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Natural Abundance ^{15}N Nuclear Magnetic Resonance Spectroscopic Evidence for the Structural Relationship between the Peptide Antibiotics Thiostrepton and Siomycin-A

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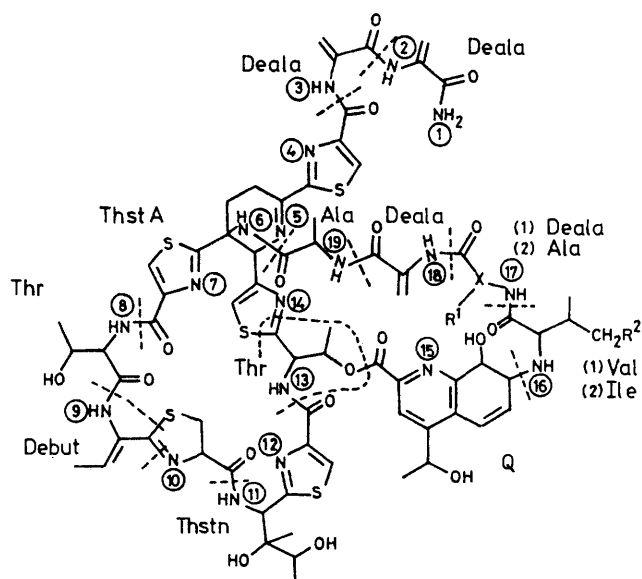
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Summary Nitrogen-15 n.m.r. spectral comparison between the antibiotics thiostrepton and siomycin-A has permitted further evidence for the structure of the latter to be obtained.

As a result of chemical degradations¹ and a ^{13}C n.m.r. study² the tentative structure (1) has been proposed for the peptide antibiotic siomycin-A isolated from *Streptomyces sioyaensis*.³ We now provide further evidence for the complete structure of siomycin-A by comparing its ^{15}N n.m.r. spectrum with that of the closely related antibiotic thiostrepton (2) of known constitution.^{2,4} The structural difference between the two antibiotics was previously suggested to be the substitution of an Ile-Ala dipeptide unit of (2) by a Val-Deala residue in (1).^{1,2}

Nitrogen-15 n.m.r. spectroscopy is a promising tool for studies of complex molecules of biological interest⁵⁻⁹ since the chemical shift range of nitrogen covers about 800 p.p.m.¹⁰ With the availability of n.m.r. spectrometers operating at higher fields for ^{15}N , the severe sensitivity problems associated with this nucleus can be overcome⁹ even for compounds of the size investigated here.



- (1) X = C, R¹ = CH₃, R² = H
(2) X = CH, R¹ = R² = Me

Amino-acid sequence information in peptides from ^{15}N n.m.r. spectra seems to depend on solvent.^{5,11,12} Of the 19 nitrogen atoms present in both (1) and (2), 16 have almost identical chemical shifts. As a result of the structural difference in the nature of the dipeptide units, the 2 nitrogen atoms involved in the dipeptide units and the following Deala-nitrogen are expected to exhibit different chemical shifts because of the effect of the neighbouring residues.^{11,12} On the basis of well established α -, β -, γ -, δ -, and ϵ -effects^{13,14} the Table contains tentative signal assignments

case for (1) the assignment of the lower-field shift to the Val-following Deala unit is preferred. It is not clear at the moment why the Ala-preceding Deala-nitrogen signal of (1) is shifted by 4.7 p.p.m. upfield relative to that of (2) as a result of the sequence effect.

The ^{15}N signal of the primary amide in the side chain can be easily assigned by its chemical shift⁸ without the help of the time-consuming ^1H -undecoupled spectrum. The lowest field signal of (2) in the amide region is attributed, in view of the β -effects, to that originating from its

TABLE. ^{15}N N.m.r. chemical shifts (p.p.m.) in CDCl_3 -MeOH (8:2) solution, upfield from external $^{15}\text{NO}_3^-$ in H_2O and tentative signal assignments for the antibiotics siomycin-A (1) and thiostrepton (2).^a

(1)	Nitrogen number	Assignment	(2)
55.2; 63.6; 64.1; 66.6	4; 5; 7; 10 ^b	Azine-nitrogens	54.9; 63.5; 64.1; 66.1
72.0; 85.3; 87.5	12; 14; 15		72.2; 84.8; 87.4
250.1	17	Backbone Deala following Val	—
253.1	6	Thiostreptonic acid (ThstA) unit	253.2
253.8; 253.8	2; 3	Side-chain Deala	253.9; 253.9
—	18	Backbone Deala	254.8
—	17	Ala following Ile	255.3
256.8	19	Ala	256.8
259.2	9	Dehydrobutyrin (Debut)	259.3
259.5	18	Backbone Deala preceding Ala	—
262.9; 264.5	8; 13 ^b	Thr	263.1; 264.7
265.6	11	Thiostreptine (Thstn) unit	265.4
280.4	1	Side-chain CONH_2	280.9
—	16	Ile	331.4
336.5	16	Val	—

^a ^{15}N N.m.r. spectra of the antibiotics were recorded at 36.48 MHz and 25 °C with a Bruker WH-360 Fourier transform n.m.r. spectrometer using 1 g of substance in 4 ml of CDCl_3 -MeOH (8:2) and 15 mm diameter sample tubes. Broad-band ^1H decoupling (2.7 W) was employed in the inverse-gated mode to provide a decoupled spectrum but without nuclear Overhauser enhancement. A spectral width of 14,000 Hz was acquired in 16K data points using a 50 μs (ca. 70° flip angle) pulse width. The acquisition time (with ^1H decoupling) of 0.57 s was preceded by a delay (without decoupling) of 3 s. A total of 15,500 scans were accumulated and transformed with an exponential line broadening of 0.6 Hz. ^b Specific assignments cannot be made.

for the amine and for all the amide nitrogens. As a consequence of the loss of a deshielding δ -effect the Val-nitrogen of (1) appears 5.1 p.p.m. upfield relative to the Ile-nitrogen signal of (2). From the observed chemical shifts Deala-nitrogen resonances are close to those of Ala-nitrogens. The dehydrobutyrin (Debut)-nitrogen resonates a little upfield because of one shielding γ -effect, the Thr-nitrogens further upfield because of two such effects, and the thiostreptine (Thstn)-nitrogen at the highest field because of three.

Signal assignment of the two backbone Deala-nitrogens of (1) is not unambiguous. It has been established that a given amino-acid nitrogen involved in a peptide resonates at the lowest possible field when the contributions from the preceding residue across a carbonyl group amount to two δ -effects and no shielding ϵ -effect.¹¹ Since this is the

thiostreptonic acid (ThstA) portion. The thiazole, pyridine, and C=N nitrogens could not be unambiguously differentiated.¹⁰

On the basis of the highly sensitive ^{15}N n.m.r. chemical shift identities and differences between the spectrum of (1) and (2), the earlier proposal for the structure of siomycin-A as (1)^{2,3} can be accepted with confidence. The ^{15}N signal assignment presented here will be of great help in future structural investigations of antibiotics of the thiostrepton family.

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