

N-ACYL-2-METHYLENE- β -ALANINE METHYL ESTERS FROM THE SPONGE *FASCIOSPONGIA CAVERNOSA*

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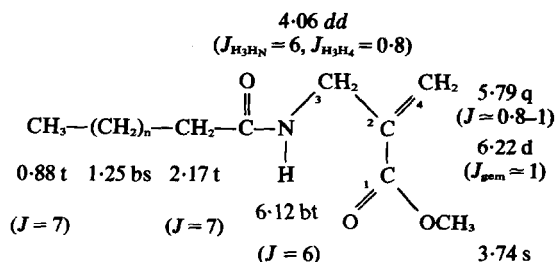
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Abstract—The isolation and structural determination of a series of N-acylated 2-methylene- β -alanine methyl esters, a new naturally-occurring β -amino acid, from the sponge *Fasciospongia cavernosa*, is reported.

In the course of our investigation of natural products derived from primitive multicellular animals living in the Red Sea, we undertook the determination of the chemical compounds found in several sponges.

Fasciospongia cavernosa was found to contain a new naturally-occurring β -amino acid (1), which appears in the sponge in a complete series of N-acylated methyl esters, in remarkably high concentrations (between 1–2%, dry weight).

From a petrol-ether extract of the fresh dry sponge, a white material (2, m.p. 67°) crystallized out directly after standing at 5° for 24–48 hr. A second closely-related oily fraction (3) *vide infra* was obtained in a pure form only after chromatography on a silica-gel column. Compound 2 showed the presence of three functional groups in the IR spectrum: an amide ($\nu = 3260$ (bonded NH), 1640, 1550), a conjugated terminal methylene ($\nu = 940$) and a conjugated esteric group (1715, 1230 cm^{-1}). The location of these groups relative to each other in the molecule could be estimated from the following NMR spectrum data, as well as from the mass spectrum data (Scheme 1), suggesting the N-acyl-2-methylene- β -alanine methyl ester structure, shown below, for 2:



In this NMR spectrum, the protons vicinal to the C-3-H (the latter giving rise to the signal at 4.06 dd ($J = 6$ and 0.8 Hz)) were determined by a double irradiation experiment or deuterium exchange of the NH-proton, cancelling the 6 Hz splitting; and by double irradiation of one of the terminal methylenic protons resonating at 5.79 d ($J = 0.8$ Hz). (The deuterium exchange of the amide proton could be achieved only at a relatively slow rate by DCl/D₂O or NaOD/D₂O solution, possibly because of an internal H-bond). Further evidence for the amino acid moiety of 2 could be obtained from its dihydro derivative 4, m.p. 55° (CH₃CN), which exhibited the following NMR spectrum (Fig 1). In this spectrum the various coupling constants were confirmed by double resonance experiments and by deuterium exchange of the NH, as in the case of 2.

According to the elemental analysis and mass spectrum (Scheme 1), it was clear that 2 was a mixture of at least five N-acyl derivatives of 1, that is, five amides of 1 with C₁₆-C₂₀ aliphatic fatty acids. In order to identify these acids, 2 was refluxed for 48 hr in saturated methanolic HCl solution until the methanolysis of the amide was complete, giving the free fatty acid methyl esters and 2-methylene- β -alanine methyl ester (5).

The five main fatty acid methyl esters were separated by gas chromatography and identified by their mass spectra, which were obtained by direct introduction of the compounds into the mass spectrometer. The compounds were found to be the C₁₆-C₂₀ normal unbranched aliphatic carboxylic acids.

Compound 5 was characterized as the hydrochloride C₅H₉O₂N·HCl, m.p. 93–95°. ν_{max} 3400, 2950, 2600, 1720, 980, 810 cm^{-1} ; NMR (D₂O): 4.00 s (OCH₃); 4.06 s (N-CH₂); 6.29 bs and 6.73 bs

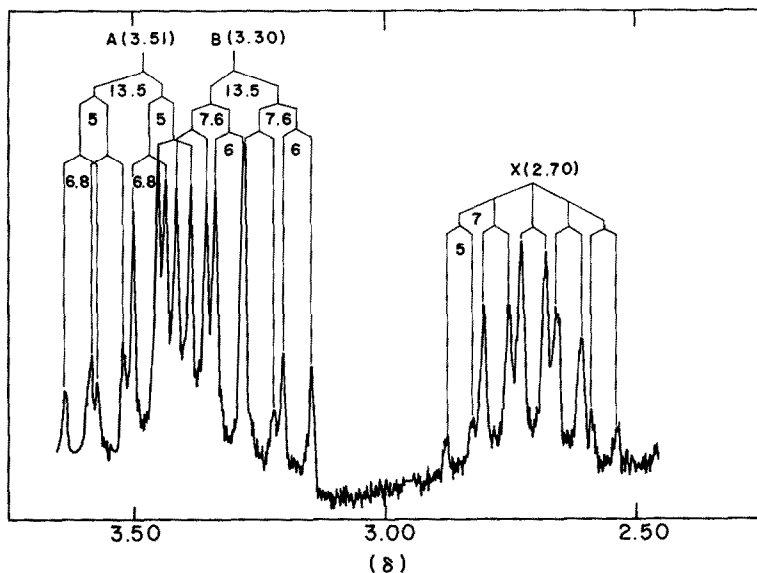
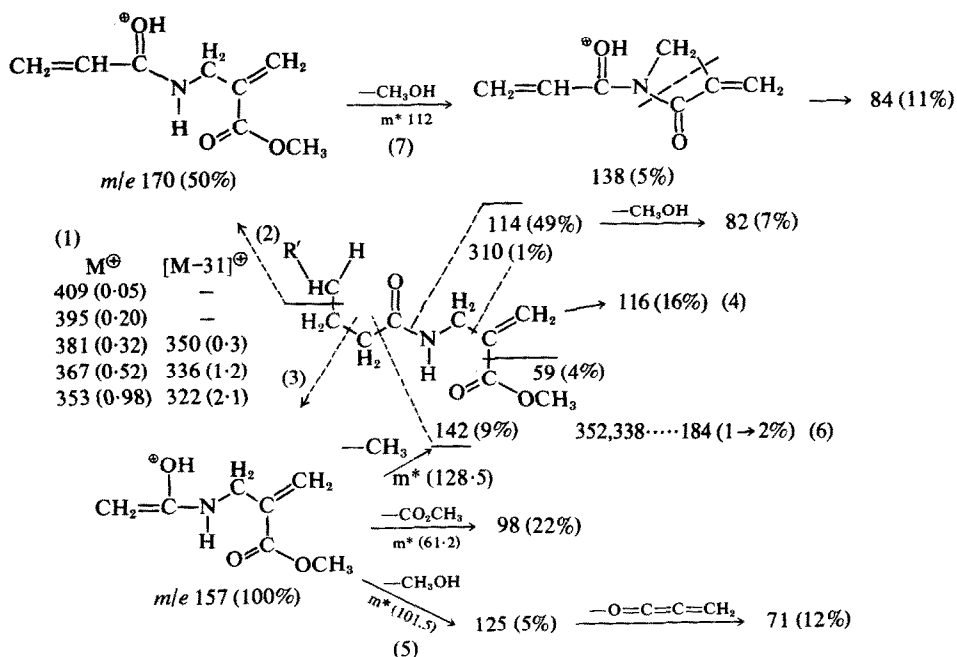


Fig 1. Partial NMR spectrum of compound 4.



SCHEME 1. Mass spectrum of 2.

- (1) The $(M-OCH_3)$ fragments of C_{20} and C_{19} are not strong enough to be seen.
- (2) Fragmentation by a mechanism similar to that known for long-chain methyl esters.¹
- (3) A "McLafferty rearrangement".
- (4) Probably the protonated amino acid methyl ester.
- (5) The metastable peak was observable only in the dihydro compound.
- (6) A series of fragments originating from the various esters by known fragmentations of long-chain methyl esters.
- (7) Deuterium exchange (NH to ND) established this fragmentation.
- (8) In the mass spectrum of 4 and N-d, 2, the expected shifts according to the proposed pattern are seen.

