

Dehydrobrachylaenolide: an eudesmane-type sesquiterpene lactone

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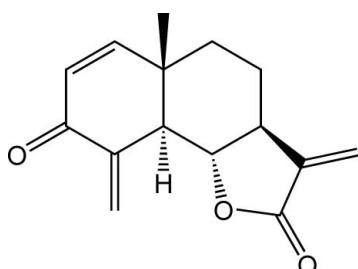
Received 26 November 2008; accepted 12 December 2008

Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(C-C) = 0.002$ Å; R factor = 0.033; wR factor = 0.095; data-to-parameter ratio = 14.1.

The three-ring eudesmanolide, $C_{15}H_{16}O_3$, is a natural product isolated from *Dicoma anomala* Sond. (Asteraceae). The compound contains an *endo-exo* cross conjugated methylenecyclohexenone ring with an envelope conformation fused with cyclohexane and *trans*-annelated with an α -methylene γ -lactone. The absolute structure was assigned by optical rotation measurements compared to those from the synthetic compound with known stereochemistry. The crystal packing is consolidated by C–H···O interactions.

Related literature

For NMR studies of this compound, see: Bohlmann & Zdero, (1982); Grass *et al.* (2004). For the chemical synthesis and confirmation of the absolute structure, see: Higuchi *et al.* (2003).



Experimental

Crystal data

$C_{15}H_{16}O_3$	$V = 1233.67$ (12) Å ³
$M_r = 244.28$	$Z = 4$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
$a = 9.5648$ (6) Å	$\mu = 0.09$ mm ⁻¹
$b = 11.1631$ (6) Å	$T = 150$ (2) K
$c = 11.5542$ (6) Å	$0.50 \times 0.50 \times 0.40$ mm

Data collection

Oxford Diffraction Excalibur2 CCD diffractometer	12604 measured reflections
Absorption correction: multi-scan (Blessing, 1995)	2294 independent reflections
$(Blessing, 1995)$	1988 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.909$, $T_{\max} = 0.963$	$R_{\text{int}} = 0.016$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.033$	163 parameters
$wR(F^2) = 0.095$	H-atom parameters constrained
$S = 1.05$	$\Delta\rho_{\max} = 0.31$ e Å ⁻³
2294 reflections	$\Delta\rho_{\min} = -0.21$ e Å ⁻³

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C6–H6···O1 ⁱ	0.98	2.39	3.360 (2)	171
C14–H14A···O1 ⁱ	0.96	2.57	3.393 (2)	143

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2006); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2006); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *PLATON* (Spek, 2003) and *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BI2330).

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supporting information

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Dehydrobrachylaenolide: an eudesmane-type sesquiterpene lactone

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S1. Comment

The title compound, a sesquiterpene lactone dehydrobrachylaenolide, was isolated from *Dicoma anomala* Sond (Asteraceae). These bi-functional *exo-endo* cross conjugated dienones are of importance as synthetic intermediates in the preparation of biologically active natural products (Higuchi *et al.*, 2003). NMR studies of the compound have been reported previously (Bohlmann & Zdero, 1982; Grass *et al.*, 2004) and the absolute stereochemistry has been confirmed as 3-oxoeudesma-1,4(15),11 (13)-triene-12,6a-olide by chemical synthesis (Higuchi *et al.*, 2003). Here we report the crystal structure. Although the absolute structure could not be elucidated by X-ray diffraction, unambiguous assignment of stereochemistry was made on the basis of the value of optical rotation ($[\alpha]^{24}_D +68^\circ$ (c 1/2, CHCl₃)) which is identical to that of the synthetic compound ($[\alpha]^{24}_D +67.9^\circ$ (c 0.16, CHCl₃)) for which the stereochemistry is known (Higuchi *et al.*, 2003) and very close to the value for the naturally isolated material ($[\alpha]^{24}_D +67^\circ$ (c 0.16, CHCl₃)) (Bohlmann & Zdero, 1982).

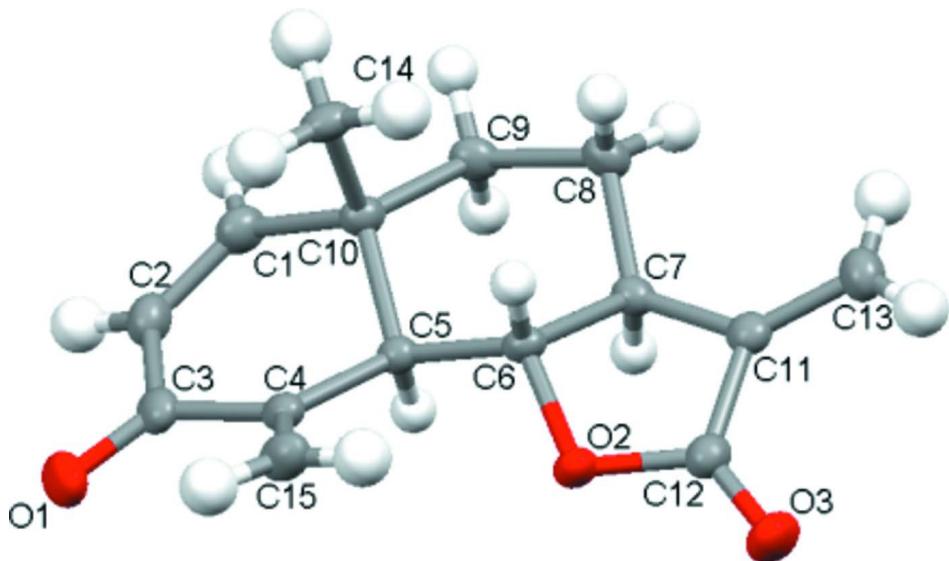
The molecular geometry and labelling scheme are shown in Fig. 1. The methylenecyclohexenone ring adopts an envelope conformation, with the C5 atom out of the plane of the ring by approximately 0.7 Å. The γ -lactone ring is twisted on C6—C7, while the cyclohexane ring adopts a chair conformation. An axial position is occupied by methyl group C14, and the methylene carbon atom C15 is in the equatorial position. A weak intramolecular interaction is formed between C15—H15B···O2. Fig. 2 illustrates the molecular packing viewed down the *c* axis. Weak intermolecular hydrogen bonds are present between atoms C6—H6···O1ⁱ and C14—H14···O1ⁱ [symmetry code (i): 1/2 - *x*, 1 - *y*, 1/2 + *z*].

S2. Experimental

The compound was isolated from *Dicoma anomala* Sond (Asteraceae), and recrystallized from propanol at room temperature.

S3. Refinement

H atoms were placed geometrically and refined in idealized positions in the riding-model approximation, with C—H = 0.93–0.98 Å with $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5U_{\text{eq}}(\text{C})$. In the absence of significant anomalous scattering effects, Friedel pairs were merged as equivalent data.

**Figure 1**

Molecular structure showing displacement ellipsoids at 50% probability for all atoms.

Dehydrobrachylaenolide

Crystal data

$C_{15}H_{16}O_3$
 $M_r = 244.28$
Orthorhombic, $P2_12_12_1$
Hall symbol: P 2ac 2ab
 $a = 9.5648 (6) \text{ \AA}$
 $b = 11.1631 (6) \text{ \AA}$
 $c = 11.5542 (6) \text{ \AA}$
 $V = 1233.67 (12) \text{ \AA}^3$
 $Z = 4$

$F(000) = 520$
 $D_x = 1.315 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 8167 reflections
 $\theta = 4.0\text{--}31.8^\circ$
 $\mu = 0.09 \text{ mm}^{-1}$
 $T = 150 \text{ K}$
Block, colourless
 $0.50 \times 0.50 \times 0.40 \text{ mm}$

Data collection

Oxford Diffraction Excalibur2 CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(Blessing, 1995)
 $T_{\min} = 0.909$, $T_{\max} = 0.963$

12604 measured reflections
2294 independent reflections
1988 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.016$
 $\theta_{\max} = 31.9^\circ$, $\theta_{\min} = 4.0^\circ$
 $h = -13 \rightarrow 13$
 $k = -15 \rightarrow 16$
 $l = -16 \rightarrow 17$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.095$
 $S = 1.05$
2294 reflections
163 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

$$w = 1/[\sigma^2(F_o^2) + (0.0708P)^2]$$

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

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

$$\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
C4	0.86180 (14)	0.08184 (11)	1.24094 (11)	0.0237 (2)
O2	0.87541 (10)	0.20733 (7)	1.00141 (7)	0.0245 (2)
C7	1.04877 (12)	0.07144 (11)	0.94161 (10)	0.0210 (2)
H7	1.1253	0.1211	0.9714	0.025*
C8	1.10043 (14)	-0.05782 (12)	0.94001 (11)	0.0246 (3)
H8A	1.1798	-0.0657	0.8883	0.029*
H8B	1.0268	-0.1109	0.9136	0.029*
C9	1.14347 (13)	-0.08979 (11)	1.06503 (11)	0.0245 (2)
H9A	1.1717	-0.1732	1.0675	0.029*
H9B	1.2238	-0.0416	1.0866	0.029*
O3	0.85244 (12)	0.31953 (9)	0.84156 (9)	0.0372 (3)
C14	0.90759 (14)	-0.16252 (11)	1.13671 (12)	0.0270 (3)
H14A	0.8694	-0.1540	1.0603	0.041*
H14B	0.8354	-0.1493	1.1930	0.041*
H14C	0.9447	-0.2419	1.1458	0.041*
C6	0.92565 (13)	0.08521 (10)	1.02492 (10)	0.0197 (2)
H6	0.8525	0.0278	1.0037	0.024*
C3	0.90807 (15)	0.04878 (12)	1.36063 (11)	0.0274 (3)
O1	0.84841 (13)	0.08529 (10)	1.44767 (9)	0.0385 (3)
C11	0.98993 (14)	0.13415 (11)	0.83767 (11)	0.0234 (2)
C1	1.08111 (15)	-0.08814 (12)	1.27617 (12)	0.0279 (3)
H1	1.1553	-0.1409	1.2867	0.033*
C5	0.97025 (12)	0.06224 (10)	1.14800 (10)	0.0197 (2)
H5	1.0491	0.1156	1.1647	0.024*
C10	1.02589 (13)	-0.06954 (10)	1.15444 (10)	0.0213 (2)
C15	0.73174 (15)	0.12080 (12)	1.22442 (14)	0.0321 (3)
H15A	0.6708	0.1272	1.2868	0.039*
H15B	0.7019	0.1416	1.1505	0.039*
C13	1.00061 (16)	0.11111 (13)	0.72528 (11)	0.0303 (3)
H13A	0.9501	0.1564	0.6722	0.036*
H13B	1.0586	0.0497	0.6996	0.036*
C12	0.89936 (14)	0.23105 (11)	0.88715 (11)	0.0263 (3)

C2	1.02875 (16)	-0.03277 (13)	1.36888 (11)	0.0304 (3)
H2	1.0696	-0.0460	1.4408	0.036*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C4	0.0291 (6)	0.0164 (5)	0.0257 (5)	-0.0021 (5)	0.0069 (5)	0.0005 (4)
O2	0.0309 (5)	0.0174 (4)	0.0253 (4)	0.0031 (3)	0.0029 (4)	0.0023 (3)
C7	0.0202 (5)	0.0206 (5)	0.0221 (5)	-0.0010 (4)	0.0013 (4)	-0.0030 (4)
C8	0.0239 (6)	0.0254 (6)	0.0244 (5)	0.0037 (5)	0.0003 (5)	-0.0048 (5)
C9	0.0200 (5)	0.0245 (6)	0.0289 (6)	0.0036 (5)	-0.0014 (5)	-0.0027 (4)
O3	0.0499 (7)	0.0262 (5)	0.0356 (5)	0.0061 (5)	0.0000 (5)	0.0081 (4)
C14	0.0277 (6)	0.0168 (5)	0.0366 (6)	-0.0020 (5)	-0.0027 (5)	0.0014 (5)
C6	0.0195 (5)	0.0148 (5)	0.0248 (5)	0.0006 (4)	0.0012 (4)	-0.0001 (4)
C3	0.0330 (6)	0.0233 (6)	0.0259 (5)	-0.0079 (5)	0.0076 (5)	0.0017 (5)
O1	0.0493 (6)	0.0366 (6)	0.0297 (5)	-0.0062 (5)	0.0165 (5)	-0.0008 (4)
C11	0.0233 (6)	0.0207 (5)	0.0261 (5)	-0.0040 (4)	0.0015 (5)	0.0001 (4)
C1	0.0288 (6)	0.0252 (6)	0.0297 (6)	0.0015 (5)	-0.0043 (5)	0.0036 (5)
C5	0.0202 (5)	0.0171 (5)	0.0217 (5)	-0.0007 (4)	0.0021 (4)	-0.0002 (4)
C10	0.0212 (5)	0.0186 (5)	0.0241 (5)	0.0011 (4)	-0.0022 (4)	0.0000 (4)
C15	0.0310 (7)	0.0263 (6)	0.0391 (7)	0.0033 (5)	0.0129 (6)	0.0040 (6)
C13	0.0314 (6)	0.0337 (7)	0.0257 (6)	-0.0050 (6)	0.0028 (6)	0.0006 (5)
C12	0.0303 (6)	0.0219 (6)	0.0266 (6)	-0.0028 (5)	-0.0002 (5)	0.0019 (4)
C2	0.0358 (7)	0.0301 (6)	0.0253 (6)	-0.0048 (5)	-0.0011 (5)	0.0048 (5)

Geometric parameters (\AA , ^\circ)

C4—C15	1.332 (2)	C14—H14B	0.960
C4—C3	1.4982 (18)	C14—H14C	0.960
C4—C5	1.5090 (16)	C6—C5	1.5067 (16)
O2—C12	1.3659 (15)	C6—H6	0.980
O2—C6	1.4708 (14)	C3—O1	1.2260 (16)
C7—C11	1.4997 (17)	C3—C2	1.473 (2)
C7—C8	1.5253 (18)	C11—C13	1.3278 (18)
C7—C6	1.5287 (16)	C11—C12	1.4991 (18)
C7—H7	0.980	C1—C2	1.334 (2)
C8—C9	1.5439 (18)	C1—C10	1.5167 (17)
C8—H8A	0.970	C1—H1	0.930
C8—H8B	0.970	C5—C10	1.5662 (15)
C9—C10	1.5437 (17)	C5—H5	0.980
C9—H9A	0.970	C15—H15A	0.930
C9—H9B	0.970	C15—H15B	0.930
O3—C12	1.2059 (16)	C13—H13A	0.930
C14—C10	1.5490 (17)	C13—H13B	0.930
C14—H14A	0.960	C2—H2	0.930
C15—C4—C3		O1—C3—C2	121.14 (13)
C15—C4—C5		O1—C3—C4	122.53 (13)

C3—C4—C5	114.71 (11)	C2—C3—C4	116.32 (11)
C12—O2—C6	107.66 (9)	C13—C11—C12	123.86 (13)
C11—C7—C8	123.60 (10)	C13—C11—C7	131.60 (13)
C11—C7—C6	99.69 (10)	C12—C11—C7	104.38 (10)
C8—C7—C6	110.63 (10)	C2—C1—C10	123.39 (12)
C11—C7—H7	107.3	C2—C1—H1	118.3
C8—C7—H7	107.3	C10—C1—H1	118.3
C6—C7—H7	107.3	C6—C5—C4	116.89 (10)
C7—C8—C9	107.08 (10)	C6—C5—C10	107.51 (9)
C7—C8—H8A	110.3	C4—C5—C10	109.63 (9)
C9—C8—H8A	110.3	C6—C5—H5	107.5
C7—C8—H8B	110.3	C4—C5—H5	107.5
C9—C8—H8B	110.3	C10—C5—H5	107.5
H8A—C8—H8B	108.6	C1—C10—C9	110.29 (10)
C10—C9—C8	113.46 (10)	C1—C10—C14	106.58 (10)
C10—C9—H9A	108.9	C9—C10—C14	110.22 (10)
C8—C9—H9A	108.9	C1—C10—C5	106.91 (10)
C10—C9—H9B	108.9	C9—C10—C5	110.69 (9)
C8—C9—H9B	108.9	C14—C10—C5	112.02 (9)
H9A—C9—H9B	107.7	C4—C15—H15A	120.0
C10—C14—H14A	109.5	C4—C15—H15B	120.0
C10—C14—H14B	109.5	H15A—C15—H15B	120.0
H14A—C14—H14B	109.5	C11—C13—H13A	120.0
C10—C14—H14C	109.5	C11—C13—H13B	120.0
H14A—C14—H14C	109.5	H13A—C13—H13B	120.0
H14B—C14—H14C	109.5	O3—C12—O2	121.23 (12)
O2—C6—C5	115.13 (9)	O3—C12—C11	129.75 (13)
O2—C6—C7	103.21 (9)	O2—C12—C11	109.00 (10)
C5—C6—C7	111.05 (10)	C1—C2—C3	121.90 (12)
O2—C6—H6	109.1	C1—C2—H2	119.1
C5—C6—H6	109.1	C3—C2—H2	119.1
C7—C6—H6	109.1		
C11—C7—C8—C9	-176.53 (11)	C15—C4—C5—C10	-124.67 (13)
C6—C7—C8—C9	-58.75 (13)	C3—C4—C5—C10	52.90 (13)
C7—C8—C9—C10	55.27 (14)	C2—C1—C10—C9	151.41 (13)
C12—O2—C6—C5	153.05 (11)	C2—C1—C10—C14	-88.95 (15)
C12—O2—C6—C7	31.88 (12)	C2—C1—C10—C5	31.01 (17)
C11—C7—C6—O2	-39.39 (11)	C8—C9—C10—C1	-173.06 (11)
C8—C7—C6—O2	-171.01 (9)	C8—C9—C10—C14	69.52 (13)
C11—C7—C6—C5	-163.29 (9)	C8—C9—C10—C5	-54.96 (13)
C8—C7—C6—C5	65.10 (12)	C6—C5—C10—C1	175.55 (10)
C15—C4—C3—O1	-20.6 (2)	C4—C5—C10—C1	-56.42 (12)
C5—C4—C3—O1	161.68 (12)	C6—C5—C10—C9	55.39 (12)
C15—C4—C3—C2	158.27 (13)	C4—C5—C10—C9	-176.57 (10)
C5—C4—C3—C2	-19.48 (16)	C6—C5—C10—C14	-68.06 (12)
C8—C7—C11—C13	-19.4 (2)	C4—C5—C10—C14	59.98 (12)
C6—C7—C11—C13	-142.26 (15)	C6—O2—C12—O3	170.70 (12)

C8—C7—C11—C12	155.99 (11)	C6—O2—C12—C11	−10.45 (13)
C6—C7—C11—C12	33.14 (12)	C13—C11—C12—O3	−21.0 (2)
O2—C6—C5—C4	58.69 (14)	C7—C11—C12—O3	163.19 (14)
C7—C6—C5—C4	175.49 (10)	C13—C11—C12—O2	160.33 (12)
O2—C6—C5—C10	−177.59 (9)	C7—C11—C12—O2	−15.53 (13)
C7—C6—C5—C10	−60.79 (12)	C10—C1—C2—C3	2.2 (2)
C15—C4—C5—C6	−2.05 (18)	O1—C3—C2—C1	169.28 (13)
C3—C4—C5—C6	175.53 (10)	C4—C3—C2—C1	−9.6 (2)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C6—H6···O1 ⁱ	0.98	2.39	3.360 (2)	171
C14—H14A···O1 ⁱ	0.96	2.57	3.393 (2)	143

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