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New spirobisanaphthalenes from an endolichenic fungus strain CGMCC 3.15192 and their anticancer effects through the P53–P21 pathway†

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Natural products from fungi have remained a rich resource for drug discovery. Here we report the isolation of three new spirobisanaphthalenes, namely sacrosomycin A–C (1–3), and three known analogues (4–6), from the ethyl acetate extract of a nonsporulating endolichenic fungus derived from *Peltigera elisabethae* var. *mauritzii*. The structures of these compounds were elucidated by IR, UV, MS, and NMR. Biological functions of these compounds were evaluated using cultured human cancer cell lines. Short-term cell growth and long-term cell survival assays show that compound 5 demonstrated the strongest cancer cell growth inhibition effect. We reveal that compound 5 induced both cell cycle arrest at the G2/M phase and cell death. Using western blotting, luciferase reporter assay and quantitative PCR (qPCR), we show that compound 5 induced up-regulation of the P53–P21 pathway, supporting the cell cycle arrest and growth inhibition effect of this compound. In contrast, these compounds did not induce cell death in a normal cell line. These results demonstrate a potential anticancer effect of this rare family of spirobisanaphthalene compounds isolated from endolichenic fungi.

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Introduction

Spirobisanaphthalenes, also called bisnaphthospiroketal, belong to a relatively rare family of bioactive natural products that contain a 1,8-dihydroxynaphthalene derived spiroketal unit linked to a second, oxidized naphthalene moiety.^{1–3} Generally, spirobisanaphthalenes are divided into three subtypes, namely, spiroxin, preussomerin and deoxypreussomerin according to the linkage type of the two naphthalene rings, with the former joined together by two oxygen bridges and one carbon–carbon bridge and the later both joined together with three or two oxygen bridges, respectively.⁴ Recently, a small group of the fourth subtype of spirobisanaphthalenes was also discovered with only eight compounds reported up to now.^{5–7}

Spirobisanaphthalenes have attracted much attention because of their unique structures and broad spectrum of biological activities including antibacterial, antifungal, herbicidal, cytotoxic and anti-tumor activities.^{3,7} The unique skeleton and broad spectrum of bioactivities of spirobisanaphthalenes demonstrate great promise for the discovery of new agents for the treatment of human diseases, especially cancer.

P53 is a transcriptional factor that regulates the expression of a cascade of genes involved in cell cycle, apoptosis, autophagy, metabolism, pluripotency, and many other cellular processes.⁸ P53 induces apoptosis through inducing expression of the pro-apoptotic BCL-2 family proteins including PUMA, BAX, NOXA, *etc.*^{9–12} On the other hand, P53 induces upregulation of P21,^{13–16} contributing largely to the cell cycle arrest function of P53.^{17,18} Hence, P53 is a critical gene in suppressing tumorigenesis.¹⁹ Consistently, P53 mutation is detected in almost half of all human cancers.¹⁶ P21 is an endogenous inhibitor of the cyclin-dependent kinase, which inhibits cell cycle progression genes such as CDC25C, CDK1, CDK2, CCNE1, CCNB1, *etc.*,^{20,21} leading to cell cycle arrest in G1 or G2/M phase.^{16,22–24}

As part of our ongoing efforts to search for new bioactive metabolites from rarely studied fungal species, we cultured a nonsporulating endolichenic fungus (CGMCC 3.15192) derived from *Peltigera elisabethae* var. *mauritzii*. From the ethyl acetate extract, we isolated three new spirobisanaphthalenes named sacrosomycin A–C (1–3) and three known analogues (4–6). Here we report the isolation, structural elucidation of sacrosomycin A–C (1–3) and biological assessment of these six

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‡ Equally to this work.



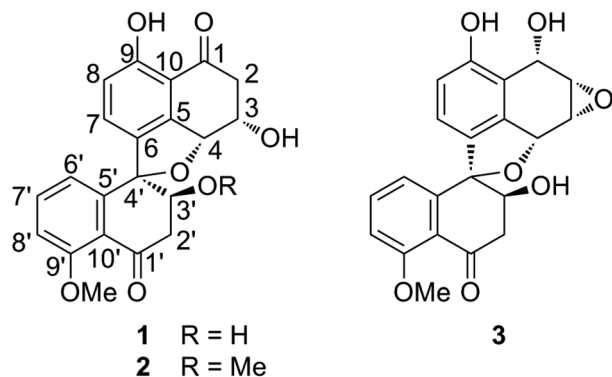


Fig. 1 Chemical structures of compounds 1–3.

compounds in the anticancer regimen. Our results show that compound 5 exhibited the strongest anticancer effect in cultured human cancer cell lines. We further reveal that this compound induced cell cycle arrest through up-regulating the P53–P21 pathway. These data demonstrate potential anticancer activity of spirobisnaphthalene compounds, laying foundation for detailed medicinal chemistry research for these compounds in the near future.

Results and discussion

Chemical identification of spirobisnaphthalenes

The fungus strain (CGMCC 3.15192) was cultured in solid state fermentation with rice at 25 °C for 47 days. The fermented material was extracted with ethyl acetate (EtOAc) and the extracts were subjected to silica gel and ODS column chromatography, which yielded five fractions (W1–W5). Preliminary screening

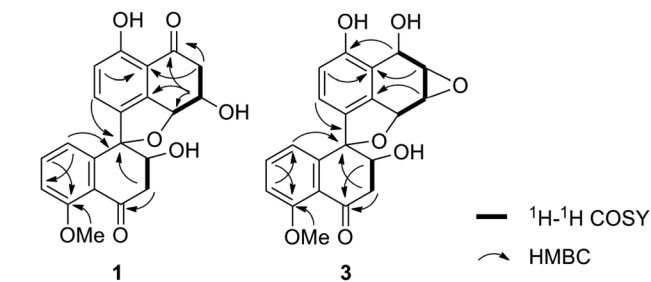


Fig. 2 Key $^1\text{H}-^1\text{H}$ COSY and HMBC correlations of compounds 1 and 3.

revealed that fractions W2 (50% MeOH in H_2O) and W3 (70% MeOH in H_2O) showed different degrees of cytotoxic activities against five cancer cell lines (HL-60, SMMC-7721, A-549, MCF-7 and SW480). Hence, fractions W2 and W3 were further subjected to silica gel, ODS, Sephadex LH 20 column chromatography and RPHPLC to yield three new spirobisnaphthalenes, named sarcosomycin A–C (1–3), along with urnucratin C (4), Plecmillin A (5), and Plecmillin C (6) (Fig. 1).

Sarcosomycin A (1) was obtained as a yellow amorphous powder. The molecular formula was established as $\text{C}_{21}\text{H}_{18}\text{O}_7$ by HRESIMS (m/z 383.1116 $[\text{M} + \text{H}]^+$, calcd for $\text{C}_{21}\text{H}_{19}\text{O}_7$: 383.1131), indicating 13° of unsaturation. The IR absorption implied the presence of hydroxyls, α,β -unsaturated ketone and aromatic ring at 3423, 1656 (broad single peak), 1594, 1474 cm^{-1} . The ^1H NMR spectrum showed the presence of a 1,2,3-trisubstituted aromatic protons at δ_{H} 6.80 (1H, d, 8.0 Hz), 7.16 (1H, d, 8.0 Hz), and 7.15 (1H, t, 8.0 Hz), a 1,2,3,4-tetrasubstituted aromatic protons at δ_{H} 6.94 (1H, d, 8.4 Hz) and 7.35 (1H, d, 8.0 Hz), and a methoxy proton at δ_{H} 3.89 (3H, s). The ^{13}C NMR spectrum

Table 1 The ^1H and ^{13}C -NMR data of 1–3 (400 MHz for ^1H and 100 MHz for ^{13}C in CD_3OD)

No	1 δ_{H} (J in Hz) δ_{C}	2 δ_{H} (J in Hz) δ_{C}	3 δ_{H} (J in Hz) δ_{C}
1		201.0	200.9
2	2.74 (br s)	44.3	43.9
3	4.56 (br s)	70.4	70.4
4	5.37 (br s)	80.9	81.3
5		147.4	147.2
6		131.6	131.3
7	7.35 (d, 8.4)	132.0	131.9
8	6.94 (d, 8.4)	119.3	119.4
9		161.7	161.6
10		114.2	114.3
1'		197.7	197.0
2'	3.08 (dd, 16.4, 10.8); 2.91 (dd, 16.4, 5.6)	45.8	42.3
3'	4.37 (m)	71.0	80.5
4'		94.2	93.6
5'		145.7	145.5
6'	6.80 (d, 8.0)	121.1	121.0
7'	7.51 (t, 8.0)	136.6	136.7
8'	7.16 (d, 8.0)	114.3	114.4
9'		161.7	161.6
10'		122.1	122.0
3'-OMe			58.4
9'-OMe	3.89 (s)	56.6	56.6
			3.88 (s)
			56.5



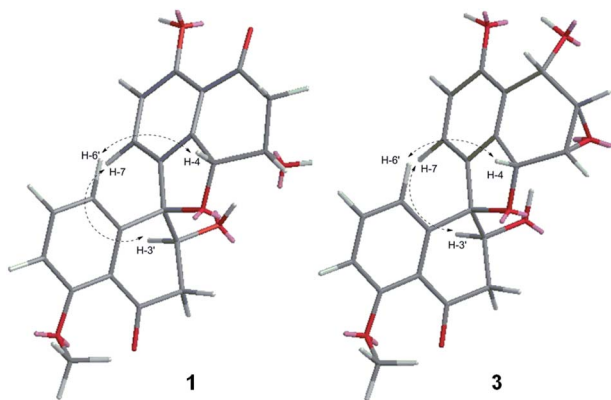


Fig. 3 Key 1D selective NOE correlations of compounds 1 and 3.

displayed two carbonyl groups at δ_C 197.7 and 201.0, twelve aromatic carbons from δ_C 114.2 to 161.8, four oxygenated carbons at δ_C 70.4, 71.0, 80.9, and 94.2, two methylene carbons at δ_C 44.3 and 45.8, and a methoxy carbon at δ_C 56.6. Comparison of the ^1H and ^{13}C NMR data of **1** with that of Plecmillin C indicated that they had similar spirobisanthralene skeleton (Table 1).⁵ The presence of one oxymethine carbon signal at δ_C 71.0 and one methylene carbon signal at δ_C 45.8 instead of a pair of olefinic carbon signals in the ^{13}C NMR spectrum of **1** suggested that the olefinic bond of α , β -unsaturated ketone was reduced and hydroxylated in compound **1**. Analyses of ^1H - ^1H COSY, HSQC and HMBC correlations established the planar structure of **1** (Fig. 2).

The relative configuration of **1** was determined by 1D selective NOE experiments, in which irradiation of H-4 (δ_H 5.37, 1H,

br s) enhanced H-6' (δ_H 6.80, 1H, d, 8.0 Hz), while irradiation of H-3' (δ_H 4.37, 1H, m) enhanced H-7 (δ_H 7.35, 1H, d, 8.4 Hz) (Fig. 3). Thus, the relative configurations of **1** were identified to be 3*S**, 4*R**, 3'*S**, 4'*R**. The CD cotton effects of **1** were consistent with that of urnucratin A, indicating the absolute configuration of **1** as 3*S*, 4*R*, 3'*S*, 4'*R* (Fig. S9†).⁶

Sarcosomycin B (**2**) was obtained as a yellow amorphous powder. The molecular formula of $\text{C}_{22}\text{H}_{20}\text{O}_7$ was determined by HRESIMS (m/z 397.1288 [$\text{M} + \text{H}$]⁺, calcd for $\text{C}_{22}\text{H}_{21}\text{O}_7$: 397.1287), indicating 13° of unsaturation. The IR absorption implied the presence of hydroxyls, α , β -unsaturated ketone and aromatic ring at 3432, 1674, 1656, 1593, and 1471 cm^{-1} . Comparison of the NMR data with that of compound **1** suggested that **2** also had a spirobisanthralene skeleton. The difference between them mainly lay in the ^{13}C resonance at C-2', C-3', and C-4' and an additional methoxy group at δ_C 58.4/ δ_H 3.15 (3H, s) in **2**. The HMBC correlation between δ_H 3.15 (3H, s) and δ_C 80.5 (C-3') located the methoxy group at C-3'. Detailed analyses of ^1H - ^1H COSY, HSQC and HMBC spectra accomplished the planar structure of **2**. The relative configuration of **2** was determined by ROESY experiment and comparison of its NMR data with that of **1**. In the ROESY spectrum, H-3' at δ_H 4.15 (1H, m) correlated with H-7 at δ_H 7.40 (1H, d, 8.0 Hz). Similar CD cotton effects of **2** with that of **1** suggested that the absolute configurations of **2** were 3*S*, 4*R*, 3'*S*, 4'*R* (Fig. S9†).

Sarcosomycin C (**3**) was obtained as a yellow amorphous powder. The molecular formula was established as $\text{C}_{21}\text{H}_{18}\text{O}_7$ by HRESIMS (m/z 383.1123 [$\text{M} + \text{H}$]⁺, calcd for $\text{C}_{21}\text{H}_{19}\text{O}_7$: 383.1131), indicating 13° of unsaturation. The IR absorption implied the presence of hydroxyls, α , β -unsaturated ketone and aromatic ring at 3424, 1669 (broad singlet), 1592, and 1468 cm^{-1} . Comparison

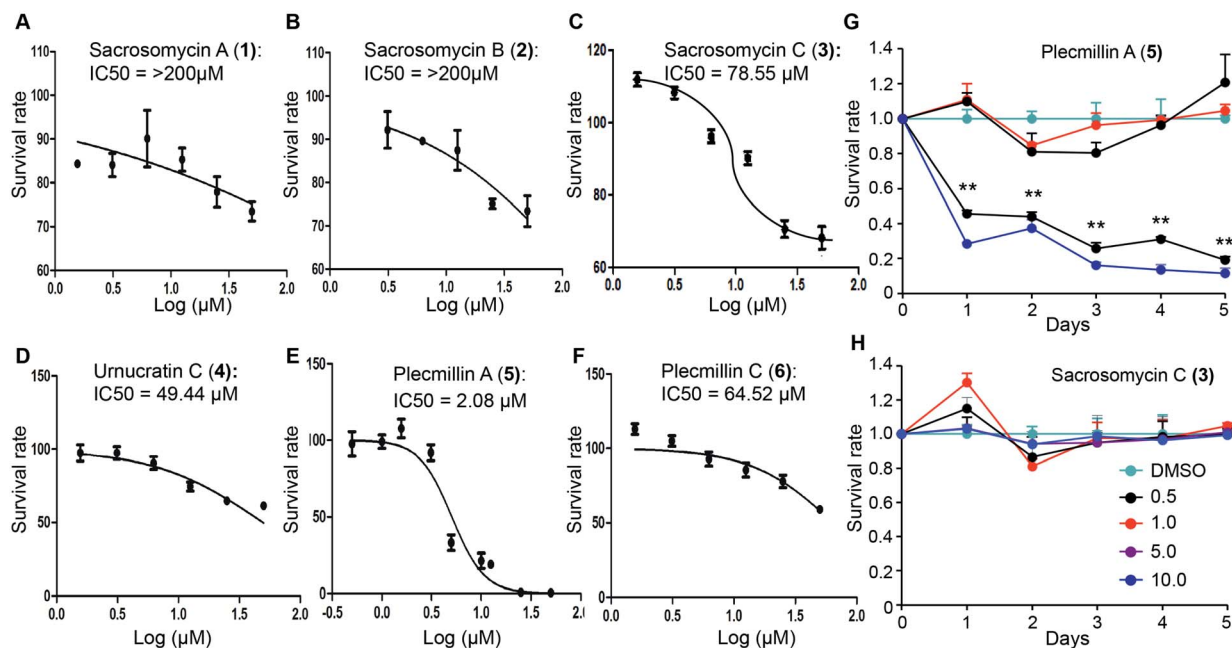


Fig. 4 Effects of compounds 1–6 on cell proliferation. (A–F) The IC_{50} values of compounds 1–6 after 48-hour treatment of HCT116 cells were assessed by MTT assay. Sacrosomycin A–C (1–3) displayed greater than 200 μM of IC_{50} . (G–H) Time- and dose-dependent growth of U2OS cells in the presence of Plecmillin A (5) or Sacrosomycin C (3). Data represent mean and standard deviation from three replicates. ** $P < 0.0001$ compared with control groups at the corresponding time points.



of the ^1H and ^{13}C NMR data with that of Plecmillin E indicated that they showed great similarity (Table 1). The main difference between them lay in the ^{13}C resonance of C-2' and C-3' and the absence of a methoxy group in **3**, suggesting that compound **3** was a demethylated derivative of Plecmillin E.⁵ The ^1H - ^1H COSY, HSQC and HMBC correlations confirmed the planar structure of **3** (Fig. 2). The relative configuration of **3** was determined based on 1D selective NOE correlations and by comparison of its NMR data with that of umucratin C. In the 1D selective NOE experiment, irradiation of H-4 (δ_{H} 5.24, 1H, br s) enhanced H-6' (δ_{H} 6.69, 1H, dd, 7.8, 0.8 Hz), while irradiation of H-3' (δ_{H} 4.34, 1H, dd, 9.4, 5.9 Hz) enhanced H-7 (δ_{H} 6.98, 1H, d, 8.4 Hz) (Fig. 3). Thus, the relative configurations of **3** were determined to be $1S^*$, $2S^*$, $3R^*$, $4R^*$, $3'S^*$, $4'R^*$. The CD cotton effects of **3** were consistent with that of Plecmillin E,⁵ indicating its absolute configurations to be $1S$, $2S$, $3R$, $4R$, $3'S$, $4'R$ (Fig. S26†). The biosynthetic origin of **1**–**6** also supported the deduction mentioned above.^{5,6}

Biological evaluation of spirobisanaphthalenes

Spirobisanaphthalenes have shown anticancer activities. To this end, we evaluated the anticancer effect of these six compounds. We found that 48 hour treatment of human colorectal HCT116

cancer cells with Plecmillin A (**5**) demonstrated the strongest growth inhibitory effect with an IC_{50} of $\sim 2.1 \mu\text{M}$, whereas the sacrosomycins were greater than $200 \mu\text{M}$ (Fig. 4A–C). Subsequently, we monitored cell proliferation over five days in the presence of increasing concentrations of Plecmillin A (**5**) or one of the new spirobisanaphthalenes, sacrosomycin C (**3**). We found that Plecmillin A (**5**) significantly inhibited cell growth at and above $5 \mu\text{M}$ (Fig. 4G). In contrast, sacrosomycin C (**3**) did not inhibit cell growth at least by $10 \mu\text{M}$ (Fig. 4H). These results are consistent with the IC_{50} data, suggesting a potential tumor suppressing role of Plecmillin A (**5**).

In our effort to determine how Plecmillin A (**5**) inhibited cancer cell growth, we measured the cell cycle profile by flow cytometry and found out that Plecmillin A (**5**) time-dependently induced accumulation of cells at the G2/M phase (Fig. 5A), suggesting that Plecmillin A (**5**) induces cell cycle arrest, which may contribute to the cancer cell growth inhibition. During analysis to understand the molecular mechanisms underlying the cell cycle arrest, we found that Plecmillin A (**5**), and to a much lesser degree, sacrosomycin C (**3**), increased the protein levels of P53 and P21 (Fig. 5B), indicating that this compound class upregulates the tumor suppressing P53–P21 pathway. On

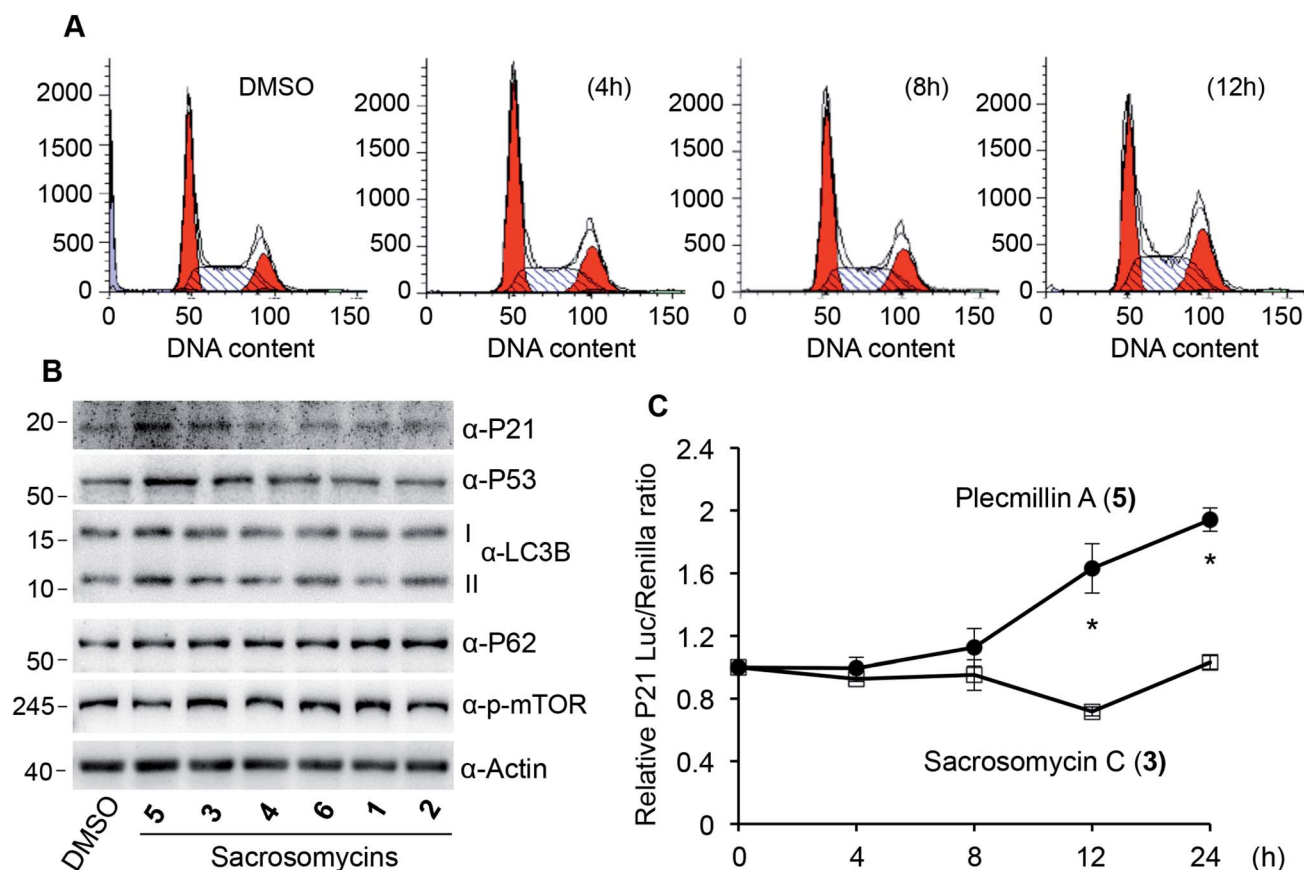


Fig. 5 Cell cycle arrest induced by Plecmillin A (**5**). (A) U2OS cells were treated with $5 \mu\text{M}$ Plecmillin A (**5**) for 0, 4, 8 and 12 hours and cell cycle profile was analyzed by flow cytometry. (B) U2OS cells were treated with $5 \mu\text{M}$ of compounds (**1**–**6**) for 6 hours, and protein expression was examined. I and II represent unlipidated and lipidated form of LC3B, respectively. (C) Luciferase reporter assay for P53-dependent transcription of CDKN1A. HEK293T cells were transfected with PGL3-P21 reporter and renilla luciferase vectors (1 : 50) for 24 hours, treated with $5 \mu\text{M}$ Plecmillin A (**5**) or Sacrosomycin C (**3**) for indicated times. Data represent mean and standard of relative luciferase/renilla ratio from three independent experiments.



the other hand, these compounds did not clearly affect the expression of LC3B or P62 (Fig. 5B), two markers for autophagy, indicating that they did not affect the autophagic pathway.

To confirm that these compounds indeed activated the P53–P21 pathway, we constructed a luciferase reporter assay to measure P53-dependent transcriptional regulation of CDKN1A, the gene encoding P21. The results show that at the dose of 5 μ M, Plecmillin A (5), but not sacrosomycin C (3), time-dependently increased the transcription level of CDKN1A (Fig. 5C). Since P21 inhibits the cyclin-dependent kinase to induce cell cycle arrest in G1 or G2/M phase,^{16,22–24} these results are consistent with the G2/M arrest by Plecmillin A (5).

To determine how spirobisanaphthalenes regulate the P53–P21 pathway and cell survival, we conducted a dose-dependent study of Plecmillin A (5) and sacrosomycin C (3). The results show that Plecmillin A (5), but not sacrosomycin C (3), greatly increased the protein level of P53 and P21 (Fig. 6A). There was a moderate increase in the MEK/ERK pathway as evidenced by a small increase in the levels of phosphorylated MEK and ERK (Fig. 6A); however, this increase in this pathway may have not effect the cell survival. Consistently, we observed a gradual increase in the level of cleaved poly (ADP-ribose) polymerase (PARP), a known cell death marker,^{25–27} by Plecmillin A (5), and to a much lesser degree sacrosomycin C (3) (Fig. 6A). Further, Plecmillin A (5), but not sacrosomycin C (3), dose-dependently reduced the long-term survival capability of U2OS cancer cells

(Fig. 6B–C). These results suggest cell death induction by higher doses of spirobisanaphthalenes.

Further, we found that Plecmillin A (5) time-dependently increased the protein levels of P53 and P21 (Fig. 7A). We noticed a weak induction of PARP cleavage (Fig. 7A), likely due to shorter incubation time than in previous experimental settings. To further confirm the induction of the P53–P21 pathway, we performed quantitative PCR (qPCR) to measure the mRNA levels of TP53 and CDKN1A. Our results show that Plecmillin A (5) induced upregulation of both genes during the first 8 h of treatment but then slightly declined at 24 h of treatment (Fig. 7B). The reduction at 24 h might be due to cell death after prolonged compound treatment. Similar to short-term growth curve results, Plecmillin A (5) time-dependently inhibited the long-term cell survival (Fig. 7C). To determine if this compound directly induces cell death, we measured dead cell population using trypan blue exclusion assay. The results show that Plecmillin A (5) time-dependently induced cell death (Fig. 7D). To understand if this compound could non-selectively kill normal cells, we measured cell death of a normal hepatocyte LO2 after treatment with Plecmillin A or Sacrosomycin C. The results show almost no cell death induction by these two compounds (Fig. 7E), indicating a relatively selective cell death induction in cancer cells by these compounds. Although the magnitude of cell death seemed to be weak, the increase was significant, supporting a chronic tumor suppressing role of this class of compound.

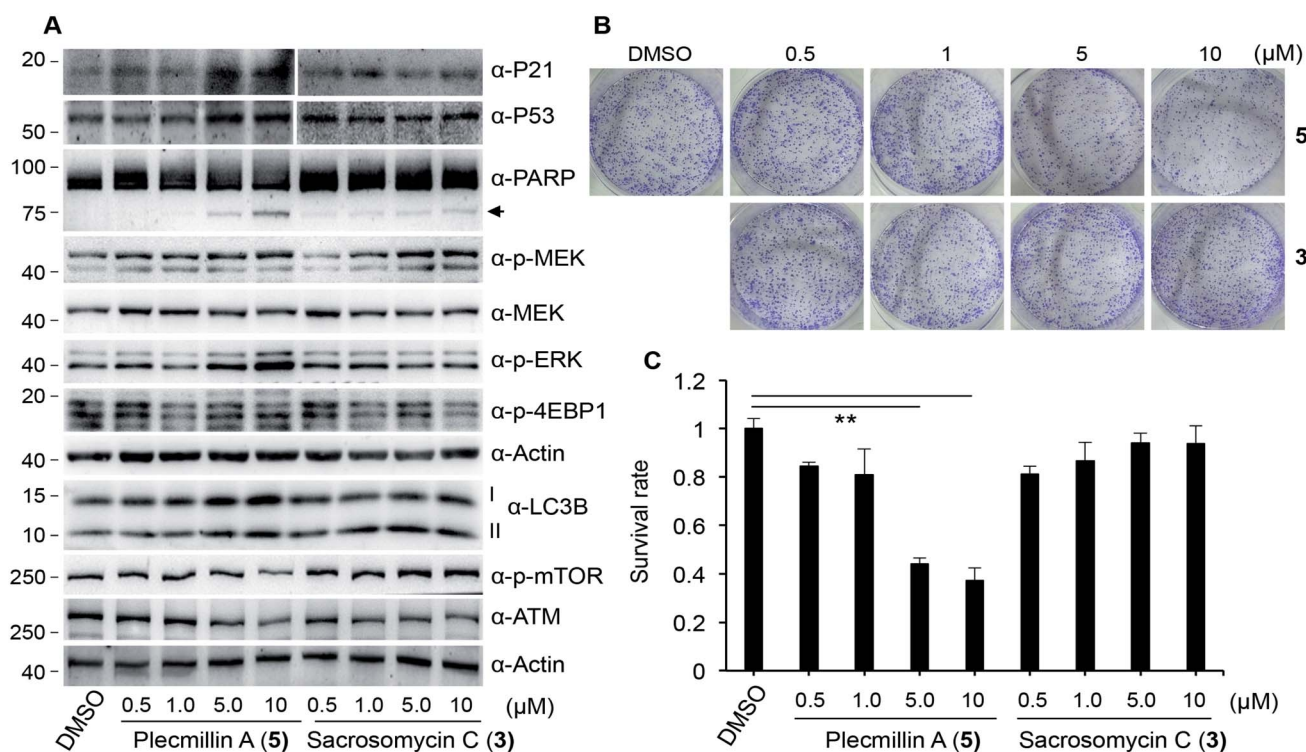


Fig. 6 Dose-dependent P53–P21 induction and cell growth inhibition by Plecmillin A (5). (A) U2OS cells were treated with increasing doses of Plecmillin A (5) or Sacrosomycin C (3) for 12 hours and protein expression was assessed. Arrow indicates cleaved PARP. (B) Representative images of clonogenic survival of cells from A. (C) Quantitation of B. Data represent mean and standard deviation from three replicates. $**P < 0.0001$ compared with control group.



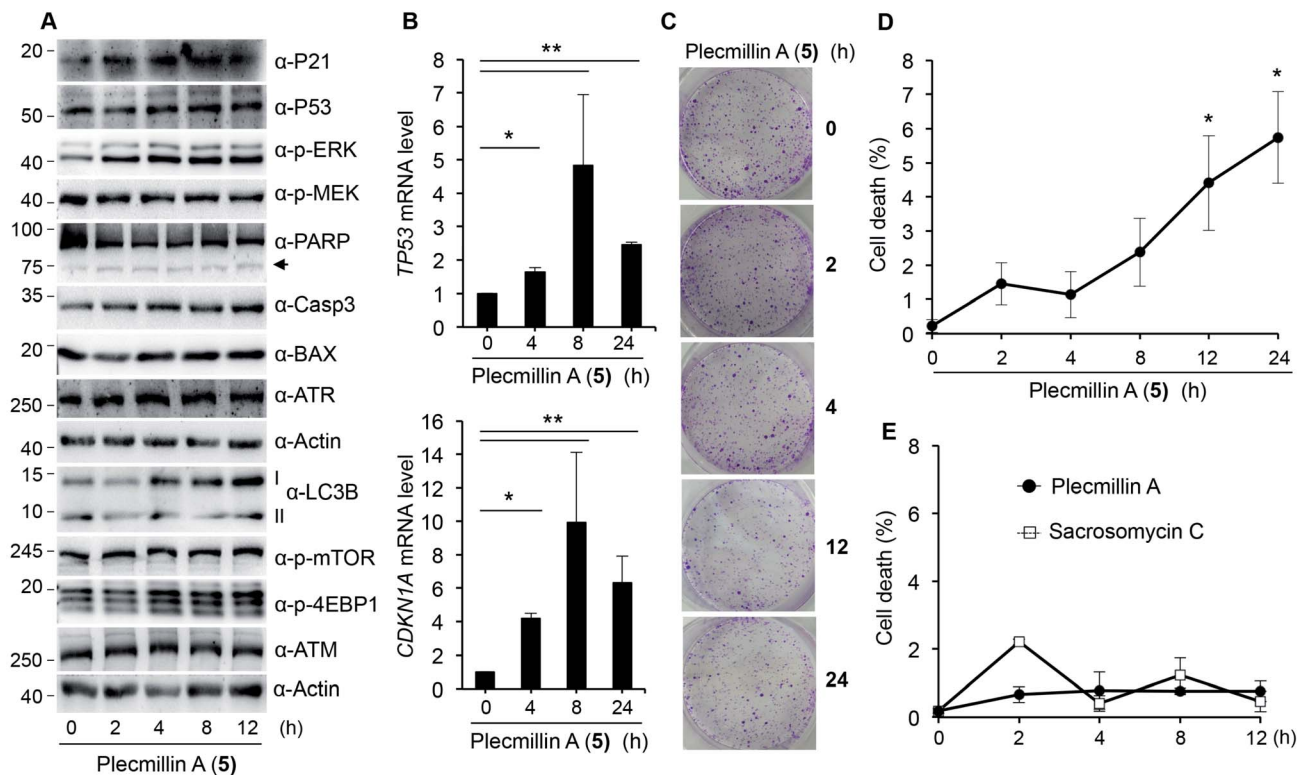


Fig. 7 Time-dependent P53–P21 induction and cell growth inhibition by Plecmillin A (5). (A) U2OS cells were treated with 5 μM of Plecmillin A (5) for 0, 2, 4, 8 and 12 hours and protein expression was assessed. Arrow indicates cleaved PARP. (B) qPCR results of TP53 and CDKN1A in U2OS cells treated with 5 μM of Plecmillin A (5) for 0, 4, 8 and 24 hours. (C) Representative images of clonogenic survival of cells from A. (D) Quantitation of C. (E) LO2 cells were treated with 10 μM of Plecmillin A (5) or Sacrosomycin C for 0, 4, 8 and 12 hours, and cell death was assessed. Data represent mean and standard deviation from three replicates. * $P < 0.01$ and ** $P < 0.0001$ compared with the control group.

Conclusion

In summary, three new and three known spirobisnaphthalenes were obtained from the extract of a cultured nonsporulating endolichenic fungus. The three novel compounds were described as two partially saturated naphthalene rings joined together by one oxygen bridge (C4'–C4) and one carbon–carbon bridge (C4'–C6), which was belonged to the fourth subtype of spirobisnaphthalenes. This is also the first report of spirobisnaphthalenes obtained from endolichenic fungus and from this family Sarcosomataceae. Anticancer activity evaluation reveals that Plecmillin A (5) displayed potent growth inhibitory effect on colon HCT116 and osteosarcoma U2OS cancer cell lines, but not on normal hepatocytes. Mechanically, Plecmillin A (5) induced the upregulation of the P53–P21 pathway, correlating with the cell cycle arrest effect by this compound. Plecmillin A (5) treatment at relatively higher doses induced cancer cell death. These biological results allow us to identify more potent spirobisnaphthalenes, for instance through chemical modification of the three novel compounds, in the anticancer regimen in the near future. In conclusion, spirobisnaphthalenes isolated from an endolichenic fungus strain (CGMCC 3.15192) demonstrated potential anticancer effect.

Experimental procedure

General experimental procedures. Optical rotations were measured on a JASCO P-1020 digital spectropolarimeter. IR spectra were obtained using a JASCO FT/IR-480 plus spectrometer. ESI-MS spectra were recorded on a Finigan LCQ Advantage MAX mass spectrometer. HRESIMS spectra were recorded using a Micromass Q-TOF and an Agilent 6210 LC/MSD TOF mass spectrometer. 1D and 2D NMR spectra were recorded using a Bruker Avance 400 MHz NMR spectrometer. Column chromatography was performed using silica gel (200–300 mesh, Qingdao) and ODS (60–80 μm , YMC). TLC was performed on pre-coated silica gel plates (silica gel GF254, 1 mm). Semipreparative HPLC was performed on a SHIMADZU LC-6AD Liquid Chromatography with SPD-20A Detector, using an ODS column [Welch Materials XB-C18 column (4.6 \times 250 mm, 5 μm), YMC-Pack ODS-A (10.0 \times 250 mm, 5 μm)]. All culture media and glassware were autoclaved prior to use. The slate cultivation was carried out in an EYELA LTI-700 incubator. The seed cultures were grown in an HZQ-Q constant temperature breeding shaker.

Fungal material. The nonsporulating endolichenic fungus strain CGMCC 3.15192 was isolated from lichen *Peltigera elisabethae* var. *mauritzii* (Gyeln.) J. C. Wei collected in the Changbai Mountain in Jilin Province of China. The fungus was



identified as a member of Sarcosomataceae according to its molecular identification by Professor Liang-Dong Guo in the Institute of Microbiology, Chinese Academy of Sciences, Beijing.

Fermentation. The fungus was cultured on slants of potato dextrose agar at 25 °C for 4 days. Agar plugs were used to inoculate Erlenmeyer flasks (250 mL), each containing 50 mL of 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 at 220 rpm for 3–4 days. Fermentation was carried out in thirty Erlenmeyer flasks (500 mL), each containing 70 g of rice. Distilled H₂O (105 mL) was added to each flask, and the contents were soaked overnight before autoclaving at 15 lb per in² for 30 min. After cooling to room temperature, each flask was inoculated with 5.0 mL of the inoculum and incubated at 25 °C for 47 days.

Extraction and isolation. The fermented material was extracted repeatedly with EtOAc (0.5 L × 30 × 4), and the organic solvent was evaporated to dryness under vacuum to afford the crude extract (32.2 g), which was fractionated by silica gel column chromatography eluted with petroleum ether and MeOH successively. The MeOH portion was separated by ODS column chromatography using MeOH–H₂O gradient elution to obtain five fractions (W1–W5). Fractions W2 (50% MeOH in H₂O, v/v) and W3 (70% MeOH in H₂O, v/v) showed different degrees of cytotoxic activities against five cancer cell lines (HL-60, SMMC-7721, A-549, MCF-7 and SW480). The fraction W2 (1.76 g) was chromatographed on a Sephadex LH-20 column eluting with CHCl₃–MeOH 1 : 1 to yield four subfractions (W2-a to W2-d). W2-b (1.15 g) was subjected to a silica gel column chromatography eluted with a stepwise elution of CHCl₃/acetone (from 98 : 2 to 0 : 100) to obtain 14 subfractions. W2-b-h (183 mg) was purified by RPHPLC (40% MeOH in H₂O, v/v) on an YMC-Pack ODS-A column (10.0 × 250 mm, 5 μm, 4.0 mL min⁻¹) to yield compounds **4** (14.5 mg, *t_R* = 23.3 min) and **6** (6.9 mg, *t_R* = 24.1 min). Compound **1** (8.2 mg, *t_R* = 25.3 min) was afforded from the subfraction W2-b-g (52.5 mg) using RPHPLC (40% MeOH in H₂O, v/v). Compound **2** (18.5 mg, *t_R* = 16.3 min) was afforded from the subfraction W2-b-i (52.5 mg) using RPHPLC (25% CH₃CN in H₂O, v/v). Compound **3** (2.5 mg, *t_R* = 15.8 min) was afforded from the subfraction W2-b-72 (50.0 mg) using RPHPLC (20% CH₃CN in H₂O, v/v). Meanwhile, the fraction W3 (1.49 g) was chromatographed on a Sephadex LH-20 column chromatography eluted with CHCl₃–MeOH 1 : 1 to yield five subfractions (W3-a to W3-e). W3-c (0.43 g) was subjected to a silica gel column chromatography eluted with a stepwise elution of CHCl₃/acetone (from 98 : 2 to 0 : 100) to obtain five subfractions. W3-c-2 (50.4 mg) was purified by RPHPLC (40% MeOH in H₂O, v/v) on an YMC-Pack ODS-A column (10.0 × 250 mm, 5 μm, 4.0 mL min⁻¹) to yield compound **5** (7.2 mg, *t_R* = 15.7 min).

Sarcosomycin A (**1**): yellow powder; [α]_D²⁵ 26.8 (*c* = 0.10, MeOH); UV (MeOH): λ_{\max} (log ϵ) = 216 nm (4.41), 255 nm (3.98), 319 nm (3.56); CD (MeOH) 214 ($\Delta\epsilon$ –10.6), 227 ($\Delta\epsilon$ +5.7), 260 ($\Delta\epsilon$ +10.9), 312 ($\Delta\epsilon$ –1.4) nm; IR (KBr): ν_{\max} = 3423, 2925, 2854,

1657, 1594 cm⁻¹; HR-ESI-MS: (*m/z* 383.1116 [M + H]⁺, calcd for C₂₁H₁₉O₇: 383.1131). ¹H and ¹³C-NMR data see Table 1.

Sarcosomycin B (**2**): yellow powder; [α]_D²⁵ 3.7 (*c* = 0.95, MeOH); UV (MeOH): λ_{\max} (log ϵ) = 218 nm (4.34), 254 nm (3.90), 323 nm (3.48); UV (CH₃CN): λ_{\max} (log ϵ) = 218 nm (4.34), 254 nm (3.90), 323 nm (3.48); CD (MeOH): 214 ($\Delta\epsilon$ –11.5), 260 ($\Delta\epsilon$ +9.7), 320 ($\Delta\epsilon$ –1.9) nm; CD (CH₃CN): 214 ($\Delta\epsilon$ –6.3), 227 ($\Delta\epsilon$ +6.7), 260 ($\Delta\epsilon$ +11.6), 315 ($\Delta\epsilon$ –1.7) nm; IR (KBr): ν_{\max} = 3433, 1674, 1656, 1593 cm⁻¹; HR-ESI-MS: (*m/z* 397.1288 [M + H]⁺, calcd. for C₂₂H₂₁O₇: 397.1287). ¹H and ¹³C-NMR data see Table 1.

Sarcosomycin C (**3**): yellow powder; [α] 5.2 (*c* = 0.11, MeOH); UV (MeOH): λ_{\max} (log ϵ) = 202 nm (–4.68), 254 nm (3.98), 317 nm (3.65); IR (KBr): ν_{\max} = 3423, 2924, 2854, 1669, 1591, 1281, 1241 cm⁻¹; HR-ESI-MS: (*m/z* 383.1123 [M + H]⁺, calcd for C₂₁H₁₉O₇: 383.1131). ¹H and ¹³C-NMR data see Table 1.

Biological materials and methods

Cell cultures. Human cancer cell lines were obtained from American Type Culture Collection (ATCC). The cell lines were cultured in Dulbecco's modified eagle medium (DMEM) (for U2OS and HEK293T) or PRMI1640 medium (for HCT116) supplemented with 10% fetal bovine serum (FBS, #FSP500, ExCellBio), 100 U mL⁻¹ penicillin and 100 μg mL⁻¹ streptomycin (#15140–122, Gibco) in 5% CO₂ and 37 °C incubator. The cells were passaged once every 3–4 days, and cells in the logarithmic phase were used for the experiment.

Measurement of cell viability. Cell viability was measured by MTT assay. For IC₅₀ determination, HCT116 cells were plated at the density of 5 × 10⁴/mL in 96 well plate overnight, treated with varying concentrations of compounds for 48 h, add 20 μL MTT (#T100896, Aladdin) into each well and cultured for 3 h, discarded the supernatant, added 150 μL DMSO to each well, and measured the OD value at 570 nm. For measuring cell growth, U2OS cells were seeded at the density of 5 × 10⁴/mL in 96 well plates for 1–5 days, and MTT assay was performed as above. The Graphpad Prism5 was used to calculate the IC₅₀ value.

Western blotting. For the detection of protein expression, 5 × 10⁵ cells were plated at 6 well plates overnight, treated with compounds for 6 h or indicated times. The cells were harvested and lysed in 50 μL lysis buffer (#P0013, Beyotime) on ice for 30 min, sonicated and centrifuged to collect the supernatant. The protein concentration in the supernatant was quantitated by BCA protein assay kit (#23225, Pierce) and equal amount of total proteins from each sample was loaded onto SDS-PAGE followed by detection using specific antibodies. Antibodies used include the anti-P53 (#GTx70214, GeneTex), anti-P21 (#sc-6246, Santa Cruz Biotechnologies), β-actin (AP0060, Bioworld), anti-GADD45a (#ab180768, abcam), anti-Apaf-1 (#ab2000, abcam), anti-MDM2 (#ab38618, abcam), anti-14-3-3 σ (#GTx100801, GeneTex), anti-PUMA (#GTx109675, GeneTex), anti-LC3B (NB100-2220, Novus Biologicals), anti-P62 (#NBPI-42821, Novus biologicals), anti-phospho-Chk1 (Ser345, CST), anti-ATR (#GTx70109, GeneTex), anti-ATM (#GTx70103, GeneTex), anti-PARP (#9542, CST), anti-Phospho-MEK1/2 (Ser217/221, CST), anti-MEK (#9122, CST), anti-P44/42 MAPK (Erk112)



(#9102, CST), anti-Bax (#2772, CST), anti-phospho-mTOR (Ser2448, CST), anti-Caspase3 (#9662, CST), anti-P-4EBP1 (Ser65, CST). All primary antibodies were diluted in TBST buffer. The relative expression levels of P53 and P21 proteins were quantitated by measuring the band intensity using the NIH Image J software.

Real-time PCR. Total RNA was extracted followed by the RNeasy plus mini kit 50 (#74134, Qiagen). The cDNA is synthesized using the Revert Aid first strand cDNA synthesis kit according to the manufacturer's procedure (#K1622, Thermo Fisher). Real-time PCR was performed on a CFX96 Real-Time PCR system (Bio-Rad, Hercules, CA) with SYBR Green (Invitrogen, #208054). Primers for RT-PCR are: TP53-F: 5'-CTTCCATTGCTTTGTCCCG-3', TP53-R: 5'-CATCTCCCAAA-CATCCCTCAC-3', CDKN1A (P21)-F: 5'-AACTAGGCGGTTGAAT-GAGAG-3', CDKN1A (P21)-R: 5'-GAGGAAGTAGCTGGCATGAAG-3', HPRT1-F: 5'-AGCTTGCTGGTGAAGGA-3', HPRT1-R: 5'-CCAACTCAACTTGAAGTCTCATC-3'.

Luciferase reporter assay. To analyze P53-dependent transcriptional regulation of CDKN1A expression, we carried out luciferase reporter assay. Briefly, 8 copies of the P53-binding site of the CDKN1A's promoter sequence were inserted into the PGL3-Basic plasmid. HEK293T cell were transfected with the P21 luciferase reporter plasmid together with a renilla control vector by Lipo2000 (#11668019, ThermoFisher). After 24 h, the cells were treated with 5 μ M compounds for 0, 4, 8, 12 and 24 h. Luciferase activities were measured followed the Dual-Luciferase® Reporter Assay System Kit (#E1910, Promega).

Cell cycle assay. U2OS cell were treated with 5 μ M Plecmillin A for 0, 4, 8 and 12 hours, harvested, washed once with PBS, fixed in 70% ethanol, stained with PI in the presence of RNase A (#C1052, Beyotime), analyzed by flow cytometry. The data were analyzed using the ModFitLT software.

Clonogenic survival assay. U2OS cells were treated with different concentrations of Plecmillin A for 12 or varying hours, trypsinized into single cell suspension, replated at the density of 500 cells per well into 6 well plates and incubated for 10 days. Colonies were stained with 0.1% crystal violet, washed under tap water, air-dried and images were taken. Quantitation of cell survival was performed either by quantitating the image intensity or dissolving the plates by 1% SDS and measuring the absorption at 570 nm. All of the experimental groups were repeated three times.

Conflicts of interest

There are no conflicts to declare.

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