

MICROBIAL CONVERSION OF
MILBEMYCINS: 13 β ,29-
DIHYDROXYLATION
OF MILBEMYCINS
BY SOIL ISOLATE
Streptomyces cavourensis

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Milbemycins are a family of sixteen-membered macrolides produced by *Streptomyces hygroscopicus* subsp. *aureolacrimosus*. They exhibit broad-spectrum insecticidal and acaricidal activity.¹⁻³⁾

In the course of our studies on the microbial conversion of milbemycins, we obtained 13 β -hydroxymilbemycins A₃ (**2b**), A₄ (**1b**), and D (**3b**) from milbemycins A₃ (**2a**), A₄ (**1a**), and D (**3a**), respectively, and 13 β ,29-dihydroxymilbemycin A₄ (**1c**) from 29-hydroxymilbemycin A₄ using *Cunninghamella echinulata* ATCC 9244.⁴⁾ During the screening for microbial conversion of milbemycin A₄ (**1a**)⁵⁾ we found that a soil isolate actinomycete,

strain SANK 67386, converted milbemycin A₄ (**1a**) to 13 β ,29-dihydroxymilbemycin A₄ (**1c**) besides 13 β -hydroxymilbemycin A₄ (**1b**).

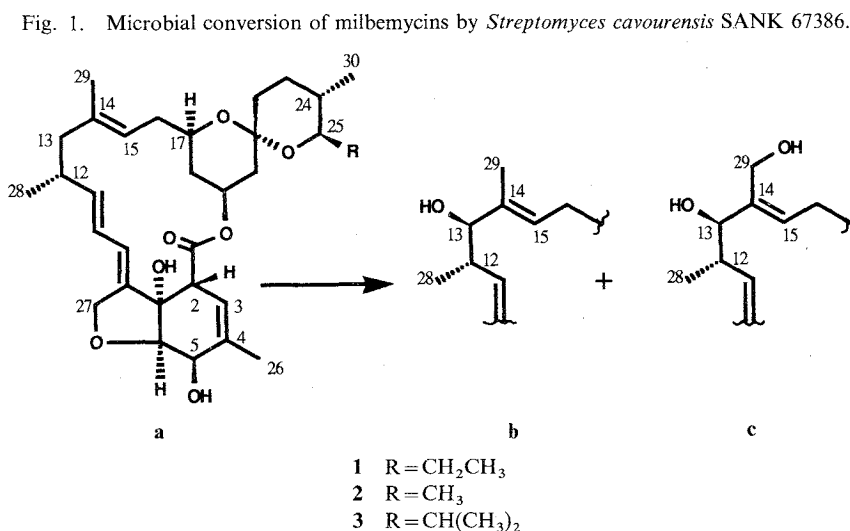
The present paper deals with the 13 β -hydroxylation and 13 β ,29-dihydroxylation of milbemycins A₃ (**2a**), A₄ (**1a**), and D (**3a**) by strain SANK 67386 (Fig. 1).

Strain SANK 67386 was isolated from solonized brown Mallee soil collected at Adelaide in Australia. On the descriptive media of SHIRLING and GOTTLIEB,⁶⁾ and WAKSMAN,⁷⁾ after 14 days at 28°C, the aerial mycelium was abundant and formed yellowish gray mass. The color of the vegetative mycelium was pale yellow to light olive gray. The strain only produced a melanoid pigment.

The substrate hyphae are long, irregularly branched, and do not fragment into short elements or formed spores. Mature spore chains are generally straight to flexuous or hooked and very long, with 10 to more than 50 spores per chain. Spores are elliptical with a smooth surface. Cell walls and whole cells were analyzed by the procedure of BECKER *et al.*,⁸⁾ and LECHEVALIER,⁹⁾ respectively. The cell wall contained LL-diaminopimelic acid and glycine as major constituents. The whole cell sugar pattern was not characteristic. The strain was considered to be a cell wall type I.

Based on morphology and analyses of the cell wall and whole cell extracts, the strain was identified as *Streptomyces cavourensis*.¹⁰⁾

The conversion of milbemycins was followed by TLC (Merck Art. No. 5715: EtOAc) and HPLC (column: Waters, Nova pak C₁₈ 8 mm × 10 cm; solvent: system I, acetonitrile-water (75:25), with



a flow rate of 1.5 ml/minute; system 2, acetonitrile-water (55:45), with a flow rate of 1.0 ml/minute; detector: UV 243 nm).

S. cavourensis SANK 67386 was cultured in five 500-ml Erlenmeyer flasks containing 100 ml MY medium composed of glucose 1.0%, Polypepton (Daigo Nutritive Chemicals) 0.5%, yeast extract (Difco) 0.3%, and malt extract (Difco) 0.3% (pH 6.3~6.5), at 28°C on a rotary shaker (200~220 rpm). After 2 days cultivation, milbemycin A₄ (5% (w/v) in 1,4-dioxane) was added to a final concentration of 500 µg/ml and cultivation was continued for seven more days. The cultured broth was then filtered and the filtrate was extracted with three 300-ml portions of EtOAc. The EtOAc extract was dried over anhydrous sodium sulfate and evaporated. The mycelium was extracted with 80% MeOH and the MeOH was evaporated, the resulting aqueous solution was extracted and evaporated in a similar manner as the filtrate. The extracts were combined and purified by silica gel chromatography (20~90% EtOAc in *n*-hexane as an eluent) to give 13β-hydroxymilbemycin A₄ (**1b**) and 13β,29-dihydroxymilbemycin A₄ (**1c**). The configurational assignment of the hydroxy group at the C-13 position was based on the comparison of the coupling constants of 13α- and 13β-hydroxy compounds in ¹H NMR spectra.^{11,12)}

Milbemycins A₃ (**2a**) and D (**3a**) were similarly subjected to the same microbial conversion by *S. cavourensis* to produce the corresponding 13β-hydroxy and 13β,29-dihydroxy derivatives. The yields are shown in Table 1. The R_f values on TLC and HPLC R_t's are listed in Table 2. The physico-chemical properties of 13β,29-dihydroxymilbemycins A₃ (**2c**) and D (**3c**) were as follows:

13β,29-Dihydroxymilbemycin A₃ (**2c**): IR (KBr) cm⁻¹ 3600~3150 (br s), 2967 (s), 2929 (s), 2873 (s), 1720 (s); ¹H NMR (270 MHz, CDCl₃) δ 5.72~5.90 (2H, m, 9-H and 10-H), 5.25~5.45 (4H, m, 3-H, 11-H, 15-H and 19-H), 4.66, 4.71 (2H, ABq, *J* = 12.2 Hz, 29-H₂), 4.13, 4.48 (2H, ABq, *J* = 12.2 Hz, 29-H₂), 4.29 (1H, d, *J* = 5.9 Hz, 5-H), 4.02 (1H, s, 7-OH), 3.96 (1H, d, *J* = 6.3 Hz, 6-H), 3.78 (1H, d, *J* = 10.3 Hz, 13-H), 3.58 (1H, m, 17-H), 3.25 (2H, m, 2-H and 25-H), 2.57 (1H, m, 12-H), 2.25~2.45 (2H, m, 16-H₂), 2.01 (1H, ddd, *J* = 1.4, 3.4 and 12.2 Hz, 20-H), 1.87 (3H, d, *J* = 2.0 Hz, 26-H₃), 1.20~1.80 (7H, m, 18-H, 20-H, 22-H₂, 23-H₂ and 24-H), 1.18 (3H, d, *J* = 6.3 Hz, 31-H₃), 1.15 (3H, d, *J* = 6.3 Hz, 28-H₃), 0.84 (3H, d, *J* = 6.8 Hz, 30-H₃), 0.75~0.90 (1H, m, 18-H); MS *m/z* 542 (M⁺ - H₂O, C₃₁H₄₂O₈), 281, 263, 237,

Table 1. Conversion of milbemycins by *Streptomyces cavourensis*.

Substrate	Concentration (µg/ml)	Conversion time (days)	Products ^a yield (%)	
			b	c
1a	500	7	36.1	16.4
2a	500	8	17.5	1.5
3a	500	7	4.3	3.3

^a **b**: 13β-Dihydroxy derivative, **c**: 13β,29-dihydroxy derivative.

Table 2. TLC R_f values and HPLC R_t's of milbemycins and conversion products.

Compound ^a	TLC R _f ^b values	HPLC R _t 's ^b (minutes)	
		System 1	System 2
1a	0.59	16.07	—
1b	0.46	3.50	10.86
1c	0.18	2.52	5.49
2a	0.59	11.80	—
2b	0.46	3.02	8.04
2c	0.18	2.27	4.52
3a	0.62	24.64	—
3b	0.48	4.59	18.26
3c	0.19	3.00	8.03

^a **a**: Substrate, **b** and **c**: products.

^b R_f values and R_t's relative to **1b**.

199, 181, 153, 129; HREI-MS Calcd for C₃₁H₄₂O₈: 542.2879, Found: 542.2879.

13β,29-Dihydroxymilbemycin D (**3c**): IR (KBr) cm⁻¹ 3600~3200 (br s), 2963 (s), 2930 (s), 2872 (s), 1716 (s); ¹H NMR (270 MHz CDCl₃) δ 5.72~5.90 (2H, m, 9-H and 10-H), 5.25~5.45 (4H, m, 3-H, 11-H, 15-H and 19-H), 4.68 (2H, br s, 27-H₂), 4.12, 4.47 (2H, ABq, *J* = 12.2 Hz, 29-H₂), 4.30 (1H, t, *J* = 6.1 Hz, 5-H), 4.01 (1H, s, 7-OH), 3.96 (1H, d, *J* = 6.1 Hz, 6-H), 3.79 (1H, dd, *J* = 4.9 and 10.3 Hz, 13-H), 3.61 (1H, m, 17-H), 3.25 (1H, m, 2-H), 3.07 (1H, dd, *J* = 2.0 and 9.3 Hz, 25-H), 2.60 (1H, m, 12-H), 2.30~2.45 (2H, m, 16-H₂), 2.16 (1H, br s, 5-OH), 2.02 (1H, ddd, *J* = 1.7, 4.9 and 12.2 Hz, 20-H), 1.87 (3H, s, 26-H₃), 1.25~1.70 (8H, m, 18-H, 20-H, 22-H₂, 23-H₂, 24-H and 31-H), 1.18 (3H, d, *J* = 6.8 Hz, 28-H₃), 1.04 (3H, d, *J* = 6.8 Hz, 32-H₃), 0.86 (3H, d, *J* = 6.8 Hz, 33-H₃), 0.81 (3H, d, *J* = 5.9 Hz, 30-H₃), 0.74~0.90 (1H, m, 18-H); MS *m/z* 570 (M⁺ - H₂O, C₃₃H₄₆O₈) 552, 309, 291, 262, 237, 226, 209, 181, 157, 139; HREI-MS Calcd for C₃₃H₄₆O₈: 570.3193, Found: 570.3188.

In our previous studies on the microbial conversion of milbemycins, 13β,24- and 13β,30-di-

hydroxymilbemycins had been obtained as minor products along with 13 β -hydroxy derivatives using *C. echinulata* ATCC 9244.⁴⁾ The C-24 and C-30 which were far from the C-13 were preferentially hydroxylated by this microorganism. In contrast, *S. cavourensis* produced 13 β ,29-dihydroxymilbemycin A₄ along with 13 β -hydroxy derivatives. It is of interest that the C-29 near to the C-13 was preferentially hydroxylated.

This study gives a new convenient method to produce 13 β ,29-dihydroxymilbemycins which will be used as metabolite reference standards in animal metabolism studies of milbemycins.

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