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Oligonucleotide synthesis under mild deprotection conditions

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Table of Contents

HPLC profiles of ODNs	2-20
MALDI MS and OD ₂₆₀ values of ODNs	21-32
Capillary electrophoresis profiles of ODNs	33-35
An example trityl assay log for ODN synthesis	36-36
Image of gel electrophoresis analysis of ODN 29j	37-37



































































 OD_{260} of ODN **29a** for the 0.52 µmol synthesis is 2.940.



 OD_{260} of ODN **29b** for the 0.52 µmol synthesis is 3.660.



 OD_{260} of ODN **29c** for the 0.52 µmol synthesis is 3.092.



 OD_{260} of ODN **29d** for the 0.52 µmol synthesis is 0.612.



 OD_{260} of ODN **29e** for the 0.52 µmol synthesis is 0.542.



 OD_{260} of ODN **29f** for the 0.52 µmol synthesis is 1.119.



 OD_{260} of ODN **29g** for the 0.52 µmol synthesis is 1.144.



 OD_{260} of ODN **29h** for the 0.52 µmol synthesis is 3.243.



 OD_{260} of ODN **29i** for the 0.52 µmol synthesis is 1.461.



 OD_{260} of ODN **29j** for the 0.52 µmol synthesis is 0.536.



 OD_{260} of ODN **29k** for the 0.52 µmol synthesis is 2.717.













An example trityl assay log of ODN synthesis

RUN NAME: KC 4107 (22 mer, Sensitive MD CE RUN DATE: 20220310 21:36:27 POSITION: COLUMN 1

Image of denatured polyacrylamide gel electrophoresis analysis of ODN **29j** in comparison with **29k**. The two ODNs have the same sequence but **29j** has a 4acC modification while **29k** does not. Polyacrylamide gel (15%, 7 M urea), 1X TBE buffer, 190 V, 45 min, stained with Gel Red.

Lane 1: 48 ng 29j. Lane 2: 48 ng 29j and 46.5 ng 29k. Lane 3: 46.5 ng 29k.

Lane 4: 24 ng 29j. Lane 5: 24 ng 29j and 23.3 ng 29k. Lane 6: 23.3 ng 29k.

Lane 7: 12 ng 29j. Lane 8: 12 ng 29j and 11.6 ng 29k. Lane 9: 11.6 ng 29k.

Lane 10: 7.2 ng 29j

Although the two ODNs could not be resolved, the single bands indicate that they do not contain any sequences that are shorter or longer than them. In addition, as presented in other parts of the paper, MALDI MS indicates that **29j** contains the 4acC modification while **29k** does not (Figure 4). Considering these and other evidences, which include CE and HPLC analyses, presented in the paper, **29j** is pure.