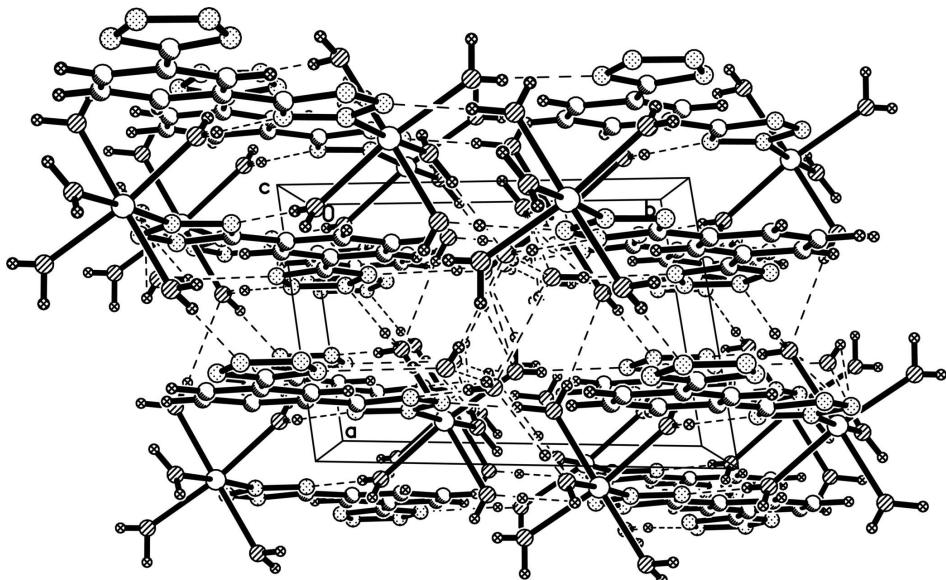
The hydrothermal reaction of Mn(NO₃)₂ (0.5 mmol) and 1,3-di(2*H*-tetrazol-5-yl)benzene (0.5 mmol) in 20 ml of distilled water at 180°C for 3 days resulted in light yellow plate crystals of the title compound, in a yield of 42%. The crystals were filtered, washed with cold EtOH and dried in air.

S3. Refinement

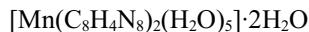
All of the H atoms on carbon atoms were positioned geometrically and refined using a riding model with C—H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2$ times $U_{\text{eq}}(\text{C})$. All of the H atoms on oxygen atoms were located from the difference Fourier map, and refined freely, except for the bond length of O5—H5A being constrained to 0.87 Å.

**Figure 1**

Molecular structure of (I) showing 50% probability displacement ellipsoids and the atom-labelling scheme.

**Figure 2**

Packing diagram viewed down the c axis,

Pentaqua[5,5'-(m-phenylene)ditetrazolato- κN^2]manganese(II) dihydrate*Crystal data*

$M_r = 393.24$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.5932 (1)$ Å

$b = 10.0711 (2)$ Å

$c = 12.9857 (3)$ Å

$\alpha = 68.296 (1)^\circ$

$\beta = 77.213 (3)^\circ$

$\gamma = 77.280 (5)^\circ$

$V = 772.10 (3)$ Å³

$Z = 2$

$F(000) = 406$

$D_x = 1.691$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 3628 reflections

$\theta = 2.7\text{--}27.9^\circ$

$\mu = 0.91$ mm⁻¹

$T = 296$ K

Plate, yellow

0.26 × 0.14 × 0.08 mm

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)

$T_{\min} = 0.798$, $T_{\max} = 0.931$

7710 measured reflections

3704 independent reflections

2846 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.022$

$\theta_{\max} = 28.1^\circ$, $\theta_{\min} = 1.7^\circ$

$h = -8\text{--}6$

$k = -13\text{--}13$

$l = -17\text{--}17$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.038$

$wR(F^2) = 0.095$

$S = 1.02$

3704 reflections

273 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

H atoms treated by a mixture of independent
and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0414P)^2 + 0.3671P]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.36$ e Å⁻³

$\Delta\rho_{\min} = -0.30$ e Å⁻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 (Å²)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|-------------|-------------|--------------|----------------------------------|
| Mn1 | 0.11390 (6) | 0.68638 (4) | 0.58413 (3) | 0.02732 (12) |
| O1 | 0.3923 (3) | 0.7734 (2) | 0.57901 (16) | 0.0388 (4) |
| O2 | -0.1155 (4) | 0.8808 (2) | 0.58408 (19) | 0.0402 (5) |

| | | | | |
|-----|-------------|--------------|---------------|-------------|
| O3 | 0.0794 (3) | 0.6043 (2) | 0.76359 (15) | 0.0361 (4) |
| O4 | -0.1679 (4) | 0.6027 (2) | 0.59456 (18) | 0.0411 (5) |
| O5 | 0.3064 (5) | 0.4784 (2) | 0.5795 (2) | 0.0508 (6) |
| O6 | 0.2643 (3) | 0.5392 (2) | 0.12086 (18) | 0.0370 (4) |
| O7 | 0.6605 (4) | 0.3559 (3) | 0.11514 (17) | 0.0367 (4) |
| N1 | 0.1613 (3) | 0.90491 (19) | 0.31988 (14) | 0.0244 (4) |
| N2 | 0.1633 (3) | 0.76902 (19) | 0.39242 (14) | 0.0264 (4) |
| N3 | 0.1980 (3) | 0.6792 (2) | 0.33621 (15) | 0.0300 (4) |
| N4 | 0.2200 (3) | 0.75287 (19) | 0.22651 (15) | 0.0268 (4) |
| N5 | 0.3286 (3) | 0.9471 (2) | -0.21040 (15) | 0.0276 (4) |
| N6 | 0.3651 (3) | 0.9732 (2) | -0.32077 (16) | 0.0315 (4) |
| N7 | 0.3610 (3) | 1.1128 (2) | -0.37435 (16) | 0.0318 (4) |
| N8 | 0.3223 (3) | 1.1812 (2) | -0.29911 (15) | 0.0275 (4) |
| C1 | 0.1974 (3) | 0.8914 (2) | 0.21865 (17) | 0.0205 (4) |
| C2 | 0.2128 (3) | 1.0130 (2) | 0.11148 (17) | 0.0216 (4) |
| C3 | 0.2001 (4) | 1.1526 (2) | 0.11091 (19) | 0.0306 (5) |
| H3 | 0.1796 | 1.1699 | 0.1783 | 0.037* |
| C4 | 0.2177 (4) | 1.2662 (2) | 0.0105 (2) | 0.0380 (6) |
| H4 | 0.2077 | 1.3597 | 0.0105 | 0.046* |
| C5 | 0.2501 (4) | 1.2411 (2) | -0.09028 (19) | 0.0311 (5) |
| H5 | 0.2627 | 1.3178 | -0.1576 | 0.037* |
| C6 | 0.2640 (3) | 1.1024 (2) | -0.09143 (17) | 0.0217 (4) |
| C7 | 0.2443 (3) | 0.9883 (2) | 0.00989 (17) | 0.0214 (4) |
| H7 | 0.2523 | 0.8950 | 0.0098 | 0.026* |
| C8 | 0.3031 (3) | 1.0766 (2) | -0.19900 (17) | 0.0224 (4) |
| H1A | 0.455 (5) | 0.808 (3) | 0.519 (3) | 0.044 (9)* |
| H6A | 0.254 (5) | 0.602 (4) | 0.148 (3) | 0.053 (9)* |
| H4B | -0.180 (5) | 0.511 (4) | 0.613 (3) | 0.063 (10)* |
| H7A | 0.659 (5) | 0.278 (4) | 0.135 (3) | 0.060 (12)* |
| H5B | 0.306 (6) | 0.402 (4) | 0.620 (3) | 0.068 (12)* |
| H1B | 0.378 (5) | 0.838 (4) | 0.608 (3) | 0.068 (11)* |
| H2B | -0.124 (5) | 0.933 (4) | 0.614 (3) | 0.052 (10)* |
| H3B | -0.018 (5) | 0.560 (3) | 0.800 (3) | 0.059 (10)* |
| H3A | 0.149 (5) | 0.615 (4) | 0.806 (3) | 0.065 (11)* |
| H4A | -0.261 (6) | 0.645 (4) | 0.597 (3) | 0.076 (16)* |
| H2A | -0.179 (7) | 0.908 (5) | 0.539 (4) | 0.105 (18)* |
| H6B | 0.286 (8) | 0.554 (5) | 0.070 (4) | 0.09 (2)* |
| H7B | 0.588 (10) | 0.394 (6) | 0.126 (5) | 0.11 (3)* |
| H5A | 0.396 (10) | 0.477 (7) | 0.567 (5) | 0.10 (3)* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|--------------|--------------|---------------|---------------|---------------|
| Mn1 | 0.0377 (2) | 0.02288 (18) | 0.02168 (18) | -0.00688 (14) | -0.00202 (14) | -0.00808 (13) |
| O1 | 0.0537 (12) | 0.0406 (10) | 0.0275 (9) | -0.0234 (9) | 0.0075 (8) | -0.0164 (8) |
| O2 | 0.0540 (13) | 0.0266 (9) | 0.0423 (11) | 0.0024 (8) | -0.0102 (10) | -0.0174 (9) |
| O3 | 0.0426 (11) | 0.0429 (11) | 0.0241 (9) | -0.0157 (9) | -0.0046 (8) | -0.0078 (8) |
| O4 | 0.0460 (13) | 0.0253 (10) | 0.0564 (13) | -0.0068 (9) | -0.0170 (10) | -0.0128 (9) |

| | | | | | | |
|----|-------------|-------------|-------------|--------------|--------------|-------------|
| O5 | 0.0666 (16) | 0.0254 (11) | 0.0400 (12) | 0.0045 (10) | 0.0116 (11) | -0.0056 (9) |
| O6 | 0.0484 (11) | 0.0342 (10) | 0.0310 (10) | -0.0101 (8) | -0.0005 (9) | -0.0150 (8) |
| O7 | 0.0479 (12) | 0.0276 (10) | 0.0350 (10) | -0.0079 (9) | -0.0089 (8) | -0.0082 (8) |
| N1 | 0.0311 (10) | 0.0217 (9) | 0.0191 (8) | -0.0043 (7) | -0.0027 (7) | -0.0059 (7) |
| N2 | 0.0347 (10) | 0.0215 (9) | 0.0209 (9) | -0.0044 (8) | -0.0021 (8) | -0.0062 (7) |
| N3 | 0.0428 (12) | 0.0239 (9) | 0.0223 (9) | -0.0070 (8) | -0.0005 (8) | -0.0083 (7) |
| N4 | 0.0356 (11) | 0.0240 (9) | 0.0203 (9) | -0.0063 (8) | -0.0011 (8) | -0.0077 (7) |
| N5 | 0.0333 (10) | 0.0258 (9) | 0.0236 (9) | -0.0052 (8) | -0.0022 (8) | -0.0092 (8) |
| N6 | 0.0375 (11) | 0.0343 (11) | 0.0247 (10) | -0.0074 (9) | -0.0005 (8) | -0.0135 (8) |
| N7 | 0.0380 (11) | 0.0333 (11) | 0.0219 (9) | -0.0072 (9) | 0.0009 (8) | -0.0088 (8) |
| N8 | 0.0338 (10) | 0.0269 (10) | 0.0193 (9) | -0.0048 (8) | 0.0001 (8) | -0.0072 (7) |
| C1 | 0.0197 (10) | 0.0214 (10) | 0.0202 (10) | -0.0028 (8) | -0.0008 (8) | -0.0083 (8) |
| C2 | 0.0207 (10) | 0.0219 (10) | 0.0203 (10) | -0.0010 (8) | -0.0028 (8) | -0.0066 (8) |
| C3 | 0.0427 (14) | 0.0259 (11) | 0.0222 (11) | -0.0016 (10) | -0.0027 (10) | -0.0105 (9) |
| C4 | 0.0638 (18) | 0.0197 (11) | 0.0301 (12) | -0.0058 (11) | -0.0040 (12) | -0.0101 (9) |
| C5 | 0.0448 (14) | 0.0199 (10) | 0.0230 (11) | -0.0015 (10) | -0.0051 (10) | -0.0028 (9) |
| C6 | 0.0206 (10) | 0.0238 (10) | 0.0203 (10) | -0.0029 (8) | -0.0022 (8) | -0.0076 (8) |
| C7 | 0.0223 (10) | 0.0186 (10) | 0.0228 (10) | -0.0027 (8) | -0.0022 (8) | -0.0073 (8) |
| C8 | 0.0206 (10) | 0.0235 (10) | 0.0213 (10) | -0.0039 (8) | -0.0024 (8) | -0.0058 (8) |

Geometric parameters (\AA , $^\circ$)

| | | | |
|-----------|-------------|-----------|-------------|
| Mn1—O3 | 2.1423 (18) | N1—N2 | 1.343 (2) |
| Mn1—O4 | 2.162 (2) | N2—N3 | 1.314 (2) |
| Mn1—O1 | 2.1797 (19) | N3—N4 | 1.332 (2) |
| Mn1—O2 | 2.1946 (19) | N4—C1 | 1.338 (3) |
| Mn1—O5 | 2.212 (2) | N5—N6 | 1.333 (3) |
| Mn1—N2 | 2.2857 (17) | N5—C8 | 1.336 (3) |
| O1—H1A | 0.78 (3) | N6—N7 | 1.315 (3) |
| O1—H1B | 0.85 (4) | N7—N8 | 1.343 (3) |
| O2—H2B | 0.75 (3) | N8—C8 | 1.338 (3) |
| O2—H2A | 0.73 (5) | C1—C2 | 1.476 (3) |
| O3—H3B | 0.82 (3) | C2—C3 | 1.387 (3) |
| O3—H3A | 0.84 (4) | C2—C7 | 1.394 (3) |
| O4—H4B | 0.88 (4) | C3—C4 | 1.382 (3) |
| O4—H4A | 0.67 (4) | C3—H3 | 0.9300 |
| O5—H5B | 0.75 (4) | C4—C5 | 1.385 (3) |
| O5—H5A | 0.57 (6) | C4—H4 | 0.9300 |
| O6—H6A | 0.82 (3) | C5—C6 | 1.385 (3) |
| O6—H6B | 0.61 (5) | C5—H5 | 0.9300 |
| O7—H7A | 0.73 (4) | C6—C7 | 1.393 (3) |
| O7—H7B | 0.57 (6) | C6—C8 | 1.472 (3) |
| N1—C1 | 1.335 (3) | C7—H7 | 0.9300 |
| | | | |
| O3—Mn1—O4 | 88.99 (8) | N3—N2—N1 | 109.26 (16) |
| O3—Mn1—O1 | 89.37 (8) | N3—N2—Mn1 | 120.96 (13) |
| O4—Mn1—O1 | 177.83 (8) | N1—N2—Mn1 | 129.79 (13) |
| O3—Mn1—O2 | 92.33 (8) | N2—N3—N4 | 109.71 (17) |

| | | | |
|--------------|--------------|-------------|--------------|
| O4—Mn1—O2 | 81.61 (9) | N3—N4—C1 | 104.97 (17) |
| O1—Mn1—O2 | 97.05 (8) | N6—N5—C8 | 105.15 (17) |
| O3—Mn1—O5 | 89.42 (8) | N7—N6—N5 | 109.76 (17) |
| O4—Mn1—O5 | 90.03 (11) | N6—N7—N8 | 109.04 (17) |
| O1—Mn1—O5 | 91.36 (10) | C8—N8—N7 | 105.04 (18) |
| O2—Mn1—O5 | 171.42 (11) | N1—C1—N4 | 111.28 (17) |
| O3—Mn1—N2 | 177.82 (7) | N1—C1—C2 | 124.65 (18) |
| O4—Mn1—N2 | 92.24 (8) | N4—C1—C2 | 124.07 (18) |
| O1—Mn1—N2 | 89.44 (7) | C3—C2—C7 | 119.41 (19) |
| O2—Mn1—N2 | 89.63 (8) | C3—C2—C1 | 120.24 (19) |
| O5—Mn1—N2 | 88.78 (8) | C7—C2—C1 | 120.34 (18) |
| Mn1—O1—H1A | 116 (2) | C4—C3—C2 | 120.2 (2) |
| Mn1—O1—H1B | 118 (2) | C4—C3—H3 | 119.9 |
| H1A—O1—H1B | 102 (3) | C2—C3—H3 | 119.9 |
| Mn1—O2—H2B | 131 (3) | C3—C4—C5 | 120.2 (2) |
| Mn1—O2—H2A | 115 (4) | C3—C4—H4 | 119.9 |
| H2B—O2—H2A | 113 (4) | C5—C4—H4 | 119.9 |
| Mn1—O3—H3B | 120 (2) | C6—C5—C4 | 120.4 (2) |
| Mn1—O3—H3A | 129 (2) | C6—C5—H5 | 119.8 |
| H3B—O3—H3A | 110 (3) | C4—C5—H5 | 119.8 |
| Mn1—O4—H4B | 127 (2) | C5—C6—C7 | 119.31 (19) |
| Mn1—O4—H4A | 119 (4) | C5—C6—C8 | 119.95 (19) |
| H4B—O4—H4A | 113 (4) | C7—C6—C8 | 120.73 (18) |
| Mn1—O5—H5B | 132 (3) | C6—C7—C2 | 120.44 (19) |
| Mn1—O5—H5A | 117 (7) | C6—C7—H7 | 119.8 |
| H5B—O5—H5A | 99 (7) | C2—C7—H7 | 119.8 |
| H6A—O6—H6B | 119 (5) | N5—C8—N8 | 111.01 (18) |
| H7A—O7—H7B | 120 (6) | N5—C8—C6 | 125.20 (18) |
| C1—N1—N2 | 104.79 (16) | N8—C8—C6 | 123.78 (19) |
| | | | |
| C1—N1—N2—N3 | -0.3 (2) | N4—C1—C2—C3 | -176.0 (2) |
| C1—N1—N2—Mn1 | -179.63 (15) | N1—C1—C2—C7 | -177.9 (2) |
| O4—Mn1—N2—N3 | -62.82 (18) | N4—C1—C2—C7 | 2.7 (3) |
| O1—Mn1—N2—N3 | 118.54 (17) | C7—C2—C3—C4 | 0.2 (4) |
| O2—Mn1—N2—N3 | -144.41 (18) | C1—C2—C3—C4 | 179.0 (2) |
| O5—Mn1—N2—N3 | 27.16 (18) | C2—C3—C4—C5 | -0.6 (4) |
| O3—Mn1—N2—N1 | -119.2 (19) | C3—C4—C5—C6 | 0.4 (4) |
| O4—Mn1—N2—N1 | 116.41 (19) | C4—C5—C6—C7 | 0.2 (4) |
| O1—Mn1—N2—N1 | -62.23 (19) | C4—C5—C6—C8 | -178.7 (2) |
| O2—Mn1—N2—N1 | 34.82 (19) | C5—C6—C7—C2 | -0.5 (3) |
| O5—Mn1—N2—N1 | -153.6 (2) | C8—C6—C7—C2 | 178.31 (19) |
| N1—N2—N3—N4 | 0.2 (2) | C3—C2—C7—C6 | 0.3 (3) |
| Mn1—N2—N3—N4 | 179.52 (14) | C1—C2—C7—C6 | -178.46 (19) |
| N2—N3—N4—C1 | 0.1 (2) | N6—N5—C8—N8 | 0.2 (2) |
| C8—N5—N6—N7 | -0.3 (2) | N6—N5—C8—C6 | -178.4 (2) |
| N5—N6—N7—N8 | 0.4 (3) | N7—N8—C8—N5 | 0.0 (2) |
| N6—N7—N8—C8 | -0.2 (2) | N7—N8—C8—C6 | 178.67 (19) |
| N2—N1—C1—N4 | 0.4 (2) | C5—C6—C8—N5 | 176.9 (2) |

| | | | |
|-------------|--------------|-------------|-----------|
| N2—N1—C1—C2 | −179.00 (19) | C7—C6—C8—N5 | −1.9 (3) |
| N3—N4—C1—N1 | −0.3 (2) | C5—C6—C8—N8 | −1.6 (3) |
| N3—N4—C1—C2 | 179.09 (19) | C7—C6—C8—N8 | 179.6 (2) |
| N1—C1—C2—C3 | 3.3 (3) | | |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|-----------------------------|----------|----------|-----------|---------|
| O7—H7B···O6 | 0.57 (6) | 2.31 (6) | 2.852 (3) | 162 (8) |
| O5—H5A···O5 ⁱ | 0.57 (6) | 2.41 (5) | 2.910 (6) | 148 (9) |
| O6—H6B···O7 ⁱⁱ | 0.61 (5) | 2.21 (5) | 2.814 (3) | 171 (6) |
| O4—H4A···O1 ⁱⁱⁱ | 0.67 (4) | 2.38 (4) | 3.035 (3) | 167 (5) |
| O3—H3A···O7 ⁱ | 0.84 (4) | 1.91 (4) | 2.747 (3) | 171 (3) |
| O3—H3B···O6 ^{iv} | 0.82 (3) | 1.98 (3) | 2.794 (3) | 176 (3) |
| O2—H2B···N1 ^v | 0.75 (3) | 2.06 (3) | 2.800 (3) | 173 (3) |
| O1—H1B···N6 ^{vi} | 0.85 (4) | 1.89 (4) | 2.730 (3) | 176 (3) |
| O5—H5B···N8 ^{vii} | 0.75 (4) | 2.07 (4) | 2.810 (3) | 168 (4) |
| O7—H7A···N5 ⁱⁱ | 0.73 (4) | 2.10 (4) | 2.828 (3) | 173 (4) |
| O4—H4B···N3 ^{iv} | 0.88 (4) | 1.80 (4) | 2.681 (3) | 175 (3) |
| O6—H6A···N4 | 0.82 (3) | 2.07 (4) | 2.886 (3) | 176 (3) |
| O1—H1A···N7 ^{viii} | 0.78 (3) | 1.99 (3) | 2.771 (3) | 175 (3) |

Symmetry codes: (i) $-x+1, -y+1, -z+1$; (ii) $-x+1, -y+1, -z$; (iii) $x-1, y, z$; (iv) $-x, -y+1, -z+1$; (v) $-x, -y+2, -z+1$; (vi) $x, y, z+1$; (vii) $x, y-1, z+1$; (viii) $-x+1, -y+2, -z$.