## A Novel Alkaloid Serantrypinone and the Spiro Azaphilone Daldinin D from *Penicillium thymicola*

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The novel quinazoline metabolite serantrypinone (1) has been isolated from an isolate of the microfungus *Penicillium thymicola* together with daldinin D (2), a new peracetylated spiro azaphilone derivative. The structures of 1 and 2 were elucidated by analysis of spectroscopic data, including 2D NMR, and comparison with literature data.

Recently a large number of suspected ochratoxin Aproducing *Penicillium* isolates were investigated in a chemotaxonomic study.<sup>1</sup> The study divided the isolates into two large groups representing the two species *P. verrucosum* and *P. nordicum* together with a small group of four non-ochratoxin-producing isolates. This latter group included the isolate of *P. thymicola* recently studied by our group.<sup>2</sup>

In addition to displaying different morphological characteristics the three species can be distinguished by differences in production of quinazoline and other secondary metabolites.<sup>1</sup> *P. verrucosum* produces verrucines,<sup>3</sup> *P. nordicum* produces anacines,<sup>1</sup> while *P. thymicola* produces anacine, fumiquinazoline F, and the related spiroquinazoline alantrypinone.<sup>1,2</sup> From previous work with UV-guided isolation of secondary metabolites from *P. thymicola*<sup>2</sup> we knew that the fungus often produces a minor compound with UV data almost identical to those seen for a major component, alantrypinone (**4**).<sup>2</sup> The present report describes the isolation and structure elucidation of this minor and novel quinazoline compound, serantrypinone (**1**), together with a new azaphilone derivative, daldinin D (**2**), from *P. thymicola* (IBT 5891).



The molecular composition of serantrypinone ( $C_{21}H_{16}O_4N_4$ ) was established by HREIMS ([M<sup>+</sup>] at m/z 388.1170). This result is consistent with the presence of an additional oxygen atom in **1** when compared to **4**. NMR data of **1** showed signals characteristic of an oxindole unit as in **4** 

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and a quinazoline moiety derived from anthranilic acid. HMBC experiments predicted the linkage between C-3 in the oxopiperazine ring and C-17 in the tryptophane-derived oxindole as in 4. However, a significant difference between the two compounds was revealed by the chemical shift value of C-16, indicating the oxygenation in 1. In agreement with this assignment, H-16 appeared as two doublets at  $\delta$  3.57 and 3.71 instead of the singlet at  $\delta$  1.19 in **4**. On the basis of this information, 1 can be established as a new structure resulting from the exchange of the alanine residue in 4 for a serine residue. The compound is named serantrypinone, according to the nomenclature of Penn et al.<sup>4</sup> The circular dichroism spectrum of **1** was similar to that of alantrypinone,<sup>2</sup> enabling the assignment of the corresponding absolute configuration at C-3 (3R), C-14 (14R), and C-17 (17S).<sup>3</sup>

A new azaphilone derivative, daldinin D (2), was also isolated from IBT 5891. The molecular formula of daldinin D (C<sub>21</sub>H<sub>24</sub>O<sub>10</sub>) was established by HREIMS ([M<sup>+</sup>] at m/z436.1369). The <sup>1</sup>H and <sup>13</sup>C NMR data showed three acetate groups and two trisubstituted olefinic units. The UV spectrum displayed absorptions characteristic of  $\alpha, \beta, \gamma, \delta$ conjugated carbonyl groups. These data, together with careful investigation of the H-H COSY and HMBC spectra, led to the establishment of the tricyclic spiro-acetal structure 2 almost identical to that of daldinin C produced by Daldinia concentrica.<sup>5</sup> Compounds 2 and 3 only differ with respect to their substitution at position 4. The linkage of the partial structures was inferred from the long-range correlations observed between H-1, H-4, H-9 and CH<sub>3</sub>-13 to C-3, a quaternary carbon at  $\delta$  100.9. This <sup>13</sup>C NMR chemical shift is characteristic of acetal groups. The absolute configuration of 3 was established by X-ray crystallography and chemical degradation.<sup>5</sup> The CD spectrum of 2 was similar to that of an authentic sample of 3 (see experimental data for both compounds), suggesting identical absolute configuration. The stereostructure proposed is coherent with the correlations observed in the NOESY spectrum of **2** (Figure 1).

Interestingly, the likely polyketide-derived structure of **2** resembles those of ochratoxin A produced by the closely related species *P. verrucosum* and *P. nordicum*, and citrinin produced only by *P. verrucosum*. This indicates that the three species recently demonstrated<sup>1</sup> to share the capability to produce the same pattern of volatile metabolites and

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Figure 1. Important NOE correlations observed in daldinin D (2).

verrucolone metabolites might also have very similar genes for polyketide biosynthesis.

In conclusion, the present study describes some further metabolites unique to *P. thymicola*. Detection of these compounds in either pure fungal cultures or in contaminated food and feed stuffs strongly indicates the producing microorganism to be *P. thymicola* and not one of the closely related ochratoxin A-producing species *P. verrucosum* and *P. nordicum*.

## **Experimental Section**

**General Experimental Procedures.** NMR spectra of daldinin D (2) were recorded in  $CDCl_3$  on a Varian 400 FT-NMR spectrometer at 400.0 and 100.6 MHz for <sup>13</sup>C and <sup>1</sup>H NMR spectra, respectively. NMR spectra of serantrypinone (1) were recorded in 5 mm tubes at 600.13 MHz for <sup>1</sup>H and at 150.92 MHz for <sup>13</sup>C and at 300 K, using DMSO- $d_6$ , on a Bruker DRX 600 according to Larsen et al.<sup>6</sup> The chemical shifts are given relative to DMSO, 2.50 ppm for H<sup>1</sup> and 39.5 ppm for <sup>13</sup>C. EIMS originate from a JEOL JMS-HX/HX110A tandem mass spectrometer. The circular dichroism (CD) spectra were measured on a JASCO J-710 spectropolarimeter and the UV spectra on a Hewlett-Packard 8452A diode array spectrophotometer. Analytical HPLC conditions were similar to those given by Smedsgaard,<sup>7</sup> and retention indices (RI) of fungal metabolites were calculated according to Frisvad and Thrane.<sup>8</sup>

The 3D structure of daldinin D (Figure 1) was drawn using CS Chem 3D Pro, Molecular Modeling and Analysis, Version 3.2, CambridgeSoft Corporation, MA.

**Fungal Material and Fermentation.** The *Penicillium* isolate (IBT 5891) was obtained from the IBT Culture Collection at BioCentrum-DTU, Technical University of Denmark. The fungus was cultured for 14 days in 10 conical flasks (1 L), each containing 200 mL of SYES liquid medium according to Svendsen and Frisvad,<sup>9</sup> however, without agar.

Extraction and Separation. The combined fungal mycelia were extracted twice for 16 h at room temperature with 300 mL of EtOAc and filtered through a Whatman 1PS phase separation filter before evaporation to give approximately 3.5 g of crude extract. This extract was subjected to vacuum-liquid chromatography on silica gel<sup>10</sup> to give a fraction rich in serantrypinone (1) and one rich in daldinin D (2). The fraction (420 mg) rich in 1 was further purified on a Merck Lichroprep RP-18 (25 × 310 mm, 40–63  $\mu$ m) column (H<sub>2</sub>O–MeOH, 50: 50, 20 mL/min), giving five fractions. The first fraction (42 mg) was purified on a Waters Prep Nova-Pak Porasil cartridge (8  $\times$  100 mm, 6  $\mu$ m, 60 Å) using 2 mL/min H<sub>2</sub>O-MeOH (70:30 to 30:70 in 20 min) as mobile phase to give 6 mg of pure 1. The fraction (520 mg) rich in 2 was purified on a Waters Prep Nova-Pak Porasil cartridge (25  $\times$  100 mm, 6  $\mu$ m, 60 Å) using 20 mL/min H<sub>2</sub>O-CH<sub>3</sub>CN (50:50) as mobile phase to give 41 mg of pure 2.

**Serantrypinone (1):**  $[\alpha]^{22}_{D} - 12^{\circ}$  (*c* 0.12, EtOH); UV  $\lambda_{max}$ (EtOH) nm (log ε) 218 (4.55), 267 (3.54), 278 (3.51), 290 (3.36) 305 (3.16), 318 (3.05); CD (EtOH, c 0.08),  $\Delta \epsilon$  ( $\lambda$  nm) 235 (-1.54), 246 (+0.91), 274 (-0.97), 295 (+0.35), 318 (-0.14); <sup>1</sup>H NMR  $\delta$ 8.22 (1H, br d, J = 8.0 Hz, H-10), 7.89 (1H, br t, J = 7.5 Hz, H-8), 7.75 (1H, br d, J = 8.0 Hz, H-7), 7.61 (1H, br t, J = 7.5 Hz, H-9), 7.29 (1H, br t, J = 7.5 Hz, H-22), 7.22 (1H, br d, J =7.5 Hz, H-24), 7.07 (1H, br t, J = 7.5 Hz, H-23), 6.89 (1H, br d, J = 8.0 Hz, H-21), 5.55 (1H, dd, J = 3.4, 2.0 Hz, H-14), 3.71 (1H, d, J = 11.7 Hz, H-16a), 3.57 (1H, d, J = 11.7 Hz, H-16b), 2.41 (1H, dd, J = 14.3, 3.4 Hz, H-15a), 2.36 (1H, dd, J = 14.3, 2.0 Hz, H-15b);  $^{13}\mathrm{C}$  NMR  $\delta$  176.8 (C-18), 169.6 (C-1), 158.3 (C-12), 152.0 (C-4), 146.5 (C-6), 142.4 (C-20), 134.5 (C-8), 129.6 (C-25), 128.9 (C-22), 127.6 (C-7), 127.1 (C-9), 126.1 (C-10), 123.6 (C-24), 121.8 (C-23), 120.1 (C-11), 109.6 (C-21), 64.9 3), 57.6 (C-16), 52.5 (C-17), 51.9 (C-14), 37.3 (C-15). The following HMBC correlations were observed: from C-1 to H-14 and H-15; from C-3 to H-16; from C-4 to H-14 and H-16; from C-6 to H-8 and H-10; from C-7 to H-9; from C-8 to H-10; from C-9 to H-7; from C-10 to H-8; from C-11 to H-7 and H-9; from C-12 to H-10; from C-17 to H-14, H-15, H-16, and H-24; from C-18 to H-15; from C-20 to H-22 and H-24; from C-21 to H-23; from C-22 to H-24; from C-23 to H-21; from C-24 to H-22; and from C-25 to H-15, H-21, and H-23; HREIMS m/z 388.1172  $(-0.3 \text{ mmu calcd for } C_{21}H_{16}O_4N_4); \text{ RI} = 637.^8$ 

**Daldinin D (2):** mp 154–156 °C;  $[\alpha]^{22}_{D}$  –19° (*c* 0.01, EtOH); UV  $\lambda_{max}$  (EtOH) nm (log  $\epsilon$ ) 305 (4.01), 375 sh (3.13), 455 (3.19); CD (EtOH, c 0.04),  $\Delta \epsilon$  ( $\lambda$  nm) 218 (-9.95), 302 (+8.09), 346 (-1.99), 379 (+1.62); <sup>1</sup>H NMR  $\delta$  7.62 (1H, d, J= 1.5 Hz, H-1), 6.13 (1H, d, J = 1.5 Hz, H-5), 5.74 (1H, s, H-4), 5.13 (1H, br t, J = 2.8 Hz, H-9), 3.83 (1H, ddq, J = 11.0, 2.7, 6.4 Hz, H-12), 2.14 (3H, s, CH<sub>3</sub>CO<sub>2</sub>-7), 2.12 (1H, m, H-10), 2.08 (3H, s, CH<sub>3</sub>CO<sub>2</sub>-9), 1.99 (3H, s, CH<sub>3</sub>CO<sub>2</sub>-4), 1.90 (1H, m, H-10), 1.65 (1H, m, H-11), 1.55 (1H, m, H-11), 1.50 (1H, s, H-14), 1.12 (1H, d, J = 6.4 Hz, H-13); <sup>13</sup>C NMR  $\delta$  193.8 (C-6), 191.8 (C-8), 169.6 (CH<sub>3</sub>CO<sub>2</sub>-9), 169.5 (CH<sub>3</sub>CO<sub>2</sub>-4), 169.0 (CH<sub>3</sub>CO<sub>2</sub>-7), 153.9 (C-1), 141.5 (C-4a), 122.1 (C-5), 110.6 (C-8a), 100.9 (C-3), 84.8 (C-7), 69.0 (C-12), 65.3 (C-4), 63.4 (C-9), 26.2 (C-11), 23.4 (C-10), 22.1 (C-14), 21.1 (C-13), 20.8 (CH<sub>3</sub>CO<sub>2</sub>-9), 20.6 (CH<sub>3</sub>CO<sub>2</sub>-4), 19.9 (*C*H<sub>3</sub>CO<sub>2</sub>-7). The following HMBC correlations were observed: from C-3 to H-1, H-4, H-9, and 13-CH<sub>3</sub>; from C-4 to CH<sub>3</sub>CO<sub>2</sub>-4 and H-5; from CH<sub>3</sub>CO<sub>2</sub>-4 to H-4 and CH<sub>3</sub>CO<sub>2</sub>-4; from C-4a to H-1 and H-4; from C-5 to H-4; from C-6 to CH<sub>3</sub>-14; from C-7 to H-5 and CH<sub>3</sub>CO<sub>2</sub>-7, CH<sub>3</sub>-14; from CH<sub>3</sub>CO<sub>2</sub>-7 to CH<sub>3</sub>CO<sub>2</sub>-7; from C-8 to H-1, H-4, and CH<sub>3</sub>-14; from C-8a to H-1, H-4, and H-5; from C-9 to CH<sub>3</sub>CO<sub>2</sub>-9; from CH<sub>3</sub>CO<sub>2</sub>-9 to H-9 and CH<sub>3</sub>CO<sub>2</sub>-9; from C-11 to H-9 and CH<sub>3</sub>-13; from C-12 to CH<sub>3</sub>-13; from C-13 to CH<sub>2</sub>-11; HREIMS m/z 436.1369 (-0.4 mmu calcd for  $C_{21}H_{24}O_{10}$ ; RI = 1016.8

Daldinin C (3):  $[\alpha] {}^{22}{}_{D} - 34^{\circ}$  (*c* 0.01, EtOH); CD (EtOH, *c* 0.04),  $\Delta \epsilon$  ( $\lambda$  nm) 238 (-9.79), 308 (+10.74), 350 (-1.25), 387 (+0.55).

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