

NOTES

TMC-86A, B and TMC-96, New Proteasome Inhibitors from *Streptomyces* sp. TC 1084 and *Saccharothrix* sp. TC 1094

II. Physico-chemical Properties and Structure Determination

YUTAKA KOGUCHI, JUN KOHNO, SHIN-ICHI SUZUKI,
MAKI NISHIO, KOHEI TAKAHASHI, TETSUO OHNUKI*
and SABURO KOMATSUBARA

Basic Technology Department, Discovery Research Laboratory,
Tanabe Seiyaku Co., Ltd.,
2-50 Kawagishi-2-chome, Toda-shi, Saitama 335-8505, Japan

(Received for publication September 13, 1999)

TMC-86A, B and TMC-96, new proteasome inhibitors have been isolated from the fermentation broth of *Streptomyces* sp. TC 1084 and *Saccharothrix* sp. TC 1094, respectively¹⁾. In the preceding paper, we have reported the taxonomy of producing strains, production, isolation, and biological activities of TMC-86A, B and TMC-96¹⁾. Here we describe the physico-chemical properties and structure determination of TMC-86A, B and TMC-96.

The physico-chemical properties and NMR spectral data

of TMC-86A, B and TMC-96 are summarized in Tables 1 and 2, respectively. The molecular formulas of TMC-86A, B and TMC-96 were determined as C₁₆H₂₆N₂O₆, C₂₀H₃₄N₂O₇ and C₁₈H₃₂N₂O₆, respectively, on the basis of their HRFAB-MS and ¹H and ¹³C NMR spectral data. Their IR spectra exhibited common signals for the presence of hydroxyl (3300~3400 cm⁻¹), ketone or ester (1720 cm⁻¹), and amide (1640~1650, 1540 cm⁻¹) groups.

The structure of TMC-86A was determined by the analyses of 1D and 2D NMR and MS spectra. The ¹³C NMR spectrum displayed 16 signals composed of two -CH₃, six -CH₂-, one =CH₂, two >CH-, two quaternary carbons, and three carbonyl carbons. The ¹H NMR spectrum showed four D₂O exchangeable protons. In the ¹H-¹H COSY spectrum, four sequential proton networks of -⁸CH₂-OH, -NH-⁴CH-⁵CH₂-, -NH-^{2'}CH-^{3'}CH₂-OH and -^{2''}CH₂-^{3''}CH₂-^{4''}CH₃ were observed (Fig. 1). The presence of the epoxy ring was indicated by the chemical shift of C-1 methylene (δ_C 47.9, δ_H 3.07 and 3.08), and characteristic coupling constants: ¹J_{C-H}=180.3 Hz and ^{gem}J_{H-H}=5.6 Hz. The connection of these structural fragments was deduced from the observation of the HMBC correlations as shown in Fig. 1. The correlations from C-3 (δ 206.2) to H-4, H-5, H-1 and H-8; and from C-2 (δ 62.9) to H-1 and H-8 (Fig. 1), along with the existence of the epoxy ring, indicated the presence of the epoxy- β -aminoketone moiety. A sequential loss of the assigned fragments was seen in the EI-MS

Table 1. Physico-chemical properties of TMC-86A, B and TMC-96.

	TMC-86A	TMC-86B	TMC-96
Appearance	Colorless oil	White powder	Colorless stickly solid
[α] _D ²⁰	+10° (c 0.27, H ₂ O)	+30° (c 0.41, H ₂ O)	+25° (c 0.45, MeOH)
Molecular formula	C ₁₆ H ₂₆ N ₂ O ₆	C ₂₀ H ₃₄ N ₂ O ₇	C ₁₈ H ₃₂ N ₂ O ₆
EI-MS (m/z)	342 (M) ⁺	414 (M) ⁺	372 (M) ⁺
HRFAB-MS (m/z)			
Found	343.1863 (M+H) ⁺	415.2430 (M+H) ⁺	373.2343 (M+H) ⁺
Calcd.	343.1869 for C ₁₆ H ₂₇ N ₂ O ₆	415.2446 for C ₂₀ H ₃₅ N ₂ O ₇	373.2339 for C ₁₈ H ₃₃ N ₂ O ₆
UV λ_{max} (MeOH)	End absorption	End absorption	End absorption
IR ν_{max} (KBr) cm ⁻¹	3400, 3070, 2960, 2940, 1720, 1650, 1540, 1460, 1380, 1250, 1220, 1050	3400, 3070, 2960, 2940, 1720, 1640, 1540, 1460, 1380, 1250, 1220, 1050	3300, 3070, 2960, 2930, 1720, 1640, 1540, 1465, 1385, 1370, 1215, 1045
Solubility			
soluble in	H ₂ O, DMSO, MeOH, EtOH	H ₂ O, DMSO, MeOH, EtOH	DMSO, MeOH, acetone, EtOAc, CHCl ₃
insoluble in	n-hexane	n-hexane	H ₂ O, n-hexane

Table 2. ^1H and ^{13}C NMR data of TMC-86A, B (in $\text{DMSO}-d_6$) and TMC-96 (in CDCl_3).

Position	TMC-86A		TMC-86B		TMC-96	
	δ_c	δ_H	δ_c	δ_H	δ_c	δ_H
1	47.9 t	3.07 (d, 5.6) 3.08 (d, 5.6)	47.9 t	3.06 (d, 5.4) 3.08 (d, 5.4)	49.3 t	3.10 (d, 5.0) 3.33 (d, 5.0)
2	62.9 s		62.9 s		62.2 s	
3	206.2 s		206.3 s		207.6 s	
4	50.2 d	4.55 (ddd, 3.2, 7.4, 9.9)	50.2 d	4.55 (ddd, 3.3, 7.4, 9.9)	51.4 d	4.51 (m)
5	36.6 t	2.00 (dd, 9.9, 14.4) 2.42 (dd, 3.2, 14.4)	36.5 t	2.41 (dd, 3.3, 14.5)	38.8 t	1.30 (m) 1.61 (m)
6	140.6 s		140.6 s		25.3 d	1.66 (m)
7	113.2 t	4.75 (bs) 4.78 (bs)	113.2 t	4.74 (bs) 4.78 (bs)	21.1 q	0.93 (d, 6.3)
8	59.5 t	3.39 (dd, 5.2, 12.5) 4.09 (dd, 6.6, 12.5)	59.5 t	3.39 (dd, 5.2, 12.5) 4.07 (dd, 6.5, 12.5)	61.5 t	3.75 (dd, 7.0, 12.7) 4.20 (dd, 6.4, 12.7)
9	21.8 q	1.70 (s)	21.9 q	1.70 (s)	23.3 q	0.94 (d, 6.3)
1'	170.3 s		170.3 s		171.5 s	
2'	54.5 d	4.31 (m)	54.5 d	4.30 (m)	56.2 d	4.42 (dd, 2.5, 7.5)
3'	61.6 t	3.46 (m) 3.56 (m)	61.6 t	3.46 (m) 3.54 (m)	66.4 d	4.30 (m)
4'					17.7 q	1.15 (d, 6.4)
1''	172.0 s		172.2 s		173.5 s	
2''	37.0 t	2.10 (t, 7.4)	35.2 t	2.10 (t, 7.4)	45.8 t	2.12 (m)
3''	18.5 t	1.49 (tq, 7.4, 7.4)	25.9 t	1.43 (m)	26.3 d	2.10 (m)
4''	13.5 q	0.85 (t, 7.4)	23.5 t	1.23-1.35 (m)	22.5 q	0.96 (m)
5''			43.4 t	1.23-1.35 (m)	22.5 q	0.96 (m)
6''			68.7 s			
7'', 8''			29.2 q	1.05 (s)		
4-NH		7.94 (d, 7.4)		7.95 (d, 7.4)		7.13 (d, 7.4)
8-OH		5.05 (dd, 5.2, 6.6)		5.04 (dd, 5.2, 6.5)		2.27 (t-like, ~7)
2'-NH		7.74 (d, 8.2)		7.72 (d, 8.1)		6.46 (d, 7.5)
3'-OH		4.76 (t-like)		4.74 (t, 5.6)		3.70 (d, 3.5)
6''-OH				4.01 (s)		

spectrum (Fig 1).

The ^1H and ^{13}C NMR data of TMC-86B corresponded well to those of TMC-86A except for the signals of the fatty acid moiety (C-2'' to C-4'' in TMC-86A). The *n*-butanoyl group in TMC-86A was replaced by 6-hydroxy-6-methylheptanoyl group in TMC-86B. Thus the planar structure of TMC-86B was determined to be the 6''-hydroxyl analog of eponemycin^{2,3}.

The structure determination of TMC-96 was accomplished in the same way as described above by the NMR studies involving ^1H - ^1H COSY, HMQC, HMBC and

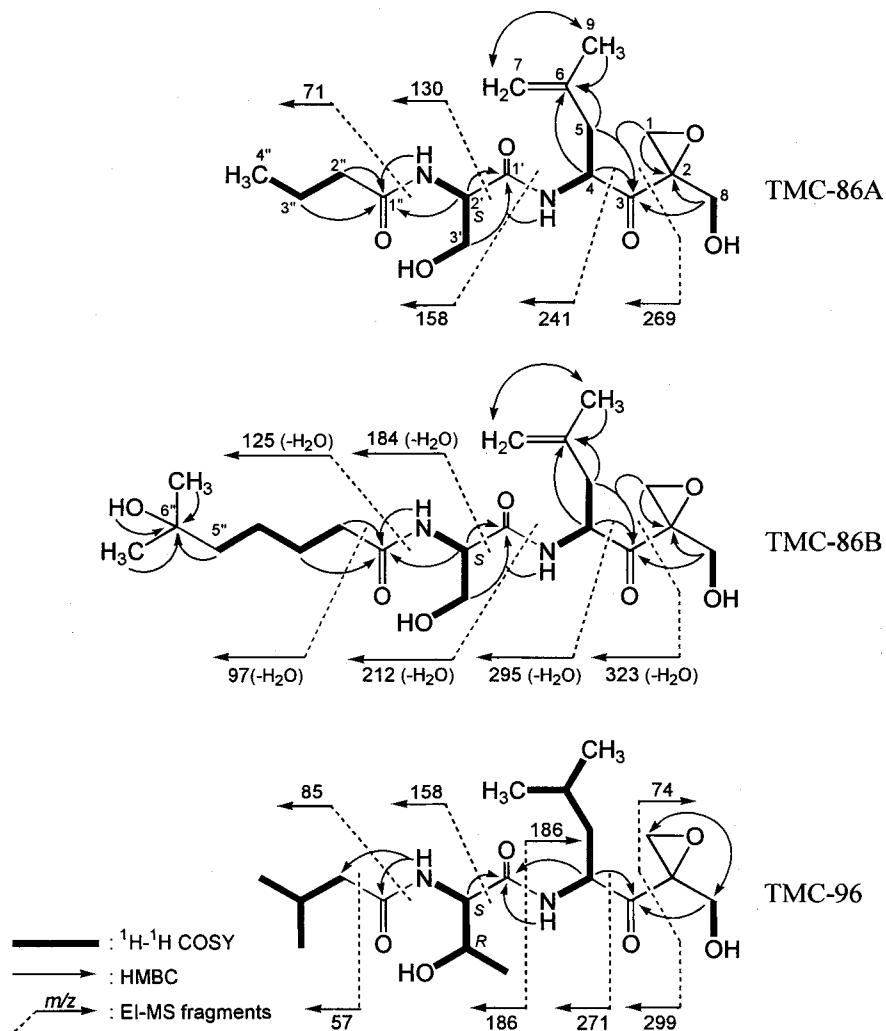
EI-MS fragmentation.

The configuration of serine in TMC-86A and B, and threonine in TMC-96 was determined to be L by chiral TLC analyses of their acid hydrolysate. The stereochemistry at C-2 and C-4 remains to be determined.

Acknowledgments

We thank Ms. NAOKO FUKUI, Ms. NORIKO OHASHI, and Ms. SONOKO SHINA for measuring spectra.

Fig. 1. Structure of TMC-86A, B and TMC-96.



References

- 1) KOGUCHI, Y.; J. KOHNO, S. SUZUKI, M. NISHIO, K. TAKAHASHI, T. OHNUKI & S. KOMATSUBARA: TMC-86A, B and TMC-96, new proteasome inhibitors from *Streptomyces* sp. TC 1084 and *Saccharothrix* sp. TC 1094. I. Taxonomy, fermentation, isolation, and biological activities. *J. Antibiotics* 52: 1069~1076, 1999
- 2) SUGAWARA, K.; M. HATORI, Y. NISHIYAMA, K. TOMITA, H. KAMEI, M. KONISHI & T. OKI: Eponemycin, a new antibiotic active against B16 melanoma. I. Production, isolation, structure and biological activity. *J. Antibiotics* 43: 8~18, 1990
- 3) HOSHI, H.; T. OHNUMA, S. ABURAKI, M. KONISHI & T. OKI: A total synthesis of 6,7-dihydroeponemycin and determination of stereochemistry of the epoxide ring. *Tetrahedron Lett.* 34: 1047~1050, 1993