Non-enzymatic template-directed synthesis of rna copolymers $^{\rm l}$

Gerald F. Joyce

The Salk Institute for Biological Studies, Post Office Box 85800, San Diego, California 92138, USA

Abstract. Cytosine-rich RNA copolymers facilitate the template-directed synthesis of complementary oligomers from mononucleotide 5'-phospho-2-methylimidazolides (Fig. 1). The efficiency of this reaction falls sharply as the ratio of cytosine to non-cytosine in the template is decreased. This is a severe limitation for self-replication because it means that any cytosine-rich polynucleotide that can serve as a good template will produce a cytosine-poor complementary strand that is unable to direct further rounds of synthesis. Studies with low-ratio random copolymer templates have shown that the efficiency can be increased by adjusting initial monomer concentrations and by providing additional activated monomers during later stages of the reaction. The oligomeric reaction products can be studied in detail using high performance liquid chromatography. It is possible to separate oligonucleotides on the basis of chain length and base composition. Thus a wealth of information is available to characterize the distribution of products over the course of the reaction and under a variety of reaction conditions.

INTRODUCTION

At the present time it is not possible to construct a selfreplicating system based on polynucleotides and activated mononucleotides. Polyguanylate forms a stable self-structure that prevents guanine-cytosine base pairing, polyuridylate forms a triple-helix with adenosine derivatives, and polyadenylate will not bind uridine mononucleotides (1).

Inoue and Orgel (2) have shown that a random polynucleotide containing a substantial excess of cytosine will facilitate the template-directed synthesis of complementary oligomers from mononucleotide 5'-phospho-2-methylimidazolides (Fig. 1).

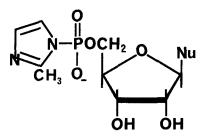


Figure 1. Structure of 2-MeImp Nu, where Nu is cytosine (C), uracil (U), adenine (A), or guanine (G)

Monomers are incorporated into newly-synthesized oligomers if and only if their complement is present in the template. However, the efficiency of this reaction falls sharply as the ratio of cytosine to non-cytosine in the template is decreased. This is a severe limitation for self-replication because it means that any cytosine-rich polynucleotide that can serve as a good template will produce a cytosine-poor complementary strand that is unable to direct further rounds of synthesis.

REACTION CONDITIONS

The problem can be illustrated by the reaction of $poly(C_3U_1)$ template with activated guanine and/or adenine mononucleotides in the presence of Mg²⁺. With guanine alone the reaction proceeds rapidly and efficiently, while with adenine alone oligomer formation does not exceed template-free controls. When both guanine and adenine are present, both are incorporated into newly-synthesized oligomers, though adenine incorporation is only about half as efficient as guanine. The highest yields are obtained with guanine alone, producing oligomers up to about the 30-mer. Since the probability of encountering 30 consecutive cytosine residues on the template is only about 10^{-4} , it seems that guanine must be forming either intrahelical (wobble) or extrahelical (loop-out) non-complementary pairs (3). Thus the relatively poor incorporation of adenine is a result of its own inefficiency as well as the overefficiency of guanine.

It has been possible to increase the efficiency of adenine incorporation both by increasing the concentration of adenine and by decreasing the concentration of guanine in the reaction mixture. Compared to the standard condition of 0.1 M 2-MeImpG and 0.1 M 2-MeImpA, doubling the concentration of 2-MeImpA to 0.2 M results in a 16% increase in the efficiency of adenine incorporation and a 6% increase in the efficiency of guanine incorporation. Presumably the greater concentration of 2-MeImpA allows it to compete more effectively at wobble sites (e.g. ³'CCU⁵'). and increases the likelihood that newly-synthesized oligomers will be able to continue beyond "difficult" sites (e.g. ³'CUU⁵'). Subsequently halving the concentration of 2-MeImpG to 0.05 M results in an additional 14% increase in the efficiency of adenine incorporation and a 6% decrease in the efficiency of guanine incorporation. The initiation of oligomer synthesis, which requires the formation of a 2-MeImpGpG dimer, is delayed under these conditions (4). Once oligomer synthesis is underway, 2-MeImpA enjoys a 4:1 concentration at wobble sites. while the amount of surviving 2-MeImpG is close to the threshold concentration necessary to sustain the reaction. If the concentration of 2-MeImpG is further reduced to 0.025 M, the efficiency of both adenine and guanine incorporation falls by over 50%.

Qualitatively similar but quantitatively less dramatic results are obtained with guanine & uracil on a poly(CA) template and with guanine & cytosine on a poly(CG) template. Unlike poly(CU), these templates do not contain wobble sites for guanine. However, cytosine and especially uracil have comparatively weak stacking interactions and thus can benefit from a concentration advantage.

HPLC ANALYSIS

The oligomeric products resulting from template-directed condensation of activated mononucleotides can be studied in detail by high performance liquid chromatography (HPLC). Using an RPC-5 column with a linear sodium perchlorate gradient (0 to 0.06 M over 90 minutes), it is possible to separate

G. F. JOYCE

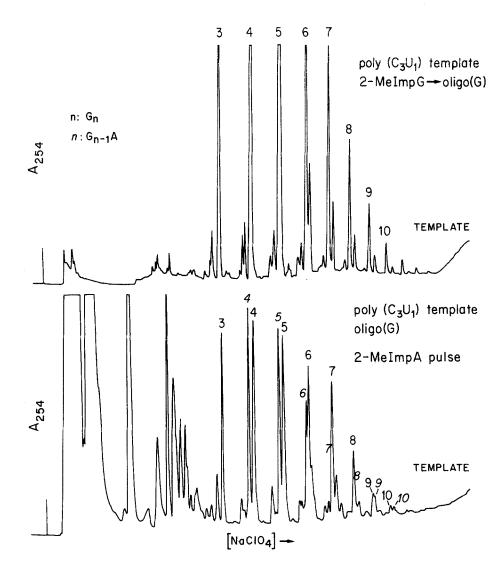
oligonucleotides on the basis of chain length (5). The elution profile is obtained by monitoring UV absorbance at 254 nm, thus providing a quantitative description of oligomer yields.

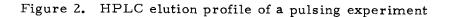
Recently, it has also become possible to determine the base composition of oligonucleotides containing guanine & adenine, guanine & uracil, and guanine & cytosine. A random copolymer template containing cytosine and one of the other three bases is used to direct synthesis of ³H-labeled oligonucleotides. The reaction is carried out in paired tubes, with $[8-{}^{3}H]$ 2-MeImpG added to the first tube and $[8-{}^{3}H]$ 2-MeImpX (the other complementary mononucleotide) added to the second. HPLC elution profiles are obtained for each tube by simultaneously monitoring UV absorbance and ³H activity. For each peak the ratio of UV absorbance to ³H-labeled G activity to ³H-labeled X activity is determined, and taking into account differences in nucleotide extinction coefficients. integer values for base composition are assigned. There is an obvious regularity in the separation of subsequent oligomers in a compositional series. In those cases where oligomers with different base composition have identical retention times (e.g. $(pG)_7$ and (pG)₆pA), this regularity is used to clarify the overlap in position.

A detailed quantitative description of the oligomeric product distribution is necessary for determining the sequence-specific efficiency of template-directed synthesis. Studies with random copolymer templates are seen as laying the groundwork for future work with known sequence templates. Given a cytosinerich oligomer template, it is now possible to determine the yield of its complement and of the various error copies simply by examining the HPLC profile. This provides a method for monitoring the time course of the reaction and for characterizing the product distribution under a variety of reaction conditions.

PULSING

"Pulsing" is a useful technique for studying the efficiency and regiospecificity of monomer addition to a newly-synthesized oligomer chain. The reaction of $poly(C_3U_1)$ template with 2-MeImpG produces a series of 3'-5' linked oligo(G)s, whose mean chain length is a function of time, temperature, pH, $[Mg^{2+}]$,





G. F. JOYCE

[template], and [2-MeImpG]. The reaction can be interrupted by dilution and subsequent acid hydrolysis of the guanosine imidazolides. Unreacted monomers and dimers are removed by gel filtration, and the solution is evaporated to dryness. The reaction can then be resumed by rehydration with radioactive 2-MeImpA, producing oligo(G)s terminated by labeled adenine residues.

The production of oligo(G_nA)s and oligo(G_nA_2)s is confirmed by examining the HPLC elution profile (Fig. 2). The site of 2-MeImpA addition to an oligo(G) primer can be determined by exhaustive alkaline hydrolysis (5). Following KOH degradation, radioactive adenine is distributed into A, Ap, and AppGp in a ratio of approximately 7:1:2, corresponding to terminal sites at the 3' end, internal sites at the 3' end, and terminal sites at the 5' end, respectively. The regiospecificity of 2-MeImpA addition at the 3' end can be determined by ribonuclease T₁ digestion (6). This enzyme cleaves only 3'-5' internucleotide linkages on the 3' side of a G residue. Following degradation with T₁, radioactive adenine is distributed into A and G² pA in a ratio of 3:1 and into ApA and G² pApA in a ratio of about 3:1. Thus the 3'-5' regiospecificity at both terminal and internal sites is about 75%.

REKINDLING

CCGCC was the first known-sequence RNA template to be used to demonstrate non-enzymatic template-directed synthesis of complementary oligomers (7). The reaction of CCGCC with 2-MeImpG and 2-MeImpC produces a host of oligo(GC)s, with GGCG and GGCGG dominating the distribution of tetramers and pentamers, respectively (Fig. 3). The identity of GGCGG was confirmed by HPLC analysis and by its hydrolysis with pancreatic ribonuclease to GGC and GG (7).

When the reaction is allowed to reach completion (after about 3 weeks), the yield of GGCG far exceeds that of GGCGG. It might seem that if the newly-synthesized oligomers could reach tetramer length, then it would be comparatively easy for the final guanine residue to be added. However, by the time an appreciable amount of GGCG has been formed, the supply of remaining 2-MeImpG has been greatly depleted by ongoing hydrolysis. At this point the reaction can be "rekindled" by removing the hydrolyzed monomers (using gel filtration) and

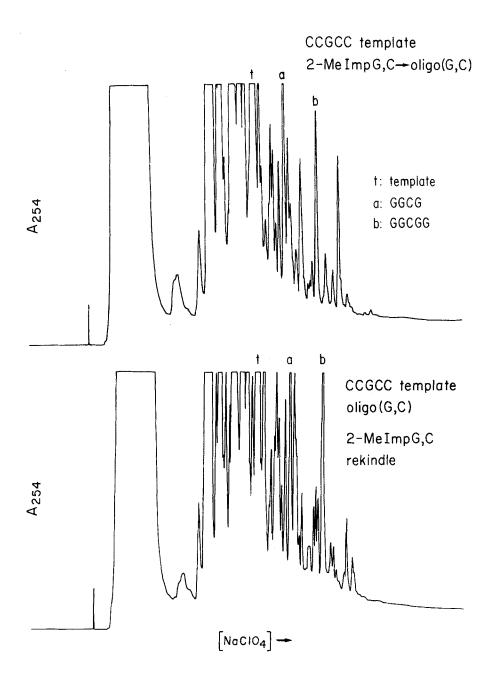


Figure 3. HPLC elution profile of a recycling experiment

replacing them with activated compounds. The result is a rapid and dramatic increase in the yield of GGCGG.

A process equivalent to rekindling can be envisioned as operating continuously in the context of a flow reactor. The reaction vessel would contain a population of GC-rich oligonucleotides and a supply of activated mononucleotides. Its semipermeable walls would allow continuous removal of unreacted monomers, while fresh monomers are added to maintain the supply of activated materials. At the present time, a major obstacle to the realization of such a system is the requirement for a method to allow periodic disruption of the association between template and complement. This method must permit each strand to begin a new round of synthesis before the strand is made inaccessible by intra- or intermolecular duplex formation.

1 This paper describes a broad preliminary survey of templatedirected synthesis. More detailed studies using RNA copolymer templates are in progress. The author wishes to thank Leslie Orgel and Tan Inoue for making this work possible, Aubrey Hill for technical assistance, and Rosemary Brown for manuscript preparation.

REFERENCES

- 1. Ts'o, P. 1974, "Basic Principles in Nucleic Acid Chemistry" (Academic Press, New York) Vol. 1, Chap. 6.
- 2. Inoue, T. and Orgel, L. E. 1983, Science 219, pp. 859-862.
- Lomant, A. J. and Fresco, J. R. 1975, Progr. Nucleic Acid Res. Mol. Biol. 15, pp. 185-218.
- 4. Fakhrai, H., Inoue, T. and Orgel, L. E., unpublished results.
- Lohrmann, R. and Orgel, L. E. 1980, J. Mol. Biol. 142, pp. 555-567.
- 6. Sulston, J. et al 1968, PNAS 60, pp. 409-415.
- 7. Brown, J. et al, unpublished results.