

Serialynic Acid, a New Phenol with an Isopentenynyl Side Chain from *Antrodia serialis*

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Abstract A new phenolic compound serialynic acid was isolated from an agar culture of the basidiomycete *Antrodia serialis*, through bioactivity-guided fractionations. It showed weak growth inhibitory activity towards phytopathogenic fungi and a dose-independent anti-*Pythium graminicola* activity.

Keywords *Antrodia serialis*, serialynic acid, isopentenynylphenol, antifungal activity, Basidiomycete

A. serialis is a brown rot fungus mainly growing on coniferous wood and constitutes one of the major timber- and woodwork-degrading fungi. A large number of triterpenoids of the lanostane and ergostane type, and simple phenolic metabolites have been isolated from the fruiting bodies of *A. cinnamomea*, which is used in Chinese folk medicines [1–6], and cytotoxic pyrrole derivatives have been found in the cultured mycelia of *A. camphorata* [7]. We report here the isolation and characterisation of **1** from *A. serialis*.

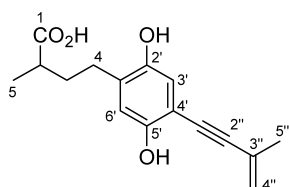
Introduction

During a systematic screening for antimicrobial and/or novel compounds from a collection of British fungi, a strain of *Antrodia serialis* (Fr.:Fr.) Donk (Coriolaceae), when cultured, was found to produce an antifungal compound in large amounts. Subsequent culturing of this strain and bioactivity-guided fractionations led to the isolation of serialynic acid (**1**).

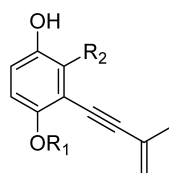
Experimental

General

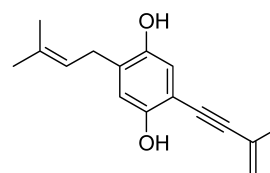
UV spectra were obtained with a Shimadzu UV-1601 spectrophotometer. NMR spectra were recorded on a Bruker Avance 400 spectrometer in CD₃OD at 30°C, with TMS as the internal standard (400 MHz for proton, and 100 MHz for carbon). Mass spectra were recorded using a



Serialynic acid (**1**)



Siccayne (**2**) $R_1=H$, $R_2=H$
Eulatinol (**3**) $R_1=CH_3$, $R_2=H$
Frustulosinol (**5**) $R_1=H$, $R_2=CH_2OH$



Culpin (**4**)

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Thermo Finnigan LCQ Classic in both positive and negative atmospheric pressure chemical ionisation (APCI) modes. TLC was run on silica gel F₂₅₄ precoated plates (Merck 5554) with CHCl₃-MeOH-H₂O, 73:24:3 (in volume), and spots were detected by UV light and bioautography [8] (see below). Final purification of **1** was carried out using a Waters HPLC system consisting of a 600 pump, a 717 plus autosampler and a 996 photodiode array detector. The column used was Genesis C₁₈ (Jones Chromatography, 10 mm i.d.×250 mm, *dp* 4 µm) at 30°C, eluting with a linear gradient of 40 to 100% MeOH in H₂O over 20 minutes at a flow rate of 4 ml/minute.

Fungal Strain, Fermentation and Isolation

The fruiting body was found growing on a windowsill near Fishguard, Wales, UK, in November, 1995 and identified by Dr. Peter Roberts of the Mycology Section, Jodrell Laboratory, Royal Botanic Gardens, Kew. The dried specimen was deposited in the mycological herbarium of the Gardens (K(M)33565). The mycelial culture (strain KC920) raised from uncontaminated tissue of the same specimen was maintained on a malt extract-agar medium under oil at 16°C.

The seed culture was grown on a medium containing malt extract (Oxoid L39, 2%) and agar (1.5%) at 25°C. The well grown culture at approx. 4 weeks old was cut into

small pieces and transferred (3-point inoculation) to forty 9 cm-diameter Petri dishes containing the same medium, and grown at 25°C for 24 days with artificial illumination (white light, 4×40 W).

The whole culture was extracted twice with CHCl₃-MeOH (2:1, v/v). The combined organic phase was concentrated *in vacuo* at 40°C, and the residue (2.81 g) was chromatographed over silica using mixtures of CHCl₃-MeOH-H₂O (100:0:0, 87:12:1, 73:24:3, 16:9:1 and 5:5:1, in volume). The fractions with antifungal activity against the test strain *Cladosporium herbarum* IMI300461 were combined (82.3 mg) and further purified by reverse phase HPLC. Serialynic acid eluted at 14.3 minutes, yielding 8.3 mg of pure compound (**1**).

The chemical structure of **1** was determined mainly by one-dimensional proton and carbon NMR at 400 and 100 MHz, respectively, and ¹H-¹H COSY, HSQC and HMBC experiments.

Physico-chemical Properties

Serialynic acid (4-[2,5-dihydroxy-4-(3-methylbut-3-en-1-ynyl)phenyl]-2-methylbutanoic acid, **1**) was obtained as a pale yellow needles from MeOH-H₂O. Mp (uncorr.) 102~104°C. APCI-MS (pos. mode) *m/z* 275 [M+H]⁺, 257, 187, 159; (neg. mode) *m/z* 547 [2M-H]⁻, 273 [M-H]⁻, 255, 229, 174. UV λ_{max} (MeOH) nm (log ε) 209

Table 1 ¹H (400 MHz) and ¹³C NMR (100 MHz) data, and HMBC correlations of serialynic acid (**1**) in CD₃OD (δ in ppm, *J* in Hz)

Position	δ (¹ H)	δ (¹³ C)	¹ H- ¹ H COSY	HMBC
1		181.0		
2	2.42 (1H, m)	40.6	H-3, H-5	C-1, 3, 4, 5
3	1.91 (1H, m)	35.0	H-3, H-4	C-1, 2, 4, 5, 1'
	1.66 (1H, m)		H-3, H-4	C-1, 2, 4, 5, 1'
4	2.56 (2H, t, 7.7)	29.2	H-3	C-2, 3, 1', 2'
5	1.18 (3H, d, 7.0)	17.8	H-2	C-1, 2, 3
1'		132.5		
2'		149.2		
3'	6.67 (1H, s)	119.3		C-4, 1', 2', 4', 5', 6', 1''
4'		109.6		
5'		152.2		
6'	6.59 (1H, br s)	118.0	H-4	C-3, 4, 1', 2', 3', 4', 5', 1'', 2''
1''		86.1		
2''		94.6		
3''		128.9		
4''	5.32 (1H, m)	121.4	H-4''	C-2'', 3'', 5''
	5.25 (1H, m)		H-4'', H-5''	C-2'', 5''
5''	1.96 (3H, dd, 1.6, 1.0)	23.8	H-4''	C-2'', 3'', 4''

Chemical shifts are shown with reference to TMS as the internal standard.

Table 2 Antifungal activity of serialynic acid (**1**)^a

Microorganisms	Concentration in $\mu\text{g/ml}$ and in (μM)		
	70 (255)	140 (511)	280 (1022)
<i>Botrytis cinerea</i>	— ^b	—	—
<i>Fusarium graminearum</i>	—	—	—
<i>Pyrenophora teres</i>	—	—	++ ^b
<i>Rhizoctonia solani</i>	—	—	—
<i>Septoria tritici</i>	—	—	—
<i>Stagonospora nodorum</i>	—	—	+ ^b
<i>Pythium graminicola</i>	+	+	+

^a Test compounds were dissolved in DMSO/isopropyl alcohol and Tween 20, and dispersed into minimal medium containing a calibrated number of spores. After incubation period (4–6 days depending on the test organisms) the growth of the test strains were visually assessed and given scores.

^b Score —: no inhibition compared to the control; ++: arbitrary level of inhibition compared to the control; +++: complete inhibition.

(4.4), 265 (4.22), 279 (4.24), 328 (4.1). ¹H (400 MHz, CD₃OD) and ¹³C (100 MHz, CD₃OD) NMR, see Table 1.

Biological Assays

The fungus *C. herbarum* IMI300461 was used as the test strain to trace the activity on the TLC bioautography [8]. The growth inhibitory activity of the compound was tested against a panel of phytopathogenic organisms (listed in Table 2), using 96-well microplates in a series of concentrations (70, 140 and 280 $\mu\text{g/ml}$).

Results and Discussion

The UV spectrum of **1** was virtually identical with that of siccayne (**2**) and eulatinol (**3**) [9], indicating the presence of a chromophore similar to 3-methyl-3-penten-1-ynyl substituted hydroquinone. The positive APCI-mass spectrum showed a protonated ion peak at 275 $[\text{M}+\text{H}]^+$, while 273 $[\text{M}-\text{H}]^-$ and 547 $[2\text{M}-\text{H}]^-$ were major ions in the negative mode, indicating the relative molecular mass to be 274. The high-resolution ESI MS of **1** gave a pseudo-molecular ion peak at m/z 292.1541 ($[\text{M}+\text{NH}_4]^+$, calcd. for C₁₆H₂₂NO₄, 292.1543, $\Delta=0.8$ ppm), corresponding to a molecular formula of C₁₆H₁₈O₄.

The ¹³C NMR spectrum revealed the presence of nine sp^2 carbons, including one carboxylic acid at δ 181.0 and two oxygenated ones (δ 152.2 and 149.2). Two sp carbons at δ 94.6 and 86.1 ppm were also detected. From the ¹H-¹H COSY, HSQC and HMBC spectra, and by comparison with literature data [9, 10], a partial structure corresponding to siccayne (**2**) was deduced. Despite the difference in

the solvents used, the ¹³C chemical shift values of the isopentenynyl moiety and the aromatic carbons 2' through to 5' were practically identical to the literature values measured in CDCl₃ [9, 10]. Serialynic acid was hardly soluble in CHCl₃ but more soluble in MeOH or Me₂CO. The two sp^2 carbons directly adjacent to the acetylenic bonds *i.e.* C-4' (δ 109.6) and C-3'' (δ 128.9) showed correlations in the HMBC spectrum with H-6' and H-4'' and -5'', respectively; therefore the chemical shift assignments given for the corresponding carbons in support of the structure of foeniculoxin [11] should be interchanged.

The remaining fragment consisting of C₅H₉O₂ was readily identified as 2-methylbutanoic acid through interpretation of the ¹H-¹H COSY, HSQC and HMBC data. This substituent was placed at the *para*-position to the isopentenynyl group on the basis of HMBC correlations between H-3' (δ 6.67) and C-4 (δ 29.2), and that no coupling was observed between the two aromatic protons (δ 6.59 and 6.67). This conclusion was further corroborated by the weak five-bond (⁵ J_{HC}) HMBC correlation observed from H-6' to C-2''. Similar long range couplings *via* conjugated triple and double bonds have been recorded in the literature [12]. The structure of **1** was therefore determined to be 4-[2,5-dihydroxy-4-(3-methylbut-3-en-1-ynyl)phenyl]-2-methylbutanoic acid. The absolute configuration of the single stereogenic centre has not been determined.

Serialynic acid belongs to a small group of simple phenolic compounds bearing isopentenynyl substructures, all of which have been isolated from fungi of both ascomycetes [9, 13, 14] and basidiomycetes [10, 15–17],

in addition to a hyphomycete [18]. They have variably been shown to have weak to no antimicrobial and phytotoxic activities [9, 10, 13, 14, 16–19]. Of these **1** possesses the closest structural resemblance to culpin isolated from *Preussia* sp. which showed a weak antimicrobial activity [14].

Serialynic acid was tested against several standard in-house test organisms at the Jealott's Hill International Research Centre, and shown to have weak to no activity against the representative strains of phytopathogenic fungi (Table 2). Interestingly, the straminipile *Pythium graminicola* was found most susceptible to the compound, and its growth was retarded at the lowest concentration tested (70 µg/ml). Frustulosinol **5** [15], isolated from a culture of *Gloeocystidiellum porosum* (Berk. & M.A. Curtis) Donk (Stereaceae) strain KC1004 (unpublished result), was devoid of activity in the assays performed against the same test organisms concurrently.

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