

Phomapyrones from blackleg causing phytopathogenic fungi: isolation, structure determination, biosyntheses and biological activity

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Abstract

The isolation and structure determination of phomapyrones D–G, three 2-pyrones and a coumarin, from a group of isolates of the fungal pathogen *Leptosphaeria maculans* (Desm.) Ces. et de Not., asexual stage *Phoma lingam* (Tode ex Fr.) Desm, is reported. As well, phomenin B, infectopyrone, and polanrazines B and C were also obtained for the first time from these isolates. In addition, based on results of incorporations of ¹³C-labeled acetate and malonate, and deuterated methionine, a polyketide pathway is proposed for the biosyntheses of phomapyrones.

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Keywords: *Brassica napus*; Canola; *Sinapis alba*; White mustard; *Brassica juncea*; Brown mustard; Cruciferae; *Leptosphaeria maculans*; *Phoma lingam*; Phytotoxin; 2-pyrones; Polyketide biosynthesis

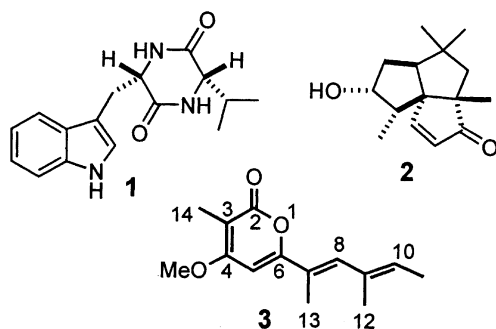
1. Introduction

Blackleg of crucifers is a plant disease caused by the fungal pathogen *Leptosphaeria maculans* (Desm.) Ces. et de Not., asexual stage *Phoma lingam* (Tode ex Fr.) Desm (Gugel and Petrie, 1992). Recent reclassification of this species introduced the new name *Leptosphaeria biglobosa* to enclose the isolates traditionally known as avirulent, weakly virulent or B group (*L. maculans*/*P. lingam* isolates least damaging), while the virulent group (most damaging) remains *L. maculans*/*P. lingam* (for a recent review see Hewlett et al., 2001). We have analyzed a large number of blackleg causing fungi and reported

the secondary metabolites produced by a new group of isolates able to infect brown mustard (*Brassica juncea* L.), a crop previously known to be resistant to blackleg disease. These new isolates were found to produce as major metabolites polanrazine A (1) (Pedras et al., 1998), phomalairdenone A (2) and phomapyrone A (3) (Pedras et al., 1999). In continuation of that work we have isolated from cultures of isolates Laird 2 and Mayfair 2, four new metabolites which we named phomapyrones D–G (4–7). As well, phomenin B (8), infectopyrone (9), and polanrazines B (10) and C (11) were isolated for the first time from these isolates. Here we report the structure determination of metabolites 4–7, and propose a biosynthetic pathway for phomapyrones based on results of incorporations of ¹³C-labeled acetate and malonate, and deuterated methionine, as well as lack of incorporation of deuterated propanoic acid.

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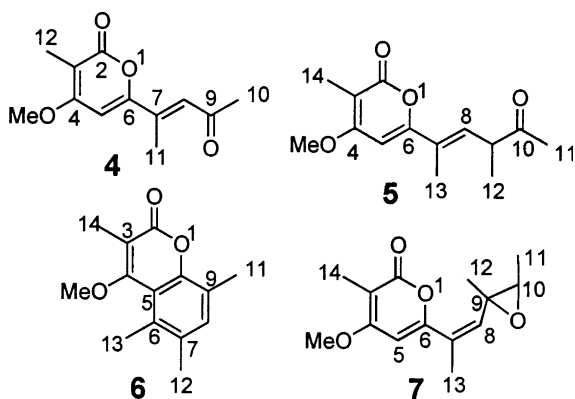
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2. Results

2.1. Isolation and structure determination

Extracts (2.3 g) of cultures of isolate Mayfair 2 (*L. maculans*/*P. lingam*) were fractionated by flash column chromatography to yield the new phomapyrones D (**4**, 10 mg), E (**5**, 3.5 mg), F (**6**, 4 mg), and G (**7**, 6.5 mg), as well as phomapyrone A (**4**, 36 mg), phomenin B (**8**, 28 mg), infectopyrone (**9**, 4 mg), and polanrazines B (**10**, 4.5 mg) and C (**11**, 21 mg).



The molecular formula of phomapyrone D (**4**) was determined to be $C_{12}H_{14}O_4$ (HREI-MS), indicating six degrees of unsaturation. Analysis of the 1H and ^{13}C NMR spectroscopic data indicated the presence of three methyl groups, one methoxy group, and eight sp^2 hybridized carbons (Tables 1 and 2). Analysis of the HMQC and HMBC data was instrumental in the assignment of the structure. Among the three carbonyl-like carbons, the carbon at δ_C 165.2 was assigned to C-2 as it displayed only a correlation with the methyl protons at δ_H 2.05 (H_3 -12). Likewise, the carbon at δ_C 164.5 displayed correlations with methyl protons at δ_H 2.05, methoxy protons at δ_H 3.95 and methine proton at δ_H 6.52 (H -5), allowing its assignment as C-4. An addi-

Table 1

1H NMR (500 MHz) chemical shifts (ppm), multiplicity, J (Hz) of phomapyrones D (**4**), E (**5**), F (**6**), and G (**7**) in $CDCl_3$

Hydrogens	4	5	6	7
5	6.52, s	6.21, s	—	6.19, s
8	7.15, s	6.51, d, 10	7.17, s	6.02, s
9	—	3.53, m	—	—
10	2.36, s	—	—	3.05, q, 6
11	2.36, s	2.17, s	2.41, s	1.46, d, 6
12	2.05, s	1.30, d, 7	2.33, s	1.50, s
13	—	2.00, s	2.61, s	1.99, s
14	—	1.97, s	2.19, s	1.96, s
OMe	3.95, s	3.94, s	3.82, s	3.88, s

Table 2

^{13}C NMR (125 MHz) chemical shifts (ppm) of phomapyrones D (**4**), E (**5**), F (**6**), and G (**7**) (in $CDCl_3$)

Carbons	4	5	6	7
2	165.2	165.2	164.5	164.8
3	106.3	103.4	114.1	103.5
4	164.5	166.0	167.1	165.5
5	97.5	93.3	116.1	95.4
6	158.2	159.4	133.3	158.9
7	139.8	128.8	131.1	127.1
8	126.5	132.8	134.6	137.7
9	199.7	48.2	123.5	60.3
10	33.0	208.0	150.6	62.4
11	13.9	28.4	16.1	14.5
12	9.4	16.9	20.9	17.9
13	—	13.3	16.7	21.0
14	—	9.2	10.9	9.1
Ome	56.7	56.6	60.7	56.6

tional correlation of the methyl group at δ_H 2.05 (H_3 -12) with a tetrasubstituted olefinic carbon at δ_C 106.3 allowed the assignment of C-3. Further correlations of the proton at δ_H 6.52 with the carbon at δ_C 158.2 (C-6), and of this carbon with a methyl group at δ_H 2.36 (H_3 -11) allowed the complete assignment of a 2-pyrone unit and established that a substituent was located at C-6. The partial structure of this substituent was deduced similarly. The long-range correlation of an olefinic proton at δ_H 7.15 (H -8) with both a carbonyl carbon at δ_C 199.7 (C-9) and an sp^2 carbon at δ_C 139.8 (C-7) suggested the presence of an α,β -unsaturated ketone in the side chain. Further, the correlation of the carbonyl carbon at δ_C 199.7 (C-9) with a methyl group at δ_H 2.36 (δ_C 33.0) allowed the complete assignment of the side chain. Finally, the configuration of the double bond was assigned as 7-*E* on the basis of NOE data, after an NOE enhancement observed on the signal of H -5 upon irradiation of the proton at δ_H 2.36 (H -11), and vice versa (no effect on signal of H -8, δ_H 7.15). Hence, the structure **4** was unambiguously assigned to phomapyrone D.

Phomapyrone E (**5**) had a molecular formula of $C_{14}H_{18}O_4$ (HREI-MS) indicating six degrees of unsaturation. The 1H and ^{13}C NMR spectra showed the presence of four methyl groups, one methoxy group and

eight sp^2 hybridized carbons. The 1H NMR spectrum displayed a spin system containing a proton at δ_H 3.53 (H-9) coupled to an olefinic proton at δ 6.51 (H-8, d, $J = 10$ Hz) and methyl protons at δ_H 1.30 (H₃-12, d, $J = 7$ Hz). As for phomapyrone D (**4**), the HMBC spectrum displayed the typical correlations of a 3,4,6-trisubstituted-2-pyrone. For instance, long-range correlations were observed between the proton at δ_H 6.21 (H-5) and carbons at δ_C 103.3 (C-3), δ_C 166.0 (C-4), and δ_C 159.4 (C-6). The long-range correlation of protons at δ_H 6.51 (H-8) and at δ_H 2.00 (H₃-13) with C-6 (δ_C 159.4) and the proton at δ_H 6.21 (H-5) with a carbon at δ_C 128.8 (C-7) suggested the connection of C-6 of 2-pyrone moiety to C-7 of the side chain. Additional long-range correlations were also observed between the methyl protons at δ_H 1.97 (H₃-14) and carbons at δ_C 165.2 (C-2) and δ_C 103.3 (C-3). Moreover, the proton at δ_H 3.53 (H-9) showed a long-range correlation with a carbonyl carbon at δ_C 208.0, which was assigned to C-10. The methyl protons at δ_H 2.17 (δ_C 28.3) showing the only long-range correlation with carbon C-10 (δ 208.0) were assigned to H₃-11. The *E* stereochemistry of the double bond at C-7 was deduced on the basis of NOE difference data; irradiation of proton at δ_H 2.00 (H₃-13) caused an NOE enhancement on signal at δ_H 6.21 (H-5), and vice versa, but it had no effect on the proton at C-8 (δ_H 6.51). Hence, the structure of phomapyrone E was assigned unambiguously as **5**.

The molecular formula of phomapyrone F (**6**) was established as $C_{14}H_{16}O_3$ on the basis of HREI-MS and NMR spectroscopic data. The 1H NMR spectrum displayed signals for four methyl groups, one methoxy group, and one sp^2 hybridized methine, accounting for the total number of protons. The proton decoupled ^{13}C NMR spectrum displayed 14 carbon resonances, of which three had chemical shift above δ_C 150 suggestive of oxygen-bearing sp^2 carbons. The HMBC spectrum displayed all characteristic correlations indicating the presence of a fully substituted 2-pyrone moiety. The carbon signal at δ_C 164.5 correlating with methyl group at δ_H 2.19 was assigned to C-2. As for phomapyrones **4** and **5**, the methyl group C-14 (δ_H 2.19, H₃-14) showed further long-range correlations with carbons at δ_C 114.1 and 167.1, which were assigned to C-3 and C-4, respectively. C-4 also exhibited a long-range correlation with the methoxy group at δ_H 3.82. Additionally, analysis of HMBC data indicated the presence of an aromatic nucleus fused to the 2-pyrone moiety along the C-5–C-10 bond, to form a coumarin ring. The positions of substitution of the remaining methyl groups at δ_H 2.61, 2.33 and 2.41 were determined by analysis of NOE difference data. Thus, irradiation of methyl protons at δ_H 2.61 caused an NOE enhancement on the methoxy group (δ_H 3.82) and methyl group at δ_H 2.33 (H₃-12), indicating the location of these methyl groups on C-6 (δ_H 2.61, H₃-13) and C-7 (δ_H 2.33, H₃-12),

respectively. Likewise, irradiation of proton at δ_H 7.17 caused an NOE enhancement on the signals of methyl protons at δ_H 2.41 (H₃-11) and 2.33 (H₃-12) indicating the methyl group at δ_H 2.41 is located at C-9. Hence, phomapyrone F has structure **6**.

The molecular formula of phomapyrone G (**7**) was established on the basis of HRMS and NMR data as $C_{14}H_{18}O_4$ indicating 6° of unsaturation. The 1H NMR spectrum displayed signals of two olefinic, one methine, and five methyl proton which accounted for the total number of hydrogens. The ^{13}C NMR spectrum displayed fourteen carbon signals, three of which were sp^2 hybridized attached to oxygen, on the basis of their chemical shifts. As for the structural assignments of phomapyrones **4–6**, HMBC data allowed the assignment of the signal at δ_C 164.8 to C-2 based on the only long-range correlation with the methyl group at δ_H 1.96. Similarly, another low field carbon at δ_C 165.5, showing a long-range correlation with the methoxy group at δ_H 3.88, was assigned to C-4. The latter also exhibited long-range correlations with the methyl group at δ_H 1.96 (H₃-14) and the proton at δ_H 6.19 (H-5). The long-range correlation of the proton H-5 with the carbon at δ_C 127.1 (C-7), together with the long-range correlation of the proton at δ_H 6.02 (H-8) with the carbon at δ_C 158.9 (C-6), indicated a substituent at C-6. The chemical shifts of two sp^3 hybridized carbons at δ_C 60.3 (C-9) and δ_C 62.4 (C-10) suggested that these carbons were attached to oxygen. Consequently, to satisfy the chemical formula an epoxide was assigned to C-9/C-10. The *Z* stereochemistry of side chain was determined on the basis of NOE data. Irradiation of the proton signal at δ_H 1.99 (H₃-13) caused an NOE enhancement on the signal at δ_H 6.19 (H-5) and δ 6.02 (H-8). Furthermore, irradiation of proton signal at δ_H 6.02 (H-8) caused an NOE enhancement on the signal at δ_H 1.99 (H₃-13) and δ 3.06 (H-10), but had no effect on the signals at δ_H 1.96 (H₃-12) and δ_H 1.47 (H₃-11). Additionally, irradiation of H-10 (δ_H 3.06) caused an NOE enhancement on the signals of H-8 (δ_H 6.02) and H₃-11 (δ 1.47). Therefore, the structure of phomapyrone G was assigned as **7**, however the configurations of the stereogenic centers C-9 and C-10 remain undetermined.

Additional pyrones **8** and **9** were isolated for the first time from isolates Mayfair 2 and Laird 2. After spectroscopic analyses their structures were determined to be identical to the known metabolites phomenin B (**8**) and infectopyrone (**9**), isolated previously from *Phoma tracheiphila* (Tringali et al., 1993) and from *Alternaria infectoria*, respectively (Larsen et al., 2003). Furthermore, dioxopiperazines **10** and **11** were also isolated from isolates Mayfair 2 and Laird 2. These metabolites were shown to be identical to polanrazines B (**10**) and C (**11**) by comparison of spectroscopic data with those of authentic samples (Pedras and Biesenthal, 2001).

Results of phytotoxicity assays using phomapyrones A (3), D (4), and phomenin B (8) indicated no activity even at relatively high concentration (10^{-3} M).

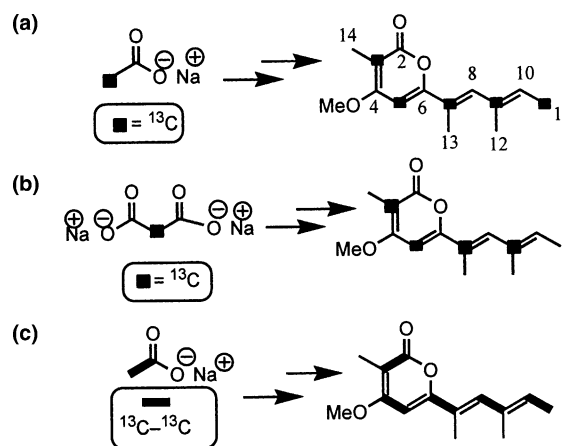
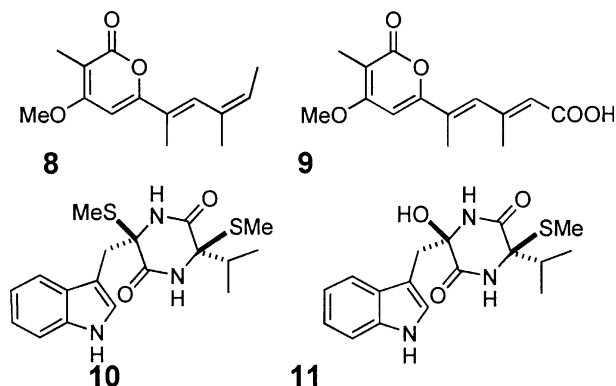


Fig. 1. Labelling pattern of phomapyrone A (3) isolated from cultures containing: (a), sodium $[2-^{13}\text{C}]$ acetate; (b), sodium $[2-^{13}\text{C}]$ malonate; and (c), sodium $[1,2-^{13}\text{C}_2]$ acetate.

2.2. Biosynthetic origin of phomapyrones

Sodium $[2-^{13}\text{C}]$ acetate, sodium $[1,2-^{13}\text{C}]$ acetate, sodium $[2-^{13}\text{C}]$ malonate, $[3,3,3-^2\text{H}_3]$ propanoic acid, and L-[Me- $^2\text{H}_3$]methionine were separately administered to cultures of isolate Mayfair 2. After extraction of the broth, and separation of the extract, phomapyrone A (3) was obtained and analyzed by both NMR spectroscopy and HRMS; on the other hand, phomapyrones 4–7 were obtained in insufficient amount for NMR studies. The results obtained from these experiments are summarized in Fig. 1 and Table 3. The distribution of ^{13}C labelling in phomapyrone A (3) was determined by comparison of the integrated and normalized intensities

of carbon signals of enriched 3 with those of the unenriched (i.e. natural abundance) sample. The ^{13}C NMR spectrum of phomapyrone A (3) isolated from administration of sodium $[2-^{13}\text{C}]$ acetate displayed 14 signals, which were assigned by comparison with the ^{13}C NMR data of the natural abundance metabolite. The signal intensities of C-3, C-5, C-7, C-9, and C-11 were enhanced relative to the other carbons (Table 3, Fig. 1(a)). Completable experiments using sodium $[2-^{13}\text{C}]$ malonate, showed that the ^{13}C NMR signals of C-3, C-5, C-7, and C-9 were more intense, whereas the ^{13}C NMR signal of C-11 was not as enhanced (Table 3, Fig. 1(b)). In addition, the ^{13}C NMR spectrum of phomapyrone A (3) resulting from incorporation of so-

Table 3

Incorporation results of feeding experiments with sodium $[2-^{13}\text{C}]$ acetate and sodium $[2-^{13}\text{C}]$ malonate based on the ^{13}C NMR of phomapyrone A (3)

Carbon	Phomapyrone A natural sample		Phomapyrone A enriched, $[2-^{13}\text{C}]$ acetate			Phomapyrone A enriched, $[2-^{13}\text{C}]$ malonate		
	SI ^a	NSI ^b	SI ^a	NSI ^b	R ^c	SI ^a	NSI ^b	R ^c
C-2	2.09	0.22	1.89	0.31	1.40	0.47	0.18	0.82
C-3	2.54	0.26	3.25	0.54	2.08 ^d	1.49	0.56	2.15 ^d
C-4	3.10	0.32	2.34	0.39	1.22	0.77	0.29	0.91
C-5	8.94	0.92	8.14	1.34	1.46 ^d	5.46	2.05	2.23 ^d
C-6	3.88	0.40	2.52	0.42	1.05	1.01	0.38	0.95
C-7	3.58	0.37	3.29	0.54	1.46 ^d	2.25	0.85	2.29 ^d
C-8	8.28	0.86	5.76	0.95	1.10	2.47	0.93	1.08
C-9	4.53	0.47	4.39	0.72	1.53 ^d	3.66	1.38	2.94 ^d
C-10	6.92	0.71	4.96	0.82	1.15	1.78	0.67	0.94
C-11	7.21	0.74	9.26	1.53	2.06 ^d	3.06	1.15	1.55 ^d
C-12	7.12	0.74	5.47	0.90	1.22	2.34	0.88	1.19
C-13	10.14	1.04	6.40	1.05	1.00	2.65	1.00	0.96
C-14	8.63	0.89	5.85	0.96	1.08	2.30	0.86	0.97
OCH ₃	9.68	1.00	6.07	1.00	1.00	2.66	1.00	1.00

^a SI = ^{13}C signal intensity of sample.

^b NSI = ^{13}C normalized signal intensity of sample relative to OCH₃.

^c R = [^{13}C normalized signal intensity of enriched sample]/[^{13}C normalized signal intensity of natural sample].

^d Enriched carbons.

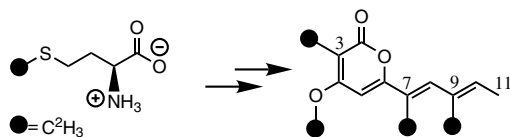


Fig. 2. Labelling pattern of phomapyrone A (3) isolated from cultures containing L-[C²H₃]methionine.

dium [1,2-¹³C]acetate displayed fourteen signals due to natural abundance ¹³C plus 10 doublets flanking carbons C-2 to C-11. The doublets due ¹³C–¹³C coupling indicated the incorporation of five intact C₂ units; however, the carbon signals of methyl groups C-12, C-13, C-14 and OMe did not show satellite peaks (Fig. 1(c)).

After the polyketide origin of phomapyrone A (3) was established, the origin of its four methyl groups, i.e. C-12–C-14 and (O)CH₃ was investigated. Thus, HREI-MS analysis of phomapyrone A (3) isolated from cultures incubated with L-[Me-²H₃]methionine showed incorporation of several deuterium atoms: M⁺ + 12 (4%), M⁺ + 9 (11%), M⁺ + 6 (30%), M⁺ + 3 (70%), relative to naturally occurring compound. However, phomapyrone A (3) isolated from cultures incubated separately with [3,3,3-²H₃]propanoic acid showed no deuterium incorporation by HREI-MS analysis. Consequently, the most likely source of the four C₁ units present in phomapyrone A (3) is L-methionine, as summarized in Fig. 2.

3. Discussion

Metabolites containing a 2-pyrone moiety have been isolated from both terrestrial and marine organisms. Phomapyrones A (3), B, and C, were obtained from cultures of weakly virulent Canadian isolates of *L. maculans*/*P. lingam* (Pedras et al., 1994), whereas phomapyrone A (3) was also reported from Polish isolates (Pedras and Biesenthal, 2001). Phomapyrone A (3) was isolated from *P. tracheiphila* by an independent group and named phomenin A (3), which was reported along with phomenin B (8) (Tringali et al., 1993). Infectopyrone (9) was reported from the phytopathogenic fungi *Alternaria infectoria* and *Stemphylium* sp. as a potential mycotoxin (Larsen et al., 2003). The chemical study of *Ercolanina funereal*, a marine mollusc, led to the isolation of several cyrcenes, one of which is identical to phomapyrone A (3) (Vardaro et al., 1992). Additional 2- and 4-pyrone polypropionates were also reported from other marine molluscs such as *Siphonaria diemenensis* (Hochlowski and Faulkner, 1983) and *Placida dendritica* (Cutignano et al., 2003). Interestingly, some pyrone metabolites from molluscs were suggested to have a defensive role (Di Marzo et al., 1993; Vardaro et al., 1992).

This investigation on the biosynthetic origin of phomapyrone A (3) produced by the fungus *L. maculans*/*P. lingam* indicates its polyketide origin. Namely, the labeling pattern obtained for phomapyrone A (3) after incorporation of [2-¹³C]malonate, [2-¹³C]acetate, and [1,2-¹³C]acetate and the lack of incorporation of propanoic acid. Furthermore, the incorporation of the four C²H₃ groups (HRMS, M + 12, 4%) derived from methionine, suggests the origin of the methyl groups at C-3, C-7, and C-9 and OMe. In conclusion, phomapyrone A (3) was found to be an acetate/malonate derived pentaketide with four methyl groups derived from methionine, as summarized in Fig. 3. These results further suggest that the new phomapyrones D–G (4–7) are of polyketide origin, resulting from a pentaketide chain as well, but phomapyrone D might result from methylation of a tetraketide or decarboxylation of a pentaketide. Previous studies of the biosynthetic origin of 2-pyrones from the fungus *Gliocladium vermoesonii* indicated a polyketide origin (Avent et al., 1992a,b), whereas another study of the biosynthetic origin of 2-pyrone and 4-pyrone metabolites from molluscs indicated a polypropionate origin (Vardaro et al., 1992). The polyketide origin of pyrones from *G. vermoesonii* is consistent with the results reported above. It might be that pyrones isolated from molluscs are polypropionates and those from fungi are polyketides.

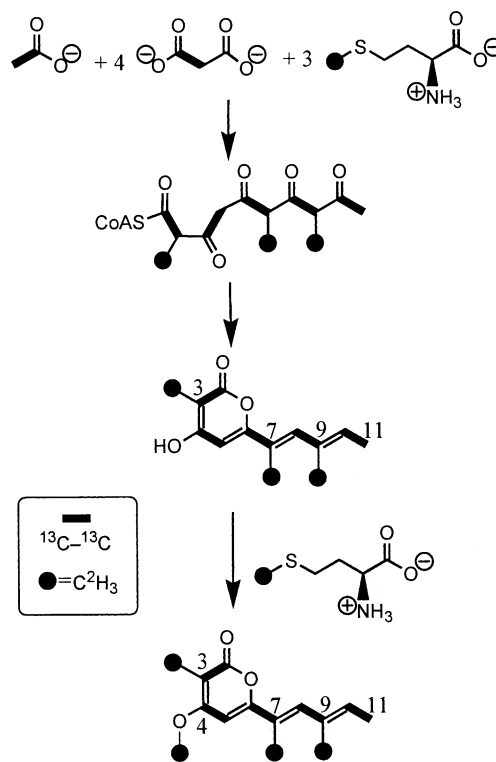


Fig. 3. Biosynthetic pathway of phomapyrone A (3) produced by *Leptosphaeria maculans*/*Phoma lingam*.

4. Experimental

4.1. General

Sodium [2-¹³C]acetate, sodium [1,2-¹³C]acetate, sodium [2-¹³C]malonate, [3,3,3-²H₃]propanoic acid, and L-[Me-²H₃]methionine were purchased from Cambridge Isotope Laboratories, Inc., Andover, MA. Analytical HPLC analysis was carried out with a high performance liquid chromatograph equipped with quaternary pump, automatic injector, and photodiode array detector (wavelength range 190–600 nm), degasser, and a Hyper-sil ODS column (5 µm particle size silica, 4.6 i.d. × 200 mm), equipped with an in-line filter. Mobile phase: H₂O–CH₃CN A:B, v/v, isocratic A:B(1:4), 10 min, gradient 1:4 to 3:2, 10 min, 3:2 to 3:1, 10 min, 3:1(A:B) to B, 10 min, at a flow rate 1.0 mL/min. Samples were dissolved either in CH₃CN or in MeOH. Other conditions are as previously reported (Pedras and Okanga, 1999).

Fungal cultures were grown as reported previously (Pedras and Biesenthal, 2001). Isolates Mayfair 2 and Laird 2 were obtained from Agriculture and Agri-Food Canada, Saskatoon, Saskatchewan, IBCN collection.

4.2. Isolation of metabolites

A total of 122 L cultures were grown in minimal medium (Pedras et al., 1997) for 6 days; the cultures were filtered, and the broth was combined. The broth obtained (110 L) was concentrated (22 L) by freeze drying, and was extracted with EtOAc (800 mL × 3/1 L broth). The combined extracts were dried (anhyd Na₂SO₄) and concentrated to dryness under reduced pressure (2.3 g). The EtOAc extract was subjected to flash Si gel cc and eluted first with CH₂Cl₂ followed by CH₂Cl₂–MeOH (97:3, 200 mL) and then with CH₂Cl₂–MeOH (90:10). All fractions collected were concentrated to dryness under reduced pressure. Fractions F₃ to F₆ were combined, subjected to prep. Si gel TLC, and developed with CH₂Cl₂–MeOH (98:2, for three times) to yield phomapyrone F (6) (4.0 mg). Fraction F₇ from FCC of EtOAc extracts was further subjected to FCC (2 cm × 15 cm, silica gel, 50 mL fractions), eluted first with CH₂Cl₂–EtOAc (95: 5, 200 mL) followed by CH₂Cl₂–EtOAc (90:10, 200 mL) and then CH₂Cl₂–EtOAc (80:20, 200 mL). Twelve fractions were collected and concentrated to dryness under reduced pressure. Fractions F₂ and F₃ were combined (39.5 mg), subjected to prep. TLC (hexane–EtOAc, 9:1, multiple development) to yield phomapyrone A (3) (36 mg) and phomenin B (8) (28 mg). Fractions F₆ to F₉ were combined, further subjected to prep. Si gel TLC (CH₂Cl₂–MeOH, 98:2) to obtain phomapyrone D (4) (10 mg). This material was further applied to prep. TLC (hexane–Et₂O, 50:50, multiple elution) to yield phomapyrone E (5) (5 mg), phomapyrone G (7) (6.5

mg), and polanrazine B (10, 4.5 mg), respectively. Fractions F₁₀–F₁₂ were combined and subjected to prep. TLC (CH₂Cl₂–MeOH, 97:3, developed three times) to yield infectopyrone (9) (4 mg) and polanrazine C (11, 21 mg).

4.3. Phytotoxicity assays

4.3.1. Phytotoxicity

Seeds of canola (*Brassica napus*) cv. Westar, white mustard (*Sinapis alba*) cv. Ochre, and brown mustard (*Brassica juncea*) were obtained from Plant Gene Resources, Agriculture and Agric-Food Canada Research Station, Saskatoon, SK. The seeds were sown in commercial potting soil mixture, and plants were grown in a growth chamber, under controlled environmental conditions, as previously reported (Pedras and Biesenthal, 2001). Phytotoxicity assays were conducted on leaves of 2-week-old plants with phomapyrones A (3), D (4) and phomenin B (8) (5 × 10^{−4} and 1 × 10^{−3} M in H₂O–MeOH, 1:1, v/v) as previously described (Pedras and Biesenthal, 2001). Plants were incubated in a growth chamber for a week.

4.4. Pressure administration experiments

Cultures of isolate Mayfair 2 (250 mL flasks with 100 mL per flask, total 2 L, 5 × 10⁸ spores per 100 mL) were incubated on a shaker at 120 rpm under fluorescent light. Each flask was administration, of 48 and 72 h with [2-¹³C]acetate at a concentration of 25 mg of sodium [2-¹³C]acetate per 100 mL of culture. After 6 days of incubation the culture was filtered, and worked up as described above. The EtOAc extract (32 mg) was separated by RP-FCC (1 cm × 10 cm, RP-C18, 20 mL fractions) and eluted first with CH₃CN–H₂O (2:3, 1 mL) followed by CH₃CN (40 mL). Ten fractions were collected and concentrated to dryness under reduced pressure. Fractions F₅ and F₆ were combined (5.0 mg) and further subjected to prep. TLC (hexane–EtOAc, 9:1, multiple elution) to yield phomapyrone A (3, 2.8 mg).

Similar experiments were carried out with sodium [2-¹³C]malonate (10 L of culture to yield phomapyrone A, 6.5 mg), [1,2-¹³C]acetate (5 L of culture to yield phomapyrone A (3), 4 mg), [3-²H₃]propanoic acid (5 L of culture to yield phomapyrone A (3), 4.5 mg), and L-[C²H₃]–methionine (5 L of culture to yield phomapyrone A (3), (3.5 mg).

4.5. Characterization data

4.5.1. Phomapyrone D (4)

HPLC *R*_t = 18.9 min for ¹H and ¹³C NMR spectra, see Tables 1 and 2; UV λ_{max} 210, 250, 350 nm; FTIR (cm^{−1}) 2996, 1681, 1357, 1013, 746; HREI-MS: *m/z*

222.0893, calc. for $C_{12}H_{14}O_4$ 222.0892, EI-MS: m/z (relative intensity) 222 (100%), 207 (17.9%), 194 (17%), 179.06 (20%), 159 (49%), 139 (24%), 57 (56%).

4.5.2. Phomapyrone E (5)

HPLC R_t = 22.2 min for 1H and ^{13}C NMR spectra, see Tables 1 and 2; UV λ_{max} 230, 335 nm; FTIR (cm^{-1}) 2970, 2927, 1692, 1554, 1355, 1170, 1013, 751; HREI-MS: m/z 250.1206, calc. for $C_{14}H_{18}O_4$ 250.1205, EI-MS: m/z (relative intensity) 250 (13%), 208 (100%), 165 (11%).

4.5.3. Phomapyrone F (6)

HPLC R_t = 30.4 min. for 1H and ^{13}C NMR spectra, see Tables 1 and 2; UV λ_{max} 210, 290 nm; FTIR (cm^{-1}) 2925, 1711, 1590, 1202, 1093; HREI-MS: m/z 232.1095, calc. for $C_{14}H_{16}O_3$ 232.1099, EI-MS: m/z (relative intensity) 232 (100%), 217 (56%), 189.09 (63%).

4.5.4. Phomapyrone G (7)

HPLC R_t = 23.0 min. for 1H and ^{13}C NMR spectra, see Tables 1 and 2; UV λ_{max} 230, 330 nm; FTIR (cm^{-1}) 2974, 2929, 1700, 1554, 1247, 1166, 1025; HREI-MS: m/z 250.1202, calc. for $C_{14}H_{18}O_4$ 250.1205, EI-MS: m/z (relative intensity) 250 (20%), 208 (66%), 139(49%), 83(28%).

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