

STRUCTURE OF AWAMYCIN,
A NOVEL ANTITUMOR
ANSAMYCIN ANTIBIOTIC

Sir:

In the course of a continuing search for novel antitumor antibiotics of microbial origin, a new antitumor antibiotic awamycin was isolated from the culture broth of *Streptomyces* sp. No. 80-217 isolated from a soil sample collected in Chiba Prefecture, Japan¹⁾. The taxonomy of the producing organism, fermentation, isolation and physico-chemical and biological properties of this antibiotic have been reported in the preceding paper¹⁾. This communication deals with the structure elucidation of awamycin.

As described in the preceding paper, physico-chemical characteristics of awamycin suggested that this antibiotic belongs to the group of naphthalenoid ansamycins having a sulphur atom in the molecule. Through the combination of elemental analysis and mass spectrometry, the molecular formula of awamycin was established as $C_{35}H_{49}O_{12}NS$ (MW 743)¹⁾.

In the ^{13}C NMR spectrum of awamycin (in $CDCl_3$), 15 singlets including two carbonyl signals [δ_C 180.4 (s) and 183.1 (s)] characteristic for the quinone carbonyls²⁾, 13 doublets and 10 quartets were observed. On the other hand, in the 1H NMR spectrum of awamycin (in $CDCl_3$), signals corresponded to 49H were observed which were classified into 10 methyl signals, 13 methine signals and 6 D_2O exchangeable signals.

The structure of the ansa moiety of awamycin was established as shown in Fig. 1 through 1H NMR homo-decoupling study and by comparison of 1H NMR data of awamycin with those of rifamycin W³⁾ and streptovaricin D^{4,5)} (Fig. 1, Table 1). The existence of a carbo-

methoxy group was suggested by IR (ν_{max}^{KBr} 1721 and 1209 cm^{-1}), 1H NMR [δ_H 3.715 (3H, s)] and ^{13}C NMR [δ_C 51.8 (q) and 172.8 (s)] spectral data and the positions of this moiety and the ansa moiety were elucidated through the 1H NMR and ^{13}C NMR comparison with those of rifamycin W³⁾ and streptovaricin D^{4,5)} (Fig. 1, Table 1). Configurations of the double bonds at C2-C3 (*cis*), C4-C5 (*trans*) and C15-C16 (*trans*) and the transoid conformation at C3-C4 have been elucidated because of the similarity of 1H NMR spectral data of awamycin [δ_H 6.440 (1H, d, $J=11$ Hz, C3-H), 6.945 (1H, dd, $J=16$ and 11 Hz, C4-H), 5.793 (1H, dd, $J=16$ and 9 Hz, C5-H) and 5.726 (1H, d, $J=9$ Hz, C15-H)] with those of rifamycin S [δ_H 6.19 (1H, d, $J=9.5$ Hz, C3-H), 6.60 (1H, dd, $J=15.0$ and 9.5 Hz, C4-H), and 5.93 (1H, dd, $J=15.0$ and 7.0 Hz, C5-H); C2-C3 (*cis*), C4-C5 (*trans*) and C3-C4 (*transoid*)⁶⁾ and protostreptovaricin I (**1**) [δ_H 5.66 (1H, d, $J=10$ Hz, C15-H; C15-C16 (*trans*)⁵⁾, respectively.

Because $C_{25}H_{39}O_5N$ out of $C_{35}H_{49}O_{12}NS$ of awamycin has been assigned to the ansa moiety (Fig. 1), it is obvious that the rest ($C_{13}H_{10}O_4S$) can be attributed to the naphthoquinone moiety. In the 1H NMR spectrum of this region, three methyl signals [δ_H 2.281, 2.377 and 3.854 (each

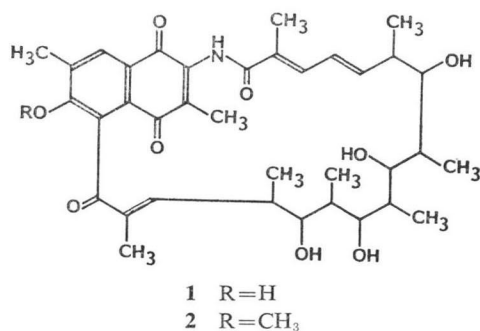


Fig. 1. Structures of the ansa regions of awamycin, rifamycin W³⁾ and streptovaricin D⁴⁾.

	R ₁	R ₂	R ₃
Awamycin	COOCH ₃	CH ₃	C=O
Rifamycin W	CH ₃	CH ₂ OH	C=O
Streptovaricin D	COOCH ₃	CH ₃	C—O

Table 1. NMR data of ansa moieties of awamycin (in CDCl₃), rifamycin W [in CDCl₃ - DMSO-*d*₆ (3: 1)] and streptovaricin D (in CDCl₃)*.

	Awamycin		Rifamycin W ³⁾	Streptovaricin D ^{4,5)}	
	δ_{H}	δ_{C}	δ_{H}	δ_{H}	δ_{C}
NH	8.36 1H, s	—	8.55 1H, s	***	—
1	—	164.6 s	—	—	169.4 s
2	—	131.7 s ^a	—	—	127.5 d
2-CH ₃	2.191 3H, s	12.1 q	2.08 3H, s	***	12.9 q
3	6.440 1H, d	128.9 d ^b	6.31 1H, d	7.66 1H, d	134.7 d
4	6.945 1H, dd	128.8 d ^b	7.00 1H, dd	6.52 1H, t	124.0 d
5	5.793 1H, dd	142.7 d ^c	6.32 1H, dd	5.83 1H, t	144.1 d
6	2.328 1H, m	42.0 d	2.61 1H, m	3.00 1H, m	41.9 d
6-CH ₃	1.240 3H, d	20.6 q	0.96 3H, d	1.26 3H, d	22.2 q
7	3.445 1H, t	82.7 d	4.46 1H, dd	3.54 1H, q	83.6 d
7-OH	4.397 1H, d	—	**	***	—
8	2.10 1H, m	38.7 d	2.06 1H, m	2.30 1H, m	38.8 d
8-CH ₃	0.890 3H, d	15.7 q ^d	1.23 3H, d	0.99 3H, d	15.7 q
9	4.212 1H, dt	76.5 d	4.81 1H, dd	4.25 1H, q	77.6 d
9-OH	4.635 1H, d	—	**	***	—
10	2.838 1H, s	46.9 d	2.54 1H, m	2.94 1H, d	47.4 d
10-COOCH ₃	3.715 3H, s	172.8 s	—	3.78 3H, s	173.0 s
	—	51.8 q	—	—	51.9 q
10-CH ₃	—	—	0.96 3H, s	—	—
11	4.146 1H, t	73.1 d	3.94 1H, dd	4.25 1H, d	73.4 d ^a
11-OH	4.089 1H, d	—	**	***	—
12	2.05 1H, m	38.3 d	2.00 1H, m	1.95 1H, m	38.5 d ^b
12-CH ₃	0.801 3H, d	9.1 q	1.03 3H, d	0.80 3H, d	9.1 q
13	3.515 1H, td	69.9 d	5.26 1H, dd	3.54 1H, d	70.4 d ^a
13-OH	2.167 1H, d	—	**	***	—
14	2.675 1H, m	37.0 d	3.12 1H, m	2.57 1H, m	37.5 d ^b
14-CH ₃	0.746 3H, d	17.4 q ^d	—	0.73 3H, d	15.1 q
14-CH ₂ OH	—	—	3.78 1H, dd	—	—
	—	—	4.08 1H, dd	—	—
	—	—	**	—	—
15	5.726 1H, d	144.4 d ^c	7.18 1H, d	5.51 1H, d	153.6 s
16	—	136.9 s ^a	—	—	***
16-CH ₃	2.038 3H, s	16.2 q ^d	2.34 3H, s	***	12.7 q
17	—	195.3 s	—	—	169.0 s

* δ ppm from TMS.** δ 3.4~4.5 (5H).

*** Data not given in the paper.

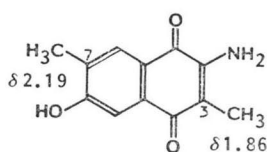
^{a-d} Assignments may be interchanged.

3H, s)] and a phenolic OH signal [δ_{H} 11.98 (1H, s, D₂O exchangeable)] were observed.

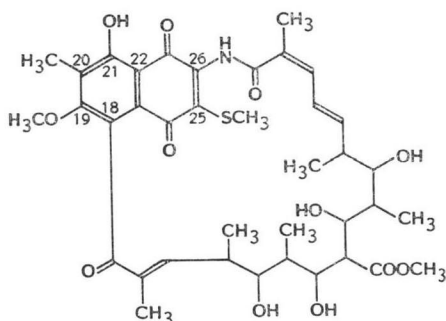
So, it was shown that the naphthoquinone moiety of awamycin is substituted with OH, CH₃, OCH₃ and SCH₃ groups in addition to the ansa moiety at C-18 and C-26.

The position of a phenolic OH (δ_{H} 11.98) was estimated to be *peri*- (C-21) to the carbonyl group of naphthoquinone because of its characteristic low field shift in the ¹H NMR spec-

trum. A signal at δ_{C} 195.3 and an IR absorption maximum at 1665 cm⁻¹, attributed to the carbonyl (C-17) of the ansa moiety of awamycin, are quite similar to those of protostreptovaricin II (2) [δ_{C} 195.8 and ν_{max} 1670 cm⁻¹ (nonhydrogen bonded), respectively] and are different from those of protostreptovaricin I (1) [δ_{C} 204.4 and ν_{max} 1640 cm⁻¹ (hydrogen bonded), respectively]⁵⁾. This fact suggests that awamycin also has an OCH₃ group at C-19 and an OH group



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at the C-21 position, like protostreptovaricin II (2). The *meta* substitution of the OCH₃ (C-19) and OH (C-21) groups is also supported by the ordinary ¹³C NMR chemical shift of C-19 and C-21 (δ_C 161.9 and 162.6).

Because positions C-18 and C-26 (ansa moiety), C-19 (OCH₃) and C-21 (OH) are occupied by the substituents, the position of the methyl group is either C-20 or C-25 and is concluded to be C-20 because the chemical shift of the methyl group (δ_H 2.281) of awamycin is similar to that of the aromatic (and not to the olefinic) methyl group of 2-amino-3,7-dimethyl-6-hydroxy-1,4-naphthoquinone (3)⁷⁾. The positions of the methyl group and the ansa moiety at C-20 and C-18 and C-26, respectively are also supported from the biosynthetic point of view of various ansamycins^{2,8)}. Because other positions of the naphthoquinone moiety of awamycin have been determined, the position of SCH₃ group is C-25.

From all the accumulated data described above, the structure of awamycin is concluded to be 4.

Acknowledgment

This work was supported by a Grant-in-Aid from

the Ministry of Education, Science and Culture, Japan.

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(Received April 30, 1985)

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