Marine Natural Products

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Leiodermatolide, a Potent Antimitotic Macrolide from the Marine Sponge *Leiodermatium* sp.**

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Marine macrolides that selectively disrupt cell cycle events continue to occupy a central position as lead compounds in the ongoing search for novel anticancer agents, [1,2] highlighted by the recent FDA approval of Halaven (eribulin mesylate, a fully synthetic analogue of the halichondrins) for the treatment of advanced breast cancer.[3] Lithistid sponges have proven to be a particularly fertile source^[4] of such biologically relevant polyketide metabolites, including dictyostatin^[5] and discodermolide. [5d,6] As part of a continued program aimed at the discovery of novel bioactive natural products from deepwater marine invertebrates, we have examined the relatively unexplored^[7] lithistid sponge *Leiodermatium*. A crude extract of Leiodermatium sp. was found to exhibit substantial activity in an assay which identifies antimitotic agents through detection of phosphonucleolin, a marker of mitosis.^[8] Bioassay-guided fractionation led to the isolation of leiodermatolide (1, Figure 1), whose unprecedented 16-membered macrolide skeleton, featuring an unsaturated side chain terminating in a δ-lactone, has been elucidated through a combination of extensive NMR spectroscopic analysis, comparative DFT GIAO NMR shift calculations, and molecular modeling. Leiodermatolide was found to exhibit potent and selective antimitotic activity (IC $_{50}$ < 10 nm) against a range of human cancer cell lines by inducing G2/M cell cycle arrest, and represents a promising new lead for anticancer drug discovery.

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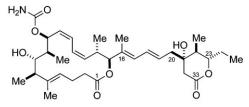
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1: Leiodermatolide

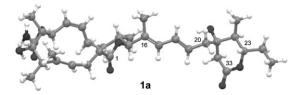


Figure 1. Structure of leiodermatolide (1) and energy-minimized conformation (1a). The absolute configurations of the C1–C16 macrolactone and C20–C33 δ -lactone regions are arbitrarily assigned.

A specimen of *Leiodermatium* (1.04 kg) was collected at a depth of 401 m off the coast of Fort Lauderdale, Florida, using the Johnson Sea Link submersible. The frozen sponge was exhaustively extracted using a mixture of EtOAc/EtOH (9:1) followed by partitioning of the dried extract residue between EtOAc and H_2O . Vacuum column chromatography of the organic partition on a silica gel stationary phase followed by reverse-phase HPLC of the active fraction led to the isolation of leiodermatolide (11.8 mg, 0.0011 % wet weight) as an amorphous white powder with $[\alpha]_D^{24} = -84.2$ (c = 0.34, MeOH). [9]

HRDART MS analysis of leiodermatolide (1) provided a $[M+H]^+$ ion at m/z 602.3705, appropriate for a molecular formula of $C_{34}H_{51}NO_8$ (calcd for [M+H], 602.3693, $\Delta=1.2$ mmu), requiring 10 sites of unsaturation. Optimum NMR signal dispersion was realized in CD_2Cl_2 ; inspection of the ^{13}C NMR spectrum revealed a total of 34 resonances in agreement with the HRMS formula. NMR resonances attributable to two ester groups ($\delta_C=172.4$, 170.4 ppm), one carbamate ($\delta_C=157.6$ ppm), and 10 olefinic carbons ($\delta_C=137.9$, 137.5, 134.2, 131.8, 130.0, 128.8, 128.5, 126.4, 125.9, 124.7 ppm) accounted for eight of the 10 sites of unsaturation, suggesting the presence of two rings.

Interpretation of the 2D DQF-COSY spectrum coupled with the edited g-HSQC spectrum led to the assignment of seven isolated spin systems (A–G, Figure 2). These spin systems and their connectivity with the remaining atoms enabled assembly into the final planar structure 1b based

Communications

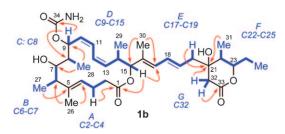


Figure 2. Isolated 1H spin systems (blue) and $^{2,3}J_{H,C}$ HMBC NMR correlations (red) leading to the planar structure $1\,b$ of leiodermatolide

upon analysis of 2J and 3J $^1H^{-13}C$ couplings observed in the g-HMBC spectrum. Accordingly, HMBC correlations between a methyl resonance ($\delta_{\rm H}$ =1.39 ppm, s, H26) and carbons C4, C5, and C6 served to place it on C5 as well as link spin systems A and B through C5. This assignment was supported by further correlations from H4 to C6/C26, and H6 to C4/C5/C26. Further correlations between H2ab and H3ab with the ester carbon observed at $\delta_{\rm C}$ =172.4 ppm allowed for C1 to be incorporated into the molecular structure, which could be linked to H15 and thus spin system D through an HMBC correlation from H15 to C1.

Unification of spin systems B, C, and D was initially hampered by the lack of scalar coupling between H8 and either H7 or H9, although this observation was subsequently exploited for purposes of stereochemical determination (see below). Spin system C thus appeared as an isolated, mutually coupled CHCH₃ system [$\delta_H = 1.72$ ppm (br q, ${}^3J = 7.2$ Hz) and 1.05 ppm (d, ${}^{3}J = 7.2 \text{ Hz}$)]. Ultimately, HMBC correlations between H28 and both C9 and C7 clearly placed C8 between these two atoms, supported by correlations from H9 to C7 and H7 to C9/C28. A carbamate was attached to C9 to account for its chemical shift ($\delta_{\rm C}$ =68.0 ppm), based on an HMBC correlation from H9 to C34 ($\delta_{\rm C}$ =157.6 ppm). The oxygenation implied by the chemical shift of C7 ($\delta_C = 78.4$ ppm) was confirmed by acetylation whereby the H7 resonance shifted from $\delta_{\rm H}$ = 3.24 to 4.88 ppm. [10] Analysis of the DQF-COSY spectrum was then sufficient to define spin system D containing H15, already related to A, thus completing the 16-membered macrocyclic core of leiodermatolide.

Correlations in the HMBC spectrum between a methyl resonance ($\delta_{\rm H}$ = 1.76 ppm, s, H30) and carbons C15, C16, and C17 served to place it on the olefinic carbon C16 as well as link spin systems D and E through C16. This assignment was supported by correlations from H15 to C17/C30, H17 to C15/ C30, and H18 to C16. The final spin systems F and G could then be connected to E through the quaternary carbon observed at $\delta_C = 72.3$ ppm (C21), whose chemical shift could be accounted for by oxygenation. HMBC correlations from C21 to H19/H20ab/H31 linked E to F, while correlations from H32ab to C21 as well as to C20 and C22 then established the C21-C32 bond. HMBC correlations from H32ab to the carbonyl carbon observed at $\delta_C = 170.4 \text{ ppm}$ (C33) allowed for incorporation of the second ester carbon into the structure, which could be connected as a δ-lactone onto C23 on the basis of a weak HMBC correlation from H23 to C33, accounting for the remaining site of unsaturation.

Leiodermatolide thus features a 16-membered macrolactone ring with an unsaturated side chain terminating in a δ -lactone ring, which together incorporate five alkenes and nine stereocenters, including a carbamate at C9 and two hydroxy groups at C7 and C21. The stereochemistry is segregated into two distinct C1–C16 macrocyclic and C20–C33 δ -lactone clusters, whose spatial isolation means that NMR methods alone are insufficient to unambigously define their relative configuration, which must likely await future synthetic studies.

Determination of stereochemical configuration within the macrolide core utilized a combination of homo- $({}^3J_{\rm H,H})$ and heteronuclear $({}^{2,3}J_{\rm C,H})^{[11]}$ *J*-based configurational analysis^[12] and extensive nOe evidence. Figure 3 depicts the results of

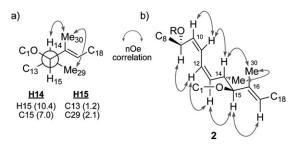


Figure 3. NMR analysis for the C9–C17 backbone of leiodermatolide. a) Rotamer and *anti* configuration determined for C14–C15; ${}^3J_{\text{H,H}}$ and ${}^{2,3}J_{\text{H,C}}$ [Hz] in parentheses. b) Key nOe contacts for C9–C17.

NMR experiments for the C9-C17 segment of leiodermatolide. Coupling constant analysis suggested H14 and H15 should be antiperiplanar [${}^{3}J(H14,H15) = 10.4 \text{ Hz}$], which together with nOe contacts between H14/H29 and H30 suggested the arrangement shown in Figure 3a. Two series of strong nOe enhancements could then be traced along either edge of what was deduced to be a pseudo-planar C9-C17 backbone sequence, as shown in 2 (Figure 3b). The nOe contacts from H10 to H11 and H12 to H13 together with the 10.3 and 11.4 Hz coupling of the respective olefinic protons indicated the Δ -10 and Δ -12 double bonds should be Zconfigured, while the nOe correlation from H15 to H17 suggested the Δ -16 trisubstituted olefin to be *E*-configured. Key nOe enhancements from H9 to H12 and H11 to H14 indicated an S-trans arrangement about C11-C12, where A(1,3)-strain would be minimized by H9 and H14 eclipsing the C10-C11 and C12-C13 olefins, respectively. Together with the nOe observed for H14 to H30, this preferred conformation has important implications for assigning the relative configuration of the C6-C9 and C14-C15 stereoclusters; the C8 chain and C15 oxygenation must reside on the same face of the pseudo-planar C9-C17 system such that the macrocycle may be closed. As such, an S configuration at C9 would translate into an S,S configuration at C14, C15.

Having established the relative configuration between the C14, C15, and C9 stereocenters, attention was turned to assignment of the C6–C9 stereotetrad. As indicated above, vicinal ¹H–¹H coupling constant analysis revealed ³*J*(H7,H8) and ³*J*(H8,H9) to approximate to 0 Hz, suggesting the corresponding torsion angles should approach 90°. Hetero-



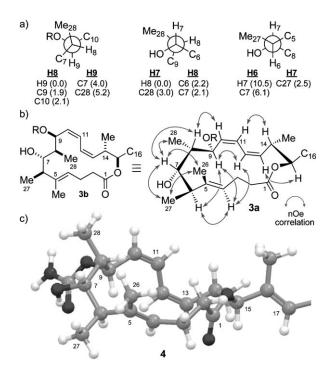


Figure 4. Configurational analysis for the C1–C16 macrocyclic region of leiodermatolide. a) Rotamers determined for C5–C10; $^3J_{\rm H,H}$ and $^{2,3}J_{\rm H,C}$ [Hz] in parentheses. b) Selected key nOe contacts for the C1–C16 macrolactone region. c) Energy-minimized conformer **4** for stereostructure **3 b.**^[14,15]

nuclear *J*-based configurational analysis^[12] was then used to determine the corresponding rotamers shown in Figure 4a, where an HSQMBC NMR experiment proved crucial to extracting necessary data the HSQC-HECADE experiment failed to provide due to the lack of scalar ¹H coupling. ^[11] The relatively large coupling between H9 and C28 and small coupling between H8 and C9 supported a *gauche* relationship between the adjacent methyl and carbamate substituents and an *anti* relationship between C7 and C10. The small values for both ²J(H8,C7) and ³J(H7,C28) indicated the mutually *gauche* arrangement shown. In the absence of an nOe contact between H6 and H7, the large vicinal coupling constant [³J(H6,H7) = 10.3 Hz] indicated an antiperiplanar relationship.

A series of nOe experiments were then carried out for the C4–C9 sequence (3a, Figure 4b), which served to define the exact rotamer for C5–C8 and consolidate the foregoing configurational analysis. A strong nOe enhancement between H4 and H6 indicated the Δ-4 double bond should be *E*-configured. The absence of nOe contacts between H27 and H28 suggested they should be 1,3-anti configured such that *syn*-pentane-type interactions in the macrocycle would be avoided. Particularly diagnostic nOe's were observed from H6 to H9 and H7 to H8, while only a weak enhancement was observed from H8 to H9, and none for H7 to H9, which together suggested that H6/H9 and H7/H8 should reside on opposite faces of the C6–C9 system. Further nOe enhancements from H8 to H10 and H7 to H26 related the orientation of this sequence to the adjacent double bonds, which was

reinforced by additional transannular nOe contacts from H4 to H9 and H12.

The combination of this data, the preferred conformation of the C9–C17 backbone derived in Figure 3 and the rotameric restrictions implied by the *J*-based analysis for the C5–C10 sequence, provided truncated macrocycle **3b** as the sole stereochemical assignment consistent with the observed NMR data. Notably, this arrangement allows for a hydrogenbonding interaction between the C7 hydroxy and C9 carbamate moieties, in agreement with the observed sharpening of ¹H resonances for the corresponding C7 acetate derivative. [10]

Molecular modeling^[14] provided the energy-minimized conformer **4** (Figure 4c) for the macrolactone structure **3b**,^[15] which appears wholly consistent with the experimental NMR data (*J*-values and nOe contacts). Notably, none of the higher-energy conformers within 10 kJ mol⁻¹ exhibited significant conformational variation about the macrolactone region (only minor flexibility in the C2–C3 region), thus supporting the validity of the nOe analysis. Similar modeling of other configurational permutations revealed in each case one or more diagnostic discrepancies with the NMR data.^[16]

Figure 5 depicts the C17–C33 side chain stereocluster of leiodermatolide, incorporating the δ -lactone ring. Vicinal ¹H–
¹H coupling constant analysis indicated the Δ -18 olefin to be *E*-configured [³J(H18,H19) = 15.1 Hz], while H22 and H23

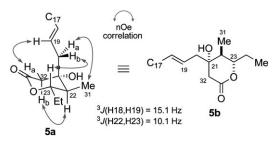


Figure 5. NMR analysis for the C17–C33 δ -lactone region of leiodermatolide, showing selected key nOe contacts.

should adopt a diaxial relationship $[^3J(\text{H22},\text{H23})=10.1~\text{Hz}]$. A series of nOe experiments again proved important in determining stereochemical relationships. In this case, an nOe contact from H23 to H20b correlated with further contacts for the olefinic H19 proton with H32a, and H20a with H31 to indicate the pseudo-axial nature of the appended C20 side chain. This was reinforced by an nOe observed between H22 and H32b, indicative of their 1,3-diaxial relationship. These data were supportive of a preferred chair-type conformation $\bf 5a$ and provided the relative δ -lactone stereostructure shown in $\bf 5b$.

With a convincing stereochemical assignment for the 16-membered macrolactone and δ -lactone regions of leiodermatolide reached by NMR spectroscopic analysis, we sought to further reinforce our findings by computational methods. Using the recently developed DP4 NMR prediction methodology of Smith and Goodman, [17a] computational DFT GIAO NMR shift comparisons of experimentally measured 1H and ^{13}C NMR resonances for leiodermatolide and those calcu-

Communications

lated for the 32 possible macrocyclic and four possible δ lactone diastereomers in isolation were carried out.[10] This analysis independently predicted a single diastereomer with > 99% probability for the macrocyclic region, with the remaining 31 stereochemical permutations being of negligible probability. Reassuringly, the predicted (6*R*,7*S*,8*S*,9*S*,14*S*,15*S*) diastereomer proved to be an exact match for that determined by our NMR analysis (3b, Figure 4b). Further DP4 computational ¹H and ¹³C NMR analysis also suggested the same δstereocluster 5 b, determined in Figure 5 lactone (21S,22S,23S), to be that for leiodermatolide with > 99%probability. Notably, this is the first application of the DP4 method to predict the stereochemistry of a complex macrolide from analysis of experimental and calculated NMR shift

The relative configurations within the macrolactone (C1-C16) and δ-lactone (C20-C33) stereoclusters had thus been determined as that shown in 1 (Figure 1), where the relationship between these regions is arbitrarily assigned. In order to reduce the remaining stereochemical permutations, we attempted to define the absolute configuration of the macrolide core through preparation and ¹H NMR analysis of the corresponding C7-OH (R)- and (S)-MTPA ester derivatives (MTPA = Mosher's acid = α -methoxy- α -trifluoromethylphenylacetic acid).[18] Whilst two distinct bis-Mosher ester derivatives were able to be prepared, NMR analysis proved inconclusive, with irregular $\Delta \delta^{SR}$ values measured either side of the C7 stereocenter. [10] Surprisingly, esterification occurred initially at C21, indicative of a highly sterically congested environment about C7, which seems likely to underlie the failure of the Mosher ester analysis itself.^[19] Determination of the full configuration will thus likely rely on the stereocontrolled synthesis of both possible diastereomers, followed by detailed NMR and chiroptical comparisons with the natural product.

Leiodermatolide exhibited potent antimitotic activity, and strongly inhibited in vitro cell proliferation in several cancer cell lines, including the human A549 lung adenocarcinoma, PANC-1 pancreatic carcinoma, DLD-1 colorectal carcinoma, NCI/ADR-Res ovarian adenocarcinoma, and P388 murine leukaemia, with IC₅₀ values of 3.3 nm, 5.0 nm, 8.3 nm, 233 nm, and 3.3 nm, respectively. Significantly reduced antiproliferative effects against the Vero monkey kidney cell line (IC₅₀ 211 nm) suggest that leiodermatolide may also possess useful selectivity for cancer cells. Cell cycle analysis in the A549 and PANC-1 cell lines confirmed that cell cycle arrest occurred at the G2/M transition. [10] Given that many compounds that induce G2/M cell cycle arrest do so through interaction with the cytoskeletal protein tubulin, the effects on microtubule structure were examined using confocal microscopy. Whilst leiodermatolide proved to have minimal effects on interphase cells, dramatic effects were observed on spindle formation in mitotic cells at concentrations as low as 10 nm (Figure 6). However, leiodermatolide neither induced nor inhibited assembly of purified tubulin in vitro. Thus, whilst the exact mechanism of action presently remains undefined, it is clearly distinct from that of other important antimitotic agents such as the epothilones, discodermolide, taxanes, and Vinca alkaloids. $^{[2,20,21]}$

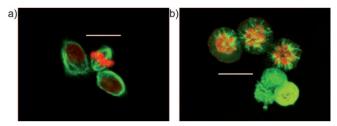


Figure 6. Immunofluorescence images of PANC-1 cells stained with anti-α-tubulin (green) and propidium iodide (red) and observed with confocal microscopy. Cells were exposed to methanol (control) or 10 nm leiodermatolide. a) Control cells display normal mitotic spindles. b) Leiodermatolide treated cells show abnormal spindle formation. Scale bars = 20 μm.

In summary, leiodermatolide (1) is a structurally unique polyketide-derived macrolide isolated from the deep-water marine sponge *Leiodermatium* sp., whose potentially novel mode of antimitotic action makes it an exciting new lead for anticancer drug discovery. Its scarce natural abundance also makes it a prime target for realizing a practical synthesis. Studies towards this end, including resolution of the remaining configurational issues, will be reported in due course.

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3223