Novel Dibenzo[*b*,*e*]oxepinones from the Freshwater-Derived Fungus *Chaetomium* sp. YMF 1.02105

Authors

Kai-Ze Shen^{1,2}, Suo Gao³, Yan-Xiu Gao^{1,2}, An-Ran Wang¹, Yao-Bo Xu¹, Rong Sun^{1,2}, Pei-Gen Hu¹, Guo-Fu Yang¹, Ai-Ji Li¹, Dan Zhong¹, Hai-Yang Liu³, Jin-Yan Dong^{1,2}

Affiliations

The affiliations are listed at the end of the article

Key words

- dibenzo[b,e]oxepinone
- Chaetomium sp.
- Chaetomiaceae
- chaetones
- cytotoxicity
- antimicrobial activity

received	Dec. 6, 2011
revised	Sept. 8, 2012
accepted	Sept. 14, 2012

Bibliography

DOI http://dx.doi.org/ 10.1055/s-0032-1327828 Published online October 24, 2012 Planta Med 2012; 78: 1837–1843 © Georg Thieme Verlag KG Stuttgart · New York · ISSN 0032-0943

Correspondence

Prof. Dr. Jin-Yan Dong School of Life Science, Southwest University Street Tiansheng 2 Beibei 400715 Chongqing People's Republic of China Phone: + 86 23 68 25 23 65 Fax: + 86 23 68 25 23 65 donjyaa@swu.edu.cn

Correspondence

Dr. Hai-Yang Liu State Key Laboratory of Phytochemistry and Plant Resources in West China Kunming Institute of Botany Chinese Academy of Sciences 132# Lanhei Road, Heilongtan Kunming 650201 People's Republic of China Phone: + 868715223246 Fax: + 868715223445 haiyangliu@mail.kib.ac.cn

Abstract

Six new dibenzo[*b,e*]oxepinone metabolites, chaetones A–F (**1–6**), as well as three known compounds, 1-hydroxy-6-methyl-8-hydroxymethyl-xanthone (**7**), citreorosein (**8**), and emodin (**9**), were obtained from a freshwater-derived fungal strain *Chaetomium* sp. YMF 1.02105. Their structures were established on the basis of extensive spectroscopic data analysis and comparison with spectroscopic data reported. Compounds **1–6** are

Introduction

Fungi are a well-known source of bioactive compounds, and the research for isolation of novel fungal metabolites that blossomed more than 40 years is still very active now [1]. At present, the search for new producers of biologically active compounds is diligently underway among fungi from unusual or specialized ecological niches, because the synthesis of new secondary metabolites and potential biologically active compounds that help them to survive and adapt to these existence conditions can be expected in these fungi with the greatest probability [2-4]. Due to different habitats from those of terrestrial fungi, the freshwater-derived fungi are of special interest. In fact, freshwater-derived fungi are being accepted as a potentially important source of novel bioactive secondary metabolites that might prove suitable for specific medicinal or agrochemical applications as many novel bioactive compounds have been gradually reported from freshwater-derived fungi Annulatascus triseptatus [5]. Dendrospora tenella [6], Decaisnella thyridioides [7], Helicodendron giganteum [8], Kirschsteiniothelia sp. [9], Massarina tunicate [10–12], Ophioceras venezuelense [13], Stachybotrys sp. [14], and others. Our research group has been interested in studies of bioactive compounds produced by freshwaterfurther additions to the small group of dibenzo[*b*, *e*]oxepinones represented by arugosins A–H. Compounds **1–7** were tested for their cytotoxic activities against A549, Raji, HepG2, MCF-7, and HL-60 cell lines. The results showed that compound **3** had significant cytotoxicity with IC₅₀ values of 1.2, 1.8, 1.9, 2.3, and 1.6 µg/mL, respectively, against the five cancer cell lines. All compounds showed modest antimicrobial activity against *Staphylococcus aureus* (ATCC 6538) in standard disk assays.

derived fungi. Previously, we have identified some unique biologically active metabolites from these fungi, such as astropaquinones [15], caryospomycins [16], colelomycerones [17], pseudohalonectrins [18], YMF 1029 A-E [19], and others. During our continuous characterization of structurally unique bioactive metabolites from freshwater-derived fungi, the organic solvent extract of the solid-substrate fermentation culture from Chaetomium sp. YMF 1.02105, which was isolated from the submerged wood collected from the River Bailong in Kunming, Yunnan Province, China, displayed antibacterial activity against Staphylococcus aureus (ATCC 6538) and inhibitory effects on the growth of two human tumor cell lines, A549 and MCF-7. Further investigation led to the discovery of six new dibenzo[b,e]oxepinone metabolites, chaetones A-F (1-6) along with three known compounds, 1-hydroxyl-6-methyl-8-hydroxymethylxanthone (7), citreorosein (8), and emodin (9) (O Fig. 1) [20-22]. We herein report the isolation and structure elucidation of these compounds, as well as their cytotoxic and antimicrobial activities.



Fig. 1 Chemical structures of compounds **1–9** isolated from *Chaetomium* sp. YMF 1.02105.

Materials and Methods

General experimental procedures

UV spectra were taken on a Shimadzu double-beam 210A spectrophometer (Shimadzu). Infrared (IR) spectra were obtained in KBr pellets with a Bio-Rad FTS-135 spectrophotometer (Bio-Rad). The nuclear magnetic resonance (NMR) spectra were recorded on DRX-500 NMR (Bruker) spectrometers, with TMS as an internal standard, and coupling constants were represented in Hertz. EI-MS data were taken on a VG Auto Spec 3000 mass spectrometer (VG). HR-MS (ESI-TOF) data were recorded with an API QSTAR Pulsar 1 spectrometer. Column chromatography was performed on silica gel G (200–300 mesh; Qingdao Marine Chemical Factory) and Sephadex LH-20 (AmershamPharmacia Biotech). Precoated silica gel plates (Qingdao Marine Chemical Factory) were used for TLC. Detection was done by spraying the plates with 5% H₂SO₄, followed by heating.

Fungal material

Chaetomium sp. was isolated by Dr. Hong Zhu from a submerged woody substrate collected from the River Bailong in Kunming, Yunnan Province, China, in April 2006. The isolate was determined to be an unidentified species of Chaetomium by one of the authors (K.Z. Shen) on the basis of morphology and sequence analysis of the ITS region of the rDNA and assigned the accession number YMF 1.02105 in the culture collection at the Laboratory for Conservation and Utilization of Bio-Resources, Yunnan University, Yunnan Province, People's Republic of China. The fungal strain was subcultured on slants of potato dextrose agar (PDA) at 25 °C for 5 days. The agar plugs were used to inoculate 250 mL Erlenmeyer flasks, each containing 50 mL of the seed media (0.4% glucose, 1% malt extract, and 0.4% yeast extract), and the final pH of the media was adjusted to 6.5 before sterilization. Flask cultures were incubated at 25 °C on a rotary shaker (190 rpm) for 5 days. Preparation fermentation was carried out in ten 500-mL Erlenmeyer flasks, each containing 100 g of rice. Distilled H₂O (100 mL) was added to each flask, and the contents were soaked overnight before autoclaving at 15 lb/in². for 30 min. The flasks were cooled to room temperature and individually inoculated with 5.0 mL of the seed culture and incubated at 25 °C under static conditions for 50 days.

Extraction and isolation

Following incubation, 300 mL of methanol was added to each flask. The suspension was filtered and the filtrate concentrated under vacuum at 50 °C. The aqueous residue (55.1 g) was partitioned into petroleum ether (bp 60–90 °C) and EtOAc layers. The EtOAc layer (13.4 g) was loaded onto a silica gel column [1800 g

silica gel G (200-300 mesh), 8.0 cm i.d. × 180 cm] using petroleum ether-EtOAc-CH₃OH gradient elution to afford six fractions (A-F) based on the TLC behavior. Fraction B (914 mg), obtained on elution with petroleum ether-EtOAc (65:35), was further subjected to passage over a Sephadex LH-20 column (3.5 cm i.d. × 180 cm) eluting with MeOH to yield subfractions B-1 and B-2. Subfraction B-1 (28 mg) was further purified by a silica gel column [5 g silica gel G (200-300 mesh), 0.8 cm i.d. × 25 cm] eluting with CHCl₃-CH₃COCH₃ (95:5) to yield **1** (8 mg). Subfraction B-2 (128 mg) was further separated into 2 (30 mg) and 3 (48 mg) by a silica gel column [10 g silica gel G (200-300 mesh), 1.0 cm i.d. × 35 cm] eluting with petroleum ether-CH₃COCH₃ (9:1), and a Sephadex LH-20 (1.6 cm i.d. \times 120 cm) column with CH₃OH. Fraction C (800 mg), obtained on elution with petroleum ether-EtOAc (4:6), was further separated on a Sephadex LH-20 column (2.0 cm i.d. × 150 cm) eluting with MeOH, followed by a silica gel column [10 g silica gel G (200-300 mesh), 1.0 cm i.d. × 35 cm] with 80% petroleum ether-acetone (8:2) to afford 4 (20 mg) and 9 (8 mg). Fraction D (200 mg), obtained on elution with petroleum ether-EtOAc-MeOH (3:7:1), was further subjected to passage over a Sephadex LH-20 column (1.6 cm i.d. × 150 cm) eluting with MeOH to yield subfractions D-1 and D-2. Subfraction D-1 (28 mg) was further purified by preparative silica gel TLC (silica gel H, 20 × 20 × 0.1 cm film thickness) developed with petroleum ether-acetone (7:3) to yield 7 (15 mg). Subfraction D-2 (42 mg) gave 8 (5 mg) after further fractionation over a silica gel column [5 g silica gel G (200-300 mesh), 0.8 cm i.d. × 25 cm] eluted with 70% petroleum ether-EtOAc (7:3). Fraction E (3.8 g), obtained on elution with petroleum ether-EtOAc-MeOH (3:7:3), was further divided into subfractions E-1 to E-4 by passage over a silica gel column [100 g silica gel G (200-300 mesh), 3.0 cm i.d. × 100 cm] with a gradient elution of CHCl₃-CH₃COCH₃ (9:1 \rightarrow 3:2). Subfraction E-2 (113 mg) was further separated over a Sephadex LH-20 gel column (1.6 cm i.d. × 150 cm) eluting with MeOH to give 5 (20 mg). Separation of subfraction E-3 (380 mg) over a silica gel column [20 g silica gel G (200-300 mesh), 1.6 cm i.d.× 40 cm], eluting with petroleum ether-acetone (1:1), afforded 6 (10 mg).

Chaetone A (1): bright yellow solid (CH₃COCH₃); UV (CH₃OH) λ_{max} (log ε) 356.0 (3.55), 280.3 (3.68), 222.0 (4.12) nm; IR (film) ν_{max} 3525, 2949, 2850, 1628, 1597, 1583, 1486, 1448, and 1035 cm⁻¹; EIMS *m/z* (rel. int) 256 [M]⁺ (100), 243 (39), 227 (54), 163 (34), 136 (48), 108 (31); HRMS (ESI-TOF) *m/z*: 279.0627 [M + Na]⁺ (calcd. for C₁₅H₁₂O₄Na, 279.0633); NMR date are given in **© Table 1**.

Chaetone B (**2**): bright yellow solid (CH₃COCH₃); UV (CH₃OH) λ_{max} (log ε) 355.0 (3.56), 282.0 (3.72), 221.5 (4.02), 202.0 (4.20) nm; IR (film) v_{max} 3425, 2959, 2921, 2850, 2839, 1631, 1597,

No.	1		2		3		4		5		6	
	δ _H	δ _C	δ _H	δ _C	δ _H	δ _C	δ _H	δ _C	δ _H	δ _C	δ _H	δ _C
1		163.7		154.0		159.5		153.9		154.5		163.4
1		(s)	2 9E (c)	(S)	2 70 (c)	(s)	2.99(c)	(s)	2.70(c)	(s)		(s)
OCH3			5.65 (5)	(a)	5.79(5)	(a)	5.00(5)	(a)	5.79(5)	(a)		
1-OH	13.50 (s)			(1)		(1)		(1)		(1)	12.20 (s)	
2	6.78 (dd,	112.6	6.73	107.9	6.44 (s)	97.9	6.90	110.5	6.87	109.4	6.33	98.2
	1.8, 9.0)	(d)	(d, 9.0)	(d)		(s)	(d, 8.9)	(d)	(d, 9.1)	(d)	(br. s)	(d)
3	/.46 (± 0.0)	137.4 (d)	7.03 (d. 0. 0)	116.5 (d)		156.2 (c)	7.06 (d. 8.0)	120.7 (d)	7.19 (d. 0.1)	(d)		165./ (c)
3-0H	(1, 9.0)	(u)	(u, 9.0)	(u)	10.30 (s)	(3)	(u, 8.9)	(u)	(u, 9.1)	(u)	8.58 (s)	(3)
4	6.85 (dd,	106.6		144.4		132.8		142.5		145.6	6.62	94.1
	1.8, 9.0)	(d)		(s)		(s)		(s)		(s)	(br. s)	(d)
4-			3.88 (s)	56.9	3.78 (s)	61.8			3.84 (s)	57.2		
0CH ₃ 4-0H				(q)		(q)	8.01			(q)		
4 011							(br. s)					
4a		155.8		149.8		156.3		147.8		150.5		157.7
6	5.24 ()	(s)	5 3 9 ()	(s)	5 22 ()	(s)	5 30 ()	(s)	5 22 ()	(s)	5 33 ()	(s)
6	5.21 (s)	//.5 (t)	5.20 (s)	//.3 (+)	5.23 (s)	//.9 (t)	5.20 (s)	//.8 (t)	5.23 (S)	//.b (t)	5.23 (s)	//.8 (t)
6a		142.1		141.1		141.9		142.4		143.0		143.5
		(s)		(s)		(s)		(s)		(s)		(s)
7	6.68	119.5	6.50	118.3	6.66	119.3	6.64	119.3	6.80	115.6	6.83	116.1
0	(d, 1.6)	(d)	(d, 1.5)	(d)	(d, 1.8)	(d)	(d, 1.6)	(d)	(d, 1.7)	(d)	(d, 1.8)	(d)
δ		147.3 (s)		146.7 (s)		147.1 (s)		147.b (s)		152.0 (s)		152.2 (s)
9	6.82	118.7	6.80	118.2	6.77	118.9	6.84	118.5	6.95	115.1	6.99	115.4
	(d, 1.6)	(d)	(d, 1.5)	(d)	(d, 1.8)	(d)	(d, 1.6)	(d)	(d, 1.7)	(d)	(d, 1.8)	(d)
10		165.3		164.4		165.9		165.1		164.8		165.0
10.011	17.01/2)	(s)	12.02 (*)	(s)	12 71 (a)	(s)	12.01 (a)	(s)	12.02 (*)	(s)	17 11 /2)	(s)
10-OH 10a	12.81 (\$)	118.2	13.02 (S)	117 7	13.71(S)	118 1	13.01 (S)	118 3	12.83 (S)	118.6	12.11(5)	119.0
Tou		(s)		(s)		(s)		(s)		(s)		(s)
11		194.8		194.1		193.8		195.4		197.1		196.8
		(s)		(s)		(s)		(s)		(s)		(s)
11a		113.4		123.7		115.2		124.5		124.8		101.9
12	2 33 (s)	(5)	2 33 (s)	(5)	2 31 (s)	(5)	2 31 (s)	(5)	4 65	(S) 63.6	4 63	(5)
12	2.55 (5)	(q)	2.55 (5)	(q)	2.31(3)	(q)	2.51 (5)	(q)	(br. s)	(t)	(br. s)	(t)
12-OH									4.50		4.47	
									(br. s)		(br. s)	

Table 1 NMR data of compounds 1-6 in CD₃ COCD₃ (δ ppm, / Hz).

1580, 1486, 1455, 1438, 1376, 1337, 1320, 1278, 1260, 1211, 1177, 1095, 1077, 1058, 1035, 974, 958, 908, 839, 813, 766, 742, 718, 674, 616, 584, 571 cm⁻¹; EIMS *m/z* (rel. int) 300 [M]⁺ (62), 285 (9), 269 (100), 254 (24), 241 (18), 229 (31), 214 (7), 169 (5); HRMS (ESI-TOF) *m/z*: 323.0902 [M + Na]⁺ (calcd. for $C_{17}H_{16}O_5Na$, 323.0895); NMR date are given in **• Table 1**.

Chaetone C (**3**): bright yellow solid (CH₃COCH₃); UV (CH₃OH) λ_{max} (log ε) 359.5 (3.53), 279.0 (3.68), 211.0 (4.19) nm; IR (film) v_{max} 3423, 2922, 2851, 1626, 1572, 1533, 1490, 1447, 1376, 1364, 1316, 1265, 1197, 1173, 1123, 1099, 1031, 976, 949, 912, 842, 751, 572 cm⁻¹; EIMS *m/z* (rel. int) 316 [M]⁺ (85), 301 (22), 285 (100), 273 (20), 270 (39), 255 (27), 245 (37), 229 (11), 213 (12), 185 (17), 161 (25); HRMS (ESI-TOF) *m/z*: 339.0829 [M + Na]⁺ (calcd. for C₁₇H₁₆O₆Na, 339.0844); NMR date are given in **○ Table 1**.

Chaetone D (**4**): bright yellow solid (CH₃COCH₃); UV (CH₃OH) λ_{max} (log ε) 355.0 (3.49), 280.0 (3.68), 222.0 (3.91) nm; IR (film) ν_{max} 3419, 2954, 2924, 2852, 2837, 1631, 1599, 1581, 1480, 1447,

1428, 1395, 1360, 1327, 1276, 1228, 1190, 1177, 1092, 1035, 1022, 1012, 980, 961, 916, 862, 828, 801, 747, 725, 711, 672, 601, 573 cm⁻¹; EIMS *m/z* (rel. int) 286 [M]⁺ (70), 271 (8), 255 (100), 227 (27), 215 (49), 187 (11), 149 (30); HRMS (ESI-TOF) *m/z*: 287.0951 [M + H]⁺ and 309.0741 [M + Na]⁺ (calcd. for $C_{16}H_{14}O_5$ 287.0921 and $C_{16}H_{14}O_5$ Na, 309.0738, respectively); NMR date are given in **• Table 1**.

Chaetone E (5): orange yellow solid (CH₃COCH₃); UV (CH₃OH) λ_{max} (log ε) 354.0 (3.43), 278.0 (3.62), 202.0 (4.18) nm; IR (film) ν_{max} 3525, 2926, 2853, 1698, 1666, 1631, 1575, 1485, 1460, 1434, 1402, 1327, 1274, 1259, 1207, 1181, 1031, 1143, 1090, 1067, 968, 912, 840, 818, 786, 548 cm⁻¹; EIMS *m/z* (rel. int) 316 [M]⁺ (57), 285 (100), 270 (20), 245 (25); HRMS (ESI-TOF) *m/z*: 339.0839 [M + Na]⁺ (calcd. for C₁₇H₁₆O₆Na, 339.0844); NMR date are given in **© Table 1**.

Chaetone F (**6**): orange yellow solid (CH₃COCH₃); UV (CH₃OH) λ_{max} (log ε) 356.0 (3.48), 279.3 (3.68), 202.0 (4.18) nm; IR (film) ν_{max} 3518, 2921, 2851, 1687, 1623, 1485, 1459, 1431, 1304, 1255,

Compd.	IC ₅₀ (μg/ml	IC ₅₀ (µg/mL)							
	A549	Raji	HepG2	MCF-7	HL-60				
1	89.2	78.7	95.6	75.6	84.1				
2	50.7	44.5	36.8	63.2	76.1				
3	1.2	1.8	1.9	2.3	1.6				
4	33.5	40.2	28.4	53.1	56.8				
5	14.5	16.7	13.2	25.7	26.4				
6	8.1	7.8	6.7	5.9	9.1				
7	> 100	>100	> 100	> 100	>100				
DDP ^b	2.8	2.1	2.6	2.4	2.1				

 Table 2
 IC₅₀ values of the tested compounds toward different tumor cell lines^a.

^a Each data point was the average value of two experimental measurements ^b DDP, an antitumor agent, was used as a positive control

1213, 1162, and 1028 cm⁻¹; EIMS *m/z* (rel. int) 288 [M]⁺ (26), 272 (35), 257 (100), 242 (5), 217(11); HRMS (ESI-TOF) *m/z*: 311.0523 [M + Na]⁺ (calcd. for C₁₅H₁₂O₆Na, 311.0531); NMR date are given in **© Table 1**.

Bioassay

The purity (>95%) of the tested compounds for the evaluation of the cytotoxic and antimicrobial activities was determined by HPLC.

The cytotoxic activities *in vitro* of compounds **1–7** against A549, Raji, HepG2, MCF-7, and HL-60 tumor cell lines were determined by the MTT assay as reported previously [23]. *Cis*-platinum (DDP; Sigma) was used as a positive control. The results are shown in **• Table 2**.

Antimicrobial assays were conducted using the agar diffusion method according to a literature procedure [24]. The target microbes included *Staphylococcus aureus* (ATCC6538), *Enterococcus faecalis* (ATCC 19433), *Streptococcus mutans* (ATCC 25175), *Aspergillus fumigatus* (ATCC 10894), *Candida albicans* (ATCC 10231), and *Geotrichum candidum* (AS2.498). Ciprofloxacine (for injection, purchased from Yichang Humanwell Pharmaceutical Co. Ltd.) was used as a positive control. Test compounds were absorbed onto individual paper disks (6 mm diameter) at 50 µg/ disk.

Results and Discussion

▼

Chaetone A(1) was isolated as a bright yellow solid. Its molecular formula, C₁₅H₁₂O₄, was determined on the basis of the positive HRESIMS at *m*/*z* 279.0627 [M + Na]⁺, (calcd. 279.0633), and this was supported by the ¹H, ¹³C, and DEPT NMR data. The IR spectrum showed the presence of OH (3525 cm⁻¹) and conjugated carbonyl (1628 cm⁻¹) groups. The UV spectrum showed an absorption band at λ_{max} 356.0, 280.3, and 222.0 nm, which was consistent with the absorptions of a dibenzophenone chromophore as previously reported [20,25]. The ¹H NMR spectra (**C** Ta**ble 1**) showed the signals for an aryl methyl group ($\delta_{\rm H}$ 2.33, s, CH₃-12), an oxygenated methylene ($\delta_{\rm H}$ 5.21, s, CH₂-6), and two hydrogen-bonded phenolic OH protons ($\delta_{\rm H}$ 13.50 and 12.81, OH-1 and OH-10, respectively). These data also showed the presence of five aromatic protons, and their coupling constants as well as COSY couplings indicated that 1 had a 1,2,3-trisubstituted aromatic ring [$\delta_{\rm H}$ 6.78 (dd, I = 1.8, 9.0 Hz, H-2), 7.46 (t, I = 9.0 Hz, H-3), 6.85 (dd, J = 1.8, 9.0 Hz, H-4)] and a 1,2,3,5-tetrasubstituted aromatic ring $[\delta_{H} 6.68 (d, J = 1.6 \text{ Hz}, \text{H-7})$ and 6.82 (d, J = 1.6 Hz, H-)9)]. The ¹³C NMR and DEPT experiments (**C** Table 1) revealed the presence of an aryl methyl group ($\delta_{\rm C}$ 21.6), an aryl ketone carbonyl (δ_{C} 194.8), an oxygenated methylene carbon (δ_{C} 77.5), and 12 aromatic carbon resonances. All of these data are very similar to those of arugosin F [20], except that the ${}^{1}H/{}^{13}C$ signals for the hemiacetal moiety of arugosin F had been substituted for signals of an oxomethylene group in the spectra of 1, which was consistent with the molecular formula. Supporting evidence for this assignment was obtained from the HMBC correlations from OH-1 to C-1, C-2 and C-11a, H-2 to C-11a, C-4 and C-11, H-3 to C-1, C-2, C-4, and C-4a, H-4 to C-2, C-3, and C-11a, H₂-6 to C-4a, C-6a, C-7, C-10a, and C-11, H-7 to C-6a, CH3-12, C-9, and C-10a, H-9 to C-10, CH₃-12, C-10a, and C-7, CH₃-12 to C-7, C-8, and C-9, and finally OH-10 to C-10, C-9, and C-10a (OFig. 2). In addition, the NOESY correlations between protons CH₃-12 and H-9 and H-7, OH-10 and H-9, and OH-1 and H-2 also indicated the positions of the methyl and hydroxyl groups in 1. On the basis of the above evidence, the structure of chaetone A (1) was elucidated as 1,10dihydroxy-8-methyldibenzo[b,e]oxepin-11 (6H)-one.

Chaetone B (2) was isolated as a bright yellow solid and showed similar UV absorptions to those of 1. The molecular formula of 2 was determined to be $C_{17}H_{16}O_5$ by HRESIMS analysis (*m/z*, 323.0902 [M + Na]⁺, calcd. 323.0895). The comparison of the spectroscopic data of 2 with those of 1 (O Table 1) showed similarities except that the trisubstituted phenyl ring C in 1 was replaced by the 1,2,3,4-tetrasubstituted phenyl group in 2, which contained two *ortho*-coupled protons at $\delta_{\rm H}$ 6.73 (d, *J* = 9.0 Hz, H-2) and 7.03 (d, J = 9.0 Hz, H-3) and two methoxy groups ($\delta_{\rm H}/\delta_{\rm C}$ 3.85/56.8 and 3.88/56.9). Furthermore, the positions of two methoxy substitutions in the 1,2,3,4-tetrasubstituted phenyl moiety at C-1 and C-4 were readily elucidated by the HMBC correlations of H₂-6 with C-4a, C-6a, C-7, C-10a, and C-11, OCH₃-1 with C-1 and C-2, H-2 with C-1, C-4, C-4a, C-11a, and C-11, H-3 with C-1, C-4, C-4a, and C-11a, and OCH₃-4 with C-3 and C-4 as well as the NOESY cross-peaks of OCH₃-1 ($\delta_{\rm H}$ 3.85, s) with H-2 ($\delta_{\rm H}$ 6.73, d), and H-3 ($\delta_{\rm H}$ 7.03, d) with OCH₃-4 ($\delta_{\rm H}$ 3.88, s) (\bigcirc Fig. 2). On the basis of the spectroscopic data, chaetone B(2) was elucidated as 10hydroxyl-1,4-dimethoxy-8-methyldibenzo[b,e]oxepin-11(6H)one.

Chaetone C (**3**) was isolated as a bright yellow solid. The molecular weight of **3** was 16 mass units larger than that of chaetone B (**2**), suggesting the presence of an additional hydroxyl group. The ¹H NMR and ¹³C NMR data of **3** (**• Table 1**) clearly indicated that this hydroxyl group was attached at C-3. Supporting evidence for this assignment was obtained from the chemical shifts of C-1 (δ_C 159.5, C), C-2 (δ_C 97.9, CH), C-3 (δ_C 156.2, C), C-4 (δ_C 132.8, C), C-4 (δ_C 156.3, C), and 4-OCH₃ (δ_C 61.8) in **3** [C-1 (δ_C 154.0, C), C-2 (δ_C 107.9, CH), C-3 (δ_C 116.5, CH), C-4 (δ_C 144.4, C), C-4 (δ_C 149.8, C), and 4-OCH₃ (δ_C 56.9) in **2**, respectively]. Moreover, the presence of a hydroxyl group at C-3 was corroborated by the HMBC



Fig. 2 Key HMBC and NOESY correlations for compounds **1–6**.

correlations from H₂-6 to C-4a, C-6a, C-7, C-10a and C-11, OCH₃-1 to C-1, H-2 to C-1, C-3, C-4, C-11 and C-11a, and OCH₃-4 to C-3, C-4, and C-4a as well as the NOESY correlation between OCH₃-1 ($\delta_{\rm H}$ 3.79, s) and H-2 ($\delta_{\rm H}$ 6.44, s) (**•** Fig. 2), but none between OCH₃-4 and H-2. Analysis of 2D NMR data led to the assignment of proton and carbon resonances of **3** (**•** Table 1). On the basis of the spectroscopic data, chaetone C (**3**) was elucidated as 3,10-dihydroxy-1,4-dimethoxy-8-methyldibenzo[*b*,*e*]oxepin-11(6H)-one.

Chaetone D (4), obtained as a bright yellow solid, gave similar ¹H and ¹³C NMR spectroscopic data (**Cable 1**) to those of **2**, except for the absence of a signal due to the methoxy group on position C-4 and the presence of an exchangeable proton signal at $\delta_{\rm H}$ 8.01 (br. s). Analysis of the ¹H and ¹³C NMR spectroscopic data of 4 revealed signals of 14 protons and 16 carbon atoms. The molecular formula of **4** was determined as $C_{16}H_{14}O_5$ by HRMS (ESI-TOF) analysis (*m/z*, 287.0951 [M + H]⁺ and 309.0741 [M + Na]⁺, calcd. 287.0921 and 309.0738, respectively). This suggested that for 4, a hydroxyl group replaced the methoxyl group in the molecule of **2**. The NOESY correlation between the 1-OCH₃ protons ($\delta_{\rm H}$ 3.88, s) with H-2 ($\delta_{\rm H}$ 6.90, d, J=8.9 Hz), but none between 1-OCH₃ protons with H-3 ($\delta_{\rm H}$ 7.06, d, J = 8.9 Hz), confirmed the position of the hydroxyl group at C-4 in 4 (O Fig. 2). The IR and UV spectra of 4 were also similar to those of 2. On the basis of the above spectroscopic data, chaetone D was elucidated as 4,10-dihydroxy-1-methoxy-8-methyldibenzo[b,e]oxepin-11(6H)-one.

Chaetone E (**5**) possessed the same molecular formula ($C_{17}H_{16}O_6$) as that of chaetone C (**3**) and also showed similar UV to those of **3**, suggesting **5** was an isomer of **3**. Analysis of the ¹H, ¹³C and HMQC NMR data for **5** (**• Table 1**) revealed the presence of structural features similar to those found in **2**, except that the aryl methyl group ($\delta_H/\delta_C 2.33/21.7$) in **2** was replaced by the signals for a hydroxymethyl unit [$\delta_H/\delta_C 4.65$ (br.s, H₂-12)/63.6] and an exchangeable proton [$\delta_H 4.50$ (br.s, OH-12)] in **5**, and this observation was supported by HMBC correlations from the new oxygenated methylene protons at $\delta_H 4.65$ (H₂-12) to C-7 ($\delta_C 115.6$), C-8 ($\delta_C 152.0$), and C-9 ($\delta_C 115.1$), and the exchangeable proton at $\delta_H 4.50$ (OH-12) to C-12 ($\delta_C 63.6$) and C-8 (**• Fig. 2**). On the basis of the above spectroscopic data, chaetone E was elucidated as 10-hydroxyl-8-(hydroxymethyl)-1,4-dimethoxydibenzo[*b*,*e*]oxepin-11(6*H*)-one.

Chaetone F (**6**) was isolated as a bright yellow solid, displaying a UV spectrum that had a close similarity to those of **1–5**. The molecular formula of **6** was determined to be $C_{15}H_{12}O_6$ by HRESIMS analysis (*m*/*z*, 311.0523 [M + Na]⁺, calcd. 311.0531), which indicated 10 degrees of unsaturation. On the basis of comparison of

the ¹H NMR and ¹³C NMR data of **6** with those of **5** (**• Table 1**), the 1,2,3,5-tetrasubstituted benzyl system was found to remain in **6**, but another 1,2,3,4-tetrasubstituted benzyl moiety in **5** was replaced by the new 1,2,3,5-tetrasubstituted phenyl moiety [6.62 (br.s, H-4) and 6.33 (br.s, H-2)] in **6**. Furthermore, the substitution patterns of the new phenyl ring were deduced by the HMBC correlations from H₂-6 to C-4a, C-6a, C-7, C-10a, and C-11, OH-1 to C-1, C-11a, and C-2, H-2 to C-1, C-3, C-4, and C-11a, and H-4 to C-4a, C-11a, C-3, and C-2, as well as the chemical shift values of C-1 (δ_C 163.4, C) and C-3 (δ_C 165.7, C) (**• Fig. 2**). On the basis of the spectroscopic data, chaetone F (**6**) was elucidated as 1,3,10trihydroxy-8-(hydroxymethyl)dibenzo[*b*,*e*]oxepin-11(6H)-one.

The other three compounds were identified as 1-hydroxy-6-methyl-8-hydroxymethylxanthone (7), citreorosein (8), and emodin (9) shown in **• Fig. 1**, respectively, by comparison of their spectroscopic data with published values [20–22].

The isolated compounds (1–7) were evaluated for their effects on five tumor cell lines, A549, Raji, HepG2, MCF-7, and HL-60, using the MTT assay. As shown in **O Table 2**, compounds 1–6 displayed in vitro inhibitory activities against the five tumor cell lines to various degrees. Among them, compound 3 showed the most potent cytotoxicity against all evaluated cell lines with IC50 values of 1.2, 1.8, 1.9, 2.3, and 1.6 $\mu g/mL$, respectively, which were even stronger than DDP, while compound 6 exhibited moderate growth inhibition with IC₅₀ values of 8.1, 7.8, 6.7, 5.9, and 9.1 μ g/ mL, respectively. However, the xanthone-type dibenzophenone 7 was inactive (O Table 2). On the basis of the structures of compounds 1-6, the appearance of hydroxyl and methoxyl groups in the above-mentioned compounds markedly increased the activities, suggesting that the hydrophobicity of the dibenzo[*b*,*e*]oxepin core in these compounds may be important to cytotoxic activities for arugosin-type dibenzophenones.

In addition, compounds **2–6** showed activities against *Staphylococcus aureus* (ATCC 6538) when tested at 50 µg/disk in standard disk assays, affording inhibitory zones ranging from 11 to 15 mm (ciprofloxacin: 20 mm zone of inhibition at 50 µg/disk). However, none of these compounds exhibited antibacterial activity against *Enterococcus faecalis* (ATCC 19433) and *Streptococcus mutans* (ATCC 25175) or antifungal activity against *Aspergillus fumigatus* (ATCC 10894), *Candida albicans* (ATCC 10231), and *Geotrichum candidum* (AS2.498) at 50 µg/disk. These bioactivity data could provide valuable information for future synthetic and pharmacologic studies to identify cytotoxic/antimicrobial compounds that are more potent and selective against cancer cells/pathogens.

Chaetones A–F (1–6) are new members of the dibenzo[*b*,*e*]oxepinone class of metabolites, and they are structurally related to the arugosins with arugosin F as the closest precedent [20]. However, 1–6 differ significantly from arugosin F and the other arugosins by the presence of an oxygenated methylene unit rather than a hemiacetal methine moiety. It has been suggested that the known arugosins result from the biological oxidation of anthraquinone/anthrone precursors [26,27] and biosynthetic studies of similar compounds support this hypothesis [28,29]. Additionally, biosynthetic studies suggested that the bi- and tricyclic arugosins and compounds from the xanthone family are biogenetically related. Thus, judging from the fact of the co-occurrence of the dibenzo[b,e]oxepinone metabolites 1-6, xanthone 1-hydroxy-6-methyl-8-hydroxymethylxanthone (7), as well as anthraquinones, citreorosein (8) and emodin (9), the biosynthesis of 1-6 probably proceeds in a manner similar to that of known arugosins.

To our knowledge, the dibenzo[*b*,*e*]oxepinones are rare in the dibenzophenone class of compounds, and only twelve dibenzo[*b*,*e*] oxepinone metabolites have so far been reported from fungi in nature, which include arugosins A-E from Aspergillus variecolor [26], A. rugulosus [30], and A. silvaticus [31], arugosin F from Ascodesmis sphaerospora [20], arugosins G and H from Emericella nidulans var. acristata [32], arugosin I from Penicilium sp. [25], massarinin A from Massarina tunicate [12], pestalone from Pestalotia sp. [33], and leptosphaerin D from Leptosphaeria sp. [34]. These metabolites were commonly reported to possess antibacterial/antifungal/antitumor/cytotoxic activities. Among the known natural dibenzo[b,e]oxepinones, pestalone is probably the most remarkable representative. Besides a moderate in vitro cytotoxicity against various tumor cell lines (mean $GI_{50}=6.0 \mu M$), pestalone was reported to exhibit highly potent antibiotic activity against methicillin-resistant Staphylococcus aureus (MIC = 37 ng/mL) and vancomycin-resistant Enterococcus faecium (MIC = 78 ng/mL). Consequently, pestalone is considered as a particularly promising molecule with antibiotic properties [35-37]. Thus, our finding that the freshwater-derived fungus YMF 1.02105 could produce dibenzo[b,e]oxepinones whose structures, close to that of pestalone, could be of significant biomedical importance. Further studies are required to examine the dibenzo [b,e]oxepinones biosynthetic capacities of this fungus and to obtain sufficient quantities of 1-6 and their analogues in order to fully evaluate their biological activities.

Acknowledgements

 $\pmb{\nabla}$

This work was financially supported by the National Natural Science Foundation of China (20862019 and 31270091), the Special Fund for Basic Scientific Research of Central Colleges, Southwest University (XDJK2009B029; XDJK2010C090), and the Scientific Research Foundation of Southwest University to Dr. J.-Y. Dong (Swu10904).

Conflict of Interest

▼

The authors declare no conflict of interest.

Affiliations

- ¹ School of Life Science, Southwest University, Beibei, Chongqing, People's Republic of China
- ² Laboratory for Conservation and Utilization of Bio-Resources, and Key Laboratory for Microbial Resources of the Chinese Ministry of Education, Yunnan University, Kunming, People's Republic of China
- ³ State Key Laboratory of Phytochemistry and Plant Resources in West China, Kunming Institute of Botany, Chinese Academy of Sciences, Kunming, People's Republic of China

References

- 1 *Claudio A, Raffaele P, Mara F, Antonio E, Angelo V.* Structure-activity relationships of derivatives of fusapyrone, an antifungal metabolite of *Fusarium semitectum.* J Agric Food Chem 2004; 52: 2997–3001
- 2 *Gloer JB.* The chemistry of fungal antagonism and defence. Can J Bot 1995; 73: 1265–1274
- 3 Gloer JB. TOR: environmental and microbial relationships. In: Wicklow DT, Soderstrom BE, editors. TOR-mycota. Heidelberg: Springer Verlag; 1997: 249–268
- 4 *Grabley IS, Thiericke R, Zeeck A.* TOR: the chemical screening approach. In: Grabley S, Thiericke R, editors. TOR-drug discovery from nature. Berlin: Springer Verlag; 1999: 124–148
- 5 Li C, Nitka MV, Gloer JB. Annularins A–H: new polyketide metabolites from the freshwater aquatic fungus Annulatascus triseptatus. J Nat Prod 2003; 66: 1302–1306
- 6 Oh H, Kwon TO, Gloer JB, Marvanová L, Shearer CA. Tenellic acids A–D: new bioactive diphenyl ether derivatives from the aquatic fungus Dendrospora tenella. J Nat Prod 1999; 62: 580–583
- 7 Jiao P, Swenson DC, Gloer JB, Campbell J, Shearer C. Decaspirones A–E, bioactive spirodioxynaphthalenes from the freshwater aquatic fungus Decaisnella thyridioides. J Nat Prod 2006; 69: 1667–1671
- 8 Mudur SV, Swenson DC, Gloer JB, Campbell J, Shearer CA. Heliconols A–C: antimicrobial hemiketals from the freshwater aquatic fungus Helicodendron giganteum. Org Lett 2006; 8: 3191–3194
- 9 *Poch GK, Gloer JB, Shearer CA*. New bioactive metabolites from a freshwater isolate of the fungus *Kirschsteiniothelia* sp. J Nat Prod 1992; 55: 1093–1099
- 10 *Oh H, Gloer JB, Shearer CA.* Massarinolins A–C: new bioactive sesquiterpenoids from the aquatic fungus *Massarina tunicata*. J Nat Prod 1999; 62: 497–501
- 11 *Oh H, Swenson DC, Gloer JB, Shearer CA*. Massarilactones A and B: novel secondary metabolites from the fresh-water aquatic fungus *Massarina tunicata*. Tetrahedron Lett 2001; 42: 975–977
- 12 Oh H, Swenson DC, Gloer JB, Shearer CA. New bioactive rosigenin analogues and aromatic polyketide metabolites from the freshwater aquatic fungus Massarina tunicate. J Nat Prod 2003; 66: 73–79
- 13 *Reátegui RF, Gloer JB, Campbell J, Shearer CA.* Ophiocerins A–D and ophioceric acid: tetrahydropyran derivatives and an africane sesquiterpenoid from the freshwater aquatic fungus *Ophioceras venezuelense.* J Nat Prod 2005; 68: 701–705
- 14 *Xu XM, De Guzman FS, Gloer JB, Shearer CA.* Stachybotrins A and B: novel bioactive metabolites from a brackish water isolate of the fungus *Stachybotrys* sp. J Org Chem 1992; 57: 6700–6703
- 15 Wang L, Dong JY, Song HC, Shen KZ, Wang LM, Sun R, Wang CR, Gao YS, Li GH, Li L, Zhang KQ. Three new naphthoquinone pigments isolated from the freshwater fungus, Astrosphaeriella papuana. Planta Med 2009; 75: 1–5
- 16 Dong JY, Zhu YH, Song HC, Li R, He HP, Liu HY, Huang R, Zhou YP, Wang L, Cao Y, Zhang KQ. Nematicidal resorcylides from the aquatic fungus Caryospora callicarpa YMF1.01026. J Chem Ecol 2007; 33: 1115–1126
- 17 Dong JY, Song HC, Li JH, Wang CR, Tang YS, Sun R, Shen KZ, Wang LM, Zhou YP, Zhang KQ. Unusual 1,2-naphthodiketones from a fresh-water fungus YMF 1.01029. Chem Biodivers 2009; 6: 569–577
- 18 Dong JY, Zhou YP, Li R, Zhou W, Li L, Zhu YH, Huang R, Zhang KQ. New nematicidal azaphilones from the aquatic fungus *Pseudohalonectria adversaria* YMF 1.01019. FEMS Microbiol Lett 2006; 264: 65–69
- 19 Dong JY, Song HC, Li JH, Tang YS, Sun R, Wang L, Zhou YP, Wang LM, Shen KZ, Wang CR, Zhang KQ, Ymf 1029 A–E, preussomerin analogues from a fresh-water-derived fungus YMF1.01029. J Nat Prod 2008; 71: 952–956
- 20 Hein SM, Gloer JB, Koster B, Malloch D. Arugosin F: a new antifungal metabolite from the coprophilous fungus Ascodesmis sphaerospora. J Nat Prod 1998; 61: 1566–1567

- 21 Yang YC, Lim MY, Lee HS. Emodin isolated from Cassia obtusifolia (Leguminosae) seed shows larvicidal activity against three mosquito species. J Agric Food Chem 2003; 51: 7629–7631
- 22 Murakami H, Kobayashi J, Masuda T, Morooka N, Ueno Y. ω-Hydroxymodin, a major hepatic metabolite of emodin in various animals and its mutagenic activity. Mutat Res 1987; 180: 147–153
- 23 Denizot F, Lang R. Rapid colorimetric assay for cell grow and survival modifications to the tetrazolium dye procedure giving improved sensitivity and reliability. J Immunol Methods 1986; 89: 271–277
- 24 Wicklow DT, Joshi BK, Gamble WR, Gloer JB, Dowd PF. Antifungal metabolites (Monorden, Monocillin IV, and Cerebrosides) from Humicola fuscoatra traaen NRRL 22980, a mycoparasite of Aspergillus flavus sclerotia. Appl Environ Microbiol 1998; 64: 4482–4484
- 25 *Lin ZJ, Zhu TJ, Fang YC, Gu QQ, Zhu WM*. Polyketides from *Penicillium* sp. JP-1, an endophytic fungus associated with the mangrove plant *Aegice-ras corniculatum*. Phytochemistry 2008; 69: 1273–1278
- 26 Chexal KK, Holker JS, Simpson TJ. The biosynthesis of fungal metabolites. Part VI. Structures and biosynthesis of some minor metabolites from variant strains of Aspergillus variecolor. J Chem Soc [Perkin I] 1975: 549–554
- 27 *Turner WB, Aldridge DC.* Fungal metabolites II. New York: Academic Press; 1983: 156–167
- 28 Holker JSE, Lapper RD, Simpson TJ. The biosynthesis of fungal metabolites. Part IV. Tajixanthone: ¹³C-nuclear magnetic resonance spectrum and feedings with [1⁻¹³C]- and [2⁻¹³C]-acetate. J Chem Soc [Perkin I] 1974: 2135–2140

- 29 Nishida H, Tomoda H, Okuda S, Omura S. Biosynthesis of purpactin A. J Org Chem 1992; 57: 1271–1274
- 30 Ballantine JA, Francis DJ, Hassall CH, Wright JLC. The biosynthesis of phenols. Part XXI. The molecular structure of arugosin, a metabolite of a wild-type strain of Aspergillus rugulosus. J Chem Soc C 1970: 1175–1182
- 31 Kawahara N, Nozawa K, Nakajima S, Kawai K. Studies on fungal products. Part 15. Isolation and structure determination of arugosin E from Aspergillus silvaticus and cycloisoemericellin from Emericella striata. J Chem Soc [Perkin I] 1988: 907–911
- 32 Kralj A, Kehraus S, Krick A, Eguereva E, Kelter G, Maurer M, Wortmann A, Fiebig H, Konig GM. Arugosins G and H: prenylated polyketides from the marine-derived fungus Emericella nidulans var. acristata. J Nat Prod 2006; 69: 995–1000
- 33 *Cueto M, Jensen PR, Kauffman C, Fenical W, Lobkovsky E, Clardy J.* Pestalone, a new antibiotic produced by a marine fungus in response to bacterial challenge. J Nat Prod 2001; 64: 1444–1446
- 34 *Lin J, Liu S, Sun B, Niu S, Li E, Liu X, Che Y.* Polyketides from the ascomycete fungus *Leptosphaeria* sp. J Nat Prod 2010; 73: 905–910
- 35 Rahman H, Austin B, Mitchell WJ, Morris PC, Jamieson DJ, Adams DR, Spragg AM, Schweizer M. Novel anti-infective compounds from marine bacteria. Mar Drugs 2010; 8: 498–518
- 36 Scherlach K, Hertweck C. Triggering cryptic natural product biosynthesis in microorganisms. Org Biomol Chem 2009; 7: 1753–1760
- 37 *Villa FA, Gerwick L.* Marine natural product drug discovery: leads for treatment of inflammation, cancer, infections, and neurological disorders. Immunopharm Immunother 2010; 32: 228–237