

STRUCTURES OF SIOMYCIN-B
AND -C AND THIOSTREPTON-B
DETERMINED BY NMR
SPECTROSCOPY AND CARBON-13
SIGNAL ASSIGNMENTS OF
SIOMYCINS, THIOSTREPTONS,
AND THIOPEPTIN-B_a

Sir:

For several years, we have been studying the total chemical structures of thiostrepton (TST), a sulfur-containing peptide antibiotic isolated from

*Streptomyces azureus*¹⁾ and siomycin-A (SIM-A) and -D₁ (SIM-D₁), the major antibiotic and a minor antibiotic produced by *S. siyoensis*, respectively²⁾, using ¹H, ¹³C, and ¹⁵N NMR spectroscopies³⁻⁶⁾; much of the TST structure was revealed by X-ray crystallographic analysis⁷⁾.

The isolation and physicochemical properties of siomycin-B (SIM-B) and -C (SIM-C), minor components of the SIM complex, have also been reported but without elucidation of their structures⁸⁾.

Fig. 1. ¹³C FT NMR spectra of (a) SIM-A (I), (b) SIM-B (II), and (c) SIM-C (III), and (d) ¹³C PRFT NMR spectra of SIM-A (I) in CDCl₃-CD₃OD (4: 1) at 25 MHz.

FT measurement conditions: see footnote a of Table I. ¹³C PRFT measurement conditions: spectral width, 5000 Hz; pulse width, 11 μs (90°); pulse interval, 0.1 s; repetition time, 3 s; number of data points, 4K; number of transients, 17K.

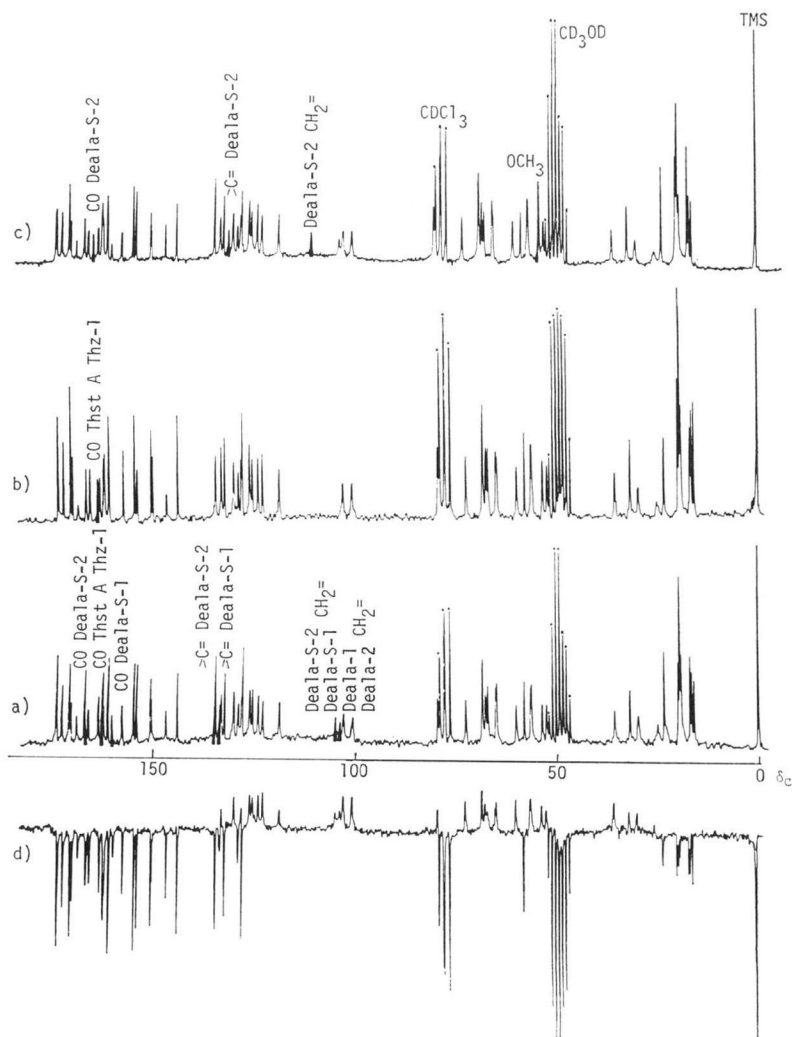
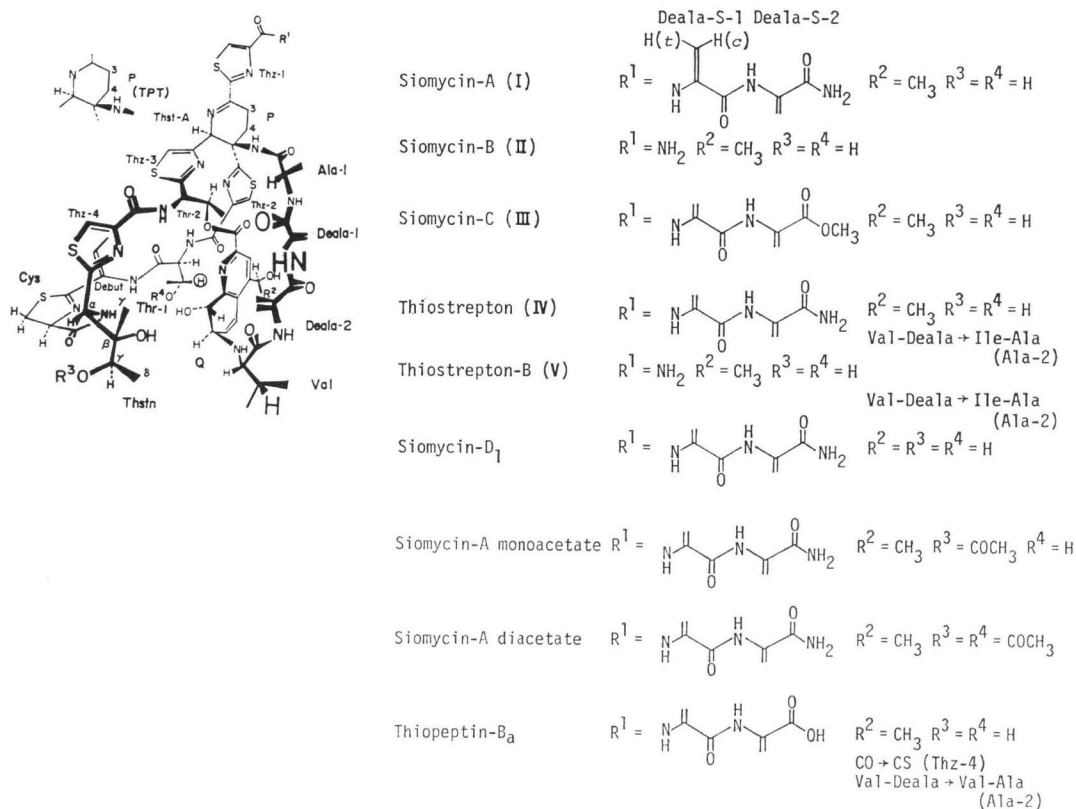


Fig. 2. Chemical structures of SIM-A (I) and related compounds.



In this communication, we wish to report the chemical structures of SIM-B and -C elucidated by 270-MHz ^1H and 25-MHz ^{13}C NMR spectroscopy, together with the structure of thiostrepton-B (TST-B), which was newly isolated from the culture broth of *S. azureus* and might be an artifact from TST. We also report here the ^{13}C signal assignments of TST and SIM as well as thiopeptin-B_a (TPT)³ since we can now compare the ^{13}C signals of several derivatives.

In a previous paper³, we reported almost all the ^1H NMR signal assignments for SIM-A, TST, and TPT at 270 MHz in CDCl_3 , $\text{CDCl}_3\text{-CD}_3\text{OD}$ (4:1), and CD_3COOD . In the 270-MHz ^1H spectra of SIM-B in these solvents, all signals arising from the Deala-S-1 and -S-2 residues disappeared completely, and some signals due to protons spatially proximate to the side chain were shifted: $\delta_{\text{H}}(\text{CDCl}_3)$ Ala βCH_3 1.47, CONH 6.56, ThstA P-3a 2.89, and P-3e 3.35. The other signals remained essentially unchanged from those of SIM-A. The 25-MHz ^{13}C spectrum in $\text{CDCl}_3\text{-CD}_3\text{OD}$ lacked six ^{13}C signals including those of

two $=\text{CH}_2$ at δ_{C} 104.0 and 105.1 in the Deala-S-1 and -S-2 residues in comparison with that of SIM-A; further, one signal at δ_{C} 162.7 was shifted downfield by +0.9 ppm, whereas the other signals remained unchanged from those of SIM-A (Fig. 1). From these observations, the structure of SIM-B was determined to be II.

The ^1H spectra of SIM-C in the solvents mentioned above showed a distinct OMe singlet at $\delta_{\text{H}}(\text{CDCl}_3)$ 3.92 and considerable shifts of the Deala-S-1 and -S-2 residue signals compared with those of SIM-A: $\delta_{\text{H}}(\text{CDCl}_3)$ Deala-S-1 $\beta=\text{CH}_2(t)$ 6.74, $\beta=\text{CH}_2(c)$ 5.53, Deala-S-2 $\beta=\text{CH}_2(t)$ 6.06, $\beta=\text{CH}_2(c)$ 5.23, and CONH 8.58. The other signals remained essentially unchanged from those of SIM-A. In the ^{13}C spectrum, the OMe signal appeared at δ_{C} 53.4 and signals of the Deala-S-2 residue were shifted considerably, whereas the other signals remained unchanged (Fig. 1). From these findings, the structure of SIM-C was assessed as III.

In a similar manner, the structure of TST-B was determined easily by comparisons of ^1H and

Table 1. ¹³C Chemical shift data.^{a)}

Carbons		SIM-A	SIM-B	SIM-C	SIM-D ₁	SIM-A-Ac	SIM-A-Ac ₂	TST	TST-B	TPT	Assignment Bases
Ala-1	αNCH	52.6	52.5	52.5	52.7	52.6	52.4	52.5	52.5	52.6	d, C, D
	βCH ₃	19.6 ^{b)}	19.6 ^{b)}	19.6 ^{b)}	19.6 ^{b)}	19.6 ^{b)}	19.5 ^{b)}	19.5 ^{b)}	19.4 ^{b)}	19.5 ^{b)}	q
	CONH	164.0	163.8	163.9	163.7	163.7	163.7	163.7	164.0	163.4	s, D
Deala-1	α=C	132.9	132.6	132.9	132.9	132.7	132.7	133.1	132.9	132.8	s, D, R-2
	β=CH ₂	103.0	103.0	103.1	103.0	102.9	102.5	103.0	103.4	103.1	t, R-2
	CONH	162.5	162.2	162.5	162.5	162.3	162.0	162.3	162.2	162.1	s, D
Deala-2	α=C	134.9 ^{e)}	134.9	135.0	134.9 ^{e)}	134.9 ^{e)}	134.9 ^{e)}	—	—	—	s, C, D
	β=CH ₂	100.9	100.6	100.9	100.6	100.6	100.2	—	—	—	t, C, R-2
	CONH	161.3	161.2	161.5	161.6	161.2	161.3	—	—	—	s, C, D
Ala-2	αNCH	—	—	—	—	—	—	49.9	50.0	49.9	d, C
	βCH ₃	—	—	—	—	—	—	18.3	18.9	19.2	q, C
	CONH	—	—	—	—	—	—	169.3	169.5	169.4	s, C, D
Val	αNCH	68.3	68.4	68.5	68.7	68.5	68.6	—	—	68.3	d, C, J
	βCH	31.8	31.7	31.8	31.8	31.8	31.7	—	—	31.7	d, C
	γCH ₃	17.1	16.9	17.0	16.9	16.9	16.9	—	—	17.0	q, C
	γ'CH ₃	19.2 ^{b)}	19.2 ^{b)}	19.2 ^{b)}	19.2 ^{b)}	19.2 ^{b)}	19.1 ^{b)}	—	—	19.2 ^{b)}	q, C
	CONH	174.0	173.7	174.1	173.7	173.9	173.1	—	—	173.5	s, C, D
Ile	αCH	—	—	—	—	—	—	66.6	66.1	—	d, C
	βCH	—	—	—	—	—	—	39.2	39.1	—	d, C
	γCH ₃	—	—	—	—	—	—	15.9	16.1	—	q, C
	γCH ₂	—	—	—	—	—	—	25.3	25.4	—	t, C
	δCH ₃	—	—	—	—	—	—	11.6	11.8	—	q, C
	CONH	—	—	—	—	—	—	173.9	173.9	—	s, C
Q	2=C	144.4 ^{d)}	144.2 ^{d)}	144.4 ^{d)}	144.2 ^{d)}	144.2 ^{d)}	143.9 ^{d)}	144.1 ^{d)}	144.1 ^{d)}	144.0 ^{d)}	s, A
	3=CH	123.0	123.0	123.3	125.1	123.1	122.6	122.9	122.9	122.9	d, A, J
	4=C	153.9	154.2	154.5	148.7	154.3	154.3	153.9	154.1	153.8	s
	5=CH	124.3	124.2	124.4	124.8	124.1	124.6	123.7	123.7	123.8	d, J
	6=CH	130.2	130.0	130.3	130.4	130.1	129.0	130.5	130.6	130.5	d, A, J
	7NCH	60.1	59.9	60.1	60.4	60.0	60.3	59.6	59.6	59.5	d
	8OCH	68.0	67.6	67.8	67.9	67.8	67.9	67.8	68.1	66.8	d
	9=C	155.2 ^{d)}	155.0 ^{d)}	155.3 ^{d)}	155.4 ^{d)}	155.1 ^{d)}	155.6 ^{d)}	155.2 ^{d)}	155.2 ^{d)}	155.2 ^{d)}	s, A
	10=C	128.3	128.0	128.3	129.7	128.1	128.8	127.7	127.8	127.7	s, A
	11OCH	65.0	64.9	65.0	61.7	64.9	64.9	64.9	64.7	65.0 ¹⁾	d, C
	12CH ₃	23.0	23.1	23.1	—	23.2	23.3	23.0	23.1	23.1	q, C
	COO	170.6 ^{e)}	170.3 ^{e)}	170.6 ^{e)}	170.7 ^{e)}	170.7 ^{e)}	170.6 ^{e)}	170.4 ^{e)}	170.3 ^{e)}	170.6 ^{e)}	s
Thr-2	αNCH	56.2	56.2	56.2	56.3	56.2	56.2	56.2	56.2	61.9	d, C, D, J
	βOCH	72.6	72.5	72.7	72.7	72.5	72.4	72.6	72.6	72.7	d, J
	γCH ₃	19.5 ^{b)}	19.4 ^{b)}	19.5 ^{b)}	19.6 ^{b)}	19.6 ^{b)}	19.4 ^{b)}	19.3 ^{b)}	19.4 ^{b)}	19.3 ^{b)}	q, C

Table 1. (Continued)

Carbons		SIM-A	SIM-B	SIM-C	SIM-D ₁	SIM-A-Ac	SIM-A-Ac ₂	TST	TST-B	TPT	Assignment Bases
Thstn	α NCH	53.7	53.7	53.7	53.8	52.6	53.0	53.7	53.6	54.0	d, A, D
	β OC	77.9	77.8	77.9	77.9	76.6	76.2	77.7	77.8	77.5	s, A, D
	γ OCH	68.6	68.4	68.6	69.0	70.5	70.4	68.5	68.5	68.6	d, A, J
	γ CH ₃	18.8	19.1	19.0	19.2	20.2	20.0	19.0	19.3	19.0	q, A
	δ CH ₃	16.5	16.5	16.5	16.6	14.3	14.1	16.5	16.5	16.5	q, A
Thz-4	2SC=N	167.3	167.0	167.4	167.4	167.3	167.4	167.1	167.0	168.0	s
	4NC=	150.8	150.7	150.9	150.8	150.6	150.8	150.7	150.7	149.5	s
	5SCH=	126.0 ^{f)}	126.0 ^{f)}	126.2 ^{f)}	126.0 ^{f)}	126.0 ^{f)}	126.0 ^{f)}	125.7 ^{f)}	125.9 ^{f)}	124.7 ^{f)}	d, J
	CONH	162.7	162.4	162.7	162.5	162.5	162.6	162.6	162.3	191.1	s, D
Cys	α NCH	79.7	79.5	79.7	79.7	79.7	79.8	79.7	79.5	79.4	d, R-2
	β SCH ₂	35.4	35.4	35.4	35.4	34.6	34.8	35.3	35.3	35.3	t, A, R-2
	CONH	172.7	172.4	172.7	172.4	170.4	171.0 ^{h)}	172.6	172.4	172.3	s, A, D
Debut	α =C	129.1	129.1	129.3	129.4	129.1	128.9	129.2	129.2	129.1	s
	β =CH	133.3	133.0	133.3	133.1	133.2	133.6	133.1	133.0	133.2	d, A, J
	γ CH ₃	15.7	15.7	15.7	15.7	15.6	15.5	15.6	15.7	15.6	q, R-1
	SC=N	171.0 ^{e)}	170.7 ^{e)}	171.0 ^{e)}	170.9 ^{e)}	171.2 ^{e)}	171.0 ^{e)}	170.8 ^{e)}	170.8 ^{e)}	170.9 ^{e)}	s
Thr-1	α NCH	56.4	56.5	56.5	56.6	56.5	54.6	56.4	56.4	56.4	d, A, C, R-1
	β OCH	67.1	67.1	67.2	67.3	67.0	69.0	67.1	67.1	66.8	d, A, C, J, R-1
	γ CH ₃	19.3	19.2	19.5	19.4	19.4	16.4	19.3	19.3	19.3	q, A, C
	CONH	166.3	166.0	166.3	166.2	166.2	165.1	166.2	166.1	166.1	s, A, D
ThstA	P2C=N	163.0	162.5	163.0	162.9	162.8	162.8	162.8	162.6	58.8 ^{j)}	s, C, D
	3CH ₂	25.3	25.4	25.4	25.5	25.3	25.5	25.0	25.1	34.0	t, C
	4CH ₂	29.6	29.7	29.7	30.3	29.7	30.2	29.7	29.9	29.2	t, C
	5NC	58.0	58.1	58.1	58.5	58.1	58.4	58.1	58.1	59.5	s, C, D
	6CH	65.0	64.9	65.0	65.0	65.0	65.2	64.9	65.0	62.6	d, C
Thz-1	2SC=N	168.9	168.6	168.7	169.0	168.9	169.0	169.0	168.8	168.8	s, D
	4NC=	157.7 ^{g)}	157.6 ^{g)}	158.0 ^{g)}	157.8 ^{g)}	157.8 ^{g)}	157.9 ^{g)}	157.8 ^{g)}	157.7 ^{g)}	156.7 ^{g)}	s
	5SCH=	125.6 ^{f)}	125.4 ^{f)}	125.7 ^{f)}	125.4 ^{f)}	125.5 ^{f)}	125.9 ^{f)}	125.4 ^{f)}	125.4 ^{f)}	125.4 ^{f)}	d, J
	CONH	162.7	163.6	162.7	162.5	162.5	162.6	162.6	163.5	162.1	s, C, D
Thz-2	2SC=N	171.0	170.7	171.0	170.8	171.0	171.0	170.8	170.8	173.1	s
	4NC=	147.3 ^{g)}	147.0 ^{g)}	147.3 ^{g)}	147.7 ^{g)}	147.1 ^{g)}	146.7 ^{g)}	147.2 ^{g)}	147.0 ^{g)}	146.4 ^{g)}	s, A
	5SCH=	128.3	128.1	128.3	128.1	128.1	128.0	128.4	128.1	128.2	d, J
	CONH	161.6 ^{g)}	161.4 ^{g)}	161.7 ^{g)}	161.7 ^{g)}	161.4 ^{g)}	161.8 ^{g)}	161.5 ^{g)}	161.4 ^{g)}	161.4 ^{g)}	s, D
Thz-3	2SC=N	174.3	173.8	174.1	174.0	173.9	174.0	174.1	174.1	174.1	s
	4NC=	150.8 ^{g)}	150.4 ^{g)}	150.9 ^{g)}	150.8 ^{g)}	150.6 ^{g)}	150.7 ^{g)}	150.6 ^{g)}	150.4 ^{g)}	155.6 ^{g)}	s
	5SCH=	118.9	118.9	119.1	119.4	118.8	119.1	118.8	118.8	118.8	d, J
Deala-S-1	α =C	135.1 ^{e)}	—	135.0	135.1 ^{e)}	135.0 ^{e)}	135.0 ^{e)}	135.0	—	135.0	s, C, D
	β =CH ₂	104.0	—	104.1	104.0	103.9	103.9	104.0	—	103.5	t, C, R-2
	CONH	160.3	—	160.0	160.2	160.1	160.2	160.3	—	160.2	s, C, D

Table 1. (Continued)

Carbons	SIM-A	SIM-B	SIM-C	SIM-D ₁	SIM-A-Ac	SIM-A-Ac ₂	TST	TST-B	TPT	Assignment Bases
Deala-S-2 $\alpha=C$	133.8	—	131.4	134.0	133.9	134.2	133.9	—	135.3	s, C, D
$\beta=CH_2$	105.1	—	110.8	104.7	104.9	104.8	105.0	—	107.1	t, C, R-1
CO	166.9	—	165.0	166.7	166.7	166.7	166.8	—	166.4	s, C, D
OCH ₃	—	—	53.4	—	—	—	—	—	—	q, C
Acetyl CH ₃	—	—	—	—	21.3	21.0	—	—	—	q, C
CH ₃	—	—	—	—	—	21.4	—	—	—	q, C
CO	—	—	—	—	169.5	169.1	—	—	—	s, C
CO	—	—	—	—	—	169.8 ^{b)}	—	—	—	s, C

^{a)} ¹³C NMR spectra were recorded on a JEOL FX 100 FT NMR spectrometer at 25.05 MHz in CDCl₃-CD₃OD (4 : 1) with TMS as an internal reference (δ_c 0).

FT measurement conditions were: spectral width, 5,000 Hz; pulse width, 5 μ s (50°); repetition time, 1 s; number of data points, 8K; number of transients, 4K; 10-mm spherical cell; concentration, 0.085 mmol/ml; 60°C.

Abbreviations are as follows: Deala, dehydroalanine; Debut, dehydrobutyrine; P, piperidine ring; Q, quinaldic acid precursor; Thstn, thiostreptine residue; ThstA, thiostreptonic acid unit; Thz, thiazole ring. For assignment bases: s, singlet; d, doublet; t, triplet; q, quartet; A, acetylation shifts; C, chemical-shift comparisons; D, deuterium substitution isotope shifts; J, Jr values in SFORD spectra; R-1, reference 12; R-2, reference 13.

^{b-h)} Assignments may be interchanged in each column.

ⁱ⁾ Triplet.

^{j)} Doublet.

¹³C spectra between TST (IV) and TST-B (V). TST-B has no side chain, like SIM-B.

We also reinvestigated the ¹³C signal assignments of SIM-A and TST which were very tentatively examined in a previous paper⁴⁾. The assignments of signals were first done by single-frequency off-resonance decoupling (SFORD) and partially-relaxed FOURIER transform (PRFT) techniques to differentiate protonated and non-protonated carbon signals (see Fig. 1d).

The assignments of CH and CH₂ signals were easy from the chemical-shift comparisons of the compounds examined as well as those of TPT and SIM-D₁⁶⁾, and the residual *J*-values in their SFORD spectra, because the ¹H chemical shifts are known³⁾. In a similar manner, CH₃ signals were assigned, but some could not be because of mutual signal overlappings.

Non-protonated carbon signals were assigned with the aid of deuterium substitution effects besides the chemical shift comparisons. Deuterium substitution isotope shifts arising from the deuterium exchange of the CONH groups⁹⁾ in the CDCl₃ - CD₃OD (4:1) solvent system at various temperatures were observed. All COND

and -CND signals were slightly shifted upfield from the corresponding CONH and -CNH signals. In particular, the ThstA P-2 (N=C) signal was considerably shifted upfield due to the imine-enamine tautomerism which causes exchange of P-3-CH₂ hydrogens for deuteriums. In fact, the P-3-CH₂ signal gradually disappeared in CDCl₃ - CD₃OD (4:1).

Acetylation shifts¹⁰⁾, in the ¹³C spectra of SIM-A monoacetate and diacetate¹¹⁾ were useful for assigning the signals due to the Thstn and Thr-1 residues. Moreover, the acetylation caused a conformational change of residues in the vicinity. Thus, the signals of the Cys residue in both acetates and those of some of the Q residue in diacetate were affected.

Table 1 lists the ¹³C signal assignments of SIM-A, -B, -C, TST, and TPT including SIM-D₁ and SIM-A mono- and diacetates.

Thus, we determined the chemical structures of TST-B, newly isolated from *S. azureus*, and of SIM-B and -C. The ¹³C signals of SIM, TST, and TPT were reassigned as listed in Table 1.

KAZUO TORI*
 KATSUYA TOKURA
 YOHKO YOSHIMURA
 YOSHIHIRO TERUI
 KEI OKABE
 HIDEO OTSUKA

Shionogi Research Laboratories,
 Shionogi & Co., Ltd.,
 Fukushima-ku, Osaka, 553 Japan

KAZUHIRO MATSUSHITA

Application Center,
 Scientific Instrument Project,
 JEOL Ltd.,
 Nakagami, Akishima, Tokyo, 196
 Japan

FUYUHIKO INAGAKI
 TATSUO MIYAZAWA

Department of Biophysics and
 Biochemistry, Faculty of Science,
 The University of Tokyo,
 Bunkyo-ku, Tokyo, 113 Japan

(Received October 22, 1980)

References

- 1) VANDEPUTTE, J. & J. D. DUTCHER: Thiostrepton, a new antibiotic. II. Isolation and chemical characterization. *Antibiot. Ann.* 1955/1956: 560~561, 1956
- 2) NISHIMURA, H.; S. OKAMOTO, M. MAYAMA, H. OTSUKA, K. NAKAJIMA, K. TAWARA, M. SHIMOHIRA & N. SHIMAOKA: Shiomycin, a new thiostrepton-like antibiotic. *J. Antibiotics, Ser. A* 14: 255~263, 1961
- 3) TORI, K.; K. TOKURA, Y. YOSHIMURA, K. OKABE, H. OTSUKA, F. INAGAKI & T. MIYAZAWA: ^1H NMR spectral evidence for the structure and conformation of peptide antibiotic siomycin-A. *J. Antibiotics* 32: 1072~1077, 1979
- 4) TORI, K.; K. TOKURA, K. OKABE, M. EBATA, H. OTSUKA & G. LUKACS: Carbon-13 NMR studies of peptide antibiotics, thiostrepton and siomycin A: The structure relationship. *Tetrahedron Lett.* 1976: 185~188, 1976
- 5) OLESKER, A.; L. VALENTE, L. BARATA, G. LUKACS, W. E. HULL, K. TORI, K. TOKURA, K. OKABE, M. EBATA & H. OTSUKA: Natural abundance ^{15}N NMR spectroscopic evidence for the structural relationship between the peptide antibiotics thiostrepton and siomycin-A. *J. Chem. Soc., Chem. Comm.* 1978: 577~578, 1978
- 6) TOKURA, K.; K. TORI, Y. YOSHIMURA, K. OKABE, H. OTSUKA, K. MATSUSHITA, F. INAGAKI & T. MIYAZAWA: The structure of siomycin-D₁, peptide antibiotic isolated from *Streptomyces sioyaensis*. *J. Antibiotics* 33: 1563~1567, 1980
- 7) ANDERSON, B.; D. C. HODGKIN & M. A. VISWAMITRA: The structure of thiostrepton. *Nature* 225: 233~235, 1970
- 8) EBATA, M.; K. MIYAZAKI & H. OTSUKA: Studies on siomycin. I. Physicochemical properties of siomycins A, B, and C. *J. Antibiotics* 22: 364~368, 1969
- 9) a) FEENEY, J.; P. PARTINGTON & G. C. K. ROBERTS: The assignment of carbon-13 resonances from carbonyl groups in peptides. *J. Magn. Resonance* 13: 268~274, 1974
 b) LADNER, H. K.; J. J. LED & D. M. GRANT: Deuterium isotope effects on ^{13}C chemical shifts in amino acids and dipeptides. *J. Magn. Resonance* 20: 530~534, 1975
 c) NEWMARK, R. A. & J. R. HILL: Assignment of primary and secondary amide carbonyl resonances in carbon-13 NMR. *J. Magn. Resonance* 21: 1~7, 1976
- 10) TERUI, Y.; K. TORI & N. TSUJI: Esterification shifts in carbon-13 NMR spectra of alcohols. *Tetrahedron Lett.* 1976: 621~622, 1976
- 11) TOKURA, K.; Unpublished results
- 12) a) WALKER, J.; A. OLESKER, L. VALENTE, R. RABANAL & G. LUKACS: Total structure of the polythiazole-containing antibiotic micrococcin P. A ^{13}C nuclear magnetic resonance study. *J. Chem. Soc., Chem. Comm.* 1977: 706~708, 1977
 b) BYCROFT, B. W. & M. S. GOWLAND: The structures of the highly modified peptide antibiotics micrococcin P₁ and P₂. *J. Chem. Soc., Chem. Comm.* 1978: 256~258, 1978
- 13) a) SAKAKIBARA, H.; H. NAGANAWA, M. OHNO, K. MAEDA & H. UMEZAWA: The structure of althiomycin. *J. Antibiotics* 27: 897~899, 1974
 b) BYCROFT, B. W. & R. PINCHIN: Structure of althiomycin, a highly modified peptide antibiotic. *J. Chem. Soc., Chem. Comm.* 1975: 121~122, 1975