

2-Demethylmonensins A and B
Novel Minor Congeners of Monensins
from *Streptomyces cinnamomensis*

STANISLAV POSPÍŠIL*, PETR SEDMERA
 and VLADIMÍR HAVLÍČEK

Institute of Microbiology,
 Academy of Sciences of the Czech Republic,
 Vídeňská 1083, 142 20 Prague 4, Czech Republic

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In the course of the study of metabolites produced by various mutant strains of *Streptomyces cinnamomensis*, we have found that besides monensins¹⁾ A (1) and B (2) and traces of 3-*O*-demethylmonensins²⁾ A (3) and B (4), some of the mutants excrete small amount of new substances exhibiting standard color reaction with vanillin reagent³⁾. They were isolated and structures determined by FAB-MS, ¹H and ¹³C NMR spectroscopy as 2-demethylmonensins A (5) and B (6).

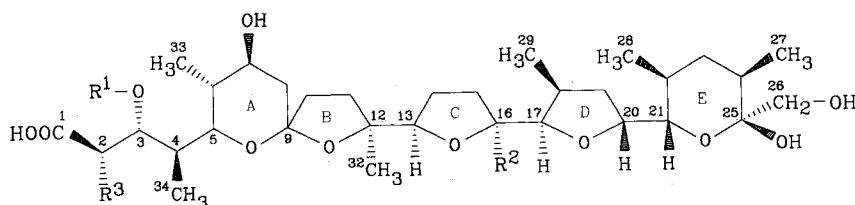
The strain *Streptomyces cinnamomensis* TR-26 (Collection of IM Prague) was cultivated under submerged conditions on soybean meal medium⁴⁾ for 6 days at 34°C on an orbital shaker (162 rpm). At the end of the fermentation, the broth was made alkaline by 0.1 M NaOH to pH 8.5. Cells were separated by centrifugation. Monensins were reextracted from the methanolic extract of the mycelium into CHCl₃. The concentrated substances were dried over anhydrous Na₂SO₄, solvent was removed *in vacuo*, and the residue was subjected to column chromatography (silica gel; heptane-ethyl acetate-methanol 5:4:1, v/v/v). Fractions containing new compounds were further chromatographed by preparative TLC on silica gel⁵⁾; R_f values in the system heptane-ethyl acetate-methanol (5:4:1, v/v/v; silica gel plate) were: 1-Na 0.44, 2-Na 0.38, 5-0.29, 6-0.33. The last purification step involved Sephadex LH-20 chromatography using methanol as a mobile phase and crystallisation from diethyl ether-H₂O. Melting points: 5

80~83°C, 6 73~76°C.

Both compounds 5 and 6 were isolated as free acids. The absence of stoichiometric sodium was confirmed by atomic absorption spectroscopy using the Na salt of 1 as a standard. The commonly used procedure for salt preparation (shaking with saturated NaHCO₃ solution) failed in this case.

First-order positive-ion FAB mass spectrum of (5) revealed the sodiated molecule at *m/z* 679 (sodium was trapped from matrix), *i.e.* at a mass of 14 daltons lower, than that of the sodium salt of monensin A (1). The first fragmentation step of the [M+Na]⁺ ion involves loss of carboxyl and the side chain attached to the ring A. The daughter ion spectrum of this ion upon collisional activation is completely identical to that of molecular ion of 3-*O*-demethylmonensin A (3). Therefore, the possible sites of demethylation are C-2 or C-3. Unfortunately, our efficient negative ion fast atom bombardment tandem mass spectrometry protocol⁶⁾ is applicable to sodium salts only and cannot be used for 5 and 6.

The distribution of ¹³C chemical shifts in 5—eight methyls, ten methylenes (one of OCH₂ type), twelve methines (seven OCH), and five quaternary carbons (one C=O, 2 × -OCO-, 2 × -C-O-) closely resembles that of monensin A. However, one methyl is missing and there is one CH₂ more. According to ¹H NMR, the molecule of 5 contains one aliphatic methoxyl, one primary, five secondary, and one tertiary methyl. Thus, the missing methyl belongs to the secondary type. COSY spectrum reveals spin system CH₂CH(O-)CH(CH₃)CH(O-)-CH(CH₃)CH(O-)CH₂, -CH₂-CH₂, -OCHCH₂CH₂, -OCHCH(CH₃)CH₂CH(O-)CH(O-)CH(CH₃)CH₂-CH(CH₃)- neatly fitting the structure of 1 having a CH₂ group at position 2 instead of -CH(CH₃) one. Except the side chain atoms, both ¹H and ¹³C NMR parameters of corresponding protons or carbons in pairs 1~5 and 2~6 are very close. The difference in C-1 chemical shifts (Table 3) is striking but not without precedences; such values were observed with laidlomycin⁷⁾ (176.46 ppm),



	R ¹	R ²	R ³
Monensin A (1)	³⁶ CH ₃	³¹ CH ₃ ³⁰ CH ₂	³⁵ CH ₃
Monensin B (2)	³⁶ CH ₃	³¹ CH ₃	³⁵ CH ₃
3- <i>O</i> -Demethylmonensin A (3)	H	CH ₃ CH ₂	CH ₃
3- <i>O</i> -Demethylmonensin B (4)	H	CH ₃	CH ₃
2-Demethylmonensin A (5)	CH ₃	CH ₃ CH ₂	H
2-Demethylmonensin B (6)	CH ₃	CH ₃	H

Table 1. Selected proton signals (ppm, 400 MHz, CDCl₃) of **1**, **5**, and **6**.

Proton	1 ¹⁷⁾	5	6
2	2.616 dq	2.611 dd 2.457 dd	2.563 dd 2.445 dd
3	3.216 dd	3.703 ddd	3.869 ddd
4	2.183 m	2.208 m	2.168 m
5	4.070 dd	4.013 dd	3.929 dd
6	2.167 ddq	1.992 ddq	2.069 ddq
7	3.860 dd	3.856 dd	3.716 dd
8	1.980 dd 1.700 dd	1.980 dd 1.729 dd	1.952 dd 1.725 dd
13	3.450 dd	3.465 dd	3.521 dd
14	1.690 m 1.570 m	1.714 m 1.665 m	1.664 m 1.618 m
15	1.900 m	1.892 m 1.709 m	2.021 m 1.850 m
17	4.090 d	4.033 d	3.943 d
18	2.333 m	2.234 m	2.237 m
19	2.200 m 1.380 m	2.161 m 1.383 m	2.209 m 1.411 m
20	4.333 ddd	4.327 ddd	4.336 ddd
21	3.940 dd	3.935 dd	3.939 dd
22	1.270 m	1.270 m	1.272 m
24	1.500 m	1.498 m	1.492 m
26	3.678 d 3.496 d	3.663 d 3.502 d	3.763 d 3.485 d
27	0.877 d	0.875 d	0.874 d
28	0.856 d	0.856 d	0.840 d
29	0.927 d	0.978 d	0.909 d
30	1.550 dq 1.370 dq	1.556 q	—
31	0.948 t	0.948 t	1.176 s
32	1.496 s	1.481 s	1.447 s
33	0.890 d	0.886 d	0.899 d
34	1.099 d	1.069 d	1.064 d
35	3.367 s	3.348 s	3.343 s
36	1.259 d	—	—

25-methoxymonensin A (178.96 ppm) and 25-methoxy-26-deoxymonensin B⁸⁾ (178.00 ppm). It confirms the existence of a free carboxyl group in **5** and **6**. Similarity of proton-proton coupling constants involving side chain and A-ring protons between **1** and **5** or **6** (Table 2) allows to reject the non-closed form⁹⁾.

Compound **6** gave a [M + Na]⁺ ion at *m/z* 665.3 in positive-ion FAB mass spectrum. Its ¹³C NMR spectrum, closely resembling that of monensin B (**2**) contains signals of 34 carbons (Table 3). Indeed, there are two tertiary methyls in the ¹H NMR spectrum (Table 1) in addition to all common features discussed above. Therefore, **6** is the lower homologue of **5**, i.e., 2-demethylmonensin B.

According to diffusion method and comparing with the corresponding monensins **1** and **2**, the 2-demethyl derivatives had approx. 200 times lower activities against *Bacillus subtilis*.

From the biogenetic point of view, monensins are classified as oligoketides. The basic skeleton is assembled by sequential coupling of precursors derived from acetate, propionate and (in the case of monensin A)

Table 2. Selected proton-proton coupling constants [Hz] in **1**, **5**, and **6**.

i, j	1 ¹⁷⁾	5	6
2d, 2u	—	13.2	16.0
2d, 3	10.2	9.9	9.4
2u, 3	—	4.8	3.1
2, 36	6.7	—	—
3, 4	2.0	2.3	2.3
4, 5	11.5	11.4	10.2
4, 34	6.8	6.9	6.8
5, 6	1.9	2.3	2.8
6, 7	≈2	2.6	2.4
6, 33	7.1	7.2	7.2
7, 8d	3.5	3.0	3.3
7, 8u	>2	2.7	2.7
8d, 8u	14.5	n.d.	n.d.
13, 14d	n.d.	5.1	4.9
13, 14u	n.d.	10.4	10.5
17, 18	4.0	4.0	3.8
18, 29	6.8	7.0	6.9
19d, 20	5.8	5.8	5.9
19u, 20	10.5	10.6	10.5
20, 21	2.6	2.9	2.7
21, 22	10.6	10.4	10.3
22, 28	6.6	6.5	6.4
24, 27	6.3	6.3	6.2
26d, 26u	11.3	11.1	11.7
30d, 30u	14.5	n.o.	—
30, 31	7.4	7.3	—

d Downfield, u upfield (for magnetically nonequivalent methylene protons); n.o. not observed; n.d. not determined.

butyrate^{10,11)}. During the biosynthesis of the 2-demethylmonensins, the last biosynthetic unit—propionate—is replaced by acetate. By feeding of precursors to cultural medium of *Streptomyces cinnamomensis* it is possible to influence the ratio of monensins A and B produced. *E.g.* valine¹²⁾, butyrate and isobutyrate yielded higher proportion of monensin A, propionate caused shift of the biosynthesis in favour of monensin B¹³⁾. On the other hand, acetate at low concentrations (0.05 to 0.2%, w/v) did not significantly influence the total production of monensins A and B or their proportion though the butyrate unit of monensin A was found to be produced also by a condensation of two acetate subunits^{11,14)}. High concentration of acetate completely inhibited the production of antibiotic. However, in no case a higher proportion of 2-demethylmonensins was observed; *e.g.*, with *S. cinnamomensis* TR-26 these compounds amounted 1~2% of the total monensin production only.

Addition of 2-demethylmonensins A and B to the known monensins A~D¹⁾ completes the number of this family members to six. The structural diversity among these compounds is due to the use of different building blocks during their biosynthesis. The question whether there is one polyketide synthase or more of them remains open^{15,16)}.

Table 3. Carbon signals (ppm, 100 MHz, CDCl₃) of 1, 2, 5, and 6.

Carbon	1 ¹⁷⁾	5	2 ¹⁷⁾	6
1	181.40 s	177.88 s	181.33 s	172.94 s
2	45.12 d	39.08 t	45.63 d	39.05 t
3	83.16 d	77.46 d	83.03 d	77.21 d
4	37.58 d	39.15 d	37.47 d	39.05 d
5	68.46 d	68.09 d	68.31 d	67.22 d
6	34.92 d	35.24 d	34.91 d	35.46 d
7	70.61 d	70.38 d	70.55 d	70.77 d
8	33.61 t	33.19 t	33.58 t	33.60 t
9	107.15 s	106.96 s	106.96 s	107.80 s
10	39.34 t	39.15 s	39.34 t	38.49 t
11	33.35 t	33.44 t	33.22 t	32.48 t
12	85.37 s	85.14 s	85.32 s	84.36 s
13	82.64 d	82.48 d	81.59 d	82.48 d
14	27.31 t	27.28 t	27.42 t	27.41 t
15	29.97 t	29.91 t	30.40 t	29.69 t
16	86.02 s	85.82 s	83.91 s	85.44 s
17	85.04 d	84.89 d	86.47 d	86.39 s
18	34.46 d	34.33 d	34.60 d	34.39 d
19	33.35 t	33.20 t	33.05 t	32.31 t
20	76.53 d	76.40 d	76.65 d	76.78 d
21	74.58 d	74.53 d	74.61 d	74.14 d
22	31.92 d	31.79 d	31.85 d	32.73 d
23	35.76 t	35.62 t	35.76 t	35.74 t
24	36.61 d	36.44 d	36.51 d	37.37 d
25	98.44 s	98.24 s	98.27 s	97.13 s
26	65.02 t	64.85 t	64.91 t	65.47 t
27	16.12 q	16.06 q	16.05 q	15.23 q
28	16.77 q	16.74 q	16.76 q	16.29 q
29	14.63 q	14.61 q	14.14 q	14.11 q
30	30.69 t	30.64 t	—	—
31	8.26 q	8.19 q	23.42 q	24.49 q
32	27.50 q	27.48 q	27.65 q	27.58 q
33	10.53 q	10.62 q	10.51 q	10.53 q
34	11.05 q	10.62 q	11.01 q	10.26 q
35	16.77 q	—	16.88 q	—
36	58.00 q	56.52 q	57.91 q	56.70 q

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