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Supporting Information

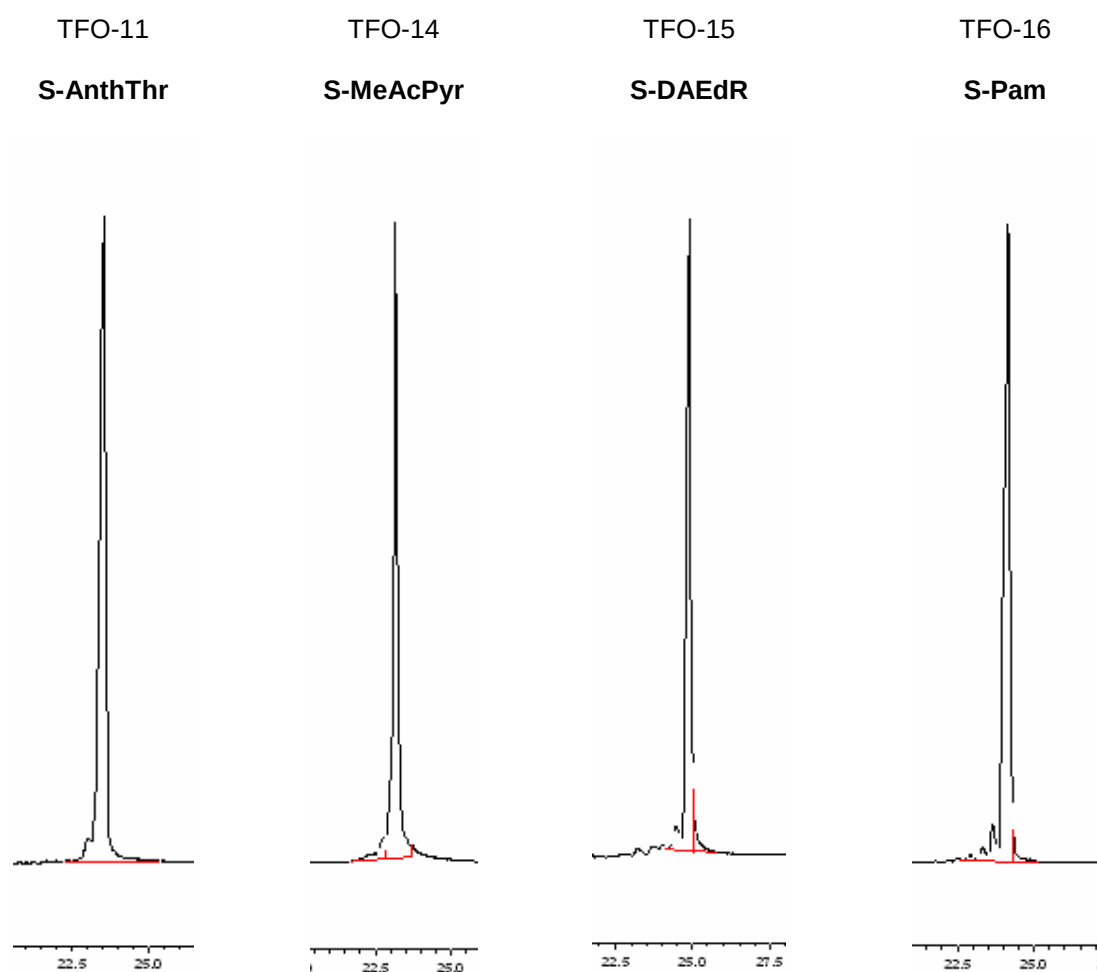
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Supporting Information

for

Potent Triple Helix Stabilization by 5',3'-Modified Triplex Forming Oligonucleotides

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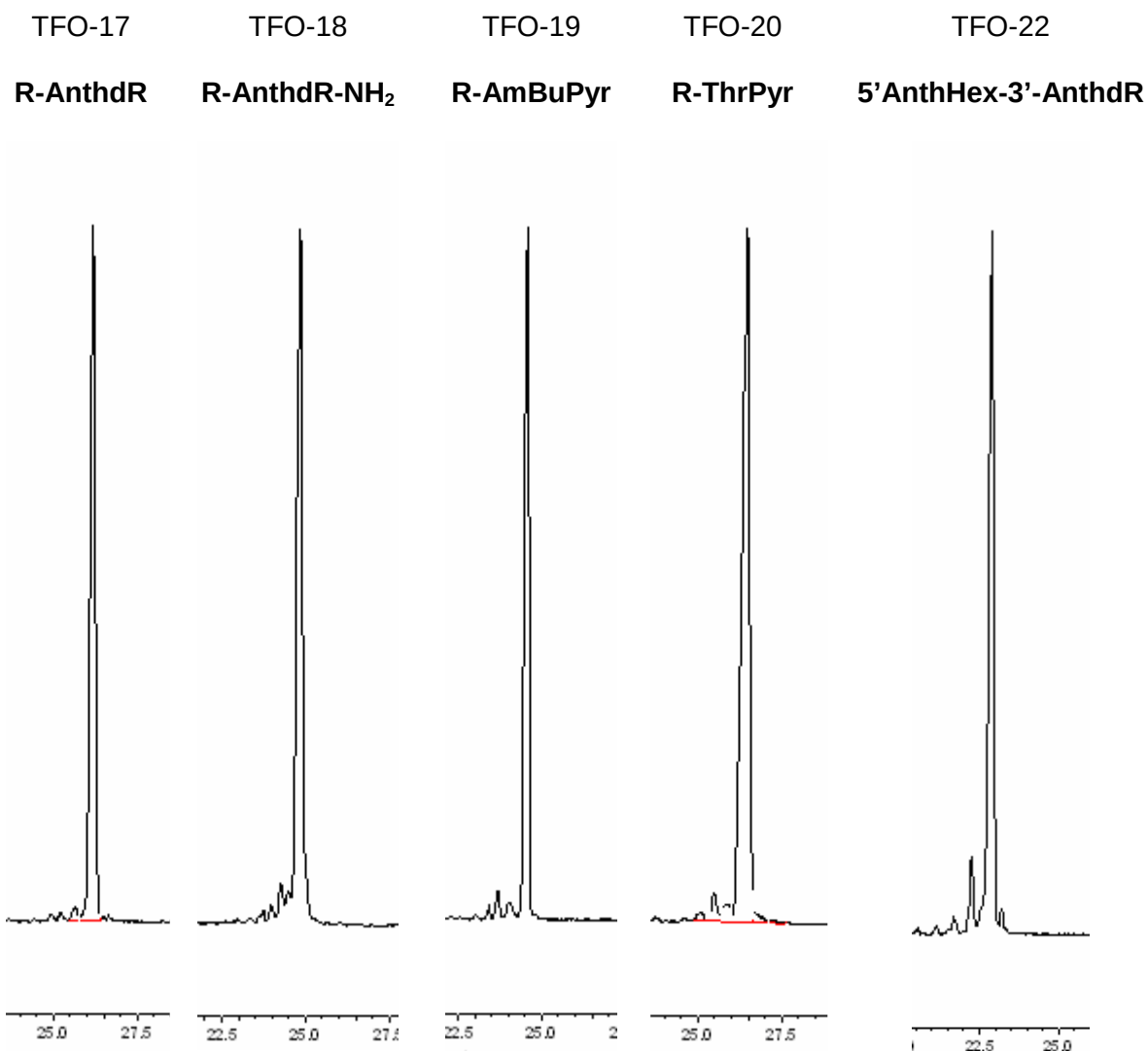
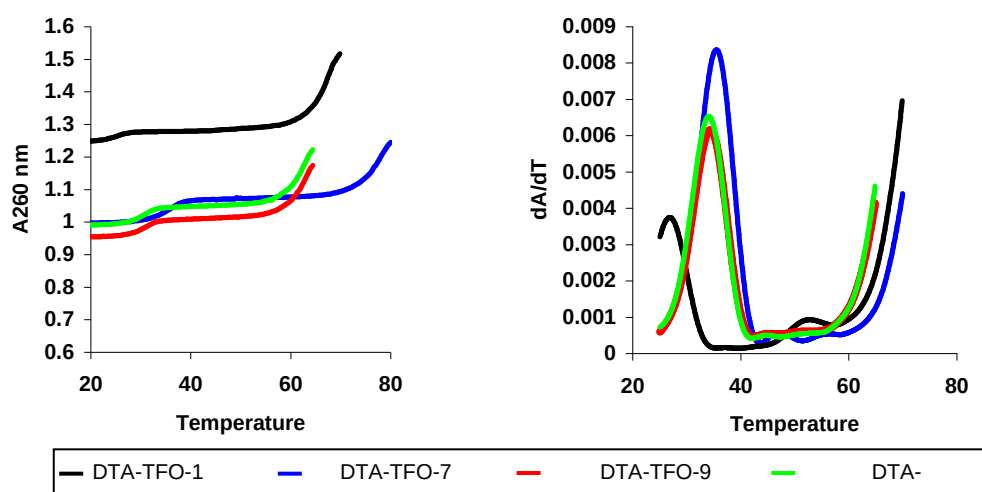
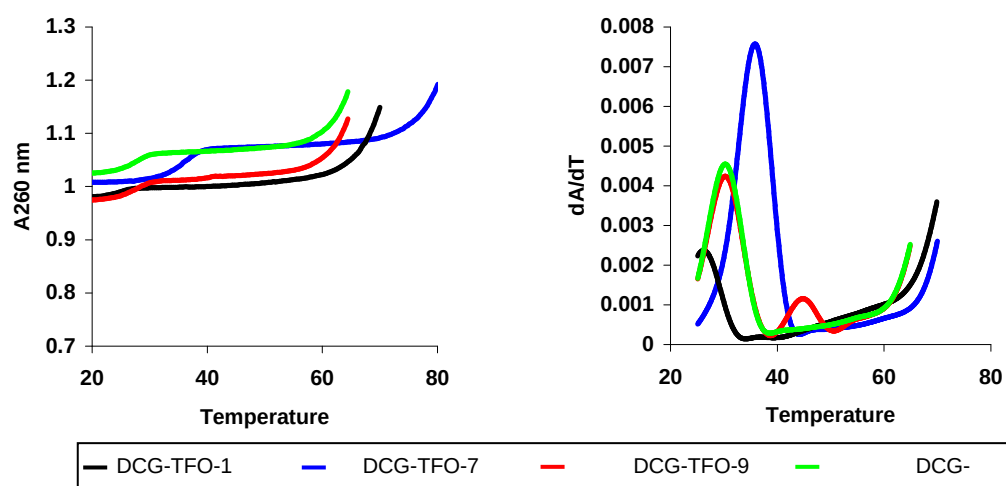


Figure S1. Representative Capillary electrophoresis (CE) analysis of modified oligonucleotides (TFO-11, TFO14-20 and TFO-22) performed after HPLC purification. The purity of oligonucleotides was confirmed by injection (0.4 OD/100 μ L) of each sample individually on ssDNA 100-R Gel. Tris-Borate 7M Urea were used (kit N°477480) on a Beckman coulter P/ACE™ MDQ Capillary Electrophoresis system using 32 Karat software. UV-254, injection voltage 10.0kV and separation voltage 9.0 kV (45.0 min duration). The x-axis shows time in min and the y-axis is UV absorbance at 254 nm.

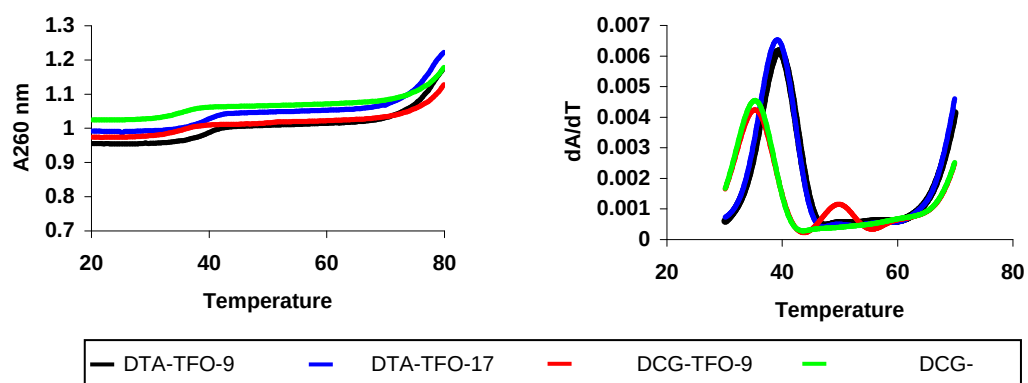
A)



B)



C)



D)

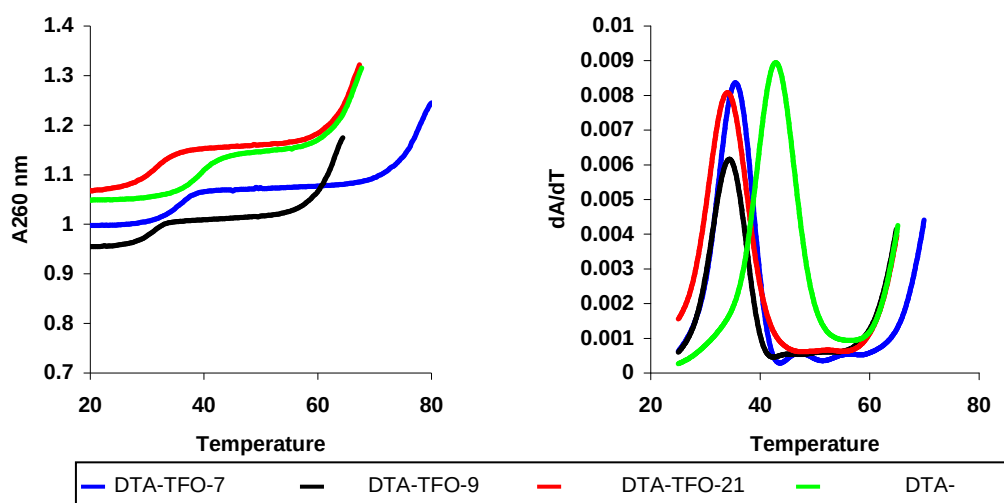


Figure S2. Representative UV melting curves and derivatives. 3'-Anthraquinone TFOs with TA and CG base pair duplexes. Spectra were recorded on a Cary 400 Scan UV-visible spectrophotometer (Varian) at a ratio of 3/1 (TFO/Duplex) in 10 mM phosphate buffer, 200 mM NaCl at pH 6.2, measured at 260 nm. T_m values were calculated using Cary Win UV thermal application software. A) Effect of 3'-modification on TA recognition. UV melting study performed between DTA and TFO-1 (T-P), TFO-7 (T-AnthdR), TFO-9 (S-AnthdR), TFO-17 (R^{An} -AnthdR) at pH 6.2. B) Effect of 3'-modification on CG recognition. UV melting study performed between DCG and TFO-1 (T-P), TFO-7 (T-AnthdR), TFO-9 (S-AnthdR), TFO-17 (R^{An} -AnthdR) at pH 6.2. C) Direct comparison of 3'-anthraquinone TFOs on TA or CG base pairs. UV melting study was performed between DTA and DCG and TFOs 9 (S-AnthdR) & 17 (R^{An} -AnthdR) at pH 6.2. D) Effect of 5' and 5'-3'-anthraquinone on TA recognition. UV melting study performed between DTA and TFO-7 (T-AnthdR), TFO-9 (S-AnthdR), TFO-21 (5'-Anth-Hex) and TFO-22 (5'-AnthHex-3'-AnthdR) at pH 6.2.

Table S1. Mass spectra of triplex forming oligonucleotides (TFOs) were recorded by negative mode electrospray on a Fisons VG platform spectrometer in water with tri-isopropylamine (2 μ L of 1% solution in MeOH). Duplexes were analysed by MALDI-TOF using a ThermoBioAnalysis Dynamo MALDI-TOF spectrometer in positive ion with oligo-dT as reference.

Code number	Sequence	MW required	MW found
DTA	5' -GTGTTAGGAAGAGAAAAAGAAGGT ^H 3' -CACAAATCCTTCTCTTTTTCTT ^H ATCCA	16961	16960
DAT1	5' -GTGTTAGGAAGAGAAAAAGAAAGGT ^H 3' -CACAAATCCTTCTCTTTTTCTT ^H ATCCA	16961	16958
DAT2	5' -GTGTTAGGAAGAGAAAAAGAAAGGT ^H 3' -CACAAATCCTTCTCTTTTTCTT ^H ATCCA	16961	16959
DCG	5' -GTGTTAGGAAGAGAAAAAGAAAGGT ^H 3' -CACAAATCCTTCTCTTTTTCTT ^H ATCCA	16962	16948
DGC1	5' -GTGTTAGGAAGAGAAAAAGAAAGGT ^H 3' -CACAAATCCTTCTCTTTTTCTT ^H ATCCA	16962	16959
DGC2	5' -GTGTTAGGAAGAGAAAAAGAAAGGT ^H 3' -CACAAATCCTTCTCTTTTTCTT ^H ATCCA	16962	16963
TFO-1	TCCTTCTCTTTTTCTTT-P	5475	5482
TFO-2	TCCTTCTCTTTTTCTTC-P	5460	5461
TFO-3	TCCTTCTCTTTTTCTTS-P	5625	5630
TFO-4	TCCTTCTCTTTTTCTTR ^{An} -P	5632	5633
TFO-5	TCCTTCTCTTTTTCTTC-AnthdR	5864	5866
TFO-6	TCCTTCTCTTTTTCTTC-ThrPyr	5761	5764
TFO-7	TCCTTCTCTTTTTCTTT-AnthdR	5881	5885
TFO-8	TCCTTCTCTTTTTCTTT-ThrPyr	5776	5783
TFO-9	TCCTTCTCTTTTTCTTS-AnthdR	6028	6035
TFO-10	TCCTTCTCTTTTTCTTS-AnthdR-NH ₂	6143	6147
TFO-11	TCCTTCTCTTTTTCTTS-AnthThr	5886	5896
TFO-12	TCCTTCTCTTTTTCTTS-AmBuPyr	6050	6054
TFO-13	TCCTTCTCTTTTTCTTS-ThrPyr	5922	5926
TFO-14	TCCTTCTCTTTTTCTTS-MeAcPyr	5936	5945
TFO-15	TCCTTCTCTTTTTCTTS-DAEdR	5793	5802
TFO-16	TCCTTCTCTTTTTCTTS-Pam	5940	5948
TFO-17	TCCTTCTCTTTTTCTTR ^{An} -AnthdR	6036	6033
TFO-18	TCCTTCTCTTTTTCTTR ^{An} -AnthdR-NH ₂	6150	6151
TFO-19	TCCTTCTCTTTTTCTTR ^{An} -AmBuPyr	6059	6063
TFO-20	TCCTTCTCTTTTTCTTR ^{An} -ThrPyr	5931	5935
TFO-21	AnthHex-TCCTTCTCTTTTTCTTT-P	5976	5979
TFO-22	AnthHex-TCCTTCTCTTTTTCTTT-AnthdR	6377	6379
TFO-23	AnthHex-TCCTTCTCTTTTTCTTT-ThrPyr	6272	6276