

Engineering a Cell-surface Aptamer Circuit for Targeted and Amplified Photodynamic Cancer Therapy

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Supporting information

Gel electrophoresis to demonstrate the catalytic effect of C sequence. Polyacrylamide gel electrophoresis was performed on a 10% native gel in TBE buffer (89 mM Tris-HCl, 89 mM boric acid, 2 mM EDTA) with 5 mM MgCl₂ and run for 60 min at 100 V. Gels were then stained using StainsALL to image the positions of DNA strands. Five samples were prepared as follows: lane 1, purified A₁; lane 2, purified A₂; lane 3, mixture of A₁ and A₂ only; lane 4, preannealed A₁ and A₂, which will form the A₁₂ duplex; lane 5, the mixture of A₁, A₂ and C (0.5× concentration of A₁) preincubated for 30 min.

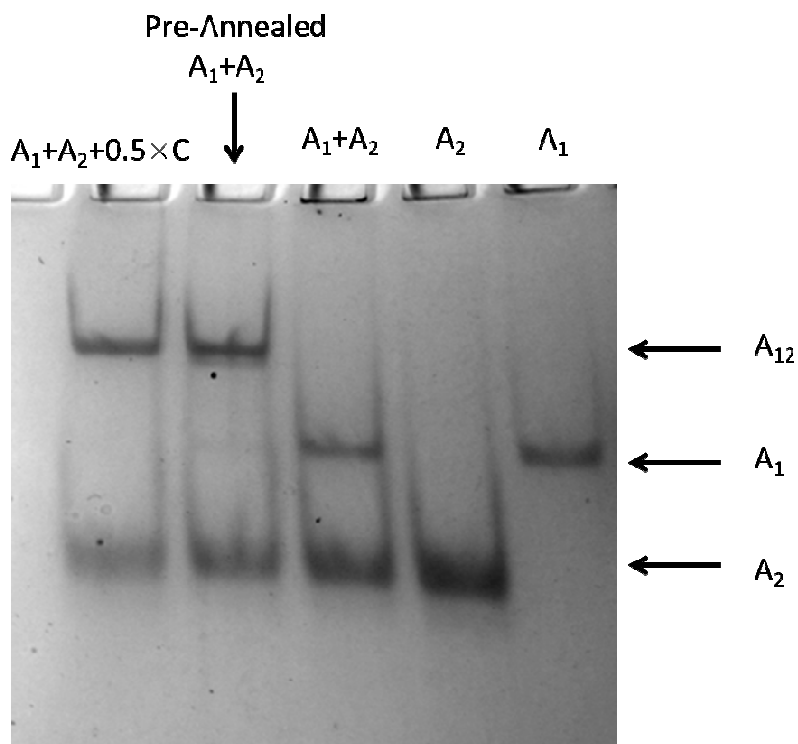


Figure S1. Image of the PAGE gel proving the catalytic effect of C sequence.

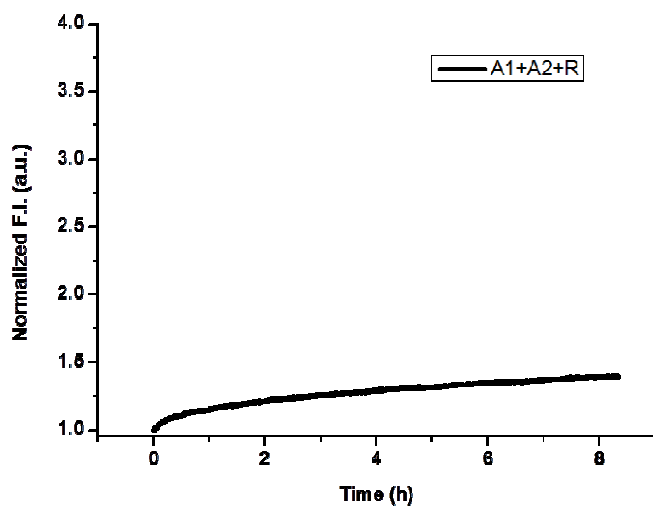


Figure S2. Fluorescence kinetics describing the leakage reaction of A_1 , A_2 and R_{12} in the Fluor buffer. The data were normalized to the initial fluorescence intensity of the circuit.

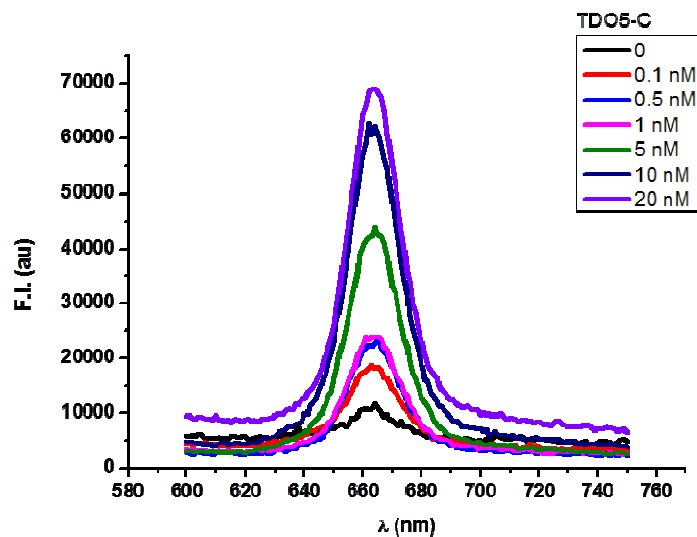


Figure S3. The fluorescence spectra of mixtures containing 100 nM A_1 , 100 nM A_2 and 150 nM Ce6-modified R_{12} with different concentrations of TDO5-C in buffer.

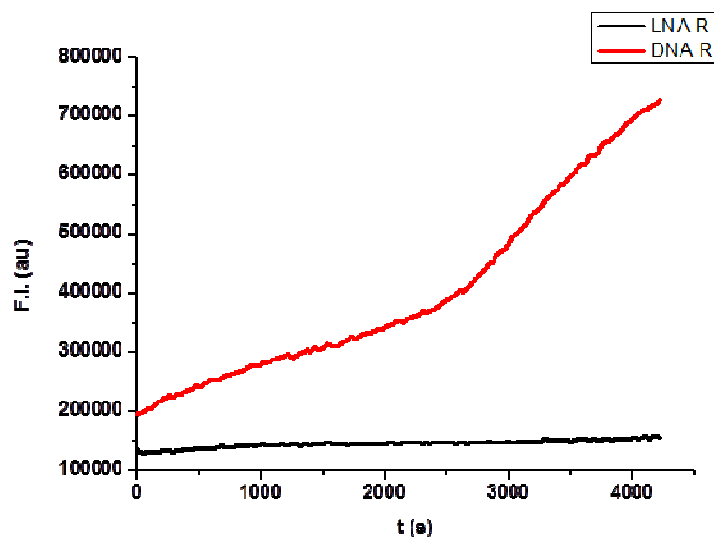


Figure S4. The fluorescence kinetics of LNA-DNA hybrid R_{12} and pure DNA R_{12} with 100 K Ramos cells in washing buffer.

Sequence Name	Sequence
C (c*b*a*)	CGACATCT_AACCTAGC_TCACTGAC
A₁ (abcd*c*b*e*)	GTCAGTGA_GCTAGGTT_AGATGTCG_CCATGTGTAGA_CGACAT C_TAACCTAGC_ACTTGTCATAGAGCAC
A₂ (cdc*b*d*)	AGATGTCG_TCTACACATGG_CGACATCTAACCTAGC_CCATGTG TAGA
R₁ (eb)	Ce6 (FAM) GTGCTCTATGACAAGT_GCTAGGTT
R₂ (b)	ACTTGTCATAGAGCAC BHQ2 (DABCYL)
TDO5	AACACCGTGGAGGATAGTTCGGTGGCTGTTTCAGGGTCTCCTCCC GGTG
TDO5-C	CGACATCTAACCTAGCTCACTGAC_TTTTTTTTTTTTTTTTTT_AAC ACCGTGG AGGATAGTTCGGTGGCTGTTTCAGGGTCTCC TCCC GTG

Table S1. Sequence of oligonucleotides used in this work. Domains are separated by underscores. LNA bases are indicated by bold and underscores.