

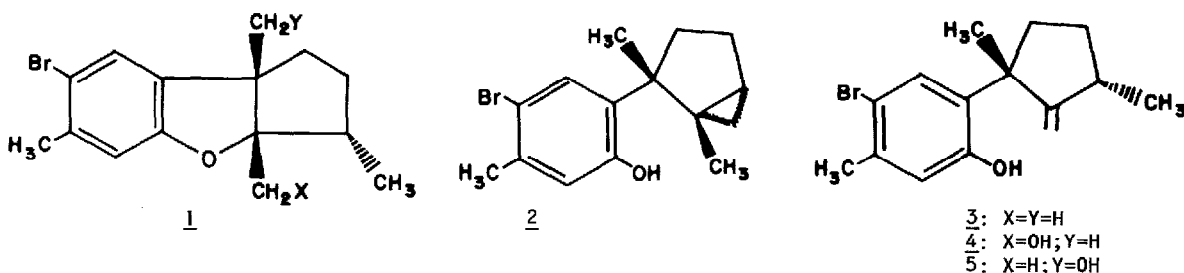
APLYSINOL FROM LAURENCIA DECIDUA: CRYSTAL STRUCTURE AND ABSOLUTE STEREOCHEMISTRY

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During the course of our extensive search for antibacterial substances¹ and halometabolites² among the samples collected on the 1974 Alpha Helix Baja Expedition¹⁻⁴ we have investigated by gas chromatography-mass spectrometry (gc-ms) the constituents of several species of the Laurencia genus.⁵ Laurencia decidua (AHBE-18-III-74-2-106)⁶ proved to be of special interest, since it contained a number of brominated sesquiterpenes other than the usual laurinterol (1), isolaurinterol (2), and aplysin (3).^{3,5}



One of the compounds tentatively identified by gc-ms appeared to be aplysinol. The gc-mass spectrum contained a molecular ion at m/e 310 ($C_{15}H_{19}^{79}BrO$) and fragment ions at m/e 277 ($M - CH_3 - H_2O$) and m/e 279 ($M - CH_2OH$), but no fragment ion due to loss of isoprene (or hydroxyisoprene) characteristic of laurinterol.³ Aplysinol was previously isolated from Aplysia kurodai⁷ and from Laurencia okamurai,⁸ supporting ingestion of red algae of this genus by the Aplysia sea hare. Stimulated by the tentative gc-ms identification, we elected to attempt to isolate the compound, and to establish its structure and absolute stereochemistry by X-ray crystallography, for two reasons. First, of the numerous brominated aromatic sesquiterpenes isolated from the genus Laurencia, the complete stereochemistry has been rigorously established by X-ray crystallography for only laurinterol (1),⁹ although chemical arguments have been advanced for the stereochemistry of aplysin (3) based on its formation from 1 in acid.⁹ Second, aplysinol had been isolated earlier and the tentative structure 4 assigned, based on biogenetic considerations, although the isomeric structure 5 could not be ruled out.⁷ These alternative structures were also pointed out in recent reviews.^{10,11}

L. decidua was extracted as previously described² and the toluene layer was chromatographed over silica gel, with fractions characterized by gc-ms. From a benzene eluate a

semisolid material was recovered and partially dissolved in acetonitrile, while undissolved material was dissolved in chloroform; on standing overnight the chloroform solution deposited a few crystals due to evaporation. The crystals were identified as *aplysinol* by *m.p.* as well as gas chromatographic coinjection and mixture *m.p.* with an authentic sample.^{1,2}

An X-ray structure determination was undertaken. The crystals were found to be *monoclinic*, $a = 13.753(5)$, $b = 7.319(4)$, $c = 13.746(4)$ Å, $\beta = 90.27(4)^\circ$, space group, $P2_1$ with four molecules per unit cell. A small, colorless crystal approximately $0.3 \times 0.1 \times 0.05$ mm was used to collect data on a Syntex $P2_1$ diffractometer ($\text{CuK}\alpha$, $\lambda = 1.54178$ Å). Heavy atom methods were used to solve the structure; the Patterson map revealed that four bromine atoms in the unit cell were related by near-perfect 4_2 symmetry. Anomalous dispersion corrections were applied to the scattering due to the bromine atoms and all non-hydrogen atoms were refined by full-matrix least squares methods incorporating anisotropic temperature factors. The hydrogen atoms were given isotropic temperature factors. The final R -factors on 1503 observed reflections were R_1 (unweighted) = 0.076, R_2 (weighted) = 0.055 for the enantiomer shown below. The other possible enantiomer refined to R -factors of $R_1 = 0.080$, and $R_2 = 0.059$. The atomic co-ordinates are listed in Table I. A stereoscopic drawing of the two molecules in the asymmetric unit is shown in Figure 1. The hydroxyl oxygen atoms form hydrogen bonds which join the two enantiomeric molecules together into a spiral column around the 2_1 axis in the y -direction. The molecules in this column have no close contacts in the x - or z -direction. The closest Br---Br distance is 4.27 Å. A stereoscopic diagram showing the packing is given in Figure 2.

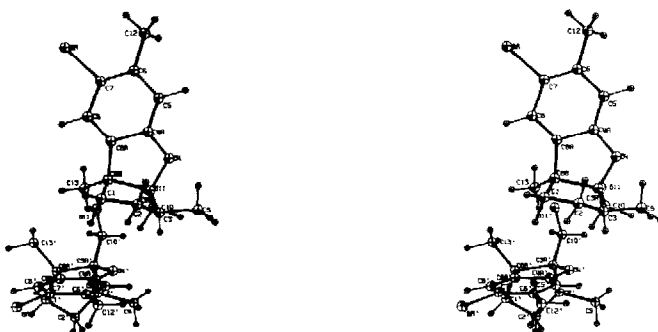


Figure 1. Stereoscopic drawing of the two molecules in the asymmetric unit.

Table I: Final Atomic Coordinates (Standard Deviations in Parentheses)

ATOM	x	y	z	ATOM	x	y	z
Br	0.8350(2)	0.50000(0) ^a	0.8886(2)	Br ¹	0.1107(2)	0.7490(5)	0.8351(2)
O(4)	0.5345(9)	-0.0565(15)	0.7858(9)	O(4 ¹)	0.2156(8)	0.2012(16)	0.5366(8)
O(11)	0.4573(9)	0.0160(19)	0.5927(8)	O(11 ¹)	0.4083(8)	0.2716(21)	0.4607(8)
C(1)	0.417(1)	0.321(3)	0.876(1)	C(1 ¹)	0.124(1)	0.569(2)	0.413(1)
C(2)	0.399(2)	0.144(3)	0.934(1)	C(2 ¹)	0.065(1)	0.396(2)	0.397(1)
C(3)	0.376(1)	0.013(3)	0.859(1)	C(3 ¹)	0.145(1)	0.255(3)	0.379(1)
C(3a)	0.447(1)	0.041(2)	0.768(1)	C(3a ¹)	0.228(1)	0.294(3)	0.446(1)
C(4a)	0.607(1)	0.057(3)	0.810(1)	C(4a ¹)	0.188(1)	0.308(2)	0.609(1)
C(5)	0.701(1)	-0.001(3)	0.835(1)	C(5 ¹)	0.169(1)	0.268(3)	0.703(1)
C(6)	0.773(2)	0.138(3)	0.859(1)	C(6 ¹)	0.144(1)	0.381(3)	0.770(1)
C(7)	0.745(2)	0.318(3)	0.859(1)	C(7 ¹)	0.149(1)	0.568(3)	0.744(1)
C(8)	0.648(2)	0.377(3)	0.831(1)	C(8 ¹)	0.167(1)	0.627(3)	0.648(1)
C(8a)	0.580(1)	0.251(4)	0.808(1)	C(8a ¹)	0.193(1)	0.503(3)	0.582(1)
C(8b)	0.474(1)	0.265(3)	0.783(1)	C(8b ¹)	0.213(1)	0.513(3)	0.474(1)
C(9)	0.382(2)	-0.188(4)	0.889(2)	C(9 ¹)	0.105(2)	0.046(3)	0.378(1)
C(10)	0.402(1)	-0.000(3)	0.677(1)	C(10 ¹)	0.325(1)	0.237(3)	0.403(1)
C(12)	0.881(2)	0.090(3)	0.875(1)	C(12 ¹)	0.124(1)	0.331(3)	0.873(1)
C(13)	0.457(2)	0.381(3)	0.699(1)	C(13 ¹)	0.299(1)	0.641(3)	0.457(1)

^aThis coordinate was held constant to define the origin.

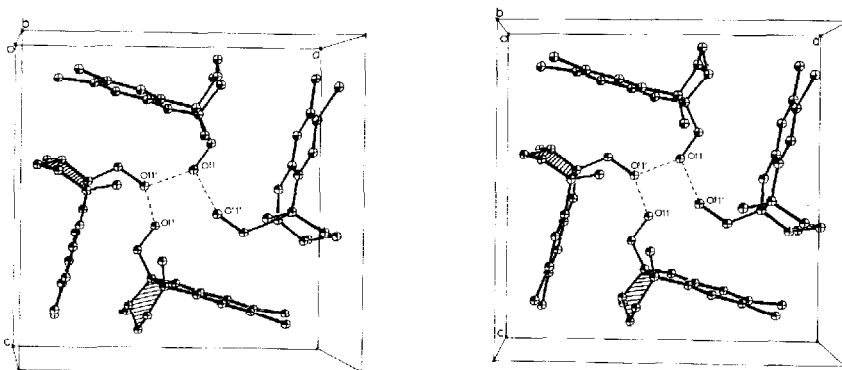


Figure 2. Stereoscopic view of the packing looking along the b-axis. Parts of the two reference molecules have been shaded. The dashed lines indicate hydrogen bonding.

The bond lengths and angles are not unusual except for the C(8b)-C(3a) bond (1.69(3) Å) and the C(8b¹)-C(3a¹) bond (1.66(3) Å). There is severe steric hindrance around these two quaternary substituted atoms; in fact, the C(13)-C(10) distance measured by Dreiding models,

using the normal bond lengths, would be 2.85 Å, well below the outer limit calculated using pair-wise Lennard-Jones potential interactions and the coefficients from Scott and Scheraga.¹³ The observed lengthening of the C(8b)-C(3a) bond partly releases this energy strain and the actual C(13)---C(10) non bonded distance is 3.45(3) Å.

These results confirm the structure 4 and the absolute stereochemistry of aplysinol is seen to be the same as that of laurinterol, in keeping with a related biosynthetic pathway.

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