Five new secondary metabolites from an endophytic fungus *Phomopsis* sp. SZSJ-7B

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Two previously undescribed lactones, phomolides A and B (1 and 2), and three new sesquiterpenoids, phomenes A–C (3–5), together with one known compound, colletotricholide A (6), were isolated from the endophytic fungus *Phomopsis* sp. SZSJ-7B. Their chemical structures, including the absolute configurations, were comprehensively established by extensive analyses of NMR, high-resolution electrospray ionization mass spectrometry, electronic circular dichroism powered by theoretical calculations, and X-ray diffractions. Moreover, the cytotoxic and antibacterial activities of compounds 1–6 were also evaluated, and the results demonstrated that compound 2 showed significant antibacterial effects towards methicillin-resistant *Staphylococcus aureus* and *S. aureus* strains with minimum inhibitory concentration as low as 6.25 μg/ml, which was comparable to that of the clinical drug vancomycin. Moreover, all compounds showed no cytotoxic activity.

KEYWORDS
endophytic fungus, *Phomopsis*, secondary metabolites, antibacterial activity, cytotoxic activity

1 Introduction

Endophytes play an important role of producing biologically meaningful natural products and can be considered as a strategically promising bio-resource of significantly economic potential for the agrochemical and pharmaceutical industries (Gouda et al., 2016). It is well documented that the genus *Phomopsis* can generate structurally diverse and pharmaceutically active secondary metabolites (Huang et al., 2008; Yu et al., 2008; Hemtasin et al., 2011), including xanthones (Ding et al., 2013), α-pyrones (Cai et al., 2017), steroids (Hu et al., 2017), sesquiterpenes (Xie et al., 2018), diterpenes
(Wei et al., 2014), triterpenes (Li et al., 2008), oblongolides (Bunyapaiboonsri et al., 2010), pyrenocines (Hussain et al., 2012), alkaloids (Chen et al., 2019), cytochalasins (Yan et al., 2016), etc. Moreover, these intriguing natural compounds shared various biological activities such as antimutator (Pavao et al., 2016), immunosuppressive (Wei et al., 2014), antifungal (Krohn et al., 2011), antioxidant (Chen et al., 2018), antibacterial (Jouda et al., 2016), anti-inflammatory (Xu et al., 2019b), and α-glucosidase inhibitory effects (Huang et al., 2018).

Our group pursued continuous research commitments towards discovering structurally fascinating and biologically significant natural products from endophytic fungi in recent years, and a series of metabolites with excellent antibacterial and antimutator activities from endophytic fungi of the genus Phomopsis have been reported (Xu et al., 2019a; Chen et al., 2020; Liu et al., 2021). In continuation of our ongoing endeavors, a strain of Phomopsis sp. SZSJ-7B isolated from the fresh leaves of Alpinia shengzhen, which is a beautiful horticultural plant (Zingiberaceae), was chosen as the appealing target for the chemical constituent investigation. Preliminary thin-layer chromatography and high-performance liquid chromatography (HPLC) screenings of the strain SZSJ-7B were conducted, and the experimental data showed that its ethyl acetate (EtOAc) extracts exhibited a remarkable diversity of secondary metabolites. A further systematical chemical study of the strain led to the isolation of five previously undescribed metabolites including two lactones, phomolides A and B, and three sesquiterpenoids, phomenes A–C. Herein the details of the extraction, purification, structure elucidation, and their biological evaluation are described.

2 Results and discussion

Compound 1 was isolated as a white solid. Its molecular formula C_{11}H_{13}O_{4} was deduced from high-resolution electrospray ionization mass spectrometry (HRESIMS) m/z 209.0815 [M + H]^+ [calculated (calcd) for C_{11}H_{13}O_{4}, 209.0808], indicating six degrees of hydrogen deficiency. The infrared (IR) spectrum of 1 showed obvious absorption bands at 3,325, 1,699, and 1,022 cm⁻¹ and revealed the presence of corresponding hydroxyl and carbonyl functionalities together with the ether bonds. The ¹H nuclear NMR data (Table 1) of 1 showed two singlet methyl groups (δ_H 2.12 and 2.57) and an upfield doublet methyl group (δ_H 1.65, d, ¹J = 5.2 Hz). Its ¹³C NMR (Table 1) and heteronuclear single quantum coherence (HSQC) spectra showed 11 carbon signals including six quaternary carbons (δ_C 163.2, 161.9, 159.1, 143.0, 120.6, and 104.5), two oxymethine (δ_C 98.7 and 97.7), and three methyl groups (δ_C 18.7, 16.1, and 10.0).

Moreover, the significant heteronuclear multiple bond correlation (HMBC) correlations (Figure 1) from H3-11 to C-1 (δ_C 104.5), C-7 (δ_C 120.6), and C-8 (δ_C 143.0), H3-10 to C-6 (δ_C 161.9), C-7, and C-8, and H-5 to C-3 and C-7 strongly suggested the existence of a penta-substituted benzene ring. In addition, on the basis of the ¹H–¹H correlation spectroscopy (COSY) fragment of C-1/C-11, the obvious HMBC correlations from H3-11 to C-1 (δ_C 97.7) coupling with H-1 to C-2 (δ_C 120.6), C-7, and C-8, and H-5 to C-3 and C-7 strongly suggested the existence of a benzene ring. Moreover, these intriguing natural compounds shared various biological activities such as antitumor (Pavao et al., 2016), immunosuppressive (Wei et al., 2014), antifungal (Krohn et al., 2011), antioxidant (Chen et al., 2018), antibacterial (Jouda et al., 2016), anti-inflammatory (Xu et al., 2019b), and α-glucosidase inhibitory effects (Huang et al., 2018).

### Table 1

| No. | δ₁H (J in Hz) | δ₁C |  |
|-----|---------------|-----|---|
| 1   | 5.64, q, (5.2) | 97.7, CH |  |
| 2   | 163.2, C      | 31.8, CH₂ | 13 |
| 3   | 104.5, C      | 31.3, CH₂ | 15 |
| 4   | 6.32, s       | 159.1, C |  |
| 5   | 161.9, C      | 38.5, CH | 1' |
| 6   | 6α            | 34.8, CH₂ | 3' |
| 7   | 2.44, m       | 1.83, m |  |
| 8   | 143.0, C      | 2.12, m |  |
| 9   | 2.57, s       | 16.1, CH₃ |  |
| 10  | 2.12, s       | 10.0, CH₃ |  |
| 11  | 1.65, δ, (5.2) | 18.7, CH₃ |  |
| 12  | 5.46, s       | 100.8, CH |  |

In continuation of our ongoing endeavors, a strain of Phomopsis sp. SZSJ-7B isolated from the fresh leaves of Alpinia shengzhen, which is a beautiful horticultural plant (Zingiberaceae), was chosen as the appealing target for the chemical constituent investigation. Preliminary thin-layer chromatography and high-performance liquid chromatography (HPLC) screenings of the strain SZSJ-7B were conducted, and the experimental data showed that its ethyl acetate (EtOAc) extracts exhibited a remarkable diversity of secondary metabolites. A further systematical chemical study of the strain led to the isolation of five previously undescribed metabolites including two lactones, phomolides A and B, and three sesquiterpenoids, phomenes A–C. Herein the details of the extraction, purification, structure elucidation, and their biological evaluation are described.
result, the experimental ECD spectrum perfectly matched with the calculated ECD spectrum of 1S configuration for 1, showing the same clear Cotton effect at 205 nm. Thus, the absolute configuration of 1 was determined to be 1S (Figure 3), and it was revealed as a new, natural, rarely occurring acetal derivative, which was given the trivial name phomolide A.

Compound 2 was isolated as a colorless oil. Its molecular formula C23H28O5 was deduced from its HRESIMS spectrum with a molecular ion peak observed at m/z 385.2007 [M + H]+ (calcd for C23H29O5, 385.2010), which chemologically implied 10 degrees of hydrogen deficiency. Moreover, the IR spectrum of compound 2 showed a series of characteristic absorption bands at 3,357, 1,714, and 1,020 cm⁻¹, which were attributed to hydroxyl and carbonyl functional moieties as well as ether bonds. The ¹H NMR data (Table 1) of 2 showed typical proton resonances for three methyl groups at δH 0.95 (s, H3-14), 0.84 (d, J = 6.5 Hz, H3-15), and 2.57 (s, H-8′), an oxygenated methine moiety at δH 5.46 (s, H-12), and two aromatic protons at δH 6.35 (d, J = 2.0 Hz, H-4′) and δH 6.50 (d, J = 2.0 Hz, H-6′) together with an olefinic proton at δH 5.34 (m, H-9). The ¹³C NMR (Table 1) and HSQC spectra exhibited 24 carbon signals comprising three methyls, six methylenes, six methines, and nine quaternary carbons. The ¹H–¹H COSY spectrum of 2 revealed the existence of two independent spin systems of H2-1/H2-2/H2-3/H-4/H3-15 and H2-6/H-7/H2-8/H-9.

FIGURE 1
¹H–¹H correlation spectroscopy and key heteronuclear multiple bond correlation correlations of compounds 1–5 and key nuclear Overhauser effect spectroscopy correlations of compounds 2–4.

FIGURE 2
Structures of compounds 1–6.
Compound 2 was further conclusively revealed as a novel meroterpenoid with eremophilan and acetophenone units conjugating as an acetal skeleton after the careful comparison of 1D NMR data of 2 with those of the known compound colletotricholide A (Zhao et al., 2020), which was also co-isolated from this strain. The main difference in NMR spectra between 2 and colletotricholide A (6) was attributed to the lack of a methyl group in 2 at C-6 position. This speculation could be further verified by the 1H and 13C NMR signals for H-6' (δH 6.50) and C-6' (δC 114.4) in 2 and the key HMBC correlations from H-6' to C-1' (δC 164.1) and C-4' (δC 99.8). Therefore, the planar structure of 2 was identified as shown in Figure 2.

The relative configuration of 2 was determined by the nuclear Overhauser effect spectroscopy (NOESY) experiment (Figure 1). As shown in Figure 1, the NOESY correlations of H-6β with H-8β, H3-14, and H3-15 revealed that these protons were on the same face and assumed as β-oriented, while the correlations of H-6α with H-8α together with H-12 with H-7 and H-8α indicated that H-7 and H-12 were α-oriented. The CD spectrum of 2 showed positive Cotton effects at 211 and 268 nm, which were very similar with those in the CD spectrum of the known compound cyclodebneyol (Burden et al., 1986) tentatively revealed that compound 3 shared the same planar structure as that of the previously reported natural product cyclodebneyol. Moreover, the key HMBC correlations from H3-14 to C-3 (δC 30.9) and C-5 (δC 39.3); H15 to C-6 (δC 32.2), C-4 (δC 38.0), and C-10 (δC 153.7); and H3-13 to C-7 (δC 45.1) and C-12 (δC 78.6), combined with the COSY fragments a and b, further confirmed the aforementioned conclusion (Figure 1).

The obvious differences in the chemical shifts of both 1H and 13C NMR data between 3 and cyclodebneyol suggested that these two compounds ought to be a pair of closely related stereoisomers. Moreover, the cross-peaks of H3-14/H-6β, H3-15/H-6β, and H3-15/H-6β in the NOESY spectrum were clearly distinguished; thus, it could be readily speculated that the three methyls H3-13, H3-14, and H3-15 in 3 directed on the same side in its 6/6/5 fused ring skeleton and assumed as β-oriented. However, the proton chemical shift of H-7 was heavily overlapped with H-2β so the NOESY correlations of H-7/H-4 could not conclusively determine the orientation of protons H-7 and H-4 to further completely confirm the final relative configuration of 3.

In order to absolutely determine the relative configuration of C-7 for 3, the gauge-independent atomic orbital (GIAO) density functional theory (DFT) 13C NMR calculations (McWeeny, 1961; Ditchfield, 1972) towards the structures 3a and 3b were
performed at the \( \omega B97x-D/6-31G^* \) (Chai and Head-Gordon, 2008) (IEFPCM, CDCl\(_3\)) level, and the calculation data were then compared with their experimental \(^{13}\)C NMR data following the reported sorted training set (STS) protocol (Li et al., 2020). According to the linear regression analysis of \(^{13}\)C NMR chemical shifts, the values of the correlation coefficient \((R^2)\) were 0.9989 for 3a and 0.9800 for 3b (Figure 4). Moreover, the resulting \(P_{\text{rel}}\) value of 3a is 100%, and the mean absolute error (MAE), root mean square error (RMSE), and \(P_{\text{mean}}\) values of 3a showed that the calculated \(^{13}\)C NMR data match the experimental data very

**TABLE 2** Calculated \(^{13}\)C chemical shifts (CDCl\(_3\)) fitting with the experimental data of compounds 3a, 3b, 4a, and 4b following the STS protocol.

| Exptl. | 3  | 4  | Exptl. | 4  |
|--------|----|----|--------|----|
|        | 3a | Dev | 3b | 3a | Dev | 4a | Dev | 4b | Dev |
|        |    |     |    |    |     |    |     |    |     |
| 32.1   | 32.07 | 0.03 | 32.87 | 0.67 | 32.4 | 32.61 | 0.21 | 32.43 | 0.03 |
| 31.43  | 28.74 | 1.06 | 27.08 | 4.22 | 29.9 | 26.00 | 3.90 | 28.66 | 1.24 |
| 25.58  | 29.32 | 1.58 | 29.77 | 2.61 | 30.9 | 29.53 | 1.37 | 29.54 | 1.36 |
| 28.29  | 37.68 | 0.32 | 43.67 | 4.34 | 38.5 | 42.78 | 4.28 | 39.90 | 1.40 |
| 42.34  | 40.00 | 0.70 | 42.28 | 1.64 | 39.1 | 38.58 | 0.52 | 39.19 | 0.09 |
| 40.94  | 32.42 | 0.22 | 35.00 | 1.37 | 30.4 | 35.41 | 5.01 | 31.99 | 1.59 |
| 33.57  | 47.18 | 2.08 | 49.33 | 2.96 | 43.4 | 44.00 | 0.60 | 44.03 | 0.63 |
| 48.06  | 74.93 | 0.17 | 78.33 | 2.26 | 75.5 | 75.38 | 0.12 | 75.44 | 0.06 |
| 77.36  | 118.92 | 3.32 | 123.75 | 7.66 | 115.6 | 121.20 | 5.60 | 119.66 | 4.06 |
| 123.26 | 152.23 | 1.47 | 146.34 | 7.61 | 154.6 | 150.01 | 4.59 | 152.02 | 2.58 |
| 146.09 | 81.55 | 0.25 | 78.38 | 4.39 | 79.4 | 79.36 | 0.04 | 79.00 | 0.40 |
| 77.41  | 75.86 | 2.74 | 81.08 | 1.54 | 78.6 | 76.83 | 1.77 | 77.44 | 1.16 |
| 80.14  | 21.44 | 0.64 | 24.16 | 1.82 | 27.7 | 26.34 | 1.36 | 26.15 | 1.55 |
| 22.62  | 15.88 | 0.18 | 15.93 | 1.39 | 15.8 | 16.64 | 0.54 | 16.11 | 0.31 |
| 14.31  | 20.77 | 0.47 | 19.19 | 2.69 | 20.3 | 17.74 | 2.56 | 20.56 | 0.26 |
| 17.61  | MAE\(^a\) | 1.02 | MAE\(^a\) | 3.14 | MAE\(^a\) | 2.16 | MAE\(^a\) | 1.11 |
|        | RMS\(^b\) | 1.42 | RMS\(^b\) | 3.77 | RMS\(^b\) | 2.89 | RMS\(^b\) | 1.54 |
|        | \(P_{\text{mean}}\) | 32.34\% | \(P_{\text{mean}}\) | 0.58\% | \(P_{\text{mean}}\) | 3.12\% | \(P_{\text{mean}}\) | 31.18\% |
|        | \(P_{\text{rel}}\) | 100.00\% | \(P_{\text{rel}}\) | 0.00\% | \(P_{\text{rel}}\) | 0.00\% | \(P_{\text{rel}}\) | 100.00\% |

\(^{a}\)Absolute error.

\(^{b}\)Root mean square.
well, indicating that 3a or its enantiomer is the correct structure for 3 (Table 3). With the aforementioned informative results, the relative structure of 3 was thus unambiguously established as shown in Figure 2.

Moreover, the absolute configuration of 3 was also determined by the time-dependent density-functional theory (TDDFT) calculated circular dichroism (CD) spectrum at the mPW1PW91/SVP level. As shown in Figure 5, the calculated ECD curve of 4S,5R,7S,8R,11R-3 perfectly matched with the experimental ECD curve, which strongly suggested that compound 3 shared an absolute configuration of 4S,5R,7S,8R,11R. Therefore, the structure of compound 3 was completely established and given the trivial name phomene A.

Compound 4 was obtained as a yellow oil with the same molecular formula as 3, which was determined by HRESIMS ion peak at m/z 237.1854. Obviously, the 13C NMR spectroscopic data (Table 2) and HSQC spectrum of 4 collectively suggested 15 carbon signals, and all of them showed very similar chemical shifts to those of 3. The little differences between the chemical shifts of 3 and 4 in NMR spectra strongly implied that they should be a pair of diastereoisomers sharing the same planar structure. The further careful analysis of the NOESY spectrum of 4 could efficiently establish its relative configuration. Compared with the NOESY spectrum of 3, a clear NOESY correlation of H-2a/H-4 and H3-14/H3-15 was observed, illustrating that the critical protons H-8 and H3-13 were on the same side in the fused ring system and assumed as α-oriented. In addition, the cross-peaks of H-2β/H-4 and H-1β/H3-14 in the NOESY spectrum clearly demonstrated that the two methyls H3-14 and H3-15 were β-oriented. Therefore, the relative structure of 4 was tentatively assigned as a C-11 epimer of 3, as shown in Figure 2, although the same stereochemical issue clouded the H-7 chirality as that of 3.

In order to further confirm the relative configuration of 4, we also carried out a 13C NMR calculation for 3. As a result, the Pmean and Prel parameters as well as MAE and RMS values further showed that 4b or its enantiomer should be the correct structure for 4, as shown in Table 3. Moreover, the absolute configuration of 4 was determined to be 4S,5R,7S,8R,11S based on the experimental ECD spectrum, which was highly similar to the calculated ECD spectrum (Figure 5). Thus, the absolute configuration of compound 4 was fully confirmed and given the trivial name phomene B.

Compound 5 was isolated as yellow crystals. Its molecular formula of C15H26O2 was deduced by the HRESIMS spectrum with a protonated ion peak discovered at m/z 239.2004 [M + H]+ (calcd for C15H27O2, 239.0606), indicating three degrees of hydrogen deficiency. The IR spectrum of 5 revealed an obvious absorption band at 3,315 cm⁻¹, indicating the presence of a series of free hydroxyl functionalities. The 1H NMR data

| No.  | 3a | 3b | 4a | 4b | 5a | 5b |
|------|----|----|----|----|----|----|
| δH (J in Hz) | δC | δH (J in Hz) | δC | δH (J in Hz) | δC |
| 1α   | 2.00, d, (12.1) | 32.1, CH2 | 2.01, d, (11.9) | 32.4, CH2 | 1.99, m | 53.1, CH |
| 1β   | 2.29, td, (4.3, 12.1) | 2.31, td, (4.4, 11.9) | 2.29, CH2 | 2.31, CH2 | 1.99, m | 53.1, CH |
| 2α   | 1.25, m | 29.8, CH2 | 1.26, m | 29.9, CH2 | 1.31, m | 46.3, CH2 |
| 2β   | 1.85, m | 1.88, m | 1.88, m | 1.88, m | 1.49, m | 1.49, m |
| 3α   | 1.48, m | 30.9, CH2 | 1.49, m | 30.9, CH2 | 1.31, m | 46.3, CH2 |
| 3β   | 1.54, m | 38.0, CH2 | 1.53, m | 38.5, CH2 | 1.56, m | 21.5, CH2 |
| 4    | 39.3, C | 39.1, C | 39.1, C | 39.1, C | 2.14, ddd, (2.4, 9.4, 15.9) | 49.6, CH2 |
| 5    | 0.94, m | 32.2, CH2 | 1.91, m | 30.4, CH2 | 1.91, m | 36.7, C |
| 6α   | 1.76, m | 1.76, m | 1.76, m | 1.76, m | 1.76, m | 29.2, CH2 |
| 6β   | 1.89, m | 45.1, CH2 | 1.88, m | 43.4, CH2 | 1.43, m | 41.9, CH |
| 7α   | 1.74, m | 1.74, m | 1.74, m | 1.74, m | 1.74, m | 42.4, C |
| 7β   | 4.50, t, (5.3) | 75.1, CH | 4.18, t, (5.2) | 75.5, CH | 2.61, ddd, (9.4, 15.9) | 37.6, CH2 |
| 9    | 5.55, d, (5.3) | 115.6, CH | 5.50, d, (5.2) | 115.6, CH | 1.54, m | 18.6, CH2 |
| 10α  | 153.7, C | 154.6, C | 1.49, m | 1.49, m | 1.49, m | 23.4, CH2 |
| 10β  | 81.8, C | 79.4, C | 1.79, m | 1.79, m | 1.79, m | 23.4, CH2 |
| 11α  | 3.73, d, (9.6) | 82.6, CH2 | 3.66, d, (8.9) | 78.6, CH2 | 1.24, s | 68.2, CH2 |
| 11β  | 3.87, d, (9.6) | 78.6, CH2 | 3.66, d, (8.9) | 78.6, CH2 | 1.24, s | 68.2, CH2 |
| 13   | 1.33, s | 20.8, CH2 | 1.46, s | 27.7, CH2 | 1.22, s | 25.4, CH2 |
| 14   | 0.82, d, (6.6) | 15.7, CH2 | 0.84, d, (6.4) | 15.8, CH2 | 1.22, s | 25.4, CH2 |
| 15a  | 0.95, s | 20.3, CH2 | 1.01, s | 20.3, CH2 | 3.37, d, (10.8) | 68.2, CH2 |
| 15b  | 0.95, s | 20.3, CH2 | 1.01, s | 20.3, CH2 | 3.37, d, (10.8) | 68.2, CH2 |
(Table 3) of 5 showed three singlet methyl groups ($\delta_1$ 1.24, 1.24, and 1.26), a hydroxymethyl moiety ($\delta_3$ 3.37, 3.54), and various kinds of saturated aliphatic protons ranging from $\delta_1$ 1.22 to 2.61. According to the $^{13}$C NMR (Table 3) and HSQC data of compound 5, 15 carbon signals were resolved, namely, three methyl moieties ($\delta_1$ 22.2, 23.4, and 25.4), six methylene groups ($\delta_1$ 18.6, 21.3, 29.2, 37.6, 46.3, and 68.2), and three methine functionalities ($\delta_1$ 41.8, 49.3, and 53.1) together with three quaternary carbons ($\delta_1$ 36.6, 42.3, and 74.0). With careful consideration of the molecular formula of 5, this informative data strongly indicated that compound 5 might be a tricyclic sesquiterpenoid derivative.

In the $^1$H–$^1$H COSY spectrum, the obvious correlations of H-1/H$_2$-11/H$_2$-10 and H$_2$-3/H$_2$-4/H-5/H-8/H$_2$-7 suggested the presence of two independent spin fragments a (C-1/C-11/C-10) and b (C-3/C-4/C-5/C-8/C-7). With reference to fragment a, the HMBC correlations from H$_3$-13 to C-1 ($\delta_1$ 53.1), C-9 ($\delta_1$ 42.4), and C-10 ($\delta_1$ 37.6) evidently confirmed the presence of a cyclobutane ring (ring A). Meanwhile, the HMBC correlations from H$_2$-14 to C-5 ($\delta_1$ 49.6), C-6 ($\delta_1$ 36.7), C-7 ($\delta_1$ 29.2), and C-15 ($\delta_1$ 68.2), H$_2$-15 to C-5, C-6, and C-7 coupling with the COSY fragment C-5/C-6/C-7 further established the other cyclobutane ring (ring C) with a hydroxymethyl functionality attached at the C-6 position. The seven-membered ring B could be conveniently constructed by the HMBC correlations from H-1 to C-3 ($\delta_1$ 21.5) and C-8 ($\delta_1$ 41.9), H-3 to C-5, H-5 to C-9, H$_2$-12 to C-1, C-2 ($\delta_1$ 74.0), and C-3 on the basis of the COSY fragment C-3/C-4/C-5/C-8. Lastly, the key HMBC correlations of H$_2$-11 to C-2, H-7 to C-9, and H$_2$-4 to C-6 could further establish the connection of the 4/7/4 fused ring system (rings A–C). Therefore, the planar structure of 5 was then determined. The relative and absolute configurations of compound 5 were unambiguously confirmed by the X-ray single crystal diffraction on the CuKα with a Flack parameter of -0.05 (12). Finally, the absolute configuration was thus designated as 1R,2R,5R,6S,8R,9R, as shown in Figure 6, and given the trivial name phomene C.

At this stage, compounds 1–6 were evaluated for antimicrobial activities against the bacteria Escherichia coli, S. aureus, and methicillin-resistant S. aureus (MRSA) (Table 4). The biological screening results showed that the new compound 2 showed very potent antimicrobial activities to Gram-positive stain S. aureus and MRSA with a minimum inhibitory concentration (MIC) value at 6.25 µg/ml for both. In addition, the known compound 6 also exhibited significant antimicrobial activities, with a MIC value of 1.56 µg/ml towards S. aureus and 0.78 µg/ml towards MRSA, respectively, which were very close to those of the positive control vancomycin (0.78 µg/ml for S. aureus and 0.78 µg/ml for MRSA). However, we have only tested the bacteriostatic potential of the compounds, and their bactericidal activity is still unknown and deserves further extensive exploration.

Furthermore, compounds 1–6 were also tested for their cytotoxicity against a panel of human cancer cell lines including SF-268, MCF-7, HepG-2, and A549 and normal cell line LX-2 (Table 5). However, all the tested compounds were found to be devoid of antitumor activity even at the concentration of 100 µM. The aforementioned biological screening results collectively pointed that the acetal meroterpenoids 2 and 6 might be severed as promising lead compounds towards antibacterial innovative drug development.

Conclusively, numerous excellent efforts towards the secondary metabolites of the plant endophytic fungi have successfully clarified that the endophytic fungi shared the outstanding ability to produce pharmaceutically meaningful natural products (Debbab et al., 2013; Mousa and Raizada, 2013; Brader et al., 2014) or similar bioactive metabolites as their hosts (Stierle et al., 1993; Kusari et al., 2011; Kusari et al., 2012). In recent years, the extraction and the isolation of bioactive leading natural products, with aim of discovering innovative drugs from the plant endophytic fungi, are emerging as a hot research topic for both natural product and medicinal chemists (Praptiwi et al., 2018; Tanapichatsakul et al., 2018; Adeleke and Babalola, 2020). However, the endophytic...
fungi of the *Alpinia shengzhen* plant have not been reported. This study reported the isolation and identification of the endophytic fungi of *Alpinia shengzhen* for the first time (Hu et al., 2011). As a result, six strains of endophytic fungi, namely, *Cladorrhinum* sp. SZSJ-2 (Madrid et al., 2011), *Phyllosticta capitalensis* SZSJ-3 (Arafat, 2018), *Nigrospora oryzae* SZSJ-5 (Hudson, 1963), and two strains of *Phomopsis* sp. SZSJ-7B and 7C (Zhang et al., 2000) together with *Annulohypoxylon stygium* SZSJ-7A (Hsieh et al., 2005), were isolated from the plant tissues of *Alpinia shengzhen* by section culture because of their abundant secondary metabolites and notable biological activities. These results informatively suggested that *Alpinia shengzhen* could be applied as a promising bioresource for the discovery of medicinal fungi.

In this study, a chemical investigation of the endophytic fungus *Phomopsis* sp. SZSJ-7B of *Alpinia shengzhen* was performed for the first time to discover novel lead drug molecules with chemically diverse structures and biologically significant activities. Although previous studies revealed a huge number of intriguing novel natural products isolated from the genus *Phomopsis* (Silva et al., 2006; Yang et al., 2013; Fan et al., 2020; Gong et al., 2020; Yang et al., 2020), five novel secondary metabolites and one known natural product were also successfully isolated from this genus in this time. The six natural products shared four different structure types, all of which were isolated from this genus for the first time, thus greatly enriching the structural types of natural compounds from *Phomopsis* sp. (Udayanga et al., 2011; Xu et al., 2021). Moreover, this chemical research effort can also strongly provide chemo-logical reference and experimental guidance for future studies towards the endophytic fungi and the secondary metabolites of other ginger plants.

The acetal skeleton represents a ubiquitous and intriguing family of structurally special architecture, which is prevalent in numerous biologically meaningful natural products (Pettit et al., 2015) and many pharmaceutically significant clinical drugs, such as aquamox for hypertension (Verdel et al., 2006) and cevimeline for parasympathetic nerves (Mavragani and Moutsopoulos, 2007). In this study, phomolides A and B were characterized with a natural, rarely occurring ester-acetal skeleton, which is constructed by a carboxylic acid and a phenol hydroxyl functionality (Chen et al., 2021) together with an aldehyde fragment (Ahmed Laskar and Younus, 2019), giving rise to a distinctive difference with the common acetal moiety formatted with two alcoholic hydroxyl groups and an aldehyde. To our knowledge, there were only four examples of natural products sharing this skeleton reported in previous literatures (Zhang et al., 2020; Choi et al., 2021). The discovery of phomolides A and B further enriches the

### TABLE 4 Antibacterial activity of compounds 1–6 (minimum inhibitory concentration, μg/ml).

| Compounds       | *Escherichia coli* | *Staphylococcus aureus* | Methicillin-resistant *S. aureus* |
|-----------------|--------------------|-------------------------|-----------------------------------|
| Vancomycin/kanamycin | 3.12               | 0.78                    | 0.78                              |
| 1               | >100               | 100                     | 100                               |
| 2               | >100               | 6.25                    | 6.25                              |
| 3               | >100               | 100                     | 100                               |
| 4               | >100               | 100                     | 100                               |
| 5               | >100               | 100                     | 100                               |
| 6               | >100               | 1.56                    | 0.78                              |
structural types and the members of these ester-acetal compounds. Moreover, the biological evaluations showed that phomolide B exhibited a significant inhibitory activity against S. aureus and MRSA, which suggested that these ester-acetal compounds might continuously serve as potent innovative impetus for the further extensive research and medicinal exploration towards the development of novel antibacterial drugs (Rossiter et al., 2017; Wu et al., 2019; Dandawate et al., 2019; Dai et al., 2020).

Phomenes A and B possess a typical eremophilane-type sesquiterpene skeleton with a fascinating 6/6/5 fused ring system. The first example of this eremophilane sesquiterpene, named cyclodebney, was reported in 1986, which was successfully isolated from tobacco necrosis virus (TNV) Nigitiutua debneyi and then illustrated to show a potent antifungal activity. Furthermore, Le isolated three other new eremophilane-type sesquiterpenes from Sarcographa tricosa (Le et al., 2013) as characteristic secondary metabolites. Until now, there were about seven eremophilane sesquiterpenes with this 6/6/5 fused ring system isolated from medicinal plants (Wang et al., 2016; Shao et al., 2016) and fungi (Chang et al., 2017). However, phomenes A and B are the first two examples of eremophilane-type sesquiterpenes isolated from Phomopsis sp., and they further increase the structural diversity of secondary metabolites in this genus.

Phomene C is a tricyclic sesquiterpenoid derivative with a very intriguing 4/7/4 fused ring scaffold, which is a fascinating type of sesquiterpenoid skeleton rather rarely occurring in nature. Up to now, only two examples of this sesquiterpenoid, named koraiol and frabenol, have been previously isolated from the oleoresin of Pinus koraiensis and Fimetariella rabenhorstii, respectively (Khan et al., 1979; Tao et al., 2011). Moreover, the sesquiterpene alcohol 5,8-cyclocaryophyllan-4-ol, which was detected in Cangerana oil (Weyerstahl et al., 1996), also shared a closely similar carbon skeleton as that of koraiol and frabenol with a 4/7/4 fused ring system. In this study, the discovery of the new sesquiterpenoid phomene C further broadened the structural diversity and enriched the family members of this type of sesquiterpenoid.

### 3 Conclusion

In conclusion, this study firstly conducted a systematic chemical investigation on the secondary metabolites of the endophytic fungi Phomopsis sp. SZSJ-7B from Alpinia shengzhen and successfully resulted in the isolation of two novel acetal lactones (phomolides A and B), three undescribed sesquiterpenes (phomenes A–C), and a known lactone (colletotricholide A). All of these types of compounds were also isolated from Phomopsis sp. for the first time, which greatly enriched the structural diversity of the secondary metabolites of the genus. The structures of the new compounds were fully characterized by a combination of spectroscopic methods, X-ray diffraction, and quantum chemistry calculations. The biological activity screening clarified that both compounds 2 and 6 exhibited significant antibacterial activities towards MRSA and S. aureus strains with MIC values as low as 6.25 μg/ml, which were comparable to those of the positive control vancomycin without any notable cytotoxicity, thus illustrating its significant potential in the development of innovative antibacterial drugs. Moreover, further investigations on structure–activity relationship and antibacterial mechanism directed toward this goal are currently underway and will be reported in due course.

### 4 Experimental

#### 4.1 General experimental procedures

IR data were measured on a Shimadzu IR Affinity-1 spectrometer (Shimadzu, Kyoto, Japan). UV and optical rotation data were obtained by a Shimadzu UV-2600 spectrophotometer (Shimadzu, Kyoto, Japan) and an Anton Paar MCP-500 spectropolarimeter (Anton Paar, Graz, Austria). The ECD spectra were measured with Applied Photophysis Chirascan. The NMR spectra (1D and 2D) data were collected on a Bruker Avance-500 spectrometer with tetramethylsilane as an internal standard (Bruker, Fällanden,
The HRESIMS spectra were acquired with a Thermo MAT95XP high-resolution mass spectrometer (Thermo Fisher Scientific, Bremen, Germany). The single crystal data were collected on an Agilent Xcalibur Novasingle-crystal diffractometer equipped with CuKα radiation. A Hitachi Primaide [Hitachi Instruments (Dalian) Co., Ltd.] equipped with a diode array detector using a preparative YMC ODS C18 column (20 × 250 mm, 5 μm) was used for preparative HPLC separation. Sephadex LH-20 (GE Healthcare, Uppsala, Sweden), silica gel (200–300 and 60–100 mesh, Puke, Qingdao, China), and C18 reversed-phase silica gel (40-75 μm, Fuji, Kasugai, Japan) were used for column chromatography. All solvents were of analytical grade (Guangzhou Chemical Regents Company, Ltd., Guangzhou, China).

4.2 Fungal material

The fungal strain *Phomopsis* sp. SZSJ-7B was isolated from the fresh leaves of *Alpinia shenzheng* collected in the South China Botanical Garden in Guangzhou City, Guangdong Province of China in September 2020. Using BLAST to search the GenBank database, SZSJ-7B (GenBank accession number: OP623444.1) has 100% similarity with *Phomopsis* sp. MJ53 (GenBank accession number: KM203620.1). The strain is preserved at the Key Laboratory of South China Agricultural Plant Molecular Analysis and Genetic Improvement, South China Botanical Garden in Guangzhou City.

4.3 Fermentation, extraction, and isolation

The prepared fresh mycelium of the strain was inoculated into each of five 500-ml Erlenmeyer flasks containing 200 ml PDB medium (200 g potato, 20 g dextrose, 3 g KH2PO4, 1.5 g MgSO4, and 10 mg vitamin B in 1 L H2O) and then incubated at 28°C on a rotary shaker at 180 rpm for 5 days to obtain the seed culture. Fermentation was performed in 3-L Fernbach flasks, each containing 1.5 L PDB medium. After having been disinfected at 121°C for 30 min in an autoclave and cooled to room temperature, each flask was inoculated with 30 ml of the seed cultures and incubated at 28°C for 30 days. After cultivation, the mycelia were extracted with EtOAc for three times, and the crude extract (10 g) was obtained. The crude extract was subjected to silica gel using gradient elution with petroleum ether–EtOAc–methanol (MeOH) (v/v/v, 50:10–0:1) to afford six main fractions (Fr.1–Fr.6).

Fr.1 (880 mg) was isolated on silica gel and eluted with ether–EtOAc gradient (v/v, 100:0–2:1) to afford five sub-fractions (Fr.1-1 to Fr.1-5). Fr.1-1 (205 mg) was isolated on silica gel and eluted with ether–CHCl3 gradient (v/v, 5:1–2:1) to obtain four sub-fractions (Fr.1-2-1 to Fr.1-2-4). Fr.1-2-3 (20 mg) was isolated on silica gel and eluted with ether–EtOAc gradient (v/v, 15:1–5:1) to obtain two sub-fractions (Fr.1-2-3-1 to Fr.1-2-3-2). Fr.1-2-3-2 (10 mg) was further purified by the preparative HPLC system with CH3CN–H2O (80:20) as eluent to afford compound 1 (3.8 mg, tR = 7.0 min) and compound 2 (2.8 mg, tR = 8.0 min).

Fr.3 (1.8 g) was separated by Sephadex LH-20 CC eluted with CHCl3–MeOH (v/v, 1:3) to afford three sub-fractions (Fr.3-1 to Fr.3-3). Fr.3-2 (205 mg) was isolated on silica gel and eluted with ether–EtOAc gradient (v/v, 1000:1–1:1) to obtain seven sub-fractions (Fr.3-2-1 to Fr.3-2-7). Fr.3-2-6 (28.4 mg) was isolated on silica gel and eluted with ether–CHCl3 gradient (v/v, 10:1–1:1) to obtain compound 4 (3.6 mg).

Fr.4 (857 mg) was separated into nine subfractions (Fr.4-1 to Fr.4-9) on octadecyl-silylated silica gel column chromatography (ODS CC) with MeOH–H2O (v/v, 50:50–100:0). Fr.4-4 (109 mg) was isolated on silica gel and eluted with ether–EtOAc gradient (v/v, 100:0–1:1) to obtain compound 3 (3.4 mg).

Fr.5 (322 mg) was separated into five subfractions (Fr.5-1 to Fr.5-5) on ODS CC with MeOH–H2O (v/v, 30:70–100:0). Fr.5-3 (13.8 mg) was isolated on silica gel and eluted with ether–EtOAc gradient (v/v, 5:1–0:1) to obtain three sub-fractions (Fr.5-3-1 to Fr.5-3-3). Fr.5-3-2 (10.0 mg) was eluted isocratically with CHCl3–MeOH (50:1) to afford compound 5 (4.8 mg).

**Phomolide A**: white amorphous powder; [α]D25 + 0.07 (c 0.1, MeOH); UV (MeOH) λmax (log ϵ): 214 (3.38), 242 (2.56), 243 (2.73), and 282 (2.97) nm; IR (KBr): 3,325, 2,943, 2,833, 1,662, 1,448, 1,022, 970, and 667 cm−1; HRESIMS: [M + H]+ (calcd for C138H147O92, 209.0815); 1H (500 MHz) and 13C (125 MHz) NMR data (see Table 1).

**Phomolide B**: white amorphous powder; [α]D25 + 4.16 (c 0.1, MeOH); UV (MeOH) λmax (log ϵ): 240 (2.78) and 266 (3.30) nm; IR (KBr): 3,360, 2,933, 2,833, 1,714, 1,680, 1,585, 1,456, 1,166, 1,020, and 669 cm−1; HRESIMS: m/z 209.0815 [M + H]+ (calcld for C138H147O92, 209.0808); 1H (500 MHz) and 13C (125 MHz) NMR data (see Table 1).

**Phomolide C**: white amorphous powder; [α]D25 + 2.17 (c 0.1, MeOH); UV (MeOH) λmax (log ϵ): 200 (3.30) nm; IR (KBr): 3,363, 2,927, 2,858, 1,739, 1,653, 1,516, 1,454, 1,377, 1,251, 1,024, 923, 675, and 597 cm−1; HRESIMS: m/z 237.1851 [M + H]+ (calcld for C138H147O94, 237.1849); 1H (500 MHz) and 13C (125 MHz) NMR data (see Table 3).

**Phomolide D**: yellow oil; [α]D25 + 0.1 (c 0.1, MeOH); UV (MeOH) λmax (log ϵ): 200 (3.30) nm; IR (KBr): 3,363, 2,927, 2,858, 1,739, 1,653, 1,516, 1,454, 1,377, 1,251, 1,024, 923, 675, and 597 cm−1; HRESIMS: m/z 237.1851 [M + H]+ (calcld for C138H147O94, 237.1849); 1H (500 MHz) and 13C (125 MHz) NMR data (see Table 3).

**Phomolide E**: yellow oil; [α]D25 + 4.21 (c 0.1, MeOH); UV (MeOH) λmax (log ϵ): 200 (3.30) nm; IR (KBr): 3,361, 2,929, 2,858, 1,739, 1,653, 1,516, 1,454, 1,379, 1,024, 952, 667, and 597 cm−1; HRESIMS: m/z 237.1854 [M + H]+ (calcld for C138H147O94, 237.1849); 1H (500 MHz) and 13C (125 MHz) NMR data (see Table 3).
Phomene C: yellow crystal; \( \alpha = 0.063 \) (c 0.1, MeOH); UV (MeOH): \( \lambda_{	ext{max}} (\log e) \): 223 (1.60) and 237 (1.67) nm; IR (KBr): 3,315, 2,947, 2,858, 1,651, 1,375, 1,112, 1,029, 912, 665, and 603 cm\(^{-1}\). HRESIMS: \( m/z \) 239.2004 [M + H]\(^+\) (calcd for C\(_{15}\)H\(_{27}\)O\(_2\), 239.2006); \( ^1\)H (500 MHz) and \( ^1\)C (125 MHz) NMR data (see Table 3).

### 4.4 Quantum chemistry calculations

Conformational search of structures was performed by Crest (Pracht et al., 2020), with 4 kcal/mol energy window. Optimization and frequency calculation of the obtained conformer were performed on B3LYP/TZVP (Grimme et al., 2011; Tsuzuki and Uchimaru, 2020) (IEFPCM, C\(_{6}\)D\(_{6}\) and MeOH) level of theory. DFT GIAO \( ^{13}\)C NMR calculation was calculated on the \( \alpha \)b97x/d6-31G* (IEFPCM, C\(_{6}\)D\(_{6}\)) level, and the data processing followed the reported STS protocol. The calculated shielding tensors of conformers were Boltzmann-averaged based on Gibbs free energy. The average calculated ECD curve of each conformer was Boltzmann-averaged based on their Gibbs free energy. Theoretical ECD calculations were performed by Gaussian 16 software package (Frisch et al., 2016).

The calculated ECD curve of each conformer was Boltzmann-averaged based on their Gibbs free energy. The average calculated ECD curve of 1 was adjusted by blue shifting for 20 nm. All DFT calculations were performed by Gaussian 16 software package (Frisch et al., 2016).

### 4.5 X-ray crystallographic data

Crystal data for 6 C\(_{15}\)H\(_{26}\)O\(_2\) (M = 238.36 g/mol): trigonal, space group P\(_3\)\(_2\)\(_1\) (no. 145), a = 13.6894 (2) Å, \( c = 6.60290 (10) \) Å, \( V = 1,071.60 \) (4) Å\(^3\), \( Z = 3 \), \( T = 100.00 \) (10) K, \( \mu (\text{Cu}K\alpha) = 0.552 \) mm\(^{-1}\), Dcalc = 1.108 g/cm\(^3\); 7,072 reflections measured (7.456 \( \leq \Theta \leq 148.49 \)) and 2,779 unique (\( R_{	ext{int}} = 0.0298 \), \( R_{\text{sigma}} = 0.0367 \)), which were used in all calculations. The final \( R_1 \) was 0.0422 \( [I > 2\sigma(I)] \) and \( wR_2 \) was 0.1085 (all data). Flack parameter = -0.05 (12). The crystallographic data for 5 reported in this paper has been deposited in the Cambridge Crystallographic Data Centre (deposition number: CCDC 2192981). Copies of these data can be obtained free of charge via https://www.ccdc.cam.ac.uk.

### 4.6 Cytotoxicity and antimicrobial assays

#### 4.6.1 Cytotoxicity assays

Cytotoxicity was evaluated by the sulforhodamine B assay (Skehan et al., 1990) against five human cancer cell lines (SF-268, MCF-7, HepG2, A549, and LX-2). As a result, none of the compounds showed good cytotoxic activity.

#### 4.6.2 Antimicrobial assays

The antibacterial activities for compounds 1–6 were evaluated against three bacteria embodying S. aureus (CMCC 26003), methicillin-resistant S. aureus (JCSC 3063), and E. coli (ATCC 8739). All of the bacteria were purchased from Guangdong Institute of Microbiology (Guangzhou, China). The MICs were determined by the broth microdilution method in 96-well plates as described in previous literature (NCCLS, 1999; CLSI, 2012; Li et al., 2014). Positive control was vancomycin or kanamycin. All test samples were dissolved in dimethyl sulfoxide and diluted with culture medium.

### Data availability statement

The data presented in the study are deposited in the Cambridge Crystallographic Data Centre, accession number: CCDC 2192981.

### Author contributions

YC, HW, and XK conducted the experiment and collected the experimental data. YC performed the experiments of compound isolation. ZS and ZZ carried out the ECD calculations. YC, ZZ, and HT finished the structure identification of the isolated compounds. MK, WP, JT, and YZ evaluated the activities of all the isolates. YC and HT interpreted the data and wrote the paper. YC, ZZ, and HT revised the manuscript. ZZ and HT conceived and designed the experiments. All authors contributed to the article and approved the submitted version.

### Funding

Financial support for this research was provided by the National Natural Science Foundation of China (82173711), Youth Innovation Promotion Association of CAS (2020342), Natural Science Foundation of Guangdong Province (2019A1515011694), Natural Science Foundation of Hunan Province (no. 2021JJ30917), High-tech Industry Science and Technology Innovation Project of Hunan Province (2020GK4083), Postgraduate Research and Innovation Project of Hunan Province (CXY20210341), Postgraduates Innovation Program of Central South University (Nos. 2021zzts0978, 2021zzts0994, and 2022zzts0899), and the Open Sharing Fund for the Large-Scale Instruments and Equipment of Central South University.

### Acknowledgments

We sincerely thank Ms. Xuan Ma of South China Sea Institute of Oceanology for the X-ray measurements.
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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