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Synthesis and Properties of Oligonucleotide Chimeras Containing 5'-Amino-2'-deoxy-2'-fluoroarabinonucleosides

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Synthesis and Properties of Oligonucleotide Chimeras Containing 5'-Amino-2'-deoxy-2'-fluoroarabinonucleosides

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ABSTRACT

Oligonucleotide analogues comprised of 2'-deoxy-2'-fluoro- β -D-arabinose units joined via P3'-N5' phosphoramidate linkages (2'F-ANA^{5'N}) were prepared for the first time. Among the compounds prepared were a series of 2'OMe-RNA-[GAP]-2'OMe-RNA 'chimeras', whereby the "GAP" consisted of DNA, DNA^{5'N}, 2'F-ANA or 2'F-ANA^{5'N} segments. The chimeras with the 2'F-ANA and DNA gaps exhibited the highest affinity towards a complementary RNA target, followed by the 5'-amino derivatives, i.e., 2'F-ANA > DNA > 2'F-ANA^{5'N} > DNA^{5'N}. Importantly, hybrids between these chimeras and target RNA were all substrates of both human RNase HII and *E.coli* RNase HI. In terms of efficiency of the chimera in recruiting the bacterial enzyme, the following order was observed: gap DNA > 2'F-ANA > 2'F-ANA^{5'N} > DNA^{5'N}. The corresponding relative rates observed with the human enzyme were: gap DNA > 2'F-ANA^{5'N} > 2'F-ANA > DNA^{5'N}.

Key Words: 5'-Amino-2'-deoxy-2'-fluoroarabinonucleosides; Thermal stability; RNase H.

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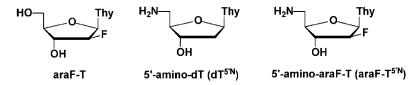


Figure 1. Structure of the nucleosides synthesized for this work.

Antisense oligonucleotides (AONs) suitable for use as therapeutics must bind tightly to their complementary RNA targets, elicit degradation of the RNA by the cellular enzyme RNase H, and have the general physiochemical characteristics of a drug-like molecule (solubility, stable to degradation, etc). A number of modified AONs have been developed, but very few of them are able to activate RNase H. We have previously shown that 2'-deoxy-2'-fluoro-β-D-arabinonucleic acids (FANA) can form stable complexes with complementary RNA and activate RNase H. [1,2] The present report describes the antisense characteristics of oligonucleotides constructed from 5'-amino-2'-deoxy-2'-fluoroarabinonucleoside units (2'F-ANA 5'N). Since substitution of the 5'-OH with a 5'-NH₂-group has very little influence on the conformation of the sugar ring, [3,4] we speculated that this modification would confer better nuclease resistance without compromising the ability of the AON to bind to RNA or elicit RNase H activity.

To test this hypothesis, the key starting material, 5'-amino-2'-deoxy-2'-fluoro-arabinothymidine (araF- $T^{5'N}$, Fig. 1) was prepared starting from araF-T. Thus, araF-T was converted to its 5'-tetrachlorophthalimide derivative, which upon treatment with ethylenediamine afforded araF- $T^{5'N}$ in good yields. [5,6] $dT^{5'N}$ was prepared by the same procedure, [5] and together with araF- $T^{5'N}$, were converted to their 5'-O-monomethoxytrityl-3'-O-cyanoethylphosphoramidite derivatives. To assess the effect of $5'O \rightarrow 5'N$ substitutions on the stability of dT/rA hybrids, oligodeoxy-thymidylates containing a single araF- $T^{5'N}$ or $dT^{5'N}$ unit were prepared. T_m data obtained demonstrated a significant destabilization effect ($\Delta T_m = -2$ to $-3^{\circ}C/$ substitution). Next, the affinity of oligonucleotide chimeras was assessed (Table 1).

Table 1. Melting temperature (T_m) of hybrids formed between RNA and oligonucleotide chimeras (gapmers).

AON #	AON composition	T _m (°C)
1	$U_{\text{me}}U$	40
2	$U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}$	27
3	$U_{me}U_{m$	44
4	$U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}u_{me}ara(T_f{}^nT_f{}^nT_f{}^nT_f{}^nT_f{}^nT_f{}^n)U_{me}U_{me}U_{me}U_{me}U_{me}U_{me}$	32

 U_{me} corresponds to 2'-O-methyluridine; dT, dT^n , $araT_f$, and $araT_f^n$ correspond to 2'-deoxy-thymidine, $dT^{5'N}$, araF-T, and $araF-T^{5'N}$ nucleosides, respectively (see Fig. 1). All melting experiments were done with rA_{18} as target, in a buffer: 140 mM KCl, 5 mM, Na_2HPO_4 , 1 mM MgCl₂, pH 7.2.

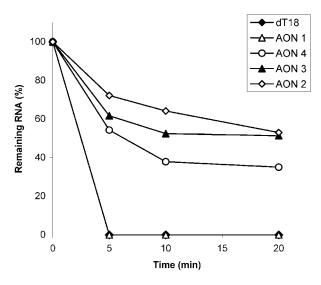


Figure 2. RNase H-mediated degradation of RNA (rA₁₈). The hybrids AON/RNA were exposed to human RNase HII as previously described.^[7] The base sequence of the AONs is given in Table 1.

These AONs contained two 2'OMe-rU₆ 'wings' flanking a hexathymidylate 'gap', namely dT_6 , $(dT^{5'N})_6$, araF-T₆ and araF- $(T^{5'N})_6$ (Table 1). Again, T_m data demonstrated a destabilizing effect upon substituting a 5'O with a 5'NH group $(\Delta T_m - 2^{\circ}C)$.

Figure 2 shows that all AON chimeras induced target RNA cleavage by human RNase HII. The ability of the various AONs to elicit RNase HII activity followed the order: DNA > araF-T^{5'N} > araF-T > dT^{5'N} demonstrating that while 5'O \rightarrow 5'N substitutions in the DNA strand is detrimental to RNase HII activity, this was not the case for 2'F-ANA strands for which a small increment in enzymatic activity was observed.

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