

## Biosynthesis of Betaenone B, Phytotoxin of *Phoma betae* Fr.

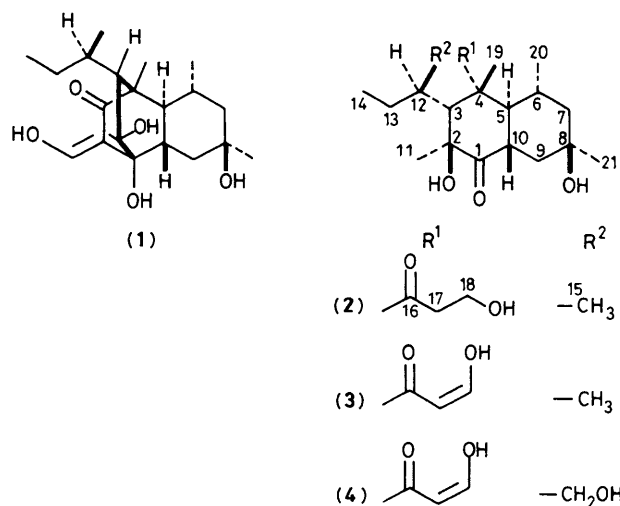
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The incorporation of [1-<sup>13</sup>C]-, [2-<sup>13</sup>C]-, [1,2-<sup>13</sup>C]-acetate, and [Me-<sup>13</sup>C]methionine, indicated that betaenone B was biosynthesized from eight acetate units *via* a polyketide pathway, and all C-methyl groups except the C-14 methyl were derived from methionine.

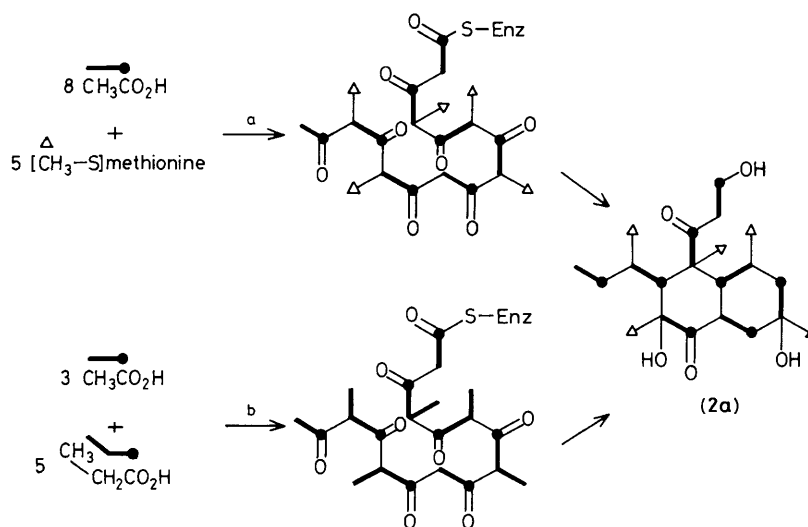
In previous papers, we have reported the isolation of betaenones, A (1), B (2), and C, (3), the phytotoxins of *Phoma betae* Fr.,<sup>1,2</sup> the fungus that causes leaf spot disease on sugar beet. Their structures were established by spectroscopic methods, chemical correlations, and X-ray crystallographic analysis. Betaenones A, (1), and C, (3), which have a unique enolic β-ketoaldehyde moiety, possess stronger phytotoxic activity than the others and (3) especially exhibited a high inhibitory effect on the RNA and protein syntheses of starfish gastrula,<sup>2</sup> and on the root elongation of rice seedlings.<sup>2</sup>

From a structural point of view, we assumed that betaenones are biosynthesized through a polyketide pathway. When the microbial metabolites of the polyketide have branching C-methyl groups, there are two possibilities, (a) the alkylation of the polyketide by a C<sub>1</sub> pool, or (b) the condensation of propionate and acetate to give the polyketide (Scheme 1).<sup>3</sup> However no fungal metabolite produced by route (b) has been reported, although homo-orsellinic acid<sup>4</sup> and aurovertin<sup>5</sup> have been biosynthesized using propionate as a starter unit instead of acetate. In this communication, we describe the biosynthetic pathway to the structurally distinct betaenone B, (2), and the origin of the branching methyl groups. We believe that this report is the first biosynthetic study on (2) or structurally related phytotoxins, *e.g.*, stemphyloxin I, (4),<sup>6</sup> and diplodiatoxin.<sup>7</sup>



In the <sup>1</sup>H n.m.r. spectrum of betaenone B, (2), all proton signals were clearly assigned by proton decoupling experiments with the exception of two kinds of methyl groups

(3 Me-C-, 2 Me-CH), which were assigned using the nuclear



Scheme 1

Table 1.  $^{13}\text{C}$  N.m.r. data for betaenone B, (2).

C-	$\delta_c^a$ /p.p.m.	$J_{cc}$ /Hz	Enrichment <sup>b</sup>
1	216.9	36.8	●
2	77.3	36.8	★
3	57.3	35.3	●
4	52.9	38.2	★
5	46.6	33.8	●
6	29.1	33.8	★
7	47.7	38.2	●
8	68.7	37.5	★
9	41.5	42.7	●
10	40.3	42.7	★
11	23.9		△
12	35.8	35.5	★
13	25.1	34.6	●
14	13.6	34.6	★
15	23.4		△
16	217.6	38.2	●
17	43.9	38.2	★
18	54.8	38.2	●
19	20.4		△
20	21.5		△
21	31.1		△

<sup>a</sup> Relative to  $\text{Me}_4\text{Si}$ . <sup>b</sup> Enrichment from  $[1-^{13}\text{C}]$ acetate (●),  $[2-^{13}\text{C}]$ acetate (★), and  $[\text{Me}-^{13}\text{C}]$ methionine (△).

Overhauser enhancement method and comparison with the spectrum of betaenone D.<sup>2</sup> On the basis of these observations, 17 signals in the  $^{13}\text{C}$  n.m.r. spectrum of (2) were unambiguously assigned by chemical shift consideration, the INEPT method, and selective proton decoupling. The remaining 4 signals (2 C=O, 2 Me-CH) were assigned by consideration of long-range selective proton decoupling. Thus, irradiation of 17-H changed the signal pattern of C-16, and a similar relationship was observed between 15-H and C-12, and further confirmed by a labelling experiment using  $[1,2-^{13}\text{C}]$ acetate. Prior to this, preliminary experiments with  $[1-^{14}\text{C}]$ acetate as the precursor established conditions which give good incorporation (7.90%).

The  $^{13}\text{C}$ -labelled phytotoxins were prepared in feeding experiments with 90% enriched sodium  $[1-^{13}\text{C}]$ ,  $[2-^{13}\text{C}]$ ,  $[1,2-^{13}\text{C}]$ -acetate and 98.57% enriched  $[\text{Me}-^{13}\text{C}]$ methionine. The labelled precursors were administered to the potato medium containing 2% sucrose (4 days after inoculation). After a further 12 day fermentation, the crude phytotoxins were extracted with EtOAc and chromatographed on a silica

gel column using  $\text{CHCl}_3$ -MeOH (98:2) and benzene-MeOH (92:8) as eluents.

The  $^{13}\text{C}$  n.m.r. spectrum of  $[1-^{13}\text{C}]$ acetate-derived betaenone B showed eight enhanced signals, attributed to C-1, C-3, C-5, C-7, C-9, C-13, C-16 and C-18, whereas the spectrum of betaenone B derived from  $[2-^{13}\text{C}]$ acetate showed enhanced signals representative of C-2, C-4, C-6, C-8, C-10, C-12, C-14, and C-17. Incorporation of eight acetate units with  $[1,2-^{13}\text{C}]$ acetate was revealed by detecting eight pairs of carbon-carbon coupling signals as indicated in (2a).

Use of  $[\text{Me}-^{13}\text{C}]$ methionine gave strong signal enhancement for five methyl carbons; C-11, C-15, C-19, C-20, and C-21. These results confirmed that the methyl group branches on the polyketide chain are derived from methionine (Table 1).

We conclude that betaenone B was biosynthesized through pathway (a) rather than (b). This conclusion is closely related to the other polymethylated fungal polyketides, e.g., citreoviridin<sup>8</sup> and radiclonic acid.<sup>9</sup>

Though the biosynthetic pathway linking the betaenones (1), (2), and (3) is not clear at present, betaenone C, (3), would be a precursor of betaenone A, (1), since the latter is formed by intramolecular aldol condensation of the former.<sup>1</sup>

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