

Complete Analysis of the Fast Fourier Transformed 100 MHz ^1H Nuclear Magnetic Resonance Spectra of Oxidized and Reduced Pyridine Mononucleotides

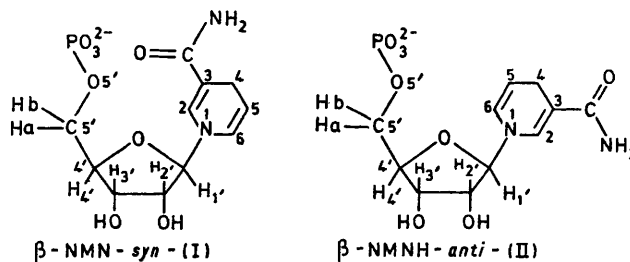
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Summary Complete analysis of the fast Fourier transformed ^1H n.m.r. spectra of oxidized and reduced pyridine mononucleotides show significant differences in their solution conformation.

THE ^1H 8K fast Fourier transformed (FFT) 100 MHz n.m.r. spectrum of oxidized pyridine mononucleotide, β -NMN (I), along with the LAOCN III computer simulated ribose region is shown in the Figure. The chemical shifts and coupling constants at two pH values are shown in the Table. The previous assignment¹ of the C (1') H resonance of reduced coenzyme (*e.g.* β -NMNH; II) was found to be erroneous. The chemical shifts of the various protons of β -NMN (0.4—0.0025M) and β -NMNH (0.1—0.01M) showed a concentration dependence of less than 1—2 Hz indicating

that these nucleotides do not undergo intermolecular stacking. The magnitude of the coupling constants



$J_{5a}^{-31\text{P}}$ and $J_{5b}^{-31\text{P}}$ for β -NMN and β -NMNH (Table) show, according to Tsuboi *et al.*,² that the torsional isomer

TABLE

Chemical shifts and coupling constants of β -NMN and β -NMNH; 0.1 M-solution 100% D_2O . Data from 100 MHz spectra.
Chemical shifts (δ /Hz) Coupling constants (J/Hz)

Nucleotides	Proton	$\Delta\delta$			Nuclei Coupled	Coupling constants (J/Hz)	
		pH 8.3	pH 4.0	pH 8.3 \rightarrow 4.0		pH 8.3	pH 4.0
β -NMN	2-H	640.9	628.7	12.2	2-4	1.5	1.5
	4-H	580.9	581.2	0	2-5	N.d.	N.d.
	5-H	512.6	512.6	0	2-6	N.d.	N.d.
	6-H	615.6	610.9	4.7	4-5	8.2	8
	1'-H	300.4	304.4	-4.0	4-6	1.5	1.5
	2'-H	148.1	139.0	9.1	5-6	6.3	6.3
	3'-H	129.2	127.0	2.2	1'-2'	5.7	5.2
	4'-H	142.0	147.9	-5.9	2'-3'	5.0	5.0
	5a'-H	101.3	112.8	-11.4	3'-4'	2.1	2.6
	5b'-H	84.2	97.3	-13.1	4'-5a'	2.6	2.4
					4'-5b'	2.0	2.3
					5a'- ^{31}P	4.2	4.4
					5b'- ^{31}P	4.4	5.1
				5a'-5b'	12.1	12.1	
β -NMNH ^b	2-H	397.5	396.1	1.4	2-6	1.7	1.6
	6-H	304.6	300.2	4.4	5-6	8.2	8.1
	5-H	184.3	183.7	0.6	4-6	1.6	1.8
	2 \times 4-H	-11.7	-12.0	0	4-5	3.5	3.4
	1'-H	170.4	171.9	-1.5	2-5	Not resolved	
	2'-H	115.1	109.3	3.8	2-4		
	3'-H	106.4	104.7	1.7	1'-2'	6.8	6.8
	4'-H	90.0	92.2	-2.2	2'-3'	5.4	5.4
	5a'-H	67.5	79.5	-12.0	3'-4'	1.9 ^c	1.8
	5b'-H ^d	67.5	79.5	-12.0	4'-5a' ^d	4.0	3.0
					4'-5b'		
					5a'- ^{31}P ^d	4.8	5.1
					5b'- ^{31}P		

^a Chemical shifts are accurate to at least 0.2 Hz; coupling constants to 0.1 Hz. For β -NMNH (low pH), chemical shifts are accurate to 0.3 Hz, coupling constants to 0.2 Hz. ^b pH 8.1 and 5.0. ^c From 220 MHz n.m.r. spectrum. ^d No difference in chemical shifts observable between 5a'-H and 5b'-H even at 220 MHz.

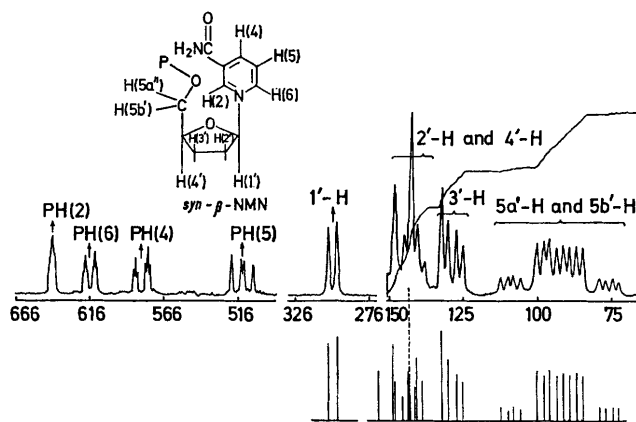


FIGURE. 8K Fourier-transformed 1H n.m.r. spectrum of 0.1M-solution of β -NMN in '100%' D_2O , pH 8.3; number of pulses 500; pulse width 40 μs ; sampling frequency 2000 $Hz s^{-1}$. P = pyridine. Chemical shifts are expressed in Hz downfield from $Me_4N^+Cl^-$; 100 MHz n.m.r. system. The spectrum was recorded using a Digilab FTS-NMR-3-system interfaced to a Varian HA-100D NMR spectrometer, with a ^{19}F lock. The total memory of the data system was 132K permitting a maximum of 32K transforms (double precision 32 bits per word length). Bottom part shows the computer simulated spectrum of the ribose region of β -NMN. One of the transitions visible in the simulation was obscured by the HDO resonance.

constrained to the O(5')-C(5') bond of β -NMN and β -NMNH is predominantly *gauche-gauche*. Calculations³ of the torsional isomer population around the C(5')-C(4') bond from the $J_{4'-5a'}$ and $J_{4'-5b'}$ values show that over 90% of β -NMN molecules exist in the *gauche-gauche* form constrained to the C(5')-C(4') bond whereas in the case of β -NMNH at pH 8.0 only 50% of the molecules exhibit *gauche-gauche* conformation. A qualitative application of Karplus equation to the $J_{1'-2'}$, $J_{2'-3'}$, and $J_{3'-4'}$ (Table) values of β -NMN and β -NMNH indicate that the ribose fragment of both compounds exists as an equilibrium mixture of 3'-*endo*- and 2'-*endo*-structures, the 2'-*endo*-population being larger for β -NMNH. The pH profiles for β -NMN (pH range 2-9, increments of 0.5 pH units) reveal that the protonation of the phosphate group (the pK of the phosphate group was determined by ^{31}P n.m.r. measurements by the method in Sarma and Mynott⁴) perturbs both pyridine 2-H and 6-H resonance, the perturbation of 2-H being about three times larger than that of 6-H. Thus the data clearly indicate that β -NMN prefers to exist in the *syn*-form (I), as has been suggested by Sarma and Kaplan⁵ from crude pH studies. Qualitative calculations show that about 78% of the molecules exist in the *syn*- and 22% in the *anti*-conformation. No extensive ^{31}P and 1H n.m.r. studies as a function of pH is possible for β -NMNH since this nucleotide decomposes at pH values below 6.0. However, the FFT mode of operation enabled us to obtain high quality spectra very quickly at pH 5.0. The data for β -NMNH (Table) indicate that this nucleotide preferentially exists in the *anti* conformation (II), contrary to the predictions of Sarma and Kaplan⁵ from

ring current considerations. The data in the Table make possible a complete analysis by ^{13}C FFT n.m.r. methods of the ^{13}C n.m.r. spectra of β -NMN and β -NMNH by proton off-resonance spin-decoupling.⁶

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