

# Biosynthesis of agglomerin A: stereospecific incorporation of pro-*R*- and pro-*S*-hydrogens at *sn*-C-3 of glycerol into the branched C<sub>3</sub> moiety

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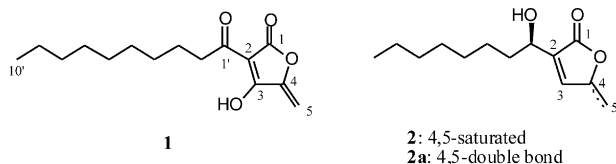
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**Abstract**—The biosynthetic origin of the C<sub>3</sub> branched unit of agglomerin A has been investigated. Feeding of *sn*-(3*R*)- and *sn*-(3*S*)-[3-<sup>2</sup>H]glycerols to *Enterobacter agglomerans* PB-6042 followed by <sup>2</sup>H NMR analysis of the resulting agglomerin A revealed that pro-*R* and pro-*S* hydrogens at *sn*-C-3 of glycerol were incorporated stereospecifically into 5*E* and 5*Z* hydrogens of agglomerin A, respectively. These results imply that the immediate precursor of the C<sub>3</sub> branched unit is not pyruvate, but 1,3-bisphosphoglyceric acid or its biological equivalent.

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Natural compounds having 2-penten-4-olide and related skeletons have been found in a variety of organisms.<sup>1</sup> This class of secondary metabolites can be classified into two structure types in terms of the oxidation state at C-3, that is, compounds with a 3-OH group (tetronic acid) and compounds with 3-H. We previously investigated the biosynthesis of a 3-H type compound, acaterin (**2**), and showed that glycerol is incorporated into the branched C<sub>3</sub> moiety (C-3, -4 and -5 positions) in such a manner that *sn*-C-1 of glycerol becomes C-3 of **2**.<sup>2</sup> Further, pro-*R* and pro-*S* hydrogens at *sn*-C-3 of glycerol become 5*E* and 5*Z* hydrogens, respectively, at C-5 of 4-dehydroacaterin (**2a**).<sup>2,3</sup> Furthermore, during this transformation hydrogens at *sn*-C-1 of glycerol were lost completely.<sup>4</sup> On the basis of these findings we proposed the immediate precursor of the branched C<sub>3</sub> unit of **2** should be a glyceric acid equivalent, and a tetronic acid type-intermediate would be involved in the formation of 3-H type compounds.<sup>2</sup> Furthermore, it has recently been established that biosynthesis of **2** involves coupling of octanoate and a C<sub>5</sub> unit corresponding to the lactone part.<sup>5</sup>

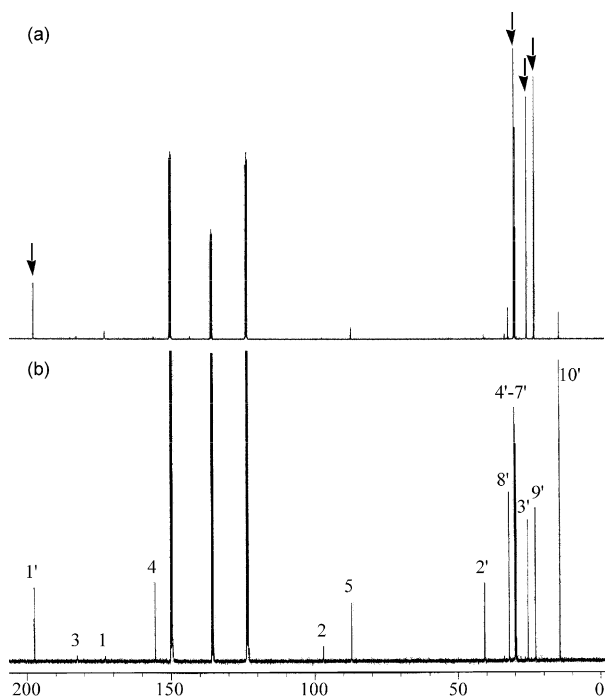


Agglomerins A, B, C and D are structurally simple tetronic acid derivatives, isolated from the fermentation broth of *Enterobacter agglomerans* PB-6042.<sup>6</sup> In order to compare the origin of the branched C-3 moiety of 3-OH type compounds with that of 3-H type compounds, we have now studied the biosynthesis of agglomerin A (**1**).

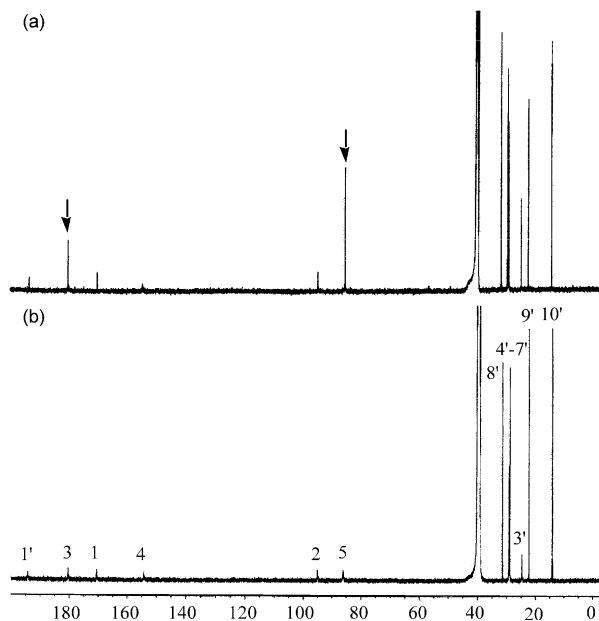
Feeding of [1-<sup>13</sup>C]acetate to *E. agglomerans* PB-6042 resulted in the <sup>13</sup>C enrichment at C-1', C-3', C-5', C-7' and C-9' of **1**,<sup>7</sup> indicating that the branched C<sub>3</sub> unit is not derived from acetate (Fig. 1). Feeding of [1,3-<sup>13</sup>C<sub>2</sub>]glycerol afforded **1** enriched at the C-3 and C-5 positions, thus confirming that a glycerol metabolite is a precursor of the C<sub>3</sub> unit (Fig. 2). Feeding of [*sn*-3,3-<sup>2</sup>H<sub>2</sub>]glycerol followed by <sup>2</sup>H NMR analysis of the resulting **1** showed that 5*Z* and 5*E* hydrogens of **1** were deuterium-labeled (Fig. 3c), indicating that *sn*-C-3 of glycerol becomes C-5 of **1** whereas *sn*-C-1 becomes C-3. Compound **1** obtained upon feeding [*sn*-1,1-<sup>2</sup>H<sub>2</sub>]glycerol did not exhibit a <sup>2</sup>H peak at the positions of 5*Z* and 5*E* hydrogens in <sup>2</sup>H NMR analysis. The observed direction of the glycerol incorporation is the same as found in the acaterin biosynthesis.

The key feeding studies of chirally <sup>2</sup>H-labeled glycerols, *sn*-(3*R*)-[3-<sup>2</sup>H]- and *sn*-(3*S*)-[3-<sup>2</sup>H]-glycerols<sup>4,8</sup> were then carried out.<sup>9</sup> The <sup>2</sup>H NMR spectrum of **1** derived from *sn*-(3*R*)-[3-<sup>2</sup>H]glycerol showed a <sup>2</sup>H signal at δ 5.35 (Fig. 3A), while **1** obtained from *sn*-(3*S*)-[3-<sup>2</sup>H]glycerol exhibited a <sup>2</sup>H signal at δ 4.82 (Fig. 3b).<sup>10</sup> These NMR data clearly indicate that the 5*E* and 5*Z* hydrogens of

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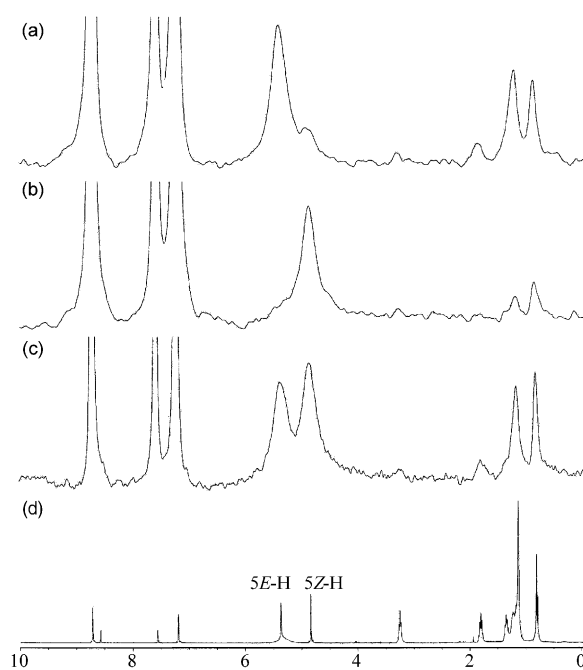


**Figure 1.**  $^{13}\text{C}$  NMR (100 MHz,  $\text{Py}-d_5$ ) spectra of agglomerin A (**1**). a: obtained upon feeding  $[1-^{13}\text{C}]$ acetate, b: non-labeled sample.

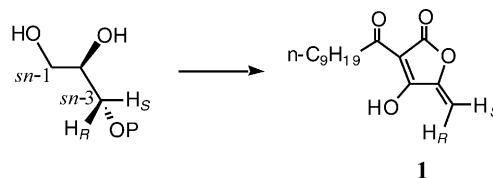


**Figure 2.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectra of agglomerin A (**1**). a: obtained upon feeding of  $[1,3-^{13}\text{C}_2]$ glycerol, b: non-labeled sample.

agglomerin A originate from pro-*R* and pro-*S*-hydrogens, respectively, at *sn*-C-3 of glycerol (Scheme 1). The fate of *sn*-C-3 hydrogens is consistent with that found in acaterin biosynthesis. The stereospecific incorporation of the chirally labeled glycerols ruled out pyruvate as the immediate  $\text{C}_3$  precursor, and 1,3-bisphosphoglyceric



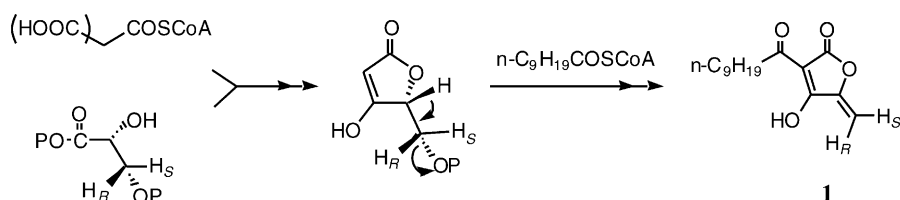
**Figure 3.**  $^2\text{H}$  NMR (61 MHz,  $\text{Py}$ ) spectra of agglomerin A (**1**). a: derived from *sn*-(3*R*)- $[3-^2\text{H}]$ glycerol, b: derived from *sn*-(3*S*)- $[3-^2\text{H}]$ glycerol, c: derived from  $[sn-3,3-^2\text{H}_2]$ glycerol, d:  $^1\text{H}$  NMR (400 MHz,  $\text{Py}-d_5$ ) spectrum of **1** (5*E* and 5*Z* hydrogens resonate at  $\delta$  5.37 and 4.84, respectively).



**Scheme 1.** Metabolic fate of *sn*-3 hydrogens in the conversion to agglomerin A (**1**).

acid or its biological equivalent is proposed as the most likely structure of the  $\text{C}_3$  precursor. Attempted feeding of  $^{13}\text{C}$  or  $^2\text{H}$ -labeled  $\text{C}_{10}$  and  $\text{C}_{12}$  fatty acids did not afford useful information on the construction of the agglomerin A carbon skeleton, since none of them was incorporated into **1**.

In conclusion, the present studies have provided evidence that a glyceric acid equivalent serves as the immediate biosynthetic precursor of the branched  $\text{C}_3$  unit of **1**, as opposed to pyruvate proposed earlier for this class of compounds. It is highly likely that a common mechanism operate in the construction of the  $\text{C}_5$  lactone moiety, irrespective of 3-*H* and 3-*OH* type 2-penten-4-olide natural compounds. Scheme 2 represents a postulated biosynthesis of **1**, starting with coupling of 1,3-bisphosphoglyceric acid with acetyl CoA or malonyl CoA. Reaction of a hypothesized  $\text{C}_5$  lactone with decanoate and anti-elimination of phosphoric acid would yield **1**.



**Scheme 2.** Postulated biosynthesis of agglomerin A (**1**).

### Acknowledgements

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- The relative enrichment of the <sup>13</sup>C signals, normalized to C-10':17 (C-1), 1.2 (C-2), 5.5 (C-3), 0.5 (C-4), 2.3 (C-5), 8.1 (C-1'), 0.7 (C-2'), 19 (C-3'), 1.2 (C-4'), 14 (C-5'), 11 (C-6' and C-7'), 2.0 (C-8'), 19 (C-9').
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- Four 500 mL-baffled Erlenmeyer flasks each containing the <sup>2</sup>H-labeled glycerol (40 mg) and the medium (100 mL) which is composed of 1.0% glucose, 0.5% yeast extract, 0.7% CaCO<sub>3</sub>, pH 7.2 adjusted by the addition of dil. HCl, were autoclaved. This was inoculated with *E. agglomerans* PB-6042 and the cultures were incubated at 25 °C in the light for 72 h on a rotary shaker at 200 rpm. The fermentation broth was processed as described in ref 6 to give **1** (7 mg) after reversed-phase HPLC purification.
- The deuterium signals observed at higher region in Figure 3 are probably due to the incorporation of [2-<sup>2</sup>H]acetate arising from [3-<sup>2</sup>H]pyruvate, a [<sup>2</sup>H]glycerol metabolite.