

Bioassay and Ultrapformance Liquid Chromatography/Mass Spectrometry Guided Isolation of Apoptosis-Inducing Benzophenones and Xanthone from the Pericarp of *Garcinia yunnanensis* Hu

GANG XU,[†] CHAO FENG,[‡] YAN ZHOU,[†] QUAN-BIN HAN,[†] CHUN-FENG QIAO,[†]
 SHENG-XIONG HUANG,[†] DONALD C. CHANG,[§] QIN-SHI ZHAO,[⊥]
 KATHY Q. LUO,^{*‡} AND HONG-XI XU^{*†}

Chinese Medicine Laboratory, Hong Kong Jockey Club Institute of Chinese Medicine, Shatin, Hong Kong, People's Republic of China, Bioengineering Graduate Program, Department of Chemical Engineering, and Department of Biology, Hong Kong University of Science and Technology, Clear Water Bay, Kowloon, Hong Kong, People's Republic of China, and State Key Laboratory of Phytochemistry and Plant Resources in West China, Kunming Institute of Botany, Chinese Academy of Sciences, Kunming 650204, People's Republic of China

Bioassay and ultrapformance liquid chromatography/photodiode array/mass spectrometry (UPLC/PDA/MS) guided isolation of the apoptosis-inducing active metabolites on HeLa-C3 cells from the pericarp of *Garcinia yunnanensis* (Guttiferae) yielded five active compounds, including the new garciyunnanins A (1) and B (2). The structures of the compounds were elucidated by comprehensive nuclear magnetic resonance and mass spectrometry analysis. Garciyunnanin B (2), featured with a natural tetracyclic xanthone skeleton derived from a polyisoprenylated benzophenone, is structurally interesting since it can be seen as an evidence of the previously described cyclization of garcinol by 2,2-diphenyl-1-picrylhydrazyl (DPPH). Garciyunnanin A (1) contains a 3-monohydroxy benzophenone skeleton, which is rarely found in *Garcinia* species. Both new compounds induce HeLa-C3 cells into apoptosis after 72 h of incubation at 15 μ M. It is noteworthy that oblongifolin C (4), the major constituent of this plant, has proved to be the most active one among the isolates for inducing apoptotic cell death in cervical cancer derived HeLa-C3 sensor cells.

KEYWORDS: *Garcinia yunnanensis*; apoptosis-inducing activity; UPLC/MS; benzophenones; xanthones

INTRODUCTION

Garcinia is a plant genus of the Guttiferae family native of Asia, Southern Africa, and Polynesia, with a total of 21 species distributed in China (1). The fruits of many plants in this genus are edible such as *G. mangostana*, *G. subfalcata*, *G. esculenta*, and *G. yunnanensis*, etc. (1). A number of polyisoprenylated benzophenones and polyisoprenylated xanthones have been isolated from *Garcinia* plants to date in which many are purified from the fruits with garcinol being the best representative (2–7).

Garcinol, a polyisoprenylated benzophenone purified from *G. indica* fruits and other species, has been reported to possess important biological activities such as antibiotic and antiulcer activities (8) and the ability to suppress colonic aberrant crypt foci (ACF) formation and inhibit histone acetyltransferases (HATs), which modulate gene expression (2), ability of induction of apoptosis through cytochrome *c* release, and activation of caspases in human leukemia HL-60 cells (9), and anti-inflammatory (10, 11) and anticarcinogenic properties (10, 11). Aiming at identifying structurally interesting and bioactive metabolites from the *Garcinia* species, we have started a series of studies on the chemical structures and pharmaceutical properties of several *Garcinia* species including *G. xipshuanbannaensis* (12), *G. kola* (13), *G. lancilimba* (14), *G. xanthochymus* (15), and *G. hanburyi* (16–19). Analytical studies of the chemical constituents of *Garcinia* plants using high-speed countercurrent chromatography (HSCCC), ultrapformance liquid chromatography/mass spectrometry (UPLC/MS), and high-performance

* Corresponding authors. Tel.: +852-3406-2873 (H.-X.X.), 852-2358-8434 (K.Q.L.). Fax: +852-3551-7333 (H.-X.X.), 852-3106-4857 (K.Q.L.). E-mail: xuhongxi@hkjicm.org (H.-X.X.), qluo@ust.hk (K.Q.L.).

[†] Hong Kong Jockey Club Institute of Chinese Medicine.

[‡] Department of Chemical Engineering, Hong Kong University of Science and Technology.

[§] Department of Biology, Hong Kong University of Science and Technology.

[⊥] Kunming Institute of Botany, Chinese Academy of Sciences.

liquid chromatography (HPLC) were also carried out in our laboratory (20–23).

Ultraperformance liquid chromatography/photodiode array/mass spectrometry (UPLC/PDA/MS) is a powerful tool for the phytochemical analysis since it can quickly separate the compounds with high efficiency and provide on-line UV and MS structural information (21, 24). All the xanthenes and benzophenones isolated from *Garcinia* plants in our laboratory have been analyzed using the UPLC/PDA/MS equipment. On the basis of these analytical results, we have constructed a database containing the uniform UPLC method and MS information of 20 benzophenones and 36 xanthenes obtained from *Garcinia* plants. To accelerate the discovery of new and bioactive compounds, an apoptotic assay and UPLC/PDA/MS guided isolation process were developed. In this method, each step of the extraction and fractionation process was guided by the UPLC/PDA/MS analysis and the cell-based apoptotic assay using HeLa-C3 cells that contain a fluorescent caspase sensor. This method enabled us to focus on the fractions and compounds with clear apoptotic effects. More importantly, for compounds with positive apoptotic effects and previously known chemical structures, their identities can be easily determined by comparing their UPLC/PDA/MS profiles with those in the database.

In this study, we used this combinative method to isolate anticancer compounds with apoptotic activities from *G. yunnanensis* Hu, a key protected plant distributed in the southwestern part of the Yunnan province in China. The fruits of this plant, called “Xiaoguliangguo” by local people, are sweet and tasty (1). During our initial apoptotic activity survey, different parts of this plant, including the twig, leaves, and fruit, were studied. The results indicated that the acetone-soluble fraction of the fruit displayed the highest apoptotic activity against HeLa-C3 cells. Further investigations on the different extracts of the pericarps and seeds indicated that the acetone extract of pericarp was more effective than others. We report herein the bioassay and UPLC/PDA/MS guided isolation, structural elucidation, and apoptotic activity evaluation of the metabolites from the pericarps of *G. yunnanensis*.

MATERIALS AND METHODS

Instrumentation. Optical rotations were measured with a Horiba SEPA-300 high-sensitivity polarimeter (Horiba, Kyoto). Ultraviolet absorption spectra were recorded by a UV-2401 PC spectrophotometer (Shimadzu, Kyoto). IR spectra were obtained from a Bio-Rad FtS-135 spectrometer (Bio-Rad, California). NMR spectra were measured on a Bruker AV-400 spectrometer with TMS as the internal standard. Mass spectrometry was performed on a Waters Q-TOF Premier (Micromass MS Technologies, Manchester, U.K.) equipped with an electrospray ionization (ESI) source in positive ion mode. Column chromatography (CC) was performed with silica gel 60 (200–300 mesh, Merck), Sephadex LH-20, and reversed-phase C-18 silica gel (250 mesh, Merck). Precoated thin-layer chromatography (TLC) sheets of silica gel 60 GF254 were used. An Agilent 1100 series equipped with Alltima C-18 column (4.6 mm × 250 mm) was used for HPLC analysis, and the preparative Alltima C-18 column (22 mm × 250 mm) was used in the sample preparation.

UPLC Analysis. UPLC was performed using a Waters ACQUITY UPLC system (Waters Corp., MS), equipped with a binary solvent delivery system, autosampler, and a PDA detector. The chromatography was performed on a Waters ACQUITY BEH C₁₈ column (100 mm × 2.1 mm, 1.7 μm, Waters Corp., Irel mL/min.) The column and autosampler were maintained at 35 and 10 °C, respectively. The mobile phase consisted of (A) 0.1% formic acid in water and (B) acetonitrile containing 0.1% formic acid. The UPLC eluting conditions were optimized as follows: isocratic at 75% B (0–0.5 min), linear gradient from 75% to 85% B (0.5–3 min), isocratic at 85% B (3–6 min), linear

Table 1. ¹³C and ¹H NMR Data (100 and 400 MHz, CD₃OD) for 1 and 2

	garciyunnanin A (1)		garciyunnanin B (2)	
	δ _C	δ _H	δ _C	δ _H
1	194.3		195.8	
2	120.0		120.6	
3	191.7		177.1	
4	69.8		66.4	
5	51.2		51.6	
6	42.0	1.84, m	39.4	1.79, m
7	42.8	2.01, m	43.7	2.09, m
		1.45, t (12.8)		1.49, m
8	64.4		66.4	
9	209.5		207.2	
10	199.1		173.9	
11	140.9		118.3	
12	116.2	7.04, s	151.2	
13	158.5		104.1	6.81, s
14	120.5	6.94, overlap	155.3	
15	129.8	7.15, t (7.8)	147.3	
16	121.3	6.95, overlap	109.5	7.44, s
17	26.6	2.68, m	26.6	3.00, m
		2.64, m		2.87, m
18	121.7	4.82, m	120.8	4.70, m
19	134.5		135.2	
20	26.2	1.63, s	26.1	1.49, s
21	16.1	1.67, s	18.6	1.82, s
22	16.1	0.81, s	18.2	0.90, s
23	37.4	1.68, m	37.4	1.92, m
				1.56, m
24	25.2	1.97, m	23.7	1.80, m
25	125.6	5.05, m	124.7	4.92, m
26	132.3		133.1	
27	26.0	1.66, s	26.1	1.59, s
28	18.2	1.59, s	17.8	1.21, s
29	30.1	2.08, m	30.1	1.98, m
		1.78, m		1.76, m
		5.03, m		4.92, m
30	123.7		123.3	
31	134.4		138.4	
32	25.9	1.68, s	40.7	1.96, m
33	18.3	1.57, s	16.5	1.54, s
34	31.4	2.45, m	31.3	2.52, m
35	121.1	5.10, m	120.7	4.96, m
36	135.1		135.2	
37	26.2	1.69, s	26.1	1.59, s
38	17.9	1.66, s	18.2	1.69, s
39			27.5	2.52, m
40			125.1	5.02, m
41			132.3	
42			25.9	1.64, s
43			17.6	1.56, s

gradient from 85% to 95% B (6–7 min), isocratic at 95% B (9–12 min), and linear gradient from 95% to 75% B (12–13 min). The flow rate was 0.5 mL/min.

Plant Material. The pericarp of *Garcinia yunnanensis* Hu were collected in Luxi of Dehong prefecture, Yunnan province, China in 2006. The plant material was identified by Dr. Chunfeng Qiao. A herbarium sample (CMED-0477) was deposited in the Hong Kong Jockey Club Institute of Chinese Medicine.

Extraction and Isolation. The air-dried and powdered pericarp (9.0 kg) was extracted with acetone (20 L) at room temperature for three times. The extracted solution was evaporated under reduced pressure to yield a dark green residue (1.2 kg). The residue was chromatographed on silica gel eluted by CHCl₃, EtOAc, and acetone sequentially. The CHCl₃ fraction was tested to be the active fraction and evaporated in vacuum to give a residue (750 g), part of which (400 g) was subjected to silica gel column eluted with a gradient hexane/acetone system (100:0 to 0:100, v/v). Four fractions (I–V) were obtained on the basis of TLC analysis. The first three fractions were all tested to have apoptotic effects against HeLa-C3 cells. Fraction I was purified by reversed-phase C-18 silica gel to afford 4 (oblongifolin C, 5.0 g). Fraction II was separated using silica gel and eluted with gradient petroleum ether/acetone system (10:0 to 0:10) to produce four subfractions. The result of caspase activity

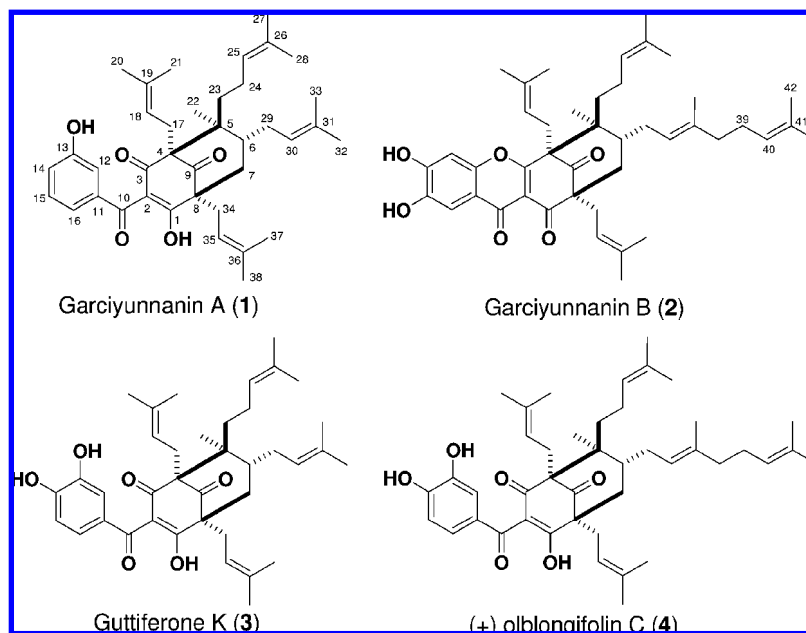


Figure 1. Structures of the new compounds (**1** and **2**) and two major constituents (**3** and **4**) of *Garcinia yunnanensis*.

assay indicated that the second and third subfractions were positive. The second subfraction was separated by preparative HPLC on an Alltima C-18 column eluted with CH₃CN in 0.1% acetic acid (0.1% acetic acid/CH₃CN, 5/95) to yield **3** (guttiferone K, 3.0 g) and **7** (oblongifolin D, 5.0 mg). The third subfraction was also separated by preparative HPLC (CH₃CN/H₂O, 85/15) to afford **4** (oblongifolin C, 6.0 mg). Fraction III was chromatographed on reversed-phase silica gel eluted with a gradient of CH₃OH/H₂O (from 8:2 to 10:0) to give five subfractions in which the second and the third subfractions demonstrated apoptotic activities. The second subfraction was then chromatographed over preparative HPLC on an Alltima C-18 column (CH₃CN/H₂O, 9/1) to give **1** (garciyunnanin A, 7.0 mg), **5** (oblongifolin A, 5.0 mg), and **8** (diethylhexyl phthalate, 10.0 mg). The third subfraction was further separated in a similar way, eluted with CH₃CN/H₂O (8:2) to give **2** (garciyunnanin B, 6.0 mg) and **6** (oblongifolin B, 9.0 mg).

Physical data for garciyunnanin A (**1**) are the following: yellow oil; [α]_D^{26.2} -3.0° (CHCl₃; *c* 0.11); IR (KBr) ν_{\max} 3424, 2965, 2925, 2856, 1727, 1670, 1623, 1581, 1446, 1378, 1287, 1116, 1074, 997 cm⁻¹; UV (CH₃OH) λ_{\max} (log ϵ) 204.2 (1.72), 262.0 (1.31) nm; ¹H and ¹³C NMR spectral data, see **Table 1**; positive HRESIMS *m/z* 587.3742 [M + H]⁺, (calcd for C₃₈H₅₀O₅, 587.3736).

Physical data for garciyunnanin B (**2**) are the following: yellow oil; [α]_D^{26.2} $+18.1^\circ$ (CHCl₃; *c* 0.12); IR (KBr) ν_{\max} 3425, 2967, 2923, 1734, 1687, 1620, 1465, 1390, 1291, 1146, cm⁻¹; UV (CH₃OH) λ_{\max} (log ϵ) 204.5 (1.77), 263.0 (1.44), 287.0 (1.32), 369.5 (1.11) nm; ¹H and ¹³C NMR spectral data, see **Table 1**; positive HRESIMS *m/z* 669.4146 [M + H]⁺, (calcd for C₄₃H₅₇O₆, 669.4155).

Bioassay. All the tested samples were dissolved in dimethyl sulfoxide (DMSO) to make stock solutions. The concentration of each stock is at least 1000 times higher than the working concentration. HeLa-C3 cells, which can detect apoptotic cell death involving caspase activation, were cultured in minimum essential medium (MEM) containing 10% fetal bovine serum, 100 U/mL penicillin, 100 mg/mL streptomycin, in a 5% CO₂ humidity incubator at 37 °C. In order to test a sample with apoptotic activity, one well of a 96-well plate was seeded with 7500 HeLa-C3 cells suspended in 100 μ L of culture medium as a sample well. Another well of the plate was added with only 100 μ L of culture medium as a background well. After 12 to ~16 h of incubation, the plate was read by a Perkin-Elmer Victor plate reader with excitation wavelength at 440 \pm 8 nm and emission wavelength at 486 \pm 8 nm for CFP and 535 \pm 8 nm for YFP to obtain the data for time point "0 h". Then immediately, the old medium was removed and 100 μ L of freshly prepared culture medium containing the testing sample at a designed working concentration was added for both the sample well and the corresponding background well. Culture medium containing

0.1% DMSO was used as a negative control, whereas 500 nM paclitaxel was used as a positive control. After that, the plate was read repeatedly at indicated time points. The data acquisition duration was up to 72 h. The YFP/CFP emission ratio was then calculated. If the YFP/CFP emission ratio was decreased down below 3, then a significant apoptosis was considered to occur. All the screening of samples was conducted in triplicate, and all experiments were repeated for three times.

RESULTS AND DISCUSSION

The pericarp of *G. yunnanensis* was extracted with acetone and then chromatographed on silica gel eluted with CHCl₃, EtOAc, and acetone sequentially. The CHCl₃ fraction was tested to be the best one by our cell-based caspase activity assay. The UPLC/PDA/MS analysis of the CHCl₃ fraction revealed eight compound peaks (**Figure 1**). Then, by comparing the retention time, UV, and MS information with that of authentic compounds in our database, the compound peaks **4–7** were identified to be oblongifolins A–D (**25**). The accuracy of these identifications was later confirmed by the isolation of the pure compounds as shown in the Extraction and Isolation Procedures section. However, compounds **1**, **2**, and **3**, with the [M + H]⁺ ion peak at 587.3742, 669.4146, and 603.3686, respectively, were absent from the database. Further isolation and purification using a series of chromatographic separations finally yielded the bioactive components garciyunnanin A (**1**) and garciyunnanin B (**2**) together with guttiferone K (**3**) and diethylhexyl phthalate (**8**) (**26**, **27**). In addition, compounds **4–7** were also isolated. Both new compounds can induce HeLa-C3 cells into apoptosis after 72 h of incubation at 15 μ M. It is noteworthy that among the five active components (**1–5**) tested, the major constituent, oblongifolin C (**4**), was the most potent (**Figure 4**).

Compound **1** was obtained as a yellow oil; its molecular formula was determined to be C₃₈H₅₀O₅ by high-resolution electrospray ionization mass spectroscopy (HRESIMS) (found [M + H]⁺ 587.3742, calcd 587.3736) and ¹³C NMR spectrum. The IR spectrum displayed bands for hydroxyl (3424 cm⁻¹) and carbonyl (1727, 1670, 1623 cm⁻¹) groups. The ¹H NMR spectrum (**Table 1**) displayed that **1** possessed a 1,3-disubstituted benzene ring, four olefinic protons, one methyl on sp³ carbon, and eight vinyl methyl groups. The ¹³C NMR spectrum of **1** exhibited the presence of six aromatic carbons, a conjugated

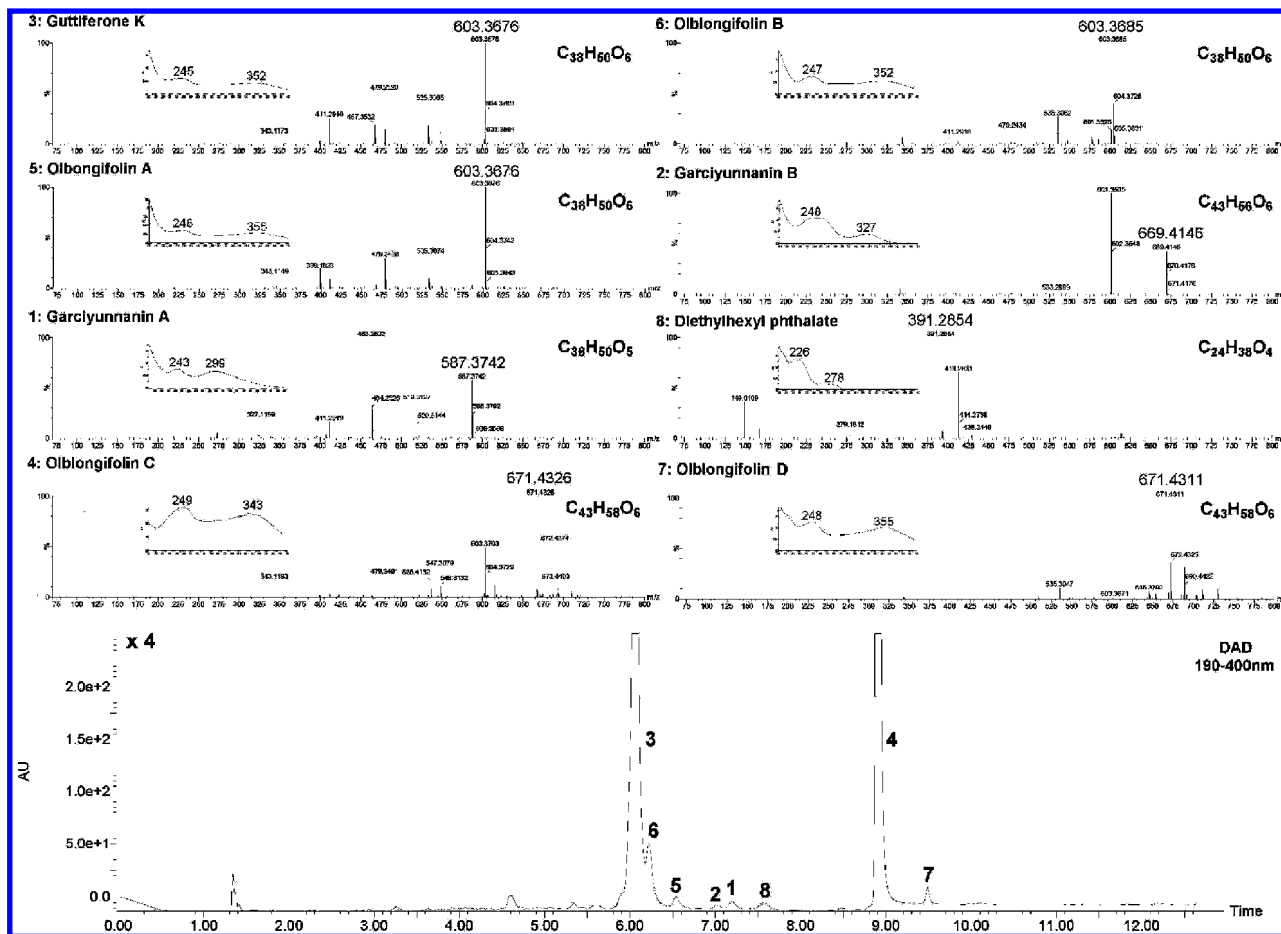


Figure 2. UPLC/PDA/MS analysis of the CH₃Cl part of the acetone extract together with the HRESIMS and UV information of the isolated compounds.

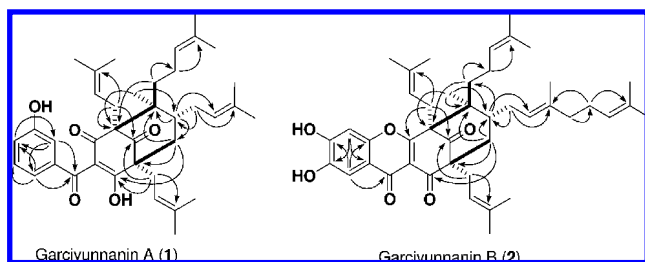


Figure 3. Key HMBC correlations of compounds 1 and 2.

carbonyl at δ_C 199.1 (C-10), a nonconjugated carbonyl at δ_C 209.5 (C-9), an enolized 1,3-diketone group (δ_C 194.3, C-1; δ_C 120.0, C-2; δ_C 191.7, C-3), two quaternary carbons at δ_C 69.8 (C-4) and 64.4 (C-8), and 25 other signals assignable to four isoprene units and another C5 unit. Considering a number of benzophenones have been isolated from *Garcinia* species, along with the spectrometric features discussed above, **1** could be ascribed to be a typical benzophenone derivative (2, 25, 26).

The ¹H and ¹³C NMR data (Table 1) of **1** is very similar to those of guttiferone K (**3**), which was also isolated as one of the major constituents in this study (26). The only difference observed in the ¹H and ¹³C NMR spectra was the presence of one more aromatic methine (δ_C 120.5; δ_H 6.94, overlap) in **1** instead of the quaternary aromatic carbon at δ_C 152.6 in **3**. In addition, the chemical shift value of C-13 was also downshifted from δ_C 146.5 in **3** to δ_C 158.5 in **1**. Therefore, compound **1** was deduced to be the 14-dehydroxyl derivative of guttiferone K (**3**). In the heteronuclear multiple bond correlation (HMBC) spectrum, the correlations from H-12 (δ_H 7.04, s) to C-10, C-13,

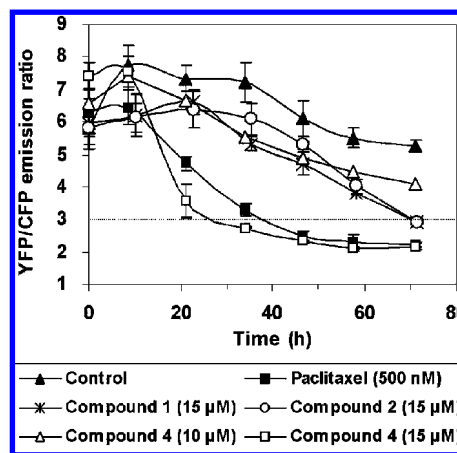
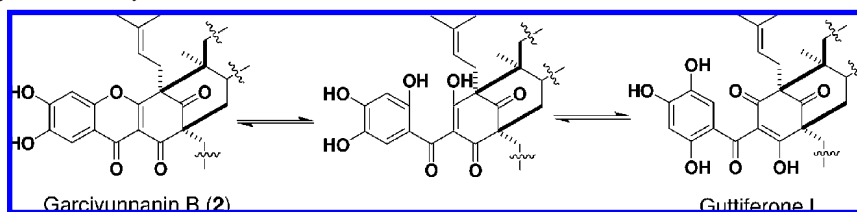


Figure 4. Time-dependent YFP/CFP emission ratio changes of **1**, **2**, and **4**.

C-14, and C-16, from H-14 (δ_H 6.94, overlap) to C-12, C-15, and from H-16 (δ_H 6.95, overlap) to C-10, C-12, C-14, and C-15 confirmed that the aromatic ring in **1** was 1,3-disubstituted (Figure 3). In addition, the HMBC correlations between H-6/C-5, C-7, C-29, and C-30; H-7/C-1, C-5, C-6, and C-8; H-17/C-4, C-9, C-18, and C-19; H-22/C-4, C-5, C-6, and C-23; H-23/C-4, C-22, and C-24; H-24/C-23, C-25, and C-26; and H-29/C-5, C-30, and C-31 were also found, which further confirmed the core fragment (a 2,2-dimethylbicyclo[3,3,1]-nonane ring system) of **1** and the locations of four prenyl groups, respectively.

Scheme 1. Plausible Biogenetic Pathway of **2**Table 2. Apoptosis-Inducing Effects of Compounds 1–8^a

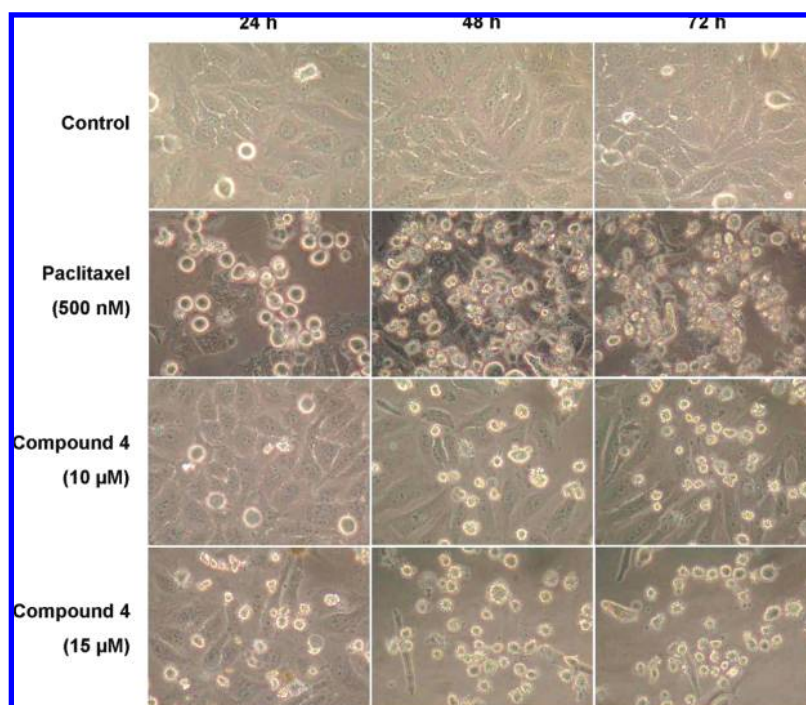
compd	apoptosis-inducing effects at	
	20 $\mu\text{g/mL}$	10 $\mu\text{g/mL}$
1	–	+
2	+	+
3	+	–
4	+	+
5	+	–
6	–	–
7	–	–
8	–	–

^a “+” indicates the compound could decrease YFP/CFP emission ratio below 3 in 72 h.

The relative stereochemistry of **1** was deduced to be consistent with that of guttiferone K (**3**). The obvious nuclear Overhauser effect (NOE) correlations between CH₃-22 and CH₂-17/H-7 α found in the nuclear Overhauser enhancement spectroscopy (NOESY) spectrum indicated that CH₃-22 must be in the α -orientation like CH₂-17. As mentioned in the literature (2, 25), the large coupling constant between H-7 α (δ_{H} 1.45, t, J = 12.8 Hz) and H-6 further revealed the axial orientation for H-6 and H-7 α together with the equatorial asset of the isoprenyl group at C-6. In addition, the ¹³C NMR chemical shift of C-6 at δ_{C} 42.0 suggested that H-6 was β -oriented, since the signal of H-6 with α -orientation is located at a lower field (δ_{C} 46.0–48) (26, 28). On the basis of all the above evidence, the structure of **1** was established and named as garciyunnanin A.

Compound **2**, named garciyunnanin B, was isolated as yellow oil. The HRESIMS (m/z 669.4146 [M + H]⁺, calcd 669.4155)

indicated that **2** has a molecular formula of C₄₃H₅₆O₆, which was supported by the ¹³C NMR spectrum (Table 1). Analysis of the ¹H and ¹³C NMR of **2** exhibited the presence of a 1,2,4,5-tetrasubstituted benzene ring, a conjugated carbonyl carbon at δ_{C} 173.9 for C-10, a 2,2-dimethylbicyclo[3,3,1]-nonane ring system, and five prenyl groups. Careful comparison of the NMR data of **2** with another major constituent of this plant, (+)-oblongifolin C (**4**) (25), indicated that they are very similar to each other except for the signals for the aromatic ring and the enolized 1,3-diketone group due to C-1, C-2, and C-3. Different from the trisubstituted benzene ring in **4**, the aromatic ring in **2** was determined to be a 1,2,4,5-tetrasubstituted benzene ring based on the analysis of its ¹H and ¹³C NMR data. Two singlets (δ_{H} 6.81 and 7.44) found in the aromatic region of the ¹H NMR spectrum were assignable to H-13 (δ_{H} 6.81) and H-16 (δ_{H} 7.44), respectively, which can be confirmed by the HMBC correlations from both H-13 and H-16 to C-10, C-11, C-12, C-14, and C-15 (Figure 3). An ether bridge between C-3 or C-1 and C-12 through an oxygen atom was determined by MS and NMR data analysis, which can be seen as the most interesting characteristic of its structure. Because the enolized 1,3-diketone can undergo keto–enol tautomerism, the formation of the epoxy group could occur between C-12 and C-3 or between C-12 and C-1. In (+)-oblongifolin C (**4**), C-3 of the enolized 1,3-diketone group was a carbonyl and C-1 was substituted by a hydroxyl group, whereas in **2**, the carbonyl group (δ_{C} 195.8) was assigned to C-1 based on its HMBC correlations with H-7 together with the HMBC correlations from H-17 to the olefinic carbon at δ_{C} 177.1 (C-3), C-4, and C-5. The relative stereochemistry of C-4,

Figure 5. Cell morphology changes during the course of compound **4** treatment.

C-5, C-6, C-7, and C-8 were deduced to be the same with those of **1** based on the comparative analysis of the ^1H and ^{13}C NMR data with those of **1** and oxyguttiferone K (**3**, **26**), together with the NOE correlations between $\text{CH}_3\text{-22}$ and $\text{CH}_2\text{-17/H-7}\alpha$ in the NOESY spectrum. Therefore, the structure of **2** was determined and named as garciyunnanin B.

As mentioned above, the most interesting characteristic of the structure of garciyunnanin B (**2**) is the epoxy linkage between C-3 and C-12, which was the second example of natural xanthone possessing this tetracyclic skeleton. Just as the isolation of the first tetracyclic xanthone, oxyguttiferone K, the natural formation of garciyunnanin B (**2**) also confirmed the previously described cyclization of garcinol by 2,2-diphenyl-1-picrylhydrazyl (DPPH) (**3**, **10**, **11**). From a biogenetic view, garciyunnanin B (**2**) should derived from a 12,13,14-trihydroxy benzophenone compound such as guttiferone L (**Scheme 1**) (**26**). As to garciyunnanin A (**1**), it is also noteworthy since its 12-monohydroxyl polyprenylated benzophenone skeleton is seldom found in *Garcinia* species.

All the isolated *G. yunnanensis* extracts and compounds were tested for their apoptotic activities using genetically engineered HeLa-C3 cells that can produce a fluorescent biosensor capable of detecting caspase-3 activation (**29**). These cells can emit green light under normal growth conditions and change to blue light when caspase-3 is activated during apoptosis to cleave the sensor protein inside the cells. This color change allows one to use a fluorescent plate reader to directly detect the activation level of caspase-3 in HeLa-C3 cells during the course of the compound treatment in a noninvasive way (**30**).

On the basis of our previous results, the emission ratio of YFP (yellow fluorescent protein)/CFP (cyan fluorescent proteins) is usually between 6 and 8 in normal cells, and this ratio will decrease to a value below 3 if a compound can activate caspase-3 and kill cancer cells. Therefore, any compound that can reduce the YFP/CFP emission ratio to a value below 3 will be considered positive in activating apoptosis. As shown in **Table 2**, among the eight tested compounds, **2**, **3**, **4**, and **5** were positive in activating apoptosis in HeLa-C3 cells at a concentration of 20 $\mu\text{g/mL}$. At 10 $\mu\text{g/mL}$, compounds **2** and **4** were still positive, whereas **3** was not, suggesting that **3** is less potent than **2** and **4**. It is worth mentioning that compound **1** tested positive for apoptosis at 10 $\mu\text{g/mL}$, but not at 20 $\mu\text{g/mL}$. We have examined cell morphology changes under compound **1** treatment at 20 $\mu\text{g/mL}$ and found that those cells displayed typical features of necrotic cell death, such as prominent nuclear membrane, cell round-up, and membrane rupture. Since HeLa-C3 cells can only specifically detect apoptosis but not necrosis, a negative result was expected for compound **1** at higher concentration.

The investigation on the time-dependent kinetics of caspase activation for compounds **1**, **2**, and **4** indicated that **4** could fully activate caspase-3 within 30 h of compound addition at 15 μM ($= 9.5 \mu\text{g/mL}$), while **1** and **2** required 72 h to reduce the YFP/CFP emission ratio to a level just below 3 (**Figure 4**). This result suggests that compound **4** is the most potent apoptotic inducer among the tested compounds from *G. yunnanensis*. We further examined the apoptotic effect of compound **4** at a lower concentration and found that, at 10 μM , it can partially activate caspase (**Figure 4**). In addition, cell morphology observation (**Figure 5**) showed that many cells became detached and died off at 48 h after the treatment of compound **4**, which further supports the results of our caspase activity assay.

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