Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

Poly[(μ_2-azido-κ^2N^1:N^1)[μ_2-5-(8-quinolyloxymethyl)tetrazolato-κ^4N^1,O,N^5:N^4]manganese(II)]

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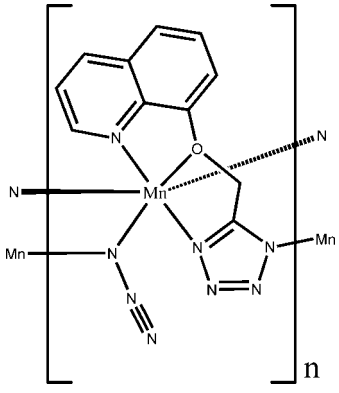
Received 1 July 2008; accepted 18 July 2008

Key indicators: single-crystal X-ray study; T = 293 K; mean σ(C–C) = 0.004 Å; R factor = 0.047; wR factor = 0.106; data-to-parameter ratio = 16.2.

In the structure of the title compound, $[Mn(C_{11}H_8N_5O)(N_3)]_n$, the Mn atoms are hexacoordinated by five N atoms and one O atom. The coordination polyhedron of the Mn atom is a slightly distorted octahedron. The Mn atoms are connected by azide anions with a μ_2-1,1 mode and by 5-(8-quinolyloxymethyl)tetrazolate ligands in a μ_2-$\eta^1(N)$,η^3-(N,N,O) fashion to form a two-dimensional framework parallel to the (100) plane. Geometric parameters of the organic ligand are in the normal ranges and the dihedral angle between the quinoline ring system and the tetrazole unit is 7.41 (15)°. The structure involves intra- and intermolecular C—H···N hydrogen bonds.

Related literature

For the use of tetrazole derivatives in coordination chemistry, see: Wang *et al.* (2005); Xiong *et al.* (2002). For the crystal structure of a tetrazole derivative, see: Wang & Ye (2007); For the synthesis of 8-cyanatoquinoline, see: Luo & Ye (2008).



Experimental

Crystal data

$[Mn(C_{11}H_8N_5O)(N_3)]$
M_r = 323.19
Monoclinic, $P2_1/c$
a = 10.431 (2) Å
b = 14.431 (3) Å
c = 8.589 (2) Å
β = 90.676 (18)°
V = 1292.8 (5) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 1.03 mm^{-1}
T = 293 (2) K
0.20 × 0.16 × 0.12 mm

Data collection

Rigaku, SCXmini diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2005)
T_{min} = 0.820, T_{max} = 0.886
13382 measured reflections
3074 independent reflections
2472 reflections with $I > 2\sigma(I)$
R_{int} = 0.054

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.046
$wR(F^2)$ = 0.105
S = 1.11
3074 reflections
190 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.32 e Å^{-3}
$\Delta\rho_{min}$ = −0.41 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
C2—H2B···N8^{i}	0.97	2.50	3.223 (4)	131
C5—H5A···N3^{ii}	0.93	2.49	3.413 (4)	173

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

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; 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*.

This work was supported by a Start-up Grant awarded to Dr Heng-Yun Ye by Southeast University.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2100).

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supporting information

Acta Cryst. (2008). E**64**, m1060 [doi:10.1107/S1600536808022617]

Poly[(μ_2-azido-κ^2N^1:N^1)[μ_2-5-(8-quinolyloxymethyl)tetrazolato-κ^4N^1,O,N^5:N^4]manganese(II)]

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S1. Comment

In the past five years, we have focused on the chemistry of 5–substituted tetrazole because of their multiple coordination modes as ligand to metal ions and the construction of novel metal–organic framework (Wang *et al.* 2005; Xiong *et al.* 2002). As part of our on going studies of the chemistry of tetrazole, we determined the crystal structure of the title compound, *catena*–[(μ_2–1,1–azido)–(μ_2– η^1(*N*), η^3–(*N,N,O*)–((tetrazol–5–yl)methoxy) quinoline)–Manganese], **I** (Fig. 1).

As shown in Fig. 1, Mn1 is hexa–coordinated by five N and one O atoms, of which two N atoms and one O atom are from one organic ligand (tetrazol–5–yl)methoxy–quinoline, one N atom is from the tetrazole unit of another symmetry–related organic ligand (symmetry code: (iv) *x*, 3/2-*y*, 1/2+*z*) and two N atoms are from two azido anions which are symmetry–related (symmetry code: (iii) 1-*x*, 2-*y*, -*z*). The coordinated geometry of Mn1 is a distorted octahedron. The N4, N5, O1 and N6 atoms form the equatorial plane with mean deviation 0.1615Å of the plane (N4, N5, N6, O1 and Mn1). Geometry parameters of organic ligand are in normal ranges (Wang & Ye, 2007), dihedral angle of quinoline unit and the tetrazole unit is 7.41 (15)°. The Mn atoms are connected by azido anions and by (tetrazol–5–yl)methoxy–quinoline) ligands to form two–dimensional net framework parallel to the (1 0 0) plane (Fig.2). Beside the van der Waals forces, the crystal structure of **I** is also stabilized by intermolecular C—H···N^{ii} hydrogen bonds. Symmetry code: (ii) *x*-1, *y*, *z* (Fig. 3).

S2. Experimental

The precusor organic compound 8–cyanatoquinoline is was synthesized by using a similar procedure described by us before (Luo & Ye, 2008).A mixture of the organic ligand (34 mg, 0.2 mmol), NaN_3 (20 mg, 0.3 mmol), $MnCl_2$(25 mg, 0.2 mmol) and water (1 ml) sealed in a glass tube was maintained at 423 K. Yellow crystals suitable for X–ray analysis were obtained after 2 days.

S3. Refinement

All H atoms were positioned geometrically and refined using a riding model with d(C—H)$_{methine}$ = 0.98Å, d(C—H)$_{aryl}$ = 0.93Å, U_{iso} = 1.2U_{eq}(C).

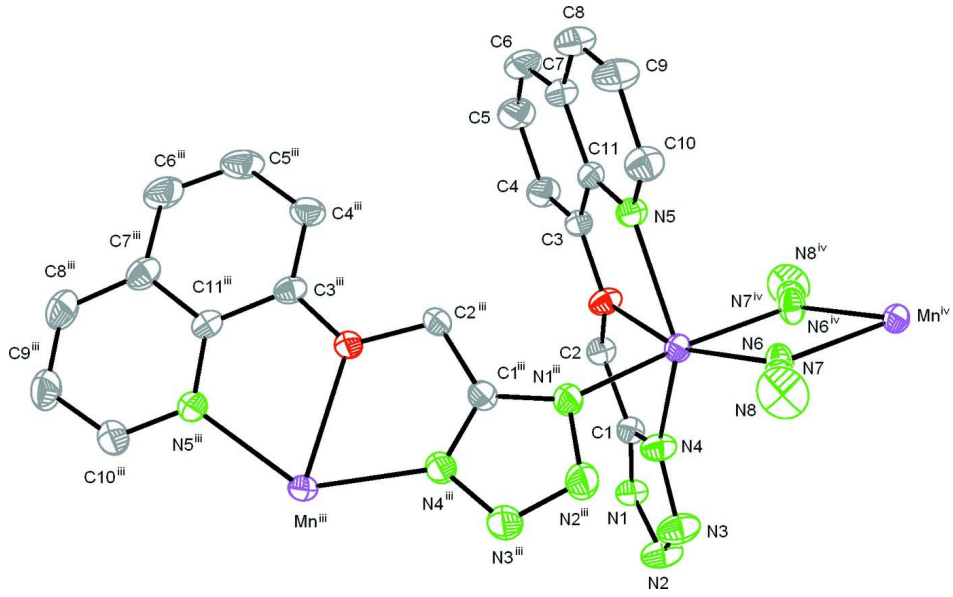

Figure 1

The fragment structure of **I** showing the atom numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. The hydrogen atoms are presented as small spheres of arbitrary radius. Symmetry codes: (iii) 1-*x*, 2-*y*, -*z*; (iv) *x*, 3/2-*y*, 1/2+*z*.

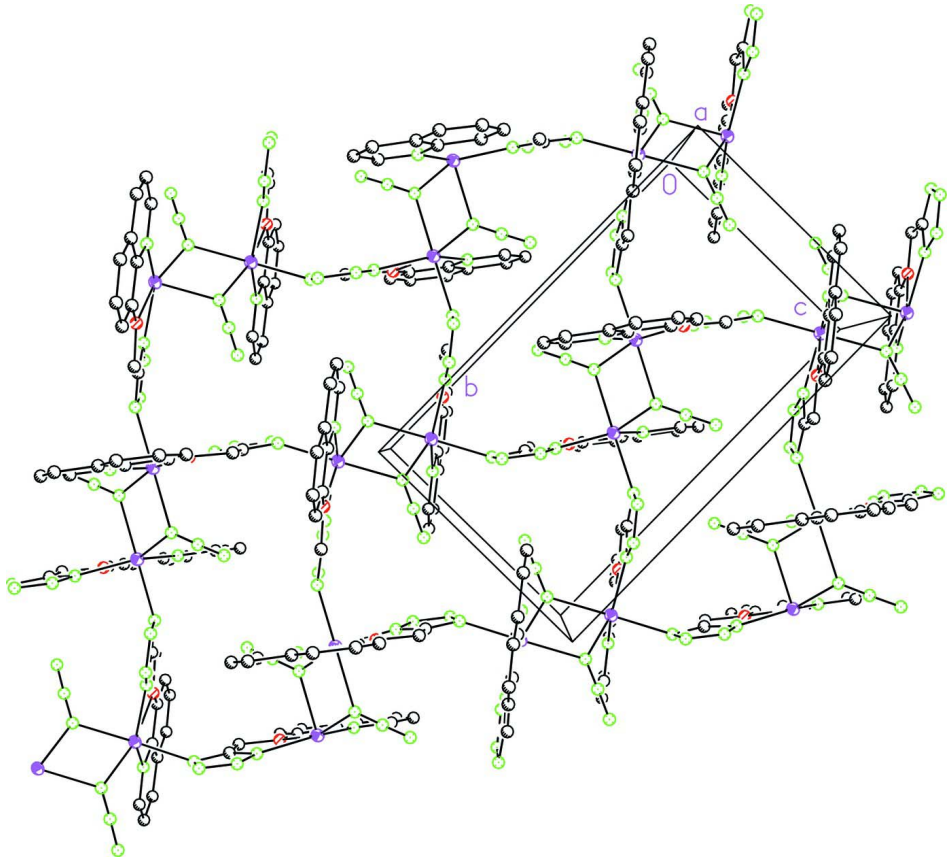

Figure 2

Two–dimensional net framework of the title compound view along *a* axis and all hydrogen atoms are omitted for clarity.

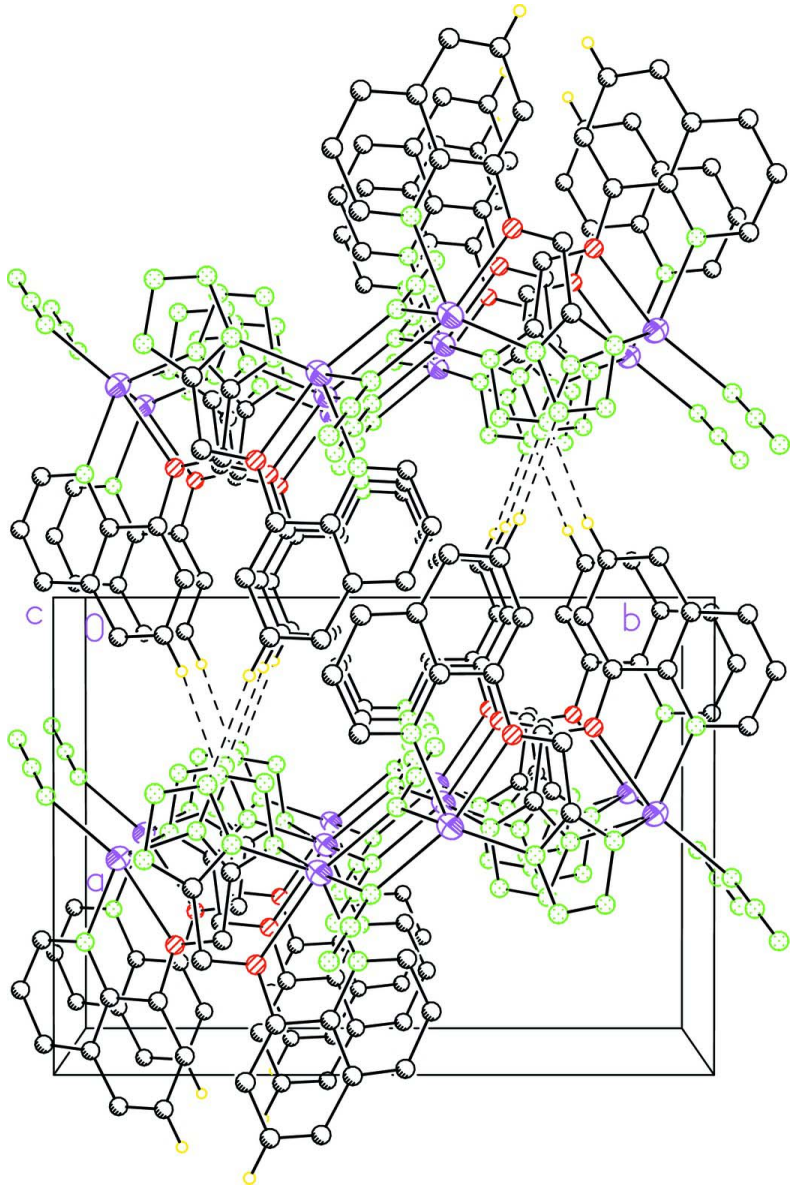

Figure 3

Crystal packing of the title compound viewed along the *c* axis. Hydrogen atoms not included in intermolecular hydrogen bonds are omitted for clarity. Dashed lines show the intermolecular hydrogen bonds.

Poly[(μ_2-azido-κ^2N^1:N^1)[μ_2-5-(8-quinolyloxymethyl)tetrazolato- κ^4N^1,O,N^5:N^4]manganese(II)]

Crystal data

[$Mn(C_{11}H_8N_5O)(N_3)$]
M_r = 323.19
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
a = 10.431 (2) Å
b = 14.431 (3) Å
c = 8.589 (2) Å
β = 90.676 (18)°
V = 1292.8 (5) Å^3
Z = 4
F(000) = 652
D_x = 1.661 Mg m^{-3}
Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 3074 reflections
θ = 2.8–27.9°
μ = 1.03 mm^{-1}

T = 293 K
Block, yellow

0.20 × 0.16 × 0.12 mm

Data collection

Rigaku, SCXmini diffractometer
Radiation source: Fine-focus sealed tube
Graphite monochromator
Detector resolution: 13.6612 pixels mm^{-1}
ω scans
Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2005)
T_{min} = 0.820, T_{max} = 0.886

13382 measured reflections
3074 independent reflections
2472 reflections with $I > 2\sigma(I)$
R_{int} = 0.054
θ_{max} = 27.9°, θ_{min} = 2.8°
h = −13→13
k = −19→19
l = −11→11

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.046
$wR(F^2)$ = 0.106
S = 1.11
3074 reflections
190 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-atom parameters constrained
$w = 1/[\sigma^2(F_o^2) + (0.0431P)^2 + 0.2319P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max}$ = 0.001
$\Delta\rho_{max}$ = 0.32 e Å^{-3}
$\Delta\rho_{min}$ = −0.41 e Å^{-3}

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 (Å^2)

	x	y	z	U_{iso}*/U_{eq}
Mn1	0.44828 (3)	0.90619 (3)	0.09904 (4)	0.03138 (13)
C1	0.4312 (2)	0.72913 (17)	−0.1025 (3)	0.0347 (6)
C2	0.2897 (2)	0.74195 (18)	−0.0956 (3)	0.0361 (6)
H2A	0.2548	0.7575	−0.1974	0.043*
H2B	0.2484	0.6860	−0.0588	0.043*
C3	0.1497 (2)	0.84401 (18)	0.0504 (3)	0.0369 (6)
C4	0.0389 (3)	0.8016 (2)	0.0034 (4)	0.0489 (7)
H4A	0.0410	0.7513	−0.0642	0.059*
C5	−0.0791 (3)	0.8355 (2)	0.0595 (5)	0.0636 (9)
H5A	−0.1549	0.8068	0.0280	0.076*
C6	−0.0843 (3)	0.9086 (2)	0.1577 (5)	0.0611 (9)
H6A	−0.1631	0.9291	0.1939	0.073*
C7	0.0296 (3)	0.9541 (2)	0.2056 (4)	0.0476 (7)
C8	0.0335 (3)	1.0319 (2)	0.3054 (4)	0.0598 (9)

H8A	−0.0425	1.0565	0.3429	0.072*
C9	0.1473 (3)	1.0708 (2)	0.3471 (4)	0.0622 (9)
H9A	0.1498	1.1213	0.4144	0.075*
C10	0.2607 (3)	1.0341 (2)	0.2875 (3)	0.0506 (7)
H10A	0.3380	1.0617	0.3161	0.061*
C11	0.1485 (2)	0.92186 (18)	0.1507 (3)	0.0350 (6)
N1	0.4916 (2)	0.66357 (15)	−0.1804 (3)	0.0402 (5)
N2	0.6184 (2)	0.67775 (19)	−0.1497 (3)	0.0551 (7)
N3	0.6309 (2)	0.74859 (19)	−0.0569 (3)	0.0569 (7)
N4	0.5125 (2)	0.78285 (16)	−0.0251 (3)	0.0431 (6)
N5	0.2634 (2)	0.96208 (15)	0.1926 (2)	0.0372 (5)
N6	0.5787 (2)	1.01876 (16)	0.1316 (3)	0.0417 (5)
N7	0.6346 (2)	1.04671 (15)	0.2407 (3)	0.0415 (5)
N8	0.6923 (3)	1.0756 (2)	0.3458 (3)	0.0701 (9)
O1	0.27145 (16)	0.81654 (12)	0.0115 (2)	0.0398 (4)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Mn1	0.0286 (2)	0.0303 (2)	0.0353 (2)	−0.00417 (15)	0.00011 (15)	0.00057 (15)
C1	0.0352 (14)	0.0325 (13)	0.0364 (13)	−0.0005 (10)	−0.0019 (11)	−0.0022 (10)
C2	0.0345 (14)	0.0334 (13)	0.0403 (14)	−0.0042 (11)	−0.0015 (11)	−0.0079 (11)
C3	0.0272 (12)	0.0390 (14)	0.0445 (15)	0.0012 (10)	0.0001 (11)	0.0068 (11)
C4	0.0340 (15)	0.0462 (17)	0.067 (2)	−0.0052 (12)	−0.0026 (13)	−0.0041 (14)
C5	0.0276 (15)	0.061 (2)	0.102 (3)	−0.0051 (14)	−0.0040 (16)	0.004 (2)
C6	0.0306 (15)	0.067 (2)	0.086 (3)	0.0049 (14)	0.0069 (16)	0.0028 (18)
C7	0.0363 (15)	0.0509 (17)	0.0558 (18)	0.0087 (13)	0.0078 (13)	0.0063 (14)
C8	0.0510 (19)	0.063 (2)	0.066 (2)	0.0169 (16)	0.0150 (16)	−0.0072 (17)
C9	0.057 (2)	0.063 (2)	0.066 (2)	0.0136 (17)	0.0090 (17)	−0.0211 (17)
C10	0.0492 (18)	0.0498 (18)	0.0529 (18)	−0.0006 (14)	0.0013 (14)	−0.0121 (14)
C11	0.0291 (13)	0.0397 (14)	0.0362 (14)	0.0020 (10)	0.0031 (10)	0.0072 (10)
N1	0.0336 (12)	0.0408 (13)	0.0461 (13)	0.0030 (9)	−0.0019 (10)	−0.0086 (10)
N2	0.0372 (13)	0.0645 (17)	0.0633 (17)	0.0066 (12)	−0.0057 (12)	−0.0228 (13)
N3	0.0349 (13)	0.0648 (17)	0.0708 (18)	0.0001 (12)	−0.0065 (12)	−0.0225 (14)
N4	0.0322 (12)	0.0428 (13)	0.0541 (14)	0.0026 (10)	−0.0034 (10)	−0.0125 (11)
N5	0.0346 (11)	0.0379 (12)	0.0392 (12)	0.0015 (9)	0.0021 (9)	−0.0013 (9)
N6	0.0456 (13)	0.0389 (12)	0.0405 (13)	−0.0140 (10)	−0.0067 (10)	0.0060 (10)
N7	0.0433 (13)	0.0348 (12)	0.0464 (14)	−0.0020 (10)	0.0019 (11)	0.0010 (10)
N8	0.083 (2)	0.073 (2)	0.0529 (17)	−0.0068 (16)	−0.0253 (16)	−0.0143 (14)
O1	0.0290 (9)	0.0403 (10)	0.0501 (11)	−0.0026 (8)	0.0035 (8)	−0.0124 (8)

Geometric parameters (Å, º)

Mn1—N6	2.135 (2)	C6—C7	1.415 (4)
Mn1—N4	2.184 (2)	C6—H6A	0.9300
Mn1—N1^{i}	2.188 (2)	C7—C11	1.411 (4)
Mn1—N5	2.247 (2)	C7—C8	1.412 (4)
Mn1—N6^{ii}	2.272 (2)	C8—C9	1.358 (5)

Mn1—O1	2.3682 (18)	C8—H8A	0.9300
C1—N4	1.322 (3)	C9—C10	1.399 (4)
C1—N1	1.322 (3)	C9—H9A	0.9300
C1—C2	1.490 (4)	C10—N5	1.322 (3)
C2—O1	1.430 (3)	C10—H10A	0.9300
C2—H2A	0.9700	C11—N5	1.376 (3)
C2—H2B	0.9700	N1—N2	1.361 (3)
C3—C4	1.364 (4)	N1—Mn1^{iii}	2.188 (2)
C3—O1	1.376 (3)	N2—N3	1.302 (3)
C3—C11	1.416 (4)	N3—N4	1.361 (3)
C4—C5	1.414 (4)	N6—N7	1.169 (3)
C4—H4A	0.9300	N6—Mn1^{ii}	2.272 (2)
C5—C6	1.353 (5)	N7—N8	1.157 (3)
C5—H5A	0.9300		

N6—Mn1—N4	119.03 (9)	C7—C6—H6A	119.9
N6—Mn1—N1^{i}	96.42 (8)	C11—C7—C8	116.5 (3)
N4—Mn1—N1^{i}	89.22 (9)	C11—C7—C6	119.2 (3)
N6—Mn1—N5	103.19 (9)	C8—C7—C6	124.3 (3)
N4—Mn1—N5	137.42 (8)	C9—C8—C7	120.5 (3)
N1^{i}—Mn1—N5	91.43 (8)	C9—C8—H8A	119.7
N6—Mn1—N6^{ii}	79.85 (9)	C7—C8—H8A	119.7
N4—Mn1—N6^{ii}	89.90 (9)	C8—C9—C10	119.1 (3)
N1^{i}—Mn1—N6^{ii}	175.15 (8)	C8—C9—H9A	120.4
N5—Mn1—N6^{ii}	92.44 (9)	C10—C9—H9A	120.4
N6—Mn1—O1	161.79 (8)	N5—C10—C9	123.2 (3)
N4—Mn1—O1	69.03 (7)	N5—C10—H10A	118.4
N1^{i}—Mn1—O1	100.11 (8)	C9—C10—H10A	118.4
N5—Mn1—O1	68.97 (7)	N5—C11—C7	122.7 (3)
N6^{ii}—Mn1—O1	84.01 (7)	N5—C11—C3	118.7 (2)
N4—C1—N1	111.6 (2)	C7—C11—C3	118.6 (3)
N4—C1—C2	122.5 (2)	C1—N1—N2	105.2 (2)
N1—C1—C2	125.9 (2)	C1—N1—Mn1^{iii}	132.27 (18)
O1—C2—C1	104.99 (19)	N2—N1—Mn1^{iii}	115.27 (17)
O1—C2—H2A	110.7	N3—N2—N1	109.1 (2)
C1—C2—H2A	110.7	N2—N3—N4	108.8 (2)
O1—C2—H2B	110.7	C1—N4—N3	105.3 (2)
C1—C2—H2B	110.7	C1—N4—Mn1	121.64 (17)
H2A—C2—H2B	108.8	N3—N4—Mn1	132.69 (18)
C4—C3—O1	125.4 (3)	C10—N5—C11	117.9 (2)
C4—C3—C11	121.5 (2)	C10—N5—Mn1	121.83 (19)
O1—C3—C11	113.0 (2)	C11—N5—Mn1	120.26 (17)
C3—C4—C5	118.8 (3)	N7—N6—Mn1	132.73 (19)
C3—C4—H4A	120.6	N7—N6—Mn1^{ii}	126.11 (18)
C5—C4—H4A	120.6	Mn1—N6—Mn1^{ii}	100.15 (9)
C6—C5—C4	121.6 (3)	N8—N7—N6	178.1 (3)
C6—C5—H5A	119.2	C3—O1—C2	120.2 (2)
C4—C5—H5A	119.2	C3—O1—Mn1	118.92 (15)

C5—C6—C7	120.2 (3)	C2—O1—Mn1	120.41 (14)
C5—C6—H6A	119.9		

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

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
C2—H2*B*···N8^{iv}	0.97	2.50	3.223 (4)	131
C5—H5*A*···N3^{v}	0.93	2.49	3.413 (4)	173

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