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## **Supplementary Information**

## Programmable CRISPR/Cas12a Activity by Adjusting Guide RNA ConformationforNon-nucleic Acid Markers Analysis andLogic Gate Applications

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Oligonucleotide name	Sequence (5' to 3') description			
crRNA	UAAUUUCUACUAAGUGUAGAUGGUUGAUAACUAAACCUGG			
	G			
Apt-T1	ACCTGGGGGAGTACCAAGTACGAA			
Apt-T2	TAATTTCTACT <u>GGTGCGGAGGAAGGT</u>			
blocker1	TTCGTACTTAGTAGAAATTA			
blocker2	TACACTT/iHS-SH/AGTAGAA			
blocker2a	TACACTTAGTAGAA			
blocker3	TACACTTA80x0-GTAGAAA			
blocker3a	TACACTTAGTAGAAA			
S	CCCAGGTTTAGTTATCAACC			
FQ	FAM-TTATT-BHQ			
blocker1a	TTCGTCTACACTTAGTAGAAATTA			
blocker1b	<i>TTCGT</i> CTACTTAGTAGAAATTA			
blocker1c	TTCGTTTAGTAGAAATTA			
Apt-T1a	ACCTGGGGGAGTACCAAGTGTAGACGAA			
Apt-T1b	ACCTGGGGGAGTACCAAGTAGACGAA			
Apt-T1c	ACCTGGGGGAGTACCAAACGAA			
ATP G1	CTAAAACCTGGGGGGAGTATTGCGGAGGAAGGT			
ATP G2	ACCTGGGGGAGTATTGCGGAGGAAGGTTGATAA			

Table S1. DNA oligonucleotides sequences used in this work.

The underlined part of Apt-T1 and Apt-T2 is the split aptamer of ATP, the underlined part of ATP G1 and ATP G2 is the complete aptamer of ATP. The middle of the blocker 2 was decorated with the disulfide bond (iHS-SH). The middle of the blocker 3 was decorated with the 8-oxoadenine (80xo-G). The 3' and 5' ends of FQ were modified with FAM and BHQ, respectively.



**Fig. S1.** Effect of length of blocker 1 on the fluorescence signal. 13, 15, 17, 19 nt blocker 1 was used.Gray: fluorescence background signal intensity without adding the target; Blue: The intensity of the fluorescent signal when the target is added.



**Fig. S2.** Optimal experiment.(A) Effect of the concentration of FQ on the fluorescence signal of the biosensor in the presence (black) and absence (red) of 10  $\mu$ M ATP.(B) Effect of the reaction time on the fluorescence signal of the biosensor in the presence (black) and absence (red) of 10  $\mu$ M ATP. (C) Effect of the reaction temperature on the fluorescence signal of the biosensor. (D) Effect of the concentration of Switch 1 on thefluorescence signal of the biosensor in the presence (black) and absence (red) of 10  $\mu$ M ATP.



Fig. S3. Schematic illustration of the proposed universal biosensing strategy for GSH detection.



Fig. S4. PAGE characterizes the feasibility of the universal biosensor for GSH detection.



Fig. S5. Fluorescence spectra of the biosensor obtained upon the different control conditions.



Fig. S6.Fluorescence intensity of "AND" logic gate and "OR" logic gate.(A) Fluorescence intensity of "AND" logic gate with ATP and Fpg as inputs. (B) Fluorescence intensity of "OR" logic gate with ATP and Fpg as inputs.(C) Fluorescence intensity of "OR" logic gate with ATP and GSH as inputs.



**Fig. S7.** The feasibility validation of this approach to implement an arbitrary logic circuit. (A) The fluorescent peak intensity of the "AND" logic gate by varying different input units. (B) The fluorescent peak intensity of the "OR" logic gate by varying different input units. (C) The fluorescent peak intensity of the three-input two-layered logic gate by varying different input units.

Detection methods	LOD(µM)	Detection range(µM)	References
Fluorescence	19	125-2000	[1]
Fluorescence	2	5-2500	[2]
Fluorescence	1	5-600	[3]
Colorimetric	0.1225	0.2-3	[4]
Electrochemiluminescence	0.64	0.64-1000	[5]
Fluorescence	0.046	0-12.8	This work

Table S2. Comparison of different methods for ATP determination.

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