

5,6-DIMETHOXYSTERIGMATOCYSTIN, A NEW METABOLITE FROM *ASPERGILLUS MULTICOLOR*

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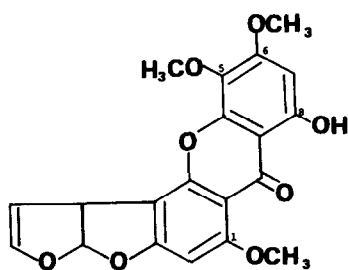
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(Received in Japan 23 May 1977; received in UK for publication 20 June 1977)

In the course of our investigation of fungal metabolites we have isolated a new metabolite containing a bisdihydrofuran ring system from mycelial mats of *Aspergillus multicolor* Sappa together with sterigmatocystin and averufin. We wish to report the structure of the new metabolite.

The metabolite (I), $C_{20}H_{16}O_8$, was crystallized from acetone to give pale yellow needles, mp 253-254°C, $[\alpha]_D^{20} -363^\circ$ ($c=1.0$, $CHCl_3$), m/e 384 (M^+). IR ν_{max}^{KBr} cm^{-1} : 3125, 1655, 1635, 1610. UV λ_{max}^{EtOH} nm (ϵ): 233(27200), 248(34000), 275sh(7700), 330(19200). Its UV spectrum is closely similar to that of sterigmatocystin. In the NMR spectrum of I in $CDCl_3$ signals at δ 4.80(1H, d.d., $J=2, 2.5$ and 7 Hz), 5.41(1H, d.d., $J=2$ and 2.5 Hz), 6.46(1H, d.d., $J=2$ and 2.5 Hz) and 6.75(1H, d, $J=7$ Hz) showed the presence of a bisdihydrofuran ring system in I. Three methoxyl



(I)

signals were observed at δ 3.91, 3.94 and 3.98. Two singlet signals at δ 6.28(1H) and 6.32(1H) were assignable to the aromatic protons of xanthone nucleus. Hydrogenation of I gave a dihydro-derivative (II), mp 241-242°C. In the NMR spectrum of I a singlet signal at δ 13.20 was assignable to the hydrogen-bonded proton of phenolic hydroxyl group. Acetylation of I gave a monoacetate (III), mp 188-189°C.

Methylation of I gave a monomethyl ether (IV), mp 257-258°C. Demethylation of II gave a dihydroxy-derivative (V), mp 238-240°C, whose NMR spectrum showed two singlet signals at δ 11.90 and 12.18 assignable to the protons of hydrogen-bonded hydroxyl groups. Alkaline degradation of I with 15% potassium hydroxide in ethanol gave a hydroxy-xanthone (VI), mp 221-222°C. Its NMR

spectrum showed three signals at δ 7.15(1H, m), 7.66(1H, m) and 8.17(1H, m) assignable to the protons of β -substituted furan ring. Methylation of VI gave a dimethyl ether (VII), mp 225-226°C. An aromatic proton at δ 6.32 in I showed downfield shift (0.18 ppm) on acetylation of the hydroxyl group. This is characteristic of the proton *ortho* to a phenolic hydroxyl group.

The locations of three methoxyl groups and the bisdihydrofuran ring in I were inferred from NMR spectra. Three methoxyl signals of I in C_6D_6 showed upfield shift (δ 3.20, 3.25, 3.73). Two strongly shifted signals at δ 3.20 and 3.25 were assignable to the protons of methoxyl groups *ortho* to the aromatic protons. In the NMR spectrum of IV a newly appeared methoxyl group signal at δ 3.93 in $CDCl_3$ was shifted upfield by 0.43 ppm in C_6D_6 ; this shift value agreed with that reported for 8-*O*-methylsterigmatocystin.¹⁾ This fact indicates the presence of a hydroxyl group at the position 8 in I. These support that a hydroxyl group locates on an aromatic nucleus not bearing the bisdihydrofuran ring system. In the NMR spectrum of VII in C_6D_6 the upfield shifted four methoxyl signals were observed at δ 3.18(3H), 3.25(3H) and 3.46(3Hx2). Based on the above-mentioned data, the assumed structure (I) is deduced to the metabolite.

The structure of monoacetate (III) has confirmed by a single-crystal X-ray analysis. The crystals were monoclinic, space group C2, with $z=4$ in a cell of dimensions $a=19.76$, $b=5.09$, $c=20.41$ Å, $\beta=105.7^\circ$. The structure was solved by Patterson and Fourier methods and refined by

least-squares method to $R=0.135$.

A stereoscopic view of monoacetate (III) is shown in Fig.1. The two asymmetric centers in figure are based on absolute configuration of *p*-bromobenzoate of sterigmatocystin.²⁾

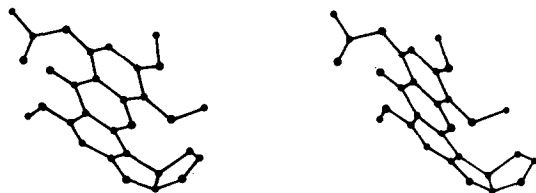


Fig. 1

The authors wish to thank Mr. Norio Nakajima for the assistance of X-ray analysis, and Mr. Isao Tanaka and Prof. Tamaichi Ashida of Nagoya University for their permission to use the computer programs.

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