

# Enzymatic synthesis of polymers containing nicotinamide mononucleotide

Rihe Liu and Leslie E. Orgel\*

The Salk Institute for Biological Studies, PO Box 85800, San Diego, CA 92186-5800, USA

Received May 15, 1995; Revised and Accepted August 9, 1995

## ABSTRACT

Nicotinamide mononucleoside 5'-diphosphate in its reduced form is an excellent substrate for polynucleotide phosphorylase from *Micrococcus luteus* both in *de novo* polymerization reactions and in primer extension reactions. The oxidized form of the diphosphate is a much less efficient substrate; it can be used to extend primers but does not oligomerize in the absence of a primer. The cyanide adduct of the oxidized substrate, like the reduced substrate, polymerizes efficiently. Loss of cyanide yields high molecular weight polymers of the oxidized form. Terminal transferase from calf thymus accepts nicotinamide mononucleoside 5'-triphosphate as a substrate and efficiently adds one residue to the 3'-end of an oligodeoxynucleotide. T4 polynucleotide kinase accepts oligomers of nicotinamide mononucleotide as substrates. However, RNA polymerases do not incorporate nicotinamide mononucleoside 5'-triphosphate into products on any of the templates that we used.

## INTRODUCTION

It is now well-established that RNA catalysts, ribozymes, can bring about many hydrolysis and group-transfer reactions (1–6). It has been speculated that the nicotinamide co-enzymes may be fossils of an RNA world, in which they were co-factors that extended the range of RNA-catalyzed reactions to include redox reactions (7,8). While it has usually been supposed that the co-enzymes were attached to RNA by hydrogen-bonding and other non-covalent interactions, the incorporation of nicotinamide mononucleotide (NMN<sup>+</sup>) within or at the end of RNA molecules is also feasible. The possibility that nicotinamide could substitute for adenine or guanine in a Watson–Crick base-pair (Fig. 1) suggests that template-directed incorporation of NMN<sup>+</sup> into an oligonucleotide might be possible.

In this paper we show that polymers of NMN<sup>+</sup> and of reduced nicotinamide mononucleotide (NMNH) can be prepared using polynucleotide phosphorylase from *Micrococcus luteus* (PNPase). In addition, we show that a single NMN<sup>+</sup> residue can be added to the 3'-terminus of an oligodeoxynucleotide using terminal transferase from calf thymus. We have not been able to

demonstrate template-directed reactions of NMN<sup>+</sup> derivatives with RNA polymerases or in non-enzymatic systems.

## MATERIALS AND METHODS

### Materials

All chemicals were reagent grade and were used without further purification. Nicotinamide mononucleotide was prepared from nicotinamide adenine dinucleotide (NAD<sup>+</sup>) by non-enzymatic hydrolysis with Zr<sup>4+</sup> (9). Adenosine 5'-phosphoro(2-methyl)imidazole (2-MeImpA), uridine 5'-phosphoro(2-methyl)imidazole (2-MeImpU), cytidine 5'-phosphoro(2-methyl)imidazole (2-MeImpC), guanosine 5'-phosphoro(2-methyl)imidazole (2-MeImpG) and nicotinamide mononucleoside 5'-phosphoro(2-methyl)imidazole (2-MeImpNMN<sup>+</sup>) were prepared by a published procedure (10).

Polynucleotide phosphorylase was obtained from Boehringer Mannheim, terminal deoxynucleotidyl transferase from Bethesda Research Laboratories, T4 polynucleotide kinase from New England Biolabs and T3 RNA polymerase, T7 RNA polymerase, ribonuclease A, ribonuclease T1, ribonuclease P1, and phosphodiesterase I from United States Biochemicals. [ $\alpha$ -<sup>32</sup>P]UTP, [ $\alpha$ -<sup>32</sup>P]GTP and [ $\gamma$ -<sup>32</sup>P]ATP were obtained from Amersham.

### Oligonucleotides

Oligodeoxyribonucleotides were synthesized on an Applied Biosystem Model 391A DNA synthesizer using 2-cyanoethyl phosphoramidites. Oligomers terminated by a ribonucleotide at their 3'-ends were synthesized and deprotected following the standard DNA synthesis procedures, using a solid support derivatized with a ribonucleotide (Biosearch). All oligonucleotides were purified on preparative 20% polyacrylamide gels. The concentrations of single-stranded oligomers were determined from their absorption at 260 nm using the following molar extinction coefficients: 15 400 (A), 11 700 (G), 7300 (C) and 8800 (T). <sup>32</sup>P-labelling was carried out by standard methods (11) using [ $\gamma$ -<sup>32</sup>P]ATP and T4 polynucleotide kinase.

### Chromatography

Low molecular weight products were separated by thin layer chromatography (TLC) on Silica gel 60 F<sub>254</sub> from Merck Darmstadt. The solvent systems were: (A) n-PrOH:NH<sub>4</sub>OH:H<sub>2</sub>O 22:4:14; (B) 95% EtOH:1 M NH<sub>4</sub>OAc, at pH 5.0, 7:3. Larger

\* To whom correspondence should be addressed



scale separations were carried out by descending paper chromatography on Whatman 3 MM paper using solvent systems A and B.

### Gel electrophoresis

Electrophoresis was carried out using 0.75 mm thick polyacrylamide gels (10–20%), and run at 1000 V (–15 mA) for 3–5 h. The gels were autoradiographed with Kodak XAR-5 film at –70°C.

### NMR analysis

NMR spectra were recorded on a GE QE-300 NMR using D<sub>2</sub>O (pD 10.8) as solvent. The [<sup>1</sup>H] chemical shifts were referenced to 3-(trimethylsilyl) propionate (TMSP) and the [<sup>31</sup>P] chemical shifts were referenced to H<sub>3</sub>PO<sub>4</sub>.

### Synthesis of 5'-diphosphate and 5'-triphosphate of NMN<sup>+</sup> (NMNDP<sup>+</sup> and NMNTP<sup>+</sup>)

NMNDP<sup>+</sup> and NMNTP<sup>+</sup> were prepared from NMN<sup>+</sup> by the procedure described by Hoard and Ott (12). The reactions were monitored by TLC (NMN<sup>+</sup>: R<sub>f</sub> = 0.28; NMNDP<sup>+</sup>: R<sub>f</sub> = 0.12; NMNTP<sup>+</sup>: R<sub>f</sub> = 0.09, solvent B). The products were separated on a column of DEAE–Sephadex using a linear gradient of triethylammonium hydrogen carbonate as eluent. The resulting oil was dissolved in methanol and added to a 0.2 M solution of NaClO<sub>4</sub> in acetone. The precipitated sodium salt was collected by centrifugation, washed with acetone and ether and dried over phosphorous pentoxide. The purity of the products was established by paper chromatography using the Hanes–Isherwood reagent as visualizing agent (13).

NMNDP<sup>+</sup>: UV (H<sub>2</sub>O, pH 5.0) λ<sub>max</sub> = 266 nm (log ε = 3.62)

[<sup>1</sup>H] NMR (D<sub>2</sub>O, pD 10.8) δ = 4.35 (m, 5'-H<sub>2</sub>); 4.54–4.70 (m, 2'-H, 3'-H, 4'-H); 6.20 (d, J = 4.6 Hz, 1'-H); 8.31 (pt, 5-H); 8.96 (d, J = 8.1 Hz, 4-H); 9.31 (d, J = 6.1 Hz, 6-H); 9.48 (s, 2-H).

[<sup>31</sup>P] NMR (D<sub>2</sub>O, pD 10.8) δ = –5.25 (d, J = 22 Hz, β-P), –10.12 (d, J = 22 Hz, α-P).

NMNTP<sup>+</sup>: UV (H<sub>2</sub>O, pH 5.0) λ<sub>max</sub> = 266 nm (log ε = 3.72).

[<sup>1</sup>H] NMR (D<sub>2</sub>O, pD 10.8) δ = 4.34 (m, 5'-H<sub>2</sub>); 4.54–4.70 (m, 2'-H, 3'-H, 4'-H); 6.15 (d, J = 5.3 Hz, 1'-H); 8.31 (pt, 5-H); 8.92 (d, J = 8.0 Hz, 4-H); 9.27 (d, J = 6.2 Hz, 6-H); 9.43 (s, 2-H).

[<sup>31</sup>P] NMR (D<sub>2</sub>O, pD 10.8) δ = –4.46 (d, J = 20 Hz, γ-P), –20.93 (t, J = 20 Hz, β-P), –10.67 (d, J = 10 Hz, α-P).

### Reduction of NMNDP<sup>+</sup> and NMNTP<sup>+</sup> and preparation of cyanide adduct of NMNDP<sup>+</sup>

The reduced products NMNDPH and NMNTPH were prepared using a modified version of the procedure of Gutcho and Stewart (14). 100 μl of the reduction buffer containing 200 mM Tris–HCl, pH 9.0 and 1.2% (w) sodium hydrosulfite was added to ~4 O.D. of the nucleotide. The reaction mixture was heated at 50°C for 5 min and immediately cooled on ice. Excess reducing agent was removed by aerating vigorously for 5 min at 0°C. The solution was evaporated to dryness and the residue redissolved in an appropriate amount of water. The 4-cyano adduct of NMNDP<sup>+</sup> (NMNDP–CN) was prepared by adding 20 μl 0.2 M KCN to 10 μl 0.1 M NMNDP<sup>+</sup> (pH 7.0). The survival of these products was monitored by UV spectroscopy (340 nm) (15).

### Enzymatic dephosphorylation of nicotinamide nucleotides and oligomers of NMN<sup>+</sup>

The mono-, di- and tri-phosphates of nicotinamide nucleoside and the oligomers formed in polynucleotide phosphorylase reaction mixtures were dephosphorylated using calf intestine alkaline phosphatase (CIAP) or phosphodiesterase I (PDE I).

*CIAP reaction.* A mixture containing 0.5 μmol (2.50 O.D.) of the nucleotide (NMN<sup>+</sup>, NMNDP<sup>+</sup>, NMNTP<sup>+</sup> or oligomers), 1× phosphorylation buffer (50 mM Tris–HCl, pH 8.5, 0.1 mM EDTA) and 10 U of CIAP in a total volume of 50 μl was incubated for an appropriate time at 37°C.

*PDE I reaction.* The corresponding nucleotide (2.50 O.D.) was incubated for an appropriate time with 10 U of phosphodiesterase I in a Tris buffer (Tris–HCl, 100 mM, pH 9.0; NaCl, 100 mM; MgCl<sub>2</sub>, 14 mM) at 25°C.

The reactions were monitored by TLC in solvent B.

### Polynucleotide phosphorylase reactions

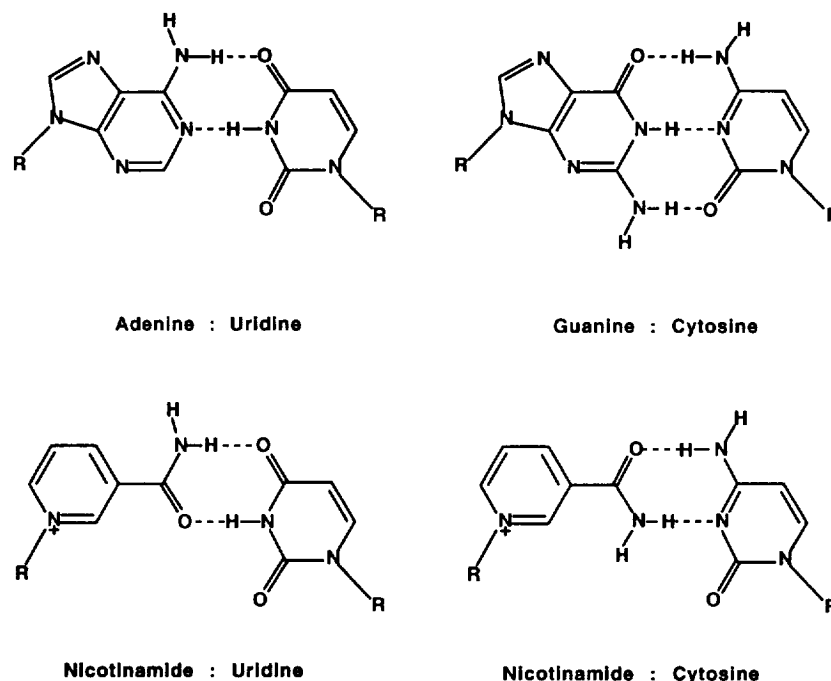
The primer-initiated polynucleotide phosphorylase catalyzed reactions were performed as follows. Each polymerization reaction contained ~2 nM of 5'-<sup>32</sup>P-labelled oligonucleotide as primer and an appropriate concentration of nicotinamide nucleotide in 100 mM Tris–HCl, pH 9.0 (at 37°C), 25 mM MgCl<sub>2</sub> or MnCl<sub>2</sub> and 0.5 mM EDTA. The concentration of polynucleotide phosphorylase was 20 U/ml. After an appropriate time at 37°C, the reactions were quenched by addition of 7.5 μl of 0.5 M EDTA, pH 8.0 and then loaded onto an analytical polyacrylamide gel (15% for the oxidized form and 10% for the reduced form and cyanide adduct).

The *de novo* polymerization reactions were carried out under the above conditions, but without primer. The reaction mixtures were extracted with phenol/chloroform to remove the enzyme, passed through a Nensorb column and then lyophilized. The solid was dissolved in TE buffer in preparation for kination. A mixture containing an aliquot of the polymeric product (0.05 O.D.), 1× kinase buffer, 20 μCi [γ-<sup>32</sup>P]ATP, and 20 U polynucleotide kinase in a total volume of 50 μl was incubated at 37°C for 1.5 h. The reaction was stopped by addition of 2 μl 0.5 M EDTA, pH 8.0 and the product was desalted using a Nensorb column. Another aliquot of the polymeric product was first dephosphorylated by CIAP, purified on a Nensorb column and then kinated using the same procedure as above. The final products were analyzed on a 10% polyacrylamide gel.

### Nuclease digestion of products

*Ribonuclease T1 digestion.* An aliquot of the purified polymeric product (2000 c.p.m.) was dissolved in a Tris buffer (5 μl) containing Tris–HCl (0.05 M, pH 7.5) and EDTA (2 mM, pH 8.0). The solution was incubated at 37°C for 5 min. Ribonuclease T1 (1 μl, 5 U/μl) was added and the reaction mixture was incubated at 37°C for 1 h. The resulting solution was mixed with loading buffer (5 μl) and then analyzed on a 15% polyacrylamide gel.

*Ribonuclease A digestion.* The purified polymeric product (2000 c.p.m.) was dissolved in an acetate buffer (10 μl, sodium acetate, 50 mM, pH 5.0; tRNA, 0.5 mg/ml). After adding ribonuclease A (1 μl, 18 U/μl), the solution was incubated at 37°C for 1 h.



**Figure 1.** Schematic comparison of nicotinamide and purines.

Samples of the resulting solution were evaporated to dryness, redissolved in 3  $\mu$ l of water, mixed with 3  $\mu$ l of loading buffer and subjected to electrophoresis on 15% polyacrylamide gels.

**Phosphodiesterase I digestion.** A sample of the purified polymeric product (2000 c.p.m.) was incubated in a Tris buffer (5  $\mu$ l, Tris-HCl, 100 mM, pH 9.0; NaCl, 100 mM; MgCl<sub>2</sub>, 14 mM) with 1 U of phosphodiesterase I at 25°C for 1 h. The reaction was stopped by addition of 1  $\mu$ l of 0.2 M EDTA, pH 8.0 and analyzed on a 15% polyacrylamide gel as described above.

### Terminal transferase reactions

Terminal deoxynucleotidyl transferase catalyzed addition reactions were carried out according to a procedure of Deng and Wu (16). An appropriate amount of 5'-labelled primer ([<sup>32</sup>P]-(dA)<sub>6</sub>) was incubated with NMNDP<sup>+</sup> (various concentrations) and terminal transferase (0.8 U/ $\mu$ l) at 37°C for 2.5 h in 2 mM cobalt chloride buffer, pH 7.0. The reactions were stopped by addition of an appropriate amount of EDTA and samples were then subjected to electrophoresis on 20% polyacrylamide gels.

### Transcription reactions

*In vitro* transcription reactions were carried out according to the procedure given by Milligan *et al.* (17). The transcription templates were created by heating a 1:1.2 mixture of complementary single strands (top strand and 18mer promoter) in TE buffer to 90°C for 2 min and cooling quickly on ice. Each reaction contained 500 nM template and 150 U of T3 or T7 RNA polymerase in 20  $\mu$ l of 40 mM Tris-HCl, pH 8.0 (at 37°C), 20 mM MgCl<sub>2</sub>, 0.5 mg/ml BSA, 5 mM DTT, 20 U RNase inhibitor, 1 mM standard nucleoside triphosphates, 5 mM nicotinamide nucleoside triphosphate and 2  $\mu$ Ci [ $\alpha$ -<sup>32</sup>P]UTP or [ $\gamma$ -<sup>32</sup>P]GTP. Reaction mixtures were incubated at 37°C for 2.5 h, quenched by

addition of 2.4  $\mu$ l of 0.5 M EDTA, pH 8.0 and loaded on an analytical 20% denaturing polyacrylamide gel.

### Non-enzymatic template-directed reactions

Template-directed reactions were carried out by a modification of a published procedure (18). First, a solution (15  $\mu$ l) containing 20  $\mu$ mol NaCl, 4  $\mu$ mol MgCl<sub>2</sub>, and 5'-<sup>32</sup>P-labelled hairpin oligonucleotide (~240 000 c.p.m., ~0.2 pmol) was evaporated to dryness. The residue was dissolved in 14  $\mu$ l of 0.286 M 2,6-lutidine-HCl buffer, pH 8.0 (at 0°C). The template was annealed by heating at 50°C for 3 min, cooling for 3 min at room temperature and finally cooling to 0°C on an ice-bath. Next, 6  $\mu$ l of a freshly prepared solution of the relevant 5'-phosphoro(2-methyl)imidazolides was added and the mixture incubated at 0°C for 7 days. The concentration of 2-MeImpG was usually 20 mM and the concentration of the activated nicotinamide nucleotide 100 mM. The reaction was stopped by addition of 2.4  $\mu$ l of 0.5 M EDTA, pH 8.0, to 2  $\mu$ l of the reaction mixture. After mixing with 4.4  $\mu$ l of loading buffer, the final products were analyzed on a polyacrylamide gel.

## RESULTS

NMNDP<sup>+</sup> and NMNDP<sup>+</sup> were identified by UV, [<sup>1</sup>H]NMR and [<sup>31</sup>P]NMR spectroscopy. The maximum wavelengths of UV absorption of NMNDP<sup>+</sup> and NMNDP<sup>+</sup> are identical to that of NMNDP<sup>+</sup> (19). The [<sup>31</sup>P] chemical shifts and coupling constants are similar to those of normal nucleoside diphosphates and triphosphates (data above) (20).

The oxidized forms of the pyridine coenzymes are susceptible to base-catalyzed decomposition, whereas the reduced pyridine nucleotides are labile in acid (21–23). The stabilities of NMNDP<sup>+</sup> and NMNDP<sup>+</sup> were studied under the conditions used in the polynucleotide phosphorylase reaction (25 mM Mg<sup>2+</sup> or Mn<sup>2+</sup>, pH 9.0, 37°C, 0.5–24 h). The results of TLC and UV spectroscopy show that little decomposition of NMNDP<sup>+</sup> occurs within



**Figure 2.** Autoradiogram of the gel electrophoretogram of the products from the polymerization of NMNDP<sup>+</sup> by polynucleotide phosphorylase primed by p(rA)<sub>5</sub> after 24 h: lane 1 and lane 12, <sup>32</sup>P-labelled primer. Substrates: lane 2, 40 mM UDP; lanes 3–10, 5, 10, 20, 30, 40, 50, 60 and 100 mM NMNDP<sup>+</sup>, respectively; lane 11, 4 mM UDP and 16 mM NMNDP<sup>+</sup>. The concentration of PNPase was 20 U/ml.

the incubation time, but we observed ~30% decomposition of NMNDPH. In the latter case, a new UV absorption band is observed at 280 nm, presumably due to the presence of a compound in which water is added to the 5–6 double-bond of the nicotinamide ring (24).

### Polymerization of NMNDP<sup>+</sup> by polynucleotide phosphorylase

Primed polymerization of NMNDP<sup>+</sup> was successful in the presence of Mg<sup>2+</sup> or Mn<sup>2+</sup>, but Mg<sup>2+</sup> was more efficient than Mn<sup>2+</sup> in promoting the formation of long oligonucleotides (data not shown). In typical experiments 2–30 residues of NMN<sup>+</sup> are added to the primer. The abnormally large separations between successive bands on the gel indicate that monomers with positive charge are incorporated. The distribution of the product lengths is highly sensitive to the concentration of the substrate (Fig. 2). The optimal concentration of NMNDP<sup>+</sup> for the formation of longer oligomers is ~25 mM. When the NMNDP<sup>+</sup> concentration is >60



**Figure 3.** Autoradiogram of the gel electrophoretogram of the products from the polymerization of NMNDPH by polynucleotide phosphorylase using p(rA)<sub>5</sub> as primer after 4 h: lane 1, <sup>32</sup>P-labelled primer; lane 2, primer incubated with PNPase; lane 3, substrate, 1.5 mM UDP; lane 4, substrate, 1.5 mM UDP in reduction buffer; lane 5, substrate, 1.5 mM NMNDP<sup>+</sup>; lane 6, substrate, 1.5 mM NMNDPH. The reduction buffer was 1.2% (w) sodium hydrosulfite in 0.20 M Tris, pH 9.0. The concentration of PNPase was 20 U/ml.

mM, the majority of the products are very short. When it is significantly <5 mM, there is very little reaction (data not shown).

Extension of the following primers was studied: (rA)<sub>2</sub>, (rA)<sub>5</sub>, (dA)<sub>4</sub>rA and (dA)<sub>9</sub>rA. The optimal reaction conditions are different for different primers. It is found that (rA)<sub>5</sub> is the most efficient primer. When (rA)<sub>2</sub> is used as primer, the major products contain only two or three NMN<sup>+</sup> residues. The addition of NMN<sup>+</sup> to the 3'-terminus of a primer is rapid. When (rA)<sub>5</sub> is used as primer, the starting material disappears within 1 h at 37°C. The yield of longer products increases slowly as the incubation time is extended (data not shown).

In the absence of a primer, we did not detect the *de novo* polymerization of NMNDP<sup>+</sup>.

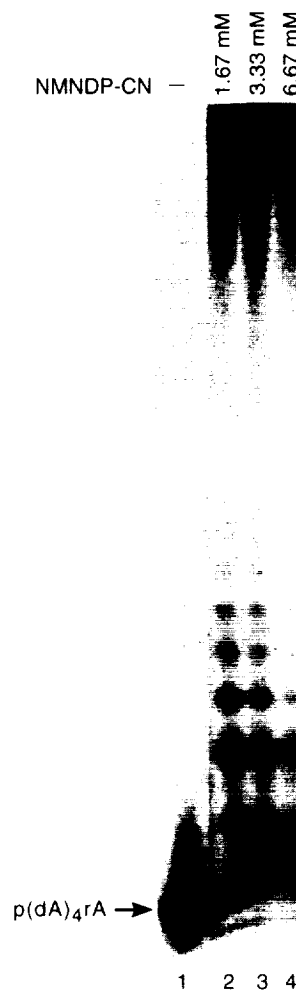


**Figure 4.** Autoradiogram of the products from the *de novo* polymerization of NMNDPH by polynucleotide phosphorylase. The polymeric products were kinased directly (lanes 1 and 2) or after dephosphorylation with alkaline phosphatase (lanes 3 and 4).

### Polymerization of NMNDPH by polynucleotide phosphorylase

Figure 3 illustrates the polymerization of NMNDPH by polynucleotide phosphorylase in the presence of (rA)<sub>5</sub> as primer. In contrast to NMNDP<sup>+</sup>, NMNDPH is an excellent substrate even at very low concentrations and the length of the products formed is much greater than with NMNDP<sup>+</sup>. After phenol/chloroform extraction of the product, we found that even on 10% denaturing polyacrylamide gels, a significant amount of the material remained in the wells. It can be seen from Figure 3 that the distribution of the polymer sizes is broad and that the majority of the products are large polymers (>200 bases). The polymerization rate of NMNDPH in the presence of primer is rapid. The primer disappears within half an hour (data not shown).

NMNDPH is also an excellent substrate for polynucleotide phosphorylase in the absence of a primer. Long oligomeric products that do not move away from the origin of a paper chromatogram are obtained in good yield. However, polymeric products are not detected in this *de novo* reaction before 8 h, indicating that the initiation of polymerization is slow.

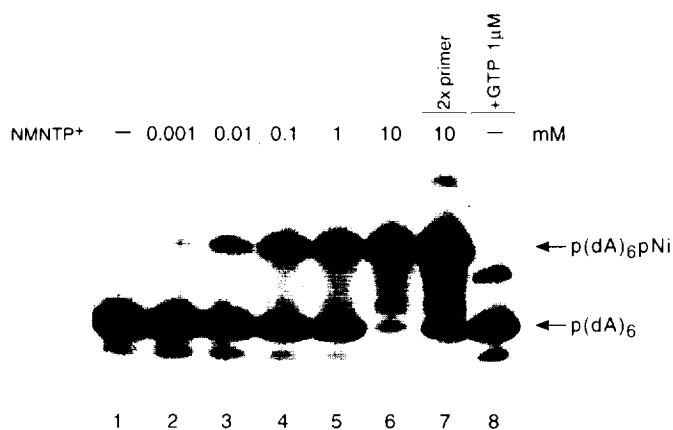


**Figure 5.** Autoradiogram of the products from the primed polymerization of NMNDP-CN by polynucleotide phosphorylase after 1 h: lane 1, <sup>32</sup>P-labelled primer p(dA)<sub>4</sub>rA; Substrate: 1.67 mM (lane 2), 3.33 mM (lane 3) and 6.67 mM (lane 4) NMNDP-CN. The ratio of the concentration of NDP and MgCl<sub>2</sub> was maintained at 2. The concentration of PNPase was 20 U/ml.

Further evidence for the *de novo* synthesis of high molecular weight poly(NMNDPH) was obtained by high-resolution gel electrophoresis. The purified products of the polymerization reaction are kinased directly and efficiently by T4 polynucleotide kinase (Fig. 4), presumably due to the Pi exchange activity of the kinase (25). Essentially the same pattern of products is obtained if the polymeric products are first treated with alkaline phosphatase, reisolated, and then kinased.

### Polymerization of the cyanide adduct and the hydration product of NMNDP<sup>+</sup> by polynucleotide phosphorylase

The gel electrophoretogram in Figure 5 illustrates results of the polymerization of NMNDP-CN in the presence of primer. The pattern of products is similar to that obtained from NMNDPH. As in the case of NMNDPH, retention of material at the origin of the electrophoretogram indicates the presence of very high molecular



**Figure 6.** Autoradiogram of the gel electrophoretogram of the products from the addition of NMNTP<sup>+</sup> to oligodeoxynucleotide p(dA)<sub>6</sub> by terminal deoxynucleotidyl transferase after 2.5 h: lane 1, <sup>32</sup>P-labelled primer. NMNTP<sup>+</sup> concentration: 1 μM (lane 2), 10 μM (lane 3), 100 μM (lane 4), 1 mM (lane 5) and 10 mM (lane 6); lane 7, 10 mM NMNTP<sup>+</sup> with double the concentration of primer; lane 8, 1 μM GTP. The concentration of TdT was 0.8 U/μl.

weight material. Large polymers were obtained even when we used low concentrations (1.67 mM) of NMNDP-CN as substrate. A broad distribution of polymer sizes was also observed. The rate of polymerization in the presence of primer was even faster than with NMNDPH. Large polymers were present after 5 min and the reaction was essentially complete within 20 min (data not shown).

The separation between bands in the electrophoretograms (Fig. 5) corresponded to that obtained with NMNDP<sup>+</sup> rather than NMNDPH, indicating that cyanide dissociates from the polymer prior to the electrophoretic separation. This method, therefore, provides an efficient route to high molecular weight poly(NMN<sup>+</sup>).

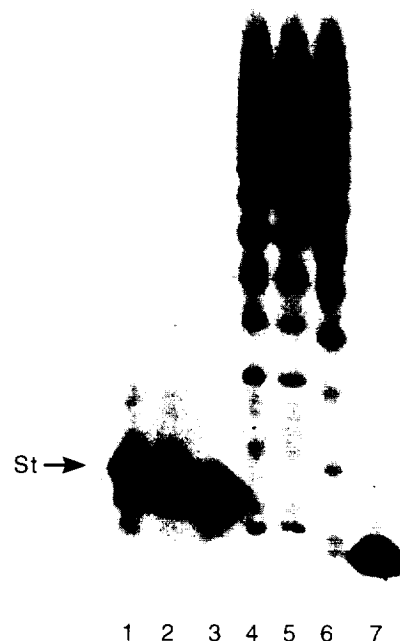
Gel electrophoresis and paper chromatography show that the hydration product of NMNDP<sup>+</sup> is not a substrate of polynucleotide phosphorylase either in the presence or the absence of primer (data not shown).

#### Addition of NMNTP<sup>+</sup> to oligodeoxynucleotides by terminal transferase

Terminal transferase catalyzes the addition of one or two NMN<sup>+</sup> residues to an oligodeoxynucleotide primer. The yield of the products is highly dependent on the concentrations of the substrate and primer. When the concentration of the NMNTP<sup>+</sup> is high enough (10 mM) and the ratio of primer to substrate is appropriate, almost all of the primer is elongated (Fig. 6). The spacing between bands is consistent with the incorporation of monomers carrying a positive charge. The yield of the di-adduct was always small. The purified product terminated by a NMN<sup>+</sup> residue can be used to prime the further incorporation of standard nucleotides, but the efficiency is lower than with a standard oligodeoxynucleotide primer (data not shown).

NMNTPH and NMNTP-CN are unstable at pH 7 and cannot, therefore, be used as substrates for terminal deoxynucleotidyl transferase.

primer	No NDP			NMNDP <sup>+</sup>			
	+	+	-	+	+	-	-
p(dA) <sub>4</sub> rA	+	+	-	+	+	-	-
p(dC) <sub>4</sub> rC	-	-	+	-	-	+	+
RNase A	-	+	+	-	+	-	+

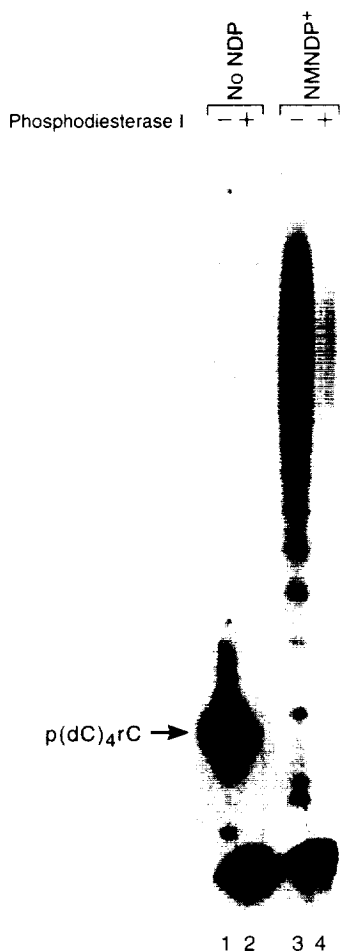


**Figure 7.** Ribonuclease A analysis of the polymeric products from the primed polymerization of NMNDP<sup>+</sup> by polynucleotide phosphorylase. The nature of the primer and the substrate for polymerization is indicated at the top of the Figure. St is the primer (dA)<sub>4</sub>rA.

#### Nuclease digestion of products

We attempted to hydrolyze the products of NMN<sup>+</sup> addition to different primers with ribonuclease T1, ribonuclease A and phosphodiesterase I. Ribonuclease T1 is an enzyme that cleaves 3'-5'-phosphodiester linkages adjacent to G residues, generating products terminated with 3'-phosphate and 5'-hydroxyl groups. The products obtained by extending a (dA)<sub>4</sub>rA or (dC)<sub>4</sub>rC primer with NMN<sup>+</sup> residues could not be degraded with ribonuclease T1 (data not shown), showing that NMN<sup>+</sup> does not substitute for a purine nucleotide in the ribonuclease T1 catalyzed reaction.

Ribonuclease A cleaves RNA adjacent to a pyrimidine residue generating 3'-phosphate- and 5'-hydroxyl-terminated products. The products of addition of NMN<sup>+</sup> to a (dA)<sub>4</sub>rA primer are not cleaved by ribonuclease A (Fig. 7, lane 5) but the products of addition to a (dC)<sub>4</sub>rC primer are completely degraded (Fig. 7, lane 7). Thus NMN<sup>+</sup> is accepted by RNase A as a nucleotide analog when it is downstream from a pyrimidine. However, it is not recognized as a pyrimidine analog when it is upstream of a second nicotinamide residue.



**Figure 8.** Phosphodiesterase I analysis of the polymeric products from the primed polymerization of NMNDP<sup>+</sup> by polynucleotide phosphorylase.

Phosphodiesterase I is an exonuclease that degrades RNA from the 3'-terminus. Products in which primers are extended with NMN<sup>+</sup> residues, are completely degraded by the enzyme (Fig. 8). Thus internucleotide bonds between pairs of NMN<sup>+</sup> residues are cleaved by phosphodiesterase I.

#### ***In vitro* transcription reactions and non-enzymatic template-directed reactions**

We attempted to replace each of the nucleoside triphosphates by NMNTP<sup>+</sup> or NMNTPH in a variety of transcription reactions using T3 or T7 RNA polymerase. We found no evidence for incorporation of the analogs into products. An extensive series of reactions using a variety of hairpin templates (18) gave no indication that the 2-methylimidazolide derivatives of NMN<sup>+</sup> in the oxidized or reduced form undergo non-enzymatic template-directed reactions.

#### **DISCUSSION**

NMNDP<sup>+</sup> and NMNTP<sup>+</sup> are readily prepared from NMN<sup>+</sup> by standard procedures (12). Since polynucleotide phosphorylase catalyzes a reversible polymerization of nucleoside 5'-diphos-

phates to give polynucleotides and inorganic phosphate, it is important to obtain the nucleoside 5'-diphosphate substrates of polynucleotide phosphorylase free of inorganic phosphate. The separation of NMNDP<sup>+</sup> from inorganic phosphate on a preparative scale is difficult. Elution from charcoal is inefficient. We achieved an excellent recovery of NMNDP<sup>+</sup>, free of inorganic phosphate, by using a long DEAE-Sephadex column and eluting with triethylammonium bicarbonate at a low flow rate.

The patterns of products obtained from the primer-initiated oligomerization of NMNDP<sup>+</sup> with polynucleotide phosphorylase (PNPase) are often complicated and the gels are often badly smeared (Fig. 2). We believe that this is due to the relative instability of the nucleotides and their oligomers under neutral or slightly alkaline conditions. NMNDP<sup>+</sup> is fairly stable, but some decomposition by loss of nicotinamide from the nucleotide is expected at pH 9 (23). The modification of one or more residues in an oligomer of NMN<sup>+</sup> would give a large number of isomeric products each with a characteristic mobility. This we believe is the origin of multiple bands and of some of the smearing. Loss of nicotinamide during the electrophoretic separation must account for further smearing. Consistent with the latter suggestion, a better-defined gel was obtained at pH 7.6 in the cold room (data not shown).

To confirm this suggestion we isolated from a gel the mono-adduct formed by incubating (dA)<sub>6</sub> with NMNTP<sup>+</sup> in the presence of terminal transferase. On re-electrophoresis a new band appeared that migrates close to (dA)<sub>6</sub> between the positions of (dA)<sub>6</sub> and (dA)<sub>7</sub>. The position on the gel is consistent with a ribo-adduct of (dA)<sub>6</sub>, formed by the loss of nicotinamide from the NMN<sup>+</sup> mono-adduct. Furthermore, a smear extends from the mono-adduct band to the new product band consistent with decomposition on the gel (data not shown).

The cyanide adduct of NMNDP<sup>+</sup>, a close analog of NMNDPH, is polymerized by PNPase to give high molecular weight products. The direct synthesis of high molecular weight poly(NMN<sup>+</sup>) from NMNDP<sup>+</sup> cannot be achieved with PNPase. However, since NMNDP-CN undergoes polymerization readily and cyanide addition is reversible, high molecular weight poly(NMN<sup>+</sup>) is readily obtained by incubating poly(NMNDP-CN) in the absence of cyanide.

NMNDPH and NMNDP-CN are better substrates than NMNDP<sup>+</sup> for PNPase. This was unexpected. The bond from N<sub>1</sub> of the nicotinamide ring to C'<sub>1</sub> of the ribose in NMNH or NMN-CN would deviate significantly from the plane of the nicotinamide ring (26). Thus NMNH or NMN-CN should have a different geometry from that of standard nucleotides. We anticipated, therefore, that the enzyme would discriminate against the reduced form. Clearly the enzyme is in fact permissive. The relatively ineffective polymerization of NMNDP<sup>+</sup> cannot be attributed entirely to the positive charge on the ring, since the 5'-diphosphate of 7-methyl guanosine is polymerized efficiently by the enzyme (data not shown).

T4 polynucleotide kinase is known to add phosphate to oligonucleotides even if the 5'-terminal base is heavily modified (27). It is, therefore, not surprising that the enzyme will accept poly(NMNH) as a substrate. The direct labelling of the products from the *de novo* polymerization of NMNDPH by polynucleotide kinase, without the need for preliminary dephosphorylation, must be the result of the well-known phosphate-transfer reaction catalyzed by the enzyme. Since this is a reaction of oligonucleotides terminated by a 5'-monophosphate, we may conclude that the products of the *de novo* polymerization of NMNDPH are 5'-monophosphates rather than 5'-diphosphates (28-30).



The introduction of NMN<sup>+</sup> residues into RNA is of considerable interest in the context of *in vitro* RNA evolution. It should be possible to select ribozymes with redox function from random pools of oligodeoxynucleotides terminated by an NMN<sup>+</sup> residue. The use of terminal transferase to introduce a single NMN<sup>+</sup> residue at the end of a presynthesized oligodeoxynucleotide should provide the substrates for this kind of selection.

NMN<sup>+</sup> can, in principle, form hydrogen bonds similar to those formed by adenylic acid or guanylic acid (Fig. 1). However, we were unable to incorporate NMNTP<sup>+</sup> into RNA on DNA templates using T3 or T7 RNA polymerase under standard conditions, or in the presence of lowered concentrations of the standard nucleoside triphosphates. Attempts to incorporate NMN<sup>+</sup> into oligonucleotides in non-enzymatic template-directed reactions were also unsuccessful. These results suggest that NMN<sup>+</sup>, if it is a molecular fossil of the RNA world, was not an informational unit of a polymer. At first it may have functioned via non-covalent attachment to RNA. Alternatively it could have been attached covalently to RNA by a ribozyme.

## ACKNOWLEDGEMENTS

This work was supported by NSCORT/EXOBIOLGY Grant No. NAGW-2881 and grant No. NAGW-1600 from the National Aeronautics and Space Administration. We thank Dr Heinz-Peter Muth for communicating preliminary data on the synthesis of NMNDP<sup>+</sup> and NMNTP<sup>+</sup>, Aubrey R. Hill, Jr for technical assistance and Sylvia H. Bailey for manuscript preparation.

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