

NOTE

Ophiosetin, a new tetramic acid derivative from the mycopathogenic fungus *Elaphocordyceps* ophioglossoides

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Members of the subgroup of tetramic acids (pyrrolidine-2,4-diones) that feature substituted aliphatic bicyclic ring systems, such as equisetin from Fusarium equiseti¹ and altersetin from Alternaria sp.,² show potent antibacterial activities against Gram-positive bacteria, and coniosetin from Coniochaeta ellipsoidea³ and CJ-17572 from Pezicula sp.4 further exhibit inhibitory activity against multidrug-resistant Staphylococcus aureus and Enterococcus faecalis. In addition to their antimicrobial activity, equisetin-related compounds display various notable biological activities, such as cytotoxicity against P388 tumor cell lines (paecilosetin from *Isaria farinosa*⁵), phytotoxicity against five plant species (trichosetin from the dual culture of Trichoderma harzianum and Catharanthus roseus callus⁶) and HIV (human immunodeficiency virus)-1 integrase inhibitory activity (equisetin from Fusarium heterosporum and phomasetin from Phoma sp.7). This wide range of biological activities makes this class of compounds highly attractive for the discovery of novel bioactive compounds. During our continuous screening for new metabolites from filamentous fungi, 8-10 an HPLC-diode array detection analysis of the n-BuOH extract of Elaphocordyceps ophioglossoides showed two compounds with maximum absorbances at \sim 250 and 290 nm, which were indicative of the presence of tetramic acid moiety.^{5,11,12} As previous available reports mentioned only limited metabolites from E. ophioglossoides, including the antifungal antibiotic ophiocordin¹³ in 1977 and an antitumor polysaccharide in 1984, ¹⁴ we initiated the isolation of the compounds. A detailed analysis of the fungal extract led to the identification of equisetin and a new equisetin analog, which we named ophiosetin (1 and 2, Figure 1). This is the first report on either equisetin or an equisetin-related compound produced by this species. In the antimicrobial assay against bacteria, yeasts and filamentous fungi, ophiosetin (1) showed a markedly weaker antibacterial activity compared with the very closely related analogs, equisetin (2) and

paecilosetin (3). We herein describe the isolation, structure elucidation and biological activity of ophiosetin (1).

MATERIALS AND METHODS

General procedures

UV spectra were recorded on a Hitachi U-3000 spectrophotometer (Hitachi, Tokyo, Japan). NMR spectra were obtained on a JEOL ECP 400 and ECS 400 spectrometer (Jeol, Tokyo, Japan). The $^1\mathrm{H}$ and $^{13}\mathrm{C}$ chemical shifts were referenced to the solvent signals ($\delta_{\mathrm{H}}.3.30$ and $\delta_{\mathrm{C}}.49.1$ in CD₃OD). HR (high-resolution) FAB-MS were recorded on a JEOL JMS-700 spectrometer. HPLC analyses for screening were carried out on an Agilent HP1100 system (Agilent, Tokyo, Japan) using a Cosmosil 5C18-AR-II column ($100\times4.6~\mathrm{mm}^2$ i.d.; Nacalai Tesque, Kyoto, Japan). The solvent used to dissolve 1 and 2 was methanol or chloroform unless stated otherwise. All chemicals, media and reagents were purchased from Wako (Osaka, Japan), unless stated otherwise.

Fermentation and isolation

The fungus, *E. ophioglossoides* HF272, was isolated from a soil specimen collected at the Tsuchiyu Hot Spring in Fukushima, Japan. Ascospores released from the specimen were transferred to SMY and incubated at $25\,^{\circ}\text{C}$ for several days. After mycelia had developed on SMY slants, the strain was kept at $-30\,^{\circ}\text{C}$ as a stock culture. The isolated strain was identified as *E. ophioglossoides* according to its morphology by one of the authors (FI) and is deposited at the culture collection of the National Institute of Fruit Tree Science (Ibaraki, Japan) as strain HF272.

The seed culture was prepared as follows: The mycelium of *E. ophioglossoides* HF272 grown on a slant culture was inoculated into test tubes, each containing 5 ml of the seed medium, SMY (maltose 4%, yeast extract 1%, peptone 1%) and cultivated at 25 °C for 5 days. The seed culture (5 ml) was transferred into 500-ml baffled flasks containing 250 ml of the production medium (SMY supplemented with Diaion HP-20 (Mitsubishi Chemical, Tokyo, Japan) 1%) and the flasks were cultured at 25 °C for 21 days under a static condition.

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Figure 1 Structure of ophiosetin (1), equisetin (2) and paecilosetin (3).

Table 1 NMR spectroscopic data of ophiosetin (1) in CD₃OD at −80 °C

Position	$\delta_{\mathcal{C}}^{a}$	$\delta_H{}^{b}$	DEPT	<i>HMBC</i> ^b	COSY
1	197.3	_	Co	H-12	
2	49.8	_	C_0	H-12	
3	45.8	3.34 (br s, ¹ H)	CH	H-4, H-5, H-12, H-13, H-14	H-4, H-13
4*	128.2	5.46 (m, ¹ H)	CH		
5*	131.0	5.47 (m, ¹ H)	CH		
6	39.4	1.89 (m, ¹ H)	CH	H-5	H-11
7**	31.0	a: 1.10 , b: 1.86 (m, ¹ H)	CH_2	H-16	<i>a</i> : H-7b
8	42.2	1.62 (m, ¹ H)	CH	H-16	
9	37.6	a: 0.87 (ddd, J=12, 12, 12 Hz, ¹ H), b: 1.96(m, ¹ H)	CH ₂	H-16	<i>a</i> : H-9b, H-8
10**	28.9	a: 1.10, b: 2.07 (m, ¹ H)	CH ₂		a: H-10b, H-11
11	41.3	1.69 (m, ¹ H)	CH	H-12, H-5	H-6, H-10b
12	13.9	1.46 (br s, ³ H)	CH ₃	H-11, H-3, H-2	
13	132.1	5.20 (m, ¹ H)	CH	H-15	
14	130.6	5.21 (m, ¹ H)	CH	H-15	H-15
15	18.6	1.54 (d, <i>J</i> =5.5 Hz, ³ H)	CH ₃	H-14	H-14
16	68.3	3.39 (m, ² H)	CH_2		
2'	177.6	_	C_0	<i>N</i> -CH ₃ , H-5'	
3′	102.1	_	C_0		
4′	192.9	_	C_0	H-5′, H-6′	
5′	69.0	3.70 (br.s, ¹ H)	CH	N-CH₃	H-6′
6′	58.8	a: 3.89 (dd, J=12.4, 2.7 Hz, ¹ H), b: 3.95 (dd, J=12.4, 3.2 Hz, ¹ H)	CH ₂		H-5′
N-CH ₃	26.8	3.02 (s, ³ H)	CH ₃	H-5′	

^{*, **} exchangeable.

The whole cell broth $(250\,\text{ml}\times4~\text{flasks})$ of strain HF272 after 21 days of cultivation was extracted with ethyl acetate (without saturation with water earlier, 125 ml per flask) by stirring for 3 h. The mixture was separated by filtration using Miracloth (Calbiochem, La Jolla, CA, USA) and the filtrate was extracted three times, each with 250 ml of ethyl acetate. The organic layer was separated from the aqueous layer in an extraction funnel by solvent partition, dried over anhydrous Na₂SO₄ and evaporated to provide \sim 334 mg of extract per 11 of culture. A portion of the crude extract (200 mg) was subjected to reversed-phase column chromatography using a Sep-Pak Vac 35-cc (10 g) C_{18} cartridge (Waters, Milford, MA, USA) with a step gradient of CH₃CN-H₂O (0:1, 2:8, 4:6, 8:2 and 1:0 v/v). Compound 1 (58.8 mg g⁻¹ extract) and compound 2 (8.5 mg g⁻¹ extract) were purified by reversed-phase HPLC on a Shiseido Capcell-Pak C_{18} column (Shiseido, Tokyo, Japan) (5 µm; 250×10 mm² i.d.) at 254 nm with 40% CH₃CN+0.1% TFA and with 70% CH₃CN+0.1% TFA, respectively.

Ophiosetin (1). Ophiosetin (1) is a pink-brown oily solid; $[α]^{22}_D-244^\circ$ (c 0.02,CHCl₃); UV (MeOH) $λ_{max}$ (log ε) 204 (3.42), 252 (4.03), 293 (4.22); 1 H, 13 C NMR, heteronuclear multiple bond coherence (HMBC), see Table 1; HRFAB-MS m/z [M+H] $^+$ 390.2295 (calcd for C_{22} H₃₂NO₅, 390.2280).

Equisetin (2). Equisetin (2) is a pink-brown oily solid; $[\alpha]^{22}_D-145^\circ$ (c 0.02,CHCl₃); UV (MeOH) $\lambda_{\rm max}$ (log ε) 204 (4.29), 250 (3.94), 295 (4.01); $^1{\rm H}$ NMR data, UV and specific optical rotation were consistent with the data reported in literature; 12,15 HRFAB-MS m/z [M+H]⁺ 374.2340 (calcd for $C_{22}H_{32}NO_4$ 374.2331).

Antimicrobial assay

Antimicrobial activities of 1, 2 (from *E. ophioglossoides*) and 3 (from *Isaria farinosa*) were determined by the standardized two-fold broth dilution methods recommended by the Clinical and Laboratory Standards Institute (CLSI, Wayne, PA, USA; formerly the National Committee for Clinical and Laboratory Standards, NCCLS). ¹⁶ The MIC was defined as the lowest drug concentration resulting in complete inhibition of growth after 18 h of incubation at 37 °C (bacteria) or 35 °C (fungi). The following bacteria were used as indicator strains: *S. aureus* American Type Culture Collection (ATCC) 25923, *E. faecalis* ATCC 29212, *Escherichia coli* ATCC 25922 and *Pseudomonas aeruginosa* ATCC 27853. The following yeasts and filamentous fungi were used: *Candida albicans* OUT 6266, *Saccharomyces cerevisiae* ATCC 9804, *Aspergillus niger* ATCC 6275, *Rhyzopus oryzae* ATCC 10404 and *Geotrichum candidum* NBRC 4598. The following standard antibiotics were used as reference compounds: chloramphenicol (antibacterial) and amphotericin B (antifungal).

^aRecorded at 100 MHz. ^bRecorded at 400 MHz.

The producing strain was isolated from soil collected at the Tsuchiyu Hot Spring in Fukushima, Japan. This fungus was identified as E. ophioglossoides (previously known as Cordyceps ophioglossoides) on the basis of morphological criteria. HRFAB-MS of 1 revealed an ion [M+H]⁺ at m/z 390.2295, corresponding to the molecular formula C₂₂H₃₂NO₅ (calcd for C₂₂H₃₂NO₅, 390.2280), whereas 2 revealed an ion at m/z 374.2340, corresponding to the molecular formula C₂₂H₃₂NO₄ (calcd for C₂₂H₃₂NO₄, 374.2331). The ¹H NMR spectra of 1 and 2 were complicated by the occurrence of several subsets or broadenings of NMR signals, which supported the presence of a tetramic acid moiety because the ¹H NMR spectra of tetramic acids frequently display several tautomeric forms.^{6,7,12} Measurement of ¹H NMR at -80 °C slightly resolved the broadening of signals but significantly improved the measurement of ¹³C NMR of 1 (Supplementary Figures S1 and S2). As in the case of trichosetin, 6 the signals corresponding to C1, C2' and C4' could only be observed when measurement was taken at -80 °C. Henceforth, all NMR measurements with the exception of NOESY were taken at -80 °C. NOESY measurement at room temperature was satisfactory and, therefore, measurement at -80 °C was not required. Physical and spectroscopic data, including ¹H NMR, UV, mass spectral fragmentations and specific rotation of 2, were in complete agreement with or consistent with those of equisetin. ^{7,12,15} A careful comparison of the spectral data between 1 and 2 showed that compound 1 was a new analog of equisetin. Evaluation of the EI mass spectral fragmentation patterns of 1 and 2 suggested the addition of one oxygen atom in 1. The ions at m/z 170 (C₇H₈NO₄) that would be expected to arise from α -cleavage between the bridging carbonyl and hydrocarbon domain¹² were detected in both compounds, whereas the m/z 203 ion corresponding to the carbocyclic domain of 2 was not observed in 1. Instead, an m/z219 ion was observed, thereby suggesting the addition of one oxygen atom in the carbocyclic domain of 1. Interpretation of the ¹H, ¹³C and 2D NMR data of 1 (Table 1) showed that this compound differs from equisetin only by the hydroxylation of the C-16 methyl group attached to C-8 of the decalin moiety. This was supported by the absence of methyl doublets at δ 0.92 in the ¹H NMR spectrum and by the presence of a signal at δ 3.39, which is in agreement with the presence of a hydroxymethylene group. Furthermore, DEPT 135 analysis confirmed the absence of a methyl group (C-16), but the presence of a methylene group at δ 68.3 in 1 (Supplementary Figure S3). HMBC correlation between C-7, C-8, C-9 and C-16 further confirmed the proposed structure (Figure 2). Ha-9 showed ddd (*J*=12, 12, 12 Hz) on the basis of ${}^{2}J_{H,H}$ coupling with Hb-9 and ${}^{3}J_{H,H}$ couplings with H-8 and H-10. The latter coupling values (12, 12 Hz) indicate axial-axial relationships for H-8/Ha-9 and Ha-9/H-10 axial (Figure 3). The relative configuration of the decalin core of ophiosetin (1) was also determined by a NOESY experiment. The NOESY correlation peaks between Ha-9 and the bridge-head proton H-11, as well as between Ha-9, Hb-7 and H-16, indicate the *syn* relationship of these protons. The second bridgehead proton, H-6, which is located on the other side of the molecule, shows a correlation signal to the methyl group, H₃-12. No NOESY correlation can be detected between H-6 and H-11, which indicates a 6,11-trans ring fusion of the two six-membered rings. The cross peak observed between H-13 and H-15 indicates a 13E configuration. The NOESY-derived stereochemistry and conformation of ophiosetin is depicted in Figure 3. NOESY correlations established

that the relative configuration of the bicyclic subunit was identical to

that in equisetin. As the specific optical rotation ($[\alpha]^{22}_D$ -244°) is the

same in sign to that of equisetin, it has been tentatively assumed that

the absolute configuration in the bicyclic part of the molecule is also

the same as in equisetin. The absolute configuration of equisetin has

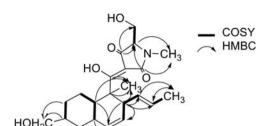


Figure 2 COSY and HMBC correlations of 1.

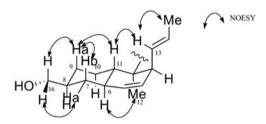


Figure 3 NOESY correlations of 1.

been previously reported, 12 which was also supported by total synthesis of the compound.¹⁷ The stereochemistry of phomasetin $([\alpha]^{22}_{D}+93.9^{\circ})$, an enantiomer homolog of equisetin, has also been described previously.⁷

The modification at the methyl group of C-16 has never been reported in any equisetin-related compound. To evaluate the effect of the hydroxyl group on the biological characteristics of 1, its antimicrobial activity was assayed. Equisetin (2) and all related compounds that have been tested so far show potent antibacterial activity against Gram-positive bacteria. 1-4 Consistent with previous reports, in our assay, compounds 2 and 3 exhibited antibacterial activity against Gram-positive S. aureus (MIC 4 µg ml⁻¹ for both 2 and 3) and E. faecalis (MIC $4 \mu g ml^{-1}$ for 2 and $2 \mu g ml^{-1}$ for 3). In addition, compounds 2 and 3 also showed moderate activity against Gramnegative E. coli (MIC 8 µg ml⁻¹). However, we could not detect any inhibitory activity of compound 1 even at a concentration of 128 µg ml⁻¹ against all other tested bacterial strains, except for a weak activity against E. faecalis (MIC 128 μg ml⁻¹). As for the antifungal activity, compound 3 exhibited weak activity (MIC $16 \,\mu \mathrm{g} \,\mathrm{ml}^{-1}$) against R. oryzae and A. niger, but compounds 1 and 2 did not show any antifungal activity at 32 μg ml⁻¹. This finding indicates that the modification at C-16 of the decalin moiety in compound 1 results in a drastic decrease in biological activity compared with that of equisetin, whereas it remains unclear at present whether the decreased biological activity is due to the more polar nature of 1, which may hinder the passage through the cell membrane or affinity toward the target molecules.

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Supplementary Information accompanies the paper on The Journal of Antibiotics website (http://www.nature.com/ja)