

Supplementary Materials

Sensitive aptasensing of tobramycin through a rational design of catalytic hairpin assembly and hybridization chain reaction amplification monomers for CRISPR/Cas12a activation

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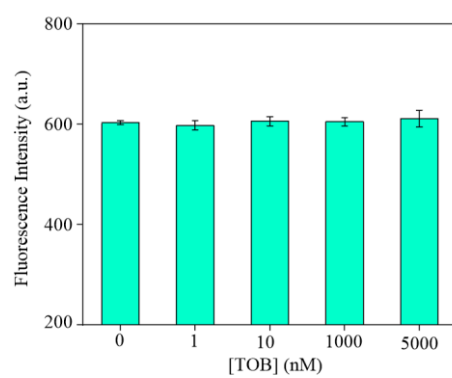


Figure S1 Dependence of the fluorescence intensity on the TOB concentration in the absence of AP.

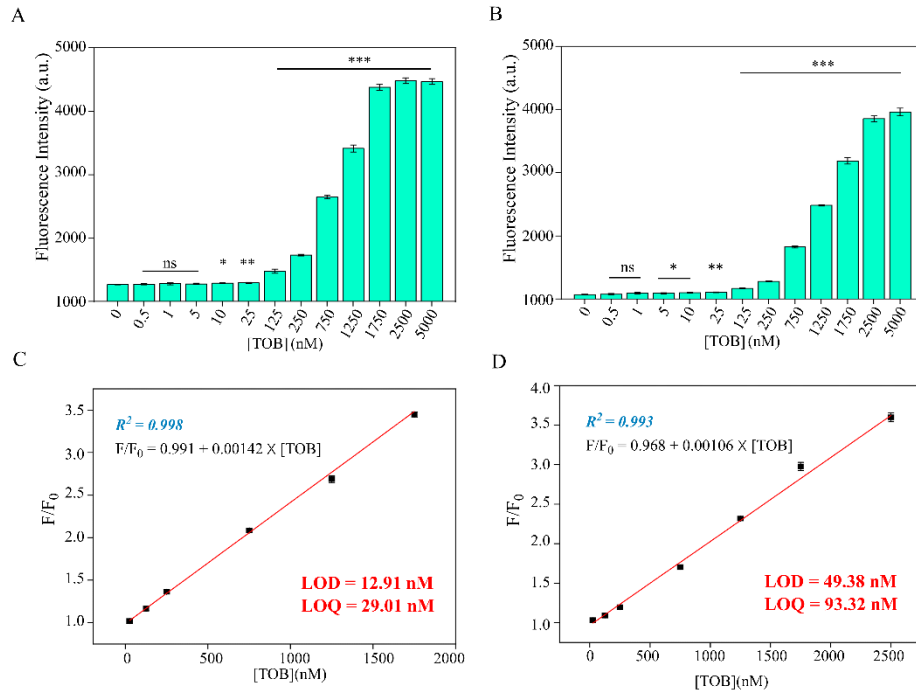


Figure S2 Analytical performance of the aptasensor for the detection of TOB in real samples. (A) Dependence of the fluorescence intensity on the TOB concentration in beef samples. (B) Dependence of the fluorescence intensity on the TOB concentration in milk samples. * $p \leq 0.05$, ** $p \leq 0.01$, *** $p \leq 0.001$, ns: no significant difference. (C) Linear relationship between the F/F_0 value and the TOB concentration in beef samples. (D) Linear relationship between the F/F_0 value and the TOB concentration in milk samples. The error bars represent the standard deviation of three repetitive measurements.

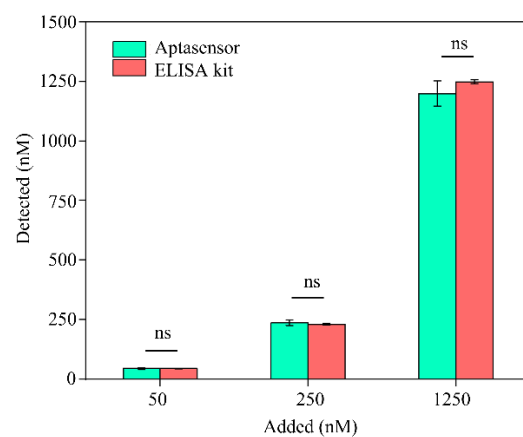


Figure S3 Comparison of the proposed aptasensor with a commercially available ELISA kit for detecting TOB in milk samples. ns: no significant difference. Data are presented as means and error bars represent the standard deviations of three repetitive measurements.

Table S1. Sequence of the oligonucleotides used in APF-CRISPR for ATP detection.

Name	Sequence (5'-3')
H1 _{SL} -HCR	CACACCGACTCCAGGTTTCGTATAGTACAGCGAAACCTGGAGTC
H2 _{SL} -HCR	ACAGCGAAACCTGGAGTCGGTGTGGACTCCAGGTTTCGCTGTACTATA
H1 _{SL} -CHA	CACACCGACTCCAGGTTTCGTATAGTACAGCGAAACCTGGAGTC
H2 _{SL} -CHA	CAGGTTTCGCTGTACTATACGAAACCTGGAGTCCTTTCGTATAGTACAGC
H1 _{PF} -HCR	GAGGAACAGGTTTCGTATAGTACAGCGTGATTTGCGCTGTACTATACGAAACCTG
H2 _{PF} -HCR	GCTGTACTATACGAAACCTGTTCTCCAGGTTTCGTATAGTACAGCGAAATCAC
H1 _{PF} -CHA	CACACCGACTCCAGGTTTCGTTTCGCTGTACTATACGAAACCTGCGAAACCTGGAGTC
H2 _{PF} -CHA	GTTTCGAGGTTTCGTATAGTACAGCGAAACGAAACCTGGAGTCGCTGTACTATACGA AACCTG
Initiator _{SL} -HCR	CGAAACCTGGAGTCGGTGTG
Initiator _{SL} -CHA	CGAAACCTGGAGTCGGTGTG
Initiator _{PF} -HCR	GCTGTACTATACGAAACCTGTTCTC
Initiator _{PF} -CHA	CGAAACCTGGAGTCGGTGTG
AP (ID: 1 bp, AD: 7 bp)	GACTAGGCACTAGTCCGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 2 bp, AD: 7 bp)	GACTAGGCACTAGTCCAGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 3 bp, AD: 7 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 4 bp, AD: 7 bp)	GACTAGGCACTAGTCCACAAGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 5 bp, AD: 7 bp)	GACTAGGCACTAGTCCACACCGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 6 bp, AD: 7 bp)	GACTAGGCACTAGTCCACACAAGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 3 bp, AD: 6 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAG
AP (ID: 3 bp, AD: 7 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 3 bp, AD: 8 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAGT
AP (ID: 3 bp, AD: 9 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAGTGC
AP (ID: 3 bp, AD: 10 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAGTGCC
AP (ID: 3 bp, AD: 11 bp)	GACTAGGCACTAGTCCACCGAAACCTGGAGTCGGTGTGGACTAGTGCCT
TS	CAGGTTTCGTATAGTACAGCGAAATCAC
NTS	GTGATTTGGCTGTACTATACGAAACCTG
crRNA	UAAUUUCUACUAAGUGUAGAUGCUGUACUAUACGAAACCUG
FQ probe	FAM - TTTATT - BHQ1

Note: The underlined sequences represent the complementary region. The sequences of PAM and its complimentary sequences are in purple font. The sequences in green font are the target sequence (TS) of CRISPR/Cas12a. In the AP sequence, the aptamer of TOB is in red font and HCR initiator is in blue font. ID means the duplex region in the initiator domain, and AD means the duplex region in the aptamer domain. The optimized AP sequence is shown with a yellow background.

Table S2. Comparison of the detection performances of the aptasensor for TOB determination

Signal	Amplification Strategy	Linear range	Limit of detection (LOD)	Samples	References
Fluorescence	-	80 nM – 2 μ M	21.86 nM	Human serum	[1]
Fluorescence	-	0.1 – 6 μ M	0.063 μ M	Human serum and Milk	[2]
Fluorescence	-	50 – 500 nM	15.3 nM	Milk	[3] [★]
Fluorescence	-	1000 – 8125 pM	860 pM	Milk	[4] [★]
Colorimetric	-	0.1 – 100 nM	70 pM	Milk	[5] [★]
Colorimetric	-	40 – 200 nM	23.3 nM	Milk and egg	[6] [★]
PEC	-	5 – 50 nM	4.28 nM	Milk	[7] [★]
PEC	-	0.01 – 50 ng/ml	4.27 pg/ml	Milk and water	[8] [★]
ECL	-	1 pM – 1 μ M	18 pM	Human serum	[9] [★]
SERS	-	6.67 – 300 nM	1.26 nM	Beef and mutton	[10] [★]
LFA	-	0.1 – 4 nM	0.02 nM	Milk and honey	[11] [★]
Electrochemical	HCR	5 – 1000 nM	3.51 nM	Milk	[12] [★]
Fluorescence	HCR	0.3 – 50 μ M	17.37 nM	Milk	[13] ^{△★}
Fluorescence	SDA	10 – 300 pM	3.719 pM	Milk and water	[14]
Colorimetric	SDA	20 – 800 nM	12.24 nM	Milk and water	[15]
Electrochemical	SDA	10 – 200 nM	5.13 nM	Milk	[16] [★]
Fluorescence	SDA&HCR	0.5 – 30 nM	0.06 nM	Milk and water	[17] [★]
Fluorescence	HCR	0.125 – 25 nM 25 – 2500 nM	92.87 pM	Milk and beef	This work

[△]Magnetic separation[★]Nanomaterials labeling

Photoelectrochemical (PEC)

Electrochemiluminescence (ECL)

Surface-enhanced Raman Scattering (SERS)

Lateral Flow Assay (LFA)

Hybridization Chain Reaction (HCR)

Strand Displacement Amplification (SDA)

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