Ophiobolin Sesterterpenoids and Pyrrolidine Alkaloids from the Sponge-Derived Fungus Aspergillus ustus

- by Hong-Bing Liu^a), RuAngelie Edrada-Ebel^b), Rainer Ebel^c), Yao Wang^d), Barbara Schulz^e), Siegfried Draeger^e), Werner E. G. Müller^f), Victor Wray^g), Wen-Han Lin*^h), and Peter Proksch*^d)
- a) Key Laboratory of Marine Drugs, Chinese Ministry of Education, School of Medicine and Pharmacy, Ocean University of China, Qingdao 266003, P. R. China
 - b) Strathclyde Institute of Pharmacy and Biomedical Sciences, University of Strathclyde, The John Arbuthnott Building, 27 Taylor Street, Glasgow G40NR, U.K.
 - c) Department of Chemistry, University of Aberdeen, Meston Building, Meston Walk, AB243UE, Aberdeen, U.K.
- ^d) Institut für Pharmazeutische Biologie und Biotechnologie, Heinrich-Heine-Universität Düsseldorf, Universitätsstrasse 1, Geb. 26.23, D-40225 Düsseldorf
 - (phone: +49-211-8114163; fax: +49-211-8111923; e-mail: proksch@uni-duesseldorf.de)
 - c) Institut für Mikrobiologie, Technische Universität Carolo-Wilhelmina zu Braunschweig, Spielmannstrasse 7, D-38106 Braunschweig
 - f) Institut für Physiologische Chemie und Pathobiochemie, Johannes-Gutenberg-Universität, Duesbergweg 6, D-55128 Mainz
 - g) Helmholtz Center for Infection Research, Inhoffenstrasse 7, D-38124 Braunschweig
 - ^h) State Key Laboratory of Natural and Biomimetic Drugs, Peking University, Beijing 100083,
 - P. R. China (phone: +86-10-82806188; fax: +86-10-82806188; e-mail:whlin@bjmu.edu.cn)

Chemical examination of the fungus Aspergillus ustus isolated from the Mediterranean sponge Suberites domuncula yielded the five new ophiobolin-type sesterterpenoids 1–5 and the two new pyrrolidine alkaloids 6 and 7, together with the known compound aurantiamine and cerebroside D. The structures of the new compounds were unambiguously elucidated on the basis of extensive spectroscopic-data analysis (1D- and 2D-NMR, MS, and UV) and comparison with literature data. All compounds were evaluated for their cytotoxicity against murine lymphoma cell line L5178Y.

Introduction. – In the last decade, marine microorganisms, such as bacteria, microalgae, and fungi, have become increasingly attractive as sources of new bioactive natural products [1][2]. Fungi have proven to be particularly prolific sources of new compounds compared to other microbial sources isolated from the sea. Several hundred new compounds have been described from marine-derived fungi making this group of microorganisms a prime target for marine-natural-product chemists. The highest incidence for discovery of new compounds lies apparently in the group of sponge-associated fungi that have received much attention in recent years [1][3]. There is nevertheless a continuing debate on the nature of sponge—fungal interactions since many if not most of these fungi are already well-known from the terrestrial environment including genera such as *Aspergillus, Penicillium, Cladosporium, Phoma*, and *Fusariu*. Thus, a true marine origin of these fungal strains is frequently doubted [4][5]. It is indeed possible that many sponge-associated fungi originated from

terrestrial habitats (e.g., soil) from which they were washed and survived as dormant spores in filter-feeding sponges. Attempts to detect fungal mycelia growing inside sponges have usually failed. On the other hand, the sponge Suberites domuncula has been shown to have receptor proteins for fungal-cell-wall components, such as $(1 \rightarrow 3)$ - β -D-glucan, providing evidence that at some time in its life cycle (or in its evolutionary history), the sponge, if not colonized by, at least has come into contact with fungi [6].

Regardless of the true nature of these interactions, sponge-derived fungi continue to be of interest as sources for new secondary constituents. An example is provided by a marine strain of *Aspergillus ustus* that had been isolated from the Mediterranean sponge *Suberites domuncula* [7]. This fungus is know from soil samples [8], as a pathogen on higher plants [9], but also from marine habitats such as Mangrove swamps [10], and from *S. domuncula* [7] [11]. We were attracted to the below-described marine isolate of *A. ustus* mainly due to the structurally diverse terpenoids produced when cultivated *in vitro*. Previously, we have reported seven new drimane sesquiterpenoids from *A. ustus* [11]. Continued investigation now afforded five new sesterterpenes of the ophiobolin group and two new pyrrolidine alkaloids (*Fig. 1*) along with the known compounds aurantiamine [12] and cerebroside D [13]. The structure elucidation of the new compounds by one- and two-dimensional NMR and by HR-MS is described.

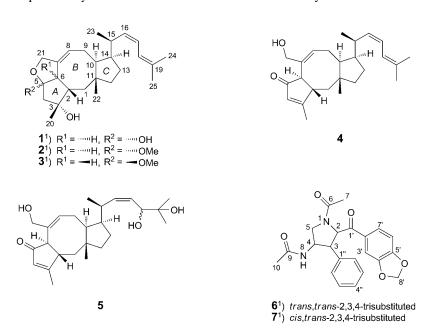


Fig. 1. Compounds 1-7 isolated from Aspergillus ustus

Results and Discussion. – The crude AcOEt extract of the fermentation broth of *A. ustus* was subjected to chromatographic separation including semi-prep. HPLC to yield

¹⁾ Trivial or arbitrary atom numbering; for systematic names, see Exper. Part.

five new ophiobolins $\mathbf{1} - \mathbf{5}^1$) and two new pyrrolidine alkaloids $\mathbf{6}^1$) and $\mathbf{7}^1$), along with the known compounds aurantiamine and cerebroside D.

Compound 1 had the molecular formula C₂₅H₃₈O₃ as determined from HR-ESI-MS $(m/z 409.2720 ([M+Na]^+))$ in association with NMR data. The ¹H-NMR spectrum (*Table 1*) exhibited four olefinic H-atoms at $\delta(H)$ 5.19 (dd, J = 9.5, 9.5 Hz, H–C(16)), 5.33 (br. s, H–C(8)), 6.03 (d, J = 11.7 Hz, H–C(18)), and 6.07 (dd, J = 11.7, 9.5 Hz, H-C(17)), four Me ss at $\delta(H)$ 0.85 (Me(22)), 1.10 (Me(20)), 1.70 (Me(24)), and 1.78 (Me(25)), one Me d at δ (H) 0.91 (d, J = 6.7 Hz, Me(23)), and two exchangeable Hatoms at $\delta(H)$ 4.09 (s, OH–C(3)) and 5.86 (s, OH–C(5)). The ¹³C-NMR spectrum (Table 2) displayed in total 25 resonances, including those of six olefinic C-atoms, one ketal C-atom ($\delta(C)$ 116.9 (C(5))) and two O-bearing C-atoms, in addition to those of five Me groups and eleven sp 3 C-atoms. The gross structure of **1** was very similar to that of the known compound ophiobolin H = (1R,2aS,6aS,7R,9aR,10aS,10bS)-7-[(1S,2Z)-1]1,5-dimethylhexa-2,4-dien-1-yl]-1,2,4,6,6a,7,8,9,9a,10,10a,10b-dodecahydro-1,9a-dimethyl-2aH-3-oxacyclopenta[5,6]cycloocta[1,2,3-cd]pentalene-1,2a-diol), previously isolated from the same fungus, based on 2D-NMR data analysis (Fig. 2) and by comparison with literature data [14][15]. However, the ROESY interactions H-C(6)/H-C(10), $H-C(10)/H_a-C(9)$, H-C(2)/Me(22), and $Me(22)/H_B-C(9)$, and the absence of a ROESY correlation H-C(6)/H-C(2) (Fig. 3), suggested the same orientation of H-C(6) and H-C(10), and of H-C(2) and Me(22), respectively. Thus, rings A/B and B/C(10)C are trans-fused. The ROESY cross-peaks OH–C(3)/H–C(6), and OH–C(5)/H–C(6) confirmed the *cis* geometry of rings A/D and furthermore indicated OH–C(5) to be spatially close to H–C(6) (Fig. 3). Thus, compound 1 was identified as the new $(5\alpha,6\alpha)$ isomer of ophiobolin H.

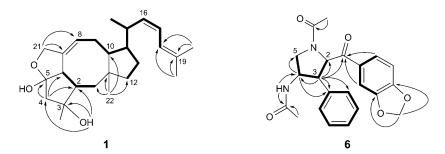


Fig. 2. Key COSY (\longrightarrow) and HMBCs (H \rightarrow C) of compound $\mathbf{1}^1$) and $\mathbf{6}^1$)

The molecular formula of **2** was determined as $C_{26}H_{40}O_3$ based on HR-ESI-MS data $(m/z \ 423.2880 \ ([M+Na]^+))$. The 1H - and ^{13}C -NMR data of **2** (*Tables I* and 2) corresponded to those of **1**, except for the presence of an additional Me group as a MeO substituent $(\delta(H)\ 3.07\ (s,3\ H);\delta(C)\ 49.0)$. The HMBC data allowed the placement of the MeO group at C(5). The ROESY signals of **2**, similar to those of **1**, indicated that **2** is $(5\alpha.6\alpha)$ -5-O-methylophiobolin H.

Table 1. ¹*H-NMR Data* (600 MHz, (D₆)DMSO) of Compounds $\mathbf{1} - \mathbf{5}^1$). δ in ppm, J in Hz^a).

	1	2	3	4	5 ^b)
H_a -C(1)	$1.46-1.50^{\circ}$	$1.45-1.50^{\circ}$)	1.31 (m)	1.07 $(t, J = 12.9)$	1.04 $(t, J = 12.9)$
	$1.46 - 1.50^{\circ}$	$1.45 - 1.50^{\circ}$)	1.44 (dd, J = 14.2, 3.2)	1.95 (dd, J = 13.1, 3.3)	1.95 (dd, J = 13.2, 3.5)
H-C(2)	1.50-1.55 (m)	1.50-1.55 (m)	2.00(m)	2.75 (br. d, J = 13.2)	2.75 (br. $d, J = 12.6$)
or H-C(4)	1.97 $(d, J = 14.1)$	2.07 (d, J = 14.2)	2.05(d, J = 13.2)	5.84 (d, J = 1.4)	5.84 (d, J = 1.3)
	1.93 $(d, J=14.1)$	1.71 $(d, J = 14.2)$	1.81 $(d, J = 13.2)$		
H-C(6)	2.91 (d, J = 9.8)		3.01 (d, J = 10.7)	3.35°)	3.35°)
	5.33 (br. s)		5.47 (d, J = 7.1)	5.48(d, J=4.7)	5.55(d, J = 4.4)
	2.15(m)		2.38 (dd, J = 13.6, 8.5)	2.32 (br. s)	2.62 (<i>m</i>)
H_{β} -C(9)	1.75 (m)		1.67 (m)	1.86 (m)	1.81 (<i>m</i>)
	2.22 (m)	2.19 (m)	1.49(m)	2.55(m)	2.50 ^d)
	1.34 (m)		1.31(m)	1.36 (m)	1.35 (m)
H_{β} -C(12)	1.39 (m)	1.38 (m)	1.31(m)	1.44 (m)	1.43 (<i>m</i>)
H_a -C(13)	1.57 (m)		1.64(m)	1.54 (m)	1.53 (m)
	$1.20 \ (m)$	1.17 (m)	1.49(m)	1.23 (m)	1.23 (<i>m</i>)
	1.82 (m)		2.02 (m)	1.85 (m)	1.81 (m)
	2.50^{d})		2.67 (m)	2.55 (m)	2.55 (<i>m</i>)
	5.19 (dd, J = 9.5, 9.5)			5.19(m)	5.27 (m)
H-C(17)	6.07 (dd, J = 11.7, 9.5)	6.06 (dd, J = 11.7, 9.5)	5.98 (dd, J = 11.3, 9.8)	6.05°)	5.23 (m)
	6.03 (d, J = 11.7)		6.01 $(d, J = 11.7)$	6.04°)	3.93 (dd, J = 8.5, 6.0)
	1.10 (s)	1.11 (s)	1.09(s)	2.00 (s)	2.00 (s)
H_a -C(21)	4.29 (d, J = 11.7)		4.52 (d, J = 11.3)	3.70 (d, J = 12.0)	3.70 (d, J = 11.0)
H_{β} —C(21)	4.37 (d, J = 11.7)		4.27 (d, J = 12.0)	3.94 (br. s)	3.99 (m)
Me(22)	0.85(s)	0.84 (s)	0.86(s)	0.96 (s)	0.97 (s)
Me(23)	0.91 (d, J = 6.7)	0.90 (d, J = 6.6)	0.82 (d, J = 6.9)	0.88(d, J = 6.6)	0.88 (d, J = 6.6)
	1.70 (s)	1.69 (s)	1.67 (s)	1.69 (s)	1.01 (s)
Me(25)	1.78 (s)	1.77(s)	1.76 (s)	1.78 (s)	1.02 (s)
	OH-C(3): $4.09(s)$	OH $-C(3)$: 4.19 (s)	OH $-C(3)$: 3.78 (s)	OH-C(21): 4.34 (br. s)	OH–C(21): $4.33 (dd, J = 9.8, 3.1)$
	OH–C(5): $5.86(s)$	MeO-C(5): 3.07 (s)	MeO-C(5): 3.08 (s)		OH–C(18): 4.31 $(d, J = 6.0)$

^a) The ¹H-NMR data were assigned by 2D-NMR (COSY, HMBC, and ROESY). ^b) The chemical shift of OH–C(19) was not assigned. ^c) Overlapping with solvent signals.

Table 2. ¹³C-NMR Data (150 MHz, (D₆)DMSO) of Compounds $\mathbf{1}-\mathbf{5}^1$). δ in ppm.

	1 ^a)	2	3	4	5
C(1)	41.8 (t)	41.7 (t)	35.7 (t)	45.4 (t)	45.6 (t)
C(2)	50.7(d)	49.8 (d)	50.0(d)	50.0(d)	50.0(d)
C(3)	80.2 (s)	80.2(s)	77.9(s)	179.0(s)	179.1 (s)
C(4)	53.6 (t)	48.0(t)	50.7(t)	129.4 (t)	129.4 (t)
C(5)	116.9(s)	119.8 (s)	119.7(s)	207.3(s)	207.5(s)
C(6)	55.2 (d)	54.3 (d)	50.9 (d)	51.6 (d)	51.7 (d)
C(7)	137.4 (s)	136.6 (s)	141.2 (s)	134.3 (s)	133.6 (s)
C(8)	116.0 (d)	120.1 (d)	120.2 (d)	127.4(d)	127.7(d)
C(9)	27.2(t)	27.5(t)	24.2(t)	28.2(t)	28.1 (t)
C(10)	41.8 (d)	41.9(d)	55.2 (d)	42.9(d)	43.1 (d)
C(11)	43.2 (s)	43.2 (s)	43.0 (s)	44.6 (s)	44.7 (s)
C(12)	45.0(t)	44.8 (t)	43.1 (t)	43.8 (t)	43.8 (t)
C(13)	27.2(t)	27.2(t)	26.0(t)	27.2(t)	27.2 (t)
C(14)	51.3 (d)	51.6(d)	46.7 (d)	51.4 (d)	51.4 (d)
C(15)	32.3 (d)	32.1 (d)	34.5 (d)	31.8(d)	31.9 (d)
C(16)	136.0 (d)	136.0 (d)	137.9(d)	136.8 (d)	138.9 (d)
C(17)	123.2 (d)	123.1 (d)	121.5 (d)	122.9 (d)	128.2 (d)
C(18)	120.1 (d)	120.1 (d)	120.3 (d)	120.2(d)	73.6 (d)
C(19)	134.9 (s)	134.9(s)	134.4 (s)	134.9(s)	71.5 (s)
C(20)	26.4 (q)	26.5 (q)	26.3 (q)	16.9(q)	16.9(q)
C(21)	b)	73.6 (t)	72.3(t)	64.2(t)	64.5 (t)
C(22)	22.8 (q)	22.7(q)	18.2 (q)	22.4(q)	22.5(q)
C(23)	21.2 (q)	21.1 (q)	19.9 (q)	21.1 (q)	20.9 (q)
C(24)	18.0 (q)	17.9 (q)	17.9 (q)	26.1 (q)	26.1 (q)
C(25)	26.1 (q)	26.1 (q)	26.1 (q)	17.9 (q)	25.1(q)
MeO-C(5)	-	$49.0 \ (q)$	49.4 (q)	-	-

^a) The ¹³C-NMR data were obtained from HMBC. ^b) The chemical shift data of C(21) of **1** was not assigned.

Compound **3** had the same molecular formula as **2** from its HR-ESI-MS data. The 1 H- and 13 C-NMR, 1 H, 1 H-COSY, HMBC, and ROESY data revealed that **3** was closely related to ophiobolin H [15] and thus differed with regard to the relative configuration at C(5) and C(6) from the core structure of **1** and **2** (*Fig. 3*). The only structural difference between **3** and ophiobolin H was the presence of a MeO group in **3** which was located at C(5) based on the HMBC cross-peak between MeO (δ (H) 3.08 (s, 3 H)) and C(5). Accordingly, compound **3** was identified as 5-O-methylophiobolin H.

Compound **4** had the molecular formula $C_{25}H_{36}O_2$ according to HR-ESI-MS $(m/z 391.2610 \ ([M+Na]^+))$ and NMR data. A comparison of its 1H - and ^{13}C -NMR data $(Tables\ 1\ and\ 2)$ with those of **1** revealed that it shared a partial structure with the latter compound. In the ^{13}C -NMR spectrum of **4**, three C-atom signals at $\delta(C)\ 179.0\ (C(3))$, 129.4 (C(4)), and 207.3 (C(5)) characteristic of an α,β -unsaturated ketone similar to ophiobolin $G\ (=(3aS,6aS,7R,9aR,10aS)-7-[(1S,2Z)-1,5-dimethylhexa-2,4-dien-1-yl]-3,3a,6,6a,7,8,9,9a,10,10a-decahydro-1,9a-dimethyl-3-oxodicyclopenta[<math>a,d$]cyclooctene-4-carboxaldehyde) were detected [14][15]. Comparison of the 1H - and ^{13}C -NMR data of **4** with those of ophiobolin $G\ [15]$ indicated that the two compounds differed by the

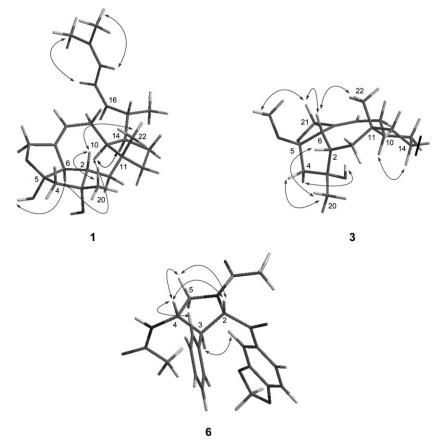


Fig. 3. Key ROESY correlations of compounds 11), 31), and 61)

presence of an OH–CH₂(21) group (δ (H) 3.94 and 3.70; δ (C) 64.2) in **4** instead of the terminal CH(21)=O group in ophiobolin G. The 1 H, 1 H-COSY data (CH₂(21)/H–C(8) and CH₂(21)/OH–C(21)), and the HMBCs CH₂(21)/C(6) and H–C(8)/C(21) supported this assignment. The ROESY correlations H–C(2)/Me(22) and H_{β}–C(1), and H–C(6)/H–C(10) and H $_{\alpha}$ –C(1), and the absence of a correlation H–C(6)/H–C(2) indicated the *trans*-junction of rings A/B. Hence, compound **4** was determined as (6α)-21,21-O-dihydroophiobolin G.

Compound **5** had the molecular formula $C_{25}H_{38}O_4$ as evident from the HR-ESI-MS $(m/z\ 425.2670\ [M+Na]^+)$). The NMR data of **5** $(Tables\ 1\ and\ 2)$ were very similar to those of **4**, with the exception of the side chain where two hydroxylated C-atoms were observed. The 1H , 1H -COSY and HMBC data allowed to position of the two OH groups at C(18) and C(19). Compound **5** was thus determined as (6α) -18,19,21,21-O-tetrahydro-18,19-dihydroxyophiobolin G.

Compound **6** was isolated as a white powder, and its molecular formula $C_{22}H_{22}N_2O_5$ was determined from the HR-ESI-MS $(m/z\ 395.1620\ [M+H]^+)$). The UV spectrum

showed absorption maxima at 235, 283, and 319 nm, characteristic of a conjugated chromophore. The ¹H-NMR spectrum (*Table 3*) exhibited eight aromatic H-atoms, two Me groups $(\delta(H))$ 1.72 (Me(10)) and 2.00 (Me(7))), and one exchangeable H-atom at $\delta(H)$ 8.12 (d, J = 8.5 Hz, H-N(8)). The ¹³C-NMR spectrum (Table 3) displayed 22 Catoms, including one ketone C=O group (δ (C) 197.0 (C(1'))), two amide C=O groups $(\delta(C) 167.9 (C(6)))$ and $(\delta(C) 167.9 (C(6)))$, twelve aromatic C-atoms for two benzene moieties, one acetal C-atom ($\delta(C)$ 101.6 (C(8'))), two Me groups, and four sp³ C-atoms. The ¹H, ¹H-COSY and HMBC data of 6 (Fig. 2) indicated the presence of a pyrrolidine moiety, in which a Ph group was linked at C(3), while a 3,4-(methylenedioxy)phenone moiety was attached to C(2). The presence of an Ac group at the N-atom was deduced from the HMBC H–C(5)/C=O (Ac). In addition, the presence of an acetamide group at position C(4) was confirmed by the HMBC H–C(4)/C=O (δ (C) 169.2). The ROESY correlations (Fig. 3) H–C(4)/H–C(2), H_b –C(5) and H–C(2"), and H–C(2)/ H_b –C(5) indicated that H-C(4) and H-C(2) had the same orientation which was opposite to that of H-C(3). Compound 6 was named aspergillamide A. Structurally similar pyrrolidine alkaloids such as preussin are known from Aspergilli and have been reported previously, e.g., from A. ochraceus [16].

Table 3. ^{1}H - and ^{13}C -NMR Data ((D₆)DMSO, 500 and 125 MHz, resp.) of Compounds 6 and 7^{1}). δ in ppm, J in Hz.

	6		7	
	$\delta(C)$	$\delta(\mathrm{H})$	$\delta(C)$	$\delta(\mathrm{H})$
H-C(2)	61.8 (d)	5.88 (d, J = 9.1)	64.4 (d)	5.32 (d, J=9.5)
H-C(3)	50.1(d)	3.80 (dd, J = 11.5, 9.3)	54.1 (<i>d</i>)	3.18 (dd, J = 9.9, 9.9)
H-C(4)	48.3(d)	5.12 (m)	53.3 (d)	4.62(m)
$CH_2(5)$	51.1 (t)	$4.14 (dd, J = 8.7, 8.7, H_a),$	51.7 (t)	$4.10 (dd, J = 8.7, 8.7, H_a)$
		$3.35 (m, H_b)$		$3.40 (m, H_b)$
C(6)	167.9(s)		167.4(s)	
Me(7)	21.5(q)	2.00(s)	21.4(q)	1.98(s)
H-N(8)	_	8.12 (d, J = 8.5)	_	8.15 (d, J = 8.2)
C(9)	169.2(s)		168.9(s)	
Me(10)	22.4(q)	1.72 (s)	22.2(q)	1.68(s)
C(1')	197.0(s)		195.5 (s)	
C(2')	131.2 (s)		130.3(s)	
H-C(3')	107.2(d)	7.04 (d, J = 1.3)	107.0(d)	7.08 (br. s)
C(4')	146.8 (s)		145.8(s)	
C(5')	150.7(s)		151.1 (s)	
H-C(6')	107.3(d)	6.76 (d, J = 8.2)	107.4(d)	6.78 (d, J = 8.2)
H-C(7')	124.5(d)	7.25 (dd, J = 8.2, 1.3)	124.1 (d)	7.14 (br. $d, J = 8.2$)
$CH_2(8')$	101.6 (t)	6.02(s)	101.7(t)	6.05, 6.04 (2s)
C(1")	134.1 (s)		137.1 (s)	
H-C(2",6")	128.6 (d)	7.12 (d, J = 7.6)	128.3 (d)	7.27 ^a)
H-C(3'',5'')	127.7(d)	7.06 (dd, J = 7.6, 7.2)	127.7(d)	7.22 ^a)
H-C(4")	126.9 (d)	7.01 (dd, J = 7.2, 7.2)	127.2(d)	7.22 – 7.27 ^a)

a) Overlapping signals.

Compound **7** had the same molecular formula as **6** as established by HR-ESI-MS data $(m/z 395.1620 ([M+H]^+))$. ¹H- and ¹³C-NMR, ¹H, ¹H-COSY, HMQC, and HMBC Data revealed that the gross structure of **7** is identical to that of **6**. However, ROESY correlations (Fig. 3) between H–C(4)/H–C(2") and H–C(2)/H–N(8) confirmed that **7** is the C(2) epimer of **6**, for which we propose the name aspergillamide B.

All of the compounds were tested (at a concentration of $10 \,\mu\text{g/ml}$) for cytotoxic activity against murine L5178Y lymphoma cells. However, none of the compounds reduced survival of cells by more than 10-20% compared to controls.

Financial support by the *Bundesministerium für Bildung und Forschung (BMBF)* to *P. P.* and *B. S.* and of *MOST* to *W. L.* is gratefully acknowledged. *H. L.* wishes to thank the *Alfried Krupp von Bohlen and Halbach Foundation* for a scholarship.

Experimental Part

General. Solvents were distilled before use, and spectral-grade solvents were used for spectroscopic measurements. CC=Column chromatography. HPLC: Dionex-P580 system coupled to a photodiodearray detector (UVD340S); routine detection at 235, 254, 280, and 340 nm; column Eurospher-10 C_{18} (125 × 4 mm (i.d.); Knauer, Germany); linear gradient 0.02% H_3PO_4 in H_2O and MeOH. Semi-prep. HPLC: LaChrom-Merck-Hitachi machine (pump L-7100); column Eurospher-100 C_{18} (8 mm length; Knauer, Germany); flow rate 5.0 ml/min. UV Spectra: L-7400 UV detector; λ_{max} in nm). Optical rotations: Perkin-Elmer-241-MC polarimeter. 1D- and 2D-NMR Spectra: Bruker-ARX-500 or Avance-DMX-600 spectrometers; at 500 or 600 MHz (^{1}H) and 125 or 150 MHz (^{13}C); δ in ppm rel. to Me₄Si as internal standard, J in Hz; DUL plus probes. ESI-MS: Finnigan-LCQ-Deca mass spectrometer; in m/z (rel. %). HR-ESI-MS: Micromass-Q-Tof mass spectrometer; in m/z (rel. %).

Fungal Material. The fungus Aspergillus ustus, internal strain 8009, was isolated from the marine sponge Suberites domunculus, which had been collected from the Adriatic Sea and then kept in an aquarium until fungi were isolated from sponge segments as previously described [7]. The fungus was identified both according to morphological and molecular attributes [7].

Cultivation and Extraction. For production of secondary metabolites, the fungus was cultivated at 22° for 21 d on both biomalt agar [4] and on barley-spelt solid media [7]. For initial analysis of the natural products, the cultures were lyophilized and extracted with AcOEt, and the dried residues were defatted by petroleum ether extraction.

Isolation. The AcOEt extract of A. ustus (3.66 g) was fractionated by VLC (silica gel, CH₂Cl₂ with increasing amounts of MeOH). Fraction 4 (CH₂Cl₂/MeOH 98:2; 681 mg) was further purified by CC (Sephadex LH-20, CH₂Cl₂/MeOH 1:1) and then subjected to prep. HPLC (MeOH/H₂O 80:20): **1** (2.4 mg), **2** (2.5 mg), **3** (3.6 mg), and **4** (1.9 mg). Fr. 6 (CH₂Cl₂/MeOH 96:4; 262 mg) was further purified by CC (Sephadex LH-20, CH₂Cl₂/MeOH 1:1) and prep. HPLC (MeOH/H₂O 80:20): **7** (1.9 mg) and aurantiamide (6.0 mg). Fr. 7 (CH₂Cl₂/MeOH 94:6; 377 mg) was further purified by CC (Sephadex LH-20, 100% MeOH) and prep. HPLC (MeOH/H₂O 80:20): **5** (2.2 mg) and **6** (6.0 mg). Fr. 8 (CH₂Cl₂/MeOH 90:10; 366 mg) was further purified by CC (Sephadex LH-20, 100% MeOH): cerebroside D (10.0 mg).

 $(5\alpha,6\alpha)$ -Ophiobolin H (= (1R,2aR,6aS,7R,9aR,10aS,10bR)-7-[(1S,2Z)-1,5-Dimethylhexa-2,4-dien-1-yl]-1,2,4,6,6a,7,8,9,9a,10,10a,10b-dodecahydro-1,9a-dimethyl-2aH-3-oxacyclopenta[5,6]cycloocta[1,2,3-cd]pentalene-1,2a-diol; 1): White amorphous powder. [α] $_{D}^{20}$ = +25 (c = 0.1, CHCl $_{3}$). UV (HPLC, mobile phase): 242. 1 H- and 13 C-NMR: Tables 1 and 2. HR-ESI-MS: 409.2720 ([M+Na] $^{+}$, C_{25} H $_{38}$ NaO $_{3}^{+}$; calc. 409.2719)

 $(5\alpha,6\alpha)$ -5-O-Methylophiobolin H (= (1R,2aR,6aS,7R,9aR,10aS,10bR)-7-[(1S,2Z)-1,5-Dimethyl-hexa-2,4-dien-1-yl]-2,2a,4,6,6a,7,8,9,9a,10,10a,10b-dodecahydro-2a-methoxy-1,9a-dimethyl-1H-3-oxacy-clopenta[5,6]cycloocta[1,2,3-cd]pentalen-1-ol; **2**): White amorphous powder. $[\alpha]_D^{30} = +99$ (c = 0.1, CHCl₃). UV (HPLC, mobile phase): 236. 1 H- and 1 3C-NMR: *Tables 1* and 2. HR-ESI-MS: 423.2880 ($[M+Na]^+$, $C_{26}H_{40}NaO_3^+$; calc. 423.2875).

5-O-Methylophiobolin H (= (1R,2aS,6aS,7R,9aR,10aS,10bS)-7-[(1S,2Z)-1,5-Dimethylhexa-2,4-dien-1-yl]-2,2a,4,6,6a,7,8,9,9a,10,10a,10b-dodecahydro-2a-methoxy-1,9a-dimethyl-1H-3-oxacyclopenta-[5,6]cycloocta[1,2,3-cd]pentalen-1-ol; **3**): White amorphous powder. [a] $_D^{20}$ = +33 (c = 0.1, CHCl $_3$). UV (HPLC, mobile phase): 242. 1 H- and 13 C-NMR: Tables 1 and 2. HR-TOF-MS: 423.2860 ([M+Na] $^+$, C_{26} H $_{40}$ NaO $_3^+$; calc. 423.2875).

(6a)-21,21-O-Dihydroophiobolin G (= (3aR,6aS,7R,9aR,10aS)-7-[(1S,2Z)-1,5-Dimethylhexa-2,4-dien-1-yl]-6,6a,7,8,9,9a,10,10a-octahydro-4-(hydroxymethyl)-1,9a-dimethyldicyclopenta[a,d]cycloocten-3(3aH)-one; **4**): White amorphous powder. [a] $_D^{20}$ = +49 (c = 0.1, CHCl $_3$). UV (HPLC, mobile phase): 236. $_1^1$ H- and $_1^3$ C-NMR: *Tables 1* and 2. HR-ESI-MS: 391.2610 ([M+Na] $_1^+$, C_{25} H $_{36}$ NaO $_2^+$; calc. 391.2613).

 (6α) -18,19,21,21-O-Tetrahydro-18,19-dihydroxyophiobolin G (= (3aR,6aS,7R,9aR,10aS)-7-[(1S,2Z)-4,5-Dihydroxy-1,5-dimethylhex-2-en-1-yl]-6,6a,7,8,9,9a,10,10a-octahydro-4-(hydroxymethyl)-1,9a-dimethyldicyclopenta[a,d]cycloocten-3((3aH)-one; **5**): White amorphous powder. [$(a_1)^{20}_{10} = +27$ ($(c=0.1, CHCl_3)$). UV (HPLC, mobile phase): 232. 1 H- and 1 3C-NMR: Tables 1 and 2. HR-ESI-MS: 425.2670 ([$(M+Na)^+$, $(C_2SH_{38}NaO_4^+$; calc. 425.2668).

Aspergillamide A = N-[(3\$, 4\$, 5\$)-1-Acetyl-5-(1,3-benzodioxol-5-ylcarbonyl)-4-phenylpyrrolidin-3-yllacetamide;**6**): White amorphous powder. [<math>a] $_D^{20} = -76$ (c = 0.1, MeOH). UV (HPLC, mobile phase): 235, 283, 319. 1 H- and 13 C-NMR: *Tables 1* and 2. HR-ESI-MS: 395.1620 ([M+H] $^{+}$, $C_{22}H_{23}N_{2}O_{5}^{+}$; calc. 395.1607)

Aspergillamide $B = N-[(3S,4S,5R)-1-Acetyl-5-(1,3-benzodioxol-5-ylcarbonyl)-4-phenylpyrrolidin-3-yl]acetamide; 7): White amorphous powder. [<math>\alpha$] $_D^{20} = +45$ (c = 0.1, MeOH). UV (HPLC, mobile phase): 233, 282, 316. 1H - and 13 C-NMR: $Tables\ 1$ and 2. HR-ESI-MS: 395.1620 ([M+H] $^+$, $C_{22}H_{23}N_2O_5^+$; calc. 395.1607)

Cell Proliferation Assays. Cytotoxicity was tested against L5178Y cells by using the MTT (= 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-2*H*-tetrazolium bromide) assay as described previously [11].

REFERENCES

- [1] P. Proksch, W. E. G. Müller, 'Frontiers in Marine Biotechnology', Horizon Bioscience, Norfolk, 2006.
- [2] G. M. König, S. Kehraus, S. F. Seibert, A. Abdel-Lateff, D. Müller, Chem. Bio. News 2006, 7, 229.
- [3] T. S. Bugni, C. M. Ireland, Nat. Prod. Rep. 2004, 21, 143.
- [4] U. Höller, A. D. Wright, G. F. Matthee, G. M. König, S. Draeger, H.-J. Aust, B. Schulz, Mycol. Soc. Phytochem. 2000, 104, 1354.
- [5] J. Kohlmeyer, B. Volkmann-Kohlmeyer, Mycol. Res. 2003, 107, 386.
- [6] S. Perovic-Ottstadt, T. Adell, P. Proksch, M. Wiens, M. Korzhev, V. Gamulin, W. E. G. A. Müller, J. Eur. Biochem. 2004, 271, 1924.
- [7] P. Proksch, R. Ebel, R. Edrada, F. Reibe, H. B. Liu, A. Diesel, M. Bayer, X. Li, W. H. Lin, V. Grebenyuk, W. E. G. Müller, S. Draeger, A. Zuccaro, B. Schulz, *Bot. Mar.* 2008, 51, 209.
- [8] M. A. Hayes, S. K. Wrigley, I. Chetland, E. E. Reynolds, A. M. Ainsworth, D. V. Renno, M. A. Latif, X. M. Cheng, D. J. Hupe, P. Charlton, A. M. Doherty, J. Antibiot. 1996, 49, 505.
- [9] W. A. Ayer, L. M. Pena-Rodriguez, J. Nat. Prod. 1987, 50, 408.
- [10] Z. Lu, Y. Wang, C. Miao, P. Liu, K. Hong, W. Zhu, J. Nat. Prod. 2009, 72, 1761.
- [11] H. Liu, R. Edrada, R. Ebel, Y. Wang, B. Schulz, S. Draeger, W. E. G. Müller, V. Wray, W. Lin, P. Proksch, J. Nat. Prod. 2009, 72, 1585.
- [12] T. O. Larsen, J. C. Frisvad, S. R. Jensen, Phytochemistry 1992, 31, 1613.
- [13] R. D. Sitrin, G. Chan, J. Dingerdissen, R. M. DeBrosse, G. Roberts, S. Rottschaefer, D. Staiger, J. Valenta, K. M. Snader, R. J. Stedman, J. R. E. Hoover, J. Antibiot. 1988, 41, 469.
- [14] H. G. Cutler, F. G. Crumley, R. H. Cox, J. P. Springer, R. F. Arrendale, R. J. Cole, P. D. Cole, J. Agric. Food Chem. 1984, 32, 778.
- [15] H. Wei, T. Itoh, M. Kinoshita, Y. Nakai, M. Kurotaki, M. Kobayashi, Tetrahedron 2004, 60, 6015.
- [16] R. E. Schwartz, J. Liesch, O. Hensens, L. Zitano, S. Honeycutt, G. Garrity, R. A. Fromtling, J. Onishi, R. Monaghan, J. Antibiot. 1988, 41, 1774.

Received July 27, 2010