

DNA-templated control of chirality and efficient energy transport in supramolecular DNA architectures with aggregation-induced emission

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Supporting Information

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1. Materials and methods

Solvents and reagents

Used Chemicals had the purification grade “for synthesis”, used solvents for synthesis, optical spectroscopy or analysis the grade “HPLC” or “pro analysis”. Water for the sample preparation was deionized and ultrafiltrated by a *Millipore Direct 8/16* from MERCK MILLIPORE. Unmodified and Atto dye-modified D-configured DNA-strands were obtained HPLC-purified and lyophilized from METABION. The L-configured DNA-strands were synthesized, described in the DNA synthesis part.

NMR spectroscopy

The NMR spectra were recorded on a *Bruker Advanced 400* or *Bruker Advanced 500*. The samples were dissolved in 0.5 mL denatured solvent from EURISOTOP. The chemical shifts are given in parts per million (ppm) relative to the standard tetramethylsilan (TMS). The spectra were calibrated against the ¹H- residues of the incompletely deuterated solvents.

Mass spectrometry

FAB-Mass spectra were measured on a FINNIGAN *MAT95 spectrometer*. MALDI-TOF spectra were recorded on a SHIMADZU *AXIMA Confidence spectrometer*. ESI-mass spectra recorded on a THERMOFISHER SCIENTIFIC *Q Exactive (Orbitrap)*.

Optical spectroscopy

Absorption spectra were recorded on a *Lambda 750* from PERKIN ELMER with a *PTP-6+6 Peltier System*. The fluorescence was determined on a *Fluoromax-4* from HORIBA SCIENTIFIC with an *AC 200 thermostat* from THERMO SCIENTIFIC. All samples were excited at 341 nm. Absolute fluorescence quantum yields were measured with a *Quantaaurus QY C11347* from HAMAMATSU ($\lambda_{exc} = 389$ nm).

Circular dichroism was measured on a JASCO *J-810 Spectropolarimeter* with the *peltier-element PTC-423S* (100 nm/min, 4 accumulations). FDCD spectra were recorded with a JASCO *J-1500 CD spectrometer* (0.2 nm resolution, 4 accumulations) equipped with a filter system and a *PML-534 detector* in a perpendicular positioning to the excitation light path. The Long pass filter was chosen according to the emission spectrum of the Tpe chromophore.

2. Synthesis

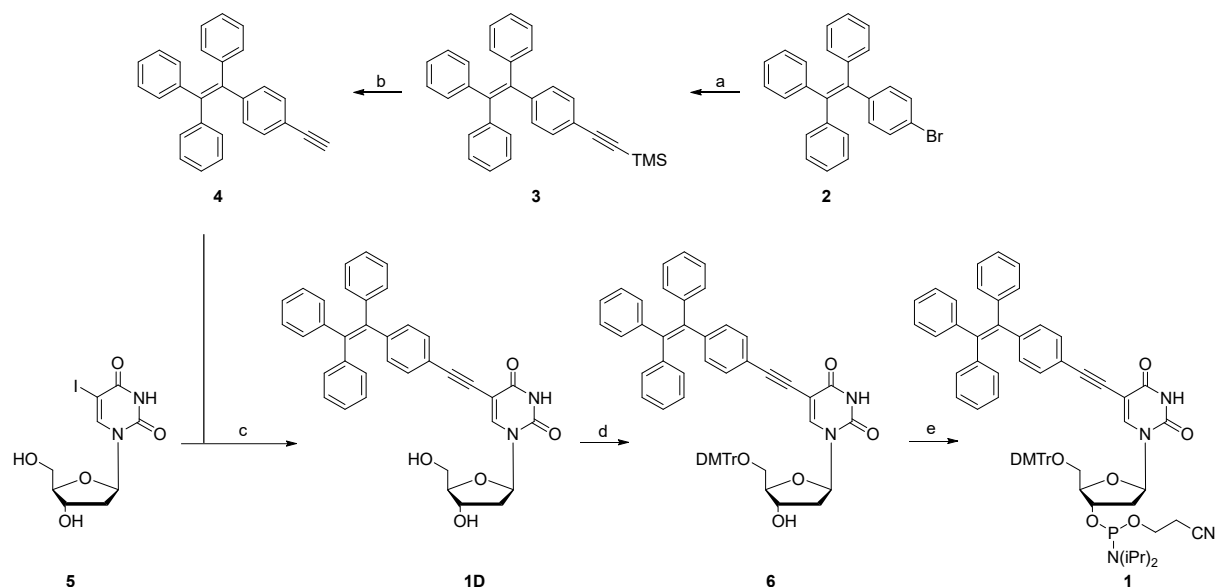


Figure S1. a) TMS-acetylene, PdCl₂(PPh₃)₂, CuI, PPh₃, THF/Et₃N, 60 °C, 19 h, 84%; b) KOH, THF/MeOH, r.t., 19 h, quant.; c) Pd(PPh₃)₄, CuI, DMF, Et₃N, 60 °C, 19 h, 81%; d) DMTrCl, pyridine, r. t., 19 h, 70%; e) iPr₂NP(Cl)OCH₂CH₂CN, DIPEA, DCM, r.t., 3h, 88%.

The synthesis of the acetylene-modified Tpe **4** was prepared according to the literature. ^[1]

Synthesis of Tpe-modified 5-Iodo-2'-deoxyuridine **1D**

A mixture of 0.40 g **4** (1.12 mmol, 1.00 eq.), 0.44 g 5-Iodo-2'-deoxyuridine **5** (1.23 mmol, 1.10 eq.) and 0.04 mg CuI (0.22 mmol, 0.20 eq.) was dissolved in 5 mL dry DMF and 0.39 mL triethylamine (2.81 mmol, 2.50 eq.) was added under argon. The reaction mixture was degassed through injection of argon under stirring for 10 min. After addition of 0.26 g Pd(PPh₃)₄ (0.22 mmol, 0.20 eq.) the reaction mixture was stirred by 60 °C for 19 h. Afterwards the solvents were removed under reduced pressure. The crude product was dissolved in DCM, washed with brine and dried over sodium sulfate. The solvent was again removed under reduced pressure. The crude product was purified by flash-column chromatography (silica, DCM:MeOH = 1:0 – 20:1). The desired product **1D** could be obtained as a white solid in a yield of 81% (0.53 g, 0.91 mmol).

¹H-NMR (500 MHz, DMSO): δ = 11.67 (s, 1H), 8.35 (s, 1H), 7.22 (d, *J* = 8.4 Hz, 2H), 7.19 – 7.09 (m, 9H), 7.00 – 6.94 (m, 8H), 6.12 (t, *J* = 6.5 Hz, 1H), 5.25 (d, *J* = 4.2 Hz, 1H), 5.14 (t, *J* = 4.7 Hz, 1H), 4.24 (p, *J* = 4.2 Hz, 1H), 3.80 (q, *J* = 3.3 Hz, 1H), 3.70 – 3.50 (m, 2H), 2.22 – 2.04 (m, 2H).

¹³C-NMR (126 MHz, DMSO): δ = 161.4, 149.4, 143.8, 143.6, 142.9, 142.7, 141.4, 139.8, 131.0, 130.7, 130.7, 130.6, 128.0, 127.9, 127.8, 126.8, 126.7, 120.4, 98.1, 91.7, 87.6, 84.8, 82.8, 69.9, 60.8, 40.2.

HR-MS (FAB): *m/z* calculated C₃₇H₃₀N₂O₅⁺ [*M*⁺] = 582.2155; found = 582.2154.

The L-configured nucleoside **1L** was synthesized in the same manner and the spectroscopic data was identical.

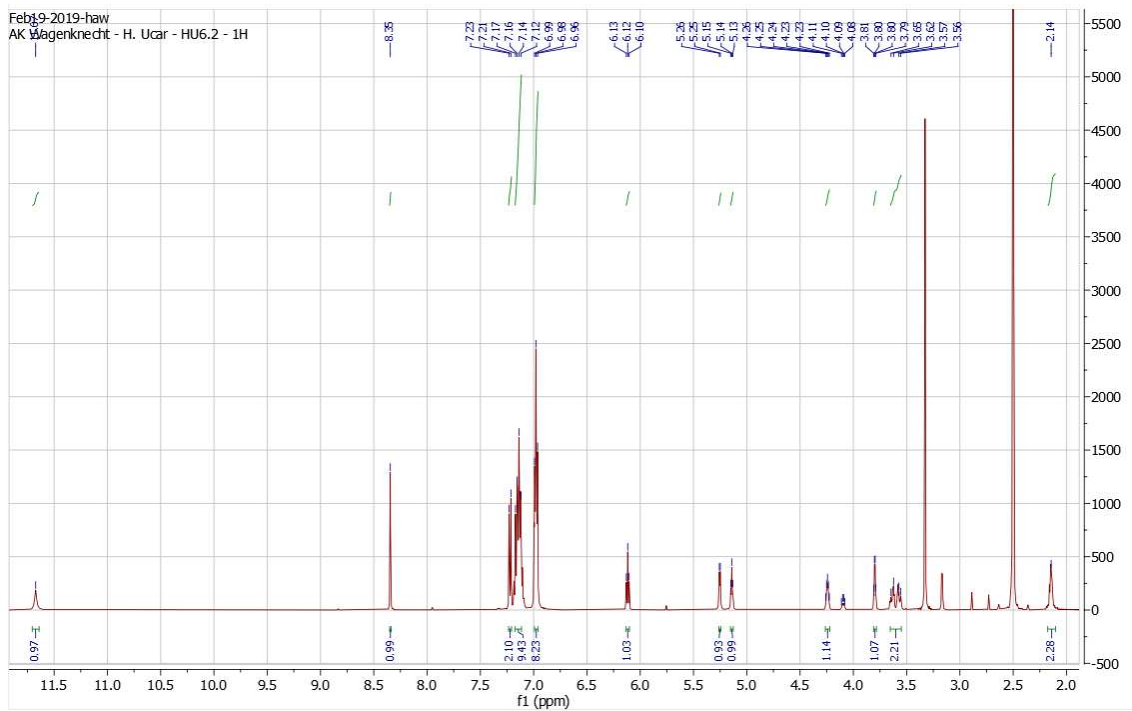


Figure S2. ^1H -NMR spectrum (500 MHz, DMSO) of **1D**.

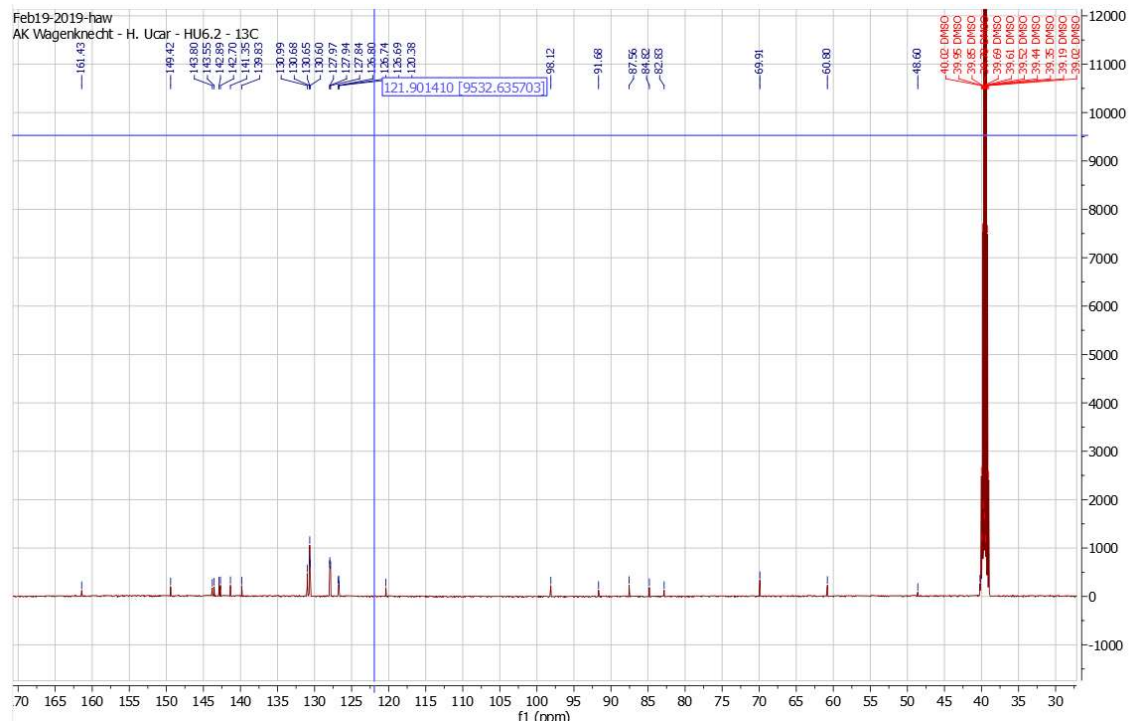


Figure S3. ^{13}C -NMR spectrum (126 MHz, DMSO) of **1D**.

hu-6-c3 #30 RT: 4.83 AV: 1 NL: 5.43E4
T: + c EI Full ms [84.44-700.44]

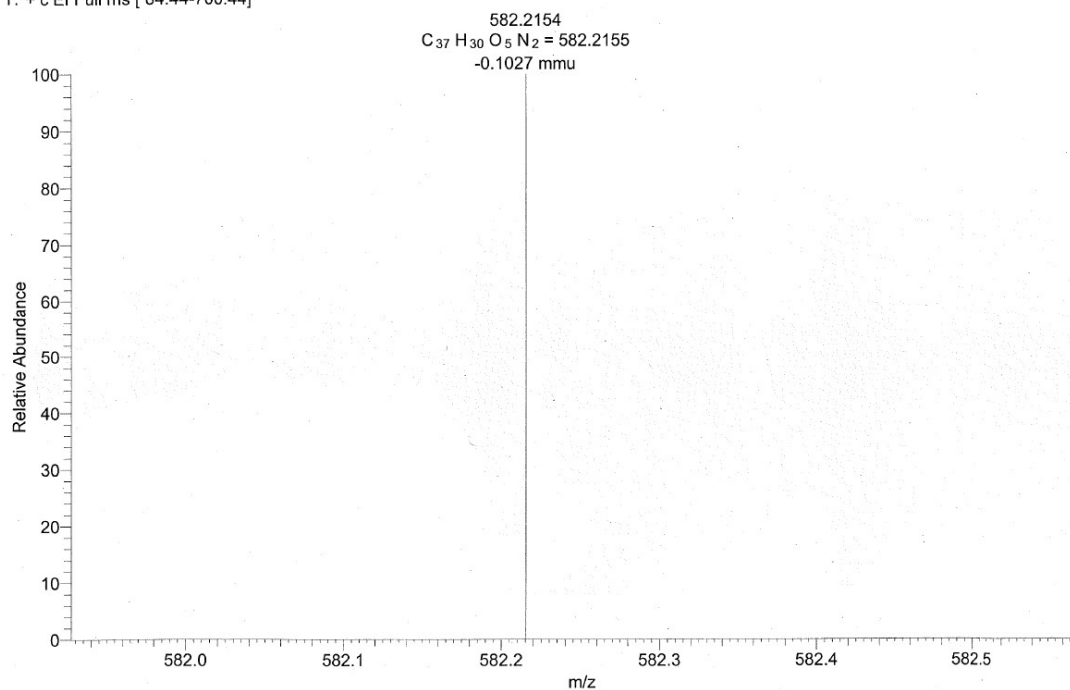


Figure S4. HR-MS (FAB) analysis of 1D.

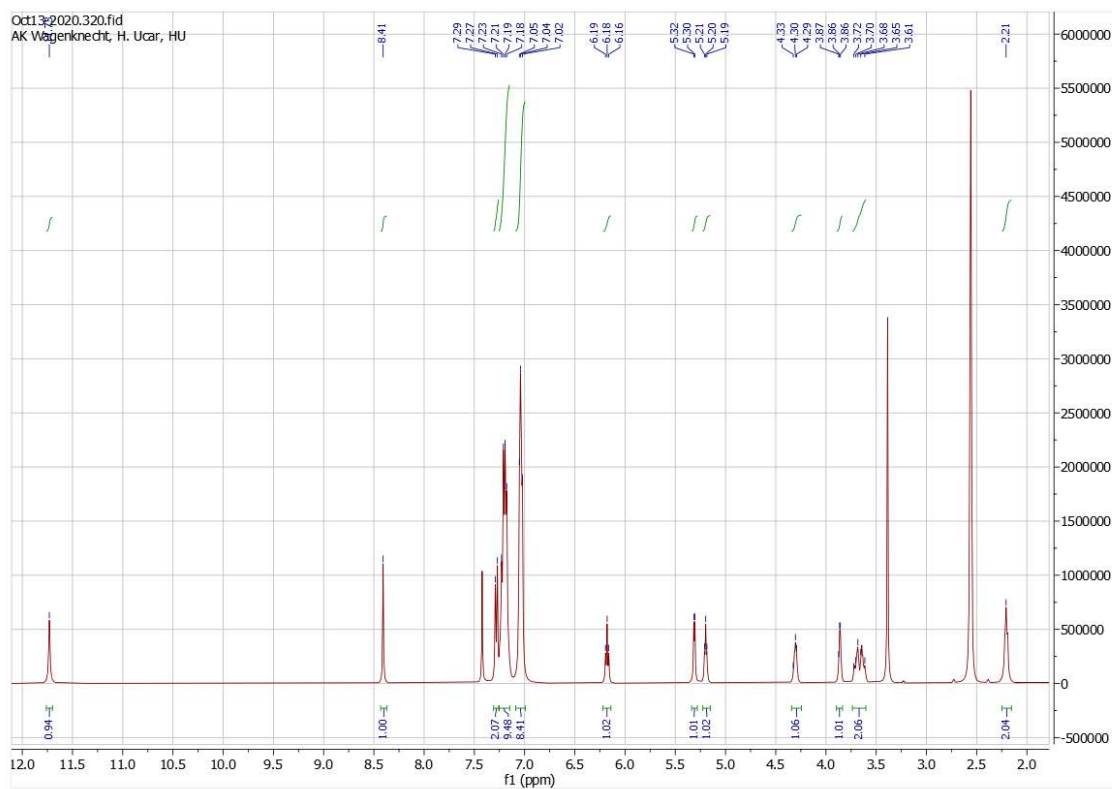


Figure S5. ¹H-NMR spectrum (400 MHz, DMSO) of 1L.

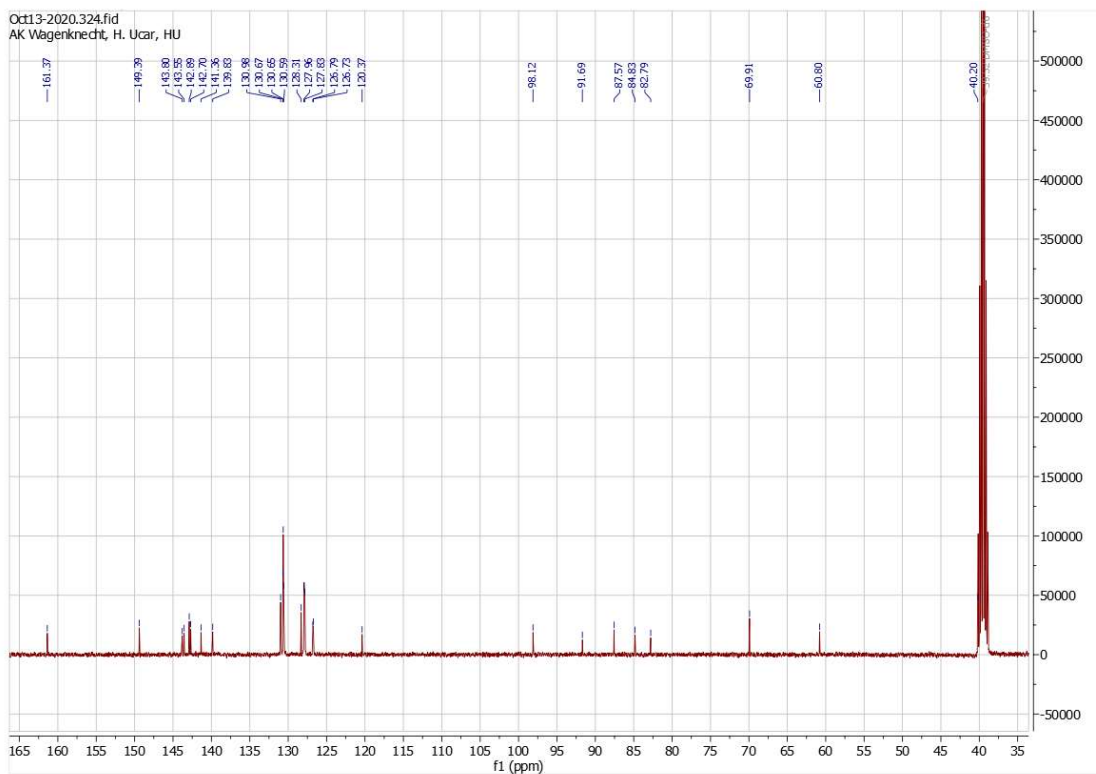


Figure S6. ^{13}C -NMR spectrum (126 MHz, DMSO) of **1L**.

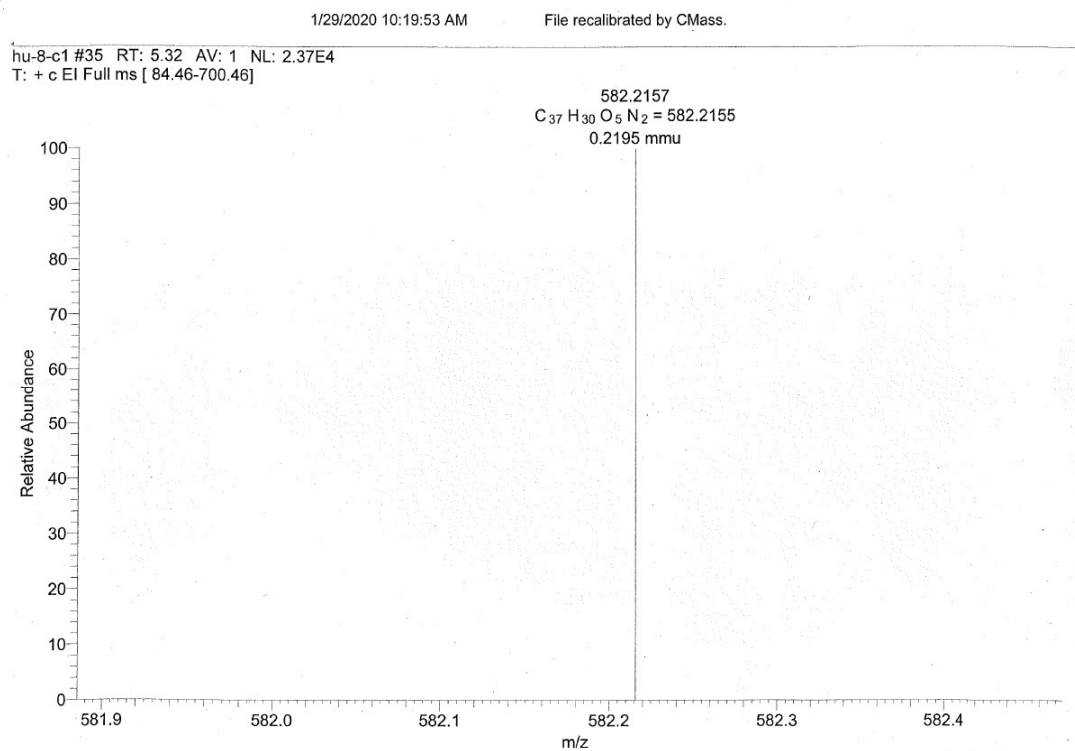


Figure S7. HR-MS (FAB) analysis of **1L**.

Compound 6

The compound **1D** (0.30 g, 0.51 mmol, 1.00 eq.) was lyophilized from benzene overnight. Afterwards it was dissolved in 5 mL dry pyridine under argon. After addition of 0.23 g 4,4'-dimethoxytrityl chloride (0.67 mmol, 1.30 eq.) the reaction mixture was stirred overnight. MeOH (2 mL) was added to the reaction and the solvents were removed under reduced pressure. The crude product was purified by flash-column chromatography (silica, DCM:MeOH = 99:1 + 0.1% NEt₃). The desired product was obtained as pale yellow solid in a yield of 70% (0.32 g, 0.36 mmol).

TLC (DCM/MeOH (1%)): R_f = 0.186

¹H-NMR (500 MHz, DMSO) δ = 11.73 (s, 1H), 8.02 (s, 1H), 7.37 (d, J = 9.0 Hz, 2H), 7.30 – 7.24 (m, 4H), 7.22 (t, J = 7.8 Hz, 2H), 7.15 – 7.10 (m, 10H), 7.02 – 6.92 (m, 6H), 6.88 (s, 4H), 6.81 (t, J = 8.9 Hz, 4H), 6.13 (t, J = 6.7 Hz, 1H), 5.34 (d, J = 4.4 Hz, 1H), 4.56 – 4.11 (m, 1H), 3.95 (dt, J = 5.5, 3.1 Hz, 1H), 3.63 (s, 6H), 3.18 (ddd, J = 44.6, 10.6, 3.9 Hz, 2H), 2.34 – 2.16 (m, 2H).

¹³C-NMR (126 MHz, DMSO) δ = 161.4, 158.0, 149.3, 144.6, 143.4, 142.9, 142.9, 142.8, 142.7, 141.2, 139.8, 135.5, 135.3, 130.7, 130.7, 130.6, 130.6, 129.7, 129.6, 128.3, 127.9, 127.9, 127.8, 127.5, 126.7, 126.6, 120.2, 113.2, 99.5, 98.5, 98.5, 91.9, 86.0, 85.9, 85.1, 82.1, 81.1, 70.5, 63.7, 55.0.

HR-MS (ESI): m/z calculated for C₅₈H₄₈N₂O₇⁺ [M⁺] = 884.3462; found = 884.3455.

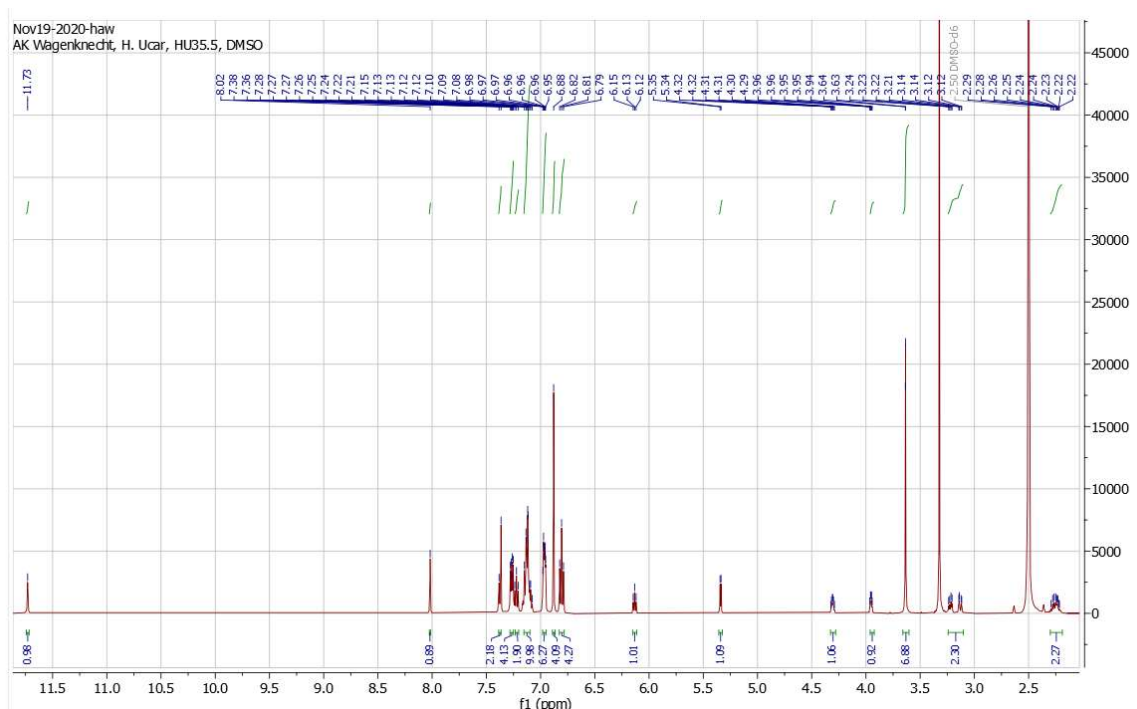


Figure S8. ¹H-NMR spectrum (500 MHz, DMSO) of **6**.

Compound 1

The compound **6** (0.30 g, 0.34 mmol, 1.00 eq.) was lyophilized from benzene overnight. Afterwards it was dissolved in 5 mL dry DCM under argon. After addition of 0.18 mL N,N-diisopropylethylamine (0.13 g, 1.02 mmol, 3.00 eq.) and 0.11 mL 2-cyanoethyl-N,N-diisopropylchlorophosphoramidite (0.12 g, 0.51 mmol, 1.50 eq.) the reaction mixture was stirred at room temperature for 3 h. The reaction mixture was purified by flash-column chromatography (silica, DCM:acetone = 5:1 + 0.1 % NEt₃). The desired product was obtained as pale yellow solid in a yield of 88 % (0.35 g, 0.30 mmol).

TLC (DCM/Aceton 5:1): R_f = 0.89

³¹P-NMR (162 MHz, MeOD): δ = 148.5, 148.2.

HR-MS (ESI): m/z calculated for C₆₇H₆₅N₄O₈P⁺ [M⁺] = 1084.4540; found = 1084.4533.

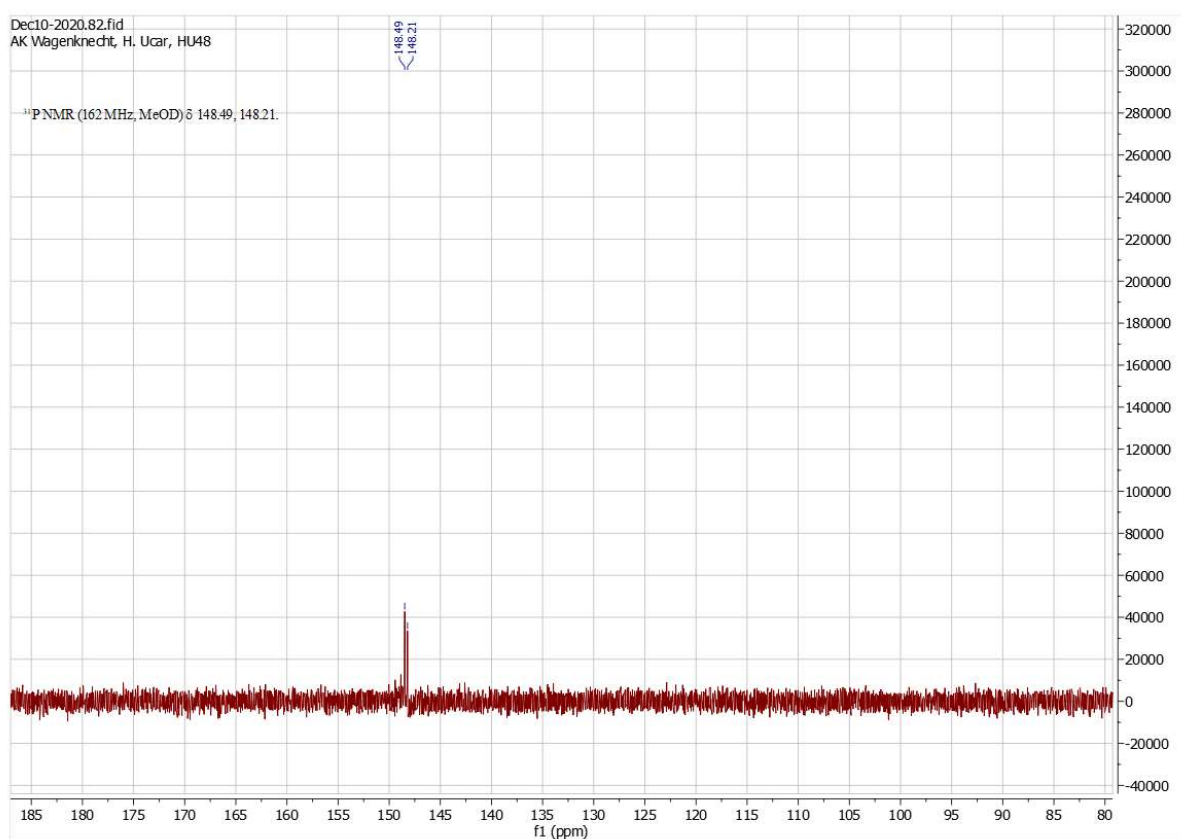


Figure S11. ³¹P-NMR spectrum (162 MHz, MeOD) of **1**.

HU48#1-20 RT: 0.02-0.36 AV: 20 NL: 1.21E6
T: FTMS + p ESI Fullms [1000.0000-1600.0000]

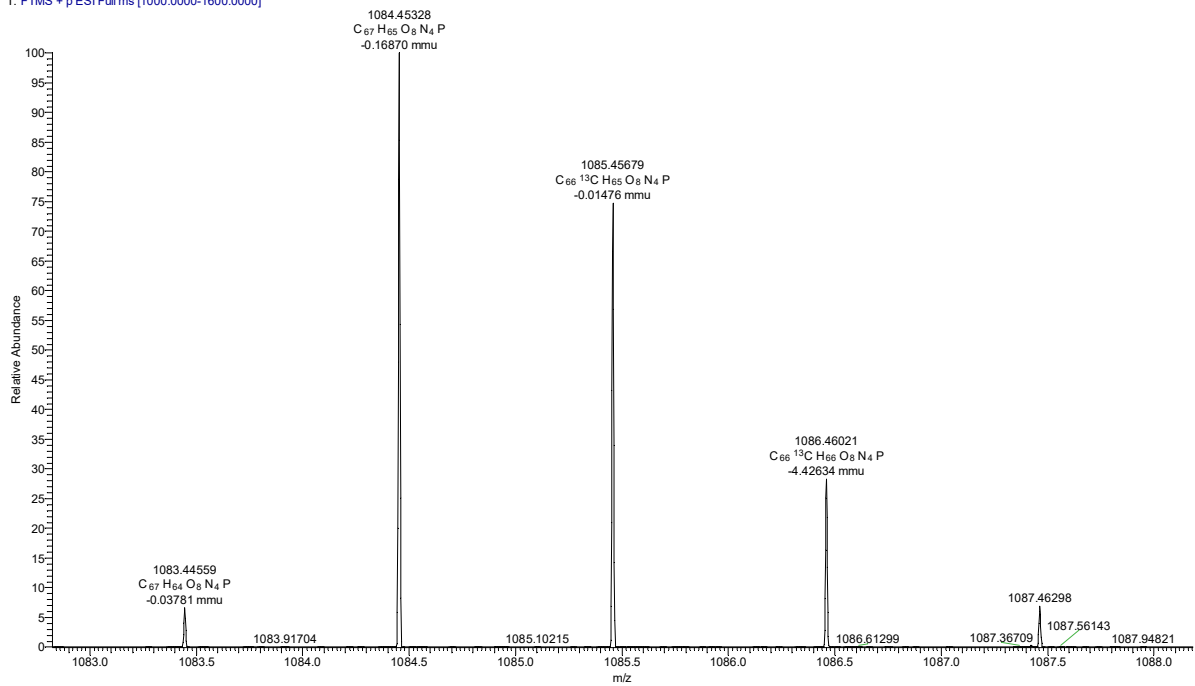


Figure S12. HR-MS(ESI) analysis of 1.

3. DNA preparation

The L-configured DNA-template strands (L-A₂₀ & L-T₂₀) and the Tpe-modified DNA strands **Tpe1a**, **Tpe1b**, **Tpe2a** and **Tpe2b** were synthesized on a *H-6 DNA/RNA Synthesizer* from K&A LABORGERÄTE using the standard protocol. The strands **Tpe1a** and **Tpe2b** were prepared with a longer coupling time of 500 s (3 pulses) and a higher concentration of the phosphoramidite (0.1 M). The β-L-deoxyadenosine/β-L-deoxythymidine phosphoramidites (0.067 M) and the CPG-columns (1 μmol, 500 Å) loaded with the first nucleobase are commercially available and were purchased from CHEMGENES. The 5'-terminal DMT protecting group was not removed during synthesis. The cleavage of the DNA strands from the CPG resin were achieved by heating the strands in 0.7 mM conc. aqueous ammonia solution (>25 %, trace select, FLUKA) to 60 °C for 16 h. The solvents were removed under reduced pressure and the DNA was purified using *Glen-Pak™ DNA Purification Cartridges* from GLEN RESEARCH with the manufacturer given standard procedure. As final step the DNA was further purified by reverse phase HPLC purification using the following conditions: A = NH₄OH buffer (50 mM), B = MeCN; gradient = 0-30 % in 30 min (L-A₂₀ and L-T₂₀); 0-50 % in 50 min (**TPE1a**, **TPE1b**, **TPE2a** and **TPE2b**). The HPLC runs were carried out on a THERMO FISCHER SCIENTIFIC (*Dionex UltiMate3000 auto sampler, software Chromeleon 7*) with a *VDSpher OptiBio PUR 300 S18-SE-column* (250 x 10 mm, 5 μm) and a flow rate of 2.5 mL/min. The DNA-strands were dissolved in water. The concentrations were determined spectrometric with a *Nanodrop ND-100* spectrophotometer by their absorbance at 260 nm. Duplexes of TPE-modified DNA were prepared by heating the chromophore-modified strands in the presence of 1.0 eq. unmodified or with atto dye modified complementary strands to 90 °C, hold this temperature for 5 min and then slowly cooling to RT.

Table S1. MS analyses and extinction coefficients of the used DNA strands.

DNA	[M] ⁺ calc. [g/mol]	[M] ⁺ found [g/mol]	ε ₂₆₀ [mM ⁻¹ cm ⁻¹]
L-A ₂₀	6199.2	6197.4	277.2
L-T ₂₀	6019.0	6020.4	158.4
TPE1a	6404.1	6403.3	203.6
TPE1b	6404.1	6405.5	203.6
TPE2a	6744.3	6749.8	219.1
TPE2b	6744.3	6745.8	219.1
1D			23.4

Confidence
Data: HU_dA20_1_HPA_0001.B6[c] 12 Oct 2020 9:35 Cal: 6kDa_HPA_15082019 26 Feb 2020 9:56
Shimadzu Biotech Axima Confidence 2.9.3.20110624: Mode 2019_Linear_neg_new, Power: 116, Blanked, P.Ext. @ 1700 (bin

%Int. 0.6 mV[sum= 288 mV] Profiles 1-475 Smooth Av 30 -Baseline 400

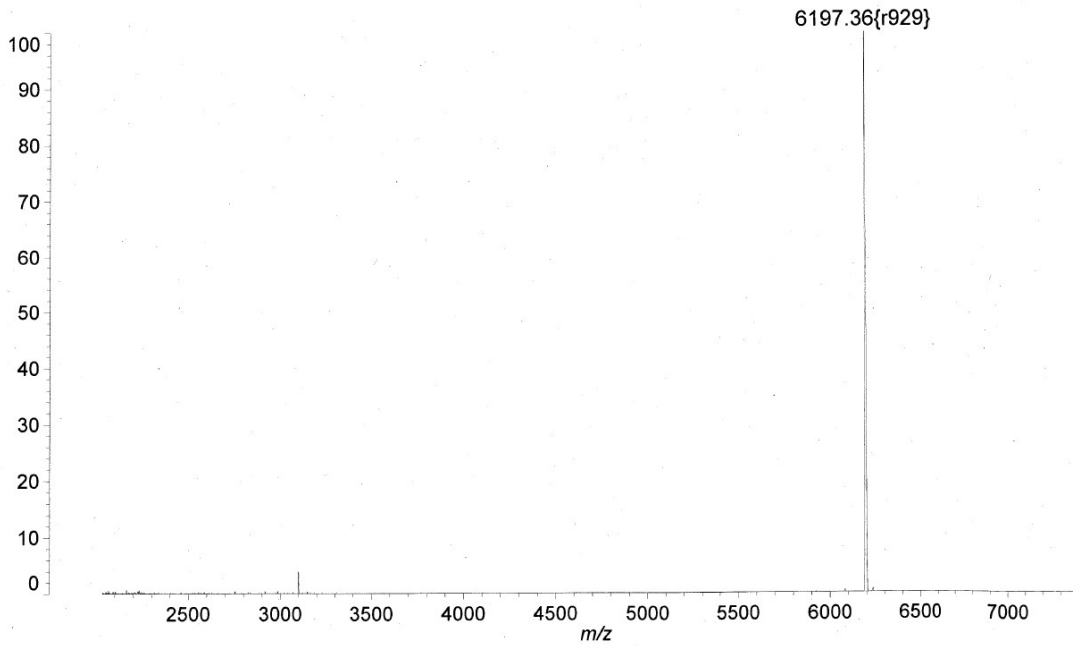


Figure S13. MS (MALDI) analysis of L-A₂₀.

Confidence
Data: SM-T20_c_HPA_0001.K4[c] 6 Jul 2020 11:42 Cal: 2-4kDa_HPA_08012019 26 Aug 2019 10:39
Shimadzu Biotech Axima Confidence 2.9.3.20110624: Mode 2019_Linear_neg_new, Power: 127, Blanked, P.Ext. @ 6019 (bin

%Int. 13 mV[sum= 934 mV] Profiles 1-73 Smooth Av 30 -Baseline 400

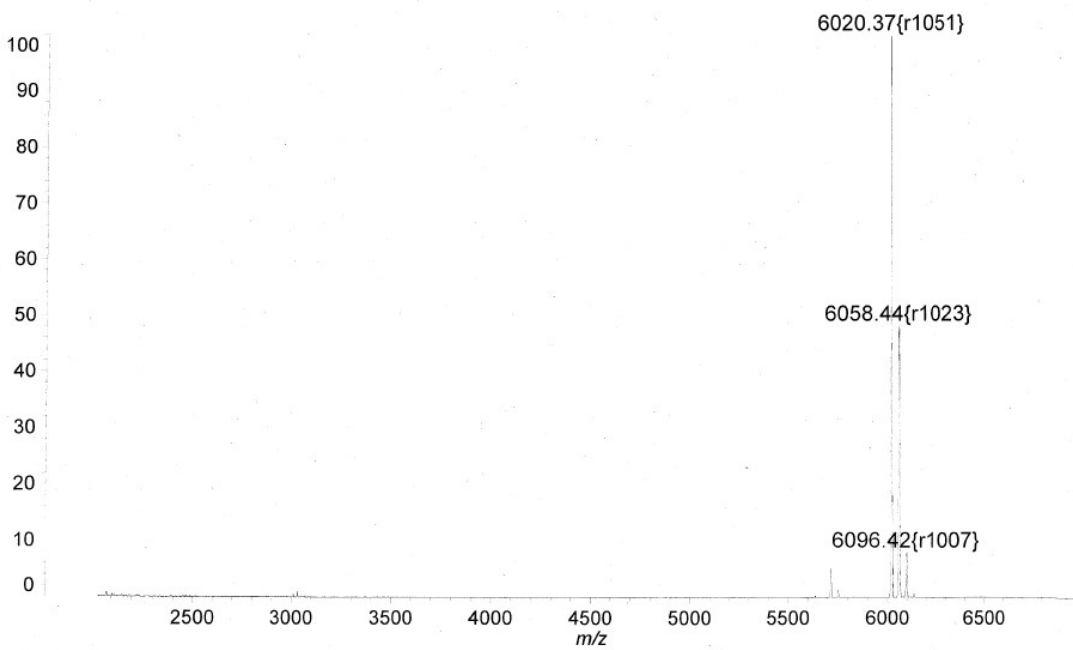


Figure S14. MS (MALDI) analysis of L-T₂₀.

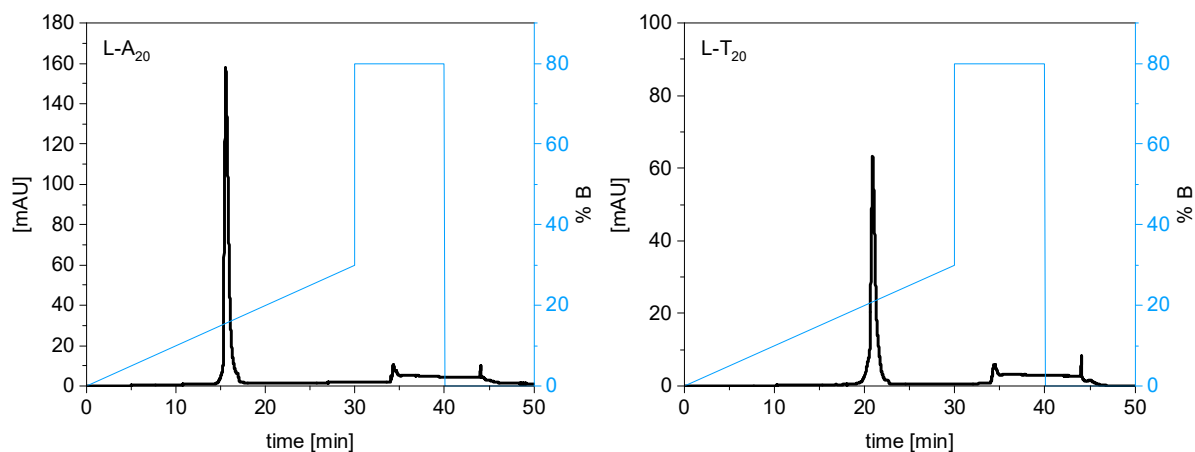


Figure S15. HPLC analyses of L-A₂₀ (left) and L-T₂₀ (right).

Confidence
 Data: HU_TPE_1_HPA_0001.C2[c] 9 Feb 2021 10:18 Cal: 6_7KDa_linneg_20 Jan 2021 10:26
 Shimadzu Biotech Axima Confidence 2.9.3.20110624: Mode 2019_Linear_neg_new, Power: 119, Blanked, P.Ext. @ 6000 (bir

%Int. 12 mV[sum= 930 mV] Profiles 1-80 Smooth Av 30 -Baseline 400

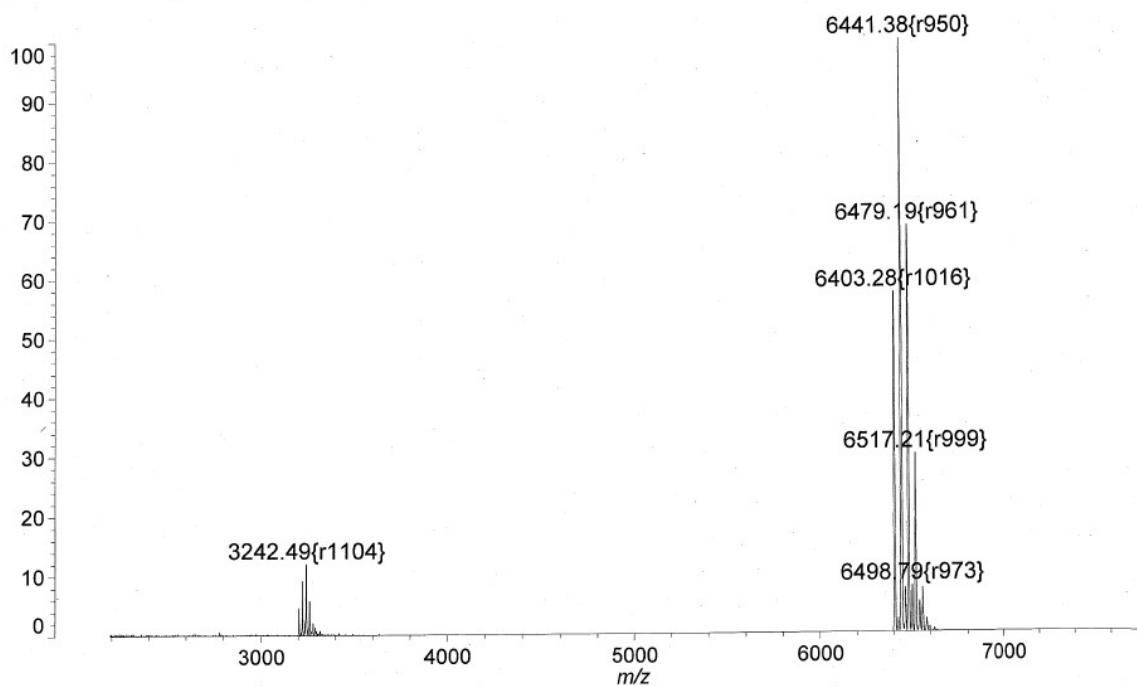


Figure S16. MS (MALDI) analysis of TPE1a.

Confidence
Data: HU_DNA_TPE_9_17_HPA_0001.G12[c] 22 Mar 2021 10:09 Cal: 6kDa_HPA_15082019 20 Jan 2021 11:25
Shimadzu Biotech Axima Confidence 2.9.3.20110624: Mode 2019_Linear_neg_new, Power: 113, Blanked, P.Ext. @ 6200 (bir

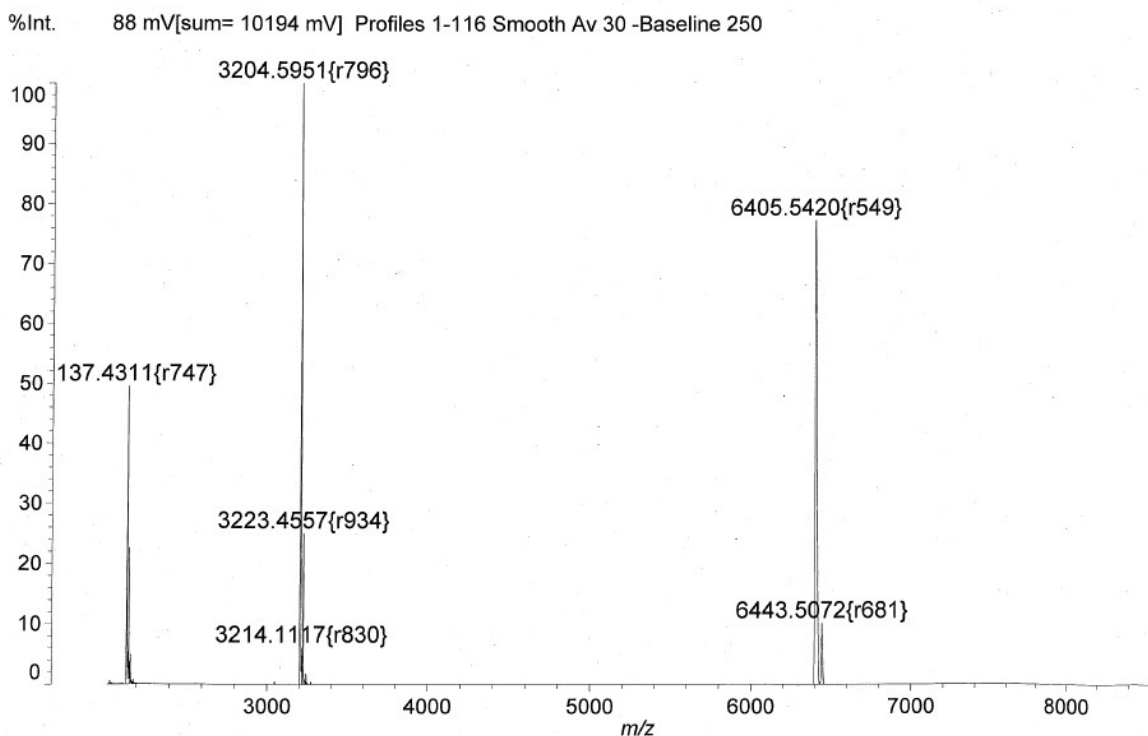


Figure S17. MS (MALDI) analysis of TPE1b.

Confidence
Data: HU_TPE4_42_HPA_0001.I17[c] 1 Feb 2021 11:46 Cal: 6_7KDa_linneg_20 Jan 2021 10:26
Shimadzu Biotech Axima Confidence 2.9.3.20110624: Mode 2019_Linear_neg_new, Power: 119, Blanked, P.Ext. @ 6000 (bir

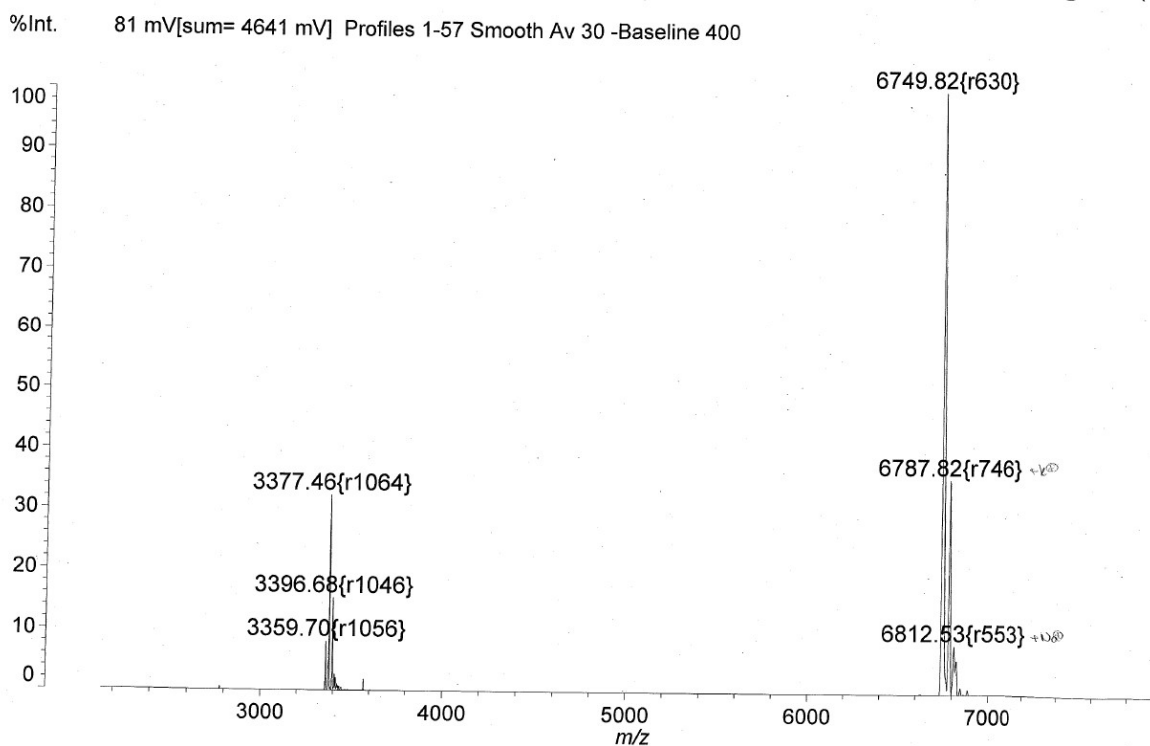


Figure S18. MS (MALDI) analysis of TPE2a.

Confidence
Data: HU_TPE5_39_HPA_0001.K3[c] 1 Feb 2021 11:51 Cal: 6_7KDa_linneg_20 Jan 2021 10:26
Shimadzu Biotech Axima Confidence 2.9.3.20110624: Mode 2019_Linear_neg_new, Power: 116, Blanked, P.Ext. @ 6000 (bir

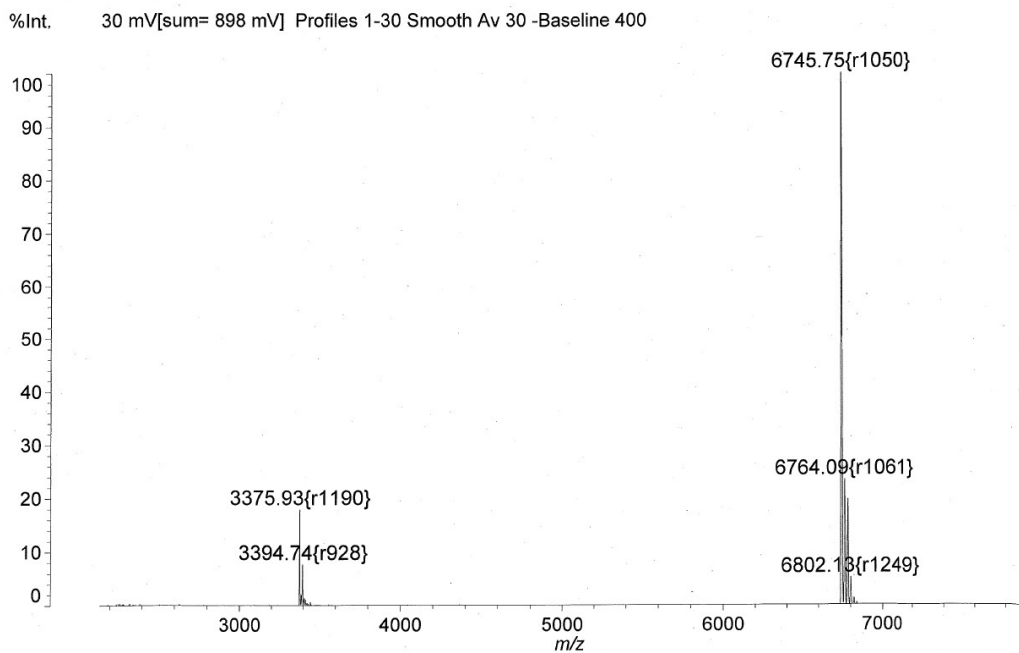


Figure S19. MS (MALDI) analysis of TPE2b.

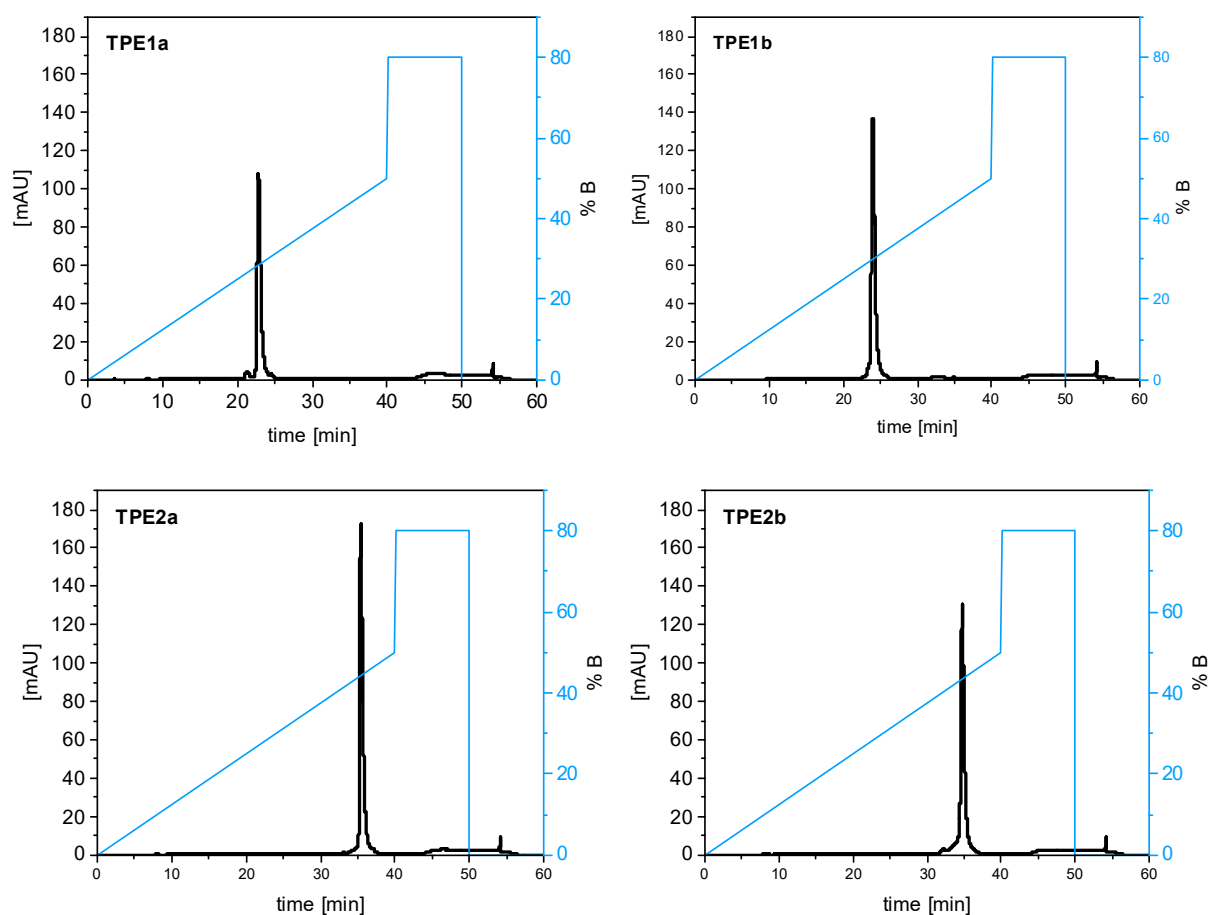


Figure S20. HPLC analysis of TPE1a, TPE1b, TPE2a and TPE2b.

4. Additional optical spectroscopy

Non-templated supramolecular assemblies

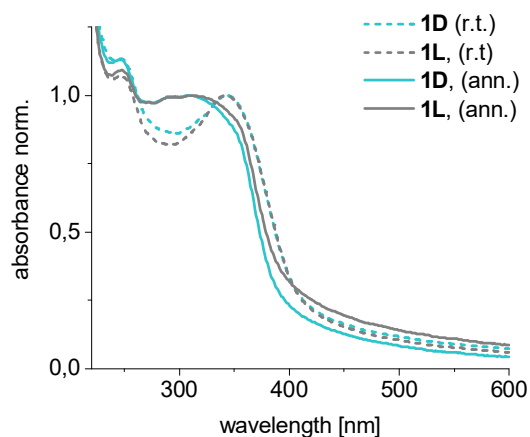


Figure S21. UV/Vis absorbance of the assemblies prepared with **1D** and **1L** in the absence of any DNA template (H_2O , 0.9% **THF**) either at room temperature (dashed) or by annealing after heating to 90 °C for 5 min (solid).

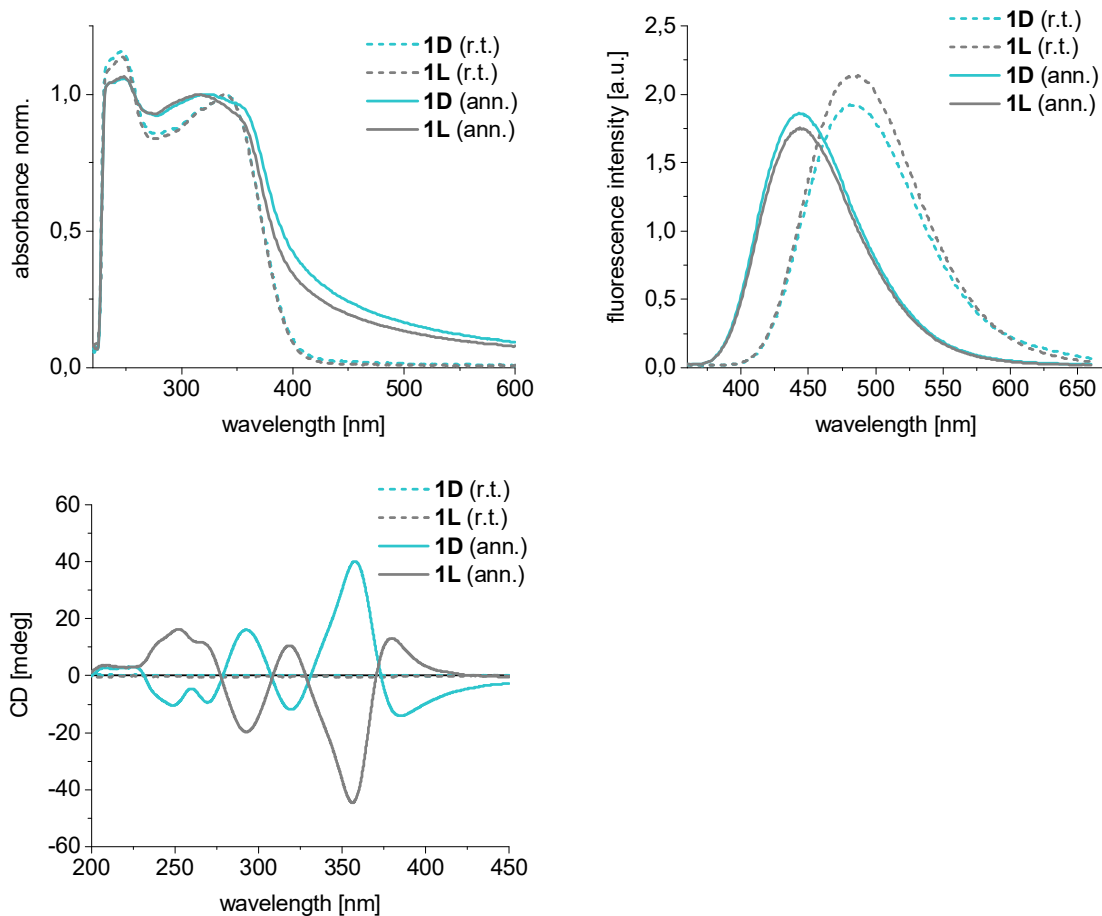


Figure S22. UV/Vis absorbance, emission and circular dichroism of the assemblies prepared with **1D** and **1L** in the absence of any DNA template (H_2O , 0.9% **DMSO**) either at room temperature (dashed) or by annealing after heating to 90 °C for 5 min (solid).

DNA-templated supramolecular architectures

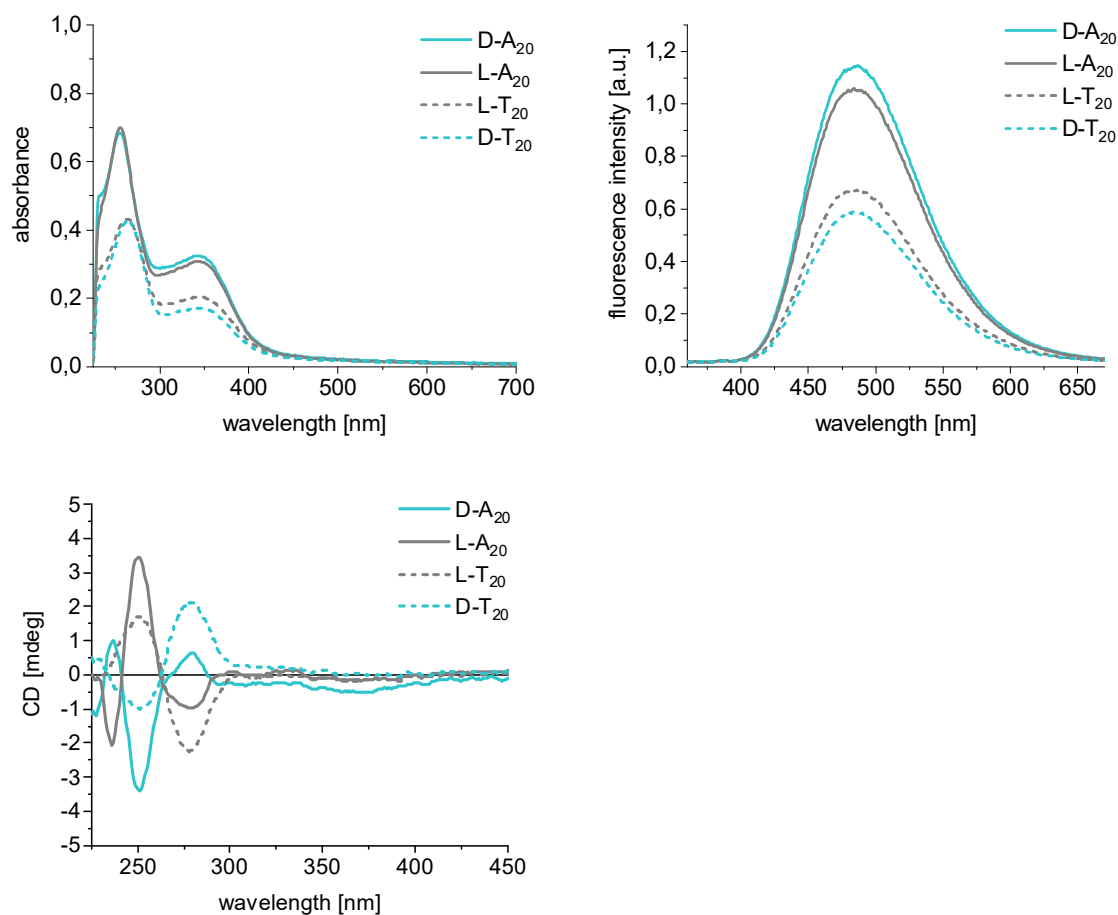


Figure S23. UV/Vis absorbance, fluorescence and circular dichroism of the DNA-templated assemblies of **1L** (1.25 μM DNA, 37.5 μM **1L**, 250 mM NaCl, 0.9 % DMSO, 1h incubated at r. t., 3 min at 16 000 g centrifuged, $\lambda_{\text{exc}} = 341$ nm).

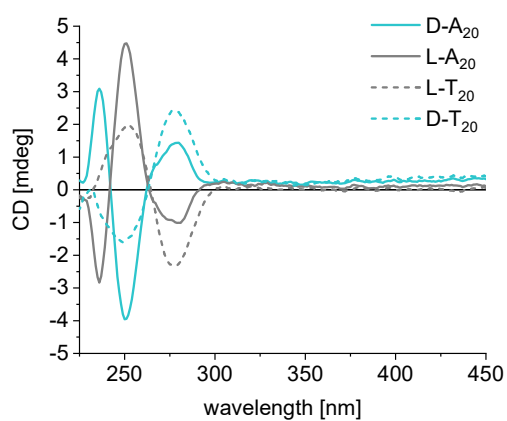


Figure S24. Circular dichroism of the DNA-templates in the absence of **1D** and **1L** (1.25 μM DNA, 250 mM NaCl, 0.9 % DMSO, 1h incubated at r. t.).

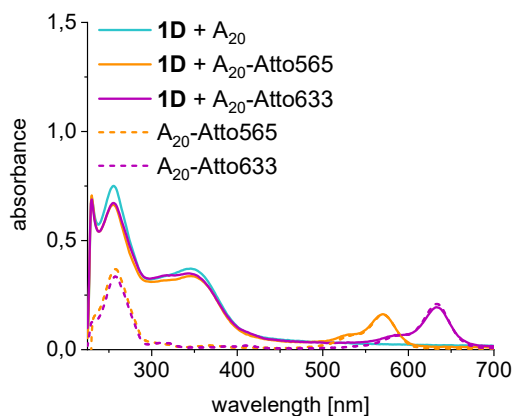


Figure S25. UV/Vis absorbance of the DNA-templated (with and without atto dye modification) assemblies of **1D** (1.25 μM DNA, 37.5 μM **1D**, 250 mM NaCl, 0.9 % DMSO, 1h incubated at r. t., 3 min at 16 000 g centrifuged).

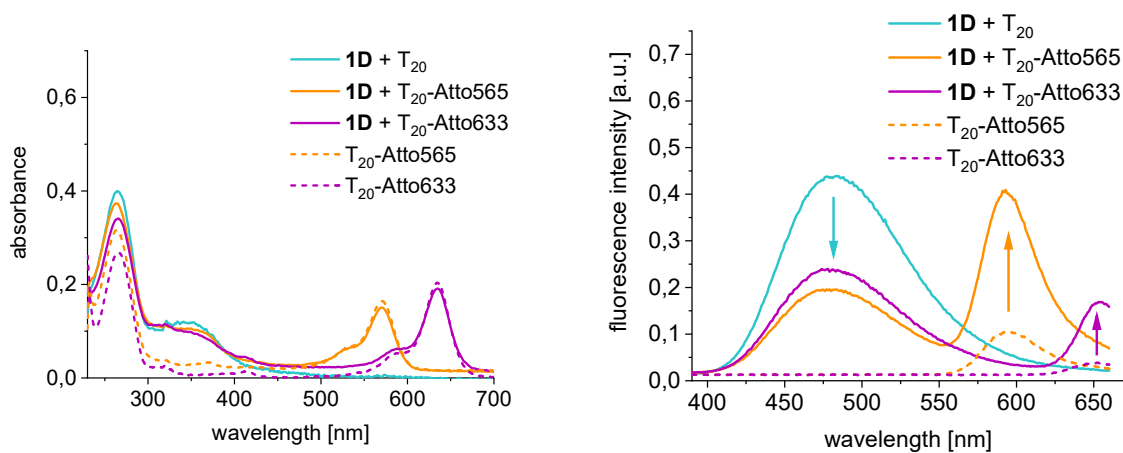


Figure S26. UV/Vis absorbance and fluorescence of the DNA-templated (with and without atto dye modification) assemblies of **1D** (1.25 μM DNA, 37.5 μM **1D**, 250 mM NaCl, 0.9 % DMSO, 1h incubated at r. t., 3 min at 16 000 g centrifuged).

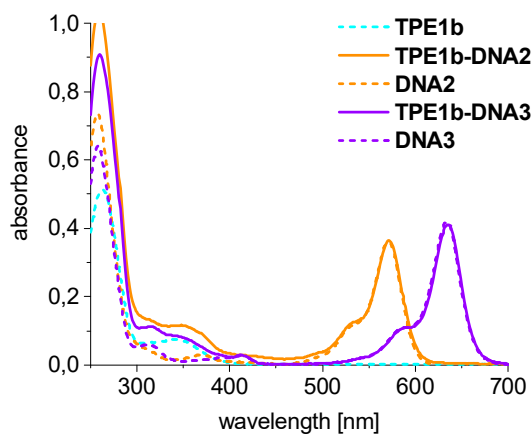


Figure S27. UV/Vis absorbance of the TPE-modified DNA single strand **TPE1b** and hybrids with atto dye modified counter strands **DNA2** and **DNA3** ($c(\text{TPE1b}) = c(\text{cs}) = 2.5 \mu\text{M}$, 250 mM NaCl, 10 mM NaPi buffer).

5. References

- [1] J. Wang, J. Mei, E. Zhao, Z. Song, A. Qin, J. Z. Sun, B. Z. Tang, *Macromolecules* **2012**, *45*, 7692-7703.