

Fumaquinone, a New Prenylated Naphthoquinone from *Streptomyces fumanus*

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Abstract A new prenylated naphthoquinone antibiotic, fumaquinone (5,7-dihydroxy-2-methoxy-3-methyl-6-(3-methyl-but-2-enyl)[1,4]naphthoquinone) was isolated from cultures of *Streptomyces fumanus* (LL-F42248). Its chemical structure was determined primarily by NMR spectroscopy. Preliminary feeding experiments indicated the naphthoquinone is of polyketide origin, while the *O*-methyl and aromatic *C*-methyl groups are derived from methionine.

Keywords prenylated naphthoquinone, actinomycete, fumaquinone, antibiotic, stable isotope labeling

Although terpenoidal compounds are commonly produced by plants and fungi, only a very limited number of isoprene-derived moieties or structures have been reported as metabolites of actinomycetes [1]. Selected examples of prenylated naphthoquinones include naphterpin [2], naphthgeranines [3], furaquinocins [4, 5], marinone [6], neomarinone [7] and the napyradiomycin [8] family of compounds. These compounds generally exhibit antitumor, antibiotic or antioxidative activity [1].

During our studies of culture LL-F42248, *Streptomyces fumanus*, that is known for producing chlorinated antibiotics [9, 10], we isolated fumaquinone (**1**), a new prenylated naphthoquinone. Here, we report the isolation and structure elucidation of **1**.

Characteristics of the producing organism and conditions

of its fermentation were described recently in a related publication [10]. As in the previous work, the polyamide resin together with the cell mass from 2 liter fermentation was extracted and purified by reversed-phase HPLC using acetonitrile/water. The fumaquinone rich fraction was then further purified by a second reversed-phase HPLC step using a linear gradient of 60~90% MeOH/H₂O with 0.01% TFA over 10 minutes to give **1** (2.7 mg, 15.1 minutes) that was characterized by standard spectroscopic methods.

Fumaquinone was obtained as an orange solid and the molecular formula was established as C₁₇H₁₈O₅ by high-resolution Fourier transform ion cyclotron resonance (HRFT-ICR) mass spectrometry (*m/z* 301.10835 [M–H]^{1–} calcd for C₁₇H₁₇O₅ 301.10815) which requires 9 degrees of unsaturation. Fumaquinone exhibited UV maxima at 224, 272, 308 and 428 nm (Fig. 2) that are characteristic for a 2-oxy-[1,4]naphthoquinone (juglone) chromophore [2~5].

The NMR spectra provided further support for a substituted naphthoquinone moiety. The ¹³C NMR (Table

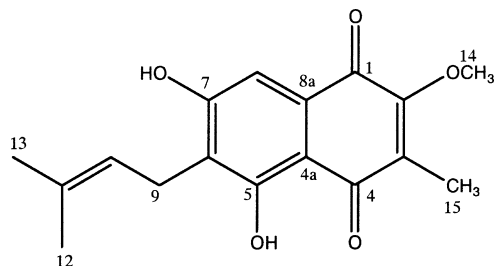


Fig. 1 Fumaquinone (**1**).

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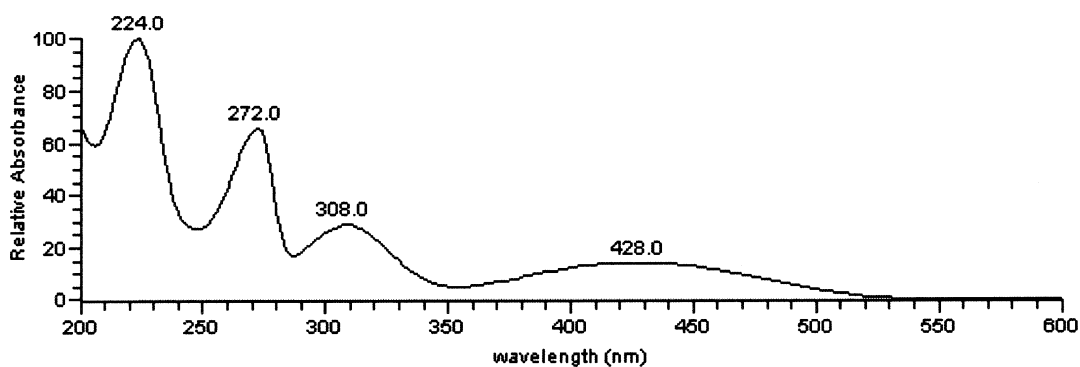


Fig. 2 UV spectrum of fumaquinone (**1**).

Table 1 NMR spectral data for **1** recorded in acetone- d_6 at 400 MHz

pos.	δ_C	δ_H (mult., $J=Hz$)	1H - 1H COSY	HMBC
1	181.0	—	—	—
2	158.9	—	—	—
3	131.6	—	—	—
4	191.0	—	—	—
4a	109.1	—	—	—
5	162.2 ^a	—	—	—
6	122.3	—	—	—
7	161.9 ^a	—	—	—
8	108.2	7.13 (s)	—	C-1, C-4a, C-6, C-7, C-8a
8a	131.7	—	—	—
9	22.7	3.39 (d, 7.4)	H-10	C-5, C-6, C-7, C-10, C-11
10	122.0	5.23 (t, 7.2)	H-9, H-12, H-13	C-9, C-12, C-13
11	132.7	—	—	—
12	18.0	1.78 (br s)	H-13	C-10, C-11, C-13
13	25.9	1.65 (br s)	H-12	C-10, C-11, C-12
14	61.3	4.07 (s)	—	C-2
15	8.8	1.99 (s)	—	C-2, C-3, C-4
5-OH	—	12.80 (s)	—	C-4a, C-5, C-6
7-OH	—	9.78 (br s)	—	—

^a Assignments can be interchanged

1) of **1** in acetone- d_6 displayed resonances for two carbonyls at δ 191.1 and 181.0 and ten olefinic and/or aromatic carbons between δ_C 108.2 and δ_C 162.2. The 1H NMR spectrum in acetone- d_6 showed nine resonances due to one aromatic proton (δ_H 7.13), one methoxy (δ_H 4.07), three methyls (δ_H 1.99, 1.78, 1.65), one methylene (δ_H 3.39) and one methine (δ_H 5.23). The HSQC data of **1** indicated that two hydrogen atoms were located on heteroatoms and were readily recognized by their chemical shifts as phenolic hydroxyl protons [5-OH (δ_H 12.80), and 7-OH (δ_H 9.78)].

The ^{13}C resonances at δ_C 191.1 and 181.0 and eight other

olefinic carbon resonances were assigned to the naphtho-1,4-quinone moiety based on the following correlations. A methine proton resonance at δ_H 7.13 was assigned to H-8 on the basis of strong HMBCs from H-8 to C-1 (δ_C 181.0), C-4a (δ_C 109.1), C-6 (δ_C 122.3), C-7 (δ_C 161.9) and C-8a (δ_C 131.7). Further, HMBCs from the methoxy protons at δ_H 4.07 to the olefinic carbon C-2 (δ_C 158.9) and from the allylic methyl at δ_H 1.99 to C-2, C-3 (δ_C 131.6) and C-4 were indicative of a methoxy and a methyl substituent on the naphthoquinone moiety. HMBC correlations from the hydroxyl proton at δ_H 12.80 to C-4a, C-5 (δ_C 162.2) and C-6 secured its position at C-5. Although the relatively broad

resonance of a hydroxyl proton at δ_{H} 9.78 did not show any HMBs, it was assigned to the C-7 hydroxyl group by exclusion which completed the assessment of the 5,7-dihydroxy-2-methoxy-3-methyl-[1,4]-naphthoquinone moiety.

The remaining resonances in ^1H and ^{13}C NMR spectra, correlated by ^1H - ^1H COSY, HSQC and HMBC data, readily identified the dimethyl allyl side chain as the remaining substituent. COSY correlations from H-9 at δ_{H} 3.39 to H-10 at δ_{H} 5.23 and allylic coupling between H-10 and H-12 (δ_{H} 1.78) as well as between H-10 and H-13 (δ_{H} 1.65) established the spin system from H-9 to H-13. The presence of the dimethyl allyl group was further corroborated by HMBC correlations from H-9 to C-10 (δ_{C} 122.0) and C-11 (δ_{C} 132.7) and from H-10 to C-9 (δ_{C} 22.7), C-12 (δ_{C} 18.0) and C-13 (δ_{C} 25.9). Mutual correlations between H-12 and C-13 and H-13 and C-12 further supported the assignment of the dimethyl allyl side chain. The dimethyl allyl side chain was placed on the naphthoquinone moiety at C-6 on the basis of HMBC correlations from H-9 to C-5 (δ_{C} 162.2), C-6 (δ_{C} 122.3) and C-7 (δ_{C} 161.9). The structure of **1** was thus shown to be 5,7-dihydroxy-2-methoxy-3-methyl-6-(3-methyl-but-2-enyl)[1,4]naphthoquinone.

Feeding experiments using L-[methyl- ^{13}C]methionine indicated that the methoxy (C-14) and the aromatic methyl (C-15) groups were derived from methionine with isotopic enrichment of >85% for each carbon. Preliminary work with [1,2- $^{13}\text{C}_2$]sodium acetate indicated that the biosynthesis of the juglone moiety follows the polyketide pathway, a finding similar to that of the furaquinocins [11] which share the juglone moiety.

Fumaquinone is the first prenylated naphthoquinone isolated from *S. fumanus*. It showed antimicrobial activity against selected Gram-positive bacteria with MIC of about 64 $\mu\text{g}/\text{ml}$.

Experimental

General Experimental Procedures

All 1D and 2D NMR spectra were recorded on a Bruker Avance DPX-400 spectrometer at 400 and 100 MHz for ^1H and ^{13}C , respectively, using a 3 mm broadband detect probe. Proton detected heteronuclear correlations were measured using HSQC (optimized for $^1J_{\text{C-H}}=140$ Hz) and HMBC (optimized for $^nJ_{\text{C-H}}=8.3$ Hz) pulse sequences. UV data were obtained on HP1100 HPLC system equipped with DAD. High resolution mass spectra (HRMS) were obtained using a Bruker (Billerica, MA) APEXII FTICR mass spectrometer equipped with an actively shielded 7.1 Tesla

superconducting magnet (MagneX Scientific Ltd., UK), an external Bruker APOLLO ESI source, and a Synrad 50W CO_2 CW laser. The molecular ion at m/z 301 was isolated using correlated sweep and then dissociated using Infrared Multi-Photon Dissociation (IRMPD). All HPLC solvents were EM Omnisolv quality and used without further purification.

Isolation and Purification Procedures

The harvested fermentation broth (2 liters) was centrifuged at 4000 rpm for 20 minutes and the polyamide resin together with the cell mass was extracted with acetone (2 \times 200 ml). The combined acetone extract was concentrated *in vacuo* and the aqueous suspension extracted with ethyl acetate (3 \times 60 ml). The ethyl acetate extract (1.5 g) was purified by reversed-phase HPLC (YMC ODS-A column, 20 \times 250 mm, 5 μm) using a gradient of 40~90% acetonitrile/water containing 0.05% TFA in water over 40 minutes at a flow rate of 7 ml/minute. Final purification of the fumaquinone rich fraction (10 mg) was achieved by reversed-phase HPLC (YMC ODS-A column, 10 \times 250 mm, 5 μm) using a gradient of 60~90% methanol/water with 0.02% TFA in water over 10 minutes and maintaining for 5 minutes at a flow rate of 4 ml/minute to give fumaquinone (**1**, 2.7 mg, 15.1 minutes) as an orange solid.

L-[methyl- ^{13}C]methionine Feeding Experiment

A solution of L-[methyl- ^{13}C]methionine at a final concentration of 10 mM was added to the production medium described previously [10]. The fermentation was harvested at five days and processed as described above.

[1,2- $^{13}\text{C}_2$]sodium Acetate Feeding Experiment

[1,2- $^{13}\text{C}_2$]sodium acetate was added to the production medium described previously [10] at a final concentration of 10 mM. The fermentation was harvested at five days and processed as described above.

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References

1. Kuzuyama T, Seto H. Diversity of the biosynthesis of the isoprene units. *Nat Prod Rep* 20: 171–183 (2003)
2. Shin-Ya K, Imai S, Furihata K, Hayakawa Y, Kato Y, Vanduyne GD, Clardy J, Seto H. Isolation and structural elucidation of an antioxidative agent, naphterpin. *J Antibiot* 43: 444–447 (1990)

3. Wessels P, Gohrt A, Zeeck A, Drautz H, Zahner H. Metabolic products of microorganisms, 260. Naphthgeranines, new naphthoquinone antibiotics from *Streptomyces* sp. *J Antibiot* 44: 1013–1018 (1991)
4. Funayama S, Ishibashi M, Anraku Y, Komiyama K, Ōmura S. Structures of novel antibiotics, furaquinocins A and B. *Tetrahedron Lett* 30: 7427–7430 (1989)
5. Ishibashi M, Funayama S, Anraku Y, Komiyama K, Ōmura S. Novel antibiotics, furaquinocins C, D, E, F, G and H. *J Antibiot* 44: 390–395 (1991)
6. Pathirana C, Jensen PR, Fenical W. Marinone and debromomarinone: Antibiotic sesquiterpenoid naphthoquinones of a new structure class from a marine bacterium. *Tetrahedron Lett* 33: 7663–7666 (1992)
7. Kalaitzis JA, Hamano Y, Nilsen G, Moore BS. Biosynthesis and structural revision of neomarinone. *Org Lett* 5: 4449–4452 (2003)
8. Fukuda DS, Mynderse JS, Baker PJ, Berry DM, Boeck LD, Yao RC, Mertz FP, Nakatsukasa WM, Mab J, Ott J, Counter FT, Ensminger PW, Allen NE, Alborn WE, Hobbs JN. A80915, A new antibiotic complex produced by *Streptomyces aculeolatus*. Discovery, taxonomy, fermentation, isolation, characterization, and antibacterial evaluation. *J Antibiot* 43: 623–633 (1990)
9. Carter GT, Nietsche JA, Goodman JJ, Torrey MJ, Dunne TS, Borders DB, Testa RT. LL-F42248 α , a novel chlorinated pyrrole antibiotic. *J Antibiot* 40: 233–236 (1987)
10. Charan RD, Schlingmann G, Bernan VS, Feng X, Carter GT. Additional pyrrolomycins from cultures of *Streptomyces fumanus*. *J Nat Prod* 68: 277–279 (2005)
11. Funayama S, Ishibashi M, Komiyama K, Ōmura S. Biosynthesis of furaquinocins A and B. *J Org Chem* 55: 1132–1133 (1990)