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Complete ^1H and ^{13}C NMR Chemical Shift Assignments for Some N-Polyformylated Aminoglycoside Antibiotics

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**COMPLETE ^1H AND ^{13}C NMR CHEMICAL SHIFT ASSIGNMENTS FOR
SOME N-POLYFORMYLATED AMINOGLYCOSIDE ANTIBIOTICS**

KEY WORDS: Aminoglycoside Antibiotics, ^1H NMR, ^{13}C NMR,
Formylation shifts.

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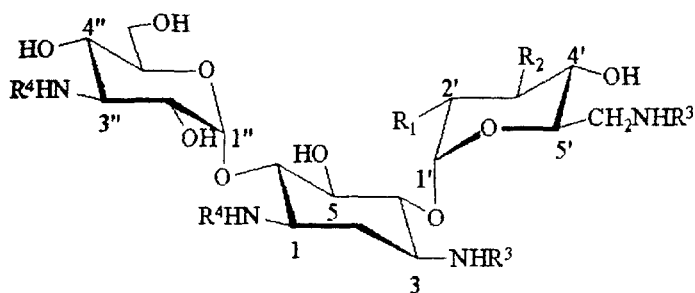
ABSTRACT

The complete ^1H and ^{13}C NMR assignments of a series of semisynthetic N-polyformyl derivatives of aminoglycoside antibiotics kanamycin A, kanamycin B, tobramycin, apramycin and sisomicin were achieved by 1D and 2D NMR methods (mainly ^1H - ^1H and ^1H - ^{13}C shift-correlation spectroscopy).

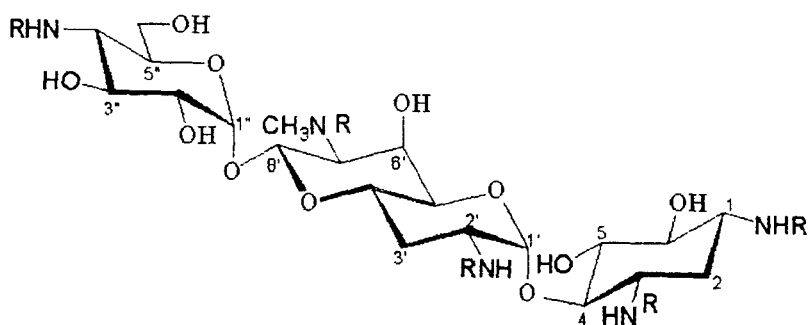
INTRODUCTION

Kanamycin A (1), kanamycin B (2), tobramycin (3), apramycin (4) and sisomicin (5) - medically important members of the aminoglycoside family of antibiotics have been studied extensively by ^1H and ^{13}C NMR methods [1 - 5].

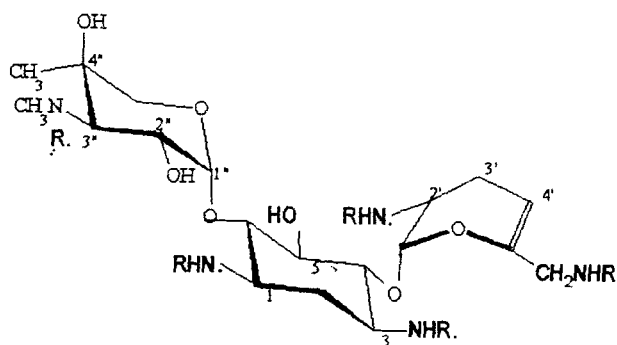
We reported the complete assignments of the ^1H and ^{13}C NMR spectra of antibiotics 1-5 and of a series of their semisynthetic N-polyacetyl- and N,O-polyacetyl derivatives [6]. The aim of this work was to achieve a complete assignments of N-polyformylated derivatives 6-11 of aminoglycoside antibiotics 1-5.



1 Kanamycin A	$\text{R}^1=\text{R}^2=\text{OH}; \text{R}^3=\text{R}^4=\text{H}$
2 Kanamycin B	$\text{R}^1=\text{NH}_2; \text{R}^2=\text{OH}; \text{R}^3=\text{R}^4=\text{H}$
3 Tobramycin	$\text{R}^1=\text{NH}_2; \text{R}^2=\text{R}^3=\text{R}^4=\text{H}$
6 Tetra-N-formylkanamycin A	$\text{R}^1=\text{R}^2=\text{OH}; \text{R}^3=\text{R}^4=\text{CHO}$
7 Penta-N-formylkanamycin B	$\text{R}^1=\text{NHCHO}; \text{R}^2=\text{OH}; \text{R}^3=\text{R}^4=\text{CHO}$
8 Tetra-N-formylkanamycin B	$\text{R}^1=\text{NHCHO}; \text{R}^2=\text{OH}; \text{R}^3=\text{CHO}; \text{R}^4=\text{H}$
9 Penta-N-formyltobramycin	$\text{R}^1=\text{NHCHO}; \text{R}^2=\text{H}; \text{R}^3=\text{R}^4=\text{CHO}$



- 4 Apramycin $R=H$
 10 Penta-N-formylapramycin $R=CHO$



- 5 Sisomicin $R=H$
 11 Penta-N-formylsisomicin $R=CHO$

EXPERIMENTAL

NMR Spectra

The NMR spectra were obtained on a Varian XL-300 spectrometer operating at 299.94 and 75.43 MHz for ^1H and ^{13}C , respectively, using 0.05–0.1 M solutions in D_2O . The proton chemical shifts are referenced to internal sodium 4,4-dimethyl-4-silapentane-1-sulphonate, and external tetramethylsilane (TMS) was used for the ^{13}C chemical shifts. The measurements were carried out at ambient temperature (ca 300 K) and pH 5–7 for the formyl derivatives **6–11**. The 2D NMR spectra were obtained using the standard Varian software: COSY, phase-sensitive COSY (COSYPS) and phase-sensitive double-quantum filtered COSY (DQ COSY) for the ^1H – ^1H correlations, HETCOR for the ^1H – ^{13}C correlations.

Typical conditions for the 1D ^1H spectra were: pulse width 32° , FT size 32K and digital resolution 0.2 Hz per point, and for the 1D ^{13}C spectra pulse width 60° , FT size 64K and digital resolution 0.5 Hz per point. The COSY experiments were typically performed with a spectral width of ca 1500 Hz, relaxation delay 1.3 s, mixing pulse width 45 or 60° , number of increments 256 or 512 and FT size 1K x 1K. The HETCOR experiments were carried out with a spectral width of ca 6000 Hz for ^{13}C and ca 1500 Hz for ^1H , relaxation delay 1.4 s, number of increments 128, FT size 4K x 256W, H-H decoupled multiplets.

Compounds

Samples of kanamycin A (**1**), kanamycin B (**2**), tobramycin (**3**) and sisomicin (**5**) were obtained commercially as sulphates and converted into the free bases via ion-exchange chromatography [8]. Apramycin (**4**) was isolated from the nebramycin complex (Antibiotics Factory, Razgrad, Bulgaria) and purified by ion-exchange chromatography [9].

The N-formylated compounds **6–11** were prepared by the method described for N-formylation of kanamycin A and B [7]. The compounds **6–11** were purified by ion-exchange chromatography (Amberlite CG-50, H^+ -

or NH_4^+ -form). The purities of the compounds were checked by thin-layer chromatography: Silica gel G-60, mobile phase - acetone : ethanol : acetic acid : water (1:1:0.5:0.5). Their structures were verified by ^1H and ^{13}C NMR studies. Compound **6**, m.p. 188-190°C (decomp.), $[\alpha]^{20\text{D}} +110^\circ$ (c 0.5, H_2O), R_f 0.69; **7**, m.p. 197-200°C (decomp.), $[\alpha]^{20\text{D}} +118^\circ$ (c 0.5, H_2O), R_f 0.72; **8**, m.p. 195-198°C (decomp.), $[\alpha]^{20\text{D}} +102.7^\circ$ (c 0.5, H_2O), R_f 0.57; **9**, m.p. 198-201°C (decomp.), $[\alpha]^{20\text{D}} +102.5^\circ$ (c 0.5, H_2O), R_f 0.78; **10**, m.p. 258-261°C (decomp.), $[\alpha]^{20\text{D}} 0^\circ$ (c 0.5, H_2O), R_f 0.50; **11**, m.p. 195-197°C (decomp.), $[\alpha]^{20\text{D}} +120.9^\circ$ (c 0.5, H_2O), R_f 0.82.

RESULTS AND DISCUSSION

The ^1H and ^{13}C chemical shifts obtained for the N-polyformylated derivatives **6-11** are presented in Tables 1 and 2. For comparison purposes, data for the parent compounds **1-5** taken from Ref. 6 are also included. An additional complication in the cases of formyl derivatives **6-11** was the observation of signals for more than one rotamer due to hindered rotation about the C-N bonds at ambient temperature, as already reported for some aminoglycoside antibiotics and their acetyl derivatives [6]. Only the signals belonging to the major rotamer (amounting to ca. 70% in the various compounds) are reported.

^{13}C chemical shifts

The ^{13}C -NMR chemical shifts of compounds **1-11** are presented in Table 1. The ^{13}C -NMR chemical shifts for antibiotics **1-5** are in agreement with the data from our earlier paper [6] and with the literature data [2-5]. As can be seen from Table 1, the N-formylation of **1-5** in most cases causes upfield shifts of the ^{13}C signals. The shift is more pronounced for the carbons in a β -position to the amino groups, in agreement with earlier observations [6]. The strongest shifts (4.5-7.5 ppm) are observed for the deoxystreptamine carbons participating in the glycosidic bonds (C-4 and C-6). Anomeric carbons with β -amino groups

TABLE I.
 ^{13}C NMR Chemical Shifts (ppm from TMS) for Compounds 1-11.

C	1	6	2	7	8	3	9	4	10	5	11
1	51.3	48.6	50.6	48.4	50.7	50.4	48.4	50.2	48.7	50.9	48.8
2	36.3	33.5	35.4	33.5	34.8	35.5	33.5	35.4	31.9	35.3	33.2
3	49.8	47.5	49.4	47.4	47.3	49.2	47.5	49.3	47.7	49.3	46.8
4	83.2	82.8	86.1	80.7	80.9	86.5	80.7	86.7	82.3	84.3	79.8
5	74.9	75.9	74.6	76.0	76.0	74.5	76.1	75.9	77.0	74.5	75.8
6	88.6	81.1	88.0	81.1	88.2	88.1	81.1	77.3	74.9	86.9	80.5
1'	100.3	99.1	100.1	98.4	98.4	99.7	97.0	100.6	98.0	99.9	96.9
2'	72.7	73.3	55.4	52.8	52.9	49.4	47.1	48.9	46.9	46.5	44.9
3'	73.8	73.0	73.6	71.7	71.7	34.9	32.3	31.8	29.6	24.6	21.2
4'	71.9	72.5	71.5	71.2	71.2	66.2	65.8	67.1	67.2	94.4	97.7
5'	73.8	71.3	72.3	71.5	71.4	73.7	72.0	70.1	70.0	148.9	145.8
6'	42.4	38.9	41.6	38.8	39.0	41.6	38.9	65.3	70.0	42.3	40.2
7'								61.4	60.9		
8'								95.5	97.2		
1"	100.8	99.1	100.1	98.7	98.7	100.0	98.5	94.5	94.8	100.6	99.7
2"	72.7	70.2	71.9	70.1	70.1	71.8	70.1	70.8	71.3	69.3	64.2
3"	55.0	53.6	54.4	53.3	53.6	54.2	53.4	73.3	69.4	63.3	53.9
4"	70.2	68.0	69.3	68.0	67.9	69.3	68.0	52.2	52.5	72.3	73.8
5"	72.9	71.3	72.3	73.0	72.9	72.1	73.1	72.5	68.9	67.7	69.3
6"	61.2	60.9	60.4	60.9	60.7	60.3	60.8	60.9	63.3	21.6	22.1
7"								32.0	33.2	36.9	33.0

TABLE 2.
1H NMR Chemical Shifts (ppm from TMS) of Compounds 1-11.

H	1	6	2	7	8	3	9	4	10	5	11
1	2.90	4.05	2.86	4.05	2.90	2.90	4.10	2.74	4.00	2.80	4.10
2ax	1.22	1.65	1.22	1.65	1.45	1.23	1.70	1.23	1.70	1.18	1.65
2eq	1.96	2.05	1.94	2.05	2.00	1.96	2.05	2.00	2.05	1.95	2.10
3	2.89	4.05	2.87	4.05	4.00	2.90	4.10	2.87	3.90	2.76	3.95
4	3.33	3.53	3.32	3.55	3.50	3.33	3.50	3.31	3.60	3.44	3.60
5	3.66	3.81	3.65	3.75	3.75	3.63	3.80	3.49	3.60	3.55	3.70
6	3.25	3.63	3.24	3.61	3.25	3.25	3.65	3.16	3.40	3.25	3.60
1'	5.33	5.34	5.35	5.38	5.33	5.16	5.27	5.16	5.20	5.35	5.43
2'	3.59	3.50	2.77	4.00	4.00	2.90	4.10	3.02	4.10	3.07	4.12
3'ax	3.70	3.69	3.56	3.70	3.65	1.61	1.70	1.67	1.80	1.95	2.00
3'eq						2.03	2.05	2.12	2.10	2.20	2.10
4'	3.31	3.28	3.30	3.38	3.25	3.54	3.55	3.78	3.85	4.88	4.85
5'	3.75	3.70	3.75	3.70	3.70	3.58		3.67	3.50		
6'	2.78	3.50	2.87	3.50	3.50	2.73	3.50	4.28	4.30	3.17	3.82
	3.00		3.07			2.97					
7'								2.68	3.65		
8'								4.92	5.40		
1''	5.04	5.17	5.03	5.20	5.16	5.05	5.30	5.37	5.40	5.07	5.27
2''	3.50	3.55	3.49	3.53	3.54	3.51	3.58	3.59	3.60	3.80	4.20
3''	3.01	4.12	2.99	4.14	4.00	3.01	4.14	3.65	3.80	2.78	4.40
4''	3.33	3.56	3.32	3.51	3.52	3.33	3.50	2.74	3.70		
5''	3.92	4.07	3.90	4.05	4.00	3.92	4.05	3.65	3.80	3.31	3.35
										4.05	4.15
6''	3.75	3.75	3.75	3.79	3.75	3.77	3.78	3.72		1.20	1.10
								3.85			
7''								2.38	3.03	2.52	2.96
											3.11

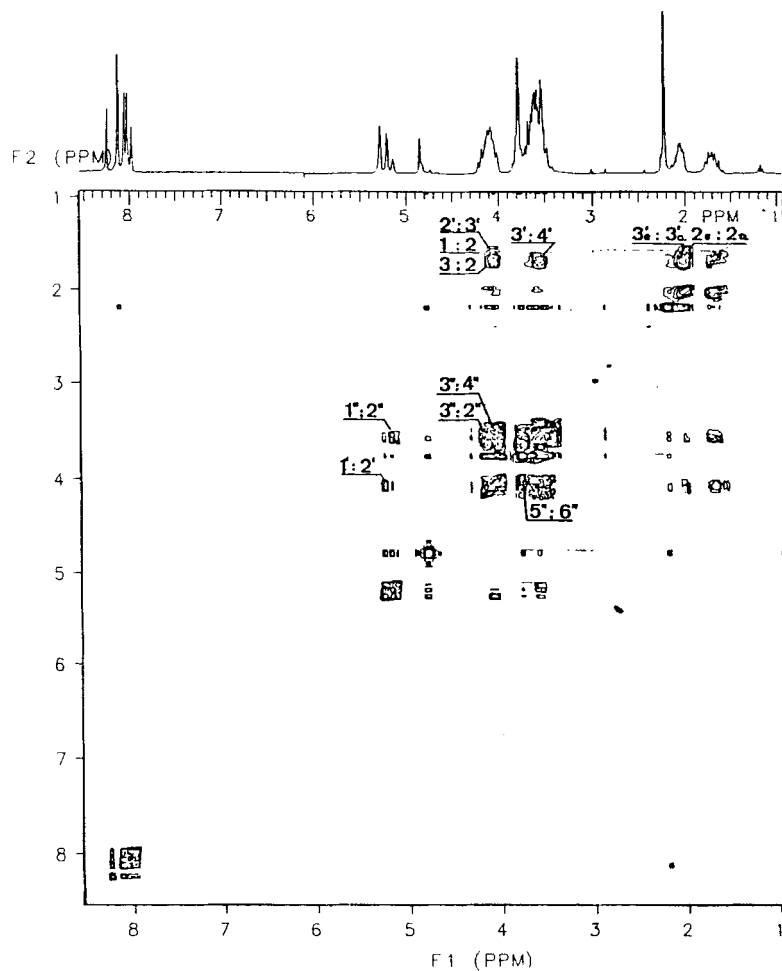


FIG. 1. 300 MHz 2D ^1H - ^1H Correlation Spectrum (COSY) of Penta-N-formyltobramycin (**9**) in D_2O with Indication for Some of the Cross-peaks.

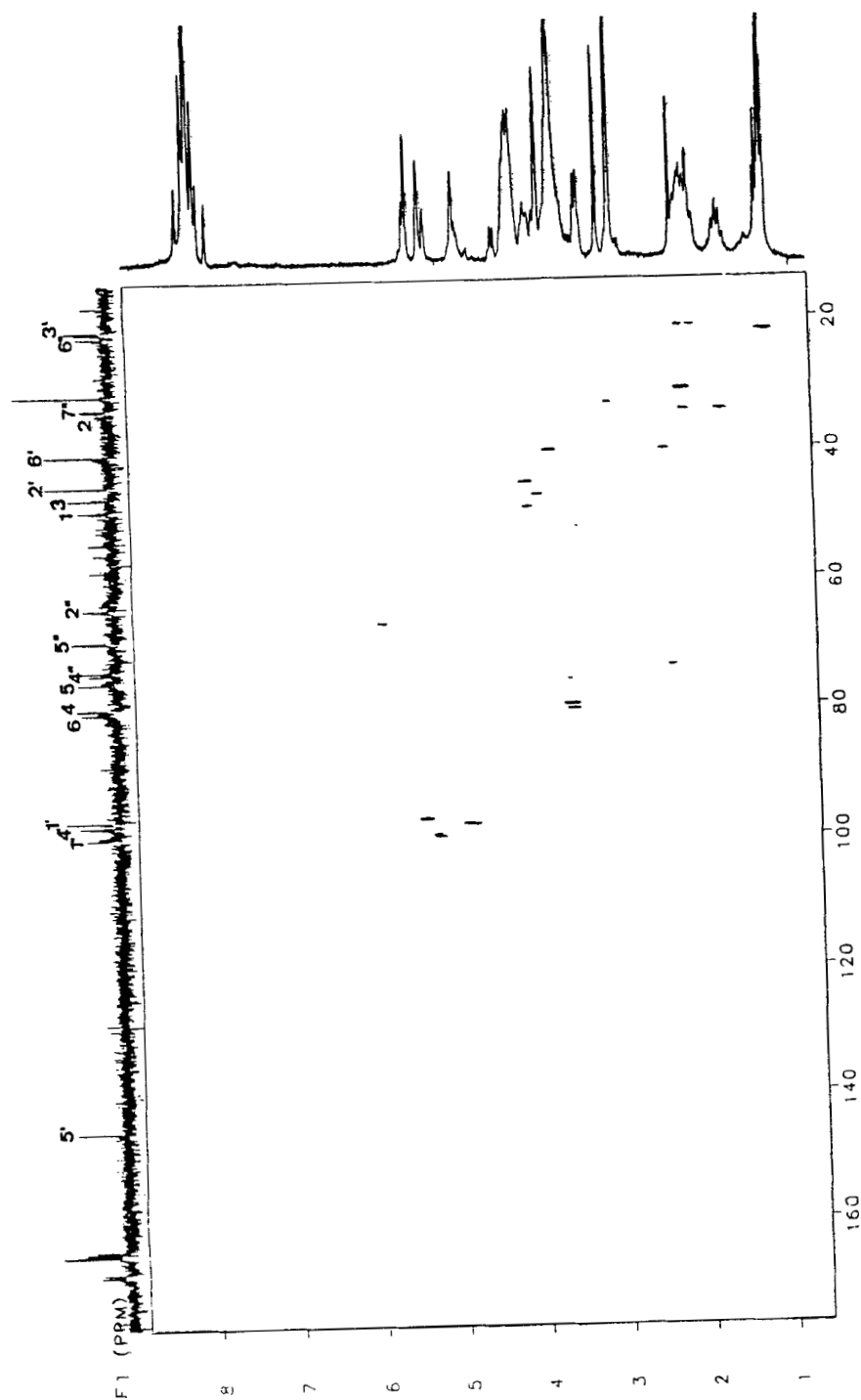


FIG. 2. 300/75 MHz 2D ^1H - ^{13}C Correlation Spectrum of Penta-N-formylisomycin (**11**) in D_2O .

are shifted upfield by 1.7 - 3 ppm, whereas C-2 is shifted by 2.0 - 2.8 ppm. The formylation effect on the α -carbons is less pronounced (0.8-3.5 ppm). It is strongest (2.7-3.5 ppm) for the methylene carbons, weaker (1.7 - 2.7 ppm) for carbons adjacent to a CH₂ and a C-OH group and weakest (0.8 - 1.4 ppm) for carbons between two C-OH groups. The data for tetra-N-formyl-kanamycin A (**6**) are in very good agreement with those published by Horii et al. [7] for some carbons of **6**.

¹H chemical shifts

The results of the analyses of the proton spectra of compounds **1-11** are presented in Table 2. The spectral assignment and multiplet analysis were based primarily on the ¹H-¹H 2D correlation spectra (COSY, FIG. 1) in the normal, phase-sensitive and double-quantum filtered versions. In the cases when the ¹³C assignments were known (**1-6**) ¹H-¹³C 2D heterocorrelated spectra (FIG. 2) were also helpful.

The N-formylation of **1-5** causes deshielding of the protons attached to α -carbons by 1.1 - 1.2 ppm for the CH, and by 0.4 - 0.6 ppm for the CH₂. Unusually large deshielding (1.6 ppm) was observed for H-3" of **10** (assignment also conformed via selective decoupling). A reliable determination of coupling constants was not possible due to the superposition of signals of the rotamers.

CONCLUSION

The results from the present investigation showed that the effect of N-formylation on the ¹H and ¹³C chemical shifts of aminoglycoside antibiotics **1-5** exhibits qualitatively similar trends as in the case of the N-acetylated derivatives. However, the magnitude of the α -shifts is larger for the formyl derivatives as compared to the acetylated compounds, whereas in the case of the β -shifts the opposite is generally true, although in the latter case the effects are less systematic.

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