Samoquasine A, a Benzoquinazoline Alkaloid from the Seeds of *Annona* squamosa

Hiroshi Morita,[†] Yumiko Sato,[†] Kit-Lam Chan,[‡] Chee-Yan Choo,[‡] Hideji Itokawa,[§] Koichi Takeya,[§] and Jun'ichi Kobayashi^{*,†}

Graduate School of Pharmaceutical Sciences, Hokkaido University, Sapporo 060-0812, Japan, School of Pharmaceutical Sciences, Universiti Sains Malaysia, 11800 Penang, Malaysia, and School of Pharmacy, Tokyo University of Pharmacy & Life Science, 1432-1 Horinouchi, Hachioji, Tokyo 192-03, Japan

Received July 10, 2000

A benzooxyquinazoline alkaloid, samoquasine A (1), has been isolated from the seeds of *Annona squamosa*, and its structure was elucidated by spectroscopic and chemical methods.

Annona squamosa L. (Annonaceae), commonly known as custard apple is a tropical fruit tree distributed mainly in the Americas and in southeast Asia. The seeds are wellknown to contain both many acetogenins with antifeedant, antimalarial, cytotoxic, and immunosuppressive activities and waxy substances containing long-chain fatty acids.¹ On the other hand, the leaves and wood are known to be a rich source of aporphine alkaloids.² Recently, we have isolated seven new cyclic peptides, cyclosquamosins A–G, from the seeds of *A. squamosa* collected in Malaysia.³ Further investigation on extracts of the seeds of the same plant resulted in isolation of samoquasine A (1), a benzooxyquinazoline alkaloid. Herein we describe the isolation and structure elucidation of 1.

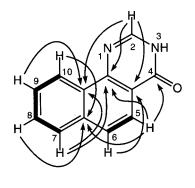
The EtOAc-soluble fraction of the methanolic extract of the seeds was partitioned between 3% tartaric acid and EtOAc. The 3% tartaric acid-soluble extract was adjusted to pH 9.0 with saturated Na₂CO₃ solution followed by partitioning with CHCl₃. The CHCl₃-soluble materials were subjected to a Si gel column (CHCl₃/MeOH, 15:1), followed by C₁₈ HPLC (MeOH/MeCN/H₂O, 1:1:3), to afford samoquasine A (**1**, 0.003%), together with a known aporphine alkaloid, liriodenine.⁴

Samoquasine A (1) has a molecular formula of $C_{12}H_8N_2O_1$, as revealed by HRFABMS (m/z 197.0728, $[M + H]^+$, $\Delta + 1.3$ mmu). IR absorptions at 3422, 1663, and 1626 cm^{-1} demonstrated the presence of hydroxyl and conjugated carbonyl groups, respectively. ¹H and ¹³C NMR data (Table 1) indicated that the molecule possesses a conjugated ketone carbonyl and 11 olefinic carbons, one of which is included in an imino group. Because 7 out of 10 unsaturations were accounted for, compound 1 was inferred to contain a naphthalene ring system and a pyrimidine ring. Interpretation of the ¹H-¹H COSY and HOHAHA spectra revealed proton connectivities from H-5 to H-6 and from H-7 to H-10. The connections between the above units and two sp² carbons (C-6a and C-10a) were assigned on the basis of ¹H-¹³C long-range correlations in the HMBC spectrum. (See Figure 1.) HMBC correlations of H-9 to C-10a ($\delta_{\rm C}$ 148.5), H-8 to C-6a ($\delta_{\rm C}$ 123.6), and H-5 to C-6a suggested a connectivity from C-5 to C-10 through C-6a and C-10a. In addition, connectivities indicated by HMBC correlations of H-10/C-6a, H-7/C-10a, H-7/C-10b, H-5/C-10b, and H-6/C-4a confirmed a 1,2-disubstituted naphtha-

[‡] Universiti Sains Malaysia.

Table 1. $^{1}\mathrm{H}$ and $^{13}\mathrm{C}$ NMR Data of Samoquasine A (1) in CDCl_3

position	δ_{H} (<i>J</i> , Hz)	$\delta_{\rm C}$	HMBC (1H)
2	9.60 s	150.5	
4		164.0	5
4a		118.7	2,6
5	7.55 d (7.3)	136.0	6
6	7.30 d (7.3)	101.6	
6a		123.6	5, 8, 10
7	8.40 dd (1.0, 8.3)	125.0	9
8	7.71 dt (1.0, 8.3)	128.7	10
9	7.87 dt (1.0, 8.3)	132.8	
10	8.12 dd (1.0, 8.3)	130.4	8
10a		148.5	2, 7, 9
10b		144.3	2, 5, 7



■ ¹H-¹H COSY & HOHAHA ----> HMBC

Figure 1. 2D NMR correlations fo samoquasine A (1).

lene skeleton. An amide carbonyl carbon, C-4 ($\delta_{\rm C}$ 164.0), and C-2 ($\delta_{\rm C}$ 150.5), assignable to the carbon between two nitrogen atoms, were included as part of a pyrimidine ring, together with the remaining two olefinic carbons, C-4a and C-10b ($\delta_{\rm C}$ 118.7 and 144.3). Thus, the singlet olefinic proton (H-2) at $\delta_{\rm H}$ 9.60, which was attached to C-2 ($\delta_{\rm C}$ 150.5), showed long-range correlations to C-4a, C-10a ($\delta_{\rm C}$ 148.5), and C-10b, while H-5 ($\delta_{\rm H}$ 7.55) showed long-range correlations to C-10b ($\delta_{\rm C}$ 164.0), completing the benzoquinazoline skeleton. Accordingly, the structure of samoquasine A (1) was elucidated as 3,4-dihydrobenzo[*h*]quinazolin-4-one.

Treatment of samoquasine A (1) with trimethylsilyldiazomethane afforded the *O*-methyl derivative (2), of which the ¹H and ¹³C NMR spectra were almost the same as those of 1, except for the existence of a methoxyl signal ($\delta_{\rm H}$ 3.72)

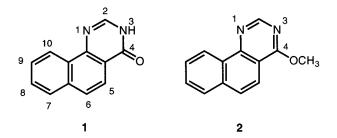
10.1021/np000342i CCC: \$19.00 © 2000 American Chemical Society and American Society of Pharmacognosy Published on Web 11/08/2000

^{*} To whom correspondence should be addressed. Tel.: +81-11-706-4985. Fax: +81-11-706-4989.

[†] Hokkaido University.

[§] Tokyo University of Pharmacy & Life Science.

in 2. The cross-peak for H₃-OMe/H-5 in the NOESY spectrum indicated that the methoxyl group was located at C-4. These data supported the proposed structure of samoquasine A (1).



This is the first isolation of a benzo[*h*]quinazoline alkaloid from a plant in the Annonaceae, although many aporphine alkaloids such as liriodenine have been reported from this family.² Samoquasine A (1) exhibited significant cytotoxicity against murine lymphoma L1210 cells, with an IC₅₀ value of 0.38 μ g/mL.

Experimental Section

General Experimental Procedures. Melting points were measured on a micromelting point apparatus and are uncorrected. UV spectra were obtained on a Shimadzu UV-1600PC spectrophotometer. IR spectra were run on a JASCO FT/IR-230 infrared spectrometer recorded as KBr pellets. ¹H and 2D NMR spectra were recorded in CDCl₃ on a 600 MHz spectrometer at 300 K, while ¹³C NMR spectra were measured at 125 MHz (Bruker AMX600 and ARX500). Chemical shifts were reported using residual CDCl₃ peaks ($\delta_{\rm H}$ 7.26 and $\delta_{\rm C}$ 77.03) as internal references. Standard pulse sequences were employed for the 2D NMR experiments. HMBC spectra were recorded using a 50-ms delay time for long-range C-H coupling with z-axis PFG. NOESY spectra were measured with a mixing time of 800 ms. FABMS were measured using a glycerol matrix.

Plant Material. The seeds of A. squamosa were collected at Penang, Malaysia, in December 1997. The botanical identification was made by one of us (K.-L.C.). A voucher specimen (no. 971201) has been deposited in the herbarium of this institute.

Extraction and Isolation. The seeds of A. squamosa (1.36 kg) were crushed and extracted with MeOH (3 L) three times to give an extract (225 g), which was treated with hexane, EtOAc, n-BuOH, and H₂O, successively. The EtOAc-soluble fraction (18.8 g) was partitioned by 3% tartaric acid to give a water-soluble fraction, which was adjusted to pH 9.0 with saturated Na₂CO₃ solution. The alkaline fraction was partitioned with CHCl₃, and the CHCl₃-soluble materials were subjected to Si gel column chromatography (CHCl₃/MeOH). A fraction that eluted with CHCl₃/MeOH (15:1) was subjected to C₁₈ HPLC (Develosil ODS-HG-5, MeOH/CH₃CN/H₂Ŏ, 1:1: 3) to give samoquasine A (1, 0.003%) and liriodenine (0.0007%).

Samoquasine A (1): colorless needles; mp >300 °C; UV (MeOH) $\bar{\lambda}_{max}$ (log ϵ) 244 (ϵ 3.7), 251 (3.6), 322 (3.1), 337 (3.1), 353 (3.1) nm; (MeOH + 1 N NaOH) λ_{max} 240 (ϵ 3.8), 287 (3.1), 325 (3.0), and 368 (3.2) nm; IR (KBr) v_{max} 3422, 2918, 2849, 1663, 1626, 1463, 1384, and 1127 cm⁻¹; FABMS *m*/*z* 197 [M + H]⁺; HRFABMS m/z 197.0728 [M + H] (calcd for C₁₂H₉N₂O, 197.0715); ¹H and ¹³C NMR data, see Table 1.

Methylation of 1. To a hexane solution (100 μ L) containing 2.0 M trimethylsilyldiazomethane was added a methanol solution (200 μ L) of **1** (1 mg), which was stirred at room temperature for 10 min. The solution was evaporated under reduced pressure, and the residue was chromatographed on a Si gel column (CHCl₃/MeOH, 20:1) to give the methyl derivative **2** (0.8 mg) as a colorless powder: ¹H NMR (CDCl₃) δ 3.72 (3H, s, OMe, C-4), 7.20 (1H, d, J = 7.4, H-6), 7.54 (1H, d, J = 7.4, H-5), 7.69 (1H, t, J = 8.0, H-8), 7.86 (1H, t, J = 8.0, H-9), 8.22 (1H, d, J = 8.0, H-10), 8.33 (1H, d, J = 8.0, H-7), and 9.78 (1H, s, H-2); FABMS m/z 211 [M + H]+.

Cytotoxic Assay. Assays were carried out by the method described previously.5

Acknowledgment. This work was partly supported by a Grant-in-Aid for Scientific Research from the Ministry of Education, Science, Sports, and Culture of Japan.

References and Notes

- (1) Hopp, D. C.; Zeng, L.; Gu, Z.-M.; McLaughlin, J. L. J. Nat. Prod. 1996, *59*, 97–99.
- Leboeuf, M.; Cave, A.; Bhaumik, P. K.; Mukherjee, B.; Mukherjee, (2)(2) Leboetti, M., Cave, A., Bhaumin, P. K., Mutheljee, B., Mutheljee, R. *Phytochemistry* **1982**, *21*, 2783–2813.
 (3) Morita, H.; Sato, Y.; Kobayashi, J. *Tetrahedron* **1999**, *55*, 7509–7518.
 (4) Buchanan, M. A.; Dickey, E. E. *J. Org. Chem.* **1960**, *25*, 1389–1391.
 (5) Kobayashi, J.; Shigemori, H.; Hosoyama, H.; Chen, Z.-S.; Akiyama, C. S. (2017).

- S.; Naito, M.; Tsuruo, T. Jpn. J. Cancer Res. 2000, 91, 638-642.

NP000342I