

Structure Elucidation of the Fumonisin, Mycotoxins from *Fusarium moniliforme*

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The structures of the fumonisins, a family of structurally related mycotoxins isolated from cultures of *Fusarium moniliforme*, were elucidated by mass spectrometry and ¹H and ¹³C n.m.r. spectroscopy as the diester of propane-1,2,3-tricarboxylic acid and either 2-acetyl-amino- or 2-amino-12,16-dimethyl-3,5,10,14,15-penta-hydroxyicosane as well as in each case the C-10 deoxy analogue; in all cases both the C-14 and C-15 hydroxy groups are involved in ester formation with the terminal carboxy group of propane-1,2,3-tricarboxylic acid.

The fungus *Fusarium moniliforme* Sheldon occurs world-wide on a variety of plant hosts and is one of the most important ear rot pathogens of maize (*Zea mays* L.).¹ An isolate of *F. moniliforme* (strain MRC 826) obtained from maize in an area of the Transkei with a high oesophageal cancer rate² was found to be not only highly toxic to experimental animals and able to induce leukoencephalomalacia in horses,³ but also hepatocarcinogenic to rats,⁴ and mutagenic to *Salmonella typhimurium*.⁵ Although *F. moniliforme* (strain MRC 826) produces the mutagen fusarin C,⁶ this compound is apparently not involved in the carcinogenicity of the fungus. In continuation of our efforts towards solving this important problem, we now report on the isolation and structure elucidation of a family of related mycotoxins, the fumonisins, from *F. moniliforme*.

Extraction of cultures of *F. moniliforme* (strain MRC826)

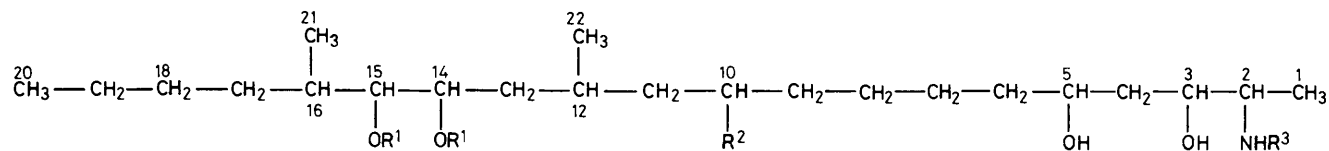
grown on sterilized maize with aqueous methanol gave an extract which was active in rat liver histopathological and cancer initiation/promotion bioassays.⁷ Subsequently, extensive fractionation of this extract, using macroreticular polystyrene resin (XAD-2), Sephadex LH-20 and reversed phase silica gel chromatography, and guided by the assays mentioned, resulted in the isolation of a mixture of fumonisin A₁ (1) and A₂ (2), as well as fumonisin B₁ (3) and B₂ (4). Treatment of the mixture of fumonisin A₁ and A₂ with an excess of diazomethane, followed by chromatography on silica gel (chloroform-methanol, 92:8 v/v) gave colourless oils of the tetramethyl ester derivatives (5) and (6), respectively.

Liquid secondary ion mass spectrometry (l.s.i.-m.s.) of tetramethylfumonisin A₁ (5), ν_{\max} . (CHCl₃) 1730 (ester CO) and 1650 (amide CO) cm⁻¹, gave the protonated molecular ion as m/z 820 ($M + H$)⁺, supported by an ion at m/z 842 (M

+ Na)⁺⁺. The multiplicities of the different resonances in the ¹³C n.m.r. spectrum of (5) were determined by generating the proton-decoupled CH, CH₂, and CH₃ subspectra using the DEPT pulse sequence.⁸ These data (see Table 1) suggested the empirical formula C₄₀H₆₉NO₁₆ for (5) which is consistent with the accurate mass measurement of the ion of highest mass in the electron impact (e.i.) mass spectrum at *m/z* 770.4321 (C₃₉H₆₄NO₁₄) [*M* - (H₂O + CH₃O)⁺]. The presence of three hydroxy groups in (5) was indicated by the formation of a tris(trimethylsilyl) derivative (*M*⁺ 1035, C₄₉H₉₃NO₁₆Si₃) on treatment with *N*-methyl-*N*-trimethylsilyl trifluoroacetamide and a triacetate (*M*⁺ 945, C₄₆H₇₅NO₁₉) on acetylation. Basic

hydrolysis with 0.05 M methanolic potassium hydroxide for 16 h at room temperature gave a neutral product (7), C₂₄H₄₉NO₆, and a salt which on acidification and methylation with an excess of diazomethane was converted into a product identical (g.c.-m.s.) to trimethylpropane-1,2,3-tricarboxylate.

The proton-proton connectivity pattern of tetramethyl fumonisin A₁ (5) was determined by two-dimensional (2D) (¹H, ¹H) correlation spectroscopy using the COSY-45 pulse sequence,⁹ and defined two distinct structural units, *i.e.* C-1—C-5 and C-10—C-17. The involvement of both the C-14 and C-15 oxygen atoms in ester linkages with the terminal



- (1) R¹ = COCH₂CH(CO₂H)CH₂CO₂H, R² = OH, R³ = C²³OC²⁴H₃
 (2) R¹ = COCH₂CH(CO₂H)CH₂CO₂H, R² = H, R³ = COCH₃
 (3) R¹ = COCH₂CH(CO₂H)CH₂CO₂H, R² = OH, R³ = H
 (4) R¹ = COCH₂CH(CO₂H)CH₂CO₂H, R² = R³ = H
 (5) R¹ = COCH₂CH(CO₂CH₃)CH₂CO₂CH₃, R² = OH, R³ = COCH₃
 (6) R¹ = COCH₂CH(CO₂CH₃)CH₂CO₂CH₃, R² = H, R³ = COCH₃
 (7) R¹ = H, R² = OH, R³ = COCH₃

Table 1. N.m.r. data for tetramethylfumonisin A₁ (5).^a

Carbon atom	δ _C ^b	δ _H ^{c,j}	<i>J</i> (HH)/Hz		Carbon atom	δ _C ^b	δ _H ^c	<i>J</i> (HH)/Hz
1	18.03 Q	1.139 d	6.8	C-15 ester: R ¹ = C ²⁵ OC ²⁶ H ₂ C ²⁷ H(C ³⁰ O ₂ C ³¹ H ₃)C ²⁸ H ₂ C ²⁹ O ₂ C ³² H ₃	25	170.88 S	—	
2	49.72 D	3.902 m			26	35.27 T ^h	2.780 dd ^h	16.7, 6.7
3	71.49 D	3.79 m			27	37.28 D ^f	2.603 dd	16.8, 6.4
4	40.22 T	1.56			28	35.10 T ^h	3.22 m	
5	68.83 D	3.82			29	173.41 S	2.732 dd ^h	16.6, 7.1
6	38.55 T ^d	1.43			30	171.69 S ^g	2.583 dd	16.7, 6.5
7	25.63 T ^e	1.37			31	52.31 Q ⁱ	—	
8	25.48 T ^e	0.96			32	51.82 Q ⁱ	3.664 s ⁱ	3.645 s ⁱ
9	37.22 T ^d	1.75						
10	68.78 D	1.64						
11	43.08 T	1.28						
12	25.17 D	5.137 ddd	11.3, 3.3, 2.3					
13	35.39 T	4.867 dd	8.8, 3.3					
14	71.24 D	1.58						
15	77.79 D	1.31		33	171.42 S	—		
16	33.64 D	0.98		34	35.10 T ^h	2.672 dd	16.8, 8.1	
17	31.84 T	1.23				2.400 dd	16.8, 5.9	
18	28.48 T	1.08		35	37.19 D ^f	3.22 m		
19	22.73 T	~1.2		36	34.93 T ^h	2.647 dd ^h	16.7, 6.5	
20	13.92 Q	0.829 t	7.1			2.583 dd	16.7, 6.5	
21	15.36 Q	0.872 d	6.8	37	173.93 S	—		
22	20.36 Q	0.912 d	6.6	38	171.66 S ^g	—		
23	170.48 S	—		39	52.15 Q ⁱ	3.645 s ⁱ		
24	23.32 Q	1.949 s		40	51.80 Q ⁱ	3.634 s ⁱ		

^a Recorded on a Bruker WM-500 spectrometer. ^b Relative to CDCl₃ at δ_C 77.00. Chemical shifts of the proton-bearing carbon atoms were correlated with specific proton resonances.^{9,11} ^c Relative to CDCl₃ at δ_H 7.24. ^{d-i} May be interchanged. ^j The NH proton appears as a doublet at δ_H 6.071 [*J*(NH, 2-H) 8.7 Hz].

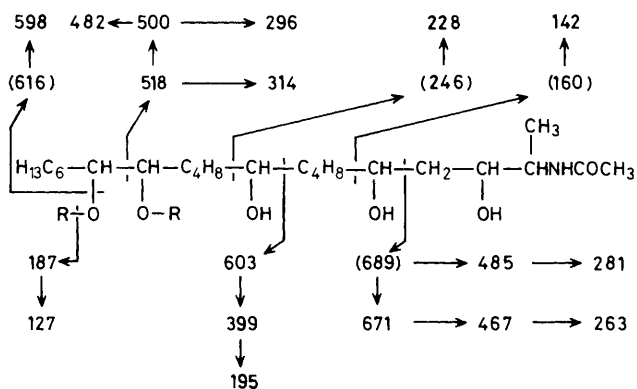


Figure 1. Fragmentation pattern observed in the e.i. mass spectrum of (5).

carboxy groups of the two propane-1,2,3-tricarboxylate moieties (as indicated by the ^1H and ^{13}C n.m.r. data)¹⁰ and the location of the methyl groups in (5) followed from the two- and three-bond (^{13}C , ^1H) connectivity pattern (Table 2) determined in a chemical shift correlation experiment adjusted to detect correlations *via* long-range (^{13}C , ^1H) couplings (nJ 5 Hz).^{9,11}

The fragmentation pattern for compound (5) (see Figure 1) in the electron impact mass spectrum was of vital importance in determining the linkage of the C-1—C-5 and C-10—C-17 structural fragments by a C_4H_8 group and placing a C_6H_{13} substituent at C-15. The base peak in the mass spectrum occurs at m/z 187 ($\text{C}_8\text{H}_{11}\text{O}_5$) and is associated with the cleavage of the ester moiety as indicated. The loss of 204 mass units from a number of fragment ions throughout the spectrum is due to the loss of a dimethyl propane-1,2,3-tricarboxylate group. The location of the hydroxy groups in (5) as indicated by the n.m.r. data is consistent with the fragment ions at m/z 671, 603, 228, and 142 and was confirmed by the fragmentation pattern of the tris(trimethylsilyl) derivative.

Tetramethylfumonisins A_2 (6), $\text{C}_{40}\text{H}_{69}\text{NO}_{15}$, contains one oxygen atom less than tetramethylfumonisins A_1 (5). The protonated molecular ion of (6) appears at m/z 804 ($M + \text{H}$)⁺ in the l.s.i. mass spectrum and the fragment ion which arises from the molecular ion in the e.i. spectrum through loss of ($\text{H}_2\text{O} + \text{CH}_3\text{O}^+$) appears at m/z 754 ($\text{C}_{39}\text{H}_{64}\text{NO}_{13}$). A comparison of the fragmentation patterns of compounds (6) and (5) shows that it is the C-10 hydroxy group of (5) which is replaced by a hydrogen atom in (6) as the bond cleavage which generates the fragment ions m/z 603 \rightarrow 399 \rightarrow 195 in the spectrum of (5) does not occur. In addition a number of bond cleavages common to both compounds generate fragment ions of 16 mass units less for tetramethylfumonisins A_2 (6).

L.s.i. mass spectrometry of fumonisins B_1 (3) ($\text{C}_{34}\text{H}_{59}\text{NO}_{15}$) gave the protonated molecular ion as m/z 722 whereas that of fumonisins B_2 (4) ($\text{C}_{34}\text{H}_{59}\text{NO}_{14}$) appears at m/z 706. In each case ions are also present at m/z values corresponding to ($M + \text{Na}$)⁺. A comparison of the ^{13}C n.m.r. data of these two compounds† indicates that once again it is the C-10 hydroxy group of fumonisins B_1 (3) (δ_{C} 67.17) which is replaced by a hydrogen atom in fumonisins B_2 (4). In addition, the ^1H and

Table 2. Two- and three-bond (^1H , ^{13}C) connectivity pattern for tetramethylfumonisins A_1 (5).

δ_{H}	Correlation signals δ_{C} /p.p.m.
0.829 (20-H)	22.73 T (C-19), 28.48 T (C-18)
0.872 (21-H)	31.84 T (C-17), 33.64 D (C-16)
	77.79 D (C-15)
0.912 (22-H)	25.17 D (C-12), 35.39 T (C-13)
1.139 (1-H)	49.72 D (C-2), 71.49 D (C-3)
1.949 (24-H)	170.48 S (C-23)
4.867 (15-H)	33.64 D (C-16), 71.24 D (C-14), 170.88 S (C-25)
5.137 (14-H)	77.79 D (C-15), 171.42 S (C-33)

^{13}C resonances of the atoms of the *N*-acetyl group present in the fumonisins A metabolites, e.g. (5): δ_{H} 1.949; δ_{C} 170.48, and 23.32, are absent from the spectra of the fumonisins B compounds.

It is of interest to note that a related mono-ester of propane-1,2,3-tricarboxylic acid and 1-amino-11,15-dimethyl-2,4,5,13,14-pentahydroxyheptadecane is a host-specific phytotoxin produced by *Alternaria alternata* (Fr.) Keissler f. sp. *lycopersici*, the causal agent of stem canker disease of tomato.^{10,12}

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† (3): δ_{C} (CD_3SOCD_3) 175.91S, 175.60S, 174.31S, 173.97S, 171.45S (C-25), 171.39S (C-33), 76.48D (C-15), 70.35D (C-14), 68.42D (C-13), 67.17D (C-10), 66.01D (C-5), 51.69D (C-2), 43.28T (C-11), 40.72T (C-4), 38.17D (C-27 and C-35), 38.19T, 38.07 (2 \times T), 37.80T (C-26), 36.54T (C-28), 36.38T (C-34), 35.72T (C-36), 32.96D (C-16), 30.96T (C-17), 28.08T (C-18), 25.43T, 25.28T, 25.20D (C-12), 22.22T (C-19), 20.19Q (C-22), 15.57Q (C-1), 15.51Q (C-21), 13.86Q (C-20).