A novel steric effect-regulated isothermal exponential amplification

technology for one-step homogeneous sensing of proteins

Wanghua Wu^a, Wufan Pan^b, Dongdong Yu^c, Zhen Yuan^a, Yazhou Qin^b, Yuxiang Lu^b, Tao Zhang*^a and Jianguang Zhou*^a

^aResearch Center for Analytical Instrumentation, Institute of Cyber-Systems and Control, State Key Laboratory of Industrial Control Technology, Zhejiang University, Hangzhou, 310027, P.R. China
 ^bDepartment of Chemistry, Zhejiang University, Hangzhou, 310027, P.R. China
 ^cHospital of Zhejiang University, Zhejiang University, Hangzhou, 310027, P.R. China

*corresponding authors: zhtao@zju.edu.cn, jgzhou@zju.edu.cn

Experimental

Materials. All oligonucleotides of HPLC purity were synthesized from Sangon Biotech Co., Ltd. (Shanghai, China). Sequences of these oligonucleotides are listed in Table S1. Bst 2.0 WarmStart DNA polymerase (8000 units mL⁻¹), Nt.BstNBI nicking endonuclease (10000 units mL⁻¹), deoxyribonucleotide triphosphates (dNTPs, 10 mM) and 10× NEBuffer 3.1 were purchased from New England Biolabs (Ipswich, MA). Anti-Digoxin mouse monoclonal antibody (IgG, 0.5 mg mL⁻¹) was purchased from GeneTex, Inc. (Irvine, CA, USA). Avidin from egg white (Bio-Ultra), Elastase from porcine pancreas and human thrombin were purchased from Sigma Aldrich (St. Louis, MO, USA). SYBR Green I (10,000×) was purchased from Invitrogen Life Technologies (Carlsbad, CA, USA). DNA-grade water (DNase- and Protease-free) was purchased from Fisher Scientific (Pittsburgh, PA, USA). The clinical serum samples of healthy person were obtained from the Hospital of Zhejiang University. All solutions were freshly prepared with DNA-grade water.

General Procedure of SER-EXPAR. First, a stock solution of 20 nM EBH, 50 nM EXPAR template, 2 U Nt.BstNBI, 0.8 U Bst 2.0 WarmStart DNA polymerase, 0.4 mM dNTPs, 0.5× SYBR Green I, 1× NEBuffer 3.1 (100 mM NaCl, 50 mM Tris-HCl, 10 mM MgCl₂, 100 μ g/ml BSA) and water was freshly prepared on ice. The final reaction mixture was obtained by distributing 22.5 μ L stock solution and 2.5 μ L target protein of varying concentrations into 8-strip PCR tubes. After vortexing and centrifugation, these PCR tubes were immediately transferred onto a Bio-rad CFX-96 real-time PCR system and incubated at 55 °C. The fluorescence of SYBR Green I was measured at 1 min intervals.

Gel Electrophoresis. In order to demonstrate the feasibility of SER-EXPAR, 15% non-denaturing polyacrylamide gel electrophoresis (PAGE) was used to characterize the interactions among avidin, Nt.BstNBI and biotin-labeled EBH. Specifically, a solution containing 1 μ M EBH₆ and 1 μ M Nt.BstNBI or 1 μ M target protein (or both) in 1× NEBuffer 3.1 (100 mM NaCl, 50 mM Tris-HCl, 10 mM MgCl₂, 100 μ g/ml BSA, pH 7.9@25 °C) was first incubated at 55 °C for 15-20 min. Afterwards, 5 μ L of each resultant were loaded onto 15% non-denaturing polyacrylamide gel and subjected to 150 V constant voltage in 0.5× TBE buffer (45 mM Tris, 45 mM borate, 1 mM EDTA, pH 8.3) for 60 min. The gel was then stained by 1 × SYBR Green I and analyzed on a Maestro Ex IN-VIVO Imaging System (CRI).

Sensing of Avidin in Human Serum. In order to demonstrate the utility of SER-EXPAR in complex biological matrix, it was performed in the presence of human serum. The serum was used, without dilution or other pretreatment,

to dilute target protein (avidin) by tenfold in the series. Subsequently, 2.5 μ L of spiked serum was introduced into SER-EXPAR system and subjected to the same procedure as described above. The results were compared with those obtained in buffer.

Name	Sequence (5'-3')		
EXPAR template	CTCACGCTACGGACGACTCTCTCACGCTAC		
EBH1	CATCACTGCAC <u>TGACTCACGTTTTTCGTGAGTCA</u> ACTCCTGTAAG		
EBH ₂	CATCACTGCAC <u>TGACTCACGTTTTTCGTGAGTCA</u> ACT		
EBH ₃	CAC <u>TGACTCACGTTTTTCGTGAGTCA</u> ACTCCTGTAAG		
EBH ₄	CAC <u>TGACTC</u> ACGTTTTTCGT <mark>GAGTC</mark> AACT		
EBH ₅	CATCACTGCACTGACTCACGTTTTTCGTGAG (biotin-T) CAACTCCTGTAAG		
EBH ₆	CATCACTGCACTGACTCACGTTTTTCG (biotin-T) GAGTCAACTCCTGTAAG		
EBH ₇	CATCACTGCACTGACTCACGTTTT (biotin-T) CGTGAGTCAACTCCTGTAAG		

Table S1. Sequences of the Oligonucleotides.^[a]

^[a]The underlined bases represent the basic sequences of hairpin probes' stem-loop structure, among which the red bases are the recognition sites of Nt.BstNBI.

Detection limit	Detection range	Time	Complexity	Ref.
0.4 nM	$0\text{nM}{\sim}200\text{nM}$	\sim 30 min	Multi-steps	1
low nanomolar	1 nM \sim 10 nM	10 min	One-step, need	2
			surface modification	
0.1 nM	$0.5\mathrm{nM}{\sim}100$	>1.5 h	Multi-steps	3
	nM			
1.07 nM	$4 nM \sim 1 \mu M$	3 h	Multi-steps	4
nanomolar	$0\mathrm{nM}{\sim}2\mathrm{nM}$	overnight	One-step	5
1.29 nM	5 nM \sim 50 nM	20-25 min	One-step	/
	Detection limit 0.4 nM low nanomolar 0.1 nM 1.07 nM nanomolar 1.29 nM	Detection limit Detection range 0.4 nM 0 nM~200 nM low nanomolar 1 nM~10 nM 0.1 nM 0.5 nM~100 0.1 nM 0.4 nM~100 1.07 nM 4 nM~1 µM nanomolar 0 nM~2 nM 1.29 nM 5 nM~50 nM	Detection limit Detection range Time 0.4 nM $0 \text{ nM} \sim 200 \text{ nM}$ $\sim 30 \text{ min}$ low nanomolar $1 \text{ nM} \sim 10 \text{ nM}$ 10 min 0.1 nM $0.5 \text{ nM} \sim 100$ $> 1.5 \text{ h}$ 0.1 nM $0.5 \text{ nM} \sim 100$ $> 1.5 \text{ h}$ 1.07 nM $4 \text{ nM} \sim 1 \mu \text{M}$ 3 h 1.07 nM $0 \text{ nM} \sim 2 \text{ nM}$ overnight 1.29 nM $5 \text{ nM} \sim 50 \text{ nM}$ $20 \cdot 25 \text{ min}$	Detection limitDetection rangeTimeComplexity 0.4 nM $0 \text{ nM} \sim 200 \text{ nM}$ $\sim 30 \text{ min}$ Multi-stepslow nanomolar $1 \text{ nM} \sim 10 \text{ nM}$ 10 min One-step, need surface modification 0.1 nM $0.5 \text{ nM} \sim 100$ > 1.5 h Multi-steps 0.1 nM $0.5 \text{ nM} \sim 100$ > 1.5 h Multi-steps 1.07 nM $4 \text{ nM} \sim 1 \mu M$ 3 h Multi-steps 1.07 nM $0 \text{ nM} \sim 2 \text{ nM}$ overnightOne-step 1.29 nM $5 \text{ nM} \sim 50 \text{ nM}$ $20-25 \text{ min}$ One-step

Table S2. Comparison of different methods based on small molecule-linked DNA for the detection of avidin.

References:

1 X. X. Chen, C. S. Lin, Y. Y. Chen, F. Luo, Y. R. Wang and X. Chen, Biosens. Bioelectron., 2016, 83, 97-101.

2 S. Ranallo, M. Rossetti, K. W. Plaxco, A. Vallée-Bélisle and F. Ricci, *Angew. Chem. Int. Ed.*, 2015, **54**, 13214-13218

15210

3 H. B. Wang, H. D. Zhang, Y. Chen and Y. M. Liu, *Biosens. Bioelectron.*, 2015, **74**, 581-586.

4 F. F. Ban, H. Shi, C. Feng, X. X. Mao, Y. M. Yin and X. L. Zhu, *Biosens. Bioelectron.*, 2016, 86, 219-224.

5 Z. Zhang, C. Hejesen, M. B. Kjelstrup, V. Birkedal and K. V. Gothelf, J. Am. Chem. Soc., 2014, 136, 11115-11120

Supplement Figures



Figure S1. (a) Four kinds of EBH with different side arm length. (b) Effects of 10nM and 20nM $EBH_1 \sim EBH_4$ on the real-time fluorescence of EXPAR; NTC represents no-EBH control.



Figure S2. (a) Three kinds of EBH labeled with biotin at different sites, *i.e.*, among, near and relatively far from the recognition sequence of Nt.BstNBI. (b) Effects of 10nM and 20nM $\text{EBH}_5 \sim \text{EBH}_7$ on the real-time fluorescence of EXPAR; NTC represents no-EBH control.



Figure S3. Real-time fluorescence curves caused by 10nM and 20nM avidin in the presence of 20nM EBH₆ (Left) and EBH₇ (Right); NTC represents no-avidin control.



Figure S4. Real-time fluorescence curves caused by 20nM, 25nM and 30nM avidin; NTC represents no-avidin control; "+" represents that the system is in the presence of 20nM EBH_6 and "-" represents that the system is in the absence of EBH_6 .



Figure S5. (a) Real-time fluorescence curves caused by various concentrations of EBH₆ without avidin; NTC represents no-EBH₆ control. (b) Real-time fluorescence curves caused by various concentrations of avidin in the presence of 50nM EBH₆; NTC represents no-avidin control.



Figure S6. Real-time fluorescence curves caused by 20 nM avidin and the protein mixture in the presence of 20 nM EBH₆. Mix represents the mixture of 200 nM IgG, 200 nM thrombin and 200 nM elastase. NTC represents no-protein control. (Overmuch proteins (*i.e.* 3*500 nM) in the buffer would influence the rate of EXPAR, thus the total amount of proteins in the mix was set 3*200 nM that had little effect on the EXPAR).