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Study on secondary metabolites of endophytic fungus *Diaporthe* sp. AC1 induced by tryptophan analogs

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Small molecule-induced fermentation of the endophytic fungus Diaporthe sp. AC1 originated from Artemisia argyi was executed to investigate its secondary metabolites. It was fermented in a culture medium containing 5-hydroxytryptophan (5-HTP), 1-methyl-L-tryptophan (1-MT), and tryptamine (TA), respectively. The antibacterial activities of crude extracts against pathogenic bacteria and pathogenic fungi were determined by using the Oxford cup method, while the cytotoxicity of crude extracts against cancer cells was determined by using the MTT method. The results showed that the secondary metabolites of Diaporthe sp. AC1 induced by 1-MT exhibited optimal antibacterial activity and tumor cytotoxicity. The induction conditions of 1-MT were optimized, and the antibacterial activities and tumor cytotoxicity of crude extracts under different induction conditions were investigated. As indicated, the optimal moment for 1-MT addition was before inoculation and its optimal concentration was 0.25 mM. Under these conditions, *Diaporthe* sp. AC1 was fermented and approximately 12 g of crude extracts was obtained. The crude extracts were then separated and purified to acquire nine monomer compounds, including three new compounds (1-3) and six known compounds (4-9). The antibacterial activities of the compounds against pathogenic bacteria and pathogenic fungi were investigated by using the microdilution method, while their cytotoxicity against cancer cells was analyzed by using the MTT method. The results demonstrated that Compound 1 exhibited moderate antibacterial activities against Verticillium dahlia, Fusarium graminearum, and Botrytis cinerea, as well as a low inhibitory activity against Listeria monocytogenes. Nevertheless, Compound 1 showed significant cytotoxicity against five cancer cells, with IC_{50} ranging from 12.26 to $52.52 \,\mu$ M. Compounds **2** and **3** exhibited negligible biological activity, while other compounds showed detectable inhibitory activities against pathogenic bacteria and cancer cells.

KEYWORDS

Diaporthe, endophytic fungi, 1-methyl-L-tryptophan, secondary metabolites, biological activity

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1. Introduction

Endophytic fungi are characterized by rich diversity of species (Gokhale et al., 2017) and capability to generate compounds with pharmaceutical and commercial values (Deshmukh et al., 2018; Gao et al., 2018), thus being regarded as a goldmine of secondary metabolites (Uzma et al., 2018; White et al., 2019). Recent studies of the whole fungal genomes have demonstrated the presence of various silent gene clusters, indicating that a large proportion of new compounds remain to be identified (Hautbergue et al., 2018). Therefore, maximized activation of silent gene clusters has been playing a key role in efficient utilization of fungal resources in the post-genomic era.

One Strain Many Compounds (OSMAC) strategy is a facile and effective strategy to identify novel active metabolites (Liu et al., 2017). By tuning the medium composition (Ranuka et al., 2014) and culture conditions (Siridechakorn et al., 2017), co-culturing with other strains (Wang et al., 2022), and adding small molecule inducers (Gakuubi et al., 2022; Li et al., 2022), the metabolic pathway of secondary metabolites can be adjusted to identify novel secondary metabolites (Pinedo-Rivilla et al., 2022). Specifically, the addition of small molecule inducers leads to significant increase in the variety of secondary metabolites. It has been demonstrated that the addition of tryptophan analog 1-methyl-L-tryptophan (1-MT) into the culture of Chaetomium globosum 1C51 could activate cryptic gene and trigger the Pictet-Spengler reaction of 1-MT and flavipin to produce new indole alkaloids (Yan et al., 2014). Additionally, tryptamine, another analog of tryptophan, has been used as a "scavenger" to capture the biosynthetic intermediates of Diaporthe sp., resulting in the identification of novel alkaloids originated from tryptamine-exposed fungal culture (Chen et al., 2021).

Diaporthe is typically isolated as endophytic fungi and secondary metabolites produced by this genus have been thoroughly investigated (Chepkirui and Stadler, 2017). The endophytic fungus Diaporthe sp. AC1 derived from Artemisia argyi can produce various active natural products, including phomopsolides, pyranones, and indole alkaloids (Gu et al., 2022). The structural diversities and biological activities of the compounds encouraged further study on the use of Diaporthe sp. AC1 for the production of new metabolites by the "OSMAC" technique. Since metabolites production of endophytic fungi can be influenced by adding small molecule inducers as mentioned earlier, three analogs of tryptophan [1-MT, tryptamine, and 5-hydroxytryptophan (5-HTP)] were added into the fermented culture of this strain, with the aim of generation of new fungal metabolites. Herein, 1-MT was screened as the optimal inducer, and three new compounds (1-3) as well as six known ones were produced by the endophytic fungus Diaporthe sp. AC1 after adding 1-MT into media for fungal cultivation.

2. Materials and methods

2.1. Chemicals and reagents

The media used for the experiments included the potato dextrose agar (PDA) medium (200 g of potato, 20 g of dextrose, and 15 g of agar in 1000 mL of deionized water), the malt extract (ME) liquid medium The organic solvent used for chromatographic separation was purchased from Tianjin Kaitong Chemical Reagent Co., Ltd., China. The silica gel (200–300 mesh) used for column chromatography was purchased from Qingdao Ocean Chemical Co., Ltd., China. Sephadex LH-20 gel was purchased from GE Healthcare. Thin-layer chromatography silica gel plates (GF254) were purchased from Qingdao Ocean Chemical Co., Ltd., China. 5-HTP, 1-MT, and tryptamine (TA) were purchased from Shanghai Aladdin Biochemical Technology Co., Ltd., China.

2.2. Fungal and bacterial materials

Diaporthe sp. AC1 was isolated from *Artemisia argyi* and identified by following a previously reported protocol (Gu et al., 2022). The ITS sequence data of this fungal strain have been submitted to the GenBank under the accession number of OL589639.

Pathogenic bacteria *Staphylococcus aureus* ATCC25923, *Listeria monocytogenes* CGMCC1.10753, *Salmonella enteritidis* ATCC13076, *Pseudomonas aeruginosa* ATCC27853 and *Escherichia coli* ATCC8739, as well as pathogenic fungi *Candida albicans* ATCC10231 were purchased from Shanghai Bioresource Collection Center (SHBCC). Pathogenic fungi *Fusarium graminearum*, *F. moniliforme, Botrytis cinerea* and *Verticillium dahlia* were isolated, identified and preserved by our laboratory (Gu et al., 2022).

2.3. Screening of tryptophan analogs

Diaporthe sp. AC1 was inoculated on PDA medium and incubated at 28°C for 5 days. Five disks (diameter = 1 cm) were then inoculated in 100 mL of ME liquid medium and shaken under 180 rpm at 28°C for 2 days. Afterwards, 5-HTP, 1-MT, and TA were separately added to the fungal cultures at 4×12 h intervals until a final concentration of 1 mM (Yan et al., 2019). After that, the fermentation broths were incubated for another 3 days. Herein, a fermentation broth without tryptophan analogs was used as the control group.

Whole culture broths (including mycelia) were extracted with equal volume of ethyl acetate for two times. Meanwhile, organic solvents were evaporated under reduced pressure to obtain crude extracts. Additionally, the antibacterial activities and cytotoxicity of the crude extracts were evaluated.

2.4. Optimization of induction conditions

According to the methods described before (Yan et al., 2019) with some modification, the addition moment and concentration of the tryptophan analogs were optimized as follows: for Groups 1 and 2, the inducer was added simultaneously with mycelia before fermentation at a concentration of 1 and 0.25 mM, respectively; for Group 3, the inducer was added after two-day fermentation at a concentration of 1 mM; for Group 4, inducer was added to culture after two-day fermentation at $4 \times 12h$ intervals until a concentration of 1 mM was reached.

2.5. Preparation of crude extract

Under optimized induction conditions, *Diaporthe* sp. AC1 was fermented to obtain crude extracts. Briefly, *Diaporthe* sp. AC1 was inoculated in the PDA medium and incubated at 28° C for 5 days. Subsequently, five disks (diameter = 1 cm) were inoculated in 100 mL of ME liquid medium and shaken under 180 rpm at 28° C for 5 days to obtain the seed solution, which was then transferred to 2.5 L of ME medium for induced fermentation. Herein, 50 L of culture (200 bottles of 2.5 L medium) was fermented. After fermentation, the crude extract was extracted and concentrated by using the method described above. Additionally, antibacterial activities and cytotoxicity of the crude extracts were evaluated.

2.6. Antibacterial activity of crude extract

As described in our previous study (Gu et al., 2022), the antibacterial activities of the crude extracts against five pathogenic bacteria and five pathogenic fungi were investigated by using the Oxford cup method. Firstly, four Oxford cups were placed uniformly on top of water agar which was poured into the Petri dish beforehand. Then, $150 \,\mu$ L of pathogen suspension ($1.0 \sim 5.0 \times 10^7 \,\text{CFU/mL}$) and $15 \,\text{mL}$ of medium (LB agar medium for bacteria, PDA medium for fungi) were poured into the Petri dish. After solidification, the Oxford cups were removed, resulting in generation of four small wells in the upper medium.

In the experimental group, $50\,\mu$ L of the crude extract solution ($20\,\text{mg/mL}$, dissolved in methanol) was added to each well. Ampicillin sodium ($20\,\text{mg/mL}$) and amphotericin B ($20\,\text{mg/mL}$) were used as the positive control of antagonistic bacteria and fungi, respectively. Then, $50\,\mu$ L of methanol was added to the negative control wells. The plates inoculated with bacteria were cultured at 37° C for 24 h, and those containing fungi were cultured at 28° C for $5{\sim}7$ days. Additionally, the diameters of inhibition zones were measured.

2.7. Cytotoxicity of crude extract

The cytotoxicity of the crude extracts against five cancer cells (human hepatoma cell line Huh-7, human cervical cancer cell line Hela, human colorectal adenocarcinoma cell line HCT-15, human lung cancer cell line A549, human breast cancer cell line MDA-MB-231) was analyzed by using the MTT method (Mosmann, 1983). Specifically, cancer cells were added to a 96-well plate $(4 \times 10^4 \text{ cell/mL})$ in DMEM (Dulbecco's Modified Eagle Medium) containing 10% FBS, 100 μ L/well) and incubated at 37°C (7% CO₂) for 20 h. Then, 100 μ L of the cell culture media containing compounds of different concentrations (50, 25, 12.5, 6.25 µg/mL) was added into different wells; the well added with 100 µL of the cell culture medium with no compound was used as the negative control, while the blank control group contained culture medium but no cells. After 48 h, the plate was rinsed once with PBS buffer (200 µL/well) and MTT solution (1 mg/mL in DMEM containing 10% FBS, 100 µL/well) was added, followed by another four-hour incubation. After that, the solution was removed and DMSO (150 µL/well) was added to dissolve the formazan crystals generated during incubation. After 10 min of shaking, the absorbance was measured at 570 nm. Each group had three replicates. The inhibition rates of the compounds were calculated by:

Inhibition rate
$$\binom{\%}{=} \left(1 - \frac{A_1 - A_0}{A_2 - A_0}\right) \times 100\%$$

where A₀, A₁, and A₂ refer to the absorbance of the blank control, the experimental group, and the negative control, respectively.

2.8. Purification and characterization of compounds

As shown in Figure 1, the crude extract was eluted by silica gel column chromatography with a gradient elution of PE-EtOAc (petroleum ether – ethyl acetate) to obtain 47 fractions (Fr. 1 ~ Fr. 47). Then, TLC was used to analyze each fraction and similar ones were combined, resulting in generation of seven subfractions (Fr. A ~ Fr.I). Among them, subfractions C, D, and E were further separated by using the C18 reverse-phase silica gel column with 50% ~ 100% methanol as the eluent. Additionally, the fractions were further separated by using the Sephadex LH-20 gel column.

The compounds were dissolved with CDCl₃ or acetone- d_6 , and identified by nuclear magnetic resonance (NMR), high-resolution mass spectrometry (HR-MS), and/or two-dimensional nuclear magnetic resonance (2D-NMR). ¹H NMR (600 MHz) and ¹³C NMR (151 MHz) spectra of the samples were obtained by using the NMR spectrometer (Bruker Avance, Germany) with tetramethylsilane (TMS) as the internal standard. Additionally, HR-MS spectra of the samples were obtained by using an AB SCIEX X500R QTOF mass spectrometer.

2.9. Antimicrobial activity of compounds

Preparation of bacterial suspension: Pathogenic bacteria were inoculated in LB liquid medium, and incubated at 37°C 200 rpm for 12 h. The bacterial culture was then diluted with LB liquid medium so that its OD₆₀₀ value reached to 0.2. The culture was further diluted 300 times to obtain a bacterial suspension in which the concentration was $1 \sim 5 \times 10^5$ CFU/mL.

Preparation of fungal suspension: Pathogenic fungi were incubated at 28°C on PDA medium for $3 \sim 7$ days, the dishes were then rinsed with sterile water to prepare spore suspension. The spores were counted using the Hemocytometer, and the concentration of spores was diluted to $1 \sim 5 \times 10^5$ CFU/mL with PDB medium.

Modified microdilution method (Zgoda and Porter, 2001) was used to evaluate the antimicrobial activities of the compounds against pathogenic bacteria and fungi. Firstly, 100 μ L of the cell suspension (containing $1 \sim 5 \times 10^5$ CFU/mL bacteria or fungal spores) was added to each well of a 96-well plate. Then, 100 μ L of the culture medium (LB medium for bacteria and PDB medium for fungi) containing compounds of different concentrations (512, 256, 128, 64, 32 μ g/mL) was added into each well. The well added with 100 μ L of culture medium but no compound was used as the negative control, while the



blank control contained culture medium but no pathogenic bacteria or compounds. Each group had three replicates. The 96-well plate inoculated with bacteria was incubated at 37°C for 24 h, while the plate containing fungi was incubated at 28°C for 48 h. The absorbance was measured at 600 nm. The inhibition rates of compounds were calculated by:

Inhibition rate
$$(\%) = \left(1 - \frac{A_1 - A_0}{A_2 - A_0}\right) \times 100\%$$

where A₀, A₁, and A₂ refer to the absorbance of the blank control, the experimental group, and the negative control, respectively.

2.10. Cytotoxicity of compounds

The cytotoxicity of the compounds against five cancer cells was determined by using the MTT method as described above. The final concentrations of the compounds ranged from 200 to $12.5\,\mu M$ with two-fold serial dilution.

3. Results and discussion

3.1. Screening of tryptophan analogs

As indicated by previous studies, tryptophan analogs 1-MT could change the metabolic pathway of *Chaetomium globosum* 1C51 and

generate novel active compounds (Yan et al., 2019), and addition of tryptamine to *Diaporthe* sp. resulted in three novel alkaloids (Chen et al., 2021). In order to further explore the active metabolites of *Diaporthe* sp. AC1, an endophytic fungus from *Artemisia argyi* with antimicrobial potential, was fermented with three tryptophan analogs (5-HTP, 1-MT, and TA), respectively.

The results in Supplementary Table S1 revealed that the yields of crude extracts from fermentation broth added with tryptophan analogs were significantly higher than that of the control group (CK), indicating that addition of three amino acid analogs has positive impacts on the yield of fermentation products. The yield of fermentation products followed the order of TA (305.0 mg/L) > 5-HTP (296.5 mg/L) > MT (253.0 mg/L).

The antibacterial activities of the crude extracts are summarized in Table 1 and Supplementary Figure S1. Compared with those induced by no tryptophan analogs, crude extracts induced by 5-HTP exhibited enhanced antibacterial activity against *L. monocytogenes* and *S. aureus*, and those induced by TA exhibited enhanced inhibitory activity against *E. coli*, *S. enteritidis*, and *P. aeruginosa*. In particular, crude extracts induced by 1-MT exhibited optimized activity, as reflected by enhanced inhibitory activity against all of the five pathogenic bacteria involved.

According to Table 2 and Supplementary Figure S2, the antifungal activities of crude extracts induced by 5-HTP and TA decreased compared with those of crude extracts induced by no tryptophan analogs. However, the crude extracts induced by 1-MT exhibited a slight increase in inhibitory activity against *C. albicans*, *F. moniliform*, and *B. cinerea*.

The cytotoxicity of crude extracts is summarized in Figure 2. All crude extracts involved in this study exhibited inhibitory activities against the five cancer cells in a dose-dependent manner. Compared with the control group, the crude extracts induced by 5-HTP and 1-MT exhibited enhanced cytotoxicity against the five cancer cells, while those induced by TA showed enhanced activity against Huh-7, HCT-15, A549, and MDA-MB-231. Additionally, crude extracts induced by 1-MT exhibited the optimal inhibitory activity, with inhibitory rates above 90 and 65% at concentrations of 50 and 25 μ g/mL, respectively.

In summary, crude extracts induced by 1-MT exhibited the optimal antimicrobial and inhibitory performance toward cancer cells. Hence, 1-MT was employed as the inducer for subsequent experiments.

3.2. Optimization of induction conditions

The yields and characteristics of crude extracts induced by 1-MT are shown in Supplementary Table S2. 1-MT-1 and 1-MT-2 denoted crude extracts induced by 1 and 0.25 mM 1-MT, respectively; 1-MT-3 represented the crude extract generated by adding 1 mM 1-MT after two-day fermentation; 1-MT-4 indicated the crude extract generated by adding 1-MT after two-day fermentation at 4×12 h intervals until a concentration of 1 mM was reached. The yields of 1-MT-2, 1-MT-4, 1-MT-1, and 1-MT-3 were 251.0, 253.0, 242.0, and 240.0 mg/L, respectively.

The inhibition activities of the crude extracts fermented under four induction conditions against pathogenic bacteria and pathogenic fungi are shown in Tables 3, 4, and Supplementary Figures S3, S4. 1-MT-2 exhibited the optimal inhibition performance, with the diameters of inhibition zones of L. monocytogenes, S. enteritidis, S. aureus, F. moniliforme, and B. cinerea being above 20 mm. Moreover, its antibacterial activities against E. coli, C. albicans, V. dahlia, and F. graminearum were also considerably high, with the diameters of inhibition zones ranging from 15.93 to 18.47 mm. The inhibition performance of 1-MT-4 was also rational, with the diameters of the inhibition zones for pathogenic bacteria and pathogenic fungi ranging from 14.83 to 19.03 mm and from 12.03 to 17.80 mm, respectively. Additionally, 1-MT-1 and 1-MT-3 exhibited poorer antimicrobial activities against pathogenic bacteria and fungi. Therefore, it can be deduced that direct addition of excessive 1-MT has negative impacts on the yield of active compounds.

Crude extracts fermented under four induction conditions showed different inhibitory activities against the five cancer cells (Figure 3). 1-MT-1, 1-MT-2, and 1-MT-4 exhibited high cytotoxicity against the five cancer cells, with inhibition rates >70% at a concentration of 32 μ g/mL. Among them, 1-MT-2 exhibited the highest cytotoxicity, with inhibition rates of Huh-7, HCT-15, A549 and MDA-MB-231>50% at a concentration of 8 μ g/mL.

Overall, the optimal induction conditions were the addition of 0.25 mM 1-MT and mycelium before fermentation.

3.3. Compounds elucidation

As demonstrated in Figure 1, Compounds 2 (4.0 mg), 3 (3.6 mg), 4 (69.3 mg), and 5 (4.4 mg) were isolated from Fr.C; Compounds 7

Number	Inhibition zone diameter (mm)					
	L. monocytogenes	E. coli	S. enteritidis	S. aureus	P. aeruginosa	
СК	15.73 ± 0.21	11.73 ± 0.21	13.10 ± 0.20	17.13 ± 0.35	12.90 ± 0.30	
5-HTP	18.17 ± 0.31	-	-	18.57 ± 0.21	12.23 ± 0.31	
1-MT	20.03 ± 0.50	16.97 ± 0.21	20.27 ± 0.32	19.43 ± 0.21	16.70 ± 0.36	
ТА	14.20 ± 0.30	14.57 ± 0.31	17.77 ± 0.25	13.73 ± 0.21	13.93 ± 0.15	
Ampicillin sodium	34.17 ± 0.35	31.93 ± 0.15	34.17 ± 0.31	34.07 ± 0.38	27.03 ± 0.42	
МеОН	-	-	-	-	-	

TABLE 1 Antimicrobial effects of four kinds of crude extracts on pathogenic bacteria.

"-" no obvious antimicrobial effect. The outer diameter of the Oxford cup was 8 mm. The concentration tested was 20 mg/mL. CK, crude extract without tryptophan analogs; 5-HTP, crude extract added with 5-HTP; 1-MT, crude extract added with 1-MT; TA, crude extract added with TA. Three replicates were carried out for each test.

TABLE 2 Antimicrobial effects of four kinds of crude extracts on pathogenic fungi.

Number	Inhibition zone diameter (mm)					
	C. albicans	V. dahliae	F. moniliforme	F. graminearum	B. cinerea	
СК	11.07 ± 0.15	13.50 ± 0.36	16.57 ± 0.68	15.80 ± 0.30	15.63 ± 0.61	
5-HTP	-	10.67 ± 0.15	12.53 ± 0.40	13.60 ± 1.35	13.87 ± 0.35	
1-MT	12.00 ± 0.26	12.83 ± 0.31	16.73 ± 0.47	12.43 ± 0.76	17.57 ± 0.35	
TA	-	-	10.83 ± 0.31	15.67 ± 0.55	-	
Amphotericin B	16.80 ± 0.30	18.77 ± 0.31	17.10 ± 0.30	14.67 ± 0.55	20.50 ± 1.40	
МеОН	-	-	-	-	-	

"-" no obvious antimicrobial effect. The outer diameter of the Oxford cup was 8 mm. The concentration tested was 20 mg/mL. CK, crude extract without tryptophan analogs; 5-HTP, crude extract added with 5-HTP; 1-MT, crude extract added with 1-MT; TA, crude extract added with TA. Three replicates were carried out for each test.

TABLE 3 Antimicrobial effects of four crude extracts induced b	by 1-MT	on pathogenic bacteria.
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Crude	Inhibition zone diameter (mm)					
	L. monocytogenes	E. coli	S. enteritidis	S. aureus	P. aeruginosa	
1-MT-1	16.93 ± 0.15	11.40 ± 0.26	13.07 ± 0.38	17.50 ± 0.87	-	
1-MT-2	22.37 ± 0.45	18.27 ± 0.21	21.23 ± 0.38	22.93 ± 0.51	13.30 ± 0.20	
1-MT-3	16.73 ± 0.21	10.97 ± 0.12	12.47 ± 0.35	17.43 ± 0.31	12.93 ± 0.15	
1-MT-4	18.63 ± 0.15	16.20 ± 0.36	18.07 ± 0.38	19.03 ± 0.21	14.83 ± 0.31	
Ampicillin sodium	33.45 ± 0.80	30.30 ± 0.72	30.97 ± 0.95	32.53 ± 0.71	23.77 ± 0.70	
МеОН	-	-	-	-	-	

"-" no obvious antimicrobial effect. The outer diameter of the Oxford cup was 8 mm. The concentration tested was 20 mg/mL. Three replicates were carried out for each test.

Crude	Inhibition zone diameter (mm)						
	C. albicans	V. dahliae	F. moniliforme	F. graminearum	B. cinerea		
1-MT-1	11.17 ± 0.25	10.83 ± 0.06	17.30 ± 0.10	15.67 ± 1.15	15.10 ± 0.20		
1-MT-2	15.93 ± 0.15	17.30 ± 1.35	21.10 ± 0.53	18.47 ± 1.31	21.83 ± 1.30		
1-MT-3	10.87 ± 0.21	13.40 ± 0.66	14.57 ± 0.31	12.77 ± 1.30	13.87 ± 0.35		
1-MT-4	12.03 ± 0.21	14.80 ± 0.30	17.80 ± 0.66	13.10 ± 0.56	13.83 ± 0.31		
Amphotericin B	16.60 ± 0.62	20.67 ± 0.47	15.90 ± 0.80	13.97 ± 0.50	21.77 ± 0.45		
МеОН	-	-	-	-	-		

"-" no obvious antimicrobial effect. The outer diameter of the Oxford cup was 8 mm. The concentration tested was 20 mg/mL. Three replicates were carried out for each test.

(7.4 mg) and **8** (66.6 mg) were isolated from Fr.D; and Compounds **1** (62.5 mg), **6** (10.0 mg), and **9** (6.9 mg) were isolated from Fr.E.

Compound 1 is reddish-brown oil with a molecular formula of $C_{10}H_{10}O_4$, as deduced from HR-ESI-MS ion peak at m/z 195.0652 [M+H]⁺ (Supplementary Figures S5). The NMR spectra can be found in Supplementary Figures S6–S11. The ¹³C NMR (Table 5) and HSQC data of Compound 1 revealed the presence of 2*H*-pyran-2-one moiety, including three methine (δ_C 145.70 (4), 114.57 (3), 104.35 (5)), and two quaternary carbons (δ_C 166.41 (6), 162.93 (2)). The data in ¹H NMR spectrum displayed singlets for two methylene (δ_H 3.00 (t, 7.3 Hz, 2H, 2'), 2.65 (t, 7.3 Hz, 2H, 1')) as well as a methyl (δ_H 2.15 (s, 3H, 5')), and that in ¹³C NMR indicated the presence of two carbonyl (δ_C 198.52 (3'), 198.14 (4')). Key HMBC correlations (Figure 4) from H-5' to C-4', C-3' established the propanedione moiety. The HMBC correlation of H-2' with C-1' and C-6 indicated that propanedione moiety was connected to the 2*H*-pyran-2-one moiety. Then the structure of Compound 1 was established.

TABLE 5 NMR data of compound 1 (600 MHz, Acetone-d₆).

No.	1				
	δ_{C}	$\delta_{\scriptscriptstyle H}$ (mult. J)			
2	162.93				
3	114.57	5.96 (dd, 9.4, 0.9 Hz, 1H)			
4	145.70	7.28 (dd, 9.4, 6.6 Hz, 1H)			
5	104.35	6.06 (dq,6.6, 0.9 Hz, 1H),			
6	166.41				
1'	28.60	2.65 (t, 7.3 Hz, 2H)			
2'	34.03	3.00 (t, 7.3 Hz, 2H)			
3'	198.52				
4'	198.14				
5'	24.31	2.15 (s, 3H)			



Compound **2** obtained as pale-yellow oil, was determined to be $C_{16}H_{19}NO_4$ by HR-ESI-MS (m/z 312.1206 [M+Na]⁺) (Supplementary Figure S12) and was in accordance with the ¹³C NMR data (Table 6). The NMR spectra were shown in Supplementary Figures S13–S18. In the ¹H NMR spectrum, the characteristic signals at $\delta_{\rm H}$ 6.91 (s, 1H, 2), 7.61 (dt, 7.9 Hz, 1H, 4), 7.11 (ddd, 8.0, 7.0, 1.0 Hz, 1H, 5), 7.23 (ddd, 8.2, 6.9, 1.1 Hz, 1H, 6), 7.29 (d, 8.2 Hz, 1H, 7) and 3.76 (s, 3H, 10) were ascribed to a 1-methylindole moiety. The ¹³C NMR and HSQC data displayed singlets for a methyl ester ($\delta_{\rm C}$ 172.33 (14), 51.85 (15)), an ethyl ester ($\delta_{\rm C}$ 24.66 (8), 65.05 (9), 172.82 (11)) and two methylene ($\delta_{\rm C}$ 29.20 (12), 28.92 (13)). Key HMBC correlations (Figure 5) from H-13 to C-11, and H-12 to C-14 determined the connection of two esters. The HMBC correlation of H-9 with C-3 finally established the structure of Compound **2**.

Compound **3** is yellow oil with a molecular formula of $C_{22}H_{22}N_2O_2$, as deduced from HR-ESI-MS at m/z 369.1573 [M+Na]⁺ (Supplementary Figure S19). Comparison the NMR data of Compound **3** and fusariumindole A (Guo et al., 2020; Table 7)



revealed that Compound **3** had two extra methyl groups ($\delta_{\rm C}$ 32.69 and $\delta_{\rm C}$ 32.56). ¹H-¹H COSY and HMBC spectra demonstrated that the two imino groups in fusariumindole A were methylated (Supplementary Figures S20–S25). Hence, Compound **3** was identified as a methyl derivative of fusariumindole A and denoted as fusariumindole D (Figure 6).

Based on the NMR data of compounds **4–9** in the Supplementary Figures S26–S37, six known compounds were identified as *N*-methyl tryptophol (**4**) (Liu et al., 2012), 3-(2-acetoxyethyl)-1-methylindole (**5**) (Leeson, 1984; Shaker et al., 2021), *N*-methyl-3a-hydroxyfuroindoline (**6**) (Hirose et al., 2005),

3,5-dimethyl-8-methoxy-3,4-dihydroisocoumarin (7) (Kamisuki et al., 2007), p-hydroxyphenethyl alcohol (8) (Jang et al., 2014), 3-phe-nylpyrazin-2(1*H*)-one (9) (El Euch et al., 2018). The structures of compounds **4–9** are shown in Figure 7.

Conventional culture of *Diaporthe* sp. AC1 led to identification of eight compounds with no indole alkaloids (Gu et al., 2022), while in this study, five derivatives of the 1-MT (Compounds 2–6) were obtained. Researches have demonstrated that 1-MT not only could elicit the production of Pictet-Spengler reaction-based alkaloids (Yan et al., 2014), but can also be converted into novel indoles through biotransformation by *Chaetomium globosum* (Yan et al., 2019). The



TABLE 6 NMR data of compound 2 (600 MHz, CDCl₃).

No.	2					
	δ_{C}	$\delta_{\scriptscriptstyle H}$ (mult. J)				
2	126.90	6.91 (s, 1H)				
3	110.33					
4a	127.84					
4	118.85	7.61 (dt, 7.9 Hz, 1H)				
5	118.89	7.11 (ddd, 8.0, 7.0, 1.0 Hz, 1H)				
6	121.65	7.23 (ddd, 8.2, 6.9, 1.1 Hz, 1H)				
7	109.22	7.29 (d, 8.2 Hz, 1H),				
7a	136.92					
8	24.66	3.09 (td, 7.2, 0.8 Hz, 2H),				
9	65.05	4.36 (dd, 9.2, 5.2 Hz, 2H)				
10	32.65	3.76 (s, 3H)				
11	172.82					
12	29.20	261.266 (m. 411)				
13	28.92	2.61–2.66 (m, 4H)				
14	172.33					
15	51.85	3.68 (s, 3H)				

results herein proved that 1-MT could also be transformed by *Diaporthe* sp. AC1. The indole structure motif occurs in a variety of natural products as well as in synthetic drugs (Hu et al., 2021). This nucleus is associated with a broad spectrum of biological profiles ranging from antifungal and antibacterial to cytotoxic activities (Zhang et al., 2014; Li et al., 2016; Han et al., 2020). Hence, the compounds were used to determine antimicrobial and cytotoxic activities.

3.4. Antimicrobial activity of compounds

The antibacterial activities of the compounds against pathogenic bacteria were relatively weak, thus only the antibacterial effect at the concentration of 512µg/mL was shown. As displayed in Figure 8, among the new compounds, Compound 1 exhibited high inhibition rate (> 90%) against L. monocytogenes, while Compounds 2 and 3 exhibited low inhibitory activities against pathogenic bacteria. For the known compounds, Compounds 4 and 9 exhibited good antibacterial activities against five pathogenic bacteria with all inhibition rates above 80%, and Compound 8 showed promising inhibitory performance on P. aeruginosa, with an inhibition rate>90%. As reported before, Compound 9 showed good inhibitory activity against S. aureus, L. monocytogenes, and S. typhimurium (El Euch et al., 2018). The results of this study demonstrated that Compound 9 also had significant antibacterial activity against S. enterica, P. aeruginosa, and E. coli, indicating that this compound has broad-spectrum antibacterial activity.

The MIC values of the compounds against the five pathogenic fungi was shown in Table 8. Compound 1 showed different inhibitory activities against four pathogenic fungi, with MIC values ranging from 128 to $512 \mu g/mL$. Compounds 2 and 3 showed inhibitory activity against only one pathogenic fungus. Compound 4 exhibited inhibitory activity against five pathogenic fungi, with MIC against *F. moniliforme*, *F. graminearum*, and *B. cinerea* being 128 $\mu g/mL$.

Overall, Compound 4 exhibited inhibition activities against both pathogenic bacteria and pathogenic fungi. However, Compounds 2, 3, 5 and 6, which are derivatives of Compound 4, showed low antibacterial activity, indicating that 11-OH is a key functional group.



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TABLE 7 NMR data of compound 3 (600 MHz, $CDCl_3$) and fusariumindole A (600 MHz, $DMSO-d_6$).

NO.		3	fusariumindole A		
	δ_{C}	$\delta_{\scriptscriptstyle H}$ (mult. J)	δ_{C}	$\delta_{\scriptscriptstyle H}$ (mult. J)	
1				10.93, (brs, 1H)	
1'				10.86, (brs, 1H)	
2		6.99 (s, 1H)	124.10	7.22, (d, 2.4, 1H)	
2'	126.92	6.74 (s, 1H)	123.20	7.16, (d, 1.8, 1H)	
3	106.88		107.00		
3'	110.45		110.00		
4a	105.54		127.10		
4a'	127.76		127.10		
4	119.13	7.60 (dd, 7.8, 1.2 Hz, 1H)	118.20	7.54, (d, 8.4 Hz, 1H)	
4'	119.01	7.57 (dd, 7.8, 1.2 Hz, 1H)	118.30	7.42, (d, 8.4 Hz, 1H)	
5	118.88	7.10 (t, 7.2 Hz, 1H)	118.28	6.96 (ddd, 8.4, 7.8, 0.6 Hz, 1H)	
5'	118.83	7.10 (t, 7.2 Hz, 1H)	118.28	6.98 (ddd, 8.4, 7.8, 0.6 Hz, 1H)	
6	121.71	7.23 (t, 7.8 Hz, 1H)	121.00	7.07 (ddd, 8.4, 7.8, 1.2 Hz, 1H)	
6'	121.56	7.22 (t, 7.8 Hz, 1H)	120.95	7.07 (ddd, 8.4, 7.8, 1.2 Hz, 1H)	
7	109.22	7.30 (d, 8.4 Hz, 1H)	111.30	7.35, (d, 8.4 Hz, 1H)	
7'	109.16	7.28 (d, 8.4 Hz, 1H)	111.30	7.35, (d, 8.4 Hz, 1H)	
7a	10/0/		136.10		
7a'	136.86		136.20		
8	31.35	3.77 (s, 2H)	31.30	3.73, (s, 2H)	
9	172.18		171.60		
10	65.08	4.37 (t, 7.2 Hz, 2H)	64.40	4.27 (t, 7.2 Hz, 2H)	
11	24.64	3.08 (t, 7.2 Hz, 2H)	24.10	3.00 (t, 7.2 Hz, 2H)	
12	32.69	3.74 (s, 3H)			
12'	32.56	3.67 (s, 3H)			

TABLE 8 Antifungal activities of compounds 1–9 and amphotericin B.

Compounds	Pathogenic fungi (MIC µg/mL)				
	C. albicans	V. dahliae	F. moniliforme	F. graminearum	B. cinerea
1	-	256	512	128	256
2	-	-	-	-	256
3	-	512	-	-	-
4	256	512	128	128	128
5	-	512	-	-	256
6	-	-	-	128	512
7	-	512	-	256	-
8	-	256	-	256	512
9	-	-	512	256	512
Amphotericin B	32	16	32	16	16

"-" MIC>512 $\mu g/mL.$ Three replicates were carried out for each test.





The bioactivity of *N*-methyl tryptophol (**4**) was few reported, but the antifungal and antibacterial activities of tryptophol have been proved (Li et al., 1994; Jin et al., 2015; Singkum et al., 2019), indicating the negligible effect of N-methyl on the activity.

3.5. Cytotoxicity of compounds

As shown in Table 9, the inhibitory effects of different compounds on cancer cells were significantly different. Compound 1 showed high cytotoxicity against all cancer cells. The IC_{50} of (S,E)-6-(4-hydroxy-3-oxopent-1-en-1-yl)-2H-pyran-2-one (an analog of Compound 1)

against A549 and MDA-MB-231 was 84.5 and 64.3 μ M, respectively (Gu et al., 2022), while in this study, the IC₅₀ of Compound 1 against the two cancer cells was 28.20 and 12.26 μ M, respectively. Structurally, the hydroxyl group in the C-4' position of (S,E)-6-(4-hydroxy-3-oxopent-1-en-1-yl)-2H-pyran-2-one changed into a carbonyl group, demonstrating that the carbonyl group of C-4' in compound 1 was extremely important for the enhanced cytotoxicity.

Compound **4** showed moderate cytotoxicity against Hela $(IC_{50} = 29.74 \,\mu\text{M})$. Among the derivatives of 1-MT (Compounds **2–6**), only Compound **4** showed relatively good inhibitory activity against the cancer cells, further demonstrating the key role of the hydroxyl group at C11 in Compound **4**.



TABLE 9 Cytotoxicity of compounds 1–9 and doxorubicin hydrochloride (DOX).

Compounds	IC ₅₀ (μM)					
	Huh-7	Hela	HCT-15	A549	MDA-MB-231	
1	29.70 ± 3.35	52.52 ± 3.62	13.60 ± 1.43	28.20 ± 1.77	12.26 ± 1.43	
2	-	-	-	-	170.68 ± 12.81	
3	-	130.66 ± 12.48	184.05 ± 9.50	184.69 ± 17.47	80.46 ± 1.56	
4	133.74 ± 13.67	29.74 ± 1.30	101.88 ± 8.50	95.80 ± 2.50	71.63 ± 1.88	
5	180.06 ± 1.34	137.76 ± 12.55	-	133.48 ± 14.32	144.82 ± 10.23	
6	-	139.43 ± 7.09	-	-	163.12 ± 3.69	
7	-	118.23 ± 9.81	-	106.66 ± 3.38	129.52 ± 6.12	
8	-	120.42 ± 16.56	-		-	
9	168.00 ± 1.44	-	-	135.88 ± 1.79	159.13 ± 9.60	
DOX	0.22 ± 0.02	0.18 ± 0.05	0.17 ± 0.01	0.23 ± 0.01	0.27 ± 0.02	

"-" IC₅₀ > 200 μ M.

4. Conclusion

By feeding 1-MT into cultures of the endophytic fungus Diaporthe sp. AC1 to increase the diversity of metabolites, three novel compounds (1-3), together with six known ones were isolated and characterized. Most of the compounds were the biotransformation products of 1-MT. In the bioactivity assay, Compound 1 showed moderate antibacterial activity against V. dahlia, F. graminearum, and B. cinerea, with MIC values being 256, 128, and 256 $\mu g/mL,$ respectively. It also showed high cytotoxicity against five cancer cells, with IC₅₀ ranging from 12.26 to 52.52 µM. Other compounds also exhibited certain antimicrobial and cytotoxic activities. Under the severe situation that antibiotic resistance is a growing public threat, this work provides an efficient and eco-friendly methodology for the development of structurally undescribe compounds, from which lead molecules can be recognized and optimized for new drugs discovery.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary material, further inquiries can be directed to the corresponding authors.

Author contributions

SZ: Writing – original draft, Investigation. QX: Writing – original draft, Investigation. CJ: Writing – original draft, Investigation. XH: Investigation, Writing – original draft. YZ: Investigation, Writing – original draft. CL: Investigation, Writing – original draft. LM: Investigation, Writing – original draft. WS: Investigation, Writing – original draft. YL: Writing – original draft, Formal analysis. ZY: Formal analysis, Writing – review & editing. FZ: Writing – review & editing, Project administration. YT: Project administration, Writing – original draft.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fmicb.2023.1254609/ full#supplementary-material

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