

PAPER

View Article Online
View Journal | View Issue



Cite this: *Org. Biomol. Chem.*, 2024, **22**, 4521

Neuroprotective azaphilones from a deep-sea derived fungus *Penicillium* sp. SCSIO41030†

Weihaio Chen,^{‡a,c} Jiahui Jiang,^{‡b} Xiaoyan Pang,^a Yingying Song,^{a,c} Zhiyou Yang,^{*b} Junfeng Wang^{ID} ^{*a,c,d} and Yonghong Liu^{*a,c,d}

Ten azaphilones including one pair of new epimers and three new ones, penineulones A–E (**1–5**) with the same structural core of angular deflectin, were obtained from a deep-sea derived *Penicillium* sp. SCSIO41030 fermented on a liquid medium. Their structures including absolute configurations were elucidated using chiral-phase HPLC analysis, extensive NMR spectroscopic and HRESIMS data, ECD and NMR calculations, and by comparing NMR data with literature data. Biological assays showed that the azaphilones possessed no antitumor and anti-viral (HSV-1/2) activities at concentrations of 5.0 μ M and 20 μ M, respectively. In addition, azaphilones **8** and **9** showed neuroprotective effects against A β _{25–35}-induced neurotoxicity in primary cultured cortical neurons at a concentration of 10 μ M. Azaphilones **8** and **9** dramatically promoted axonal regrowth against A β _{25–35}-induced axonal atrophy. Our study indicated that azaphilones could be promising lead compounds for neuroprotection.

Received 9th April 2024,
Accepted 10th May 2024
DOI: 10.1039/d4ob00586d
rsc.li/obc

Introduction

Azaphilones are an enormous family with diverse structures of fungal secondary polyketide metabolites, which possess a highly oxygenated isochroman scaffold containing a pyrone–quinone bicyclic core and a quaternary carbon center. According to the latest review on azaphilones,^{1,2} up to now, over 625 azaphilones have been discovered from more than 20 genera of fungi, such as *Penicillium*,³ *Hypoxylon*,⁴ *Pestalotiopsis*,⁵ *Chaetomium*,⁶ and so on. Besides, most azaphilone molecules exhibit a wide range of significant biological

activities including enzyme inhibition,⁷ antimicrobial,⁸ cytotoxic, antiviral, antioxidative, and anti-inflammatory activities,² due to which they attract lots of attention continuously from chemists and pharmacologists. Recently, during our ongoing search for bioactive molecules from marine-derived fungi,^{9,10} a chemical investigation of a deep-sea derived *Penicillium* sp. SCSIO41030, fermented on a rice medium, resulted in the isolation and identification of two new *p*-terphenyl derivatives with antiviral activity against herpes simplex virus and one novel 4,5-diphenyl-2-pyrone.¹¹ And further study of the strain SCSIO41030 fermented on a liquid medium led to the discovery of ten azaphilones with the same structural type of angular deflectin,² including one pair of new epimers and three new compounds, penineulones A–E (**1–5**), along with five known analogues. Herein, the details of the isolation, structural identification and biological activities of compounds **1–10** are described.

Results and discussion

After the purification of the EtOAc extract from the culture broth of *Penicillium* sp. SCSIO41030, the ¹³C NMR data of compound **P1** were found to possess two sets of similar carbon shift signals (Fig. S1†). Therefore, **P1** was subsequently subjected to separation on a chiral column of DAICEL CHIRALPAK IC to afford enantiomeric compounds penineulone A (**1**) (t_R = 12.5 min) and penineulone B (**2**) (t_R = 16.1 min) in a ratio of 1 : 0.8, using ethanol/*n*-hexane (v/v: 55/65, flow rate 1 mL min^{–1}) with 0.1% trifluoroacetic acid (TFA) as the mobile

^aCAS Key Laboratory of Tropical Marine Bio-resources and Ecology/Guangdong Key Laboratory of Marine Materia Medica/Innovation Academy of South China Sea Ecology and Environmental Engineering, South China Sea Institute of Oceanology, Chinese Academy of Sciences, Guangzhou 510301, China.

E-mail: wangjunfeng@scsio.ac.cn, yonghongliu@scsio.ac.cn

^bCollege of Food Science and Technology, Guangdong Provincial Key Laboratory of Aquatic Product Processing and Safety, Guangdong Province Engineering Laboratory for Marine Biological Products, Guangdong Provincial Engineering Technology Research Center of Seafood, Key Laboratory of Advanced Processing of Aquatic Product of Guangdong Higher Education Institution, Guangdong Ocean University, Zhanjiang 524088, China. E-mail: zyyang@gdou.edu.cn

^cUniversity of Chinese Academy of Sciences, 19 Yuquan Road, Beijing 100049, China

^dSanya Institute of Marine Ecology and Engineering, Yazhou Scientific Bay, Sanya 572000, China

†Electronic supplementary information (ESI) available: Copies of ¹H and ¹³C NMR spectra of all new products, 2D NMR spectra of selected compounds, UV and MS data, and energies of all calculated conformers. CCDC 2214106 and 2214107. For ESI and crystallographic data in CIF or other electronic format see DOI: <https://doi.org/10.1039/d4ob00586d>

‡These authors have contributed equally to this work.



phase. Compounds **1** and **2** were isolated as a light-yellow oil and have the same molecular formula $C_{19}H_{24}O_6$ with eight degrees of unsaturation deduced using the HRESIMS data and ^{13}C NMR data (Table 1). The 1D NMR data along with the HSQC experiment of **1** showed the presence of three aromatic/olefinic methines at $\delta_{C/H}$ 146.2/9.36 (s, CH-1), 107.5/6.01 (s, CH-3), and 103.7/5.41 (s, CH-4); three sp^3 methines at $\delta_{C/H}$ 54.7/4.30 (d, $J = 7.7$ Hz, CH-10), 48.6/2.66 (q, $J = 6.9$ Hz, CH-12), and 45.5/3.65 (d, $J = 7.7$ Hz, CH-7); five methyls at $\delta_{C/H}$ 53.0/3.76 (s, CH_3 -19), 21.7/1.14 (s, CH_3 -16), 19.4/2.14 (s, CH_3 -15), 16.6/1.18 (d, $J = 6.9$ Hz, CH-18), and 11.4/0.86 (t, $J = 6.5$ Hz, CH-14); and one methylene at $\delta_{C/H}$ 25.3/1.77 and 1.39 (m, CH_2 -13). Besides the above twelve corresponding hydrogen-bearing carbons, seven more carbons were present in the ^{13}C NMR spectrum, including three carbonyls (one ester group), three olefinics (one oxygenated), and one oxygenated tertiary carbon. The aforementioned NMR data resembled those of chermesinone B (**8**),¹² an azaphilone derivative obtained from a mangrove endophytic fungus *Penicillium chermesinum* (ZH4-E2). The main differences were one additional methoxyl group [$\delta_{C/H}$ 53.0/3.76 (s, CH_3 -19)] and one less degree of unsaturation in **1**, which indicated that the angular γ -lactone ring of the azaphilone skeleton was opened. The above deduction was further verified using key HMBC signals from H_3 -19 and H-10 to C-17. Detailed analysis of the COSY and HMBC data (Fig. 2) allowed the assignments of all carbon and proton resonances in **1**.

The NMR data of compound **2** (Table 1) showed great similarity to those of **1**, suggesting that they shared the same planar structure based on the same molecular formula. The remarkable distinction was the proton coupling constants between H-7 and H-10 (7.7 Hz in **1** vs. 2.7 Hz in **2**), indicating that they possessed different relative configurations at C-7 and

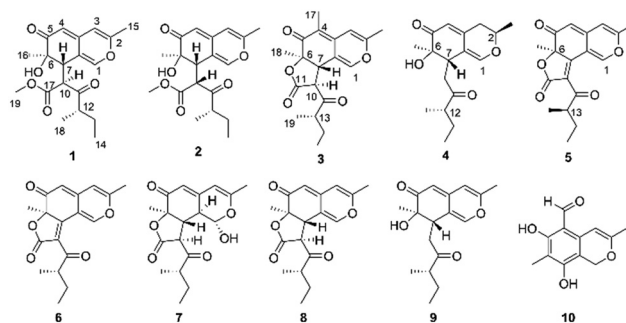


Fig. 1 Structures of compounds **1**–**10**.

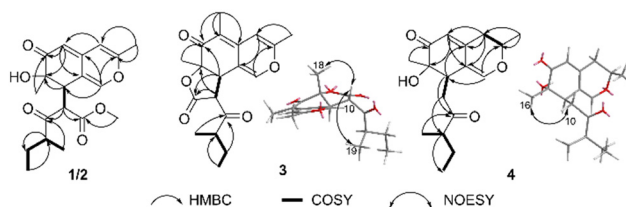


Fig. 2 Key COSY, HMBC and NOESY correlations of **1**–**4**.

C-10 (*threo* configuration for **1** and *erythro* configuration for **2**).¹³ In addition, the optical rotation values of **1** and **2** were $[\alpha]_D^{25} +21.3$ and $+28.2$, respectively, as well as their electronic circular dichroism (ECD) spectra were also almost the same (Fig. 3), indicating that both **1** and **2** were a pair of epimers. Considering the limited contribution of the flexible side chain to the ECD spectrum and the lack of NOESY signals of H-7 and H_3 -16, two truncated models **a/b** were used for the ECD calculations. The ECD spectrum of **6R,7S-b** (Fig. 3) showed the best agreement with experimental curves, which led to the determination of the **6R,7S** absolute configuration for **1** and **2**. The NMR calculations of the four candidate diastereoisomers (**1a/1b** and **2a/2b**) were then carried out using the gauge independent atomic orbital (GIAO) strategy at the B3LYP/6-31+G(d, p) level of theory with an IEPPCM solvent model of chloroform using the ORCA 5.0.3 program.¹⁴ The calculated chemical shifts of **1a/1b** and **2a/2b** were compared with the experimental values, respectively, applying total absolute deviation (TAD), mean absolute error (MAE), and DP4⁺ probability analysis as all three methods have been widely applied in addressing the

Table 1 1H and ^{13}C NMR data of compounds **1**–**3** (700, 175 MHz, TMS, δ in ppm, J in Hz)

Pos.	1^a			2^a			3^b		
	δ_C type	δ_H mult. (J)		δ_C type	δ_H mult. (J)		δ_C type	δ_H mult. (J)	
1	146.2, CH	6.91, s		147.7, CH	7.39, s		145.1, CH	7.27, s	
2	158.7, C			158.6, C			158.7, C		
3	107.5, CH	6.01, s		107.7, CH	6.01, s		104.7, CH	6.38, s	
4	103.7, CH	5.41, s		103.3, CH	5.37, s		110.6, C		
5	198.3, C			197.9, C			190.3, C		
6	73.3, C			73.2, C			82.5, C		
7	45.5, CH	3.65, d (7.7)		45.2, CH	3.43, d (2.7)		43.3, CH	3.77, d (12.0)	
8	118.1, C			116.8, C			113.7, C		
9	148.6, C			148.8, C			139.2, C		
10	54.7, CH	4.30, d (7.7)		54.9, CH	4.45, d (2.7)		54.8, CH	4.56, d (12.0)	
11	207.1, C			208.9, C			170.1, C		
12	48.6, CH	2.66, q (6.9)		47.2, CH	2.72, q (6.8)		206.9, C		
13	25.3, CH_2	1.77, m		26.0, CH_2	1.69, m		47.1, CH	2.81, h (7.0)	
		1.39, m			1.38, m				
14	11.4, CH_3	0.86, t (6.5)		11.7, CH_3	0.84, t (6.4)		24.0, CH_2	1.61, m	
								1.29, m	
15	19.4, CH_3	2.14, s		19.5, CH_3	2.14, s		11.5, CH_3	0.79, t (7.4)	
16	21.7, CH_3	1.14, s		22.4, CH_3	1.13, s		19.1, CH_3	2.15, s	
17	169.4, C			169.8, C			9.6, CH_3	1.72, s	
18	16.6, CH_3	1.18, d (6.9)		16.5, CH_3	1.16, d (6.8)		22.9, CH_3	1.40, s	
19	53.0, CH_3	3.76, s		52.7, CH_3	3.75, s		15.3, CH_3	0.92, d (7.0)	

^a Recorded in chloroform-*d*. ^b Recorded in DMSO-*d*₆.

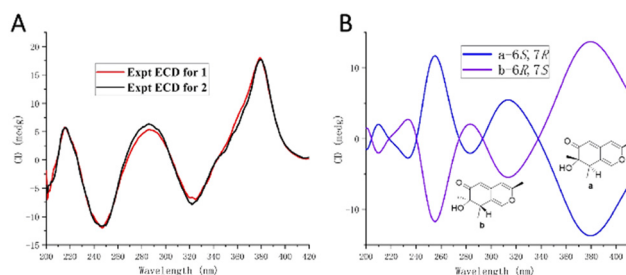


Fig. 3 (A) Experimental ECD spectra of **1** and **2** and (B) calculated ECD spectra of **a** and **b**.



stereochemical assignment of isomeric compounds.¹⁵ The lower TAD and MAE results showed that **1a** and **2a** are the most probable stereoisomers, and these results were further strongly supported by the DP4⁺ probability analysis with a high confidence level of 100% (both ¹H and ¹³C data) (Fig. S37†). Finally, the absolute configurations of **1** and **2** were defined as (6*R*,7*S*,10*S*,12*S*) and (6*R*,7*S*,10*R*,12*S*), respectively, which confirmed that they were indeed a pair of epimers at C-10.

Compound **3** was obtained as a light-yellow oil and has the molecular formula C₁₉H₂₂O₅ as deduced from HRESIMS data, suggesting nine degrees of unsaturation. The 1D NMR data of **3** shown in Table 1 reveal the same azaphilone skeleton with an angular lactone ring as that of the known analogue chermesinone B (**8**).¹²

The obvious differences were that the ¹H NMR signal of H-4 (δ_{H} 5.34) was absent and the ¹³C NMR resonance of C-4 showed an upfield shift (δ_{C} 110.6 vs. 106.1), while an additional methyl signal ($\delta_{\text{C/H}}$ 9.6/1.72, s, CH₃-17) was observed. Further analysis of HMBC signals from H₃-17 to C-4, C-5, and C-9 (Fig. 2) indicated that the additional methyl was located at C-4. Consequently, the planar structure of **3** was established as shown. The relative configurations at C-6, C-7, C-10, and C-12 in **3** were determined by a combination of a NOESY experiment and ¹H NMR coupling constant analysis. The NOESY correlations of H-10/H₃-18 and H-10/H₃-19 indicated that H-10, H₃-18, and H₃-19 were on the same side of the molecule. A large coupling constant between H-7 and H-10 ($J_{\text{H-7/H-10}}$ = 12.0 Hz) and the lack of a NOE between H-7 and H₃-18 revealed the *trans* configurations of H-7/H-10 and H-7/H₃-18. Thus, considering the limited contribution of the flexible side chain to the ECD spectrum, two truncated models **c** and **d** were employed for the ECD calculations. As shown in Fig. 4A, the Boltzmann-weighted ECD spectrum of 6*R*,7*S*,10*S*-**d** showed the best agreement with the experimental curve of **3**, which led to the determination of the 6*R*,7*S*,10*S* absolute configuration in **3**. The same as for **1** and **2**, NMR calculations of the two candidate diastereoisomers **3a/3b** were then carried out to determine the absolute configuration at C-13 using the GIAO strategy at the B3LYP/6-31+G(d,p) level of theory with an IEFPCM solvent model of DMSO. As a result, the calculated NMR data with lower TAD and MAE values and a higher DP4⁺ probability of 100% (both ¹H and ¹³C data) showed **3a** to be the most probable stereoisomer (Fig. 4B). Therefore, the absol-

ute configuration of **3** was defined as 6*R*,7*S*,10*S*,13*S* and it was named penineulone C.

Compound **4** was also obtained as a light-yellow oil and has the molecular formula C₁₇H₂₄O₄ as determined from the HRESIMS ion peak at m/z 293.1675 [M + H]⁺ (calcd for C₁₇H₂₅O₄, 293.1646), suggesting six degrees of unsaturation. The 1D NMR data of **4** shown in Table 2 reveal the presence of two olefinic methines CH-1 ($\delta_{\text{C/H}}$ 147.9/6.39, s) and CH-4 ($\delta_{\text{C/H}}$ 116.3/5.48, s), three sp³ methines CH-2 ($\delta_{\text{C/H}}$ 72.6/4.09, m), CH-7 ($\delta_{\text{C/H}}$ 40.9/3.00, d, J = 9.1 Hz), and CH-12 ($\delta_{\text{C/H}}$ 46.6/2.62, p, J = 6.6 Hz), three methylenes CH₂-3 ($\delta_{\text{C/H}}$ 34.4/2.55, m; 2.46, m), CH₂-10 ($\delta_{\text{C/H}}$ 37.7/2.86, dd, J = 18.0, 2.7 Hz; 2.76, dd, J = 18.0, 9.1 Hz) and CH₂-13 ($\delta_{\text{C/H}}$ 25.5/1.59, m; 1.35, m), and four methyls at $\delta_{\text{C/H}}$ 11.3/0.79, t, J = 7.4 Hz CH₃-14; 20.3/1.30, d, J = 6.2 Hz, CH₃-15; 20.2/1.01, s, CH₃-16 and 15.7/0.99, m, CH₃-17. Besides, the signals of the remaining five tertiary carbons including two carbonyls (δ_{C} 200.8 and 212.8), two olefinics (δ_{C} 112.2 and 150.5), and one oxygenated sp³ tertiary carbon (δ_{C} 74.5) were observed in the ¹³C NMR spectrum. The above characteristics of the NMR data indicate the same azaphilone skeleton as that of the co-isolated known analogue chermesinone A (**9**).¹²

The obvious difference between **4** and **9** was that the double bond $\Delta^{2,3}$ was reduced as one methine and one methylene, which was further verified by COSY correlations of H-2 and H₂-3 along with the HMBC signals from H₃-15 to C-2 and C-3 and from H₂-3 to C-4, C-8 and C-9 (Fig. 2). Detailed analysis of the other 2D NMR signals in Fig. 2 confirmed that the planar structure of compound **4** was a reduced product of chermesinone A (**9**).

The absolute configuration at C-12 was determined as *S*, due to the same chemical shifts at CH-12 of **4** and **9** recorded

Table 2 ¹H and ¹³C NMR data of compounds **4** and **5** (700, 175 MHz, DMSO-*d*₆, TMS, δ in ppm, J in Hz)

Pos.	4		5	
	δ_{C} type	δ_{H} mult. (J in Hz)	δ_{C} type	δ_{H} mult. (J in Hz)
1	147.9, CH	6.39, s	153.1, CH	8.63, s
2	72.6, CH	4.09, m	159.2, C	
3	34.4, CH ₂	2.55, dd (13.6, 2.8) 2.46, d (13.6)	108.1, CH	6.49, s
4	116.3, CH	5.48, s	103.9, CH	5.29, s
5	200.8, C		189.5, C	
6	74.5, C		87.4, C	
7	40.9, CH	3.00, d (9.1)	166.0, C	
8	112.2, C		110.5, C	
9	150.5, C		144.5, C	
10	37.7, CH ₂	2.86, dd (18.0, 2.7) 2.76, dd (18.0, 9.1)	122.2, C	
11	212.8, C		168.0, C	
12	46.6, CH	2.62, p (6.6)	199.8, C	
13	25.5, CH ₂	1.59, m 1.35, m	44.2, CH	3.36, q (6.5)
14	11.3, CH ₃	0.79, t (7.4)	25.5, CH ₂	1.72, m 1.29, m
15	20.3, CH ₃	1.30, d (6.2)	11.6, CH ₃	0.88, t (7.4)
16	20.2, CH ₃	1.01, s	18.9, CH ₃	2.20, s
17	15.7, CH ₃	0.99, m	24.8, CH ₃	1.59, s
18			15.6, CH ₃	0.93, d (7.0)

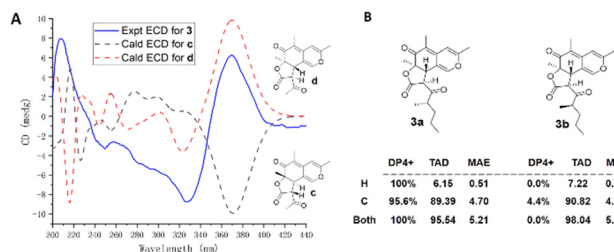


Fig. 4 (A) Experimental ECD spectra of **3** and calculated ECD spectra of **c/d**; (B) DP4⁺ probability, TAD, and MAE analysis of **3a/3b**.

in the same solvent DMSO- d_6 ($\delta_{C/H}$ 46.6/2.62 for **4** vs. 46.6/2.65 for **9**). In addition, CH₃-16 and CH₂-10 were deduced to be cofacial in compound **4** through a NOESY experiment (Fig. 2), which indicated the relative configuration of 6*R*,7*S* or 6*S*,7*R*. Subsequently, the four possible diastereoisomers, (2*R*,6*R*,7*S*,12*S*)-**4a**, (2*S*,6*R*,7*S*,12*S*)-**4b**, (2*R*,6*S*,7*R*,12*S*)-**4c**, and (2*S*,6*S*,7*R*,12*S*)-**4d**, of **4** were subjected to TDDFT ECD calculations in the MeOH model. However, the calculated ECD spectra of **4a–4d** were nearly the same and none of them showed the best agreement with the experimental curve (Fig. 5A). The TAD and MAE analysis of NMR calculations for the four candidate diastereoisomers (**4a–4d**) led to the deduction that **4** adopted the absolute configuration of (2*R*,6*R*,7*S*,12*S*)-**4a** with a DP4⁺ probability (both ¹H and ¹³C data) of 81.11% (Fig. 5B). Therefore, the absolute configuration of **4** was defined as 2*R*,6*R*,7*S*,12*S* as shown in Fig. 1 and it was named penineulone D.

Compound **5** was obtained as a light-yellow oil and has the molecular formula C₁₈H₁₈O₅ as deduced from HRESIMS data, suggesting ten degrees of unsaturation. The 1D NMR data of **5** shown in Table 2 exhibit a slight difference compared with those of the co-isolated analogue 8,11-didehydrochermesinone B (**6**),¹⁶ which is produced by an endophytic fungus *Nigrospora* sp. of *Aconitum carmichaeli*. In the same solvent of DMSO- d_6 , the most obvious differences between **5** and **6** were the chemical shifts of C-12 (199.8 vs. 200.1), CH₂-14 ($\delta_{C/H}$ 25.5/1.29, 1.72 vs. 25.7/1.25, 1.53), and CH₃-18 ($\delta_{C/H}$ 15.6/0.93, d, *J* = 7.0 Hz vs. 14.5/1.05, d, *J* = 6.7 Hz), which enlightened us that **5** and **6** may be a pair of epimers at C-13. Similar ECD spectra of **5** and **6** indicated that they were indeed not the same compound and the calculated ECD spectra of the four candidate diastereoisomers (**5a–5d**) exhibited that the ECD spectra of **5** and **6** were governed mainly by the C-6 chirality center as the 6*R* absolute configuration (Fig. 6). Consequently, the only explanation of different chemical shifts around C-13 in **5** was that **5** was a C-13 epimer of **6**, while they shared the identical 6*R* absolute configuration. Finally, compound **5** was determined as a new azaphilone with the 6*R*,13*R* absolute configuration and named penineulone E. To our knowledge, **5** represents the third azaphilone with an *R* configuration at the side chain chirality center linked to a methyl group,^{17,18} while most azaphilones have been discovered with an *S* configuration.^{1,2}

Additionally, the other five known compounds were identified as 8,11-didehydrochermesinone B (**6**),¹⁶ chermesinone C (**7**), chermesinone B (**8**), chermesinone A (**9**),¹² and anishidiol (**10**)¹⁹ by using X-ray diffraction crystal structures or compari-

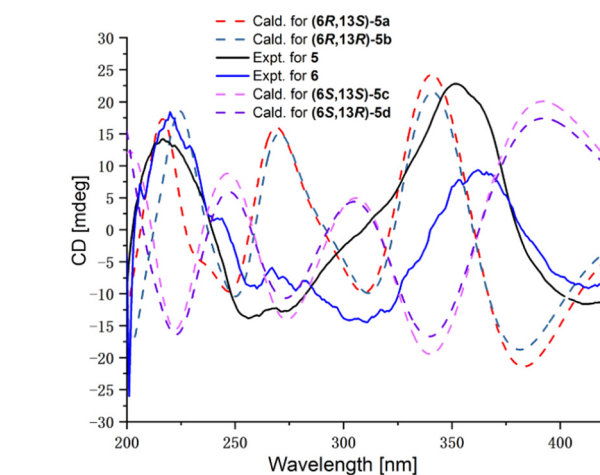


Fig. 6 Experimental and calculated ECD curves of **5**, **6** and **5a–5d**.

son of their NMR data with those reported in the literature. The colorless crystals of **8** and **9** were obtained from a mixed solvent of CHCl₃ and CH₃OH (v : v, 1 : 1). Crystallographic data for structure **8** were first reported and crystallographic data of **8** and **9** have been deposited with the Cambridge Crystallographic Data Centre under supplementary publication numbers CCDC 2214106 and 2214107† (Fig. 7).

The isolated compounds (**1–10**) were evaluated for their antitumor (C42B, H446, and H69 cell lines) and anti-viral (HSV-1/2) activities. However, the cell viabilities of the three human tumor cell lines were all over 80% after treatment with compounds **1–10** at a concentration of 5.0 μM for 72 h. And the CPE assay result showed no inhibition of **1–10** against HSV-1/2 at a concentration of 20 μM.

Neuroprotective effects of azaphilones against Aβ_{25–35}-induced neurotoxicity were assessed in primary cultured cortical neurons.²⁰ Among the ten tested compounds, **8** and **9** showed significant neuronal protective effects against Aβ_{25–35}-induced neuronal death at a concentration of 10 μM (Fig. 8). In addition, the axonal regrowth effects of **8** and **9** were evaluated (Fig. 9A). As a result, compared with the control group, the Aβ_{25–35}-treatment significantly reduced the length of axons in primary cultured cortical neurons. However, treatment with compounds **8** and **9** dramatically increased the length of pNF-H positive axons at both concentrations of 1 and 10 μM (Fig. 9B and C).

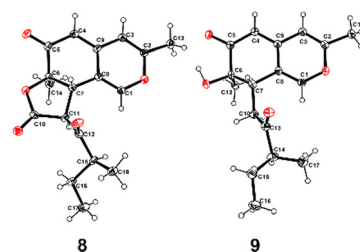


Fig. 7 X-ray diffraction crystal structures (ORTEP drawing) of **8** and **9**.

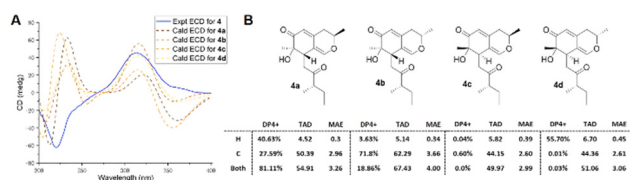


Fig. 5 (A) Experimental ECD spectra of **4** and calculated ECD spectra of **4a–4d**; (B) DP4⁺ probability, TAD, and MAE analysis of **4a–4d**.



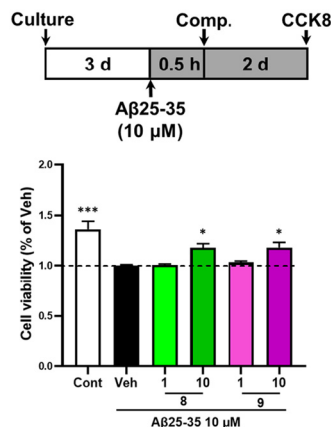


Fig. 8 Neuroprotective effects of **8** and **9** on Aβ₂₅₋₃₅-induced neuronal death in primary cultured cortical neurons. The error bars represent SEM. **p* < 0.05, ****p* < 0.001 vs. Veh, one-way ANOVA *post hoc* Dunnett's test (*n* = 3 repeated experiments).

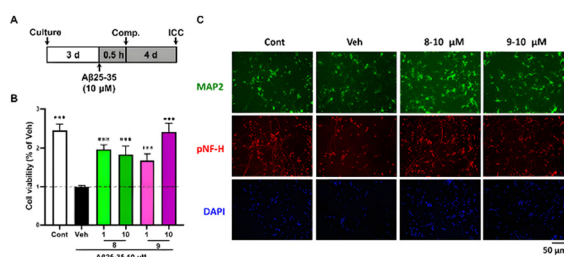


Fig. 9 Axonal regenerative effects of **8** and **9** on Aβ₂₅₋₃₅-induced axonal atrophy in primary cultured cortical neurons. A. Experimental schedule. B. Statistical results. C. Representative photos of pNF-H-positive axons and MAP2-positive neurons. The error bars represent SEM. ****p* < 0.001 vs. Veh, one-way ANOVA *post hoc* Dunnett's test.

Experimental section

General experimental procedures

Optical rotations were acquired using a PerkinElmer MPC 500 (Waltham) polarimeter. UV spectra were recorded on a Shimadzu UV-2600 PC spectrometer (Shimadzu). ECD spectra were recorded using a Chirascan circular dichroism spectrometer (Applied Photophysics). IR spectra were recorded using an IR Affinity-1 spectrometer (Shimadzu). NMR spectra were recorded using a Bruker DRX-700 spectrometer (Bruker BioSpin, Fallanden, Switzerland) at 700 MHz for ¹H-NMR and 175 MHz for ¹³C-NMR using TMS as the internal standard and chemical shifts were recorded as δ-values. HRESIMS spectra were recorded using a Bruker miXis TOF-QII mass spectrometer (Bruker). X-ray diffraction intensity data were collected using a CrysAlis PRO CCD area detector diffractometer with graphite-monochromated Cu Kα radiation (λ = 1.541 78 Å). TLC and column chromatography (CC) were performed on plates precoated with silica gel GF254 (10–40 μm) and over silica gel (200–300 mesh) (Qingdao Marine Chemical Factory, Qingdao, China), and on Sephadex LH-20 (Amersham

Biosciences, Uppsala, Sweden), respectively. Spots were detected on TLC under 254 nm UV light or by heating after spraying with 5% H₂SO₄ in EtOH. All solvents used were of analytical grade (Tianjin Fuyu Chemical and Industry Factory, Tianjin, China). Semipreparative HPLC was performed using an ODS column (YMC-pack ODS-A, 10 × 250 mm², 5 μm, 2 mL min^{−1}).

Fungal material

The fungal strain SCSIO41030 was isolated from the sediment collected at a depth of 1509 m in the South China Sea and was the same strain as reported previously.¹¹

Fermentation and extraction

A large-scale fermentation of the fungal strain SCSIO41030 was carried out at room temperature in 1 L conical flasks containing the liquid medium (300 mL per flask) composed of mannitol (20 g L^{−1}), yeast extract (3 g L^{−1}), sodium glutamate (10 g L^{−1}), glucose (10 g L^{−1}), maltose (20 g L^{−1}), corn steep liquor (1 g L^{−1}), MgSO₄·7H₂O (0.3 g L^{−1}), KH₂PO₄ (0.5 g L^{−1}) and tap water after adjusting its pH to 7.5. After static cultivation for 30 days, the whole fermented broth (20 L) was filtered through cheesecloth to separate it into filtrate and mycelia. The filtrate was extracted three times with EtOAc to yield an EtOAc solution, while the mycelia were cut into small pieces, soaked in EtOAc, and sonicated for 20 min to yield another EtOAc solution. Both EtOAc solutions were combined and concentrated under reduced pressure to obtain a black extract (16.2 g).

Isolation and purification

The EtOAc extract was subjected to vacuum liquid chromatography on a silica gel column using step-gradient elution with MeOH–CH₂Cl₂ (0–100%) to separate it into six fractions based on TLC properties. Fraction 2 was divided into three parts (Frs 2-1–2-3) followed by Sephadex LH-20 column elution with MeOH. Compounds **7** (7.2 mg, *t*_R = 18.5 min) and **8** (10.5 mg, *t*_R = 23.5 min) were obtained from Fr. 2-1 purified by semi-preparative HPLC (45% CH₃CN–H₂O, 2 mL min^{−1}). Fr. 2-2 was then purified by semi-preparative HPLC (45% CH₃CN–H₂O, 2 mL min^{−1}) to yield **P1** (3.1 mg, *t*_R 16.9 min). **P1** was subsequently subjected to a chiral column of DAICEL CHIRALPAK IC to afford enantiomeric compounds penineulone A (**1**) (2.2 mg, *t*_R = 12.5 min) and penineulone B (**2**) (1.76 mg, *t*_R = 16.1 min), using ethanol/*n*-hexane (v/v: 55/65, flow rate 1 mL min^{−1}) with 0.1% trifluoroacetic acid (TFA) as the mobile phase. Fr. 2-3 was separated by semi-preparative HPLC (52% CH₃CN–H₂O with 0.1% TFA, 2 mL min^{−1}) to afford **5** (4.2 mg, *t*_R 35.9 min), **6** (6.3 mg, *t*_R 20.0 min), and **9** (9.0 mg, *t*_R 18.2 min). Fraction 3 was separated into two parts by ODS column and Sephadex LH-20 column elution with MeOH. Subfraction 3-2 was further purified by semi-preparative HPLC (56% CH₃OH–H₂O, 2 mL min^{−1}) to yield **3** (3.1 mg, *t*_R 14.8 min) and **10** (8.2 mg, *t*_R 22.5 min). Subfraction 4-2, obtained from Sephadex LH-20 column separation of fraction 4, was also subjected to semi-preparative HPLC with 45% CH₃OH–H₂O (0.1% TFA) elution to give **4** (5.3 mg, *t*_R 18.5 min, 2 mL min^{−1}).



Penineulone A (1): light yellow oil; $[\alpha]_{\text{D}}^{25} +28.3$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 352 (3.88), 225 (3.76) nm; for ^1H and ^{13}C NMR data (DMSO- d_6 , 700/175 MHz), see Table 1; HR-ESI-MS m/z 349.1638 $[\text{M} + \text{H}]^+$ (calcd for $\text{C}_{19}\text{H}_{25}\text{O}_6$, 349.1646).

Penineulone B (2): light yellow oil; $[\alpha]_{\text{D}}^{25} +21.7$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 351 (3.98), 226 (3.96) nm; for ^1H and ^{13}C NMR data (DMSO- d_6 , 700/175 MHz), see Table 1; HR-ESI-MS m/z 349.1637 $[\text{M} + \text{H}]^+$ (calcd for $\text{C}_{19}\text{H}_{25}\text{O}_6$, 349.1649).

Penineulone C (3): light yellow oil; $[\alpha]_{\text{D}}^{25} -6.5$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 349 (3.75), 225 (3.58) nm; for ^1H and ^{13}C NMR (DMSO- d_6 , 700/175 MHz) data, see Table 1; HR-ESI-MS m/z 331.1467 $[\text{M} + \text{H}]^+$ (calcd for $\text{C}_{19}\text{H}_{23}\text{O}_5$, 331.1469).

Penineulone D (4): light yellow oil; $[\alpha]_{\text{D}}^{25} +30.1$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 348 (3.85), 226 (3.78) nm; for ^1H and ^{13}C NMR (DMSO- d_6 , 500/125 MHz) data, see Table 2; HR-ESI-MS m/z 293.1675 $[\text{M} + \text{H}]^+$ (calcd for $\text{C}_{17}\text{H}_{25}\text{O}_4$, 293.1646).

Penineulone E (5): light yellow oil; $[\alpha]_{\text{D}}^{25} -12.0$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 349 (3.68), 225 (3.98) nm; for ^1H and ^{13}C NMR (DMSO- d_6 , 500/125 MHz) data, see Table 2; HR-ESI-MS m/z 314.1167 $[\text{M} + \text{H}]^+$ (calcd for $\text{C}_{18}\text{H}_{18}\text{O}_5$, 314.1159).

ECD and NMR calculations

The preliminary conformational distribution search was performed using the Confab algorithm in OpenBabel software. The conformers with Boltzmann population of over 1% (the relative energy within 10 kcal mol $^{-1}$) were reoptimized using density functional theory (DFT) at the B97-3c level under vacuum using the ORCA 5.0.3 program.¹⁴ After that, frequency calculations following the geometry optimizations were performed to verify that all the structures correspond to energy minima and have no imaginary frequency. The overall theoretical calculation of ECD was conducted in MeOH using time-dependent density functional theory (TDDFT) at the PBE0 def2-TZVP level for the stable conformers (the relative energy within 5 kcal mol $^{-1}$). Rotatory strengths for a total of 35 excited states were calculated. The ECD spectra of different conformers were generated using the Multiwfn program²¹ with a half-bandwidth of 0.5–0.6 eV, according to the Boltzmann-calculated contribution of each conformer after UV correction. Solvent effects of methanol solution were evaluated at the same DFT level by using the CPCM method.

In the case of conformationally flexible compounds, the conformational search was done in the gas phase using the MMFF force field. All conformers within 5 kcal mol $^{-1}$ of the lowest energy conformer were subjected to further reoptimization and frequency calculations at the B3LYP/6-31G* level of theory. NMR calculations of all candidate diastereoisomers were then carried out using the gauge independent atomic orbital (GIAO) strategy at the B3LYP/6-31G*(d,p) level of theory with an IEFPCM solvent model in the ORCA 5.0.3 program.

Assay of neuroprotection effects

Primary cortical neuronal culture, cell viability assay and axonal density measurement. The primary cortical neurons were isolated from embryos at 14 days of gestation in ICR mice (Guangdong Medical Laboratory Animal Centre, Guangzhou, China), as described previously (ref. 1). Briefly, the dura mater was removed from dissected cerebral cortices. The neurons were obtained from minced cortices after trypsinization with 0.5% trypsin-EDTA (Gibco, 25300054) and were seeded in a Neurobasal medium (Gibco, 21103-049) containing 2 mM L-glutamine, 0.6% D-glucose, and 2% B27. The neurons were cultured in a 96-well plate (for the cell viability assay) or an 8-well plate (for the axonal density assay) at a density of 2.1×10^4 cells per cm 2 .

To detect neuronal viability, primary cortical neurons were cultured for 3 days and treated with A β_{25-35} (Sigma, A4559) for 0.5 h, followed by treatment with **8** and **9** (1 and 10 μM) for 48 h. Afterwards, the cell counting kit (CCK8, APEX BIO, K1018) reagent (10 μL) was added to each well for 3 h according to the manufacturer's instructions. The absorbance values of CCK8 test were measured using a 96-well ELISA microplate reader (Biotek, VT, USA) at 450 nm. A β_{25-35} was dissolved in dH $_2\text{O}$ to a final concentration of 5 mM and was incubated at 37 $^{\circ}\text{C}$ for 4 days for aggregation.

To test the axonal regenerative effects of **8** and **9** on A β_{25-35} -induced axonal atrophy, after culturing neurons for 3 days, 10 μM A β_{25-35} was pretreated for 0.5 h and then cocultured with **8** and **9** (1 and 10 μM) for 4 days. The neurons were then fixed with 4% paraformaldehyde (PFA; Solarbio, Beijing, China) at 25 $^{\circ}\text{C}$ for 1 h and immunostained for visualizing phosphorylated neurofilament H (pNF-H, 1:500; Covance, SMI-35R, CA, United States) and microtubule-2 associated protein 2 (MAP2, 1:2000; Abcam, ab32454). Alexa Fluor 594-conjugated goat anti-mouse IgG (1:300, Abcam, ab150116) and Alexa Fluor 488-conjugated goat anti-rabbit IgG (1:300, Abcam, ab150081) were used as secondary antibodies. 4',6-Diamidino-2-phenylindole (DAPI) (1 $\mu\text{g mL}^{-1}$, Biomol, Hamburg, Germany) was used for nuclear counterstaining. A fluorescence microscope system (Echo Revolve, ECHO, CA, United States) was used for image capture at a size of $480 \times 640 \mu\text{m}^2$. Ten images of each group were captured and analyzed using ImageJ (NIH), with the Neurite Tracer plugins. The average lengths of pNF-H positive axons per neuron were measured.

Conclusions

In conclusion, a further study of a deep-sea derived *Penicillium* sp. SCSIO41030 that produced *p*-terphenyl derivatives revealed ten azaphilones including one pair of new epimers and three new ones, penineulones A–E (**1**–**5**) with the same structural core of angular deflectin, which were obtained from the strain fermented on a liquid medium. Extensive NMR spectroscopic and HRESIMS data and ECD and NMR calculations were used to determine the structures of new compounds. In addition,



azaphilones **8** and **9** were found to possess neuroprotective effects against A β _{25–35}-induced neurotoxicity at a concentration of 10 μ M and significantly promoted axonal regrowth in primary cultured cortical neurons. To our knowledge, this is the first report of azaphilones with neuroprotective effects.

Author contributions

W. Chen: investigation, data curation, visualization, and writing – original draft; J. Jiang: investigation, data curation, and writing – original draft; X. Pang and Y. Song: investigation; Z. Yang: methodology, supervision, and resources; J. Wang: project administration, funding acquisition, supervision, resources, and writing – review & editing; Y. Liu: funding acquisition, supervision, and resources. All authors read and approved the final manuscript.

Conflicts of interest

The authors declare no conflict of interest.

Acknowledgements

This work was financially supported by the Guangdong MEPP Funds (No. GDNRC [2024]28), the Hainan Provincial Joint Project of Sanya Yazhou Bay Science and Technology City (2021CXLH0013, 2021JJLH0097), the National Natural Science Foundation of China (U23A201140 and 42376124), the Guangdong Local Innovation Team Program (2019BT02Y262), the Guangzhou Science and Technology Project (202201010678), the Key-Area Research and Development Program of Guangdong Province (2023B1111050008), and the Key Science and Technology Plan Projects in Nansha District (2023ZD010). We are grateful to ZH Xiao, AJ Sun, X Ma, XH Zheng, and Y Zhang in the analytical facility at SCSIO for recording spectroscopic data.

References

- 1 J. M. Gao, S. X. Yang and J. C. Qin, *Chem. Rev.*, 2013, **113**, 4755–4811.
- 2 C. Chen, H. Tao, W. Chen, B. Yang, X. Zhou, X. Luo and Y. Liu, *RSC Adv.*, 2020, **10**, 10197–10220.
- 3 Y. Shao, H. Yan, T. Yin, Z. Sun, H. Xie, L. Song, K. Sun and W. Li, *J. Antibiot.*, 2020, **73**, 77–81.
- 4 E. Kuhnert, F. Surup, E. B. Sir, C. Lambert, K. D. Hyde, A. I. Hladki, A. I. Romero and M. Stadler, *Fungal Diversity*, 2015, **71**, 165–184.
- 5 F. Cao, Z. H. Meng, P. Wang, D. Q. Luo and H. J. Zhu, *J. Nat. Prod.*, 2020, **83**, 1283–1287.
- 6 W.-Y. Zu, J.-W. Tang, K. Hu, Y.-F. Zhou, L.-L. Gou, X.-Z. Su, X. Lei, H.-D. Sun and P.-T. Puno, *J. Org. Chem.*, 2021, **86**, 475–483.
- 7 C. Huo, X. Lu, Z. Zheng, Y. Li, Y. Xu, H. Zheng and Y. Niu, *Phytochemistry*, 2020, **170**, 112224.
- 8 M. Chen, N.-X. Shen, Z.-Q. Chen, F.-M. Zhang and Y. Chen, *J. Nat. Prod.*, 2017, **80**, 1081–1086.
- 9 W. Chen, C. Chen, J. Long, S. Lan, X. Lin, S. Liao, B. Yang, X. Zhou, J. Wang and Y. Liu, *J. Antibiot.*, 2021, **74**, 156–159.
- 10 X. Pang, W. Chen, X. Wang, X. Zhou, B. Yang, X. Tian, J. Wang, S. Xu and Y. Liu, *Front. Microbiol.*, 2021, **12**, 730807.
- 11 W. Chen, J. Zhang, X. Qi, K. Zhao, X. Pang, X. Lin, S. Liao, B. Yang, X. Zhou, S. Liu, J. Wang, X. Yao and Y. Liu, *J. Nat. Prod.*, 2021, **84**, 2822–2831.
- 12 H. Huang, X. Feng, Z. Xiao, L. Liu, H. Li, L. Ma, Y. Lu, J. Ju, Z. She and Y. Lin, *J. Nat. Prod.*, 2011, **74**, 997–1002.
- 13 V. A. Cao, B.-K. Choi, H.-S. Lee, C.-S. Heo and H. J. Shin, *J. Nat. Prod.*, 2021, **84**, 1843–1847.
- 14 F. Neese, *Wiley Interdiscip. Rev.: Comput. Mol. Sci.*, 2022, **12**, e1606.
- 15 N. Grimblat, M. M. Zanardi and A. M. Sarotti, *J. Org. Chem.*, 2015, **80**, 12526–12534.
- 16 S.-P. Zhang, R. Huang, F.-F. Li, H.-X. Wei, X.-W. Fang, X.-S. Xie, D.-G. Lin, S.-H. Wu and J. He, *Fitoterapia*, 2016, **112**, 85–89.
- 17 P. S. Steyn and R. Vleggaar, *J. Chem. Soc., Perkin Trans. 1*, 1986, 1975.
- 18 C. Chen, J. Wang, H. Zhu, J. Wang, Y. Xue, G. Wei, Y. Guo, D. Tan, J. Zhang, C. Yin and Y. Zhang, *Chem. Biodivers.*, 2016, **13**, 422–426.
- 19 T. Hosoe, N. Mori, K. Kamano, T. Itabashi, T. Yaguchi and K. Kawai, *J. Antibiot.*, 2011, **64**, 211–212.
- 20 J. Deng, X. Feng, L. Zhou, C. He, H. Li, J. Xia, Y. Ge, Y. Zhao, C. Song, L. Chen and Z. Yang, *Food Res. Int.*, 2022, **158**, 111576.
- 21 T. Lu and F. Chen, *J. Comput. Chem.*, 2012, **33**, 580–592.

