

Biosynthesis of the Polyether Antibiotic Monensin-A: Stereochemical Aspects of the Incorporation and Metabolism of Isobutyrate

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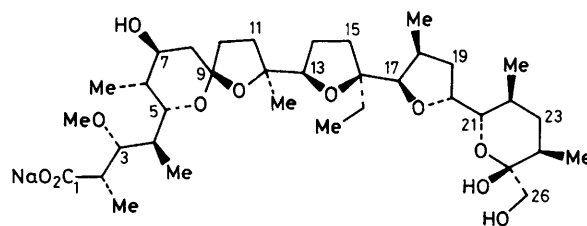
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The incorporation of stereospecifically labelled isotopomers of isobutyrate into monensin-A provides stereochemical and mechanistic information about the bioconversion of isobutyrate into methylmalonyl-CoA and n-butyryl-CoA in *Streptomyces cinnamonomensis*; a rearrangement of isobutyryl-CoA to n-butyryl-CoA occurs without loss of the α -hydrogen, and the carbonyl carbon migrates to the *pro*-(S) methyl group, and is replaced by a hydrogen atom with overall retention of configuration.

The roles of L-valine and isobutyric acid in the secondary metabolism of macrolide and polyether antibiotic producing strains of *Streptomyces* have been described by Havranak,¹ and by Omura.² In particular, the Czech group¹ have shown that monensin-A is specifically enriched at C(1), C(3), C(5), C(11), C(15), C(17), C(21), and C(23) upon incorporation of either [1-¹³C]isobutyrate or [1-¹³C]butyrate³ (see Figure 1). However, seven of these enrichments occur in units formally derived from propionate building blocks, and the highest enrichment was seen from labelled isobutyrate into the n-butyrate unit [*i.e.* C(15)–C(16)–C(32)–C(33)] in monensin-A. We report here the results of our studies with stereospecifically labelled isotopomers of isobutyric acid which confirm the existence of a pathway linking branched chain metabolism and straight chain metabolism in *Streptomyces cinnamonomensis*, and which define stereochemical and mechanistic features of the reaction(s) linking isobutyryl-CoA and n-butyryl-CoA.

The addition of sodium isobutyrate to cultures of *S. cinnamonomensis* at high concentrations (>20 mM) leads to a

significant reduction in the yield of monensin-A. In two control experiments [1-¹³C]- and [3,3'-¹³C₂]-isobutyrate were administered batchwise to 60 ml cultures, to a final concentration of 17 mM. The enriched Na-monensin-A isolated in each experiment was purified by chromatography and examined by 90 MHz ¹³C n.m.r. spectroscopy. The results shown in Table 1



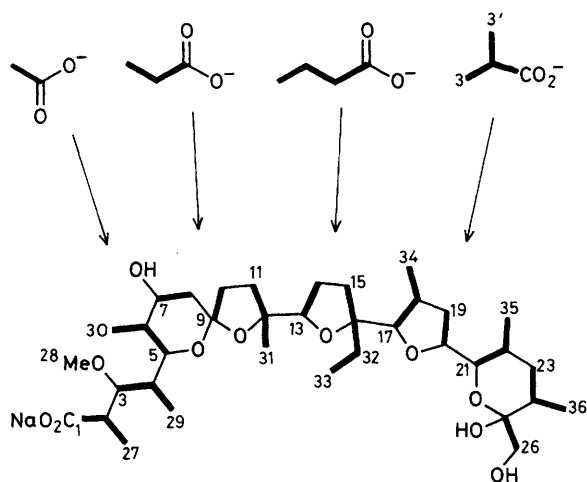
monensin - A

Figure 1

Table 1. Enrichments in monensin-A biosynthesized from [1-¹³C]- and [3,3'-¹³C₂]-isobutyrate (IBA).^a

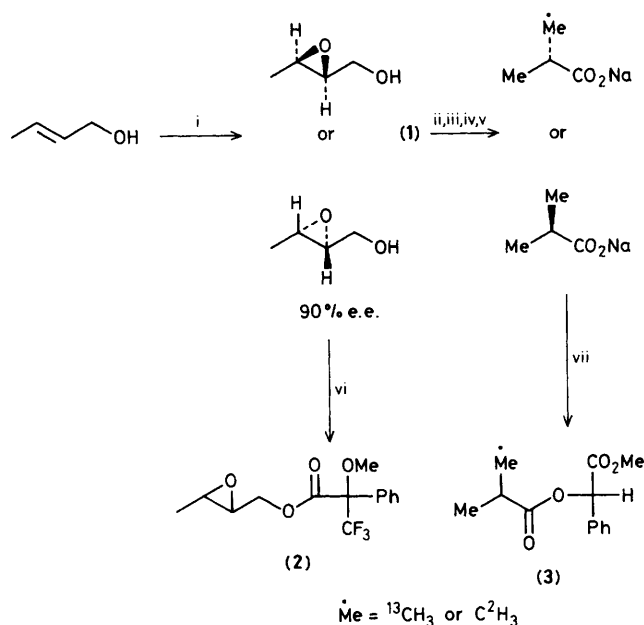
| Carbon | [1- ¹³ C]IBA | [3,3'- ¹³ C ₂]IBA |
|--------|-------------------------|--|
| C-1 | 2.3 | — |
| C-3 | 3.2 | — |
| C-5 | 2.8 | — |
| C-11 | 2.7 | — |
| C-15 | 5.7 | — |
| C-16 | — | 2.8 (<i>J</i> _{1,3} 1.83 Hz) |
| C-17 | 3.5 | — |
| C-21 | 3.2 | — |
| C-23 | 2.6 | — |
| C-27 | — | 3.1 |
| C-29 | — | 2.5 |
| C-30 | — | 2.2 |
| C-31 | — | 2.8 |
| C-32 | — | — |
| C-33 | — | 4.0 (<i>J</i> _{1,3} 1.83 Hz) |
| C-34 | — | 2.3 |
| C-35 | — | 2.2 |
| C-36 | — | 3.7 |

^a Spectra were recorded at 90.5 MHz in CDCl₃, 64 K data points, 7 s relaxation delay, 90° pulse, exponential multiplication with 0.5 Hz line broadening. The enrichment is the factor by which the height of a signal in the ¹³C{¹H} spectrum of the enriched monensin-A Na⁺ salt is increased over the height at natural abundance.

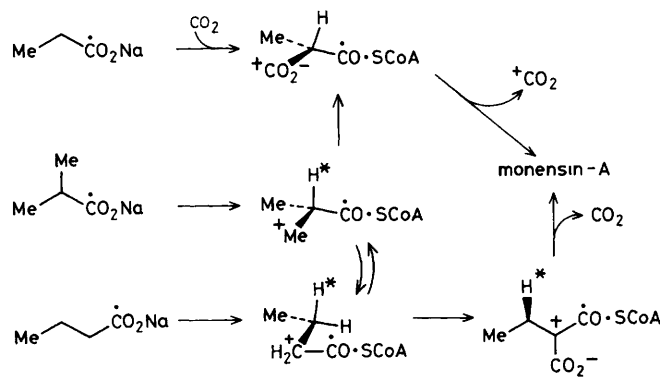
**Scheme 1**

confirm the earlier findings,¹ and show also that [3,3'-¹³C₂]-isobutyrate enriches both C(16) and C(33) (an intact unit: *J*_{1,3} 1.83 Hz), centres that are formally derived from C(2) and C(4) of n-butyrate (see Scheme 1), as well as enriching each of the methyl groups formally arising from propionate.

Although the oxidation of isobutyryl-CoA to (*S*)-β-hydroxyisobutyryl-CoA is well known,⁴ no other precedent exists for the conversion of isobutyryl-CoA into n-butyryl-CoA. In order to study stereochemical aspects of isobutyrate metabolism in *S. cinnamomensis*, (*S*)-[3-¹³C]-, (*S*)-[3-²H₃]-, and (*R*)-[3-²H₃]-isobutyrate were synthesized by the route shown in Scheme 2, which is a modification⁵ of the method developed by Aberhart.⁶ The optical purities of the epoxy-alcohols (1) were assayed by 360 MHz ¹H n.m.r. spectroscopy after conversion into the Mosher derivatives⁷ (2). The configurational purity of each labelled isobutyric acid was monitored by ¹H and either ¹³C or ²H n.m.r. spectroscopy after conversion into the mandelate derivatives (3), (90%

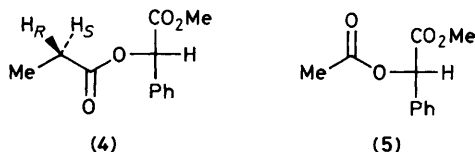


Scheme 2. Reagents: i, (+) or (-)-di-isopropyl tartrate, BuOOH, Ti(OPr)₄; ii, BuMe₂SiCl; iii, Me₂CuLi; iv, F⁻; v, NaIO₄, KMnO₄; vi, (-)-(MeO)(CF₃)CPhCOCl, *N,N*-dimethylaminopyridine, CH₂Cl₂; vii, (+)-methyl mandelate, dicyclohexylcarbodi-imide, Et₂O, *N,N*-dimethylaminopyridine.

**Scheme 3**

enantiomer excess ≡ enantiomer ratio 95:5), and these materials were subsequently used in feeding experiments to cultures of *S. cinnamomensis*.

The incorporation of (*S*)-[3-¹³C]isobutyrate led specifically to a single enrichment in monensin-A at C(16), the position formally derived from C(2) of n-butyrate. No other significant incorporations into propionate-derived units were detected by ¹³C n.m.r. spectroscopy. Taken with the labelling experiments described earlier, this result indicates that the carboxy carbon of isobutyrate migrates to the *pro*-(*S*) methyl centre, to afford an n-butyrate moiety specifically enriched at C(2). Moreover, the non-appearance of carbon-13 label in the propionate units is consistent with the conversion of (*S*)-[3-¹³C]isobutyrate into (*S*)-methylmalonyl-CoA by oxidation of the *pro*-(*S*) methyl group⁴ (see Scheme 3). These conclusions were supported by the results from two additional feeding experiments using (*S*)- and (*R*)-[3-²H₃][1-¹⁴C]isobutyrate. The specific incorporation of ¹⁴C activity in each labelled monensin-A was 8.64 and 8.34% respectively, but in the ²H n.m.r. spectrum of each, run under identical conditions, only monensin-A biosynthesized



from the (*R*) isomer showed strongly enriched signals† at δ 1.5, 1.25, and 0.9, which can be assigned unambiguously to overlapping signals from all of the C-methyl resonances.

Finally, in two separate experiments [α - ^3H , 1 - ^{14}C]- ($^3\text{H}/^{14}\text{C}$ ratio 0.84) and [α - ^2H]-isobutyrate were administered to cultures of *S. cinnamomensis*. The first experiment afforded monensin-A with a $^3\text{H}/^{14}\text{C}$ ratio of 1.31, and the second experiment gave monensin-A whose ^2H n.m.r. spectrum showed only a single broad absorption at δ 1.6, which is consistent⁸ with the location of deuterium label at C(32). To confirm this, the monensin-A (2.0 g) was degraded⁹ by Kuhn-Roth oxidation, and the acetic and propionic acids formed were collected by steam distillation and converted into the (+)-mandelate ester derivatives (4) and (5), which were then cleanly separated by chromatography. The (+)-mandelate ester of the propionate (4) was characterized by ^1H and ^2H n.m.r. spectroscopy under conditions where the diastereotopic proton resonances at C(2) are clearly resolved.¹⁰ The ^2H n.m.r. spectrum at 55.3 MHz in benzene revealed two strongly enriched signals at δ 2.45 and 2.32 in the ratio 85:15. The major signal is assigned¹⁰ to the *pro*-(2*R*) hydrogen in propionate, and it follows that the bulk‡ of the ^2H enriched propionate has the (2*R*) absolute configuration.

Integrating these results reveals stereochemical details of two metabolic pathways utilizing isobutyrate, one leading to (*S*)-methylmalonyl-CoA, and the other affording an *n*-butyrate unit by a reaction (or reactions) not requiring the loss of the α -hydrogen of isobutyrate, but involving the migration of

† The monensin-A derived from (*S*)-[3- $^2\text{H}_3$; 1- ^{14}C]isobutyrate also showed very weakly enriched ^2H n.m.r. signals at δ 1.5, 1.25, and 0.9, which are almost certainly due to the presence of approximately 5% of the (*R*)-isotopomer in the synthetically derived (*S*)-isobutyrate.

‡ The (*S*)-(+)-methyl mandelate used for derivative preparation had $[\alpha]_{\text{D}}^{20} +118^\circ$ (c 0.98, H_2O) (lit.,¹¹ $[\alpha]_{\text{D}}^{20} +134^\circ$) and for the derivative (4) $[\alpha]_{\text{D}}^{20} +91^\circ$ (c 0.96, CHCl_3) (lit.¹¹ $[\alpha]_{\text{D}}^{20} +135.5^\circ$). The appearance of the minor ^2H n.m.r. signal at δ 2.32 is therefore accounted for by a small amount of epimerization that occurred at C(2) in mandelate during formation of the derivatives.

the carboxy carbon atom, presumably activated as a coenzyme-A thioester, to the *pro*-(*S*) methyl group, with its replacement by a hydrogen atom occurring with overall retention of configuration (see Scheme 3). These pathways presumably operate in all Streptomyces and may also provide an important link between straight-chain fatty acid metabolism, and the branched-chain building blocks needed for antibiotic biosynthesis.¹² Carbon skeleton rearrangements are rare in intermediary metabolism and attempts to discover more details of this process, particularly at the enzymic level in cell-free systems, are currently underway.

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