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## ENT-NORAMBREINOLIDE FROM SIDERITIS NUTANS

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Key Word Index—Sideritis nutans; Labiateae; diterpenoid; ent-nor!mbreinolide; X-ray analysis.

Abstract—ent-Norambreinolide has been isolated from Sideritis nutans and its structure determined by X-ray analysis.

Continuing with our work on the phytochemical study of the genus *Sideritis*, endemic to the Canary Islands [1], we describe here the X-ray structure determination of *ent*norambreinolide (1), isolated from *Sideritis nutans* Svent.

This substance, molecular formula  $C_{16}H_{26}O_2$ , has IR absorptions characteristic of a lactone and two geminal methyl groups. Its <sup>1</sup>H NMR spectrum shows resonances typical of four methyls and of two hydrogens on a carbon allylic to a carbonyl group. In the mass spectrum, the molecular ion (m/z 250) readily loses a methyl group (m/z 235) or carbon dioxide (m/z 206). The loss of both gives a fragment at m/z 191. As only a small quantity of 1 was isolated, an X-ray analysis was carried out. Figure 1 shows its molecular structure. The absolute configuration was determined from the CD curve. *ent*-Norambreinolide in methanol exhibited a negative maximum at 215 nm characteristic of the structure 1. Its enantiomer has a positive Cotton effect [2].

Compound 1 has not been obtained previously as a

natural compound, but its enantiomer has been prepared synthetically by sclareol oxidation [3-5], by ozonization of  $12\alpha$ -hydroxy-13-epimanoyl oxide [6], and by degradation of a manoyloxide derivative [7].

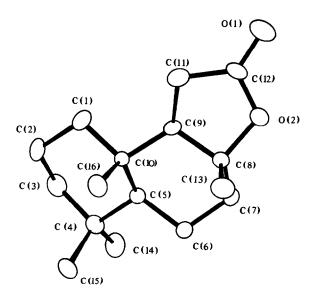


Fig 1. An ORTEP [10] drawing showing the relative molecular structure of *ent*-norambreinolide (1).

## **EXPERIMENTAL**

A general description of the procedure to isolate the terpenoid compounds of species of the genus *Sideritis* has been published previously [1].

ent-Norambreinolide (1). From 700 g of Sideritis nutans 15 mg 1 was isolated. Mp 120–122° (from petrol–EtOAc), [M]<sup>+</sup> at m/z 250.1931 (calc. for  $C_{16}H_{26}O_2$ , 250.1933). IR  $v_{\rm max}$  cm<sup>-1</sup>: 3000, 2950, 2860, 2840, 1760, 1450, 1385, 1380, 1280, 1260, 1190, 1115, 1020, 1010, 950, 925, 910. <sup>1</sup>H NMR (60 MHz, CDCl<sub>3</sub>):  $\delta$ 0.84, 0.88, 0.92 and 1.33 (each 3H, s), 2.27 (2H, s). EIMS (rel. int.): 250 [M]<sup>+</sup> (1), 235 (18), 207 (11), 206 (9), 191 (7), 150 (9), 149 (6), 137 (19), 136 (14).

Crystal data.  $C_{16}H_{26}O_2$ , MW = 250 4, monoclinic, a = 7.583 (1), b = 10.850 (2), c = 9.504 (1) A,  $\beta = 110.96$  (2)°, U = 730.2 A, Z = 2,  $Dc = 1.14 \ g/cm^3$ , F(000) = 276. Monochromated MoK $\alpha$  radiation,  $\lambda = 0.71069$  A,  $\mu = 0.8 \ cm^{-1}$  Space group  $P2_1$  from systematic absences of 0k0 for k odd and successful structure refinement.

Data were measured on an Enraf-Nonius CAD4 diffractometer using a crystal of size  $ca \ 0.2 \times 0.2 \times 0.2$  mm. Preliminary cell dimensions were found using the SEARCH and INDEX routines of the CAD4 and final values were calculated from the setting angles for 25 reflections with  $\theta \simeq 15^{\circ}$ . Intensities for  $hk \pm 1$  reflections with  $2 < \theta < 25^{\circ}$  were measured by a  $\theta/2\theta$  scan with a scan width of  $\Delta\theta = (0.8 + 0.35 \tan \theta)^{\circ}$  The scan rate for each reflection was determined by a rapid pre-scan at 10°/min1 in  $\theta$ , at which point any reflection with  $I < \sigma(I)$  was coded as unobserved. The remaining reflections were re-scanned at such a speed as to give a minimum value of  $\sigma(I)/I$  of 0.05 subject to a maximum scan time of 60 sec. Two standard reflections monitored every hour showed no significant variation. Data were corrected for Lp effects but not for absorption, and after averaging any equivalent reflections 1109 reflections with  $|F^2| > \sigma$  (F<sup>2</sup>) were used in the structure refinement. The values of  $\sigma(F^2)$  were taken as  $|\sigma^2(I) + (0.021)^2|^{1/2}/Lp$ .

The structure was solved by direct methods using the MULTAN [8] program. Refinement of non-hydrogen atoms

with anisotropic temp. factors was by full-matrix least-squares. Hydrogen atoms were placed at calculated positions (C-H 1.08 A) and held fixed with a common isotropic temp. factor of B = 6.0 A [9]. Refinement converged at R = 0.053, R' = 0.059, when the maximum shift/error was 0.01 and the weighting scheme was  $w = 1/\sigma^2$  (F). A final difference map was completely featureless.

The structure solution and refinement were done on a PDP 11/34 computer using the Enraf-Nonius Structure Determination Package. Scattering factors for neutral atoms were taken from ref. [9].

Final atom coordinates, a list of temperature factors, hydrogen atom positions, and final structure factors have been deposited at the Cambridge Crystallographic Data Centre.

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