Supporting information

End-capped HyBeacon Probes for the Analysis of Human Genetic Polymorphisms Related to Warfarin Metabolism

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| | Modification | Oligonucleotide sequence |
|----|---------------------------------|--|
| 1 | Matched sequence | 3'-CCTCTTGAACAC <u>G</u> GTCCTCAATGCTCC-5' |
| 2 | Mismatched sequence | 3'-CCTCTTGAACAC <u>A</u> GTCCTCAATGCTCC-5' |
| 3 | 3'-P | 5'-CGATFGAGGACCGFGTTCAAG-P-3' |
| 4 | 5'-TMS-3'-P | 5'-TMS-CGATFGAGGACCGFGTTCAAG-P-3' |
| 5 | 3'-AnthdR | 5'-CGATFGAGGACCGFGTTCAAG-AnthdR-3' |
| 6 | 5'-TMS-3'-AnthdR | 5'-TMS-CGATFGAGGACCGFGTTCAAG-AnthdR-3' |
| 7 | 3'-AnthdRNH ₂ | 5'-CGATFGAGGACCGFGTTCAAG-AnthdRNH2-3' |
| 8 | 5'-TMS-3'-AnthdRNH ₂ | 5'-TMS-CGATFGAGGACCGFGTTCAAG-AnthdRNH2-3' |
| 9 | 3'-AmBuPyr | 5'-CGATFGAGGACCGFGTTCAAG-AmBuPyr-3' |
| 10 | 5'-TMS-3'-AmBuPyr | 5'-TMS-CGATFGAGGACCGFGTTCAAG-AmBuPyr-3' |
| 11 | 3'-ThrPyr | 5'-CGATFGAGGACCGFGTTCAAG-ThrPyr-3' |
| 12 | 5'-TMS-3'-ThrPyr | 5'-TMS-CGATFGAGGACCGFGTTCAAG-ThrPyr-3' |

Table S1: List of oligonucleotides. Twelve oligonucleotides were prepared: two targets and ten probes labelled with two internal fluorescein-dT residues (F), a 3'-modification and/or a 5'-trimethoxystilbene (TMS). P: phosphate.

| Sequence | Modification | MW calc. | MW found |
|----------|---------------------------------|----------|-----------|
| ODN1 | Matched sequence | 8128 | 8135 |
| ODN2 | Mismatched sequence | 8113 | 8125 |
| ODN3 | 3'-P | 7589 | 7600 |
| ODN4 | 5'-TMS/ 3'-P | 8021 | 8031 |
| ODN5 | AnthdR | 8067 | 8099 |
| ODN6 | 5'-TMS-AnthdR | 8489 | 8494 |
| ODN7 | AnthdRNH ₂ | 8168 | 8169 |
| ODN8 | 5'-TMS-3'-AnthdRNH ₂ | 8603 | 8606 |
| ODN9 | 3'-AmBuPyr | 8079 | 8078/8100 |
| ODN10 | 5'-TMS-3'-AmBuPyr | 8513 | 8514 |
| ODN11 | 3'-ThrPyr | 7948 | 7950 |
| ODN12 | 5'-TMS-3'-ThrPyr | 8383 | 8386 |

Table S2: Mass spectra. Mass spectra of all targets and HyBeacons® were recorded by negative mode electrospray on a Fisons VG platform spectrometer in water with Triisopropylamine (2%).





Figure S1: Capillary electrophoresis (CE) analysis of modified oligonucleotides 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/ACETM MDQ Capillary Electrophoresis system using 32 Karat software. UV-254, inject voltage 10.0 kV and separate 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.

Figure S2: UV spectra

A) UV spectra of 3'-caps





B) UV/vis spectra of 5'-Trimethoxystilbene oligonucleotide.



Fluorescence properties of end-caps.

The above end-caps and 5'-modified TMS-oligonucleotide were excited by irradiation at 460 nm. None of them yielded a fluorescence emission signal.

Quantum yield determination of 3'-end caps (350 nm excitation)

Quantum yields (Φ) of the 3'-modifications were calculated at an optical density of 0.01 at the excitation wavelength of 350 nm for the various 3'- modifications using a solution of quinine sulphate in 1 M H₂SO₄ as a standard while compounds were dissolved in MeOH.

Determination of extinction coefficients of 3'-modifications

General Experimental

Flash chromatography was performed on silica (40-63 μ m) purchased from Fisher Scientific and thin layer chromatography was performed on Merck Kieselgel 60F₂₅₄

coated plates (0.22 mm thickness, aluminum backed). Compounds were visualized by staining with vanillin (2g of vanillin in 100 mL of EtOH/ H_2SO_4 98:2) and by ultraviolet absorbance at 254nm. ¹H and ¹³C NMR spectra were recorded on a Bruker AV300 or a Bruker DPX 400 spectrometer. All spectra were internally referenced to the appropriate residual undeuterated solvent signal. Chemical shifts are given in ppm relative to tetramethylsilane. *J* values are given in Hz and corrected to within 0.5 Hz. Multiplicities of ¹³C signals were determined using the DEPT spectral editing technique. High-resolution mass spectra were recorded in acetonitrile, methanol or water (HPLC grade) using the electrospray technique on a Bruker APEX III FT-ICR mass spectrometer.

3'-Oligonucleotide modifications

Anthraquinone and Pyrene analogues for 3'-oligonucleotide modification and the corresponding oligonucleotide synthesis resins were prepared as described previously¹ These compounds were detritylated by the general procedure outlined below to obtain extinction coefficients for calculating oligonucleotide concentrations. DMTr protected compounds (1-4) were dissolved in a solution of 3% trichloroacetic acid (TCA) in CH_2Cl_2 (oligonucleotide grade) at a concentration of 0.03 M. The mixture was stirred at room temperature until detritylation was complete (TLC CH_2Cl_2 : MeOH 90:10). The reaction mixture was then quenched with a saturated solution of NaHCO₃ and CH_2Cl_2 was added. The organic layer was washed with H_2O and brine, dried over Na_2SO_4 and evaporated to dryness. The crude product was purified by silica gel flash chromatography using an Isolute cartridge (Biotage) eluting with a gradient of MeOH in CH_2Cl_2 .

N-(6-(9,10-Dihydro-9,10-dioxoanthracen-1-ylamino)hexyl)-2'-deoxy-D-

ribofuranose-1-\beta-acetamide (1). The reaction was performed on 84 mg (0.107 mmol) in 3.5 mL of TCA 3% in CH₂Cl₂. 50 mg, 97%.

R_f CH₂Cl₂: MeOH (90:10) 0.31. **NMR** ¹**H** (400 MHz, DMSO-d6) $\delta_{\rm H}$ 9.73-9.71 (2H, t, *J*= 5 Hz, NH), 8.26-8.18 (2H, dd, *J*= 7.5 Hz, H-Ar), 7.97-7.88 (3H, m, H-Ar), 7.71-7.67 (1H, t, *J*= 8.5 Hz, H-Ar), 7.29-7.27 (1H, d, *J*= 8.5 Hz, H-Ar), 4.99-4.98 (1H, d, *J*= 4 Hz, OH), 4.74-4.71 (1H, m, OH), 4.43-4.36 (1H, m, H1'), 4.16 (1H, d, *J*= 2.48 Hz, H3'), 3.72-3.69 (1H, m, H4'), 3.46-3.37 (4H, m, CH₂), 3.21-3.10 (2H, m, CH₂), 2.48-2.29 (2H, ddd, *J*₁= 7 Hz, *J*₂= 6.5 Hz, *J*₃= 33.1 Hz, H5'), 1.92-1.88 (1H, m, H2'),

1.77-1.69 (3H, m, CH₂, H2'), 1.54-1.44 (6H, m, CH₂). **NMR** ¹³C (100 MHz, DMSOd6) δ_{C} 183.77, 182.66 (<u>C</u>O), 180.24 (<u>C</u>O-NH), 169.49 (<u>C</u>-Ar), 151.18 (<u>C</u>-Ar), 135.41, 134.26 (<u>C</u>H-Ar), 134.20 (<u>C</u>-Ar), 133.72 (<u>C</u>H-Ar), 133.21 (<u>C</u>-Ar), 128.19, 126.25 (<u>C</u>H-Ar), 126.06 (<u>C</u>-Ar), 118.35, 114.79 (<u>C</u>H-Ar), 111.70 (<u>C</u>-Ar), 87.15 (C1'), 74.72 (C4'), 71.88 (C3'), 62.26 (<u>C</u>H₂-NH), 42.6 (<u>C</u>H₂-CO), 40.74 (C2'), 38.85 (C5'), 29.53, 29.01, 26.76, 26.62 (<u>C</u>H₂). UV-Vis λ_{max} (MeOH)/nm 510 (ϵ /dm³.mol⁻¹.cm⁻¹ 2360). Φ (MeOH, 350 nm) 0.04. **m/z LRMS** [ES⁺, MeOH] 503.3 (M+Na⁺, 100%), 983.8 (2M+Na⁺, 50%). **m/z HRMS** (M+Na⁺) (C₂₇H₃₂N₂O₆Na): calc. 503.2158 found 503.2153.

N-(6-(9,10-Dihydro-9,10-dioxoanthracen-1-ylamino)-5-yl-(6-aminohexyl)hexyl)-2-deoxy-D-ribofuranose-1-β-acetamide (2). The reaction was performed on 0.114 g (0.115 mmol) in 4 mL of TCA 3% in CH_2Cl_2 . Purple solid 75 mg, 95%.

R_f CH₂Cl₂: MeOH (90:10) 0.33. **NMR** ¹**H** (400 MHz, **DMSO-d**₆) $\delta_{\rm H}$ 9.77-9.74 (2H, t, *J*= 5 Hz, N<u>H</u>), 9.51 (1H, bs, N<u>H</u>), 7.89-7.86 (1H, t, *J*= 5.2 Hz, NH), 7.73-7.69 (2H, t, *J*= 7.7 Hz, <u>H</u>-Ar), 7.53-7.51 (2H, d, *J*= 7.5 Hz, <u>H</u>-Ar), 7.24-7.22 (2H, d, *J*= 8.5 Hz, <u>H</u>-Ar), 4.97 (1H, bs, OH), 4.72 (1H, bs, OH), 4.42-4.35 (1H, m, H1'), 4.15-4.14 (1H, m, H4'), 3.71-3.68 (1H, m, H3'), 3.58-3.42 (12H, m, C<u>H</u>₂), 3.33-3.28 (2H, q, *J*= 6.5 Hz, C<u>H</u>₂-CO), 2.47-2.28 (2H, ddd, *J*₁= 6.5 Hz, *J*₂= 14.0 Hz, *J*₃= 33.1 Hz, H5'), 1.91-1.86 (1H, m, H2'), 1.78-1.45 (13H, m, C<u>H</u>₂, H2'). **NMR** ¹³C (100 MHz, DMSO-d6) $\delta_{\rm C}$ 185.2 (<u>C</u>O), 181.2 (<u>C</u>OCF₃), 170.5 (<u>C</u>-CO), 152.0 (C-NH), 136.5 (<u>C</u>H-Ar), 117.9 (<u>C</u>H-Ar), 115.2 (<u>C</u>H-Ar), 112.8 (<u>C</u>F₃), 88.2 (C1'), 75.8 (C4'), 72.9 (C3'), 63.3 (<u>C</u>H₂-CO), 43.0 (C2'), 39.3 (C5'), 29.4, 29.0, 28.9, 28.6, 26.7, 26.6, 26.3 (<u>C</u>H₂). UV-Vis $\lambda_{\rm max}$ (MeOH)/nm 525 (ε/ dm³.mol⁻¹.cm⁻¹ 1077). Φ (MeOH, 350 nm) 0.04. **m/z LRMS** [ES⁺, MeOH] 713.5 (M+Na⁺, 100%). **m/z HRMS** (M+Na⁺) (C₃₅H₄₅N₄O₇F₃Na): calc. 713.3138. found 713.3133.

1-β-O-(hexylamidopropylpyrene-1-yl)-2-deoxy-D-ribofuranose (3). The reaction was performed on 0.200 g (0.25 mmol) in 8 mL TCA 3% in CH₂Cl₂. Colourless foam, 30 mg, 24 %.

R_f CH₂Cl₂: MeOH (90:10) 0.51. **NMR** ¹**H** (400 MHz, DMSO-d₆) $\delta_{\rm H}$ 8.39-8.36 (1H, d, *J*= 9.1 Hz, NH), 8.28-8.19 (4H, m, H-Ar), 8.15-8.09 (2H, dd, *J*₁= 8.8 Hz, *J*₂= 10.2 Hz, H-Ar), 8.07-8.02 (1H, t, *J*= 7.7 Hz, H-Ar), 7.94-7.91 (1H, d, *J*= 8.0 Hz, H-Ar), 7.83-

7.79 (1H, t, *J*= 5.8 Hz, H-Ar), 4.97-4.95 (1H, dd, *J*₁= 2.6 Hz, *J*₂= 5.8 Hz, H1'), 4.84-4.82 (1H, d, *J*= 5.1 Hz, H4'), 4.65-4.62 (1H, t, *J*= 5.5 Hz, H3'), 3.95-3.86 (1H, m, OH), 3.71-3.66 (1H, m, OH), 3.60-3.47 (2H, m, CH₂), 3.41-3.24 (6H, m, CH₂), 3.09-3.02 (2H, q, *J*= 6.3 Hz, CH₂), 2.29-2.20 (3H, m, CH₂, H2'), 2.06-1.96 (2H, q, *J*= 7.4 Hz, H5'), 1.66-1.58 (1H, m, H2'), 1.47-1.37 (6H, m, CH₂). **NMR** ¹³C (100 MHz, DMSO-d6) $\delta_{\rm C}$ 172.1 (CO), 137.0, 131.3, 130.9, 129.7 (C-Ar), 128.8, 127.9, 127.6, 126.9, 126.6, 125.4, 125.2 (CH-Ar), 124.6 (C-Ar), 123.9 (C-Ar), 103.3 (C1'), 85.1 (C4'), 70.5 (C3'), 67.2 (CH₂), 61.7 (C5'), 41.5 (C2'), 38.8, 35.5, 32.7, 29.6, 28.0, 26.7, 25.9 (CH₂). UV-Vis $\lambda_{\rm max}$ (MeOH)/nm 340 (ε/ dm³.mol⁻¹.cm⁻¹ 161523), 324 nm (124267), 274 (219444) and 264 (108333). Φ (MeOH, 350 nm) 0.12. **m/z LRMS** [ES⁺, MeOH] 526.3 (M+Na⁺, 100%), 1029.7 (2M+Na⁺, 10%). **m/z HRMS** (M+Na⁺) (C₃₁H₃₇NO₅Na): calc. 526.2569 found 526.2564.

L-threoninolamidoprop-1-yl pyrene (4). The reaction was performed on 200 mg (0.295 mmol) in 10 mL of TCA 3% in CH₂Cl₂. Colourless solid 75 mg, 68%.

R_f CH₂Cl₂: MeOH (90:10) 0.46. **NMR** ¹**H** (400 MHz, DMSO-d6) δ_H 8.51-8.47 (1H, dd, J_1 = 4.3 Hz, J_2 = 9.3 Hz, <u>NH</u>), 8.37-8.11 (7H, m, <u>H</u>-Ar), 8.04-8.01 (1H, dd, J_1 =4.5 Hz, J_2 = 8 Hz, <u>H</u>-Ar), 7.56-7.53 (1H, m, <u>H</u>-Ar), 4.73 (1H, bs, OH), 4.05 (1H, bs, C<u>H</u>-Thr), 3.89 (1H, bs, C<u>H</u>-Thr), 3.66 (1H, bs, OH), 3.53 (2H, bs, C<u>H</u>₂), 3.46-3.41 (2H, m, C<u>H</u>₂), 2.47-2.46 (2H, m, C<u>H</u>₂), 2.15 (2H, bs, C<u>H</u>₂), 1.20-1.17 (3H, t, J= 5.2 Hz, C<u>H</u>₃-Thr). **NMR** ¹³C (100 MHz, DMSO-d6) δ_C 180.2 (CO), 172.2 (C-Ar), 136.6, 130.8, 130.3, 129.2, 128.1 (C-Ar), 127.4, 127.3, 127.1, 126.3, 126.0, 124.8, 124.7, 124.6 (CH-Ar), 124.0 (C-Ar), 123.4 (CH-Ar), 64.4 (<u>C</u>H-Thr), 60.6 (<u>C</u>H₂), 55.6 (<u>C</u>H-Thr), 35.1, 32.2, 27.8 (<u>C</u>H₂), 20.1 (<u>C</u>H₃). UV-Vis λ_{max} (MeOH)/nm 340 (ε/dm³.mol⁻¹.cm⁻¹ 29864), 324 (23000), 274 (40889) and 264 (20665). Φ (MeOH, 350 nm) 0.1. **m/z LRMS** [ES⁺, MeOH] 398.3 (M+Na⁺, 100%), 773.6 (2M+Na⁺, 50%). **m/z HRMS** (M+Na⁺) (C₂₄H₂₅NO₃Na): calc. 398.1732 found 398.1729.

All compounds were detritylated efficiently except 1- β -*O*-(hexylamidopropylpyrene-1-yl)-2-deoxy-D-ribofuranose **3** for which an impurity corresponding to 6hydroxyhexylamidopropylpyrene was identified by MS and NMR analysis. The same side-reaction was also observed during oligonucleotide synthesis to yield modest amounts of oligonucleotide products without 3'-pyrene. These impurities were easily removed by reversed-phase HPLC. DMTr protected compounds 1-4 were attached *via* a succinyl linkage to long chain aminoalkyl silica resin (92 μ mol/g amino-loading, Link Technologies Ltd) and used in oligonucleotide synthesis.¹



Figure S3: A) Fluorescence melting curves of all HyBeacon® analogues against matched sequence. B) Fluorescence melting curves of all HyBeacon® analogues against mismatch sequence. C) Fluorescence melting curves of ThrPyr analogue in presence and in absence of 5'-TMS in comparison to control HyBeacons®. Fluorescence melting was recorded in TaKaRa buffer 1x, 1M NaCl.

UV melting curves



Figure S4: UV melting analysis. A) Pyrene analogues and control HyBeacons® against matched sequence (ODN1). B) Pyrene analogues and control HyBeacons® against mismatched sequence (ODN2). C) Direct comparison of 5'-TMS-3'ThrPyr (ODN12) and 3'-P HyBeacons® (ODN3) towards matched and mismatched sequence. UV melting recorded in phosphate buffer, 200 mM NaCl, pH 7.0.

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