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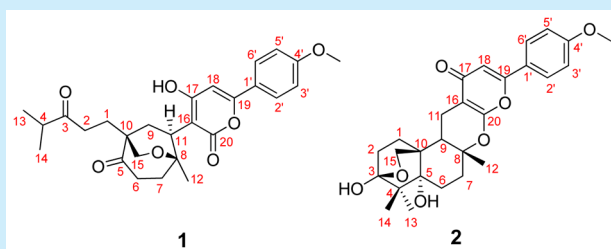
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New Meroterpenoids from *Aspergillus terreus* with Inhibition of Cyclooxygenase-2 ExpressionChih-Chuang Liaw,^{*,†,‡,§,⊥} Yu-Liang Yang,^{†,§,⊥,||} Chun-Kuang Lin,[†] Jin-Ching Lee,^{§,▽} Wen-Ying Liao,^Δ Chia-Ning Shen,^Δ Jyh-Horng Sheu,^{†,‡,§} and Shih-Hsiung Wu^{*,†,□,○}[†]Doctoral Degree Program in Marine Biotechnology, National Sun Yat-sen University, Kaohsiung 80424, Taiwan[‡]Department of Marine Biotechnology and Resources, National Sun Yat-sen University, Kaohsiung 80424, Taiwan^{||}Agricultural Biotechnology Research Center, Academia Sinica, Taipei 115, Taiwan[§]Graduate Institute of Natural Products, Kaohsiung Medical University, Kaohsiung, Taiwan[▽]Department of Biotechnology, Kaohsiung Medical University, Kaohsiung, Taiwan^ΔGenomics Research Center, Academia Sinica, Taipei 115, Taiwan[□]Department of Chemistry and Institute of Biochemical Sciences, National Taiwan University, Taipei 106, Taiwan[○]Institute of Biological Chemistry, Academia Sinica, Taipei 115, Taiwan

S Supporting Information

ABSTRACT: Two novel meroterpenoids, yaminteritrems A (1) and B (2), were isolated from *Aspergillus terreus* collected from hot spring zones in Yang-Ming Mountain, Taiwan, and cultured at 40 °C. The structures of 1 and 2 were elucidated by NMR, MS spectral and X-ray crystallographic analyses. The biosynthetic route for 1 and 2 involving the conversion of the sesquiterpene with phenyl- α -pyrone is proposed. Besides, 2 exhibited a dose-dependent inhibitory effect on COX-2 expression in LPS-stimulated RAW264.7 macrophages.



Meroterpenoids, a characteristic type of fungal metabolites, merge polyketide–terpenoid structures. Some of them have been reported as an inhibitor selective for acetylcholinesterase,¹ which can decrease the amount of acetylcholine present in the synapses between cholinergic neurons.² In our previous studies on indigenous thermophilic fungi, a large group of compounds from thermophilic *Aspergillus terreus* (Trichocomaceae) have been identified as sources of biofunctional chemical components.³ Our ongoing study on chemical investigations of the indigenous fungi are beginning to afford two novel meroterpenoids, yaminteritrems A (1) and B (2), in which the former possesses an unusual seven-member ring and the latter has a naphtho[2,1-*b*]pyrano-[3,2-*e*]pyran moiety.

We report, herein, the isolation and structure elucidation of compounds 1 and 2. The fungus, *A. terreus* (Stain No. C9408-3), collected from a hot spring zone in Yang-Ming Mountain, Taipei, was cultured at 40 °C for 7 days on potato dextrose agar (PDA) plates (400 plates) and then were extracted with ethyl acetate. The ethyl acetate extract (3.28 g, ASP-EA) was fractionated using a Sephadex LH-20 column eluted with MeOH to yield 20 fractions. Fraction 6 (ASP-EA-f6) was further separated by column chromatography on Sephadex LH-20 with MeOH and purified by RP-HPLC (Sunfire C18, 250 mm \times 4.5 mm, 1.0 mL/min, CH₃CN–H₂O, 60:40) to give compound 1 (1.7 mg, *t*_R 5.24 min). There were some particles

precipitated in Fraction 3. Then the precipitated particles were further separated by silica gel column and eluted with CHCl₃ to 30:1 CHCl₃–MeOH to yield compound 2 (28 mg).

Yaminteritrem A (1) was obtained as a yellowish oil, and its molecular formula was determined to be C₂₇H₃₂O₇ by HRESIMS on the [M + Na]⁺ (*m/z* 491.2043, calcd 491.2046 for C₂₇H₃₂O₇Na). The IR spectrum showed the presence of hydroxyl at 3406 cm^{−1} and ester/lactone carbonyl at 1692 cm^{−1}. The ¹³C NMR and DEPT spectra of 1 (Table 1) exhibited the presence of 27 carbon resonances, containing three carbonyl carbons (δ_C 215.4, 212.2, and 165.8), five quaternary aromatic carbons, four aromatic methines, one olefinic methine, one oxygen-bearing quaternary carbon (δ_C 84.7), one oxymethylene (δ_C 74.0), one quaternary carbon (δ_C 60.4), two methines, five methylenes, one methoxyl, and three methyls.

The ¹H NMR spectrum of 1 (Table 1) revealed the presence of a 1,4-disubstituted phenolic moiety (δ_H 7.78, dd, *J* = 8.9, 2.0 Hz; 7.05, dd, *J* = 8.9, 2.0 Hz), an olefinic methine (δ_H 6.55, s), one oxymethylene (δ_H 3.96, *J* = 8.7 Hz; 3.89, *J* = 8.7 Hz), one methoxyl at δ_H 3.88, and three methyls (δ_H 1.38, s; 1.06, d, *J* = 7.0 Hz; 1.03, d, *J* = 7.0 Hz).

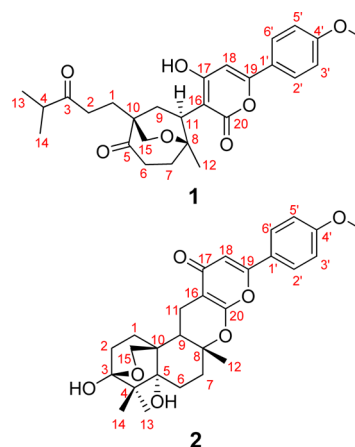
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Table 1. NMR Spectroscopic Data of **1** and **2**

no.	1		2^b	
	δ_{H} (J in Hz) ^a	δ_{C} ^b	δ_{H} (J in Hz) ^c	δ_{C} ^d
1	a 1.95 m	20.9	a 2.07 dd (9.2, 5.2)	28.9
	b 1.81 m		b 1.84 dd (9.2, 9.2)	
2	a 2.77 m	36.0	a 2.32 ddd (9.2, 9.2, 5.2)	29.6
	b 2.67 m		b 1.92 dd (9.2, 9.2)	
3		215.4		98.3
4	2.61 sep (6.8)	40.3		46.5
5		212.2		74.9
6	a 2.68 m	34.7	a 2.13 dd (11.6, 2.4)	28.4
	b 2.41 m		b 1.69 dt (11.6, 2.4)	
7	a 1.94 m	38.4	a 2.22 m	33.5
	b 1.89 m		b 1.91 d (9.6)	
8		84.7		84.0
9	a 2.78 m	19.3	2.20 m	42.2
	b 2.47 m			
10		60.5		40.5
11	2.77 m	46.7	a 2.64 m	16.3
			b 2.20 m	
12	1.38 s	20.9	1.28 s	19.9
13	1.03 d (7.0)	17.1	1.10 s	19.6
14	1.06 d (7.0)	17.1	1.14 s	21.3
15	a 3.96 d (8.7)	74.0	a 4.04 d (7.6)	66.8
	b 3.89 d (8.7)		b 3.98 d (7.6)	
16		100.9		99.6
17		166.8		180.8
18	6.55 s	95.9	6.55 s	107.5
19		158.8		159.7
20		165.8		162.8
1'		123.2		123.4
2', 6'	7.78 dd (8.9, 2.0)	126.6	7.65 dd (7.2, 2.5)	127.4
3', 5'	7.05 dd (8.9, 2.0)	114.0	6.93 dd (7.2, 2.5)	114.5
4'		162.0		162.1
OMe	3.88 s	54.4	3.83 s	55.6

^aMeasured at 400 MHz in methanol-*d*₄. ^bMeasured at 100 MHz in methanol-*d*₄. ^cMeasured at 500 MHz in CDCl₃. ^dMeasured at 125 MHz in CDCl₃.

The gross structure of **1** was elucidated by analysis of 2D-NMR data including the ¹H–¹H COSY, ROESY, HSQC, and HMBC spectra in CD₃OD. Four proton sequences, H₃-13–H-4–H₃-14, H₂-2–H₂-1, H₂-9–H-11, and H₂-6–H₂-7 were disclosed by the ¹H–¹H COSY spectrum of **1**. The HMBC experiment used to connect the above substructures was found to show the following correlations: H₂-1 to C-10; H₂-6 to C-5; H₂-7 to C-8 and C-11; H₂-9 to C-8, C-11, and C-16; H-11 to C-17 and C-20; H₃-13 and H₃-14 to C-3 and C-4; H₂-15 to C-5, C-8, C-9, and C-10. The key connectivity between C-15 and C-8 through an ether bridge was revealed by an HMBC correlation for H-15a (δ_{H} 3.96) to C-8. The presence of an $\alpha,\beta,\gamma,\delta$ -unsaturated δ -lactone moiety was indicated by HMBC cross-peaks of H-18 (δ_{H} 6.55) to C-19 (δ_{C} 158.8), C-16 (δ_{C} 100.9), and C-1' (δ_{C} 123.2). HMBC correlations for H-2'/H-6' (δ_{H} 7.78) to C-19 (δ_{C} 158.8) revealed the connectivity of C-1' to C-19. A methoxy group (δ_{H} 3.88) was positioned at C-4' (δ_{C} 162.0) based on the HMBC cross-peak between them. Thus, the gross structure of yaminteritrem A was deduced as **1** (Figure 1).

Figure 1. Structures of yaminteritrem A (**1**) and B (**2**).

The relative stereochemistry of **1** was determined by ROESY spectrum. The key ROESY correlation of H-11/H-7a suggested that H-11 and H-7a maintained a pseudo-1,3 diaxial conformation, so that H-11 should be at the opposite end of the fused ether bridge and H-11 should be in the same α orientation as Me-12 (Figure 2 and Supporting Information

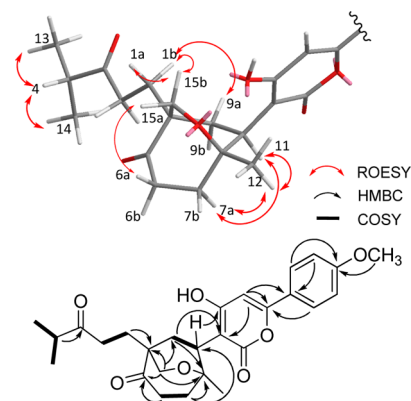
Figure 2. Selected ¹H–¹H COSY, HMBC, and ROESY correlations for compound **1**.

Figure S7). Moreover, the ROESY cross-peaks between H-15b/H-9a, and H-15a/H-6a could also be observed. Besides, due to the free rotation of the C1–C15 bond, the cross-peaks between H-15b/H-1a and H-1b can be observed.

Yaminteritrem B (**2**) was obtained as colorless powders, and its molecular formula was determined to be C₂₇H₃₂O₇ by HRESIMS on the [M + H]⁺ (*m/z* 469.2203, calcd 469.2226 for C₂₇H₃₃O₇), implying 12 degrees of unsaturation. The ¹³C NMR and DEPT spectra of **2** (Table 1) exhibited the presence of 27 carbon resonances, containing one carbonyl carbon (δ_{C} 180.8), five quaternary aromatic carbons, four aromatic methines, one olefinic methine, three oxygen-bearing quaternary carbons (δ_{C} 98.3, 84.0, 74.9), one oxymethylene (δ_{C} 66.8), one quaternary carbon (δ_{C} 46.5), one methine, six methylenes, one methoxyl, and three methyls. The ¹H NMR spectrum of **2** (Table 1) revealed the presence of the same 1,4-disubstituted phenolic moiety (δ_{H} 7.65, dd, *J* = 7.2, 2.5 Hz; 6.93, dd, *J* = 7.2, 2.5 Hz) and an olefinic methine (δ_{H} 6.55, s) as in the case of compound **1**. And, there are one oxymethylene (δ_{H} 4.04, *J* = 7.6 Hz; 3.98, d, *J* = 7.6 Hz), one methoxyl (δ_{H} 3.83), and three singlet methyls (δ_{H} 1.28, 1.14, and 1.10).

The gross structure of **2** was elucidated by analysis of 2D-NMR data including the ^1H - ^1H COSY, ROESY, HSQC, and HMBC spectra in chloroform-*d*. Three proton sequences, H_2 -1- H_2 -2, H_2 -6- H_2 -7, H_2 -9- H_2 -11, were disclosed by the ^1H - ^1H COSY spectrum of **2**. The HMBC experiment used to connect the above substructures was found to show the following correlations: H_2 -2 to C-3 and C-4; H_2 -1 to C-2, C-3, C-10, and C-15; H_2 -6 to C-5, C-7, C-8, and C-10; H_2 -7 to C-5, C-6, C-8, C-9, and C-12; H_2 -9 to C-11, C-12, and C-16; H_2 -11 to C-16 and C-20; H_3 -13 and H_3 -14 to C-3 and C-4; H_2 -15 to C-1, C-3, C-9, and C-10. Key connectivity of C-15 (δ_{C} 66.8) and C-3 (δ_{C} 98.3) through an oxygen atom was revealed by an HMBC correlation for H-15 (δ_{H} 4.04) to C-3. The presence of a pyran-4-one moiety was indicated by HMBC cross-peaks of H-18 (δ_{H} 6.55) to C-19 (δ_{C} 159.7) and C-16 (δ_{C} 99.6). The *p*-methoxyphenyl group at C-19 was established based on the HMBC cross-peaks for the signal at δ_{H} 3.88 to C-4' (δ_{C} 162.0) and H-2'/H-6' (δ_{H} 7.78) to C-19 (δ_{C} 159.7). Thus, the gross structure of yaminteritrem B was deduced as **2** (Figure 1).

The NOESY correlations of **2** between H-11b/H-15b, Me-12/H-15b, and Me-14/H-15b suggested that Me-12 should be the same orientation as CH_2 -15 (Supporting Information Figure S15). The absolute configuration of **2** was established by a single-crystal X-ray diffraction analysis using Cu $K\alpha$ radiation (Figure 3 and Supporting Information Tables S1–S5). The

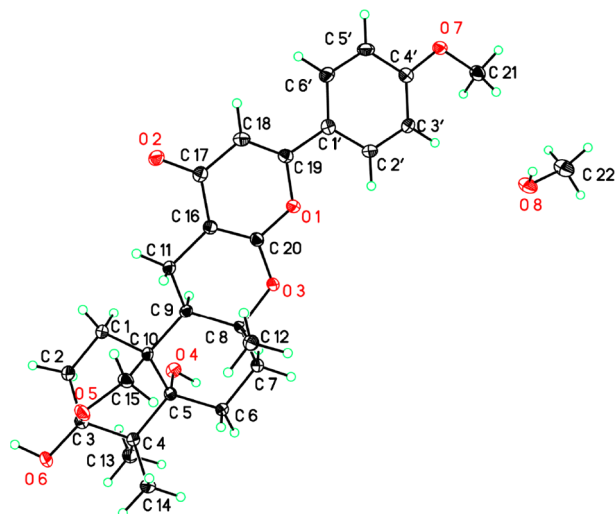
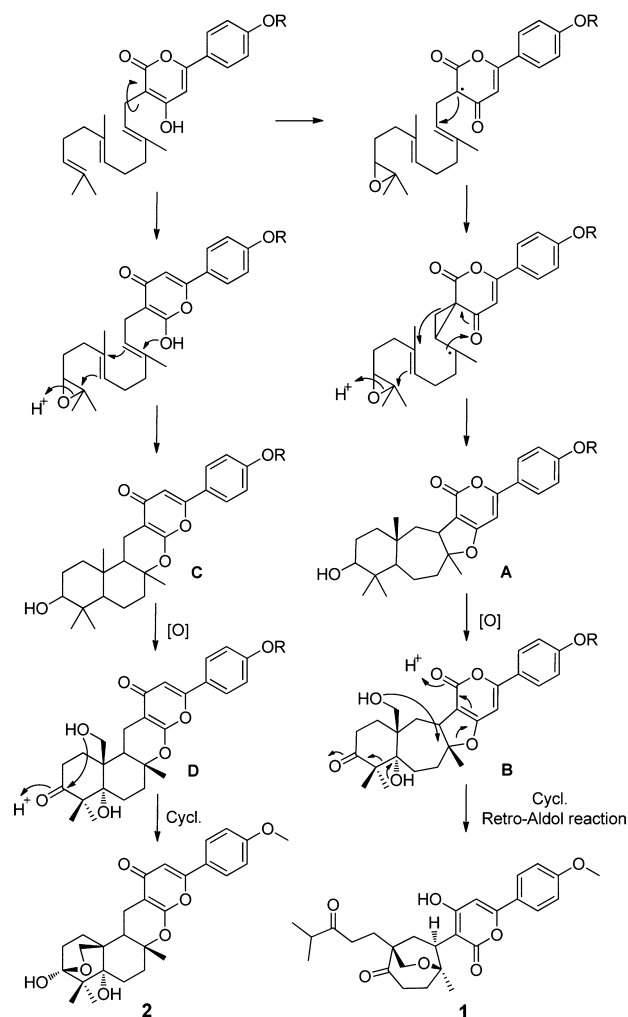


Figure 3. X-ray crystallographic structure of **2** by a single-crystal X-ray diffraction analysis using Cu $K\alpha$ radiation.

result demonstrated that the chiral centers in **2** were 3*S*, 5*R*, 8*R*, and 10*R*. Thus, the structure of **2** was assigned as and named yaminteritrem B.

A biosynthetic origin of yaminteritrems A (**1**) and B (**2**) was proposed as shown in Scheme 1. Both compounds seem to be biogenetically related to some meroterpenoids, such as arisugacin isolated *Penicillium* sp.,^{1c} terreulactone A and teritremes from *A. terreus*,⁴ and pyripyropene from *A. fumigatus*.⁵ Yaminteritrem B (**2**) might be derived from the stereospecific cyclization of the sesquiterpene with a phenyl- α -pyrone moiety through proton-initiated carbocation formation. Yaminteritrem A (**1**) might be produced from the same precursor of compound **2** through intermediates A and B. The biosynthesis pathway of compound **1** might involve the formation of an unusual cycloheptane moiety of the key intermediate B and the cleavage of the C4–C5 bond in the ring

Scheme 1. Plausible Biosynthetic Pathway for **1** and **2**



A of the terpenoid unit of the intermediate B to produce **1** via retro-Aldol reaction.

Epidemiological studies have shown that nonsteroidal anti-inflammatory drugs (NSAIDs) decrease the risk of developing Alzheimer Disease (AD).⁶ Conventional NSAIDs inhibit both isoforms of cyclooxygenase (COX) playing an important role in conversion of arachidonic acid to various prostaglandins (PG). Cyclooxygenase-2 (COX-2) is an inducible isozyme, which promotes cellular proliferation, angiogenesis, cancer invasiveness, and antiapoptosis.⁷ LPS-stimulated RAW264.7 macrophages were treated by yaminteritrem B (**2**) to investigate its anti-inflammation effect. The results indicated that **2** could reduce the LPS-induced COX-2 expression in protein and RNA levels with the EC_{50} value at 18.3 μM (Figure 4). Dexamethasone (DEX) and NS398 were used as possible controls (Figure S19).

In summary, we discovered two novel meroterpenoids with unusual fused-ring skeleton by an ether bridge linkage from the thermophilic *A. terreus*, determined the absolute stereochemistry of compound **2** with a single-crystal X-ray diffraction study, and demonstrated the inhibition of **2** on the COX-2 expression.

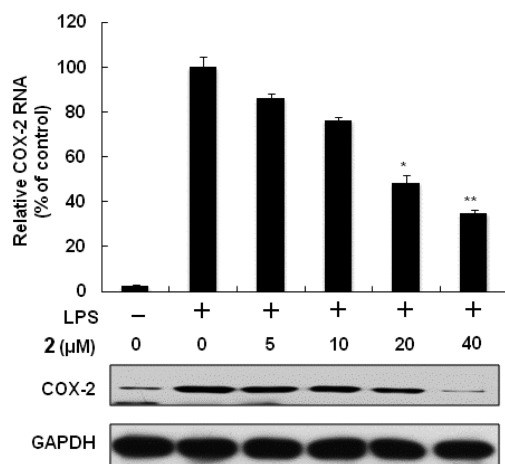


Figure 4. Inhibition effect on COX-2 expression of **2** in LPS-stimulated RAW264.7 macrophages. Cells were seeded in a 24-well plate, treated with **2** at 0, 5, 10, 20, or 40 μ M, and then incubated with 1 μ g/mL LPS for 24 h. The RNA and protein levels of COX-2 were determined by RT-qPCR or Western blotting, respectively. Data were presented as the mean \pm SD from at least triplicate observations. Asterisks indicated a significant difference compared with untreated RAW264.7 macrophages. * $P < 0.05$; ** $P < 0.01$.

■ ASSOCIATED CONTENT

📄 Supporting Information

Extraction and isolation, bioassay method, 1D and 2D selective NMR, HRESI-MS, IR and UV spectra of **1** and **2**, and the X-ray crystallographic analytical method and data for **2**. These materials are available free of charge via the Internet at <http://pubs.acs.org>.

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Notes

The authors declare no competing financial interest.

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