

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

7-Hydroxy-8-isopropyl-1,1,4a-trimethyl-4a,9,10,10a-tetrahydro-phenanthren-2(1H)-one

Ahmed Benharref,^a Essêdiya Lassaba,^{a*} Noureddine Mazoir,^a Jean-Claude Daran^b and Moha Berraho^a

^aLaboratoire de Chimie Biomoléculaires, Substances Naturelles et Réactivité, URAC16, Université Cadi Ayyad, Faculté des Sciences Semlalia, BP 2390 Bd My Abdellah, 40000 Marrakech, Morocco, and ^bLaboratoire de Chimie de Coordination, 205 route de Narbonne, 31077 Toulouse Cedex 04, France
Correspondence e-mail: elassaba@gmail.com

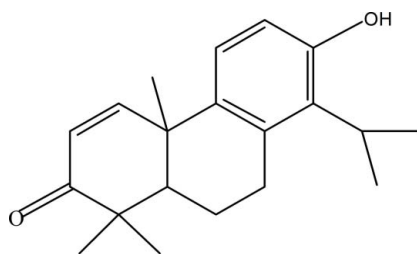
Received 7 March 2011; accepted 13 March 2011

Key indicators: single-crystal X-ray study; $T = 180$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.047; wR factor = 0.128; data-to-parameter ratio = 9.1.

The title compound, $\text{C}_{20}\text{H}_{26}\text{O}_2$, was isolated from a chloroform extract of *Tetraclinis articulata* wood. The molecule contains three fused rings which exhibit different conformations. The non-aromatic oxo-substituted ring has a screw-boat conformation, while the central ring has a half-chair conformation. In the crystal, molecules are linked to each other by intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds involving the carbonyl and hydroxy groups.

Related literature

For background to the biological activity of diterpenoids, see: Atta-ur-Rahman & Choudhary (1999); Azucena & Mobashery (2001); Panter *et al.* (2002); Ulusu *et al.* (2002). For their use in traditional medicine, see: Bellakhdar (1997) and for their medicinal properties, see: Barrero *et al.* (2003); Comte *et al.* (1995); Evidente *et al.* (1997). For the synthesis see: Zeroual *et al.* (2007). For conformational analysis, see: Cremer & Pople (1975).



Experimental

Crystal data

$\text{C}_{20}\text{H}_{26}\text{O}_2$
 $M_r = 298.41$

Monoclinic, $P2_1$
 $a = 11.6731$ (8) Å

$b = 6.4314$ (4) Å
 $c = 12.1488$ (10) Å
 $\beta = 111.592$ (9)°
 $V = 848.06$ (11) Å³
 $Z = 2$

Mo $K\alpha$ radiation
 $\mu = 0.07$ mm⁻¹
 $T = 180$ K
 $0.48 \times 0.36 \times 0.29$ mm

Data collection

Agilent Xcalibur Eos Gemini ultra diffractometer
9252 measured reflections

1893 independent reflections
1591 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.066$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.128$
 $S = 1.03$
1893 reflections
207 parameters

1 restraint
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.22$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H2}\cdots\text{O1}^i$	0.84	2.03	2.791 (3)	150

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

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

We thank the National Center of Scientific and Technological Research (CNRST) for supporting our scientific research.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FJ2405).

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supporting information

Acta Cryst. (2011). E67, o906 [doi:10.1107/S1600536811009561]

7-Hydroxy-8-isopropyl-1,1,4a-trimethyl-4a,9,10,10a-tetrahydro-phenanthren-2(1H)-one

Ahmed Benharref, Essêdiya Lassaba, Nouredine Mazoir, Jean-Claude Daran and Moha Berraho

S1. Comment

Among the diterpenoids, the class with the phenantrene skeleton may have biological activities which have been recently reported (Atta-ur-Rahman *et al.*, 1999; Azucena *et al.*, 2001; Panter *et al.*, 2002; Ulusu *et al.*, 2002). These diterpenoids are often isolated from the medicinal plants. In our study, we were interested to the medicinal plant, *Tetraclinis articulata* which is used in Moroccan traditional medicine (Bellakhdar, 1997). Some of its oxygenated compounds are effective as an antifongic (Evidente *et al.*, 1997), cytotoxic (Comte *et al.*, 1995) and inhibit various human leukocyte functions (Barrero *et al.*, 2003). In order to isolate similar compounds, we have studied the chloroform extract of *Tetraclinis articulata* wood. Thus, the extraction with chloroform using a soxhlet apparatus allow us isolate the totarolenone (7-hydroxy-8-isopropyl-1,1,4a-trimethyl-4a,9,10,10a-tetrahydro-1H-phenanthren-3-one) a derivative of totarolone (7-hydroxy-8-isopropyl-1,1,4a-trimethyl-3,4,4a,9,10,10a-hexahydrophenanthren-2-one) (Zeroual *et al.*, 2007). The structure of this new product was determined by NMR spectral analysis of ¹H, ¹³C and mass spectroscopy and confirmed by its single-crystal X-ray structure. The molecule (I) is built up from three fused six-membered rings. The non aromatic oxo-substituted ring has a screw boat conformation, as indicated by the total puckering amplitude QT = 0.490 (3) Å and spherical polar angle $\theta = 67.1$ (4)° with $\varphi = 272.2$ (3)°. While the central ring has a half chair conformation with QT = 0.545 (3) Å, $\theta = 53.4$ (3)°, $\varphi = 278.5$ (3)° (Cremer & Pople, 1975). Molecules are linked by intermolecular O—H...O hydrogen bonds (Table 1, Figure 2) involving the carbonyl and the hydroxy groups and propagate in chain parallel to the *a* axis.

S2. Experimental

50 g of wood powder *Tetraclinis articulata* was extracted with chloroform in Soxhlet apparatus during 24 h. After evaporating the solvent under reduced pressure, a residue of 3.2 g was obtained. Chromatography on a column of silica gel of this residue eluting with hexan - ethyl acetate (94 / 4), allowed us to isolate in pure form 640 mg of 7-hydroxy-8-isopropyl-1,1,4a-trimethyl-4a,9,10,10a-tetrahydro-1H-phenanthrene-3-one. Crystallization of this product was performed at room temperature in a solution of ethyl acetate.

S3. Refinement

All H atoms were fixed geometrically and treated as riding with C—H = 0.96 Å (methyl), 0.97 Å (methylene), 0.98 Å (methine) with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$ (methylene, methine) or $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}$ (methyl). In the absence of significant anomalous scattering, the absolute configuration could not be reliably determined and thus 1554 Friedel pairs were merged and any references to the Flack parameter were removed.

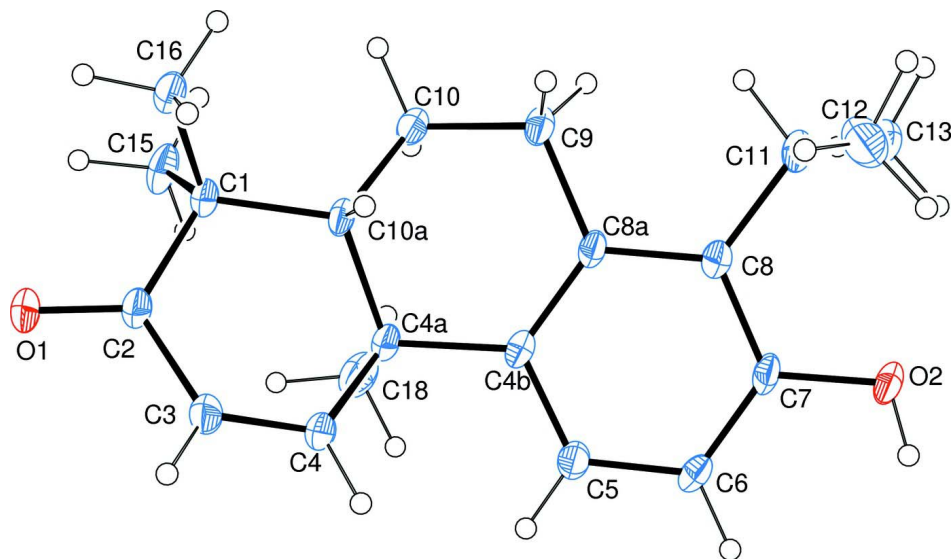


Figure 1

Molecular structure of the title compound with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii.

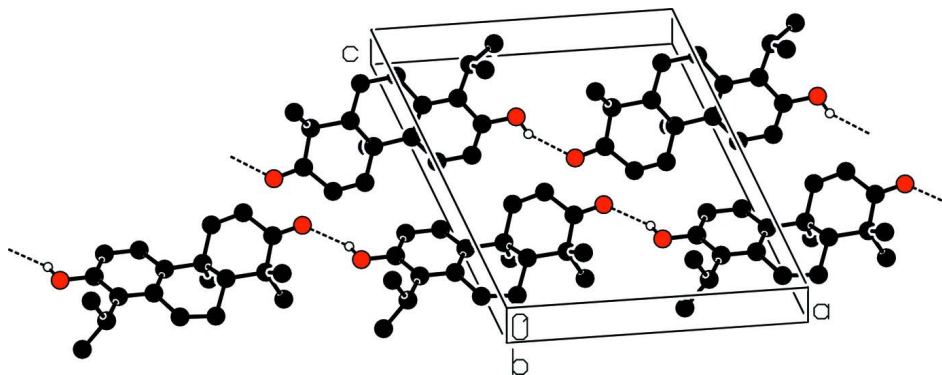


Figure 2

Partial packing view showing the O—H...O interactions (dashed lines) and the formation of a chain parallel to the *b* axis. H atoms not involved in hydrogen bonding have been omitted for clarity. [Symmetry code:(i) $x + 1, y, z$]

7-Hydroxy-8-isopropyl-1,1,4a-trimethyl-4a,9,10,10a-tetrahydro- phenanthren-2(1*H*)-one

Crystal data

$C_{20}H_{26}O_2$

$M_r = 298.41$

Monoclinic, $P2_1$

Hall symbol: $P\ 2y_b$

$a = 11.6731\ (8)\ \text{\AA}$

$b = 6.4314\ (4)\ \text{\AA}$

$c = 12.1488\ (10)\ \text{\AA}$

$\beta = 111.592\ (9)^\circ$

$V = 848.06\ (11)\ \text{\AA}^3$

$Z = 2$

$F(000) = 324$

$D_x = 1.169\ \text{Mg m}^{-3}$

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

Cell parameters from 3905 reflections

$\theta = 3.6\text{--}29.2^\circ$

$\mu = 0.07\ \text{mm}^{-1}$

$T = 180\ \text{K}$

Prism, colourless

$0.48 \times 0.36 \times 0.29\ \text{mm}$

Data collection

Agilent Xcalibur Eos Gemini ultra
diffractometer

Radiation source: Enhance (Mo) X-ray Source

Graphite monochromator

Detector resolution: 16.1978 pixels mm⁻¹

ω scans

9252 measured reflections

1893 independent reflections

1591 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.066$

$\theta_{\text{max}} = 26.4^\circ$, $\theta_{\text{min}} = 3.7^\circ$

$h = -14 \rightarrow 14$

$k = -8 \rightarrow 8$

$l = -15 \rightarrow 15$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.128$

$S = 1.03$

1893 reflections

207 parameters

1 restraint

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

$w = 1/[\sigma^2(F_o^2) + (0.0659P)^2 + 0.2266P]$

where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\text{max}} = 0.18 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.22 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	-0.3286 (2)	0.3531 (5)	0.7146 (2)	0.0290 (6)
C2	-0.4104 (2)	0.3898 (5)	0.5849 (3)	0.0314 (7)
C3	-0.3525 (3)	0.4063 (6)	0.4963 (3)	0.0364 (7)
H3	-0.4016	0.3822	0.4154	0.045 (10)*
C4	-0.2337 (3)	0.4540 (5)	0.5251 (3)	0.0324 (7)
H4	-0.2019	0.4667	0.4638	0.047 (10)*
C4A	-0.1473 (2)	0.4886 (5)	0.6526 (2)	0.0283 (6)
C4B	-0.0133 (2)	0.4386 (4)	0.6696 (2)	0.0274 (6)
C5	0.0337 (2)	0.5177 (5)	0.5876 (3)	0.0296 (6)
H5	-0.0179	0.6000	0.5237	0.036*
C6	0.1537 (2)	0.4788 (5)	0.5974 (3)	0.0303 (7)
H6	0.1846	0.5371	0.5420	0.036*
C7	0.2287 (2)	0.3544 (5)	0.6887 (2)	0.0299 (6)
C8	0.1866 (2)	0.2729 (5)	0.7742 (2)	0.0287 (6)
C8A	0.0651 (2)	0.3211 (5)	0.7647 (2)	0.0273 (6)
C9	0.0224 (2)	0.2470 (6)	0.8626 (3)	0.0351 (7)
H9A	0.0867	0.2814	0.9402	0.042*

H9B	0.0136	0.0938	0.8579	0.042*
C10	-0.0995 (2)	0.3422 (6)	0.8568 (2)	0.0336 (7)
H10A	-0.1329	0.2626	0.9079	0.040*
H10B	-0.0861	0.4875	0.8856	0.040*
C10A	-0.1905 (2)	0.3373 (5)	0.7292 (2)	0.0282 (6)
H10	-0.1817	0.1953	0.6997	0.034*
C11	0.2702 (3)	0.1345 (6)	0.8719 (3)	0.0383 (8)
H11	0.2229	0.0978	0.9231	0.046*
C12	0.3000 (3)	-0.0718 (6)	0.8246 (4)	0.0526 (9)
H12A	0.3534	-0.0452	0.7800	0.079*
H12B	0.2233	-0.1371	0.7725	0.079*
H21A	0.3421	-0.1649	0.8911	0.079*
C13	0.3870 (3)	0.2455 (7)	0.9521 (3)	0.0474 (9)
H13A	0.4396	0.2761	0.9071	0.071*
H13B	0.4317	0.1559	1.0195	0.071*
H13C	0.3646	0.3757	0.9810	0.071*
C15	-0.3538 (3)	0.5228 (6)	0.7922 (3)	0.0422 (8)
H15A	-0.4422	0.5272	0.7773	0.063*
H15B	-0.3274	0.6580	0.7727	0.063*
H15C	-0.3079	0.4913	0.8758	0.063*
C16	-0.3680 (3)	0.1419 (5)	0.7484 (3)	0.0371 (7)
H16A	-0.3634	0.0355	0.6925	0.056*
H16B	-0.4528	0.1515	0.7457	0.056*
H16C	-0.3130	0.1042	0.8286	0.056*
C18	-0.1553 (3)	0.7207 (5)	0.6809 (3)	0.0427 (8)
H18A	-0.1174	0.8047	0.6362	0.064*
H18B	-0.1117	0.7442	0.7658	0.064*
H18C	-0.2419	0.7606	0.6587	0.064*
O1	-0.52293 (18)	0.4045 (4)	0.5543 (2)	0.0440 (6)
O2	0.34674 (17)	0.3068 (4)	0.69940 (18)	0.0396 (6)
H2	0.3623	0.3635	0.6442	0.059*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0193 (12)	0.0352 (16)	0.0359 (15)	0.0034 (12)	0.0140 (11)	0.0023 (13)
C2	0.0210 (13)	0.0368 (17)	0.0389 (15)	0.0037 (12)	0.0138 (11)	0.0033 (13)
C3	0.0240 (14)	0.056 (2)	0.0299 (14)	0.0060 (14)	0.0106 (11)	0.0032 (14)
C4	0.0247 (14)	0.0416 (18)	0.0355 (15)	0.0089 (12)	0.0165 (12)	0.0062 (13)
C4A	0.0213 (13)	0.0343 (16)	0.0338 (14)	0.0029 (12)	0.0153 (11)	0.0026 (13)
C4B	0.0208 (13)	0.0313 (16)	0.0338 (14)	-0.0026 (11)	0.0145 (11)	-0.0017 (12)
C5	0.0245 (14)	0.0317 (15)	0.0345 (14)	0.0003 (12)	0.0129 (11)	0.0013 (12)
C6	0.0258 (14)	0.0365 (17)	0.0346 (14)	-0.0068 (13)	0.0182 (12)	-0.0015 (13)
C7	0.0166 (12)	0.0403 (17)	0.0354 (15)	-0.0036 (13)	0.0126 (11)	-0.0034 (14)
C8	0.0204 (13)	0.0335 (15)	0.0335 (14)	-0.0016 (12)	0.0115 (11)	0.0003 (12)
C8A	0.0180 (12)	0.0329 (15)	0.0334 (14)	-0.0017 (11)	0.0125 (11)	0.0000 (12)
C9	0.0195 (14)	0.0533 (19)	0.0340 (15)	-0.0013 (13)	0.0117 (11)	0.0041 (14)
C10	0.0235 (13)	0.0494 (19)	0.0310 (14)	-0.0024 (14)	0.0136 (11)	0.0025 (14)

C10A	0.0171 (12)	0.0396 (16)	0.0309 (14)	0.0030 (12)	0.0124 (10)	0.0005 (13)
C11	0.0193 (13)	0.055 (2)	0.0408 (16)	0.0001 (14)	0.0119 (12)	0.0067 (15)
C12	0.0414 (19)	0.049 (2)	0.061 (2)	0.0066 (16)	0.0101 (17)	0.0116 (18)
C13	0.0300 (17)	0.069 (3)	0.0414 (18)	-0.0075 (16)	0.0115 (14)	0.0022 (18)
C15	0.0279 (16)	0.053 (2)	0.054 (2)	-0.0006 (15)	0.0258 (15)	-0.0119 (17)
C16	0.0241 (14)	0.0423 (18)	0.0495 (18)	0.0017 (13)	0.0187 (13)	0.0097 (15)
C18	0.0426 (19)	0.0361 (19)	0.061 (2)	0.0025 (14)	0.0327 (17)	0.0003 (16)
O1	0.0197 (10)	0.0687 (17)	0.0459 (12)	0.0077 (10)	0.0146 (9)	0.0109 (12)
O2	0.0212 (10)	0.0626 (16)	0.0409 (12)	0.0021 (10)	0.0184 (9)	0.0044 (11)

Geometric parameters (Å, °)

C1—C2	1.531 (4)	C9—H9A	0.9900
C1—C16	1.538 (4)	C9—H9B	0.9900
C1—C15	1.539 (4)	C10—C10A	1.523 (4)
C1—C10A	1.559 (3)	C10—H10A	0.9900
C2—O1	1.230 (3)	C10—H10B	0.9900
C2—C3	1.471 (4)	C10A—H10	1.0000
C3—C4	1.336 (4)	C11—C13	1.530 (5)
C3—H3	0.9500	C11—C12	1.536 (5)
C4—C4A	1.522 (4)	C11—H11	1.0000
C4—H4	0.9500	C12—H12A	0.9800
C4A—C4B	1.535 (4)	C12—H12B	0.9800
C4A—C18	1.542 (4)	C12—H21A	0.9800
C4A—C10A	1.554 (4)	C13—H13A	0.9800
C4B—C5	1.397 (4)	C13—H13B	0.9800
C4B—C8A	1.401 (4)	C13—H13C	0.9800
C5—C6	1.385 (4)	C15—H15A	0.9800
C5—H5	0.9500	C15—H15B	0.9800
C6—C7	1.386 (4)	C15—H15C	0.9800
C6—H6	0.9500	C16—H16A	0.9800
C7—O2	1.371 (3)	C16—H16B	0.9800
C7—C8	1.403 (4)	C16—H16C	0.9800
C8—C8A	1.414 (4)	C18—H18A	0.9800
C8—C11	1.517 (4)	C18—H18B	0.9800
C8A—C9	1.526 (4)	C18—H18C	0.9800
C9—C10	1.526 (4)	O2—H2	0.8400
C2—C1—C16	106.1 (2)	C9—C10—H10A	109.8
C2—C1—C15	109.5 (2)	C10A—C10—H10B	109.8
C16—C1—C15	108.7 (2)	C9—C10—H10B	109.8
C2—C1—C10A	110.7 (2)	H10A—C10—H10B	108.3
C16—C1—C10A	108.1 (2)	C10—C10A—C4A	109.7 (2)
C15—C1—C10A	113.5 (2)	C10—C10A—C1	114.8 (2)
O1—C2—C3	120.0 (3)	C4A—C10A—C1	116.2 (2)
O1—C2—C1	121.0 (2)	C10—C10A—H10	104.9
C3—C2—C1	119.0 (2)	C4A—C10A—H10	104.9
C4—C3—C2	122.5 (3)	C1—C10A—H10	104.9

C4—C3—H3	118.8	C8—C11—C13	113.0 (3)
C2—C3—H3	118.8	C8—C11—C12	112.5 (3)
C3—C4—C4A	122.4 (2)	C13—C11—C12	111.7 (3)
C3—C4—H4	118.8	C8—C11—H11	106.3
C4A—C4—H4	118.8	C13—C11—H11	106.3
C4—C4A—C4B	111.5 (2)	C12—C11—H11	106.3
C4—C4A—C18	107.3 (3)	C11—C12—H12A	109.5
C4B—C4A—C18	108.4 (3)	C11—C12—H12B	109.5
C4—C4A—C10A	106.2 (2)	H12A—C12—H12B	109.5
C4B—C4A—C10A	109.2 (2)	C11—C12—H21A	109.5
C18—C4A—C10A	114.4 (2)	H12A—C12—H21A	109.5
C5—C4B—C8A	118.4 (2)	H12B—C12—H21A	109.5
C5—C4B—C4A	118.4 (2)	C11—C13—H13A	109.5
C8A—C4B—C4A	123.2 (2)	C11—C13—H13B	109.5
C6—C5—C4B	121.5 (3)	H13A—C13—H13B	109.5
C6—C5—H5	119.2	C11—C13—H13C	109.5
C4B—C5—H5	119.2	H13A—C13—H13C	109.5
C5—C6—C7	119.6 (2)	H13B—C13—H13C	109.5
C5—C6—H6	120.2	C1—C15—H15A	109.5
C7—C6—H6	120.2	C1—C15—H15B	109.5
O2—C7—C6	121.4 (2)	H15A—C15—H15B	109.5
O2—C7—C8	117.4 (2)	C1—C15—H15C	109.5
C6—C7—C8	121.2 (2)	H15A—C15—H15C	109.5
C7—C8—C8A	118.1 (3)	H15B—C15—H15C	109.5
C7—C8—C11	119.9 (2)	C1—C16—H16A	109.5
C8A—C8—C11	122.0 (2)	C1—C16—H16B	109.5
C4B—C8A—C8	121.1 (2)	H16A—C16—H16B	109.5
C4B—C8A—C9	120.7 (2)	C1—C16—H16C	109.5
C8—C8A—C9	118.2 (2)	H16A—C16—H16C	109.5
C8A—C9—C10	113.9 (3)	H16B—C16—H16C	109.5
C8A—C9—H9A	108.8	C4A—C18—H18A	109.5
C10—C9—H9A	108.8	C4A—C18—H18B	109.5
C8A—C9—H9B	108.8	H18A—C18—H18B	109.5
C10—C9—H9B	108.8	C4A—C18—H18C	109.5
H9A—C9—H9B	107.7	H18A—C18—H18C	109.5
C10A—C10—C9	109.2 (2)	H18B—C18—H18C	109.5
C10A—C10—H10A	109.8	C7—O2—H2	109.5

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2—H2 \cdots O1 ⁱ	0.84	2.03	2.791 (3)	150

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