

N-Methylisosaloline from *Hammada scoparia*

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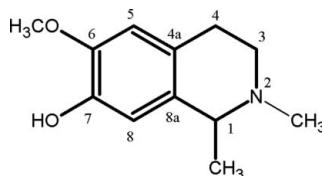
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Key indicators: single-crystal X-ray study; $T = 200\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.001\text{ \AA}$; R factor = 0.033; wR factor = 0.089; data-to-parameter ratio = 23.0.

The title compound (systematic name: 1,2-dimethyl-6-methoxy-1,2,3,4-tetrahydroisoquinolin-7-ol), $C_{12}H_{17}\text{NO}_2$, is a major alkaloid isolated from *Hammada scoparia* leaves. It belongs to the isoquinoline family and it was characterized by NMR spectroscopy and X-ray crystallographic techniques. The absolute configuration could not be reliably determined. An intermolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond is present in the crystal structure.

Related literature

For related literature on *Hammada scoparia* and isoquinoline alkaloids, see: Baker (1996); Benkrief *et al.* (1990); Carling & Sandberg (1970); El-Shazly & Wink (2003); El-Shazly *et al.* (2005); Iwasa *et al.* (2001); Jarraya & Damak (2001); Vetulani *et al.* (2001, 2003).



Experimental

Crystal data

$C_{12}H_{17}\text{NO}_2$	$V = 1089.33(14)\text{ \AA}^3$
$M_r = 207.27$	$Z = 4$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
$a = 7.5942(6)\text{ \AA}$	$\mu = 0.09\text{ mm}^{-1}$
$b = 10.8082(8)\text{ \AA}$	$T = 200(2)\text{ K}$
$c = 13.2716(10)\text{ \AA}$	$0.48 \times 0.37 \times 0.22\text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer	23859 measured reflections
Absorption correction: multi-scan (Becker & Coppens, 1974)	3132 independent reflections
$R_{\text{int}} = 0.030$	2870 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.961$, $T_{\max} = 0.988$	

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.033$	136 parameters
$wR(F^2) = 0.089$	H-atom parameters constrained
$S = 1.06$	$\Delta\rho_{\max} = 0.44\text{ e \AA}^{-3}$
3132 reflections	$\Delta\rho_{\min} = -0.24\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O2—H2 \cdots N1 ⁱ	0.84	1.90	2.6970 (10)	159

Symmetry code: (i) $x - \frac{1}{2}, -y + \frac{3}{2}, -z$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZL2131).

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supporting information

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N-Methylisosalsoline from *Hammada scoparia*

Raoudha Mezghani Jarraya, Amira Bouaziz, Besma Hamdi, Abdelhamid Ben Salah and Mohamed Damak

S1. Comment

Hammada scoparia has been reported to contain the alkaloids carnegine and *N*-methylisosalsoline as major tetrahydroisoquinoline alkaloids in addition to other minor alkaloids (Benkrief *et al.*, 1990; Jarraya & Damak 2001; El-Shazly & Wink, 2003).

The tetrahydroisoquinoline alkaloids affect the vegetative nervous system (Vetulani *et al.*, 2001 and 2003). Some of these alkaloids are known to be strong agonists at nicotinic acetylcholine receptors and it is thus likely that they serve as chemical defense compounds against insects and mammalian herbivores (El-Shazly *et al.*, 2005). Other simple isoquinoline alkaloids display potent, and often selective cytotoxicity or exhibit potential antimicrobial, antimalarial, antiviral and anti-HIV activities (Baker, 1996; Iwasa *et al.*, 2001).

The current study describes the isolation and the structure elucidation of *N*-methylisosalsoline. The structure of the title compound was established principally by two-dimensional NMR spectroscopy and through X-ray diffraction analysis although the absolute configuration could not be reliably determined.

The conformation of this compound is stabilized by an intermolecular hydrogen bond between the hydroxyl O₂—H₂ group and atom N₁ (Table 1). The molecules are assembled by intermolecular O—H···N hydrogen bonds (Table 1, Fig. 2)

S2. Experimental

The title compound was extracted from *Hammada scoparia* leaves:

Plant material *Hammada scoparia* (Pomel) Iljin= (*Haloxylon scoparium* (Pomel) Bge. = *Haloxylon articulatum* ssp *scorparium* (Pomel) Batt. = *Arthrophytum scoparium* (Pomel) Iljin), belongs to Chenopodiaceae family and is locally known as "rimth" in Sfax, Tunisia.

Leaves were carefully detached from the fresh plant, collected in June 2007 in Sfax, Tunisia, and air-dried. Voucher specimens (LCSN101) have been deposited at the " Laboratoire de Chimie des Substances Naturelles", Faculty of Science, University of Sfax, Tunisia.

Extraction and isolation of the *N*-Methylisosalsoline from *Hammada scoparia* leaves:

Air-dried leaves of *Hammada scoparia* were extracted at room temperature during 48 h with a mixture (EtOH-H₂O, 1–9, v-v). After filtration through folder filter paper Whatman N° 1, the ethanol was removed under reduced pressure and the remaining aqueous phase was acidified with HCl (pH = 3) and then defatted by extraction with CH₂Cl₂. The defatted mother liquor was made alkaline with an NH₄OH solution (pH = 10) and immediately extracted with CH₂Cl₂ to exhaustion. The latter CH₂Cl₂ extract was concentrated to yield a reddish-brown residue (total alkaloids).

The total Alkaloids (5 g) were separated, on column chromatography over silica gel 60 (0.063–0.200 mm; 160 g), using a gradient of dichloromethane-methanol as eluents. Eleven fractions were isolated according to their similarity by thin layer chromatography analyses. Further purifications gave two major pure alkaloids; the first is oily: Carnegie (1050

mg; fraction 3 eluted with dichloromethane) and the second (white rosette crystals) is *N*-methylisosaloline (545 mg; fraction 7 eluted with dichloromethane-methanol, 94–6, v–v). These alkaloids were previously isolated from *Hammada scoparia* (Carling & Sandberg, 1970; Benkrief *et al.*, 1990; Jarraya & Damak, 2001; El-Shazly & Wink, 2003). Their structures were determined on the basis of their spectral data such as UV, MS, ¹H NMR and H—H COSY, ¹³C NMR (BB, DEPT and C—H COSY, HMQC, HMBC, NOESY) and confirmed by comparison with published spectra.

N-Methylisosaloline (1-Methylcorypalline), white rosette crystals (MeOH), mp 443 K, UV λ_{max} (EtOH) nm = 207, 225, 285. λ_{max} (EtOH + OH[−]) nm = 213, 245, 300. EIMS, m/z (*rel. int.*): [M⁺] 207 (15), 193 (30), 192 (100), 177 (45), 164 (10), 149 (15), 121 (5), 96 (6), 91 (5), 77 (5), 57 (5), 42 (4). IR: (CHCl₃) ν_{max} (cm^{−1}): 3540, 2950, 2850, 2800, 1600, 1520.

Spectroscopic analysis, ¹H NMR (300 MHz, CDCl₃, p.p.m.): 1.34 (3H, d, J = 6.6 Hz, CH₃—C₁); 2.45 (3H, s, CH₃—N); 2.63 (1H, ddd, J = 11.4, 6.9, 5.1 Hz, H—C₃); 2.77 (2H, m, 2 H—C₄); 3.02 (1H, ddd, J = 11.4, 6.9, 5.1 Hz, H—C₃); 3.49 (1H, q, J = 6.6 Hz, H—C₁); 3.83 (3H, s, CH₃—O); 6.53 (1H, s, aromatic H, H—C₅); 6.63 (1H, s, aromatic H, H—C₈).

¹³C NMR (75.5 MHz, CDCl₃, p.p.m.): 58.51, C₁; 48.94, C₃; 27.36, C₄; 124.84, C_{4a}; 112.96, C₅; 145.31, C₆; 144.00, C₇; 110.57, C₈; 131.95, C_{8a}; 19.45, CH₃—C₁; 42.70, CH₃—N; 55.77, CH₃—O. The HMQC spectra showed correlations between 112.96 (C₅) and 6.53 (1H, s, aromatic H, H—C₅); 110.57 (C₈) and 6.63 (1H, s, aromatic H, H—C₈).

Suitable white X-ray quality crystals of this compound were obtained by recrystallization from methanol.

S3. Refinement

All H atoms attached to C atoms and O atom were fixed geometrically and treated as riding with C—H = 0.98 Å (Cmethine), 0.97 Å (Cmethylene), 0.96 Å (Cmethyl), 0.93 Å (Caromatic) and O—H = 0.84 Å with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{Cmethylene}, \text{Cmethine}, \text{Caromatic})$ or $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{Cmethyl}, \text{O})$.

In the absence of significant anomalous scattering, the absolute configuration could not be reliably determined and then the Friedel pairs were merged and any references to the Flack parameter were removed.

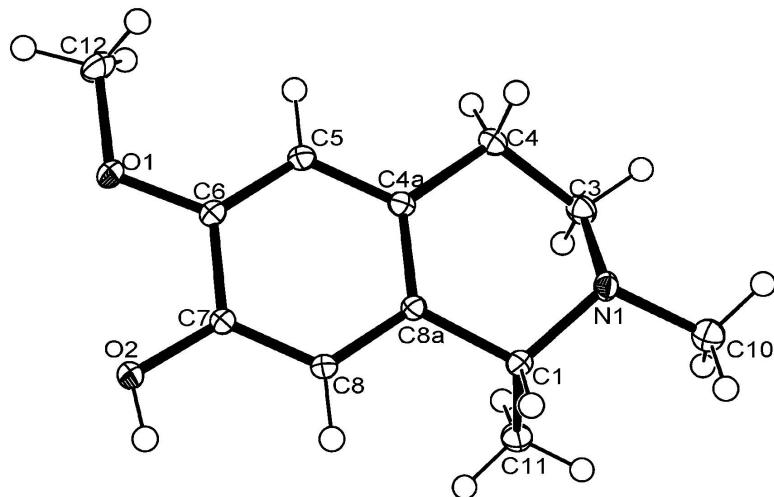
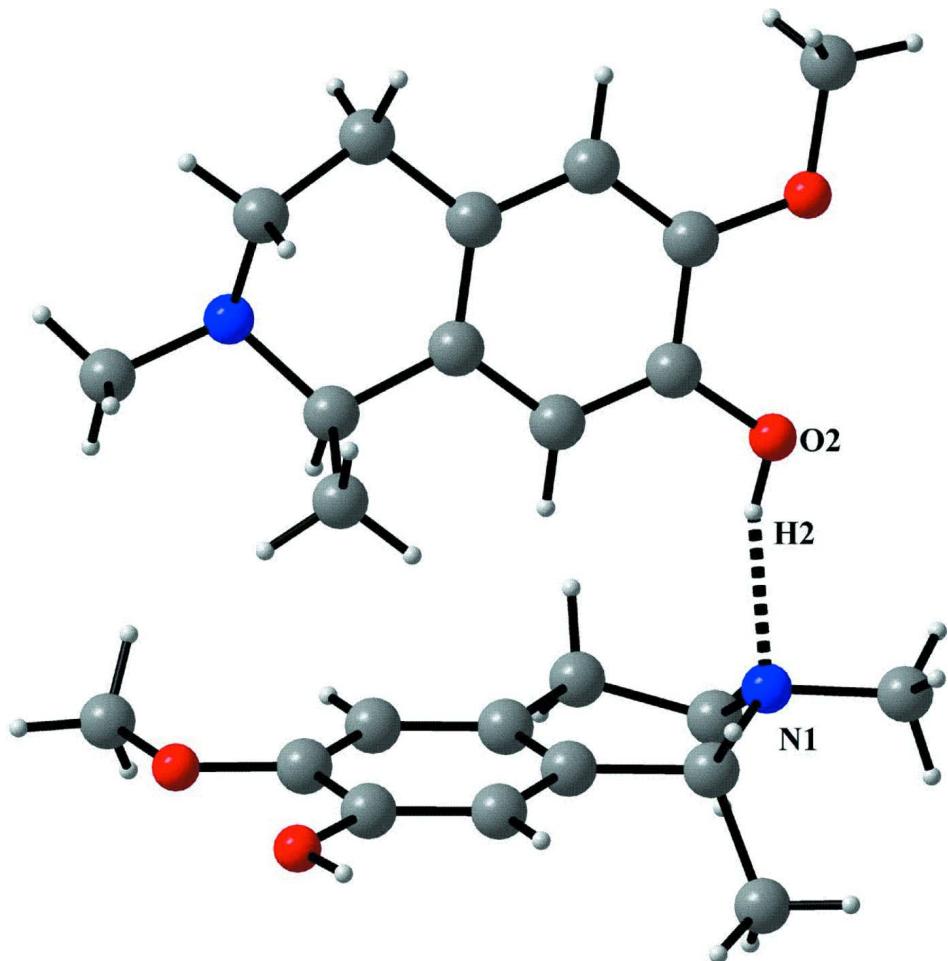


Figure 1

Molecular view of the title compound with the atom-labelling scheme. Ellipsoids are drawn at the 50% probability level. H atoms are represented as spheres of arbitrary radii.

**Figure 2**

Partial packing view showing the formation of pseudo dimer through O—H···N hydrogen bonds. Hydrogen bonds are shown as dashed lines.

1,2-dimethyl-6-methoxy-1,2,3,4-tetrahydroisoquinolin-7-ol

Crystal data

$C_{12}H_{17}NO_2$
 $M_r = 207.27$
Orthorhombic, $P2_12_12_1$
Hall symbol: P 2ac 2ab
 $a = 7.5942 (6)$ Å
 $b = 10.8082 (8)$ Å
 $c = 13.2716 (10)$ Å
 $V = 1089.33 (14)$ Å³
 $Z = 4$
 $F(000) = 448$

$D_x = 1.264$ Mg m⁻³
Melting point: 473 K
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 2100 reflections
 $\theta = 2.7\text{--}21.3^\circ$
 $\mu = 0.09$ mm⁻¹
 $T = 200$ K
Prism, colourless
0.48 × 0.37 × 0.22 mm

Data collection

Bruker SMART CCD area-detector
diffractometer
Radiation source: sealed tube

Graphite monochromator
 φ and ω scans

Absorption correction: multi-scan
 (Becker & Coppens, 1974)
 $T_{\min} = 0.961$, $T_{\max} = 0.988$
 23859 measured reflections
 3132 independent reflections
 2870 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$
 $\theta_{\max} = 37.1^\circ$, $\theta_{\min} = 2.4^\circ$
 $h = -12 \rightarrow 10$
 $k = -18 \rightarrow 18$
 $l = -22 \rightarrow 22$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.089$
 $S = 1.07$
 3132 reflections
 136 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
 $w = 1/[\sigma^2(F_o^2) + (0.0579P)^2 + 0.028P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.24 \text{ e } \text{\AA}^{-3}$

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.05540 (10)	0.85207 (7)	0.13064 (6)	0.01206 (13)
H1	0.0422	0.9013	0.0673	0.014*
C3	0.25806 (11)	0.83001 (8)	0.27194 (6)	0.01586 (14)
H3A	0.1557	0.8026	0.3124	0.019*
H3B	0.3386	0.8766	0.3167	0.019*
C4	0.35296 (11)	0.71807 (8)	0.22912 (6)	0.01475 (14)
H4A	0.4694	0.7437	0.2029	0.018*
H4B	0.3725	0.6567	0.2834	0.018*
C4A	0.24733 (10)	0.65931 (7)	0.14534 (6)	0.01156 (13)
C5	0.29165 (11)	0.54034 (7)	0.11057 (6)	0.01275 (13)
H5	0.3867	0.4973	0.1412	0.015*
C6	0.19977 (11)	0.48449 (7)	0.03258 (6)	0.01301 (13)
C7	0.05722 (11)	0.54781 (7)	-0.01256 (6)	0.01242 (13)
C8	0.01464 (10)	0.66497 (7)	0.02168 (6)	0.01194 (13)
H8	-0.0812	0.7078	-0.0084	0.014*
C8A	0.10928 (10)	0.72232 (7)	0.09969 (6)	0.01059 (12)
C10	0.14695 (12)	1.03330 (8)	0.22797 (7)	0.01802 (15)
H10A	0.1079	1.0854	0.1718	0.027*
H10B	0.2483	1.0720	0.2611	0.027*

H10C	0.0507	1.0242	0.2766	0.027*
C11	-0.12539 (11)	0.85179 (8)	0.18297 (7)	0.01757 (15)
H11A	-0.1569	0.9365	0.2022	0.026*
H11B	-0.1200	0.7998	0.2434	0.026*
H11C	-0.2144	0.8189	0.1367	0.026*
C12	0.37900 (13)	0.30406 (9)	0.03586 (8)	0.02182 (18)
H12A	0.3910	0.2239	0.0020	0.033*
H12B	0.3579	0.2908	0.1079	0.033*
H12C	0.4874	0.3520	0.0269	0.033*
N1	0.19793 (9)	0.91062 (6)	0.18960 (5)	0.01301 (12)
O1	0.23450 (9)	0.37016 (6)	-0.00665 (5)	0.01876 (13)
O2	-0.02759 (9)	0.49009 (5)	-0.08915 (5)	0.01773 (13)
H2	-0.1054	0.5372	-0.1126	0.027*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0121 (3)	0.0114 (3)	0.0127 (3)	0.0012 (2)	-0.0006 (2)	-0.0010 (2)
C3	0.0162 (3)	0.0185 (3)	0.0128 (3)	-0.0004 (3)	-0.0017 (3)	-0.0026 (3)
C4	0.0142 (3)	0.0154 (3)	0.0148 (3)	0.0000 (3)	-0.0041 (3)	-0.0009 (3)
C4A	0.0107 (3)	0.0121 (3)	0.0119 (3)	-0.0005 (2)	-0.0007 (2)	0.0007 (2)
C5	0.0119 (3)	0.0119 (3)	0.0144 (3)	0.0011 (2)	-0.0023 (2)	0.0009 (2)
C6	0.0132 (3)	0.0100 (3)	0.0159 (3)	0.0018 (2)	-0.0021 (3)	-0.0005 (2)
C7	0.0126 (3)	0.0102 (3)	0.0145 (3)	0.0007 (2)	-0.0027 (3)	-0.0004 (2)
C8	0.0116 (3)	0.0107 (3)	0.0135 (3)	0.0009 (2)	-0.0023 (2)	0.0000 (2)
C8A	0.0103 (3)	0.0100 (3)	0.0115 (3)	0.0001 (2)	-0.0002 (2)	0.0007 (2)
C10	0.0179 (4)	0.0149 (3)	0.0212 (3)	0.0006 (3)	0.0022 (3)	-0.0053 (3)
C11	0.0121 (3)	0.0191 (3)	0.0215 (3)	0.0012 (3)	0.0012 (3)	-0.0040 (3)
C12	0.0231 (4)	0.0173 (3)	0.0250 (4)	0.0099 (3)	-0.0070 (4)	-0.0022 (3)
N1	0.0132 (3)	0.0113 (3)	0.0145 (3)	-0.0006 (2)	0.0010 (2)	-0.0031 (2)
O1	0.0199 (3)	0.0113 (2)	0.0251 (3)	0.0056 (2)	-0.0085 (3)	-0.0043 (2)
O2	0.0189 (3)	0.0132 (2)	0.0211 (3)	0.0035 (2)	-0.0097 (2)	-0.0050 (2)

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

C1—N1	1.4780 (10)	C7—O2	1.3555 (10)
C1—C8A	1.5174 (10)	C7—C8	1.3837 (10)
C1—C11	1.5386 (12)	C8—C8A	1.4045 (10)
C1—H1	1.0000	C8—H8	0.9500
C3—N1	1.4703 (11)	C10—N1	1.4722 (10)
C3—C4	1.5185 (12)	C10—H10A	0.9800
C3—H3A	0.9900	C10—H10B	0.9800
C3—H3B	0.9900	C10—H10C	0.9800
C4—C4A	1.5110 (11)	C11—H11A	0.9800
C4—H4A	0.9900	C11—H11B	0.9800
C4—H4B	0.9900	C11—H11C	0.9800
C4A—C8A	1.3892 (10)	C12—O1	1.4257 (11)
C4A—C5	1.4070 (11)	C12—H12A	0.9800

C5—C6	1.3866 (11)	C12—H12B	0.9800
C5—H5	0.9500	C12—H12C	0.9800
C6—O1	1.3665 (10)	O2—H2	0.8400
C6—C7	1.4139 (11)		
N1—C1—C8A	109.97 (6)	C7—C8—C8A	121.76 (7)
N1—C1—C11	114.55 (6)	C7—C8—H8	119.1
C8A—C1—C11	111.16 (7)	C8A—C8—H8	119.1
N1—C1—H1	106.9	C4A—C8A—C8	119.42 (7)
C8A—C1—H1	106.9	C4A—C8A—C1	122.58 (7)
C11—C1—H1	106.9	C8—C8A—C1	117.99 (7)
N1—C3—C4	109.96 (7)	N1—C10—H10A	109.5
N1—C3—H3A	109.7	N1—C10—H10B	109.5
C4—C3—H3A	109.7	H10A—C10—H10B	109.5
N1—C3—H3B	109.7	N1—C10—H10C	109.5
C4—C3—H3B	109.7	H10A—C10—H10C	109.5
H3A—C3—H3B	108.2	H10B—C10—H10C	109.5
C4A—C4—C3	110.99 (7)	C1—C11—H11A	109.5
C4A—C4—H4A	109.4	C1—C11—H11B	109.5
C3—C4—H4A	109.4	H11A—C11—H11B	109.5
C4A—C4—H4B	109.4	C1—C11—H11C	109.5
C3—C4—H4B	109.4	H11A—C11—H11C	109.5
H4A—C4—H4B	108.0	H11B—C11—H11C	109.5
C8A—C4A—C5	119.05 (7)	O1—C12—H12A	109.5
C8A—C4A—C4	121.03 (7)	O1—C12—H12B	109.5
C5—C4A—C4	119.90 (7)	H12A—C12—H12B	109.5
C6—C5—C4A	121.49 (7)	O1—C12—H12C	109.5
C6—C5—H5	119.3	H12A—C12—H12C	109.5
C4A—C5—H5	119.3	H12B—C12—H12C	109.5
O1—C6—C5	125.51 (7)	C3—N1—C10	111.00 (7)
O1—C6—C7	115.09 (7)	C3—N1—C1	111.54 (6)
C5—C6—C7	119.39 (7)	C10—N1—C1	112.10 (7)
O2—C7—C8	123.81 (7)	C6—O1—C12	116.80 (7)
O2—C7—C6	117.30 (7)	C7—O2—H2	109.5
C8—C7—C6	118.87 (7)		
N1—C3—C4—C4A	-48.19 (9)	C4—C4A—C8A—C1	-0.30 (11)
C3—C4—C4A—C8A	15.68 (11)	C7—C8—C8A—C4A	-1.26 (12)
C3—C4—C4A—C5	-166.13 (7)	C7—C8—C8A—C1	178.70 (7)
C8A—C4A—C5—C6	-0.56 (12)	N1—C1—C8A—C4A	16.96 (10)
C4—C4A—C5—C6	-178.78 (7)	C11—C1—C8A—C4A	-110.97 (8)
C4A—C5—C6—O1	179.31 (8)	N1—C1—C8A—C8	-163.01 (7)
C4A—C5—C6—C7	-0.60 (12)	C11—C1—C8A—C8	69.06 (9)
O1—C6—C7—O2	-0.54 (11)	C4—C3—N1—C10	-165.48 (7)
C5—C6—C7—O2	179.38 (7)	C4—C3—N1—C1	68.74 (8)
O1—C6—C7—C8	-179.10 (7)	C8A—C1—N1—C3	-50.54 (8)
C5—C6—C7—C8	0.82 (12)	C11—C1—N1—C3	75.49 (8)
O2—C7—C8—C8A	-178.36 (8)	C8A—C1—N1—C10	-175.71 (6)

C6—C7—C8—C8A	0.10 (12)	C11—C1—N1—C10	-49.68 (9)
C5—C4A—C8A—C8	1.47 (11)	C5—C6—O1—C12	-0.82 (13)
C4—C4A—C8A—C8	179.67 (7)	C7—C6—O1—C12	179.10 (8)
C5—C4A—C8A—C1	-178.50 (7)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2···N1 ⁱ	0.84	1.90	2.6970 (10)	159

Symmetry code: (i) $x-1/2, -y+3/2, -z$.