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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.053 wR factor = 0.117 Data-to-parameter ratio = 8.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

2-(9-Hydroxy-3a,5a,5b,8,8,11a-hexamethylicosahydro-1*H*-cyclopenta[*a*]chrysen-1-yl)propanoic acid (3 β -hydroxylupan-29-oic acid)

The title compound, $C_{30}H_{50}O_3$, a lupane triterpene, was isolated from the leaves of *Ceriops decandra* (Griff.) Ding Hou. There are two crystallographically independent molecules in the asymmetric unit. In both molecules, the cyclopentane ring adopts an envelope conformation. In the crystal structure, molecules are interconnected into a two-dimensional network by intermolecular $O-H\cdots O$ hydrogen bonds.

Comment

Ceriops decandra (Griff.) Ding Hou (Rhizophoraceae) is a mangrove plant widely distributed from East Africa and Madagascar throughout tropical Asia and Queensland to Melanesia and Micronesia (Tomlinson, 1986). C. decandra has many local Thai names, e.g. Prong Khao and Prong Nu, and also a synonym of Ceriops roxburghiana Arn (Smitinand & Larsen, 1970). The bark of this plant has been used as a folk medicine for the treatment of diarrhoea, vomiting, amoebiasis and ulcers (Bamroongrugsa, 1999). An ethanol extract of the leaves has shown antinociceptive activity (Uddin et al., 2005). The title compound, 3β -hydroxylupan-29-oic acid, (I), was previously isolated from Gymnosporia wallichiana (Kulshreshtha, 1977). As part of our research on bioactive compounds from Thai medicinal plants (Chantrapromma et al., 2003, 2004; Waratchareeyakul et al., 2004; Boonnak et al., 2005), compound (I) was isolated for the first time from Ceriops decandra (Griff.) Ding Hou, collected from Phang-Nga province in the southern part of Thailand. We have undertaken the X-ray crystal structure analysis of (I) in order to establish its molecular structure and relative stereochemistry.



The asymmetric unit of (I) contains two crystallographically independent molecules, A and B, which have similar chiralities, bond lengths and angles (Fig. 1). The molecules are approximately related by a local twofold rotation axis. The bond lengths in (I) show normal values (Allen *et al.*, 1987). All the ring junctions in the lupane nucleus are *trans*-fused. In both molecules, the cyclohexane rings adopt chair conformations and the cyclopentane ring has an envelope conformation,

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved with atom C17 displaced from the C18/C22/C23/C24 plane by -0.692 (7) Å in molecule *A* and 0.674 (7) Å in molecule *B*. The hydroxyl group is equatorially attached at atom C3. The C23-C22-C25-C27 torsion angle of -55.1 (5)° [-45.7 (5)° in *B*] describes the orientation of the propanoic acid group with respect to the lupane nucleus. The C18-C22-C25-C26 torsion angle is -170.1 (4)° [-161.8 (3)° in *B*].

The molecular structure is stabilized by $C-H\cdots O$ hydrogen bonds (Table 2). $O-H\cdots O$ intermolecular hydrogen bonds link the molecules into a two-dimensional network parallel to the *ac* plane (Fig. 2).

Experimental

Air-dried leaves of *Ceriops decandra* (Griff.) Ding Hou (3.9 kg) were ground and extracted with hexane (2 × 20 l) at room temperature. The mixture was filtered and concentrated under reduced pressure to give a crude hexane extract (66.4 g). The crude extract was separated by quick column chromatography (QCC) on silica gel and eluted initially with hexane, followed by ethyl acetate and finally with methanol to give nine fractions (B1–B9). Fraction B7 (2.7 g) was subjected to column chromatography with MeOH–CH₂Cl₂ (0.2:9.8 v/ ν) and further purified by recrystallization from CHCl₃–EtOAc– MeOH (4:4:1) for a few days to obtain colourless single crystals of compound (I) [m.p. 518–520 K, $[\alpha]_D^{28}$ –42.6° (*c* = 0.125, MeOH)].

Crystal data

C ₃₀ H ₅₀ O ₃	$D_x = 1.103 \text{ Mg m}^{-3}$
$M_r = 458.70$	Mo $K\alpha$ radiation
Monoclinic, P2 ₁	Cell parameters from 14 091
a = 8.1669 (8) Å	reflections
b = 24.110 (2) Å	$\theta = 1.5 - 25.0^{\circ}$
c = 14.4666 (14) Å	$\mu = 0.07 \text{ mm}^{-1}$
$\beta = 104.057 (2)^{\circ}$	T = 293 (2) K
V = 2763.2 (4) Å ³	Plate, colourless
Z = 4	0.23 \times 0.15 \times 0.08 mm
Data collection	
Siemens SMART CCD area-	4986 independent reflections
detector diffractometer	3296 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.050$
Absorption correction: multi-scan	$\theta_{\rm max} = 25.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -9 \rightarrow 9$
$T_{\rm min} = 0.988, T_{\rm max} = 0.994$	$k = -23 \rightarrow 28$
14 091 measured reflections	$l = -14 \rightarrow 17$
D (1	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0486P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.053$	+ 0.1104P]
$wR(F^2) = 0.117$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} < 0.001$
4986 reflections	$\Delta \rho_{\rm max} = 0.16 \text{ e } \text{\AA}^{-3}$
596 parameters	$\Delta \rho_{\rm min} = -0.14 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

-			
O1A - C3A	1.452 (5)	O1B-C3B	1.445 (5)
O2A-C27A	1.234 (6)	O2B-C27B	1.236 (5)
O3A-C27A	1.264 (5)	O3 <i>B</i> -C27 <i>B</i>	1.284 (5)
O2A-C27A-O3A	122.9 (5)	O2B-C27B-O3B	123.8 (5)
O3A-C27A-C25A	118.1 (5)	O3B-C27B-C25B	116.0 (4)



Figure 1

The asymmetric unit of (I), showing 30% probability displacement ellipsoids and the atomic numbering. For clarity, H atoms have been omitted.

Table 2			
Hydrogen-bond	geometry	(Å,	°).

D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
0.82	1.91	2.705 (6)	161
0.82	1.86	2.662 (6)	165
0.82	1.98	2.724 (6)	151
0.82	1.82	2.625 (5)	163
0.98	2.52	3.262 (6)	132
0.98	2.46	3.206 (5)	133
0.97	2.49	3.222 (7)	132
0.97	2.49	3.237 (6)	134
0.96	2.56	2.957 (7)	105
0.96	2.51	2.917 (7)	105
	0.82 0.82 0.82 0.82 0.98 0.98 0.98 0.97 0.97 0.97 0.96 0.96	D=11 11.4.4 0.82 1.91 0.82 1.86 0.82 1.98 0.82 1.82 0.98 2.52 0.98 2.46 0.97 2.49 0.96 2.56 0.96 2.51	D=11 If YA D YA 0.82 1.91 2.705 (6) 0.82 1.86 2.662 (6) 0.82 1.98 2.724 (6) 0.82 1.82 2.625 (5) 0.98 2.52 3.262 (6) 0.97 2.49 3.222 (7) 0.97 2.49 3.237 (6) 0.96 2.56 2.957 (7) 0.96 2.51 2.917 (7)

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

H atoms were placed in calculated positions, with O–H distances of 0.82 Å and C–H distances in the range 0.93–0.98 Å. The $U_{\rm iso}$ values were constrained to be $1.5U_{\rm eq}$ of the carrier atom for hydroxyl and methyl H atoms, and $1.2U_{\rm eq}$ for the remaining H atoms. In the absence of significant anomalous dispersion effects, Friedel pairs were merged before the final refinement and the absolute configuration was assigned arbitrarily.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).





A view of the packing of (I). Only hydroxyl H atoms are shown. O– $H \cdots O$ hydrogen bonds are shown as dashed lines.

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