Calyciphyllines A and B, Two Novel Hexacyclic Alkaloids from *Daphniphyllum calycinum*

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ABSTRACT

Two types of *Daphniphyllum* **alkaloids with unprecedented fused-hexacyclic ring systems, calyciphyllines A (1) and B (2), have been isolated from the leaves of** *Daphniphyllum calycinum* **(Daphniphyllaceae), and the structures and relative stereochemistry were elucidated on the basis of spectroscopic data.**

Plants of the genus *Daphniphyllum* are known to produce structurally diverse alkaloids with unusual polycyclic skeletons.1,2 These unique ring systems have attracted great interest as challenging targets for total synthesis as well as biosynthetic studies.³ Heathcock and co-workers have proposed a biogenetic pathway for *Daphniphyllum* alkaloids and demonstrated a biomimetic total synthesis of several *Daphniphyllum* alkaloids.3,4

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Recently, we have isolated some novel types of *Daph* $niphyllum$ alkaloids⁵⁻¹⁰ such as daphnezomines A and B^5 with a unique aza-adamantane core, daphnezomines F and

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 $G⁶$ with a 1-azabicyclo[5.2.2]undecane ring system, daphnicyclidins $A-H$ (see Scheme 1),⁸ J, and K^9 with a unique hexa- or pentacyclic ring system, daphmanidin A^{10} with an unprecedented fused-hexacyclic skeleton from the leaves and stems of *D. teijismanni* and/or *D. humile*, and daphniglaucin A11 with a fused-polycyclic skeleton containing a 1-azoniatetracyclo $[5.2.2.0^{1.6}0.^{4.9}]$ undecane ring and a quaternary nitrogen from the leaves of *D. glaucescens*. In our continuing search for structurally unique and biogenetically interesting *Daphniphyllum* alkaloids, calyciphyllines A (**1**) and B (**2**), two types of alkaloids with unprecedented fused-hexacyclic ring systems, were isolated from the leaves of *Daphniphyllum calycinum* (Daphniphyllaceae)*.* This paper describes the isolation and structural elucidation of **1** and **2**.

The leaves of *D. calycinum* were extracted with MeOH, and the extract was partitioned between EtOAc and 3% tartaric acid. Water-soluble materials, which were adjusted at pH 9 with saturated $Na₂CO₃$, were extracted with CHCl₃. CHCl3-soluble materials were subjected to an amino silica gel column (hexane/EtOAc, $9:1 \rightarrow 1:1$ and then CHCl₃/ MeOH, $1:0 \rightarrow 0:1$), from which a fraction eluted with CHCl₃/ MeOH $(7:3)$ was purified by an ODS column (MeOH/H₂O, $2:3 \rightarrow 1:0$) followed by LH-20 (CHCl₃/MeOH 1:1) and silica gel columns (CHCl₃/MeOH/EtOAc, 8:1:1) to afford calyciphyllines A^{12} (**1**, 0.001% yield) and B^{13} (**2**, 0.0001%).

Calyciphylline A (**1**) showed the pseudomolecular ion peak at m/z 386 (M + H)⁺ in the FABMS, and the molecular formula, $C_{23}H_{31}NO_4$, was established by HRFABMS $[m/z]$ 386.2346, $(M + H)^{+}$, $\Delta + 1.4$ mmu]. IR absorptions implied the presence of ester carbonyl and ketone (1735 and 1705 cm-¹ , respectively) functionalities. 13C NMR data (Table 1) revealed 23 carbon signals due to 1 tetrasubstituted olefin, 2 carbonyls, 2 sp³ quaternary carbons, 6 sp³ methines, 8 sp³ methylenes, 2 methyls, and 1 methoxy. Among them, 2 methylenes (δ _C 74.4, δ _H 3.21 and 3.35; δ _C 67.5, δ _H 3.03 and 3.56) and 1 methine (δ _C 89.6; δ _H 3.80) were ascribed to those bearing an oxidative nitrogen.¹⁴

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⁽¹²⁾ Calyciphylline A (1): colorless solid; $[\alpha]_D$ -51° (*c* 1.0, CH₃OH); IR (neat) *ν*_{max} 3390, 2930, 1735, 1705, 1650, 1455, and 1060 cm⁻¹; ¹H and ¹³C NMR data (Table 1); FABMS m/z 386 (M + H)⁺; HRFABMS *m*/*z* 386.2346 (M + H; calcd for C₂₃H₃₂NO₄, 386.2332).

⁽¹³⁾ Calyciphylline B (2): colorless solid; $[\alpha]_D$ -58° (*c* 0.4, CH₃OH); IR (KBr) v_{max} 3280, 2930, 1740, 1460, 1270, and 750 cm⁻¹; ¹H and ¹³C NMR data (Table 1); FABMS *^m*/*^z* 358 (M ⁺ H)+; HRFABMS *^m*/*^z* 358.2392 $(M + H;$ calcd for C₂₂H₃₂NO₃, 358.2382).

Figure 1. Selected two-dimensional NMR correlations for calyciphylline A (**1**).

The ¹H-¹H COSY and HOHAHA spectra revealed connectivities of three partial structures **a** (C-2 to C-4, C-2 to C-18, and C-18 to C-19 and C-20), **b** (C-6 to C-7 and C-12, and C-11 to C-12), and c (C-13-C-17) as shown in Figure 1. HMBC correlations were observed for H-19b to C-7 (δ _C 74.4) and H-7b and H-19a to C-4 (δ _C 89.6), suggesting that C-4, C-7, and C-19 were connected to each other through a nitrogen atom. The chemical shifts of C-4, C-7, and C-19 (δ _C 67.5), which resonated at a lower field than those of carbons bearing a nitrogen, $7,14$ and a fragment ion at m/z 370 (M $- 16$)⁺ in the FABMS spectrum indicated the presence of an *N*-oxide function. The connectivity of C-21 to C-4, C-6, and C-8 through C-5 was implied by HMBC correlations for H₃-21 to C-4, C-5 (δ _C 62.7), C-6 $(\delta_C 49.6)$, and C-8 ($\delta_C 52.8$). HMBC cross-peaks for H₂-11 and H₂-17 to C-10 (δ _C 140.9) indicated connectivities of units **b** and **c** through C-10. The presence of a ketone at C-1 was suggested by the HMBC correlation for H-2 to C-1 (δ_c) 215.9). The connectivity of C-1 and C-13 to C-9 through C-8 was implied by HMBC correlations for H_2 -13 to C-1,

Figure 2. Selected two-dimensional NMR correlations for calyciphylline B (**2**).

C-8, and C-9 (δ _C 141.1). In addition, the HMBC correlation for H-15 to C-9 indicated the connectivity of C-9 to C-15. A methoxy group was attached to C-22 by HMBC correlations for H₃-23 and H-14 to C-22 (δ _C 176.1). Thus, the gross structure of calyciphylline A was assigned as **1** having an unprecedented fused-hexacyclic ring system (three five-, two six-, and one seven-membered rings) containing an *N*-oxide group as shown in Figure 1.

The relative stereochemistry of **1** was deduced from NOESY correlations as shown in Figure 1. The NOESY correlation of H-3b/H-13a suggested that the cyclohexane ring (C-1∼C-5 and C-8) took a boat form.

Calyciphylline B (**2**) was shown to have the molecular formula of C22H31NO3 by HRFABMS [*m*/*^z* 358.2392, (M ⁺ H)⁺, Δ +1.0 mmu], which was smaller than that of 1 by a CO unit. The IR absorption implied the presence of a *δ*-lactone (1740 cm-¹) functionality. 13C NMR data (Table 1) revealed 22 carbon signals due to 1 trisubstituted olefin, 1 ester carbonyl, 2 sp³ quaternary carbons, 6 sp³ methines, 9 sp3 methylenes, and 2 methyls. Among them, 1 methylene $(\delta_C$ 73.7; δ_H 3.13 and 3.54) and 2 methines (δ_C 94.4; δ_H) 4.08; δ _C 68.8; δ _H 4.09) were ascribed to those bearing an *N*-oxide group,¹⁴ while 1 sp³ quaternary carbon (δ _C 87.5) corresponded to that bearing an oxygen. Since 2 out of 8 unsaturations were accounted for, **2** was inferred to possess

^{(14) &}lt;sup>13</sup>C NMR signals for sp^3 carbons bearing an *N*-oxide are shifted about 10 ppm downfield as compared with those bearing secondary and tertiary amines: Kashiwaba, N.; Ono, M.; Toda, J.; Suzuki, H.; Sano, T. *J. Nat. Prod.* **¹⁹⁹⁸**, *⁶¹*, 253-255.

Scheme 1. Plausible Biogenetic Path for Calyciphyllines A (**1**) and B (**2**)

6 rings. The gross structure of **2** was elucidated by analyses of two-dimensional NMR data, including ¹H⁻¹H COSY,
HOHAHA HMOC and HMBC spectra in CD-OD (Figure HOHAHA, HMQC, and HMBC spectra in CD₃OD (Figure 2).

Connectivities of $C-1-C-4$, and $C-2$ to $C-18$, and $C-18$ to C-19 and C-20 (unit **^a**), C-6 to C-7 and C-12, C-10- C-12 and C-10 to C-17, and C-15-C-17 (unit **^b**), and C-13 to C-14 (unit **c**) were revealed by the ${}^{1}H-{}^{1}H$ COSY and HOH and HOH and HOH and HOH and HOH HOHAHA spectra. The presence of a hexahydroindene moiety (unit **b**) was deduced from HMBC correlations for H-7 to C-9 and C-10, and H-15 to C-10. HMBC cross-peaks for H-1 and H₃-21 to C-5 and C-8, H-6 and H-7 to C-5, and $H₂$ -4 to C-8 indicated connections of C-1 and C-4 to C-8, and C-5 to C-7 and C-8, constructing an octahydroindolizine ring system fused to a cyclopentane ring. On the other hand, the C-14 methylene (δ_c 28.1; δ_H 2.52 and 2.63) was attached to a carbonyl group at C-22 (δ_c 174.6), which was connected to C-5 (*δ*^c 87.5) through an ester linkage to form a *δ*-lactone ring, since the HMBC correlation of H-1 to C-13 was observed. These results indicated that the *δ*-lactone ring was connected to the octahydroindolizine ring at C-5 and C-8. Thus, the gross structure of calyciphylline B was assigned as **2** having an unprecedented hexacyclic ring system consisting of a hexahydroindene ring and an octahydroindolizine ring fused to a cyclopentane ring with a *δ*-lactone ring at C-5 and C-8. The relative stereostructure of **1** was deduced from cross-peaks observed in the phase-sensitive NOESY spectrum as shown in computer-generated threedimensional drawing (Figure 2).

A plausible biogenetic pathway for calyciphyllines A (**1**) and B (**2**) is proposed as shown in Scheme 1. Calyciphylline A (**1**) might be generated from the yuzurimine-type alkaloids such as yuzurimine A^{15} and macrodaphniphyllamine¹⁶ through daphniglaucin A as follows.¹¹ Cleavage of the C-1-N-1 bond

of daphniglaucin A will give the skeleton of calyciphylline A (**1**), from which daphnicyclidin A may be generated through an intermediate as proposed previously.8 On the other hand, the biogenetic origin of calyciphylline B (**2**) seems to be imine intermediate **C**, which might be produced through fragmentation reaction of the secodaphniphylline-type skeleton (**B**) derived from imine intermediate **A** proposed by Heathcock et al*.* ³ Calyciphylline B (**2**) might be generated from attack of the carbonyl group to C-5 of the intermediate **^C** and cleavage of C-4-C-5 and C-8-C-9 bonds followed by C-7-C-9 bond formation. Stereochemistry at C-6 was believed to epimerize through enamine formation during these backbone rearrangements.

Calyciphyllines A (**1**) and B (**2**) exhibited cytotoxicity against murine lymphoma L1210 cells (IC₅₀, 2.1 and 4.2 μ g/ mL, respectively) in vitro.

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Supporting Information Available: One- and twodimensional NMR spectra for compounds **1** and **2**. This material is available free of charge via the Internet at http://pubs.acs.org.

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