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Cytotoxic metabolites from the endophytic fungus *Chaetomium globosum* 7951†

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The following compounds were isolated from acetate extracts of *Chaetomium globosum* 7951 solid cultures: demethylchaetocochin C (**1**) and chaetoperazine A (**3**), two new epipolythiodioxopiperazine (ETP) alkaloids, a novel pyridine benzamide, 4-formyl-*N*-(3'-hydroxypyridin-2'-yl) benzamide (**6**), and three known ETP derivatives (**2**, **4**, and **5**). The structures of these compounds were determined using extensive spectroscopic data analysis. Compounds **1–3**, and **6**, inhibited the growth of MCF-7, MDA-MB-231, H460 and HCT-8 cells with an IC₅₀ of 4.5 to 65.0 μM.

Introduction

Novel bioactive secondary metabolites have been identified in endophytic fungi.^{1–3} The *Chaetomium* genus, belonging to the Chaetomiaceae family, contains more than 100 species derived from terrestrial and marine habitats.⁴ *Chaetomium globosum* is a species of the *Chaetomium* genus. The isolation of the cytotoxic chaetoglobosins A and B was reported by Sekita and coworkers in 1973.⁵ Since then more than 200 metabolites including chaetoglobosins, diketopiperazines, tetramic acids, bis(3-indolyl)-benzoquinones, azaphilones, pyranones, xanthenes, anthraquinones, orsellides, steroids, and terpenoids were identified in *C. globosum* cultures. Some of these metabolites exhibit cytotoxic, antibacterial, anti-malarial, and antiviral activities.⁶

Epipolythiodioxopiperazine (ETP) alkaloids, with either polysulphide bridges or thiomethyl groups, represent an important family of bioactive secondary metabolites, which are toxic to cancer cell lines.⁷ About 20 ETPs have been identified in the *Chaetomium* genus.⁶ During our search for novel and bioactive compounds from microorganisms,^{8–10} we identified the endophytic fungus, *Chaetomium globosum* 7951, which has cytotoxic activity towards human breast cancer cell lines. *Chaetomium globosum* 7951 comes from the root of *Panax notoginseng*, a traditional Chinese medicine.

Chemical investigations of the solid fermentation of the *Chaetomium globosum* 7951 strain led to the identification of 2 new ETP alkaloids, 3 known analogs, and a new pyridine benzamide. Herein, the isolation, structural determination, and cytotoxicity of these compounds are described.

Results and discussion

The molecular formula of compound **1**, a white amorphous powder, is C₃₂H₃₄N₆O₆S₄ according to the (+)-HRESIMS data, with 19 degrees of unsaturation. The IR spectrum displayed the hydroxy or amino (3359 cm⁻¹), methyl (2921 cm⁻¹), and carbonyl (1680 cm⁻¹) functionalities. The ¹H NMR spectrum (Table 1) showed two *ortho*-disubstituted benzene rings at δ_H 6.72 (1H, d), 6.77 (1H, t), 7.03 (1H, t), 7.08 (1H, t), 7.17 (1H, t), 7.22 (1H, d), 7.50 (1H, d) and 7.63 (1H, d), and a trisubstituted double bond at δ_H 7.07 (1H, s) in the lower field. In addition, four methylene groups (δ_H 3.61, 3.04, 4.04, 3.21, 4.21, 4.32, 3.73, and 3.43), one methine group (δ_H 6.07), and four isolated methyl groups (δ_H 2.12, 2.28, 2.77, and 3.11) were observed in the higher field. According to the ¹³C NMR and DEPT spectra analyses, in addition to the structural features above, there were also four carbonyls at δ_C 161.1, 164.3, 165.3, and 165.6, and five quaternary carbons at δ_C 65.0, 72.9, 73.2, 73.7, and 77.2. The spectral data, combined with the molecular formula, suggested that compound **1** is an analog of epipolythiodioxopiperazine. Extensive analysis of the NMR data indicates similarities in chemical shifts to chaetocochin C,¹¹ including the absence of one methyl group in **1**. HMBC correlations of NH-5' with C-1', C-3', C-4', C-5', and C-6', and of N-CH₃-2' with C-1', and C-3' indicated that N-CH₃-5' in chaetocochin C was replaced by NH in **1**. The ROESY correlations of H-5 with H-9', and of 3'-S-Me with 6'-S-Me, combined with the CD effects at λ_{max} nm (Δε) 240 (+7.2), 274 (−0.89) and 305 (+3.6),¹¹ and based on the similar biogenetic perspective of chaetocochin C, which revealed the absolute configuration of **1** was shown in Fig. 1.

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Table 1 NMR spectroscopic data of **1** and **3**^a

1			3		
No.	δ_{H}	δ_{C} , type	No.	δ_{H}	δ_{C} , type
1		165.3, C	1		162.7, C
2-N-Me	3.11, s	27.7, CH ₃	2-N-OMe	3.65, s	61.0, CH ₃
3		77.2, C	3	4.36, t (2.4)	63.5, CH
4		161.1, C	4		165.1, C
5	6.07, d (1.8)	80.0, CH	NH-5	8.58, s	
6a		149.4, C	6		66.5, C
7	6.72, d (7.8)	110.1, CH	6-S-Me	2.13, s	12.5, C
8	7.17, t (7.8)	130.5, CH	7	3.58, d (14.4)	33.7, CH ₂
9	6.77, t (7.8)	118.8, CH		3.20, d (14.4)	
10	7.50, d (7.8)	125.9, CH	8		107.2, C
10a		126.9, C	9	7.19, d (2.4)	125.0, CH
10b		73.2, C	NH-10	10.90, s	
11	4.04, d (15.6)	42.1, CH ₂	10a		135.6, C
	3.21, d (15.6)		11	7.28, d (7.8)	111.1, CH
12		73.7, C	12	7.02, t (7.8)	120.7, CH
13	4.32, dd (12.6, 4.8)	58.7, CH ₂	13	6.93, t (7.8)	118.3, CH
	4.21, dd (12.6, 6)		14	7.58, d (7.8)	118.9, CH
OH-13	5.93, t (6)		14a		127.9, C
1'		165.6, C	15	3.58, ov ^b	58.5, CH ₂
2'-N-Me	2.77, s	28.4, CH ₃	OH-15	4.93, t (5.4)	
3'		72.9, C			
3'-S-Me	2.12, s	12.3, CH ₃			
4'		164.3, C			
NH-5'	9.06, s				
6'		65.0, C			
6'-S-Me	2.28, s	13.9, CH ₃			
7'	3.61, d (15.4)	33.9, CH ₂			
	3.04, d (15.4)				
8'		107.4, C			
9'	7.07, s	126.8, CH			
10'a		133.3, C			
11'	7.22, d (7.8)	110.6, CH			
12'	7.08, t (7.8)	121.4, CH			
13'	7.03, t (7.8)	119.1, CH			
14'	7.63, d (7.8)	120.0, CH			
14'a		130.3, C			
15'	3.73, dd (10.8, 6)	62.8, CH ₂			
	3.43, dd (10.8, 4.8)				
OH-15'	4.90, t (6)				

^a NMR data (δ) were measured at 600 MHz for ¹H and at 150 MHz for ¹³C in DMSO-*d*₆. The assignments were based on ¹H-¹H COSY, HSQC, and HMBC experiments. ^b *J*-value was not determined due to overlapped signals.

The molecular formula of compound **3** is C₁₆H₁₉N₃O₄S according to the HRESIMS data. The IR spectrum displayed absorptions bands at 3393, 3194, 2921, and 1675 cm⁻¹, suggesting the presence of amino or hydroxyl, methyl, and carbonyl groups. The ¹H-NMR spectrum (Table 1) showed 3-substituted indole moiety signals at δ_{H} 6.94 (1H, t), 7.02 (1H, t), 7.19 (1H, d), 7.29 (1H, d) and 7.58 (1H, d), two methylene groups at δ_{H} 3.58 (1H, d), 3.20 (1H, d), and 3.56 (2H, m), a methine group at δ_{H} 4.36 (1H, t), one isolated methyl at δ_{H} 2.13 (3H, s), and a methoxyl group at 3.65 (3H, s). In addition, two carbonyls at δ_{C} 162.7 and 165.1, and a quaternary carbon at δ_{C} 66.7 were identified *via* the ¹³C-NMR spectrum. HMBC correlations of H-7 with C-7, C-8, and C-10a; NH-5 (δ_{H} 8.58) with C-1, C-6, and C-7; and S-Me-6 with C-6, revealed an α -S-methyl-substituted tryptophan residue. In addition, ¹H-¹H COSY relationships between

H-3/H₂-15/OH-15, in combination with the HMBC relationships between H-3 and H-15 with C-16, indicate a serine residue. Meanwhile, the association of NH-5 with C-3 and H-3 with C-1 in the HMBC spectrum suggests that the serine and tryptophan residues form a diketopiperazine ring. Finally, the methoxyl group is located at N-3, as indicated by the molecular formula and the chemical shift at δ_{C} 61.0. Thus, compound **3** was proposed as shown in Fig. 2. DP4+ analysis of the ¹H and ¹³C NMR data indicates 3*S**,6*S**-**3** appeared agreement with the experimental NMR data with 100% probability (Tables S3–S7†).^{12,13} Based on the common biosynthetic origin, the absolute configuration at C-3 and C-6 is probably to be the same as cyclo-L-Trp-L-Ser.¹⁴ In addition, the calculated optical rotation (OR) value⁸ (+56.1) of (3*S*, 6*S*)-**3** (Table S8†) is similar to the experimental OR value (+80.0), which supports the above speculation.



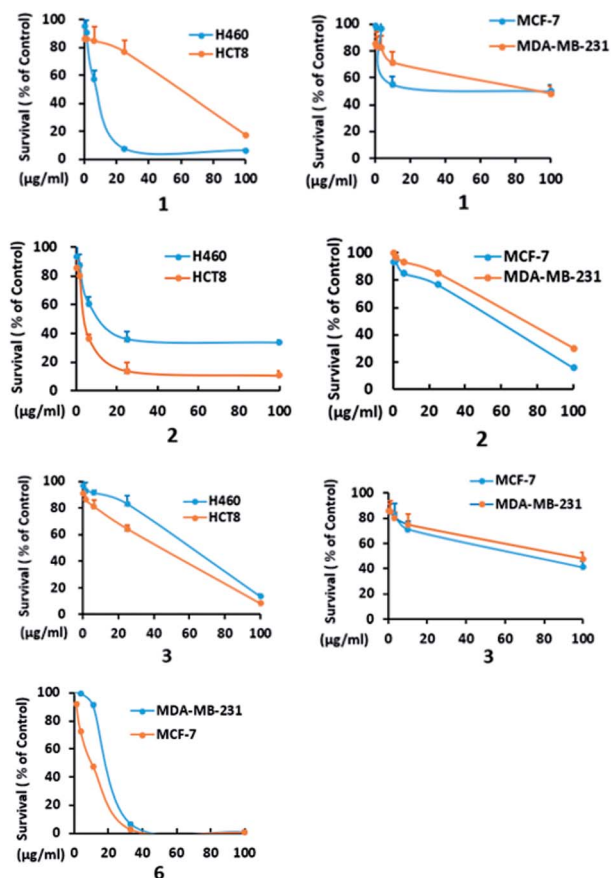


Fig. 3 H460, HCT8, MCF-7 and MDA-MB-231 cells were treated with the indicated concentrations of 1–3 and 6, and cell survival was detected by the CCK8 assay. A dose-dependent curve was depicted.

Yunnan Province, China, in 2015. The strain, which was assigned the accession no. SUB5310227, was identified using nuclear 18S rDNA sequences (GenBank: MK625020) and deposited in the Microbiology Laboratory at Shenyang Pharmaceutical University. *C. globosum* 7951, an endophytic fungus, was grown on PDA at 26 °C for 6 days, and then flushed (sterilized water) into 250 ml Erlenmeyer flasks containing 50 g rice and autoclaved (121 °C, 30 min). The fermentation was then incubated at 26 °C for 40 days.

Extraction and isolation

The cultures (10 kg) were extracted three times with methanol and then filtered. The filtrate was concentrated and three extractions were performed with equal volumes of EtOAc. The EtOAc layer was evaporated with reduced pressure resulting in a crude broth extract (12.2 g). The extract was separated into 15 fractions (A–O) using silica gel column chromatography with a CH₂Cl₂/MeOH gradient elution. Fraction H was purified using Sephadex LH-20 gel column chromatography with CH₂Cl₂/MeOH (1 : 1), resulting in subfractions H1–H8. Subfraction H5 was subjected to semi-preparative HPLC with 50% acetonitrile elution into 0.1% trifluoroacetic acid to isolate compound 1. Fraction J was purified by ODS C₁₈ with a gradient of methanol in water (10–100%) to give six subfractions (J1–J6). Subfraction

J3 was subjected to preparative TLC using CH₂Cl₂/MeOH (20 : 1) and then subjected to semi-preparative HPLC with 30% acetonitrile/H₂O (0.1% CF₃COOH) as the mobile phase to generate compound 6. Fraction K was separated into fractions using silica gel CC with CH₂Cl₂/MeOH (50 : 1), resulting in fractions K1–K3. Fraction K1 was purified with semi-preparative HPLC with 19% acetonitrile into an aqueous 0.1% trifluoroacetic acid solution, resulting in compound 3.

Demethylchaetocochin C (1). White amorphous powder; [α]_D²⁰ 61.0 (*c* 0.3, MeOH); UV (MeOH) λ_{\max} 219, 286 nm; IR ν_{\max} 3359, 3193, 2921, 2851, 1680, 1468, 1425, 1207, 1140, 1061, 1027, 722 cm⁻¹; ¹H NMR (DMSO-*d*₆, 600 MHz) data and ¹³C NMR (DMSO-*d*₆, 150 MHz) data, see Table 1. (+)-HR-ESIMS *m/z* 727.1499 [M + H]⁺ (calcd for C₃₂H₃₅N₆O₆S₄, 727.1501).

Chaetoperazine A (3). White amorphous powder; [α]_D²⁰ 80.0 (*c* 0.3, MeOH); UV (MeOH) λ_{\max} 199, 273 nm; IR ν_{\max} 3393, 3194, 2922, 2850, 1675, 1424, 1205, 1141, 801, 749, 723 cm⁻¹; ¹H NMR (DMSO-*d*₆, 600 MHz) data and ¹³C NMR (DMSO-*d*₆, 150 MHz) data, see Table 1. (+)-HR-ESIMS *m/z* 372.0994 [M + Na]⁺ (calcd for C₁₆H₁₉N₃O₄NaS, 372.0994).

4-Formyl-N-(3'-hydroxypyridin-2'-yl) benzamide (6). White amorphous powder; UV (MeOH) λ_{\max} 254, 325 nm; IR ν_{\max} 3388, 3189, 2921, 2850, 1690, 1621, 1556, 1453, 1387, 1320, 1210, 1142, 1052, 1029, 1011, 838, 724 cm⁻¹; ¹H NMR (DMSO-*d*₆, 600 MHz) data and ¹³C NMR (DMSO-*d*₆, 150 MHz) data, see Table 2. (+)-HR-ESIMS *m/z* 243.0769 [M + H]⁺ (calcd for C₁₃H₁₁N₂O₃, 243.0770).

ORs and NMR calculation of 3

See the ESI.†

Cytotoxicity assay

A CCK colorimetric assay was used to measure the cytotoxicity of compounds 1–6 was in human breast adenocarcinoma cell (MCF-7 and MDA-MB-231), human large cell lung cancer cell (H460) and human cecal adenocarcinoma cell (HCT-8). All of the cell lines were obtained from ATCC. Cells (5 × 10³ cells per mL) were added to 96-well culture dishes and grown for 24 h (5% CO₂, 37 °C) followed by the addition of fresh medium (100 µL) and the test compound. After an additional 48 h, the media was removed and fresh media with 10% CCK solution was added. The cells were incubated for 1 h (37 °C) and then the optical density at 450 nm was determined. Each assay was replicated six times. IC₅₀ values for each cell line were determined.

Conclusions

In conclusion, two new ETPs alkaloids (1, and 3), a new pyridine benzamide (6), and three known ETPs compounds were identified in the endophytic fungus *Chaetomium globosum* 7951. These new compounds moderately inhibit the human breast cancer cells (MCF-7 and MDA-MB-231) and human ileocecal adenocarcinoma (HCT-8) growth. The new compound 1 significantly exhibits cytotoxic against the human lung cancer cell (H460).



Conflicts of interest

There are no conflicts to declare.

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