Solanapyrone Analogues from a Hawaiian Fungicolous Fungus

Lori E. Schmidt,[†] James B. Gloer,^{*,†} and Donald T. Wicklow[‡]

Department of Chemistry, University of Iowa, Iowa City, Iowa 52242, and Mycotoxin Research Unit, Agricultural Research Service, USDA National Center for Agricultural Utilization Research, USDA, Peoria, Illinois 61604

Received May 30, 2007

Four new solanapyrone analogues (solanapyrones J–M; 1-4) have been isolated from an unidentified fungicolous fungus collected in Hawaii. The structures and relative configurations of these compounds were determined by analysis of 1D NMR, 2D NMR, and MS data. Solanapyrone J (1) showed antifungal activity against *Aspergillus flavus* and *Fusarium verticillioides*, while both 1 and 2 showed activity against *Staphylococcus aureus* and *Candida albicans*.

Our studies of fungicolous and mycoparasitic fungi have resulted in the discovery of many new bioactive natural products.^{1,2} Recently, a collection of such fungi obtained from the island of Hawaii have begun to afford similar results.^{3,4} In the course of this work, chemical investigation of an unidentified Hawaiian fungicolous isolate (MYC-1710) led to the isolation of four new antimicrobial compounds (1–4) that were identified as new analogues of the solanapyrones.^{5–10} Solanapyrones have been previously reported from the fungi *Alternaria solani*^{5–7} and *Ascochyta rabiei*,⁸ as well as an unidentified filamentous fungus obtained from the surface of a marine alga.⁹ Members of this group have been found to show phytotoxicity^{7–9,11} and inhibition of DNA polymerases β and λ .¹² Details of the isolation and structure elucidation of compounds 1–4 are presented here.

Results and Discussion

MYC-1710 was obtained from the surface of a black stroma of an unidentified pyrenomycete on a dead hardwood branch collected in a Hawaiian forest. The fungus was cultured by solid-substrate fermentation on rice, and the EtOAc extract of the resulting fermentation mixture showed antifungal activity against *Aspergillus flavus* and *Fusarium verticillioides* and was therefore subjected to chemical investigation, leading to the isolation of compounds 1-4.

The most abundant component (1) was found to have the molecular formula $C_{20}H_{26}O_4$ (eight unsaturations) on the basis of MS and NMR data. In the ¹H NMR spectrum (Table 1), signals were observed for an aldehyde group, an isolated olefinic or aromatic proton, and two vicinally coupled olefinic protons. One methoxy resonance, three other methyl signals (two doublets and one singlet), and multiple sp³ methine and methylene signals were also present. In addition to the aldehyde signal, the ¹³C NMR spectrum (Table 2) revealed three oxygenated olefinic or carbonyl carbon signals and two upfield-shifted sp² carbon signals suggestive of a β -oxygenated α -pyrone unit,⁵⁻¹⁰ along with two olefinic signals corresponding to the 1,2-disubstituted double bond, the methoxy group, and 11 other aliphatic carbons. These units account for six degrees of unsaturation, requiring two additional rings to be present.

The presence and substitution pattern of the α -pyrone unit was confirmed by HMBC data and by comparison of shift data with those of solanapyrone A.⁷ HMBC correlations from the methoxy signal to C-13, and from the isolated olefinic proton H-12 to C-11, C-13, and C-14, placed the upfield-shifted C-12 between the downfield-shifted C-11 and C-13. Correlations from the aldehyde proton H-17 to C-13, C-14, and C-15 allowed location of the aldehyde group adjacent to the pyrone carbonyl (at C-14). Cor-

Table 1. ¹H NMR Data [$\delta_{\rm H}$ (mult, $J_{\rm H}$)] for Compounds 1 and 4

position	1 ^{<i>a</i>}	4^{b}
2	2.83 (br m)	1.87 (m)
3	5.32 (dt, 9.6, 2.0)	5.51 (ddd, 10, 4.5, 2.5)
4	5.62 (dt 9.6, 3.6)	5.36 (br d, 10)
5	2.43 (br m)	1.79 (br m)
6 _{eq}	1.60 (br d, 13)	1.81 (m)
6 _{ax}	1.23 (m)	1.25 (m)
7_{eq}	1.50 (m)	1.76 (m)
7 _{ax}	1.48 (m)	1.41 (qt, 13, 3.8)
8 _{eq}	1.38 (br d, 12)	1.63 (br d, 13)
8 _{ax}	1.50 (m)	1.32 (dq, 4.3, 13)
9	1.51 (m)	1.40 (m)
10	2.30 (br d, 7.2)	1.94 (t, 9.9)
12	6.20 (s)	6.20 (s)
16	0.76 (d, 7.2)	0.85 (d, 6.9)
17	10.14 (s)	4.54 (s)
18	0.99 (d, 7.4)	0.62 (d, 6.7)
19	1.17 (s)	1.28 (s)
MeO	4.07 (s)	3.90 (s)

^a CDCl₃,400 MHz. ^bCDCl₃, 600 MHz.

Table 2. ¹³C NMR Data (δ_C) for Compounds 1–4

position	1 ^{<i>a</i>}	$2^{a,b}$	$3^{a,b}$	4 ^{<i>a</i>}
1	49.0	47.8	47.6	46.8
2	39.1	38.4	38.6	48.2
3	128.7	129.8	129.6	132.4
4	131.0	131.6	131.4	132.1
5	32.9	33.4	33.2	43.1
6	29.4	29.5	29.3	36.8
7	20.2	20.9	20.6	29.7
8	30.6	31.2	30.4	40.2
9	29.6	30.0	29.8	47.6
10	44.6	44.2	44.2	47.2
11	180.0	174.4	174.5	177.0
12	95.1	109.5	109.4	94.9
13	173.7	185.1	184.3	169.7
14	101.8	97.8	98.7	106.3
15	162.1	165.6	165.3	167.0
16	16.1	16.7	16.6	22.5
17	186.9	163.3	161.8	57.4
18	22.5	23.1	22.9	23.3
19	13.9	14.8	14.6	21.5
MeO	57.8			59.0
20		53.3		
21		62.1		

^{*a*} CDCl₃, 400 MHz. ^{*b*}Shifts shown for **2** and **3** correspond to the signals for the major isomeric form in solution in each case.

relations from H-12 to C-1 and from H_3 -19 to C-11 enabled connection of the pyrone ring of 1 to the remainder of the molecule at C-11.

The remaining units in the molecule were linked to form a modified decalin system by analysis of HMBC and ¹H NMR

10.1021/np070251m CCC: \$37.00 © 2007 American Chemical Society and American Society of Pharmacognosy Published on Web 07/31/2007

^{*} To whom correspondence should be addressed. Tel: 319-335-1361. Fax: 319-335-1270. E-mail: james-gloer@uiowa.edu.

[†] University of Iowa.

[‡] USDA National Center for Agricultural Utilization Research.



coupling data. The three methyl signals for H_3 -16, H_3 -18, and H_3 -19 all clearly showed the expected two- and three-bond HMBC correlations enabling construction of a sizable portion of the molecule that accounted for all of structure **1** except for the C-5–C-6–C-7 unit. Coupling of H-5 to both olefinic protons H-3 and H-4, as well as to H-10, enabled connection of C-5 to C-4 and C-10 to complete a cyclohexene ring. The two remaining methylene units (C-6 and C-7) must bridge C-5 and C-8 to complete the structure of **1**. These conclusions were supported by additional HMBC data. Structure **1** differs from that of solanapyrone A in the presence of two additional methyl groups attached at positions C-1 and C-9. The name solanapyrone J is proposed for compound **1**, and the numbering system shown is consistent with that of previously reported solanapyrones.^{5–10}

Analysis of NOESY data (Figure 1) and ¹H NMR J values enabled assignment of the relative configuration of 1. A correlation between H-2 and H-10 required these two protons to be cis to one another and pseudo-diaxial with respect to the cyclohexene ring. Similarly, a correlation between H-5 and H₃-18 places these protons on the other face of the fused ring system and 1,3-diaxial (or pseudodiaxial) with respect to the cyclohexane ring. The resulting cis ring fusion was consistent with observation of a NOESY correlation between H-5 and H-10. Further correlations of H₃-19 with H₃-16 and of H-12 with H-2 led to assignment of the remaining configuration at C-1 as shown. Other NOESY correlations were also consistent with this overall stereochemical assignment. Most of the vicinal coupling values for the cyclohexane portion of the molecule could not be discerned due to broadness and/or overlap of the signals, although the $J_{\rm H5-H10}$ value was observed (7.2 Hz). This value is a bit larger than expected for a standard cis ring fusion relationship, but was clearly smaller than that of the trans value observed for 4 (9.9 Hz; see below). The conformation adopted by 1 (and therefore the H5-H10 vicinal angle) would likely be influenced to some degree by the nearby substituents at C-1 and





Figure 1. Selected NOESY correlations for solanapyrone J (1) and solanapyrone M (4).

Table 3. ¹H NMR Data [$\delta_{\rm H}$ (mult, $J_{\rm H}$)] for Compound 2^a

position		position	
2	2.79 (br m)	12 _{minor}	5.91 (s)
3	5.29 (dt, 10, 2.0)	16	0.76 (d, 7.4)
4	5.57 (dt, 10, 3.4)	17 _{major}	8.27 (d, 13)
5	2.39 (br m)	$17_{\rm minor}$	8.42 (d, 15)
6 _{eq}	1.72 (m)	18	0.98 (d, 7.4)
6 _{ax}	1.23 (m)	19	1.04 (s)
7_{eq}	1.50 (m)	20 _{major}	3.60 (q, 5.3)
7 _{ax}	1.50 (m)	20 _{minor}	3.66 (q, 5.3)
8 _{eq}	1.34 (br d, 12)	21	3.86 (br t, 5.3)
8 _{ax}	1.50 (m)	NH _{major}	11.88 (br s)
9	1.51 (m)	NH _{minor}	10.16 (br s)
10	2.21 (br d, 7.2)	OH _{major}	2.00 (br s)
12 _{major}	5.86 (s)	OHminor	2.03 (br s)

^{*a*} CDCl₃, 300 MHz. ¹H NMR signals for the major and minor forms were not clearly resolved for the modified decalin portion of the molecule or for H₂-21, but some of the corresponding signals did appear to be broadened.

C-9, as well as the presence of the olefin, and would have more flexibility than a *trans*-fused analogue (e.g., **4**). Molecular modeling (ChemDraw Pro 9.0) suggests that the lowest energy conformation of the cyclohexane ring in **1** may be a twisted chair, resulting in a smaller H5–H10 vicinal angle (and a somewhat larger *J* value) than would be expected for a simple decalin system. On the basis of the NOESY data, solanapyrone J (**1**) was assigned the relative configuration shown, which matched that reported for solanapyrone A at relevant stereocenters C-2, C-5, and C-10.⁷ The orientation of the pyrone substituent also matches that of the other known solanapyrones.

Compound **2** was assigned the molecular formula $C_{21}H_{29}O_4N$ (eight unsaturations) on the basis of HRESIMS and NMR data. However, the ¹³C and ¹H NMR spectra (Tables 2 and 3, respectively) each showed the presence of two distinct sets of signals. Since the two constituents could not be separated, analysis was carried out on the mixture, with a focus on the major component. The ¹H and ¹³C NMR data for the major component lacked the aldehyde and methoxy signals observed in the data for **1**, but included additional resonances for a ketone group, an NH proton, two more methylene units, and an OH group. The latter signals were suggestive of an ethanolamine unit. All of the ¹H NMR δ and J values for the modified decalin system are very similar to those of 1, suggesting that this portion of the molecule is identical to the corresponding portion of 1 (including relative configuration), and the connectivity of this subunit was independently confirmed by analysis of HMBC data. Other key correlations were observed from H-12 to C-1, C-11, and nonprotonated olefinic carbon C-14, and from olefin proton H-17 to C-13, C-14, and C-20 (of the ethanolamine unit), enabling attachment of the C20-C21 ethanolamine unit at C-17 to form a cross-conjugated enamine. This structure rationalized the appearance of two sets of signals for the upper portion of the molecule due to the two possible intramolecularly hydrogen-bonded forms it can adopt. These two forms are geometric isomers of each other, but are inseparable and presumably in equilibrium via a process involving keto-enol tautomerism. The large coupling constants observed for the H-17 signals for each isomer result from hydrogen bonding of the NH proton to the carbonyl oxygen atoms at position 13 or 15, which requires a near-180° vicinal angle for the NH and H-17 protons in each of the two possible forms. Interestingly, an ethanolamine unit is also present in solanapyrone C,8 but is attached at C-13, rather than C-17, and therefore does not show this property. However, several synthetic systems similar to the upper portion of 2 have been studied by Uray et al.¹³ As in the case of 2, these researchers found that two sets of signals are observed for such enamines, and in each set, a large J_{CH-NH} value is evident. In all of the compounds surveyed, the CH signal has a slightly more downfield shift in the isomer with hydrogen bonding of the NH to the ester oxygen. In addition, as in the data for 2, this signal is consistently less intense and has a slightly larger J value than its counterpart. Thus, the signals for the two forms of 2 were assigned as shown by analogy to these synthetic model compounds. The name solanapyrone K is proposed for compound 2.

The molecular formula for compound **3** was determined to be $C_{19}H_{25}O_3N$ (eight unsaturations) on the basis of HRESIMS and NMR data. Analysis of NMR data revealed that **3** is related to **1** and **2**, but lacks the two methylene signals for the ethanolamine unit in **2**, and instead shows a second NH signal in the ¹H NMR spectrum. All of the other ¹H NMR δ and *J* values are nearly identical to those of **2**, suggesting that **3** lacks the CH₂CH₂OH unit, but is otherwise identical to **2** and is proposed to have the same relative configuration. The name solanapyrone L is proposed for compound **3**.

Compound 4 was determined to have the molecular formula $C_{20}H_{28}O_4$ (seven unsaturations) on the basis of HRESIMS and NMR data. The ¹H NMR spectrum of **4** (Table 1) is very similar to that of 1 except that the aldehyde signal is replaced by a two-proton singlet at δ 4.54 characteristic of an oxymethylene unit. HMBC correlations confirmed that the connectivity of 4 matches that of 1 except for replacement of the aldehyde with a CH₂OH unit. In addition, however, the chemical shifts and J values for several of the signals for the decalin-type portion of the molecule (Table 1) were significantly different from the corresponding signals in the ¹H NMR spectrum of **1**, suggesting that **4** possesses a different relative configuration. Analysis of NOESY data (Figure 1) and J values confirmed this suggestion and established the relative configuration of 4. A correlation between H₃-16 and H-10 required both CH₃-16 and H-10 to be in pseudoaxial orientations and cis to one another, in contrast to the situation in 1. Additional correlations of H-10 with H-6ax and H-8ax supported assignment of an axial orientation for H-10, as did large *trans*-diaxial J_{H5-H10} and J_{H9-H10} values. The large vicinal couplings (9.9 Hz each) observed between H-5 and H-10 and between H-5 and H-6ax also indicated an axial orientation for H-5 and established the presence of a trans ring fusion. Other NOESY correlations were consistent with these

assignments. These data permitted assignment of the relative configuration of **4** as shown, matching those reported for solanapyrones D and E at positions C-2, C-5, and C- 10^7 and differing from **1** at the C-2 and C-5 positions. Chemical shift differences between **1** and **4** at affected positions are consistent with similar variations observed among previously reported solanapyrones exhibiting similar stereochemical differences. The name solanapyrone M is proposed for **4**.

The occurrence of solanapyrones having these two types of relative configurations has been proposed to be due to formation of the modified decalin system via an intramolecular Diels-Alder reaction of a polyketide-derived pro-solanapyrone species to give *exo* (e.g., 1-3) or *endo* (e.g., 4) products.^{14,15} The absolute configurations of 1-4 were not independently assigned, but are presumed to match those of the previously known solanapyrones. Like all other solanapyrones for which values have been previously reported, 1-4 all display negative optical rotations, although this alone does not enable an unambiguous stereochemical assignment.

Solanapyrone J (1) showed antifungal activity against Aspergillus flavus (NRRL 6541) and Fusarium verticillioides (NRRL 25457), affording clear zones of ca. 30 mm diameter in disk assays against both organisms at 200 μ g/disk that persisted for 4 days. Upon further evaluation,¹⁶ compound **1** displayed MIC values of approximately 20 and 24 µg/mL against A. flavus and F. verticillioides, respectively, and IC₅₀ values of approximately 8.6 and 15 μ g/mL. Nystatin, an antifungal standard, afforded MIC values of approximately 10 μ g/mL in both assays and IC₅₀ values of 5 and 3 μ g/mL, respectively, using the same protocols. Compounds 2-4 were much less abundant than 1 and were not tested in these assays due to sample limitations. Compounds 1 and 2 showed antibacterial activity in standard disk assays^{17,18} against Staphylococcus aureus (ATCC 25923), with 1 affording a clear zone of inhibition of 17 mm and a zone of reduced growth of 21 mm and 2 affording a zone of inhibition of 9 mm at 50 μ g/disk after 4 days (a gentamicin sulfate standard gave a 25-mm clear zone in this assay at 25 μ g/disk). Compounds 1 and 2 were also active against Candida albicans (ATCC 14053), affording zones of inhibition of 19 and 13 mm, respectively, under the same conditions (a filipin standard gave a 23-mm clear zone in the assay at 50 μ g/disk). Compounds 3 and 4 showed no activity in either of these assays, and none of the compounds showed activity against Escherichia coli (ATCC 25922) when tested at the same level.

Experimental Section

General Experimental Procedures. ¹H and ¹³C NMR spectra were measured on Bruker AVANCE-300, DRX-400, and AVANCE-600 spectrometers. Chemical shift values were referenced to residual solvent signals for CDCl₃ ($\delta_{\rm H}/\delta_{\rm C}$, 7.24/77.0). HMQC and HMBC data were recorded on a Bruker AVANCE-600. HPLC was carried out using a Beckman System Gold HPLC instrument with a model 168 variablewavelength UV detector. HRESIMS data were recorded on a Thermo LTQ Fourier transform instrument (Washington University Mass Spectrometry Resource). Optical rotations were measured with a Rudolph automatic polarimeter, model AP III. Standards of nystatin, gentamicin sulfate, and filipin were purchased from Sigma Chemical Co.

Isolation, Cultivation, and Fermentation of Fungal Material. An isolate (MYC-1710) was obtained from the surface of a black stroma of an unidentified pyrenomycete on a dead hardwood branch collected in a montane dry forest (Ohi'a) Koloko St., Kailua-Kona, Hawaii Co., HI, on November 3, 2002. The fungus was grown on 100 g of autoclaved rice for 30 days at 25 °C. The EtOAc extract (448 mg) of the resulting fermentation mixture showed antifungal activity against *A. flavus* and *F. verticillioides*. The isolate was not taxonomically identified, as scale-up attempts revealed that the cultures and backups thereof were no longer viable, despite storage of cultures at both -20 and 5 °C. Unfortunately, efforts to extract DNA fragment that might have enabled identification or partial classification of the fungus. Efforts

to reisolate the fungus from a sample of the original substrate stromata, and from other stromata collected at the same location, all of which were continuously stored at 5 $^{\circ}$ C since collection, have thus far been unsuccessful.

Extraction and Isolation. The extract was partitioned between MeCN and hexanes (3 mL of each). The resulting MeCN fraction (360 mg) was then chromatographed on a silica gel column using a hexanes/ CH₂Cl₂/MeOH step gradient (hexanes, CH₂Cl₂, CH₂Cl₂/MeOH, 19:1, 9:1, 17:3, 4:1, 7:3, 1:1, and 1:4) to give 11 fractions. Fraction 3 (84 mg), eluted with CH₂Cl₂, was further separated by reversed-phase HPLC (50% MeCN/H₂O isocratic for 5 min, 50-100% over 35 min, and 100% for 10 min) on a HS Hyperprep BDS 8- μ m C₁₈ column (4.6 × 250 mm) at a flow rate of 2 mL/min with UV detection at 215 nm to afford solanapyrone J (1; 30 mg, $t_{\rm R}$ 28.8 min). Fraction 5 (149 mg), eluted with 19:1 CH₂Cl₂/MeOH, was separated by reversed-phase HPLC (same column as above; 30% MeCN/H2O isocratic for 5 min, 30-60% over 25 min, 60% isocratic for 20 min, and 60-100% over 10 min) to afford an additional sample of solanapyrone J (1; 8 mg, $t_{\rm R}$ 43.6 min), as well as two other subfractions, which were further purified. Subfraction 5 (3 mg) was purified by reversed-phase HPLC (65-72% MeCN/H₂O over 35 min and 72-100% over 1 min) on the same column with UV detection at 215 nm to afford solanapyrone K (2; 2 mg, t_R 25.0 min) and solanapyrone L (4; 1 mg, t_R 26.5 min). Subfraction 7 (2 mg) was further purified by reversed-phase HPLC (55% MeCN/H2O isocratic for 30 min and 55-100% over 1 min) on the same column with UV detection at 215 nm to afford solanapyrone M (3; 1 mg, $t_{\rm R}$ 26.0 min).

Solanapyrone J (1): yellow glass; $[\alpha]^{25}_{D} - 114$ (*c* 1.3, MeOH); ¹H and ¹³C NMR data, see Tables 1 and 2; HMBC data, H-2 \rightarrow C-1, 3, 4, 16; H-3 \rightarrow C-1, 2, 5, 16; H-4 \rightarrow C-2, 5, 6, 10; H-5 \rightarrow C-1, 3, 4, 6, 7, 10; H₂-6 \rightarrow C-4, 5, 7, 8, 10; H₂-7 \rightarrow C-5, 6, 8, 9; H₂-8 \rightarrow C-6, 7, 9, 10, 18; H-9 \rightarrow C-1, 7, 8, 18; H-10 \rightarrow C-1, 9, 18, 19; H-12 \rightarrow C-1, 11, 13, 14; H₃-16 \rightarrow C-1, 2, 3; H-17 \rightarrow C-13, 14, 15; H₃-18 \rightarrow C-8, 9, 10; H₃-19 \rightarrow C-1, 2, 10, 11; MeO \rightarrow C-13; HRESIMS obsd *m*/*z* 331.1902, calcd for C₂₀H₂₇O₄ (M + H)⁺, 331.1910.

Solanapyrone K (2): colorless oil; $[\alpha]^{25}_{D} - 45$ (*c* 0.12, MeOH); ¹H and ¹³C NMR data, see Tables 2 and 3; HMBC data, H-2 \rightarrow C-10, 16; H-3 \rightarrow C-1, 2, 5, 16; H-4 \rightarrow C-5; H-5 \rightarrow C-1; H₂-6 \rightarrow C-5; H₂-7 \rightarrow C-5, 9; H₂-8 \rightarrow C-7; H-9 \rightarrow C-5, 18; H-10 \rightarrow C-1, 5; H-12 \rightarrow C-1, 11, 14; H₃-16 \rightarrow C-1, 2, 3; H-17 \rightarrow C-13, 14, 15, 20; H₃-18 \rightarrow C-8, 9, 10; H₃-19 \rightarrow C-1, 2, 10, 11; H₂-21 \rightarrow C-20; HRESIMS obsd *m*/*z* 360.2166, calcd for C₂₁H₃₀O₄N (M + H)⁺, 360.2176.

Solanapyrone L (3): colorless oil; $[\alpha]^{25}_{D} -53$ (*c* 0.04, MeOH); ^{13}C NMR data, see Table 2; ¹H NMR (400 MHz, CDCl₃) δ 11.2 (br s, NH_{major}), 9.57 (br s, NH_{minor}), 8.46 (dd, 16, 8.7 Hz, H-17_{minor}), 8.33 (dd, 15 Hz, 8.9 Hz, H-17_{major}), 6.70 (br s, NH), 5.91 (s, H-12_{minor}), 5.86 (s, H-12_{major}), 5.58 (dt, 10, 3.5 Hz, H-4), 5.30 (dt, 10, 2.0 Hz, H-3), 2.89 (br m, H-2), 2.40 (br m, H-5), 2.21 (br d, 7.3 Hz, H-10), 1.73 (m, H-6_{eq}), 1.52 (m, H-9), 1.50 (m, 2H, H-7_{eq}, H-8_{ax}), 1.48 (m, H-7_{ax}), 1.34 (br d, 12 Hz, H-8_{eq}), 1.23 (m, H-6_{ax}), 1.04 (s, H₃-19), 0.99 (d, 7.3 Hz, H₃-18), 0.77 (d, 7.4 Hz, H₃-16); HRESIMS obsd *m*/z 316.1904, calcd for C₁₉H₂₆O₃N (M + H)⁺, 316.1914.

Solanapyrone M (4): colorless oil; $[\alpha]^{25}_{D} - 84$ (*c* 0.025, MeOH); ¹H and ¹³C NMR data, see Tables 1 and 2; HMBC data, H-2 \rightarrow C-1, 3, 16; H-3 \rightarrow C-2, 4; H-4 \rightarrow C-2, 3, 5, 6; H₂-6 \rightarrow C-5; H₂-7 \rightarrow C-6; H-9 → C-18; H-10 → C-1, 5, 6, 8, 9, 19; H-12 → C-1, 11, 13, 14; H₃-16 → C-2, 3; H-17 → C-13, 14, 15; H₃-18 → C-8, 9, 10; H₃-19 → C-1, 2, 10, 11; MeO → C-13; HRESIMS obsd *m*/*z* 333.2060, calcd for C₂₀H₂₉O₄ (M + H)⁺, 333.2067.

Acknowledgment. This research was supported by grants from the National Science Foundation (#CHE-0315591) and the National Institutes of Health (GM 60600). Assistance from the staff of the University of Iowa NMR and Mass Spectrometry Facilities is gratefully acknowledged. High-resolution MS data were provided by the Washington University Mass Spectrometry Facility, an NIH Research Resource (grant #P41RR0954). We are also grateful to Prof. D. Hemmes of the University of Hawaii at Hilo for guiding the collection of fungal sporocarps from Hawaii.

Supporting Information Available: ¹H NMR spectra for compounds **1**–**4** and ¹³C NMR spectra for compounds **1** and **2**. This material is available free of charge on the Internet at http://pubs.acs.org.

References and Notes

- Mudur, S. V.; Gloer, J. B.; Wicklow, D. T. J. Antibiot. 2006, 59, 500-506.
- (2) Deyrup, S. T.; Swenson, D. C.; Gloer, J. B.; Wicklow, D. T. J. Nat. Prod. 2006, 69, 608-611.
- (3) Shim, S. H.; Swenson, D. C.; Gloer, J. B.; Dowd, P. F.; Wicklow, D. T. Org. Lett. 2006, 8, 1225–1228.
- (4) Shim, S. H.; Gloer, J. B.; Wicklow, D. T. J. Nat. Prod. 2006, 69, 1601–1605.
- (5) Ichihara, A.; Tazaki, H.; Sakamura, S. *Tetrahedron Lett.* 1983, 24, 5373–5376.
- (6) Ichihara, A.; Miki, I.; Sakamura, S. Tetrahedron Lett. 1985, 26, 2453–2454.
- (7) Oikawa, H.; Yokota, T.; Sakano, C.; Suzuki, Y.; Naya, A.; Ichihara, A. Biosci. Biotechnol. Biochem. 1998, 62, 2016–2022.
- (8) Alam, S. S.; Bilton, J. M.; Slawin, M. Z.; Williams, D. J.; Sheppard, R. N.; Strange, R. M. Phytochemistry 1989, 28, 2627–2630.
- (9) Jenkins, K. M.; Toske, S. G.; Jensen, P. R.; Fenical, W. *Phytochemistry* 1998, 49, 2299–2304.
- (10) Schlörke, O. Dissertation, University of Göttingen, 2005.
- (11) Kaur, S. Plant Sci. 1995, 109, 23-29.
- (12) Mizushina, Y.; Kamisuki, S.; Kasai, N.; Shimazaki, N.; Takemura, M.; Asahara, H.; Linn, S.; Yoshida, S.; Matsukage, A.; Koiwai, O.; Sugawara, F.; Yoshida, H.; Sakaguchi, K. J. Biol. Chem. 2002, 277, 630–638.
- (13) Uray, G.; Wolfbeis, O. S.; Junek, H. J. Mol. Struct. 1979, 54, 77– 88.
- (14) Oikawa, H.; Suzuki, Y.; Naya, A.; Katayama, K.; Ichihara, A. J. Am. Chem. Soc. 1994, 116, 3605–3606.
- (15) Oikawa, H.; Katayama, K.; Suzuki, Y.; Ichihara, A. J. Chem. Soc., Chem. Commun. 1995, 1321–1322.
- (16) Jiao, P.; Swenson, D. C.; Gloer, J. B.; Campbell, J.; Shearer, C. A. J. Nat. Prod. 2006, 69, 1667–1671.
- (17) Bauer, A. W.; Kirby, W. M.; Sherris, J. C.; Turck, M. Am. J. Clin. Pathol. 1966, 45, 493–496.
- (18) Wagenaar, M. M.; Clardy, J. J. Nat. Prod. 2001, 64, 1006–1009.

NP070251M