A Fluorescence Assay for Exosome Detection Based on Bivalent Cholesterol-Anchor Triggered Target Conversion and Enzyme-Free Signal Amplification

Xiaokun Wang,† Hezhen Shang,‡ Cuiping Ma,*† Lingxin Chen*§l#

† Shandong Provincial Key Laboratory of Biochemical Engineering, College of Marine Science and Biological Engineering, Qingdao University of Science and Technology, Qingdao, 266042, PR. China.

[‡] Department of Hepatobiliary Surgery, Qingdao Chengyang District People's Hospital, Qingdao 266109, PR. China.

§CAS Key Laboratory of Coastal Environmental Processes and Ecological Remediation; Shandong Key Laboratory of Coastal Environmental Processes, Yantai Institute of Coastal Zone Research, Chinese Academy of Sciences, Yantai 264003, China.

¹The Key Laboratory of Life-Organic Analysis, Key Laboratory of Pharmaceutical Intermediates and Analysis of Natural Medicine, College of Chemistry and Chemical Engineering, Qufu Normal University, Qufu 273165, China

*School of Pharmacy, Binzhou Medical University, Yantai 264003, China.

Phone: +86-535-2109130 Fax: +86-535-2109130

*Corresponding author: mcp169@163.com (C. Ma); E-mails: lxchen@yic.ac.cn (L. Chen)

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General Experimental Section

Instruments. Native polyacrylamide gel electrophoresis (PAGE) was carried out using vertical electrophoresis systems (Beijing, China). Confocal microscopy fluorescence images were obtained by an Olympus FV1000. Transmission electron microscopy (TEM) image was acquired using a JEM-1230 instrument (JEOL, Tokyo, Japan). DS-11 Series UV-Vis Spectrophotometer (Wilmington, USA) and fluorescence spectrophotometer (F-7000, HITACHI) were used to obtain absorption and fluorescence spectral data, respectively. The annealing process was carried out by using PCR (Framingham, USA). The size and concentration of exosomes were obtained by using nanoparticle tracking analysis (NTA) at VivaCell Biosceinces with ZetaView PMX 110 (Particle Metrix, Meerbusch, Germany) and corresponding software ZetaView 8.04.02.

PAGE analysis of enzyme-free DNA circuits. For the enzyme-free DNA circuits, the sequences were designed according to previous literature with a slight modification. ¹ TNK buffer containing BC-anchor (5 nM), H1 (90 nM), H2 (135 nM), and RFQ (90 nM) was incubated for 2 h at room temperature. After different reaction products were prepared, $10 \mu L$ of each sample was mixed with 2 μL of 6×loading buffer, then 10 μL of the mixed solution was loaded into the lanes of a polyacrylamide gel. The gel was run at 180 V constant voltage for 2 min and then 135 V constant voltage for 70 min in 1×TAE buffer at room tem-perature. After the gel was stained for 1 min in EtBr solu-tion, it was photographed using the multifunctional imag-ing system.

Exosome quantification by enzyme-linked immunosorbent assay (ELISA). 30 μ L Anti-CD63 MBs were mixed with 20 μ L different concentrations of exosome and allowed to react under gentle shaking at room temperature. After 2 h, the mixture was washed three times with PBST and resuspended in PBS. 50 μ L anti-CD 63 rabbit antibody and anti-rabbit IgG-HRP were sequentially added to the immunocomplexes at an interval of 2 h. After incubation for another 1 h, the mixture was washed three times. Subsequently, 3,3',5,5'-tetramethylbenzidine (TMB) solution was added to induce the enzymatic reaction and TMB stop solution was added to terminate the reaction. Finally, the absorption spectra were measured by UV-Vis Spectrophotometer.

Table S1. Sequences of oligonucleotides used in this work.

Name	Sequence (5'-3')
I	CGACATCTAACCTAGCTCACTGACATAAGGCACGACGGCTTT-Cholesterol
I_c	Cholesterol-TTTGCCGTCGTGCCTTAT
I_{AF}	GTCAGTGAGCTAGGTTAGATGTCG-Alexa Fluor 488
H1	GTCAGTGAGCTAGGTTAGATGTCGCCATGTGTAGACGACATCTAACCTAGCCCTTGTCATAGAGCAC
H2	AGATGTCGTCTACACATGGCGACATCTAACCTAGCCCATGTGTAGA
RF	FAM-CGAGTGCTCTATGACAAGGGCTAGGTT
RQ	CCCTTGTCATAGAGCACTCG-Dabcyl

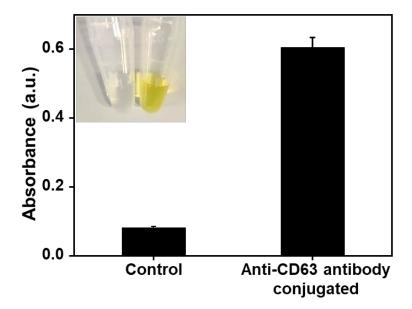


Figure S1. Absorption intensities at 450 nm for bare (control) and anti-CD63 antibody-conjugated magnetic beads.

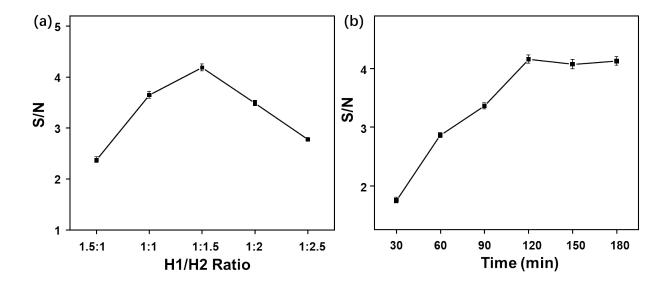


Figure S2. Optimization of experimental conditions. (a) Optimization of concentration ratios of H1 to H2. (b) Optimization of the incubation time for DNA circuitry.

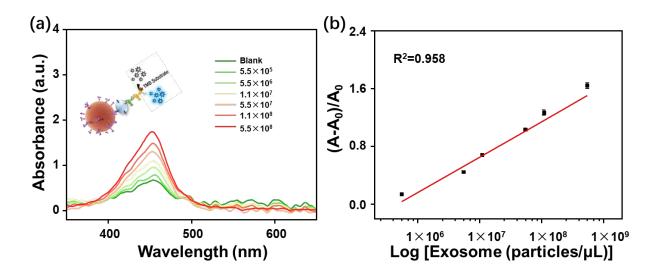


Figure S3. (a) Absorption spectra response to different concentrations of exosomes. (b) Linear relationship between the absorption intensity ratio and logarithmic values of exosome concentrations. A and A_0 refer to the absorption intensities with and without the addition of exosome, respectively.

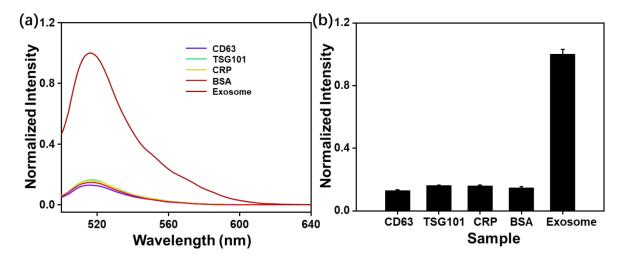


Figure S4. Evaluation of assay specificity. (a) Normalized fluorescence spectra response to CD63 (2 μ g/mL), TSG101 (2 μ g/mL), CRP (2 μ g/mL), BSA (2 μ g/mL), and exosome (1.1×10⁶ particles/ μ L), respectively. (b) Corresponding fluorescence intensity to (a).

 Table S2. Comparison among exosome detection methods.

Detection Method	Identification Target	Linear Range (particles/μL)	Limit of Detection (particles/µL)	Real Sample	Reference
Colorimetry assay	CD63	1.8×10^6 - 2.2×10^7	5.2×10^3	Serum	2
Colorimetry assay	CD63 and CD9	2.2×10^5 - 2.4×10^7	_	Serum	3
Paper-based aptasensor with LRET	CD63	$1.0 \times 10^4 \text{-} 1.0 \times 10^8$	1.1×10^{3}	_	4
SPRi-based biosensor	CD63	$3.1 \times 10^4 \text{-} 1.0 \times 10^6$	2.37×10^{4}	Serum	5
Lateral flow aptasensor	CD63	$1.0 \times 10^4 - 1.0 \times 10^8$	1.4×10^4	Serum	6
Fluorescent nanosensor	GPC-1	$7.8 \times 10^4 - 3.9 \times 10^9$	6.56×10^4	Serum	7
Fluorescent aptasensor	CD63 and bilipid layer	$7.5 \times 10^4 - 1.5 \times 10^7$	4.8×10^4	Serum	8
Fluorescent aptasensor	CD63	$1.0 \times 10^5 \text{-} 1.0 \times 10^9$	1.0×10 ⁵	Serum	9
Electrochemical aptasensor	CD63	$1.12 \times 10^2 - 1.12 \times 10^8$	96	Serum	10
Electrochemical enzymatic amplification	PSMA	1.0×10 ³ -1. 2×10 ⁵ (particles)	70	FBS	11
Fluorescence biosensor with enzyme-free amplification	CD63 and bilipid layer	$5.5 \times 10^3 - 1.1 \times 10^7$	1.29×10^3	FBS/serum/urine	This Work

Table S3. Recovery values of exosomes spiked into 20% diluted ED-FBS, 20% diluted UC-Serum, and 20% diluted UC-Urine by the proposed assay.

Samples	Added (Particles/μL)	Found (Particles/μL)	Recovery (%)	RSD (%)
	4.0×10 ⁴	3.85×10 ⁴	96.25	3.21
20% Diluted ED-FBS	4.5×10 ⁵	4.21×10 ⁵	93.56	4.41
	2.5×10 ⁶	2.52×10 ⁶	100.8	2.89
	4.0×10 ⁴	3.94×10 ⁴	98.50	3.28
20% Diluted UC-Serum	4.5×10 ⁵	4.69×10 ⁵	104.2	5.25
	2.5×10 ⁶	2.67×10 ⁶	106.8	2.84
	4.0×10 ⁴	3.69×10 ⁴	92.25	5.40
20% Diluted UC-Urine	4.5×10 ⁵	4.38×10 ⁵	97.33	3.13
	2.5×10 ⁶	2.58×10 ⁶	103.2	3.30

Table S4. Clinical information of healthy donors and cancer patients.

Clinical Condition	Number	Gender	Age
	Н1	Male	33
	H2	Male	23
	Н3	Female	33
Healthy donors	H4	Female	30
	Н5	Male	59
	Н6	Female	59
	Н7	Male	34
	P1	Male	68
Liver cancer	P2	Female	69
	Р3	Male	65
	P4	Female	52
Cholangiocarcinoma	P5	Male	70
	P6	Male	77
	P7	Female	49
Breast cancer	P8	Female	55

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