

produce triplets¹⁵ (which could presumably sensitize singlet oxygen) and also singlet oxygen directly.¹⁶ All attempts to observe, via flash spectroscopy, triplets or singlet oxygen production by trapping with β -carotene in ¹DCA*-sensitized photooxidations have thus far proved unsuccessful.¹⁷

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(17) L. T. Spada, D. S. Steichen, and C. S. Foote, unpublished results. Unfortunately, β -carotene and other singlet oxygen quenchers are too rapidly destroyed in this system to be useful in product studies.

Structures of the Didemnins, Antiviral and Cytotoxic Depsipeptides from a Caribbean Tunicate¹

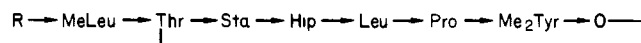
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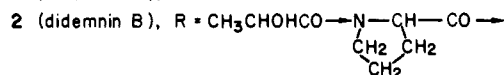
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We have recently discovered a class of depsipeptides in a Caribbean tunicate of the family Didemnidae (a species of the genus *Trididemnum*)² and have reported their isolation, separation, and biological properties elsewhere.³ In brief, the didemnins inhibit the growth of both RNA and DNA viruses, are highly cytotoxic to L1210 leukemic cells, and protect mice against P388 leukemia and B16 melanoma. For example, didemnin A shows 50% inhibition of Coxsackie virus and equine rhinovirus (both RNA) and *Herpes simplex*, type 2 (DNA), at 1.5 μ g/mL and *Herpes simplex*, type 1, at 3 μ g/mL, while didemnin B has ID₅₀ = 0.0011 μ g/mL vs. L1210 leukemic cells, T/C up to 199 vs. P388 leukemia, and T/C 160 vs. B16 melanoma. We report here the structures of these very promising new compounds 1-3, where Hip is hydroxyisovalerylpropionyl, O-CH[CH(CH₃)₂]COCH(CH₃)-CO.



1 (didemnin A). R = H



3 (didemnin C). R = CH₃CHOHCO \rightarrow

The didemnins were separated over silica gel and assigned molecular weights of 942 (didemnin A, major component), 1111 (B, minor component), and 1014 (C, trace component) from their field desorption (FD) mass spectra. Initial structural efforts centered on didemnin A, as the simplest and most abundant component. Didemnin A was concluded to be a peptide from its ¹H NMR spectrum, with broad doublets (NH) at 8.3, 7.8, and 7.5 ppm. Hydrolysis with 6 N hydrochloric acid gave 1 mol each of *N*-methylleucine (MeLeu), threonine (Thr), leucine (Leu), proline (Pro), *N,O*-dimethyltyrosine (Me₂Tyr),⁴ and statine (Sta).⁵

(1) Presented in part at the 3rd International Symposium on Marine Natural Products (International Union of Pure and Applied Chemistry and Societ  Chimique de Belgique), Brussels, Sept 16, 1980.

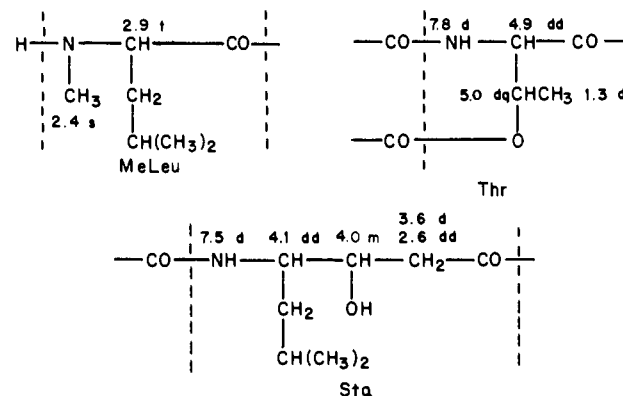
(2) Identification as family Didemnidae was carried out by Dr. Charles C. Lambert, California State College, Fullerton, and identification as a *Trididemnum* species by Dr. Fran oise Lafargue, Laboratoire Arago, Banyuls-sur-mer, France.

(3) K. L. Rinehart, Jr., J. B. Gloer, R. G. Hughes, Jr., H. E. Renis, J. P. McGovern, E. B. Swynenberg, D. A. Stringfellow, S. L. Kuentzel, and L. H. Li, *Science (Washington, D.C.)*, in press.

(4) F.-J. Marner, R. E. Moore, K. Hirotsu, and J. Clardy, *J. Org. Chem.*, **42**, 2815-2818 (1977).

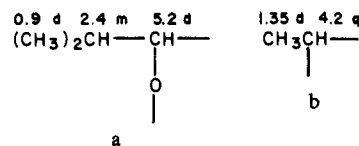
The amino acids were identified by FD mass spectrometry (FDMS) of the mixture, with M + H peaks at *m/z* 146 (MeLeu), 132 (Leu), 116 (Pro), and 210 (Me₂Tyr), an M - H₂O peak at *m/z* 101 for Thr, and M at *m/z* 175 for Sta. They were also identified by gas chromatography/mass spectrometry (GC/MS) of the amino acids' trifluoroacetyl *n*-butyl ester derivatives. In addition, they were quantitated by GC, and their identities confirmed by coinjection with derivatives of authentic samples. Statine was assigned as the allo (erythro) isomer by its coelution with the synthetic *R,S* isomer,⁵ while gas chromatography on an optically active column⁶ indicated Leu, MeLeu, Pro, Thr, and Me₂Tyr to have the L configuration.

The individual amino acids were characterized in the ¹H NMR spectrum of didemnin A by extensive spin decoupling, which established MeLeu as N-terminal, Thr as O- as well as N-acylated, and the hydroxyl group of Sta to be free, as shown. Didemnin

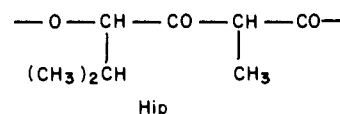


A reacts with acetic anhydride on the methylamino group of MeLeu and the hydroxyl group of Sta to give a diacetyl derivative, as judged by the product's molecular weight of 1026, two new COCH₃ groups at 1.9 and 2.1 ppm, and the shift of the appropriate protons to 2.8 (-N(Ac)-CH₃, MeLeu), 5.1 (AcNCH<, MeLeu) and 5.3 ppm (AcO-CH<, Sta).

Addition of the formulas for the amino acids' aminoacyl units (MeLeu, C₇H₁₄NO; Thr, C₄H₆NO₂; Sta, C₈H₁₅NO₂; Me₂Leu, C₁₁H₁₃NO₂; Pro, C₅H₇NO; Leu, C₆H₁₁NO) gave C₄₁H₆₆N₆O₉, 786.4891. The high resolution electron impact (HREI) mass spectrum of didemnin A gave a molecular ion peak at 942.5678. Subtraction of the two numbers yielded 156.0787 (C₈H₁₂O₃) as the unaccounted-for residue of didemnin A. The C₈H₁₂O₃ unit can be identified in the ¹H NMR spectrum as containing the units a and b and in the ¹³C NMR spectrum as containing one ketone



carbon (205.1 ppm) and one carboxyl-type carbon (one of seven near 175 ppm, in addition to the six of the amino acids). The only reasonable combination of a and b with a keto and a carboxyl carbon in a C₈H₁₂O₃ unit is as a hydroxyisovalerylpropionyl (Hip) group. All the carboxyl carbons occur as amides or esters since didemnin A does not react with diazomethane.



The order of linkage of these seven units was assigned by partial hydrolysis of didemnin A in base and acid, followed by HRFDMS

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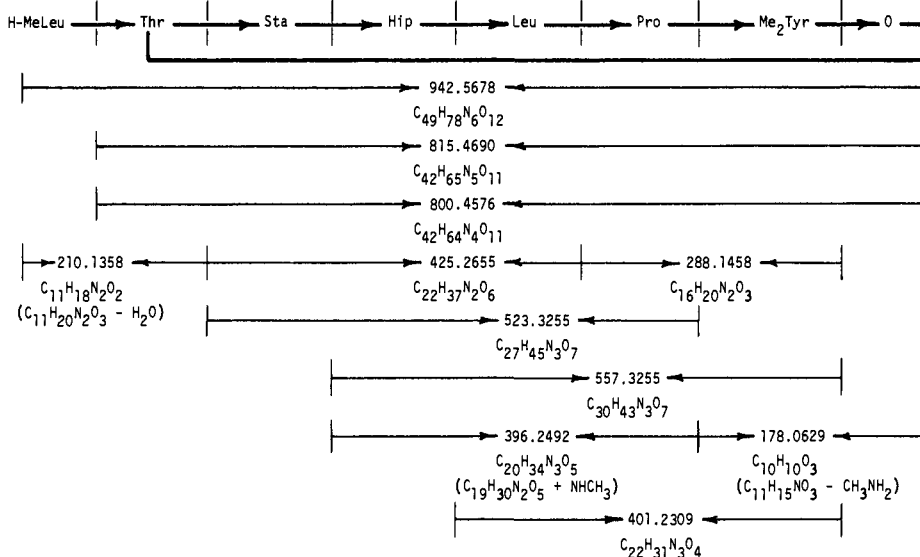
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Table I. HRFD Mass Spectral Peaks in Hydrolyzates of Didemnin A

ion	composition	assignment
576.3279 ^a	C ₃₀ H ₄₆ N ₃ O ₈	H-Hip→Leu→Pro→Me ₂ Tyr-OH + H
557.3094 ^{a-c}	C ₃₀ H ₄₃ N ₃ O ₇	H-Hip→Leu→Pro→Me ₂ Tyr-OH - H ₂ O
539.2987 ^a	C ₃₀ H ₄₁ N ₃ O ₆	H-Hip→Leu→Pro→Me ₂ Tyr-OH - 2H ₂ O
420.2492 ^{a,b}	C ₂₂ H ₃₄ N ₃ O ₅	H-Leu→Pro→Me ₂ Tyr-OH + H
386.2646 ^{b,d}	C ₁₉ H ₃₆ N ₃ O ₅	H-MeLeu→Thr→Sta-OH - H ₂ O + H
383.2214 ^{a,b}	C ₂₂ H ₂₉ N ₃ O ₅	H-Leu→Pro→Me ₂ Tyr-OH - 2H ₂ O
288.1467 ^{a-c}	C ₁₆ H ₂₀ N ₂ O ₃	[-Pro→Me ₂ Tyr-]
178.0631 ^a	C ₁₀ H ₁₀ O ₃	CH ₃ OC ₆ H ₄ CH=CHCOOH
156.0794 ^{a,b}	C ₈ H ₁₂ O ₃	Hip

^a 0.1 N NaOH, room temperature, 2 h. ^b 6 N HCl, 110 °C, 10 min. ^c Trifluoroacetic acid, room temperature, 48 h. ^d Accompanied by peaks at *m/z* 404^b and 368.^b

Scheme I. Fragment Ions Observed in the High Resolution Electron Impact Mass Spectrum of Didemnin A

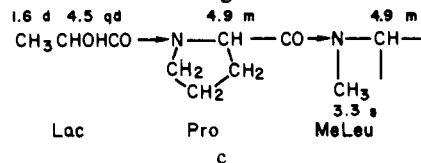


on the molecular ions of the resultant mixtures (Table I). The major peptide peaks observed due to the two lactone cleavages were C₃₀H₄₅N₃O₈ (which must contain Me₂Tyr, Hip, Pro, and Leu from its degree of unsaturation and oxygen content) and C₁₉H₃₆N₃O₅ (which must then contain MeLeu, Thr, and Sta), which together account for all the component acids of didemnin A. Smaller peptides (e.g., C₂₂H₃₄N₃O₅, Me₂Tyr, Pro, Leu; C₁₆H₂₀N₂O₃, Me₂Tyr, Pro) were also observed. On the basis of these data, Hip, the hydroxy acid, must be N-terminal in the C₃₀ peptide, while Leu could be C-terminal or between Hip and the C₁₆ dipeptide.

These peptide linkages are confirmed in the HREI mass spectrum of didemnin A (Scheme I), which contains fragment ions corresponding to the C₃₀ (Hip, Leu, Pro, Me₂Tyr), C₂₂ (Leu, Pro, Me₂Tyr), and C₁₆ (Pro, Me₂Tyr) peptides. Sequencing of the C₃₀ peptide is completed by the assignment of the *p*-methoxycinnamic acid ion (C₁₀H₁₀O₃) at *m/z* 178.0629, which shows the Me₂Tyr group to be linked as an ester, hence C-terminal in the C₃₀ peptide, and confirmed by the Hip→Leu→Pro-NHCH₃ ion at *m/z* 396.2492. Similarly, the ion at 425.2655 (C₂₂H₃₇N₂O₆), containing Sta in addition to Hip and Leu, shows Sta to be adjacent to Hip, hence C-terminal in the C₁₉ peptide. The ion at *m/z* 800.4576 (M - C₇H₁₄N₂O) shows MeLeu is linked as an amide. Thus, the structure of didemnin A is assigned as 1.

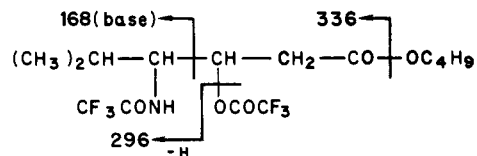
The molecular formula of didemnin B was established as C₅₇H₈₉N₇O₁₅, differing from didemnin A by only C₈H₁₁N₃O₃, by HRFDMS on the M + H ion at 1112.6442, while a major peak at *m/z* 942 in the mass spectra of B and the occurrence of nearly all of the EIMS peaks of didemnin A indicated a close similarity to the major component. Acidic hydrolysis of didemnin B to the component amino acids followed by GC of the *N*-trifluoroacetyl *n*-butyl ester derivatives indicated an additional mole of Pro, while decoupling studies of the ¹H NMR spectrum showed the unit c

as a modification of the N terminus of didemnin A. Thus, the structure of didemnin B is assigned as 2.



Similarly, the HRFD mass spectrum of the trace component, didemnin C, contains a molecular ion at *m/z* 1014.5873 (C₅₂H₈₂N₆O₁₄); its ¹H NMR spectrum contains a shifted MeLeu *N*-methyl group at 2.8 ppm; the GC/MS trace of the derivatized amino acids from its hydrolyzate is the same as that from didemnin A; and large peaks appear in the EI mass spectrum at *m/z* 172.1552 and 200.1283 [CH₃CHOHCON(CH₃)CH(CH₂-*i*-C₃H₇)R⁺, R = absent and CO]. Although lack of adequate sample has prevented more extensive studies, its structure is tentatively assigned as 3.

Each of the didemnins (A-C) is accompanied by a minor but varying amount of a lower homologue. The nature of the homology was clarified by the observation that GC traces of the derivatized amino acids always contain minor amounts of an amino acid identified by GC/MS as a lower homologue of statine, for which we propose the name norstatine [(CH₃)₂CHCHNH₂CH(OH)CH₂COOH] and whose derivative (4) fragments in the



manner shown. The lower homologous peptides can then properly be designated as nordidemins A-C, in which norstatine replaces the statine unit of 1-3.

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Regioselectivity in the Intramolecular Ene Reaction of Cyclopropene Derivatives

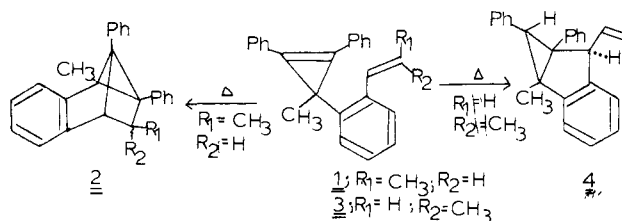
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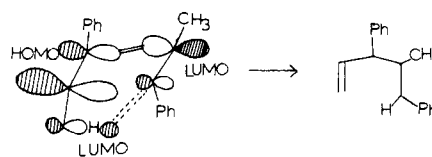
Received December 1, 1980

The thermal addition of an alkene to another olefin possessing an allylic hydrogen, the so-called "ene" reaction, is one of the most simple and versatile reactions of organic chemistry.¹ Though radical² and other mechanisms³ have been advanced, the addition is usually considered to proceed in a symmetry-allowed concerted process⁴⁻⁸ involving a six-membered cyclic transition state, unless prohibited by steric factors.⁹ Olefins with strained double bonds seem particularly prone to enter into ene reactions. For example, cyclopropene derivatives are known to undergo ready dimerization via the ene process.^{10,11} Considerable interest has recently been focused on intramolecular examples of this reaction.¹² In this communication, we wish to describe a novel regiochemical effect associated with the intramolecular ene reaction of tetrasubstituted cyclopropene derivatives.

As an extension of our studies dealing with intramolecular cycloaddition reactions of cyclopropene derivatives,¹³ we have examined the thermal and triplet sensitized behavior of a series of 3-(*o*-alkenylphenyl)-substituted cyclopropenes. Thermolysis of the trans-substituted cyclopropene **1** at 175 °C resulted in a [2 + 2] cycloaddition reaction to produce a 4.6:1 mixture of *exo* and endobenzotricycloheptene **2**.¹⁴ In marked contrast, heating

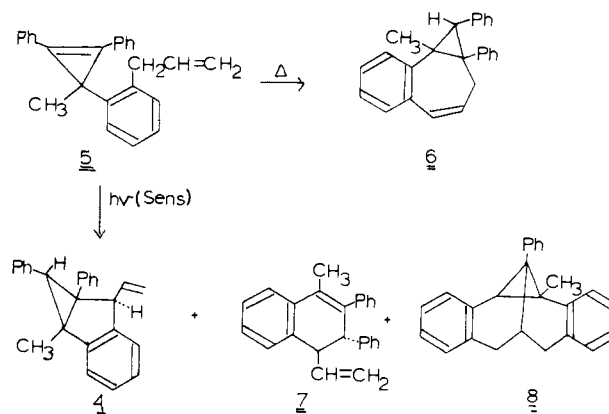


Scheme I



a sample of the isomeric *cis*-isomer **3** gave rise to benzobicyclohexane **4** in quantitative yield. We suggest that **4** most reasonably arises from **3** by a concerted ene reaction. The geometry necessary for this type of reaction is easily achieved with the *Z*-substituted cyclopropene. Although bimolecular ene reactions of cyclopropenes are known,^{10,11} the above case constitutes the first example of an intramolecular version of this reaction.

Additional examples which would establish the generality and scope of the intramolecular ene reaction were sought. With this in mind we investigated the thermolysis of the closely related 3-(*o*-allylphenyl)-substituted cyclopropene **5**. Heating a sample of **5** gave rise to the ene product **6** in quantitative yield.¹⁵ The



NMR spectrum (C_6D_6 , 90 MHz) of **6** consists of signals at δ 1.19 (s, 3 H), 2.14 (ddd, 1 H, $J = 13.3, 6.1$, and 2.0 Hz), 2.54 (dd, 1 H, $J = 13.3$ and 7.4 Hz), 2.62 (s, 1 H), 6.01 (ddd, 1 H, $J = 10.7, 7.4$, and 6.1 Hz), 6.62 (d, 1 H, $J = 10.7$ Hz), and 6.72-7.58 (m, 14 H). The sensitized irradiation of **5** was also studied and was found to produce a mixture of three products. Chromatography of the mixture on silica gel gave benzobicyclohexene **4**¹⁵ (60%) together with dibenzotricyclodecane **8** (30%) and dihydronaphthalene **7** (10%). Assignment of the minor component as dihydronaphthalene **7** was made on the basis of its NMR spectrum and by its oxidation to the corresponding aromatic hydrocarbon: NMR **7** (CDCl_3 , 90 MHz) δ 2.07 (s, 3 H), 3.55 (d, 1 H, $J = 7.5$ Hz), 3.79 (s, 1 H), 4.89-5.15 (m, 2 H), 6.21 (ddd, 1 H, $J = 17.5, 10.0$, and 7.5 Hz), 6.88-7.54 (m, 14 H). The structure of **8** was easily assigned on the basis of its straightforward spectral data:¹⁶ NMR (CDCl_3 , 90 MHz) δ 1.37 (s, 3 H), 2.22 (d, 1 H, $J = 15.5$ Hz), 2.37 (d, 1 H, $J = 15.5$ Hz), 2.60 (s, 1 H), 2.92 (t, 1 H, $J = 6.6$ Hz), 3.20 (dd, 1 H, $J = 15.5$ and 6.6 Hz), 3.42 (dd, 1 H, $J = 15.5$ and 6.6 Hz), and 6.73-7.49 (m, 13 H).

The triplet state of cyclopropene **5** can readily abstract a hydrogen from the neighboring benzylic carbon to produce a biradical intermediate which either collapses to give **4** or undergoes cyclopropyl ring opening in competition with coupling.¹⁷ The ring-opened species **10** would be expected to cyclize in a disrotatory fashion to give dihydronaphthalene **7**. The formation of dibenzotricyclodecane **8** in the sensitized irradiation represents an

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(15) The NMR spectrum of structures **4** and **6** suggest that the phenyl group is located in the *exo* position of the cyclopropyl ring. Satisfactory spectral and analytical data were obtained for each new compound.

(16) Attack of the ortho position of the triplet state of **5** on the adjacent double bond can occur in two ways. The NMR data obtained fits structure **8** better than the alternative regioisomer.

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