

Electronic Supplementary Information

DNA quadruplexes as molecular scaffolds for controlled assembly of fluorogens with aggregation-induced emission

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Supporting Figures

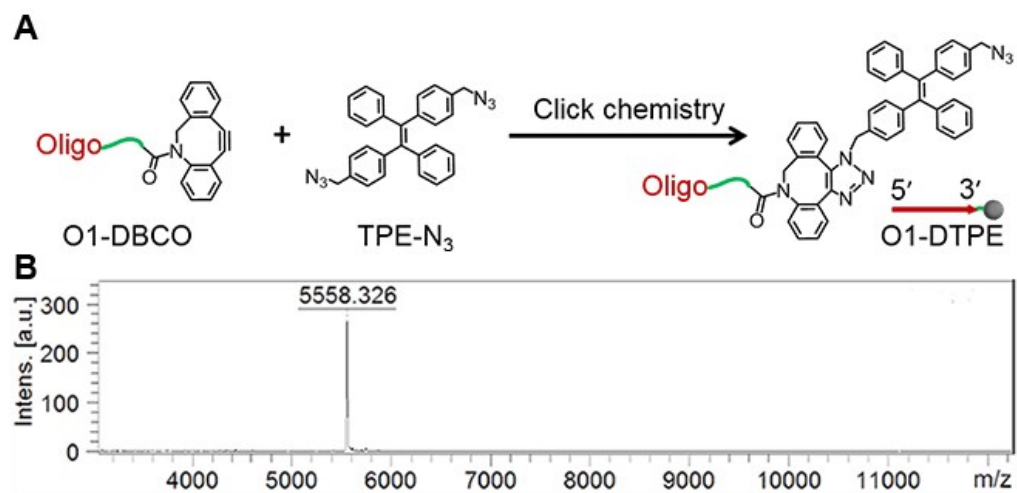


Fig. S1 (A) Synthetic scheme of O1-DTPE, (B) TOF-MS of O1-DTPE.

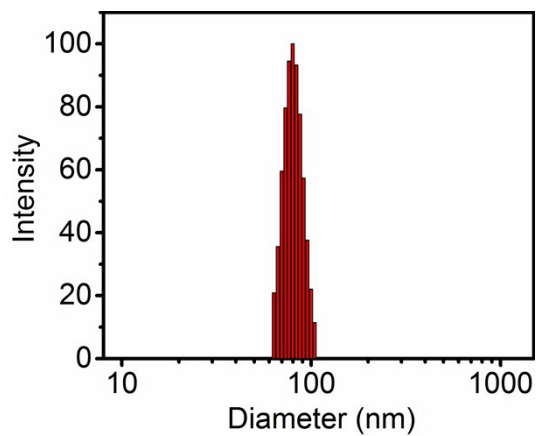


Fig. S2 Dynamic light scattering analysis of 1.0 μM TPE-N₃ in DMSO/H₂O (v/v, 1/399). No signal was observed for O1-DTPE in H₂O at 25 °C.

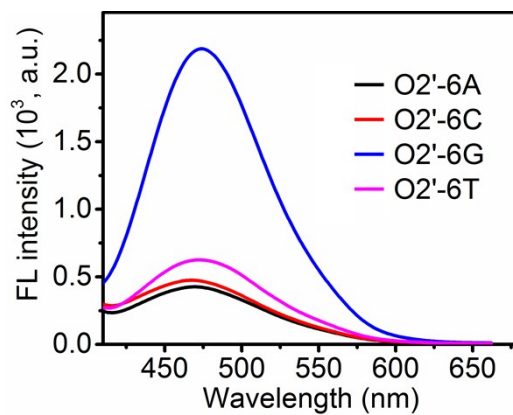


Fig. S3 Fluorescent spectra of 1.0 μM O2-DTPE in presence of indicated strands.

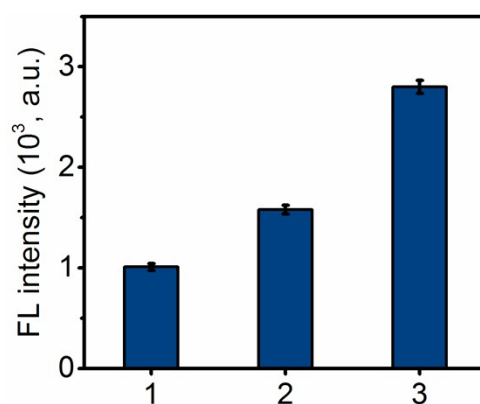


Fig. S4 Fluorescent intensity of mixture of 1.0 μM O1'-6G with (1) 0.25 μM , (2) 0.5 μM and (3) 1.0 μM O1-DTPE.

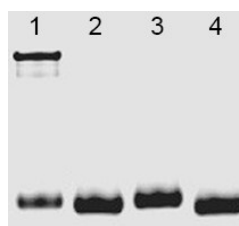


Fig. S5 PAGE results of 10 μM O1-DTPE hybridized with (1) O1'-6G, (2) O1'-6C, (3) O1'-6T and (4) O1'-6A.

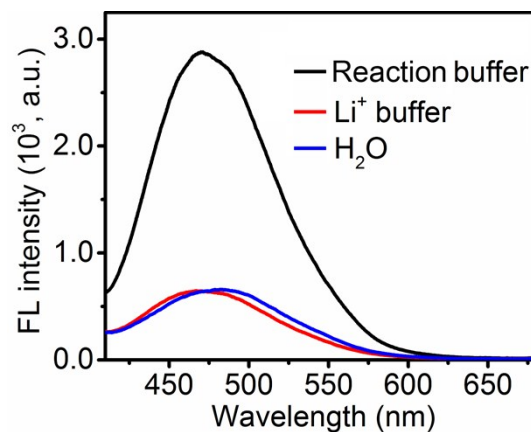


Fig. S6 Fluorescent spectra of 1.0 μM O1-DTPE hybridized with O1'-6G (1.0 μM) in TE reaction buffer (10 mM TE, pH 7.4, 100 mM NaCl, 10 mM MgCl_2), Li^+ buffer, and H_2O .

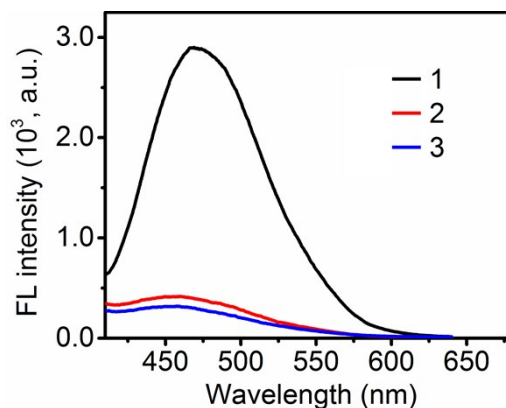


Fig. S7 Fluorescent spectra of 1.0 μM O1-DTPE mixed with (1) O1'-6G, (2) O1-6G, and (3) 15T-6G.

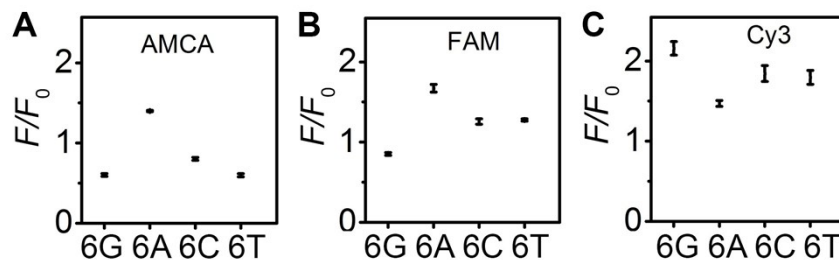


Fig. S8 Ratios of fluorescent peak intensity of 1.0 μM (A) O1-AMCA, (B) O1-FAM, and (C) O1-Cy3 after to before (F/F_0) hybridization with 1.0 μM O1' with 6G, 6A, 6C, and 6T.

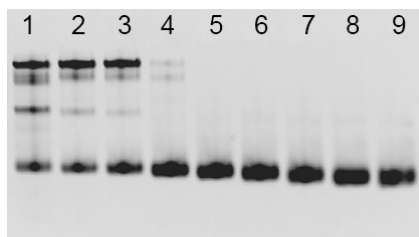


Fig. S9 PAGE results of O1-DTPE hybridized with (1) O1'-8G, (2) O1'-7G, (3) O1'-6G, (4) O1'-5G, (5) O1'-4G, (6), O1'-3G, (7) O1'-2G, (8) O1'-1G, and (9) O1'.

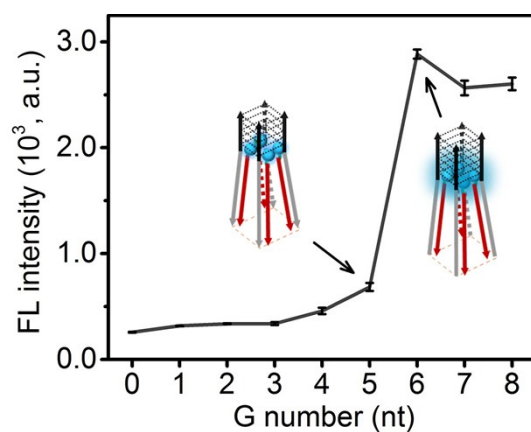


Fig. S10 Fluorescent intensity (FL) of 1.0 μM O1''-DTPE hybridized with O1' bearing 0 to 8 Gs at 3' end.

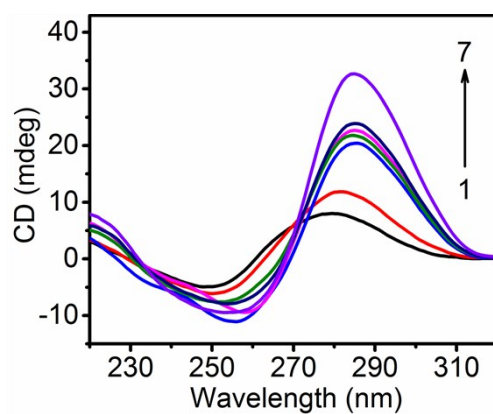


Fig. S11 CD spectra of 20 μM (1) O1, (2) O1'-2C, (3) O1'-4C, (4) O1'-8C, (5) O1'-6C, (6) O1'-10C and (7) O1'-12C in pH 4.8 TAE reaction buffer (10 mM Tris-acetate, 100 mM NaCl, 10 mM MgCl_2 , 1 mM sodium EDTA).

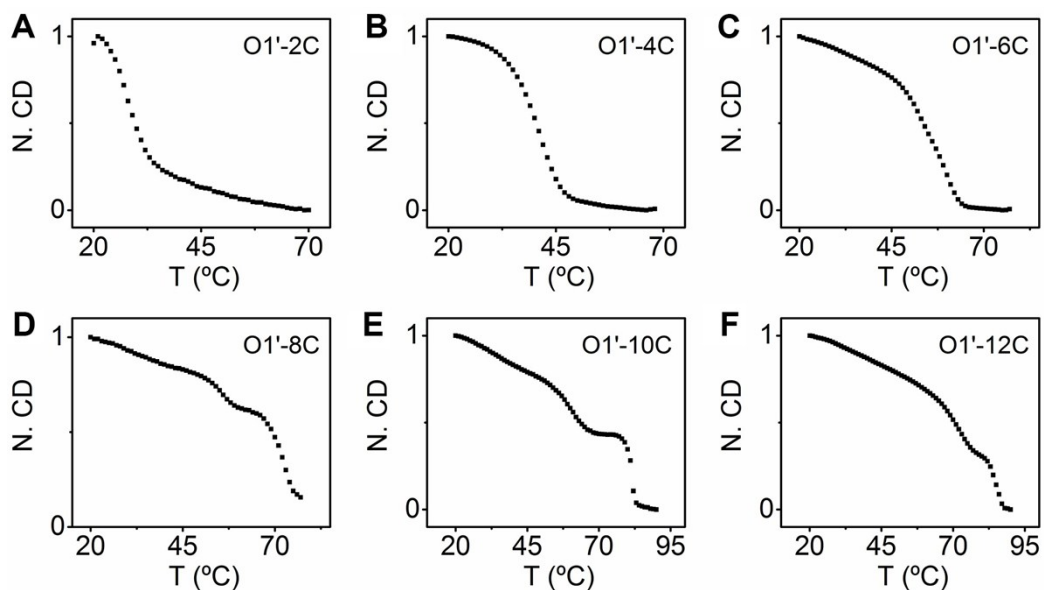


Fig. S12 Normalized CD (N. CD) melting curves monitored at 285 nm of 20.0 μM O1' bearing different numbers of Cs (O1'- $n\text{C}$, $n = 2$ to 12); temperature was increased at a rate of 2.0 $^{\circ}\text{C}/\text{min}$. Data are normalized to the highest CD value at 285 nm.

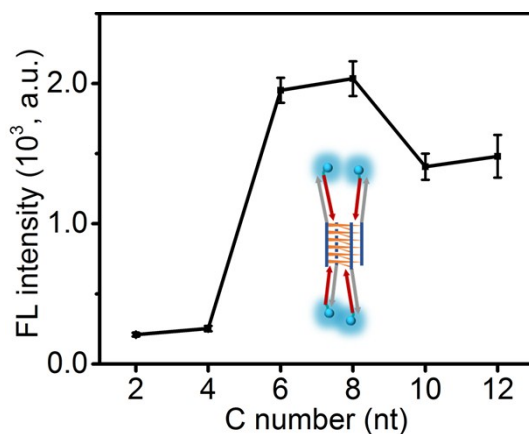


Fig. S13 Fluorescence intensity of 1.0 μM O1''-DTPE hybridized with O1'- $n\text{C}$ ($n = 0$ to 12) in pH 4.8 TAE reaction buffer.

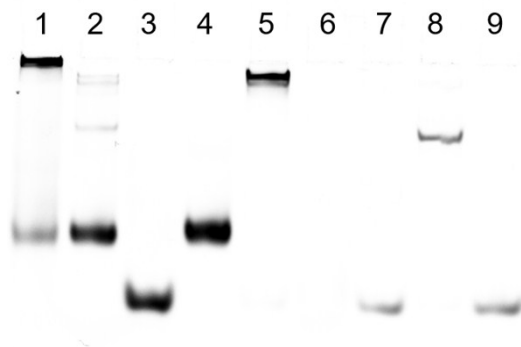


Fig. S14 PAGE results of (1) target DNA + TP'-G4-5bp + H-DNA + O1-DTPE, (2) TP'-G4-5bp + H-DNA + O1-DTPE, (3) TP'-G4-5bp + H-DNA, (4) H-DNA + O1-DTPE, (5) TP'-G4-5bp + O1-DTPE, (6) O1-DTPE, (7) H-DNA, (8) TP'-G4-5bp, (9) target DNA.

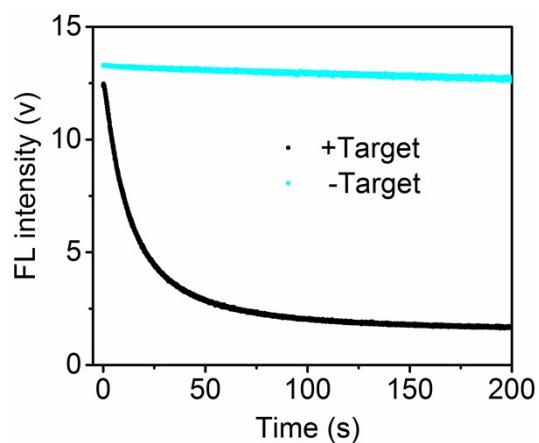


Fig. S15 Stopped-flow fluorescence over time for the mixtures of 1.0 μ M O1-FAM, H-DNA, and TP'-G4-TAMRA in presence and absence of target DNA.

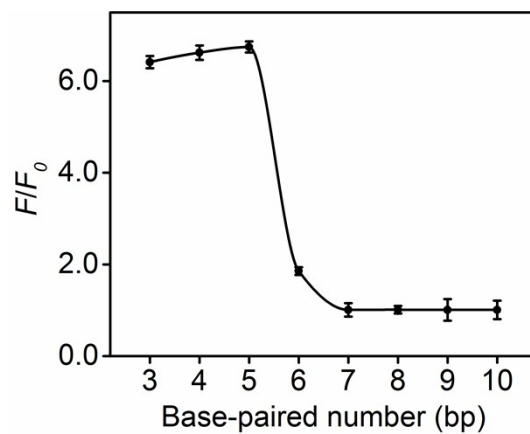


Fig. S16 Fluorescence intensity of a mixture of 1.0 μM H-DNA, target DNA, O1-DTPE, and 16-nt limbs of TP'-G4 with different numbers of base pairs (TP'-G4- n bp, $n = 3$ to 10) in pH 7.4 reaction buffer.

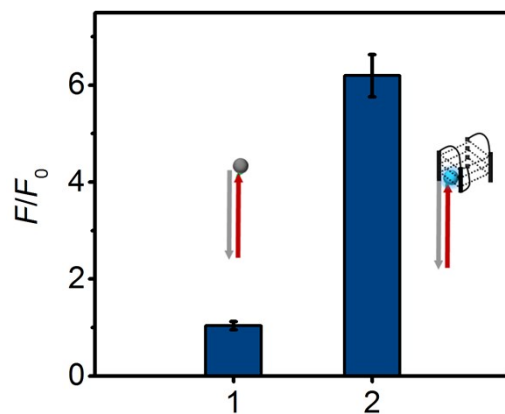


Fig. S17 Ratios of fluorescent peak intensity of 1.0 μM O1-DTPE after to before (F/F_0) hybridization with (1) O1' and (2) O1'-mG4 in TE reaction buffer.

Supporting Tables

Table. S1 Oligonucleotide sequences used in the experiments.

Name	Sequences (from 5' to 3' end)
O1-DBCO	AGTTGGAGACGTAAG-(CH ₂) ₆ NHCO(CH ₂) ₄ -DBCO
O2-DBCO	CCTACGTCTCCAACCTAAGTACGGCCCTCATTCAATACCCTACG-(CH ₂) ₆ NHCO(CH ₂) ₄ -DBCO
O1''-DBCO	DBCO-(CH ₂) ₄ CONH(CH ₂) ₆ -AGTTGGAGACGTAAG
O1'	CTTACGTCTCCAACCT
O1'-1G	GCTTACGTCTCCAACCT
O1'-2G	GGCTTACGTCTCCAACCT
O1'-3G	GGGCTTACGTCTCCAACCT
O1'-4G	GGGGCTTACGTCTCCAACCT
O1'-5G	GGGGGCTTACGTCTCCAACCT
O1'-6G	GGGGGGCTTACGTCTCCAACCT
O1'-7G	GGGGGGGCTTACGTCTCCAACCT
O1'-8G	GGGGGGGGCTTACGTCTCCAACCT
O1'-6A	AAAAAACTTACGTCTCCAACCT
O1'-6T	TTTTTTCTTACGTCTCCAACCT
O1'-2C	CCCTTACGTCTCCAACCT
O1'-4C	CCCCCTTACGTCTCCAACCT
O1'-6C	CCCCCCCTTACGTCTCCAACCT
O1'-8C	CCCCCCCCCTTACGTCTCCAACCT
O1'-10C	CCCCCCCCCCTTACGTCTCCAACCT
O1'-12C	CCCCCCCCCCCCTTACGTCTCCAACCT
O1'-5A-6G	GGGGGGAAAAACTTACGTCTCCAACCT
O1'-10A-6G	GGGGGGAAAAAAAAAACTTACGTCTCCAACCT
O1'-15A-6G	GGGGGGAAAAAAAAAAAAAAAAAACTTACGTCTCCAACCT
O1'-20A-6G	GGGGGGAAAAAAAAAAAAAAAAAAAAAAAAAACTTACGTCTCCAACCT
O1'-5T-6G	GGGGGGTTTTTCTTACGTCTCCAACCT
O1'-10T-6G	GGGGGGTTTTTTTTTCTTACGTCTCCAACCT
O1'-15T-6G	GGGGGGTTTTTTTTTTTTTTTCTTACGTCTCCAACCT

01'-20T-6G GGGGGGTTTTTTTTTTTTTTTTTTTTTTTCTTACGTCTCCA
 01'-6G-5A GGGGGGCTTACGTCTCCA
 01'-6G-10A GGGGGGCTTACGTCTCCA
 01'-6G-15A GGGGGGCTTACGTCTCCA
 01'-6G-20A GGGGGGCTTACGTCTCCA
 01'-6G-5T GGGGGGCTTACGTCTCCA
 01'-6G-10T GGGGGGCTTACGTCTCCA
 01'-6G-15T GGGGGGCTTACGTCTCCA
 01'-6G-20T GGGGGGCTTACGTCTCCA
 02'-6A AAAAAACGTAGGGTATTGAATGAGGGCCGTAAGTTAGTTGGAGACGT
 AGG
 02'-6C CCCCCCGTAGGGTATTGAATGAGGGCCGTAAGTTAGTTGGAGACGTA
 GG
 02'-6T TTTTTCGTAGGGTATTGAATGAGGGCCGTAAGTTAGTTGGAGACGTA
 GG
 02'-6G GGGGGGCGTAGGGTATTGAATGAGGGCCGTAAGTTAGTTGGAGACGT
 AGG
 01-6G GGGGGGAGTTGGAGACGTAAG
 15T-6G GGGGGGTTTTTTTTTTTTTTTTTTT
 01-AMCA AGTTGGAGACGTAAG-AMCA
 01-FAM AGTTGGAGACGTAAG-FAM
 01-Cy3 AGTTGGAGACGTAAG-Cy3
 1G-01' CTTACGTCTCCA
 2G-01' CTTACGTCTCCA
 3G-01' CTTACGTCTCCA
 4G-01' CTTACGTCTCCA
 5G-01' CTTACGTCTCCA
 6G-01' CTTACGTCTCCA
 7G-01' CTTACGTCTCCA
 8G-01' CTTACGTCTCCA
 01'-4C4T4C CCCCTTTCCCCTCTTACGTCTCCA
 TP(10)-G4 GGGGGGGTCTCCA
 TP'-G4-3bp GGGGGGTCAATACTACCTCA
 TP'-G4-4bp GGGGGGTCAATACTACCTCA
 TP'-G4-5bp GGGGGGTCAACATACTACCTCA
 TP'-G4-6bp GGGGGGTCAACCATACTACCTCA

TP'-G4-7bp GGGGGGTCAACCTATACTACCTCA
 TP'-G4-8bp GGGGGGTCAACCTCATACTACCTCA
 TP'-G4-9bp GGGGGGTCAACCTCTATACTACCTCA
 TP'-G4-10bp GGGGGGTCAACCTCTGATACTACCTCA
 H-DNA AACTATACAAGCTCTGCATTC
 Target TGAGGTAGTATCTTGTATAGTT
 TP'-G4-TAMARA GGGGGGT(TAMRA)CAACATACTACCTCA
 O1'-mG4 GGGTGGGTGGGTGGGTCTTACGTCTCCAAC

The blue bases are designed to form the nanostructure core or equivalent region in controls. FAM is 6-carboxyfluorescein, AMCA is aminomethylcoumarin, and Cy3 is cyanine 3.

Table. S2 CD melting temperatures (T_m s) of O1' bearing different numbers of Gs at the 5' end.

DNA strands	O1'	O1'-1G	O1'-2G	O1'-3G	O1'-4G	O1'-5G	O1'-6G	O1'-7G	O1'-8G
$T_m / ^\circ\text{C}$	-	-	-	< 20	44	49	53	54	54