

Antineoplastic Agents 300. Isolation and Structure of the Rare Human Cancer Inhibitory Macrocyclic Lactones Spongistatins 8 and 9†

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Two trace (0.7 and 2.2×10^{-7} % yields) potent antineoplastic macrocyclic lactones termed spongistatins 8 (**2a**) and 9 (**2b**) have been isolated from the African marine sponge *Spirastrella spinispirulifera* and found to be very potent inhibitors of glutamate-induced tubulin polymerization.

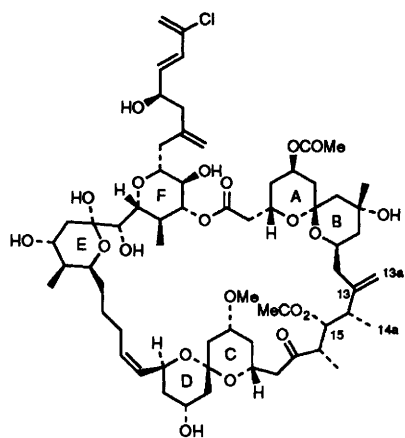
Since our early investigations,² the discovery of cytotoxic and/or antineoplastic marine animal constituents has continued to accelerate. Current examples include potentially useful substances from marine sponges,³⁻⁸ tunicates,⁹ soft coral,¹⁰ and starfish.¹¹ To date the astonishingly potent spongistatin 1 (**1**) discovered¹² in a Republic of Maldives, *Spongia sp.* has proved to be one of the most potent cancer cell growth inhibitory substances known. We later discovered spongistatins 2-7¹³⁻¹⁵ bearing variations in substituents and in the spongipyran ring system.† We now report the isolation and structural elucidation of two new and prodigiously cytotoxic macrocyclic lactones designated spongistatins 8 (**2a**) and 9 (**2b**) from the Southwest African marine sponge, *Spirastrella spinispirulifera*.

Further detailed study of P388 lymphocytic leukemia (PS cell line) active fractions prepared¹³ from *Spirastrella spinispirulifera* (2409 kg wet wt.) employing an extensive series of gel permeation and partition chromatographic separations on Sephadex LH-20 followed by HPLC (*e.g.* using reversed phase Prepex RP8, 5-20 μ , columns and water-acetonitrile gradients) afforded 1.8 mg (7.5×10^{-8} % yield, PS ED₅₀ 8×10^{-4} μ g ml⁻¹) of colourless spongistatin 8 (**2a**): mp 158-159°C; $[\alpha]_D^{22}$ -32° (*c* 0.18, MeOH); IR (film) cm⁻¹ 3439, 2936, 1736, 1653, 1602, 1383, 1252, 1178, 1090; high resolution FABMS, *m/z* 1183.5821 [M + K]⁺, C₆₁H₃₂KO₂₀ (calc. 1183.5819) and 5.4 mg (2.2×10^{-7} % yield, PS ED₅₀ 2.7×10^{-5} μ g ml⁻¹) of spongistatin 9 (**2b**): mp 164-165°C; $[\alpha]_D^{22}$ -33.3° (*c* 0.14, MeOH); IR (film) cm⁻¹ 3435, 2940, 1736, 1647, 1591, 1385, 1254, 1178, 1090; high resolution FABMS, *m/z* 1217.5425 [M + K]⁺ corresponding to C₆₁H₃₁ClKO₂₀ (calc. 1217.5429).

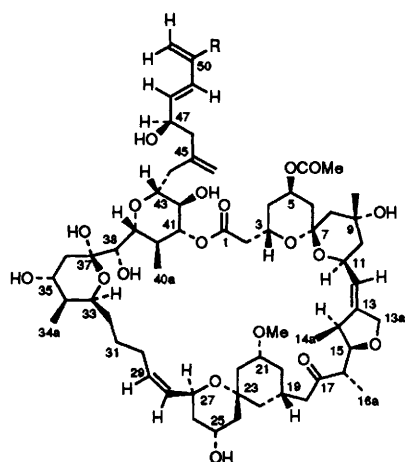
Due to the paucity of spongistatin 8 (**2a**) structural elucidation was simplified by first deducing the structure of spongistatin 9 (**2b**). Once the high resolution FABMS and high field 2D NMR interpretations were in hand for lactone **2b**, the structure of spongistatin 8 (**2a**) was completed. The THF ring of spongistatin 8 was recognized by chemical shifts at δ 4.45 (br d, *J* = 13 Hz), 4.10 (br d, *J* = 13 Hz)/70.70 and 1.95 (acetyl, s, 3H)/21.31 and 172.56 (acetyl). A signal at δ 6.33 (ddd, *J* = 10, 10, 17 Hz) indicated that C-50 was devoid of the usual spongistatin chlorine atom at that position. A series of ¹H-¹H NMR COSY experiments allowed assignment of the remaining ¹H and ¹³C NMR signals by comparison with the analogous NMR carbon data from spongistatin 9.

The structure of spongistatin 9 was determined mainly by high field NMR spectroscopy utilizing results of ¹H-¹H COSY, ¹H-¹³C COSY, APT, and HMBC NMR experiments. Both the ¹H- and the ¹³C-NMR spectra of spongistatin 9 indicated it was a member of the spongistatins by signals at δ 1.13 (3H, s)/30.17, 1.04 (3H, d)/14.59, 1.14 (3H, d)/15.10, 0.89 (3H, d)/11.52, 0.84 (3H, d)/13.00, a lactone carbonyl signal at δ 173.66 and a ketone carbonyl signal at δ 213.42. Spongistatin 9 was found to possess a THF ring by signals at δ 4.45 (br d, *J* = Hz) and 4.10 (br d, *J* = 13 Hz) corresponding to two H-13a. An acetyl group was evident by signals at δ 1.95 (3H)/21.35 and 172.61. That the acetyl group was attached to the C-5 oxygen atom was evidenced by the chemical shift of H-5 at δ 4.96. The two broad singlets at δ 5.42 and 5.33 and the lack of an ¹H signal for C-50 were indicative of a chlorine atom at that position.

Comparative testing of spongistatins **2a** and **2b** in the NCI 60 cell line *in vitro* screening panel,^{18,20} revealed an overall potency comparable to or exceeding the potency of the most active members of the series heretofore reported. The mean panel GI₅₀s, determined from quadruplicate testing of spongistatins **2a** and **2b** over three different concentration ranges (10⁻⁷, 10⁻⁸, 10⁻⁹ mol dm⁻³ upper limits; five, log₁₀-spaced



1, Spongistatin 1



2a, R = H Spongistatin 8
2b, R = Cl Spongistatin 9

concentrations in each range) were 2.3×10^{-10} and 0.4×10^{-10} mol dm⁻³, respectively (standard errors averaged less than 10% of the respective means). The distinctive mean-graph profiles (pattern of relative cellular sensitivity) produced by spongistatins **2a** and **2b** were strongly correlated (*e.g.* compare correlation coefficients^{19,20} >0.8) with the profiles obtained with other members of the series, as well as with structurally unrelated but mechanistically similar members of the important general class of tubulin-interactive antimicrotubule inhibitors.²⁰

Spongistatins **1** (**1**), **8** (**2a**) and **9** (**2b**) were found to potently inhibit the glutamate-induced polymerization of tubulin²¹ with IC₅₀ values respectively of 3.6, 5.5 and 4.2 μmol dm⁻³. By comparison the novel peptide dolastatin 10²² presumed to be the most potent inhibitor of tubulin assembly known, gave in the same series of experiments, an IC₅₀ of 2.1 μmol dm⁻³. While these spongistatins did not prevent the binding of colchicine to tubulin, they strongly inhibited the classic binding of vinblastine and nucleotide (GTP) to tubulin. Thus the spongistatins represent a completely new class of structurally unique biosynthetic products that inhibit mitosis by binding to tubulin in the Vinca alkaloid domain.²³ Very importantly, spongistatins **1** and **9** appear to be the most cancer cell growth inhibitory antimicrotubule substances discovered to date.

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Footnotes

† For the preceding contribution, refer to ref. 1.

‡ Subsequent to our first discovery¹² of spongistatin **1** and the novel spongipyran nucleus common to the series, others^{16,17} have reported structures corresponding to spongistatins **1** and **4** from different sponge sources. In order to avoid obfuscation of the literature by use of the names althohyrin (ref. 16) for spongistatin **1** (**1**) and cinachyrolide A (ref. 17) for spongistatin **4**, we recommend that the spongistatin nomenclature be used for all members of the spongipyran series.

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