Supporting Information

Aptamer Display on Diverse DNA Polyhedron Supports

Simon Chi-Chin Shiu^{1,†}, Lewis A. Fraser^{1,†}, Yifan Ding¹ and Julian A. Tanner^{1,*}

- ¹ School of Biomedical Sciences, Li Ka Shing Faculty of Medicine, The University of Hong Kong, Pokfulam, Hong Kong SAR China; <u>simon.chichin.shiu@gmail.com</u> (S.C.C.S.); <u>lewis-fraser@hku.hk</u> (L.A.F.); <u>yifand16@gmail.com</u> (Y.D.)
- * Correspondence: jatanner@hku.hk
- [†] These authors contributed equally to this work

Label	Sequence (5'-3')	Strands required for
		nanostructures
Та	biotin-	Tetrahedron 1, 2 and 3
	TTTTGCGGCTGGAGCCATGTCATCAGGGGGCACGTCTCTAGGACGCGGCCGGGTCT	
T1b	biotin-TTCATGGCTCCAGCCGCAATGACGGGACCGTGTGCCTGAGC	
T1c	biotin-TTGAGACGTGCCCCTGATGTGCTCAGGCACTGCAGGGCAAA	Tetrahedron 1
T1d	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTCGGTCCCGTCTAGACCCGGC	
	CGCGTCCTTTTTGCCCTGC	
T2b	biotin-	
	TTCATGGCTCCAGCCGCAATTATCTAGCGATCTCACCTCGTCAAGGGTAAGCCCG	Tetrahedron 2
T2c	biotin-	
	TTGAGACGTGCCCCTGATGTCGGGCTTACCCTTGACGTGTGGTAATTGTTGTGTA	
T2d	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTGGTGAGATCGCTAGATATAG	
	ACCCGGCCGCGTCCTTTACACAACAATTACCAC	
T3b	biotin-	
	TTCATGGCTCCAGCCGCAATTGATCATTGAGTGTCGGTGGCGATGTCTCCCCGATAAA	
	GCGTATGTTATAAGTCT	
T3c	biotin-	Tetrahedron 3
	TTGAGACGTGCCCCTGATGTAGACTTATAACATACGCTTTATCGGGGGTCCGGAGCCA	
	TCACACCTGGATCAGGAC	
T3d	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTGACATCGCCACCGACACTCA	
	ATGATCATAGACCCGGCCGCGTCCTTGTCCTGATCCAGGTGTGATGGCTCCGG	
C2a	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTTGGGGCCGGGCGCGCCCCGG	Aptamer with single-
	TT-biotin	stranded and double-
		stranded spacer
C2b	TTCCGGGGCGCGCCCCA	Aptamer with double-
		stranded spacer
P1a	biotin-TTATGTAACCGTGTCACTCCAGGTGTACTTCCTAAAATCGCGCGACCAG	
P1b	biotin-TTCACGGTTACATTCTGTGTTGGCACGGCGTCGTCCGGGTGGCGGTT	
P1c	biotin-TTCACCTGGAGTGTAACCGCCACCCGGACGTCGTTCGTCAAATAGTG	Square-based
P1d	biotin-TTTTTAGGAAGTTCACTATTTGACGAACGTGCTATGCCGCGCGCCT	pyramid
P1e	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTCGCCGTGCCAACACAGTCTG	
	GTCGCGCGTAGGCGCGCGCGCATAGC	
P2a	biotin-TTCGGCCGATCTGGCATAGGCTTCCCCAACGTTATCTCGCGTGGCGTACGG	
P2b	biotin-TTGATCGGCCGTTCCGCGCGTCCTGCCCTAGACGCCCCTGGGCCC	
P2c	biotin-TTGCCTATGCCTGGGCCCAGGGGGCGTCTTGGAGTCAACAGGTCGG	Pentagon-based
P2d	biotin-TTCGTTGGGGATCCGACCTGTTGACTCCTCCCGTCGACCAAAGAT	pyramid
P2e	biotin-TTCGCGAGATATATCTTTGGTCGACGGGTCTCAGCCCTGACCCCA	
P2f	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTGGGCAGGACGCGCGGATCCG	
	TACGCCTTGGGGTCAGGGCTGAG	
Pr1	GCAACAGAGCCCTATTCTGTCTCAGTCCACCGGGTTAGGAAACGCGAGTCAGT	
Pr2	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTCCCTAACGCCTGCGTGAATA	

	GGGCTCTGTTGC	Prism
Pr3	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTGATCTCAACTCAGTTACCCG	
	GTGGACTGAGAC	
Pr4	CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTACTTTCATCTGGGGCCTTTACTGAC	
	TCGCGTTTCCT	
Pr5	biotin-TAAGGCCCCAGATGATAAACCCAGGCCGGCCCT	
Pr6	TCGCTCGACAGAGATACTATTAGTACCAGGCCCGATAGGGCCCGGCCTGGGTTT	
Pr7	biotin-TCGCAGGCGTTAGGGTTCGGGGCCTGGTACTAAT	
Pr8	biotin-TACTGAGTTGAGATCTGTATCTCTGTCGAGCGA	
Pr9	CCCTAACGCCTGCGTGAATAGGGCTCTGTTGC	Pr2 without aptamer
Pr10	GATCTCAACTCAGTTACCCGGTGGACTGAGAC	Pr3 without aptamer
Apt	biotin-CTGGGCGGTAGAACCATAGTGACCCAGCCGTCTAC	

 Table S1. DNA oligonucleotides samples used in the study.

(b)



Figure S1. Formation of different DNA polyhedra. 150 nM of DNA was in each lane. (**a**) Tetrahedron 1 of theoretical height 2.65 nm. (**b**) Tetrahedron 3 of theoretical height of 10.55 nm. (**c**) Square-based pyramid. (**d**) Pentagon-based pyramid. (**e**) and (**f**) Formation of DNA prism with three aptamers as in Figure 5(c).

(a)