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WASALEXINS A AND B, NEW PHYTOALEXINS FROM WASABI: ISOLATION, SYNTHESIS, AND ANTIFUNGAL ACTIVITY

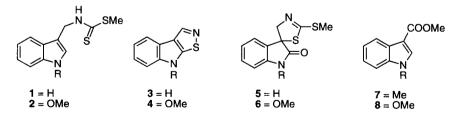
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Abstract: The chemical structure determination of two phytoalexins from wasabi (Wasabia japonica, syn. Eutrema wasabi), a plant resistant to virulent isolates of the blackleg fungus [Leptosphaeria maculans (Desm.) Ces. et de Not., asexual stage Phoma lingam (Tode ex Fr.) Desm.], as well as their synthesis and antifungal activity towards isolates of P. lingam and P. wasabiae is reported. © 1999 Elsevier Science Ltd. All rights reserved.

Phytoalexins, plant secondary metabolites synthesized de novo in response to diverse forms of stress, including fungal infection, are part of the plants' chemical and biochemical defense mechanisms.^{1,2} The phytoalexins from cruciferous plants (family *Cruciferae*, e.g. broccoli, cauliflower, cabbage, rapeseed, mustard, wasabi),³ as for example **1-6**, contain an indole or related ring and at least one sulfur atom.

Recent results established that wasabi (Wasabia japonica, syn. Eutrema wasabi) is resistant to the blackleg fungus (Leptosphaeria maculans (Desm.) Ces. et de Not., asexual stage Phoma lingam (Tode ex Fr.) Desm.), the causative agent of one of the most damaging diseases of the agronomically important oilseed crop, canola (Brassica napus and B. rapa).⁴ Consequently, it became worthwhile to establish the chemical traits responsible for the resistance of wasabi to P. lingam, as such traits are potentially transferable to canola. Subsequently, we have investigated and reported the structures of several constitutive antifungal compounds produced by wasabi, as well as the phytoalexin methyl 1-methoxyindole-3-carboxylate (8), and their activity towards isolates of P. lingam and P. wasabiae.⁵ Herein we wish to communicate studies carried out to establish the structure of two new phytoalexins, which we named wasalexins A (10a) and B (10b), produced by foliar tissue of wasabi under elicitation by P. lingam, P. wasabiae or CuCl₂.



In preliminary experiments, a time-course response of wasabi plants to abiotic elicitation was investigated. After eliciting and incubating plants, leaves were excised at intervals and extracted as previously reported;⁵ control leaves were collected and treated in similar manner. Each extract was bioassayed for

antifungal activity against *Cladosporium cucumerinum* (biodetection on TLC plates employing spores of *C. cucumerinum*) and analyzed by HPLC.⁶ The extracts of both elicited and control leaves showed similar zones of antifungal activity; however, the HPLC chromatograms of hexane extracts of elicited leaf tissue displayed two peaks with $r_T = 18.0$ and 23.8 min, which were not detectable on the chromatograms of extracts of control tissues. The structure of the compound with $r_T = 18.0$ was readily determined to be phytoalexin **8**, as previously reported.⁵ It is worthy to note that the related 1-methylindole **7** was found to be a constitutive chemical defense present in roots of a cruciferous weed (*Camelina sativa*).⁷

Nonetheless, the structure of the compound with $r_{T} = 23.8$ min, a bright yellow metabolite, could not be unambiguously assigned. The HRMS spectroscopic data indicated a molecular formula of $C_{13}H_{14}N_2O_2S_2$, further corroborated by analysis of the NMR spectroscopic data. As in compound **8**, the NMR spectra of the yellow metabolite displayed five aromatic signals ($\delta_{H}8.03$, s, 7.86, d, 7.27, dd, 7.04, dd, 6.96, d; δ_{C} 139.1, 128.8, 125.0, 122.9, 107.1), as well as signals due to a MeO group ($\delta_{H}4.01$; δ_{C} 64.2), two overlapping MeS groups ($\delta_{H}2.70$; δ_{C} 16.2), and two carbonyl or equivalent carbons (δ_{C} 164.8 and 177.5).⁸

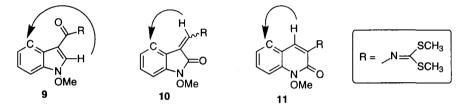
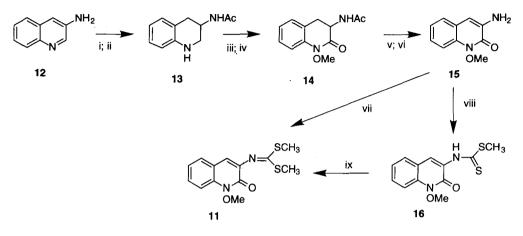


Figure 1. Proposed structures of wasabi yellow metabolite and selected ¹³C-H long-range correlation observed in the HMBC spectrum.

Analysis of HMQC and HMBC data of the yellow compound established several C-H correlations reminiscent of an indole nucleus such as 9 or 10 (Figure 1); however, structure 9 corresponded to methoxybrassenin B, a phytoalexin previously isolated from *B. oleracea*,⁹ whose NMR data did not correspond to those obtained for the yellow wasabi metabolite. Furthermore, the C-H correlation observed on the HMBC spectrum was thought more likely to correspond to a three-bond correlation $({}^{3}J_{CH})$ than to a four-bond correlation $({}^{4}J_{CH})$ and thus structure 11 appeared more plausible than 10.¹⁰ To prove the proposed structure, synthesis of the methoxyquinolone derivative 11 was undertaken.

The synthesis of compound 11 was achieved in seven (or eight) steps from 3-aminoquinoline (12), following the route shown in Scheme 1. After protection of the amino group of 3-aminoquinoline (12), the pyridine ring was reduced using sodium cyanoborohydride¹¹ to yield tetrahydroquinoline 13, which upon treatment with hydrogen peroxide employing sodium tungstate as a catalyst,¹² followed by methylation with diazomethane yielded methyl hydroxamate 14. Methyl hydroxamate 14 was then dehydrogenated with DDQ, hydrolyzed to the free amine 15, which was transformed into the desired quinoline derivative 11 upon treatment with carbon disulfide followed by methyl iodide. The spectroscopic data of compound 11, an almost white compound, were not identical to those of the bright yellow wasabi metabolite.¹³

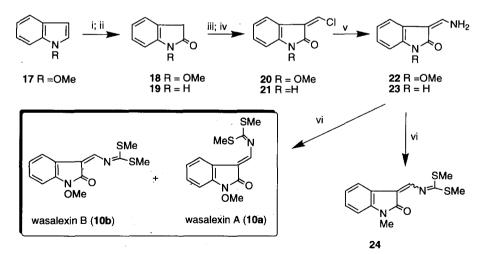
Scheme 1



(i) (Ac)₂O/py, 92%; (ii) NaCNBH₃/AcOH, 91%; (iii) Na₂WO₄•2H₂O/H₂O₂; (iv) CH₂N₂, 62%; (v) DDQ, 100°C, 14%; (vi) HCl/MeOH, 80%; (vii) NaH, CS₂; CH₃I, 30%; (viii) CS₂, Et₃N; CH₃I, 24%; (ix) NaH, CH₃I, 72%

Next, the synthesis of compound 10 was sought; however, the synthesis of analog 24 was carried out first, since the required starting material 2-oxindole (19) was commercially available. Synthesis of 24 was carried out as summarized in Scheme 2; thus, 19 was allowed to react with ethyl formate,¹⁴ the resulting enol transformed into enamine 23 via 3-chloromethyleneoxindole (21),¹⁵ which in turn yielded desired product 24 upon treatment with carbon disulfide and methyl iodide.

Scheme 2



(i) NBS/t-BuOH; (ii) Zn/AcOH, 60%; (iii) HCO₂Et/NaOEt, R = OMe 87%, R = H 81%; (iv) SOCl₂, R = OMe = H 92 %; (v) NH₄OH, R = OMe 94%, R = H 55%; (vi) THF/NaH/CS₂; MeI, R = OMe 53%, R = H 24%.

Compound 24 had a bright-yellow color and spectroscopic data similar to those of the wasabi metabolite.¹⁶ Consequently, compound 10 was then prepared; methoxyindole 17^5 was converted to the desired starting material 1-methoxy-2-oxindole (18), upon bromination with NBS followed by reduction.¹⁷ Synthesis of 10 from 18 followed a route similar to that described for 24, as shown in Scheme 2. While compound 24 was obtained as a single isomer according to NMR and HPLC data, 10 was obtained as a mixture of *E/Z* isomers (1:2), which could be separated by reversed-phase TLC.¹⁸ The spectroscopic data and HPLC retention time of the minor synthetic isomer 10 were identical to those of the natural yellow metabolite which we named wasalexin A. The spectroscopic and HPLC data obtained for the major synthetic isomer were identical to a minor compound isolated together with wasalexin A, which was named wasalexin B.¹⁸

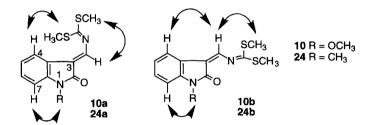


Figure 2. Selected NOE correlations observed for compounds 10a and 24a.

To complete the structure elucidation of wasalexins, assignment of the configuration of the exocyclic double bond was required. Thus, NOE experiments with wasalexin A and the analog **24a** were carried out; as expected, irradiation of the (O)CH₃ and CH₃ groups of **10a** and **24a**, respectively, caused an NOE enhancement of H-7, while irradiation of H-4 caused an enhancement of the (S)CH₃ signals and vice-versa. Conversely, no enhancements were observed for H-1' upon irradiation of H-4 and vice-versa, as indicated in Figure 2. These results allowed the unambiguous assignment of the structures of wasalexin A as **10a**, the major compound produced by elicited wasabi leaves, and wasalexin B as **10b**.¹⁹

Finally, antifungal bioassays established that wasalexin A (10a), similar to phytoalexin 8, was active against *P. lingam* utilizing a spore germination and growth assay, causing 40–50% germination inhibition relative to controls, after 24 h, whereas compound **24a** had no detectable inhibitory effect.²⁰ Interestingly, the spore germination of *P. wasabiae* was not affected significantly by wasalexin A, although the mycelial growth of cultures incubated with wasalexin A was slower. Further bioactivity studies are underway to better understand the role of these compounds in the resistance of wasabi to *P. lingam*, the major pathogen responsible for the blackleg disease of canola.

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References and Notes

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- 6. 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 Hypersil ODS column (5 μm particle size silica, 4.6 id × 200 mm), equipped with a low-dispersion column-inlet filter. Mobile phase was a linear gradient from 75% H₂O 25% CH₃CN to 100% CH₃CN, at a flow rate 1.0 mL/min over 35 min.
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- Spectroscopic data of yellow metabolite (10a): HPLC r_T = 23.8 min; ¹H NMR (CD₂Cl₂): δ 2.70 (s, 6H), 4.01 (s, 3H), 6.96 (d, J = 8 Hz, 1H), 7.04 (dd, J = 8, 7 Hz, 1H), 7.27 (dd, J = 8, 7 Hz, 1H), 7.86, (d, J = 7 Hz, 1H), 8.03 (s, 1H); ¹³C NMR (CD₂Cl₂): δ 16.0 (2 × q), 64.2 (q), 107.1 (d), 116.8 (s), 119.6 (s), 122.9 (d), 125.0 (d), 128.7 (d), 139.0 (s), 139.1 (d), 164.8 (s), 177.5 (s); HREIMS *m/z* measured: 294.0499 (294.0497 calcd. for C₁₃H₁₄N₂O₂S₂); EIMS *m/z* (% relative abundance): 294 [M]⁺ (100), 263 (30), 247 (35), 216 (49), 143 (60); CIMS (NH₃) *m/z* (% relative abundance): 295 [M+1]⁺ (97), 265 (100), 163 (34), 146 (83), 122 (38); FTIR ν_{max}: 2924, 1708, 1625, 1481, 1095, 743 cm⁻¹.
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- 16. Spectroscopic data of **24a**: ¹H NMR (CDCl₃): δ 2.69 (s, 6H), 3.25 (s, 3H), 6.78 (d, J = 8 Hz, 1H), 7.00 (dd, J = 8, 7 Hz, 1H), 7.22 (dd, J = 8, 7 Hz, 1H), 7.84, (d, J = 7 Hz, 1H), 8.08 (s, 1H); ¹³C NMR (CDCl₃): δ 15.9 (2 × q), 26.1 (q), 107.7 (d), 116.8 (s), 119.2 (s), 122.0 (d), 124.5 (d), 128.3 (d), 138.5 (s), 142.8 (d), 169.5 (s), 175.4 (s); HREIMS *m/z* measured: 278.0549 (278.0548 calcd. for

 $C_{13}H_{14}N_2OS_2$; EIMS *m/z* (% relative abundance): 278 [M]⁺ (42), 231 (100), 216 (53), 1158 (24); CIMS (NH₃) *m/z* (% relative abundance): 279 [M+1]⁺ (10), 162 (100), 148 (37); FTIR ν_{max} : 2921, 1693, 1608, 1484, 1363, 1331, 929, 746 cm⁻¹.

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- 18. Preparative TLC: RP-8 plates, $(20 \times 20 \text{ cm}, 0.25 \text{ mm}; \text{EM Science, NJ})$ acetonitrile:water (7:3). It is worthy to note that wasalexin A (10a) appeared to be stable in CH₂Cl₂ but a small amount of isomerization to 10b occurred during NMR acquisitions in CDCl₃; however, 10b could not be obtained as a single compound due to isomerization to 10a (<10%) during purification.
- 19. It is possible that wasalexin B (10b) is an artifact of the isolation process; however, considering that it was present in preliminary fractionation of leaf hexane extracts, and that wasalexin A also is isomerized, it appears more likely that both isomers are naturally occurring. Spectroscopic data of wasalexin B (10b): HPLC $r_T = 21.5 \text{ min;}^{1}\text{H} \text{ NMR} (\text{CD}_2\text{Cl}_2)$: $\delta 2.69 (s, 6\text{H})$, 3.99 (s, 3H), 6.96 (d, J = 8 Hz, 1H), 7.04 (dd, J = 8, 7 Hz, 1H), 7.27 (dd, J = 8, 7 Hz, 1H), 7.46, (d, J = 7 Hz, 1H), 7.91 (s, 1H); ¹³C NMR (CD₂Cl₂): $\delta 15.8 (2 \times \text{q})$, 63.9 (q), 107.2 (d), 114.3 (s), 119.4 (s), 120.2 (d), 122.0 (d), 128.2 (d), 137.9 (d), 139.0 (s), 161.7 (s), 173.7 (s); HREIMS *m/z* measured: 294.0497 (294.0497 calcd. for C₁₃H₁₄N₂O₂S₂).
- 20. Bioassays to determine pycnidiospores germination were carried out after modification of a method previously reported (Pedras, M. S. C.; Biesenthal, C. J. *Can. J. Microbiol.* **1998**, *44*, 547). That is, Nunclon 4-well multidish plates containing 1×10^7 spores / mL in 500 µL solutions of compound to be tested (3% Tween 80 and 2% DMSO in minimal medium) were incubated still under constant fluorescent light, at $24 \pm 1^\circ$ C. The percentage of spore germination was determined by counting about 300 spores (in randomly selected fields) for each replicate.