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Analysis of Synthetic Oligonucleotides using Quaternary Amine Functionalized, Non-porous Particles

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In addition to chromatography, Brian also has a passion for ice cream-making, and enjoys experimenting with bold, new flavors.



Introduction

Synthetic oligonucleotides (OGNs) are becoming increasingly popular as novel agents for the treatment of disease. The characterization of OGNs is important in the drug development process, and one common technique is strong anion exchange liquid chromatography. This high resolution technique for analysis for OGNs is preferred for many scientists when extensive characterization (i.e. LC-MS) isn't necessary. Another valuable benefit is that ion pair is not necessary in this technique and that n-1 failure sequences can still be separated. Although analysis of so-called "vanilla" or unmodified OGNs can be relatively straightforward, other modified OGNs present challenges. Several different modifications to the nucleic acid can be made to prevent nuclease degradation; these include backbone (e.g. phosphorothioate), ribose/deoxy-ribose sugar (e.g. bridged nucleic acid), and 2'-O substitutions. Additionally, different conjugations can be performed to either the 5' or 3' end of the oligo. Here we present several examples of different oligonucleotides, and how the subsequent mobile phase composition and gradient program may be optimized.

Materials and Methods

Crude, desalted OGNs (poly dT, Amino C12, BNA, and Phosphorothioate) were purchased from Integrated DNA Technologies (Coralville, IA, USA). Sequences, along with modifications, are indicated in **Table 1**, below. Samples were prepared at 30 µg/mL in nuclease free water. Injection volume was 2 µL. Note that mass loads for a non-porous particle will be reduced when compared to other analytical methods. A good starting point for OGNs is 1 µg for a 250 x 4.6 mm ID column.

Table 1. OGNs Sequences and Modifications

Oligo Type	Length	Sequence
Poly (dT)	12-18mer	N/A
Poly (dT)	19-24mer	N/A
Poly (dT)	19/20mer, 39/40mer, 59/60mer	N/A
Bridged Nucleic Acid	19mer	5'-TA/BNA-A/TA/BNA-meC/GT/BNA-T/TA/BNA-T/AC/BNA-G/CC/BNA-C/A-3'
Amino C12	24mer	5'AmMC125-TCGTGCTTTTGTGTTTCGCGTT-3
DNA Phosphorothioate	19mer	5'-ACT*G*A*C*T*G*A*C*T*G*A*C*G*T*A*C*T-3

Experimental Conditions

Each LC method was performed using a Clarity[®] 5 µm Oligo-SAX column on an Agilent[®] 1200 (Agilent Technologies, Santa Clara, CA USA) with an upper pressure limit of 600 bar.

Two different mobile phase systems were implemented. Anion exchange relies on the electrostatic interaction between analyte and oppositely charged stationary phase. As such, chloride anions can be used for some methods while strongly adsorbing OGNs may require perchlorate, which has a higher ionic elution strength. Gradients were also optimized depending on the particular OGN. Gradient programs are summarized in **Table 2**.

Column: Clarity 5 µm Oligo-SAX
Dimensions: 250 x 4.6 mm
Part No.: 00G-4749-E0
Mobile Phase: See Table 2
Flow Rate: 1.6 mL/min
Temperature: 30°
LC System: Agilent[®] 1200
Detection: UV-Vis @ 260 nm

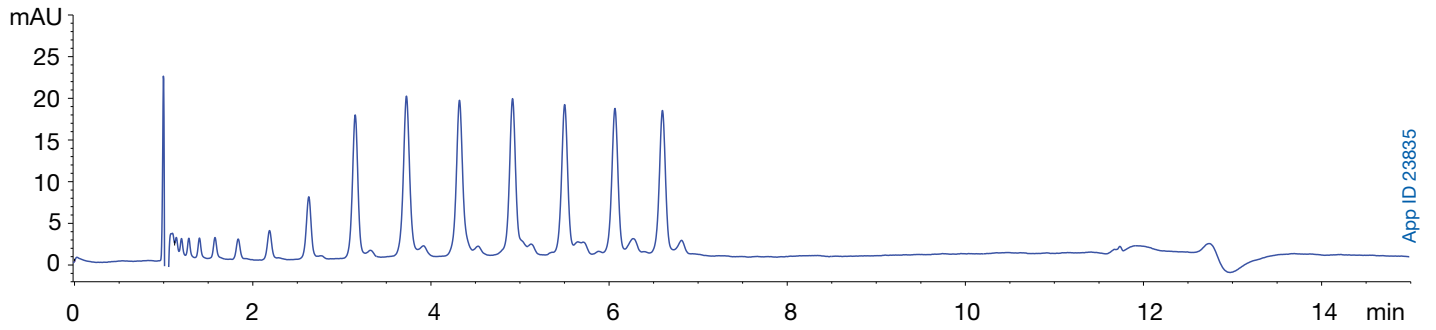
Table 2. Gradient Program

Oligo Type	Mobile Phase A	Mobile Phase B	Gradient Program
Poly (dT) 12-18mer	20 mM Tris, pH 8.0	20 mM Tris + 1.25 M NaCl, pH 8.0	32-46.6% in 10 min
Poly (dT) 19-24mer	20 mM Tris, pH 8.0	20 mM Tris + 1.25 M NaCl, pH 8.0	32-46.6% in 10 min
Poly (dT) 19/20mer, 39/40mer, 59/60mer	20 mM Tris, pH 8.0	20 mM Tris + 1.0 M NaOCl ₄ , pH 8.0	10-46.6% B in 10 min
DNA Phosphorothioate	20 mM Tris, pH 8.0	20 mM Tris + 1.0 M NaOCl ₄ , pH 8.0	5-46.6% in 10 min
Amino C12	20 mM Tris, pH 8.0	20 mM Tris + 1.25 M NaCl	32-46.6% in 10 min



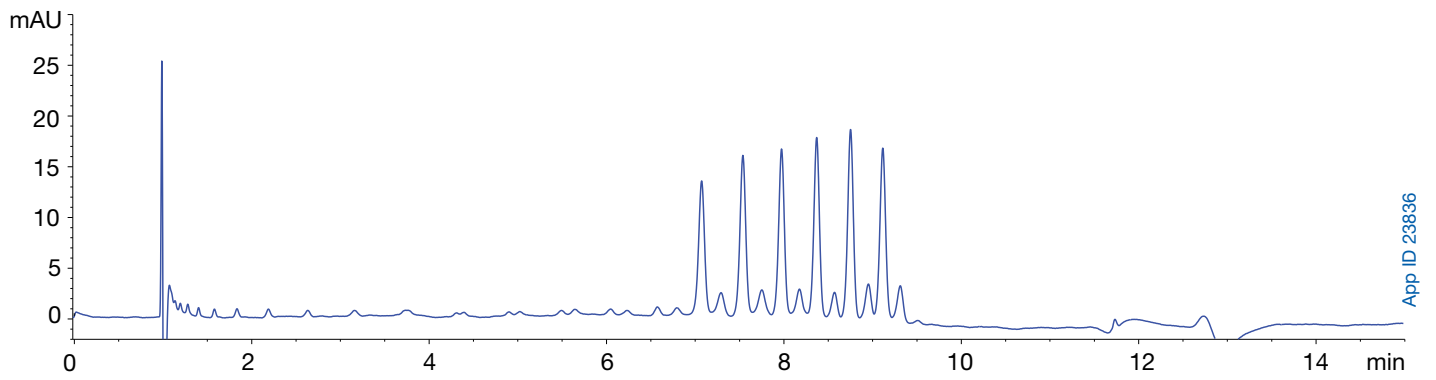
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Figure 1. Poly (dT) 12-18mer



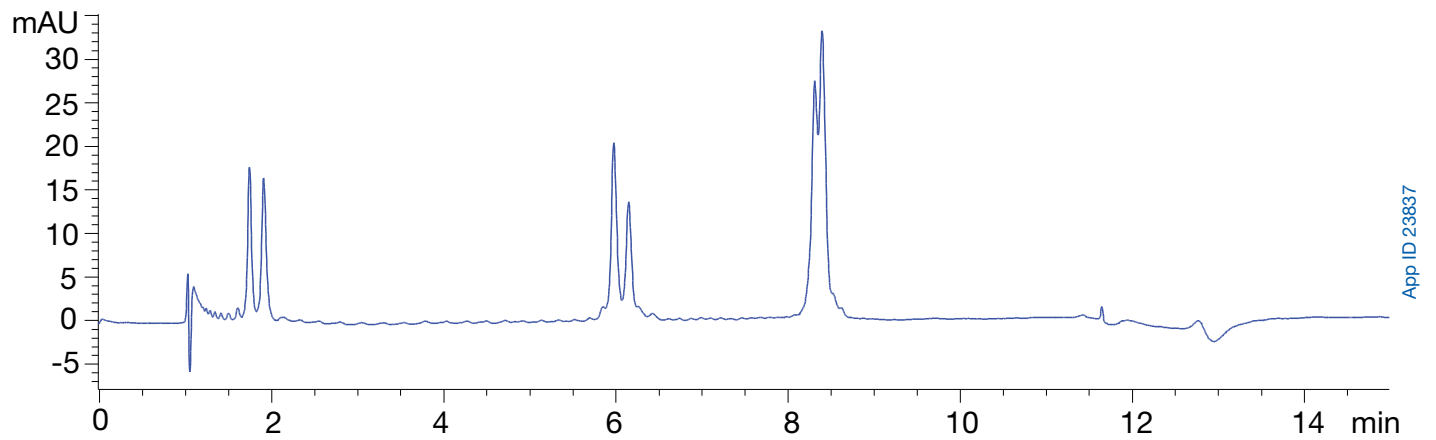
App ID 23835

Figure 2. Poly (dT) 19-24mer



App ID 23836

Figure 3. Poly (dT) 19/20mer, 39/40mer, 59/60mer



App ID 23837

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Figure 4. BNA 19mer

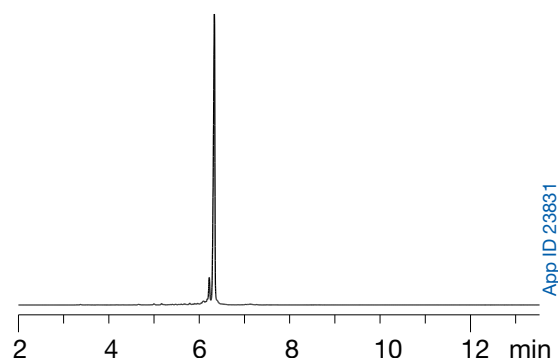


Figure 6. Amino C12 Linker DNA

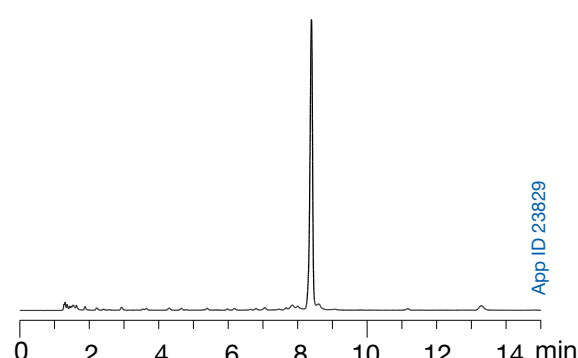
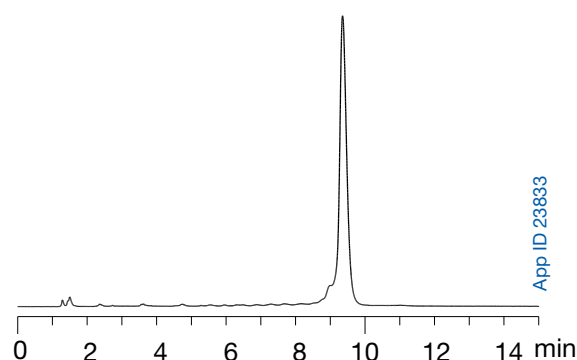


Figure 5. DNA 19mer Phosphorothioate



Results and Discussion

Figures 1 and 2 show the separation of poly dT ladders. This shows OGNs in range of 12-24mer, which are common OGN lengths in both molecular biology and therapeutic applications. Note the separation of synthesis impurities observed between 18-24mer ladder. This extent of characterization allows for separation beyond even n-1 failure sequences.

Longer OGNs typically present challenges in anion exchange. **Figure 3** shows a separation of up to 59 and 60mer. This might apply to aptamers, unique OGNs which bind to protein targets much like antibodies. Aptamers can be anywhere between 22-40mer in length (1), and the optimized chemistry of Clarity Oligo-SAX shows good proof of concept for OGNs exceeding the typically 19-24mer seen in siRNA and antisense applications.

The BNA sample observed in **Figure 4** shows good separation of n-1 failure sequence. BNA and other ribose modified OGNs present challenges since they are rigid in structure and have a higher binding affinity, thus stronger electrostatic effects(2).

Figure 5 shows reasonably good peak shape for an OGN with multiple phosphorothioate modifications. Phosphorothioates typically have higher electrostatic effects, which lend to their binding affinity and thus attractiveness for therapeutic applications (3). As such, again the perchlorate mobile phase is needed to elute the modified thioate OGN. One thing to note is that SAX column chemistries- specifically the hydrophobic coat and quaternary amine used- can have implications for column performance

for phosphorothioates. Additionally, phosphorothioates have stereochemistry which contributes to band broadening. That said, the peak shape here, although relatively broad, is more than acceptable for phosphorothioate-modified OGNs.

Finally, **Figure 6** shows good peak shape for an Amino-C12 linker conjugate. A more hydrophobic phase might show slight peak broadening.

Conclusion

With various modifications for OGNs, an optimal column chemistry like the Clarity Oligo-SAX with good ion exchange capacity and minimal secondary interactions is required. Using either standard chloride or perchlorate methods, a variety of different OGNs can be analyzed.

References

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2. Obika, Satoshi, Osamu Nakagawa, Akiko Hiroto, Yoshiyuki Hari, and Takeshi Imanishi. "Synthesis and Properties of a Novel Bridged Nucleic Acid with a P3' → N5' Phosphoramidate Linkage, 5'-amino-2',4'-BNA." *Chem. Commun.* 17 (2003): 2202-203. Print.
3. Mou, Tung-Chung, and Donald M. Gray. "The High Binding Affinity of Phosphorothioate-Modified Oligomers for Ff Gene 5 Protein Is Moderated by the Addition of C-5 Propyne or 2'-O-Methyl Modifications." *Nucleic Acids Research* 30.3 (2002): 749-758. Print.



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Ordering Information Clarity Oligo-SAX PEEK Columns

	Analytical	Analytical	Analytical	Guard
Phases	250 x 4.6	150 x 4.6	100 x 4.6	50 x 4.6
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