

Biosynthesis of Aflatoxins. Incorporation of [2-²H₃]Acetate into Aflatoxin B₁ by *Aspergillus flavus*

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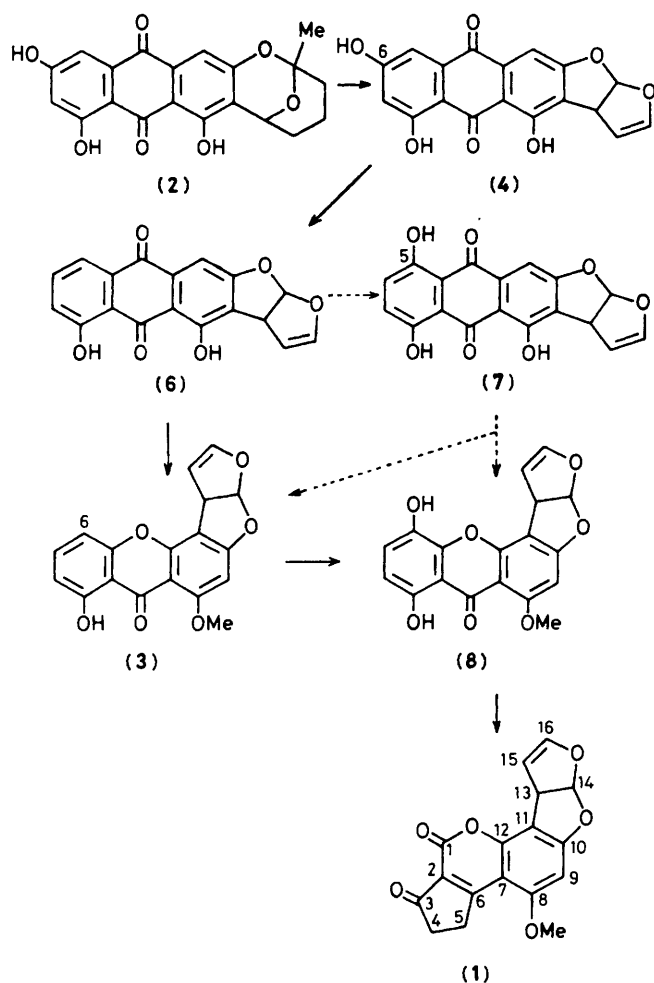
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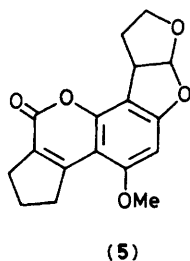
The observation of an NIH shift on incorporation of [²H₃]acetate into aflatoxin B₁ in cultures of *Aspergillus flavus* provides evidence for the timing of deoxygenation and hydroxylation steps which are an essential part of the biosynthetic pathway.

Despite extensive studies the exact sequence of intermediates on the biosynthetic pathway leading to aflatoxin B₁ (1), the

carcinogenic mycotoxin produced by the moulds *Aspergillus flavus* and *Aspergillus parasiticus*, is uncertain.¹ Even less is



Scheme 1



known of the detailed mechanisms responsible for the inter-conversion of the known intermediates. Accordingly we set out to elucidate the fate of acetate-derived hydrogen on incorporation into aflatoxin B₁ and its precursors, with a view to obtaining mechanistic information. We have recently reported ²H labelling studies on averufin (2)² and sterigmatocystin (3)³ which along with versicolorin A (4) are believed to be key intermediates on the biosynthetic pathway.¹ We now

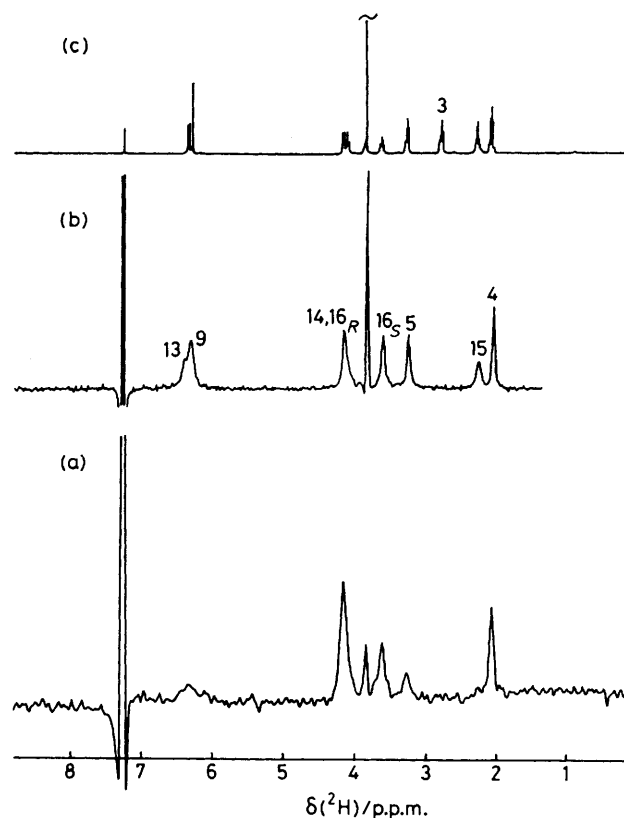


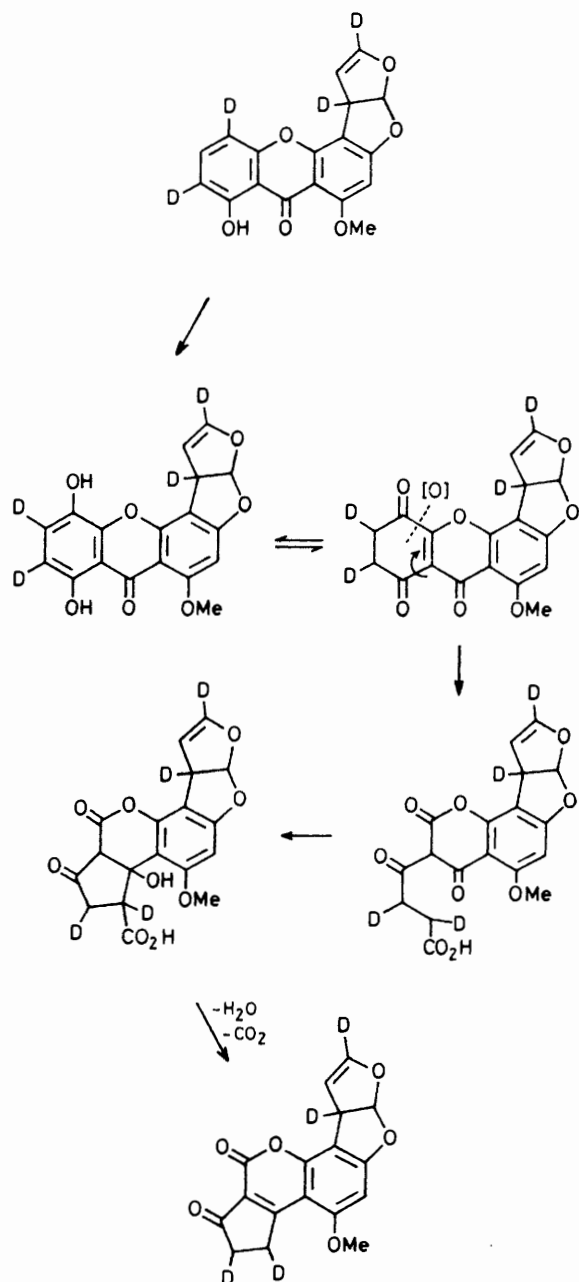
Figure 1. 55.28 MHz ²H n.m.r. spectra of 3-deoxy-aflatoxin B₂ (5) derived from (a) [2-²H₃]acetate-enriched aflatoxin B₁ and (b) from universally ²H-enriched aflatoxin B₁. (c) shows the 360.13 MHz ¹H n.m.r. spectrum of (5).

report the results of extension of these studies to aflatoxin B₁ itself.

[2-²H₃]Acetate was added to shaken cultures of *A. flavus* on low-salts medium and the aflatoxin B₁ (1) produced examined by ²H n.m.r. spectroscopy. The resulting spectrum showed signals at δ_H 6.45, 4.72, 3.35, and 2.58 p.p.m. indicating that acetate-derived hydrogen is incorporated on carbons C-9 and/or C-16, C-14, C-5, and C-4. As the signals due to H-16 and H-9 are not resolved in aflatoxin B₁ itself, the labelled sample was converted by hydrogenation⁴ into 3-deoxy-aflatoxin B₂ (5). The ²H n.m.r. spectrum of (5) [Figure 1(a)] confirmed the above observations and indicated that most of the label was on C-16 with possibly a trace of ²H incorporation at C-9. The 360.13 MHz ¹H n.m.r. spectrum of (5) [Figure 1(c)] and the 55.28 MHz ²H n.m.r. spectrum of (5) derived from universally labelled⁶ aflatoxin B₁ [Figure 1(b)] are shown for comparison.†

The crucial observation is the presence of acetate-derived hydrogen on C-5. The position is derived from the carboxy-carbon of acetate⁶ and so it must be assumed that the ²H label has migrated to C-5 from the adjacent carbon during the course of biosynthesis. The most likely mechanism would be

† Note that all the signals of interest are resolved. There is no signal for the C-3 methylene hydrogens in Figure 1(b) as these are introduced in the hydrogenation step.



Scheme 2

as a result of an NIH shift[‡] indicating that hydroxylation of the carbon adjacent to that which becomes C-5 is an essential step in the cleavage and conversion of the aromatic ring which becomes the cyclopentenone moiety in aflatoxin B₁.

One of the outstanding problems concerned with the aflatoxin biosynthetic pathway is the timing of the required loss of the 6-hydroxy-function from versicolorin A (4). One of the more plausible proposals was that versicolorin A was converted first into 6-deoxyversicolorin A (6) which was then hydroxylated to give (7), both of which are known metabolites (Scheme 1). However we have recently shown that (7) cannot be an intermediate in the formation of sterigmatocystin (3) as ²H is retained at C-6.³ The present results indicate that an intermediate which has already lost the 6-hydroxy-function present in versicolorin A (4) is hydroxylated, otherwise no NIH shift could be observed. Thus the most probable sequence of events is that versicolorin A (4) is converted *via* (6), (3), and (8) into aflatoxin B₁. A less likely sequence, (4) → (6) → (7) → (8) → (1), would mean that sterigmatocystin is *not* an obligatory intermediate and cannot yet be entirely ruled out. Cleavage of the aromatic ring and formation of the cyclopentanone ring can occur by precededented procedures as outlined in Scheme 2. Further studies to delineate the exact sequence of events between versicolorin A and aflatoxin B₁ are in progress.

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[‡] We have observed a similar NIH shift occurring during polyketide biosynthesis in studies on *O*-methyl-asparvenone,⁷ a metabolite of *Aspergillus parvulus*.