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# SYNTHESIS, BIOPHYSICAL PROPERTIES, AND STABILITY STUDIES OF MIXED BACKBONE OLIGONUCLEOTIDES CONTAINING NOVEL NON-IONIC LINKAGES

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ABSTRACT: Mixed backbone oligonucleotides (MBOs) (containing ionic and non-ionic internucleotidic linkages) in which the non-ionic segments are either methylphosphotriester (PO-OMe) or primary phosphoramidate (PO-NH<sub>2</sub>) linkages have been prepared using the recently described N-pent-4-enoyl (PNT) nucleoside phosphoramidates and H-phosphonates. Biophysical properties and stability studies suggest that these MBOs are novel antisense molecules.

During the last decade, there has been considerable progress in the development of phosphorothioate oligonucleotides as potential therapeutic and diagnostic agents, la-c and as unique tools in understanding cellular processes at the molecular level. oligonucleotide-based drug design is characterized by its universal appeal and simplicity in that it employs well-understood principles governing molecular recognition. In this context, second generation analogs exemplifying the following properties are being sought: (a) higher selectivity; (b) enhanced resistance to cellular nucleases; (c) improved cellular uptake; (d) amplification of the antisense effect by induction of RNase H; (e) unique pharmaceutical properties which enable site- and tissue-specific delivery; (f) lesser polyanion-related side effects; and (g) oral bioavailability with favorable pharmacokinetic profile. Towards have synthesized **MBOs** this goal, we which contain methylphosphotriester (PO-OMe, PS-OMe)<sup>2</sup> and phosphoramidate (PO-NH<sub>2</sub>)<sup>3a-c</sup> as the non-ionic linkages in conjunction with phosphoric diester (PO) and phosphorothioate (PS) as the ionic counterparts. Reported herein is the biophysical properties and stability studies of these MBOs.

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#### MBOs with PO-OMe and PS-OMe linkages

Our synthetic strategy involves the use of phosphoramidite chemistry on controlled-pore glass support (CPG) (Scheme 1). Briefly, the PSOMe and POOMe linkages were constructed using PNT nucleoside methoxy phosphoramidite (MEPNT) synthons and the PO or PS segment using PNT nucleoside  $\beta$ -cyanoethyl phosphoramidite (CEPNT) synthons respectively. Following the synthesis, the removal of the PNT groups, the  $\beta$ -cyanoethyl phosphate protecting groups, and cleavage of the oligonucleotide from the support were accomplished in one step by exposure to  $K_2CO_3/MeOH$  (0.05 M, rt, 30 to 36 h). The oligonucleotides were isolated by preparative polyacrylamide gel electrophoresis (PAGE) (7 M urea, 15-25 °C) using an assembly equipped for water circulation.

#### Fidelity of Oligonucleotide Synthesis

In order to ascertain the structural integrity of the PO-OMe group in the MBO, representative MBOs (having PO and POOMe linkages) were prepared and subjected to enzymatic digestion with *snake venom phosphodiesterase* and *bacterial alkaline phosphatase*. These studies revealed that the POOMe linkage was present to the extent of 97% and the presence of diester resulting from demethylation during purification and isolation were less than 3%. In the case of the MBOs having PS-OMe linkages, the expected integral ratio of the PSOMe/PS segment was observed in each case. <sup>44,b</sup>

#### Biophysical properties

The thermal denaturation temperatures ( $T_{\rm m}$ s) of complexes formed between MBOs (25-mers), and complementary DNA and RNA were determined as before. Representative data is revealed in Table 1. It is clear that in the case of the MBO analogs 1-7, the incorporation of the triester segment into GEM 91° (8) only marginally affects the  $T_{\rm m}$  values.

#### MBOs with phosphoramidate linkages

The P[O]NH<sub>2</sub> linkage is a novel non-ionic linkage as it is: (a) isosteric with phosphoric diester; and (b) is expected to make the MBO water soluble because of its potential to hydrogen-bond with water. As before, phosphoramidite chemistry was used to build the PO/PS linkage, and H-phosphonate chemistry for the PO-NH<sub>2</sub> linkage.

#### Fidelity of oligonucleotide synthesis

To ascertain that complete base and phosphate deprotection had occurred without any attendant base-modifications, enzymatic digestion of representative MBOs (having PO and PO-NH $_2$  linkages) was carried (*vide supra*). Our studies reveal that complete deprotection had been achieved while preserving the integrity of the PO-NH $_2$  linkage. <sup>6a,b</sup>

#### Biophysical properties

The thermal denaturation temperatures ( $T_{\rm m}$ s) of complexes formed between MBOs (25-mers), and complementary DNA and RNA were determined. Representative data is revealed

SCHEME 1. Synthesis of MBOs; key: (a) phosphoramidite cycle using tert-butyl hydroperoxide as the oxidizing agent and 3H-1,2-benzodithiole-3-one-1,1-dioxide<sup>6</sup> as the sulfurizing reagent; (b)  $K_2CO_3$  (0.05 M in MeOH).

TABLE 1. Comparative T<sub>m</sub> Data of GEM 91<sup>®</sup> Analogs 1-7.

| Seq.# | Sequence   | T <sub>m</sub> (°C)¶ | T <sub>mt</sub> (°C) <sup>§</sup> |
|-------|--|----------------------|-----------------------------------|
| 1     | 5' C•T•C TCG CAC CCA TCT CTC TCC TTC T<br>PSOMe PS   | 52.5                 | 62.5                              |
| 2     | 5' C-T-C-T CGC AC CCA TCT CTC TCC TTC T<br>PSOMe PS  | 51.5                 | 61.8                              |
| 3     | 5' C•T•C•T•C GCAC CCA TCT CTC TCC TTC T<br>PSOMe PS  | 50.7                 | 60.6                              |
| 4     | 5' CTC TCG CAC CCA TCT CTC TCCT T•C•T PS PS PSOMe    | 51.0                 | 62.0                              |
| 5     | 5' CTC TCG CAC CCA TCT CTC TCC T-T-C-T PS PSOMe      | 50.3                 | 61.1                              |
| 6     | 5' C•T•C TCG CAC CCA TCT CTC TCC T PSOMe PS PS PSOMe | 49.1                 | 60.5                              |
| 7     | 5' CTC TCG CAC CCA T•C•T•C TCT CCT TCT PSOMe PS      | 50.8                 | 60.3                              |
| 8     | 5' CTC TCG CAC CCA TCT CTC TCC TTC T<br>PS           | 54.0                 | 64.2                              |

<sup>¶</sup> Against complementary DNA (30-mer) (PO);  $^{\S}$  Against complementary RNA (25-mer) (PO); All  $T_{\rm ms}$  represent the average of at least two determinations.

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| TABLE 2. Comparative $T_r$ | Data of MBO Analogs 9 | - 12 of | GEM 91 <sup>®</sup> | (8) |
|----------------------------|-----------------------|---------|---------------------|-----|
|----------------------------|-----------------------|---------|---------------------|-----|

| Seq. # | Sequence  | $T_{\mathbf{m}}^{\mathbf{q}}$ | T <sub>m</sub> § |
|--------|---|-------------------------------|------------------|
| 2      | 5' C • T • C • T • C • GCA CCC ATC TCT CTC CTT CT | 61.3                          | -                |
|        | PO-NH <sub>2</sub> PO                             |                               |                  |
| 10     | 5' C • T • C • T • CGC ACC CAT CTC TCT CCT TCT    | 53.1                          | 63.3             |
|        | PO-NH <sub>2</sub> PS                             |                               |                  |
| 11     | 5' C • T • C TCG CAC CCA TCT CTC TCC TT • C • T   | 55.3                          | 63.8             |
|        | PO-NH <sub>2</sub> PS PO-NH <sub>2</sub>          |                               |                  |
| 12     | 5' CTC TCG CAC CC • A • T • C • T CTC TCC TTC T   | 52.0                          | 59.8             |
|        | PS PO-NH <sub>2</sub> PS                          |                               |                  |
| 8      | 5' CTC TCG CAC CCA TCT CTC TCC TTC T              | 55.0                          | 65.1             |
|        | PS  |                               |                  |
| 13     | 5' <u>CTC TCG CAC CCA TCT CTC TCC TTC T</u><br>PO | 64.8                          | -                |

<sup>¶</sup> Against complementary DNA (30-mer) (PO); § Against complementary RNA (25-mer) (PO). All  $T_{mS}$  represent the average of at least two determinations forward and backward.

in Table 2. Our data shows that in the case of GEM  $9.1^{\circ}$  analogs 9-1.2, the incorporation of the phosphoramidate segment into GEM  $91^{\circ}$  (8) only marginally affects the  $T_{\rm m}$  values.

### Enzymatic stability of MBOs

Preliminary studies revealed that incorporation of non-ionic linkages (PSOMe, POOMe, or PONH<sub>2</sub>) increases the stability of these MBOs against degradation by nucleases depending on the site of incorporation and the length of the non-ionic segment. 4b.6b

Thus the MBOs with methylphosphotriester and phosphoramidate linkages represent novel antisense molecules. The evaluation of biological activity, and pharmacokinetics of the MBOs is in progress, and will be reported in due course.

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