

Supporting Information

Comparative evaluation and design of G-triplex/thiflavin T-based molecular beacon

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Table S1. Oligonucleotide sequences for forming G3

Name	Sequence (5'-3')
G3	TGGGAAGGGAGGG
G3-T ₃	TGGGAAGGGAGGGTTT
G3-T ₄	TGGGAAGGGAGGGTTTT
G3-T ₅	TGGGAAGGGAGGGTTTTT
G3-T ₆	TGGGAAGGGAGGGTTTTT
G3-T ₇	TGGGAAGGGAGGGTTTTTT
G3-A ₃	TGGGAAGGGAGGGAAA
G3-A ₄	TGGGAAGGGAGGGAAAA
G3-A ₅	TGGGAAGGGAGGGAAAAAA
G3-A ₆	TGGGAAGGGAGGGAAAAAA
G3-A ₇	TGGGAAGGGAGGGAAAAAAA
G3-(AT) ₂	TGGGAAGGGAGGGATAT
G3-(AT) ₃	TGGGAAGGGAGGGATATAT
G3-(AT) ₄	TGGGAAGGGAGGGATATATAT
G3-(AT) ₅	TGGGAAGGGAGGGATATATAT

Table S2. Oligonucleotide sequences of target DNA and related DNA in this work

Name	Sequence (5'-3')
p 53 DNA	TCATCACACTGGAAGACTC
smT	TCATCACACCGGAAGACTC
dmT	TCATCATACTGGCAGACTC
ncT	ATGGGTCAGATTACGTTG A

Table S3. Oligonucleotide sequences of MB used in this work*

Name	Sequence (5'-3')
G3M53	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> TTCCC
G3M43	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> TTCCC
G3M42	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> TTCC
G3M41	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> TC
G3M35	TGGGAAG <u>GGAGGGAAAAAAGAGTCTTCCAGTGTGATGATT</u> CCCTC
G3M33	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> CCC
G3M32	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> CC
G3M26	TGGGAAG <u>GGAGGGAAAAAAGAGTCTTCCAGTGTGATGATT</u> CCCTCC
G3M25	TGGGAAG <u>GGAGGGAAAAAAGAGTCTTCCAGTGTGATGATT</u> CCCTC
G3M23	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGATT</u> CCC
G3M17	TGGGAAG <u>GGAGGGAAAAAAGAGTCTTCCAGTGTGATGAT</u> CCCTCCC
G3M16	TGGGAAG <u>GGAGGGAAAAAAGAGTCTTCCAGTGTGATGAT</u> CCCTCC
G3M15	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGAT</u> CCCTC
G3M07	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGAC</u> CCCTCCC
G3M06	TGGGAAG <u>GGAGGGAAAAAAGAGTCTTCCAGTGTGATGAC</u> CCCTCC
G3M05	TGGGAAGGGAG <u>GGAAAAAAGAGTCTTCCAGTGTGATGAC</u> CCCTC
G4-MB	AGGGTTAGGGTTAGGGTTAGGGCTTGAGGAGTCTCCAGTGTGATGA CTCAAGCCCTAAC
G4-MB-1	AGGGTTAGGGTTAGGGTTAGGGCTTGAGTCTCCAGTGTGATGAAAG CCC

* Underlined bases denote the stem of G3-based MB.

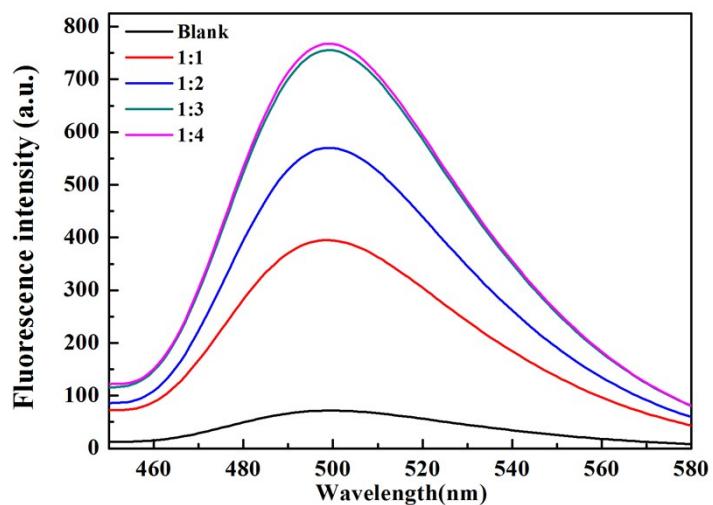


Figure S1. Fluorescence emission spectra of the G4-based MB under the different MB-to-target ratios. MB probe concentration was 0.45 μ M. Reaction media were 75 mM KCl, and 20 mM Tris-HCl (pH 8.3).

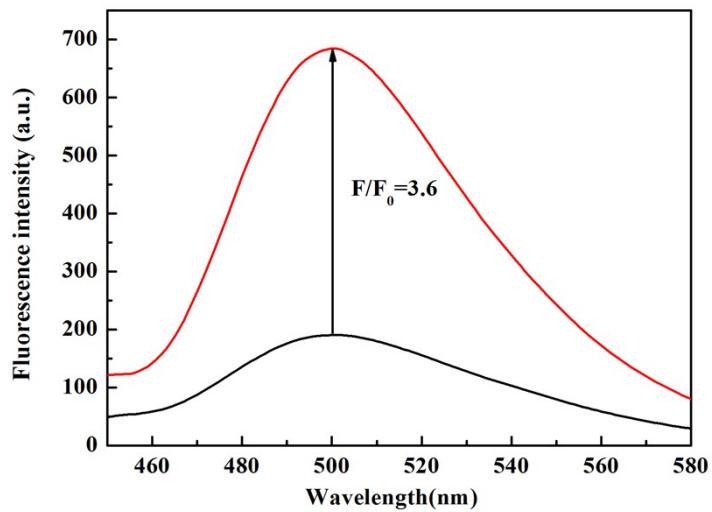


Figure S2. Fluorescence spectra of the G4-based MB-1 in the absence (black line) and presence (red line) of target p53.

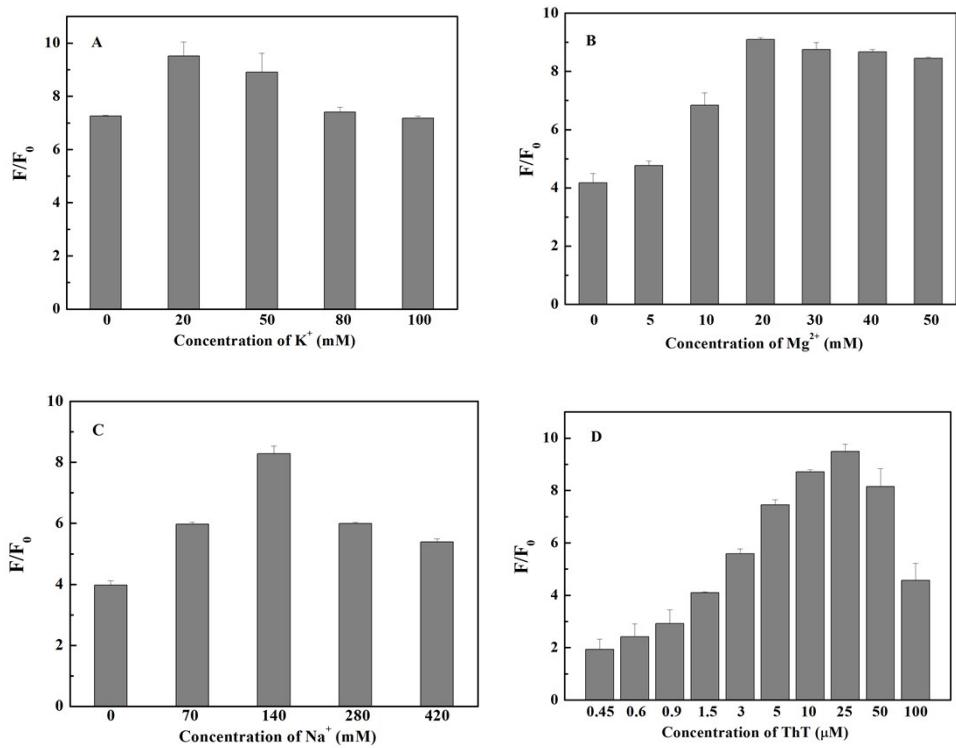


Figure S3. Effect of ions and ThT concentrations on the response of G3-based MB.

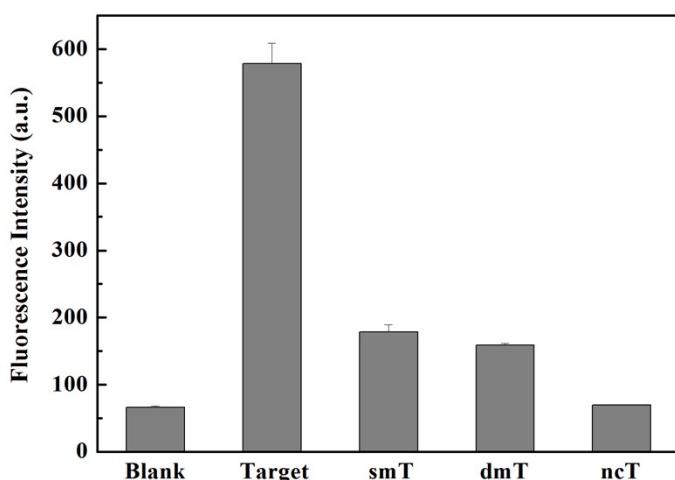


Figure S4. Fluorescence response of G3/ThT-based MB system to different DNA sequence: single-base mismatch DNA (smT), double-base mismatch DNA (dmT), random sequence (ncT), and target DNA.

Table S4. Detection of target DNA in 1% human serum samples (n=3).

Samples	Added	Found	Recovery (%)	RSD (%)
No. 1	50 nM	49 nM	98	6.4
No. 2	90 nM	89.8 nM	99	1.2
No. 3	200 nM	201 nM	101	1.2