

Structural and spectral assignments of six anthraquinone derivatives from the mangrove fungus (ZSUH-36)

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A new natural product named 6,8,1'-tri-*O*-methyl averantin(1) has been isolated together with five known anthraquinones 1'-*O*-methyl averantin(2), 6,8-di-*O*-methyl averufin (3) averufin (4), versicolorin C (5) and 6,8-di-*O*-methyl averufanin (6) from a mangrove endophytic fungus ZSUH-36 collected from the South China Sea. NMR techniques including COSY, HMQC, and HMBC were used to elucidate the structures of these compounds. We report the unambiguous assignments of the ¹H and ¹³C NMR spectra of the new compound 6,8,1'-tri-*O*-methyl averantin(1). Copyright © 2008 John Wiley & Sons, Ltd.

Keywords: NMR; ¹H NMR; ¹³C NMR; 2D NMR; anthraquinone; secondary metabolites; structure elucidation

Introduction

Anthraquinone derivatives as a class of natural products are widespread in nature. They showed strong cytotoxic activity against cancer cell lines such as L1210,^[1] LNCap,^[2] and A431.^[3] Previously, we investigated the chemistry of the fungus (ZSUH-36) and reported the isolation and structure elucidation of two anthraquinones and two xanthenes from this endophytic fungus.^[4] In a further study on the secondary metabolites of this mangrove endophytic fungus ZSUH-36, we isolated six anthraquinones which were the new natural product 6,8,1'-tri-*O*-methyl averantin (1) and five known anthraquinones, namely, 1'-*O*-methyl averantin (2),^[5] 6,8-di-*O*-methyl averufin (3),^[6] averufin (4),^[7,8] versicolorin C (5),^[8,9] and 6,8-di-*O*-methyl averufanin (6).^[10] The structures of the compounds were elucidated on the basis of the 1D- and 2D-NMR experiment results, particularly those of COSY, HMQC, and HMBC. Among the known compounds reported previously in the literature, NMR data of compound 3 were incomplete with only proton results available. Therefore, we report the NMR spectral data of compound 1 and the 2D-NMR data of compound 3. Compounds 2, 3, and 6 were isolated for the first time as marine fungus metabolites.

Results and Discussion

The new compound 6,8,1'-tri-*O*-methyl averantin (1) was obtained as a red amorphous solid. It (Fig. 1) showed a molecular ion peak at *m/z* 414 in EIMS. Its molecular formula C₂₃H₂₆O₇ was assigned on the basis of HR-EIMS (*m/z* 414.1676, calcd 414.1673) and the NMR data, which indicated that the compound has 11° of unsaturation. The ¹³C NMR and DEPT data implied that compound 1 has 1 methyl group, 3 methoxyl groups, 4 methylene carbons, 4 methine carbons, and 11 quaternary carbons including 2 carbonyl carbons at δ_C 187.3 and 182.7. The ¹H NMR spectrum and HMQC data (Table 1) revealed the presence of a hydrogen-bonded hydroxyl group [δ_H 13.99 (1H, s)], two signals of *meta*-coupled aromatic

protons at δ_H 6.85 (d, 2.5) and 7.21 (d, 2.5), one singlet due to an aromatic proton at δ_H 6.94, three methoxyl groups at δ_H 3.89, 3.87, 3.35, and four methylene groups at δ_H 1.20–1.21(4H, overlapped, H-4' and H-5'); 1.31 (1H, m, H-3'), 1.40 (1H, m, H-3'), 1.74 (1H, m, H-2'), and 1.77 (1H, m, H-2'). The ¹H and ¹³C NMR (Table 1) spectra indicated that compound 1 was similar to averantin, except that the three methoxyl groups in compound 1 were replaced by three hydroxyl groups. Detailed assignments of the carbon and proton signals were unambiguously accomplished by analysis of 2D-NMR spectral data (Table 1). The structure of the six-carbon chain moiety was confirmed by sequential correlations of H–H COSY from H-2' to H-1', H-3' and from H-6' to H-4', H-5'. In the HMBC spectrum, key correlations were observed between H-1' and C-2, indicating that the six-carbon chain moiety was connected to C-2. Furthermore, the two aromatic protons at δ_H 6.94 (H-4) and 7.21 (H-5) gave cross-peaks with the carbonyl carbon at δ_C 182.7 (C-10) and C-2 (δ_C 119.8), C-3 (δ_C 162.9), C-9a (δ_C 111.0) and C-7 (δ_C 105.1), C-8a (δ_C 115.3), respectively, which were consistent with their locations on carbons C-4 and C-5, and the carbonyl group at C-10 (δ_C 182.7). The rest of the correlations have been summarized and listed in

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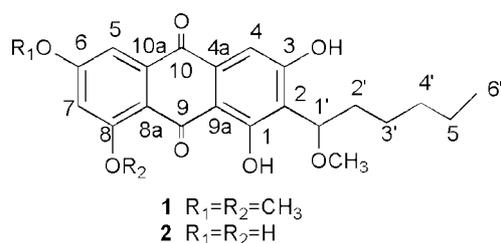


Figure 1. The structures of compounds 1 and 2.

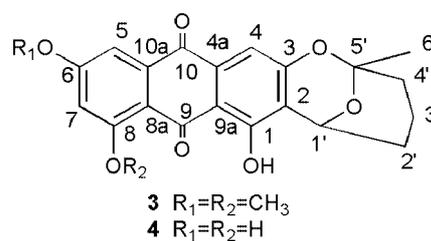


Figure 2. The structures of compounds 3 and 4.

Table 1. NMR data of compound 1 (δ in ppm, J in Hz)

C no.	¹³ C NMR (DEPT)	¹ H NMR	H-H COSY	HMBC
1	163.1			
2	119.8			
3	162.9			
4	108.1 CH	6.94(s)		C-2, C-3, C-10, C-9a
5	105.4 CH	7.21(d, 2.5)	H-7	C-7, C-10, C-8a
6	166.2			
7	105.1 CH	6.85(d, 2.5)	H-5	C-5, C-6, C-8, C-8a
8	164.4			
9	187.3			
10	182.7			
4a	134.0			
8a	115.3			
9a	111.0			
10a	138.0			
8-OCH ₃	56.9	3.89(s)		C-8
9-OCH ₃	56.5	3.87(s)		C-6
1'	79.8 CH	4.85(dd, 5.0, 8.0)	H-2'	C-2, C-3, C-7'
2'	35.2 CH ₂	1.77(m)	H-1', H-3'	C-1'
		1.74(m)	H-1', H-3'	C-1'
3'	25.7 CH ₂	1.31(m)	H-2'	C-5'
		1.40(m)	H-2'	C-4'
4'	32.3 CH ₂	1.21 (overlapped)	H-6'	C-5'
5'	23.2 CH ₂	1.20 (overlapped)	H-6'	C-4'
6'	14.2 CH ₃	0.75(m)	H-4', H-5'	C-4', C-5'
1'-OCH ₃	58.2	3.35(s)		C-1'
1-OH		13.99(s)		C-1, C-2, C-9a

Table 2. NMR data of compound 3 (δ in ppm, J in Hz)

C no.	¹³ C NMR (DEPT)	¹ H NMR	H-H COSY	HMBC
1	159.6			
2	116.8			
3	159.6			
4	107.0 CH	7.20(s)		C-2, C-3, C-10, C-9a
5	104.0 CH	7.45(d, 2.5)	H-7	C-7, C-10, C-8a
6	164.9			
7	104.8 CH	6.77(d, 2.5)	H-5	C-5, C-6, C-8, C-8a
8	162.8			
9	186.8			
10	182.6			
4a	132.5			
8a	115.3			
9a	104.0			
10a	137.6			
8-OCH ₃	56.6	4.01(s)		C-8
8-OCH ₃	56.0	3.97(s)		C-6
1'	67.1 CH	5.38(d, 3.0)	H-2'	C-1, C-2, C-3, C-3', C-5'
2'	27.4 CH ₂	1.88(m)	H-1', H-3'	C-2, C-4'
		2.10(m)		C-1
3'	16.0 CH ₂	1.53(m)	H-2', H-4'	C-5'
		1.65(m)	H-2', H-4'	C-1', C-4', C-5'
4'	35.9 CH ₂	1.84(m)	H-3'	C-3'
		2.04(m)	H-3'	C-3', C-5', C-6'
5'	100.9			
6'	27.8 CH ₃	1.58(s)		C-3, C-1', C-4'
1-OH		13.55(s)		C-1, C-2

Table 1. On the basis of these spectroscopy data the compound was identified as 6,8,1'-tri-*O*-methyl averantin (**1**).

Compound 1'-tri-*O*-methyl averantin (**2**) was isolated as a red amorphous powder with a molecular formula of C₂₁H₂₂O₇ assigned by using its EIMS and NMR data. Compared with the ¹H NMR data of **1**, compound **2** had two extra hydroxyl groups at δ_{H} 12.51, 10.15 and lacked two methoxyl groups at δ_{H} 3.89, 3.87. According to the above data this compound can be deduced as 1'-*O*-methyl averantin. This was also supported by the comparison of its ¹H and ¹³C NMR spectra with those reported in the literature.^[5] This is the first report that compound **2** was obtained from a marine fungus resource.

Compound **3** (Fig. 2) was isolated as a yellow solid. According to the LC-MS and NMR data, the molecular formula of **3** was established as C₂₂H₂₀O₇. The ¹³C NMR and DEPT spectra showed 22 carbon signals, attributable to 1 methyl (δ_{C} 27.8), 2 methoxyl

(δ_{C} 56.0, 56.6), 3 methylene (δ_{C} 16.0, 27.4, 35.9), 4 methine (δ_{C} 67.1, 104.0, 104.8, 107.0), 10 quaternary (δ_{C} 100.9, 104.0, 115.3, 116.8, 132.5, 137.6, 159.6, 159.6, 162.8, 164.9), and two ketone carbonyl (δ_{C} 182.6, 186.8) carbons. The ¹H NMR spectrum (Table 2) displayed a hydrogen-bonded hydroxyl group at δ_{H} 13.55 (1H, s), two *meta*-coupled aromatic proton signals at δ_{H} 6.77 and 7.45, one singlet due to an aromatic proton at δ_{H} 7.20, two methoxyl group at δ_{H} 4.01 and 3.97, one methyl group at δ_{H} 1.58, one methine at δ_{H} 5.38 (d, 3.0 Hz) attached to an oxygen atom, and three methylene groups at δ_{H} 1.88 (1H, m), 2.10 (1H, m); 1.53 (1H, m), 1.65 (1H, m); and 1.84 (1H, m), 2.04 (1H, m). The assignments of the carbon and proton signals were unambiguously accomplished by a further analysis of the 2-D NMR spectral data that are listed in Table 2. This was also the first report on 6,8-di-*O*-methyl averufin isolated from a marine fungus.

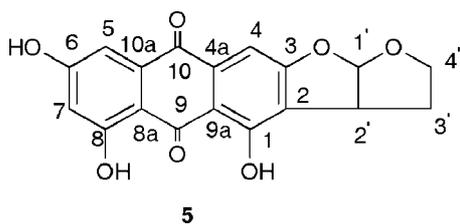


Figure 3. The structure of compound 5.

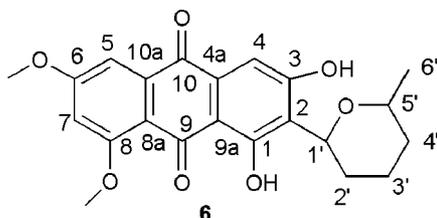


Figure 4. The structure of compound 6.

Compound 4 (Fig. 2) and compound 5 (Fig. 3) were also isolated from the fungus. Their structures were identified as averufin^[7,8] and versicolorin C^[8,9] by comparison of their spectroscopic data with those in the related literature.

Compound 6 (Fig. 4) was obtained as a red powder. The molecular formula of this compound was C₂₂H₂₂O₇ confirmed by LC-MS which showed [M – H][–] ion peak at *m/z* 397. This compound was isolated from a terrestrial fungus^[10] but this is the first report from a marine fungus in the South China Sea.

Experimental

General experimental procedures

Mass spectra were recorded on a VG ZAB-HS double focusing mass spectrometer, IR spectra were measured on a Bruker EQUINOX 55 spectrophotometer, UV spectra were obtained on a Shimadzu UV-2501PC spectrophotometer. Column chromatography was carried out on silica gel (200–300 mesh; Qingdao Haiyang Chemicals).

Fungal strain

The fungus strain (ZSUH 36) was isolated from the Shenzhen mangrove *Acanthus ilicifolius* Linn. and the specimen was stored at the Department of Applied Chemistry, Sun Yat-sen University, Guangzhou, China. Starter cultures were maintained on cornmeal seawater agar.

Fermentation, extraction, and isolation

Plugs of agar-supporting mycelium growth were cut from a solid culture medium and transferred aseptically to a 500-ml Erlenmeyer flask containing 250 ml liquid medium (glucose 10 g/l, peptone 2 g/l, yeast extract 1 g/l, sea salt 2.5 g/l).^[11] The fungus was cultured in 70 l and incubated at 28 °C, for 35 days. The broth was filtered through a cheesecloth. The mycelium (450 g) was air-dried and extracted with methanol. The filtrate was concentrated to 3 l below 60 °C and partitioned four times by shaking with an equal volume of ethyl acetate. The EtOAc extract (25.6 g) of the fungal mycelium was subjected to silica gel chromatography using a gradient from

petroleum to ethyl acetate (fractions A–F) and then from ethyl acetate to methanol (fractions G–I). Fraction D (1.2 g) was further separated by preparative TLC (silica gel GF₂₅₄) developed by 2% methanol/chloroform (v/v) to afford fractions D₁–D₄. Fraction D₁ (226.0 mg) was subjected to a further column chromatography to afford compounds 1 (2.6 mg), 3 (56.2 mg), 4 (25.5 mg), 6 (2.5 mg). Fraction D₂ (86.4 mg) was again separated by PTLC to obtain 2 (12.6 mg) and 5 (18.7 mg).

1',6,8-Tri-O-methyl averantin (1): EIMS (*m/z*): 414 (M⁺, 4), 399 (4), 382 (42), 353 (16), 343 (32), 339 (10), 325 (100), 313 (20), 310 (10).

1'-O-Methyl averantin (2): EIMS (*m/z*): 386 (M⁺, 11), 368 (13), 354 (96), 325 (39), 315 (100), 311 (40), 297 (94), 285 (84), 272 (23), 255 (11), 244 (9), 213 (8), 155 (6). ¹H NMR (500 MHz, Acetone-*d*₆): δ 12.75 (OH), 12.51 (OH), 10.15 (OH), 9.75 (OH), 7.21 (1H, d, 2.4 Hz, H-5), 7.11 (s, H-4), 6.62 (1H, d, 2.4 Hz, H-7), 4.96 (1H, dd, 5.0, 8.0, H-1'), 3.49 (s, 1'-OCH₃), 1.83 (2H, m, H-2'), 1.41 (2H, m, H-3'), 1.33 (4H, overlapped, H-4', 5'), 0.88 (t, 6.8, 6'-CH₃). ¹³C NMR (125 MHz, CDCl₃): δ 191.4 (C-9), 182.4 (C-10), 166.8 (C-6), 166.7 (C-8), 165.0 (C-1), 163.7 (C-3), 137.2 (C-10a), 135.8 (C-4a), 120.4 (C-2), 111.1 (C-8a), 111.0 (C-4), 110.6 (C-9a), 110.5 (C-5), 109.9 (C-7), 80.7 (C-1'), 59.3 (1'-OCH₃), 36.1 (C-2'), 33.3 (C-4'), 26.7 (C-3'), 24.2 (C-5'), 15.3 (C-6').

6,8-Di-O-methyl averufanin (6): ¹H NMR (500 MHz, CDCl₃): δ 13.74 (OH), 9.75 (OH), 7.38 (d, 2.4 Hz, H-5), 7.16 (s, H-4), 6.71 (d, 2.4 Hz, H-7), 5.11 (dd, 1.5, 11.1 Hz, H-1'), 3.96 (s, 5-OCH₃), 3.92 (s, 7-OCH₃), 3.68 (ddd, 5.4, 6.3, 11.2 Hz, H-5'), 1.94 (m, H-2'), 1.76 (m, H-4'), 1.40 (m, H-3'), 1.29 (d, 7.0 Hz, H-6'). LC-MS (*m/z*): 397 (M[–], 100), 382 (16), 367 (42), 353 (48), 337 (36), 312 (66), 309 (47), 297 (41).

NMR data

NMR spectra (1H, COSY, HSQC and HMBC) were recorded at 25 °C using TMS as internal reference on Varian Inova 500NB NMR spectrometer (¹H, 499.77 MHz; ¹³C, 125.7 MHz) equipped with a triple resonance inverse cold probe cooled to 25 K. The ¹H NMR and ¹³C NMR spectra provided digital resolutions of 0.25 and 0.17 Hz, respectively. The ¹H and ¹³C chemical shifts were referenced to the residual solvent peak of Acetone-*d*₆ at δ 2.05 and 30.8 ppm and CDCl₃ at δ 7.26 and 77.0 ppm, for proton and carbon respectively. All sample concentrations ranged from 5 to 15 mg ml^{–1} with a total volume of 0.5 ml for each sample. The ¹H sweep width was set at 7506 Hz for all experiments with a 45° pulse for ¹H and a ¹³C 90° pulse. The gradient COSY was acquired with 200 F1 increments with 32 scans per increment. A sinebell weighting was applied to each dimension and zero filled to 2 K points. The gradient HMQC was acquired with ¹³C sweep width of 32 000 Hz and 256 t1 increments. Each increment was acquired with 32 transients. A sinebell function was applied to the F2 dimension before zero filling to 2 K points, and a sinebell was applied to the F1 and zero filled to 2 K points before Fourier transformation. A one-bond coupling constant delay was set using 140 Hz, and WURST decoupling was applied during acquisition. The gradient HMBC was acquired using 64 transients per increment with 256 F1 increments. A sweep width of 32 000 Hz was used for the ¹³C dimension. The one-bond coupling constant of 140 Hz and a long-range coupling constant of 8 Hz were used to set the delays in the pulse sequence. A sinebell weighting was applied to both ¹H and ¹³C dimensions and zero filled to 4 and 1 K, respectively.

Acknowledgements

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