

The Biosynthetic Origin of D-Isoleucine in the Monamycins

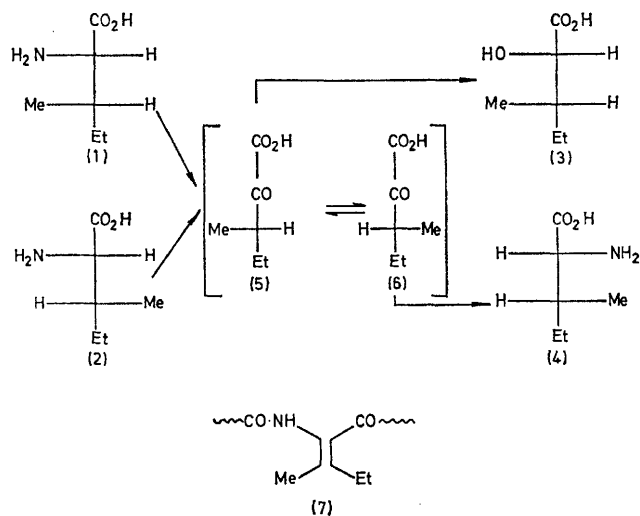
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Summary ^{14}C -Labelling studies have established that both L-isoleucine and L-alloisoleucine may serve as precursors of the D-isoleucine residues in members of the family of cyclohexadepsipeptide antibiotics known as the monamycins; possible pathways of biosynthesis of D-isoleucine from these precursors are discussed.

THE identification of D-isoleucine (4)¹ as a constituent residue of particular monamycin congeners² was the first observation to be regarded as an exception to the α -epimerisation concept formulated by Bodanszky and Perlman.³ This, together with subsequent studies^{4,5} relating to the origin of D-amino-acid residues in metabolites of microorganisms, has led us to investigate the nature of precursors of D-isoleucine in monamycins. The results of experiments with ^{14}C -labelled amino-acids are summarised in the Table.

The samples of the ^{14}C -labelled compounds were added to 20 h cultures of *Streptomyces jamaicensis* which were harvested 28 h later. The monamycins from these cultures were purified, crystallised, and then hydrolysed with 6M hydrochloric acid to give the constituent hydroxy- and amino-acids.⁶ The individual amino-acids were separated as fractions eluted from the analytical column of a Beckman 120C amino-acid analyser. Each fraction was tested for radioactivity. In all these experiments no trace of the alloisoleucine diastereoisomer was found in the hydrolysate; this included the cases where it was fed to the organism.

[U- ^{14}C]-L-Isoleucine (1) was selectively incorporated into both the D-isoleucine (4) and the L-isoleucic acid (3) components of the monamycins. Although the alloisoleucines



are not of wide occurrence in nature,^{3,7} it was of interest to determine whether the organism could use them for conversion into D-isoleucine. The L-alloisoleucine was incor-

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TABLE

Materials	Activity	% Incorporation into monamycin	Activity in D-isoleucine (d.p.m./mmol)	Activity of the amino acids in hydrolysate for comparison (d.p.m./mmol)	Activity of isoleucic acid ^a (d.p.m./mmol)
[U- ¹⁴ C]-L-Ile	50 μ C	6.3	2.1×10^6	MePro 3.5×10^4 Val 2.3×10^4	2.8×10^6
[3,3',4,5- ¹⁴ C]-DL-alloIle	20 μ C	2.4	1.5×10^6	Val 2.6×10^4 Pip 1.7×10^4	Not measured
[2- ¹⁴ C]-3-methylpent-2-enoic acid ^b	5.1 μ C	0.02	1.8×10^6	MeLeu 1.2×10^4 HyPip 1.9×10^4 Val 2.1×10^4 MePro 2.6×10^4	1.9×10^6
[3,3',4,5- ¹⁴ C]-D-alloIle	4.2 μ C	0.41	1.4×10^6	Not measured	1.9×10^6
[3,3',4,5- ¹⁴ C]-L-alloIle	5.3 μ C	2.8	8.5×10^6	Not measured	9.6×10^6

^a Based on partially purified hydroxy-acid fraction. ^b Synthesised from [2-¹⁴C]-malonate *via* the dehydrobromination of [2-¹⁴C]-2-bromo-3-methyl-pentanoic acid (ref. 9). Pip = piperazic acid, HyPip = 5-hydroxypiperazic acid.

porated selectively, as was D-alloisoleucine, but to a lesser degree than the L-isomer. Since [2-¹⁴C]-3-methylpent-2-enoic acid is not incorporated, it is unlikely that a deamination step resembling that observed for L-phenylalanine⁸ is involved in our case.

Bycroft proposed⁴ that a combined form of a dehydro-amino-acid such as (7) derived from the corresponding L-amino-acid residue, might be reduced stereoselectively *in vivo*, to the D-isomer; this would account readily in our case for the changes in chirality at the two adjacent centres. However, this does not explain for this study, the incorporation of L-isoleucine (1) into both D-isoleucine, and L-isoleucic acid. Rather, our incorporation studies suggest

that there is a common intermediate, derivable from either L-isoleucine (1) or L-alloisoleucine (2) for both L-isoleucic acid (3) and D-isoleucine (4). The most likely intermediates to satisfy these requirements are the α -keto-acids [(5), (6)] which through interconversion by inversion at C-3',¹⁰ could serve as the sources of L-isoleucic acid or a D-isoleucine. Recent studies by Lipmann¹¹ on the biosynthesis of gramicidin S and the tyrocidins are in accord with this suggestion that D-amino-acid units are incorporated directly into monamycin.

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