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## Supporting Information

# A multipedal DNA walker for amplified detection of tumor exosomes

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# **Experimental**

#### **Materials and instruments**

Iron(III) chloride hexahydrate (FeCl3•6H2O), gold(III) chloride trihydrate (HAuCl4•3H2O), sodium borohydride (NaBH4), tris (2-carboxyethyl)-phosphine hydrochloride (TCEP), ethylenediaminetetraacetic acid (EDTA), and mercaptohexanol (MCH) were purchased from Sigma (United States). Nb.BbvCl and Nb.BssSI nicking endonucleases (NEases) were purchased from New England Biolabs Ltd. (United States). Sodium acetate (NaAc) and ethylene glycol were purchased from Shanghai Aladdin Reagent Company (Shanghai, China). Dulbecco's Modified Eagle's medium (DMEM) was supplied by Gibco (Gaithersburg, United States). Fetal bovine serum (FBS) was purchased from Hangzhou Sijiqing Biological Engineering Material Co., Ltd. (Hangzhou, China). HeLa cells and MCF-10A cells were from the Institute of Biochemistry and Cell Biology, Chinese Academy of Sciences (Shanghai, China). All other reagents were of analytical grade and were used as received. Water used was purified by a Millipore system with the resistivity of 18 MΩ·cm. All DNA sequences were synthesized and purified by Sangon Biotechnology Co., Ltd. (Shanghai, China). The detailed sequences and modifications were shown in Table S2.

Ultracentrifuge Optima L-100XP (Beckman Coulter, USA) was used to isolate exosomes. JEM-1200EX transmission electron microscopy (JEOL Ltd., Japan) was used for the capture of TEM image. Hitachi S4800 scanning electron microscope (Hitachi, Japan) was used for the capture of SEM image. NanoSight LM10 (Malvern Instruments, UK) was applied for the determination of the concentration and diameter of extracted exosomes. CHI 660D electrochemical workstation (CH Instruments, China) was used for all electrochemical measurements.

#### **Extraction and characterizations of exosomes**

Cells were cultured in DMEM containing 10% FBS, 1% penicillin and streptomycin at 37°C in 5% CO<sub>2</sub> atmosphere. After reached a confluency over 90%, the cells were washed for three times with PBS. Then, 10<sup>8</sup> cells were further incubated with FBS-free DMEM for another 48 h. The culture supernatant was collected for the isolation of exosomes. Generally, it was centrifuged at 2,000 g for 20 min and then at 10,000 g for 30 min to remove cells and proteins. Next, the supernatant was further centrifuged at 110,000 g for 2 h to obtain the precipitated exosomes sediments, which were resuspended in 1 mL of PBS and stored at -80 °C before downstream analysis.

The obtained exosomes were firstly analyzed by negative staining TEM utilizing a 2% solution of phosphotungstic acid. Briefly, 10 µL of the sample solution was absorbed on carbon-coated copper grids for 5 min. Excess fluid was blotted gently by filter paper from the edge. PBS was used to wash the copper grids and phosphotungstic acid solution was then dropped. After removing the stain, the prepared copper grids were dried at room temperature and were visualized by TEM. NanoSight LM10 was then applied to analyze the concentration of obtained exosomes according to the manipulation.

### Fabrication of multipedal DNA walker

Fe<sub>3</sub>O<sub>4</sub>@Au nanoparticles were prepared according to our previous report.<sup>1</sup> Thiolated DNA Probe A dissolved in 10 mM Tris–HCl (1 mM EDTA, 10 mM TCEP, and 0.1 M NaCl, pH 7.4) was mixed with Fe<sub>3</sub>O<sub>4</sub>@Au nanoparticles (0.8 mg/mL). The final concentration of Probe A was 5 μM. After reacting for 16 h, the mixture was "aged" in salts (10 mM phosphate, 0.1 M KNO<sub>3</sub>, pH 7.0) for 24 h. Subsequently, Probe A modified Fe<sub>3</sub>O<sub>4</sub>@Au nanoparticles were purified by magnetic separation and washed by 10 mM phosphate buffer (0.25 M NaCl, pH 7.4). The pretreated cell

culture supernatant which contained exosomes was blended with Probe A modified Fe<sub>3</sub>O<sub>4</sub>@Au nanoparticles and Probe B (5 μM) for 1 h. Then, magnetic separation was performed and the collected nanoconjugates of exosomes and nanoparticles were resuspended in the buffer solution (100 mM NaCl, 50 mM Tris-HCl, 10 mM MgCl<sub>2</sub>, 100 μg/ml BSA, pH 7.9). 0.5 μM Probe C and 0.5 units/μL Nb.BbvCl were further added. After reacted at 37°C for 1 h, exosomes loaded with a number of Probe B as multipedal walker strands were obtained by magnetic separation.

### Modification of working electrode

The substrate gold working electrode was firstly treated with piranha solution (*Caution: highly corrosive*) for 5 min in order to remove any adsorbed materials. Then, the rinsed electrode was polished with silicon carbide paper (P5000) and alumina powders (1.0, 0.3, 0.05  $\mu$ m), successively. Afterward, it was sonicated in ethanol and double-distilled water for 5 min, respectively. Next, the electrode was electrochemically cleaned using 0.5 M H<sub>2</sub>SO<sub>4</sub> and rinsed carefully. After dried with nitrogen, the electrode was incubated with DNA probe D for 10 h. It was further treated with 1 mM MCH for 0.5 h to obtain the well aligned DNA monolayer. Probe D modified electrode was then immersed in the multipedal DNA walker solution obtained previously coupled with probe E (0.5  $\mu$ M) and Nb.BssSI (0.5 units/ $\mu$ L). The walking reaction was carried out at 37°C for 1.5 h. Afterward, the electrode was carefully rinsed before the following measurements.

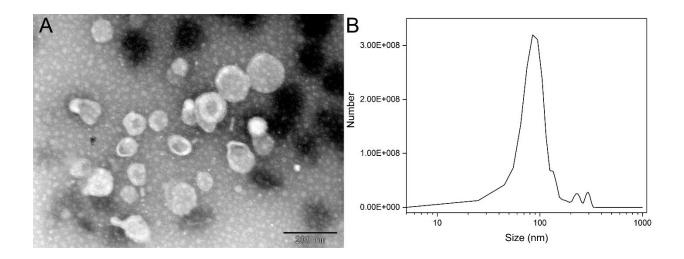
#### **Electrochemical measurements**

A traditional three-electrode system was applied, which was consisted of a platinum auxiliary electrode, a saturated calomel reference electrode and the gold working electrode. Cyclic voltammetry (CV) was scanned from 0.7 to -0.1 V with the scan rate of 50 mV/s. Electrochemical impedance spectroscopy (EIS) was performed with the parameters as follows: biasing potential, 0.217 V; amplitude, 5 mV; frequency range, 0.1 Hz to 100 kHz. Square wave voltammetry (SWV)

was carried out with a step potential of 4 mV, a frequency of 70 Hz, and an amplitude of 25 mV. The scan range was from -0.05 V to -0.55 V.

## **References**

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**Figure S1.** (A) TEM image of HeLa cell-derived exosomes. (B) Nanoparticle Tracking Analysis (NTA) of HeLa cell-derived exosomes.

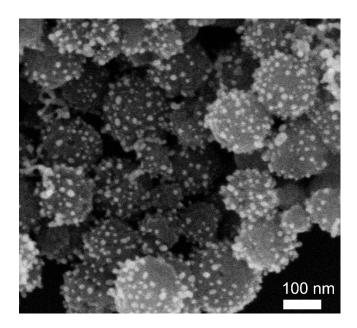
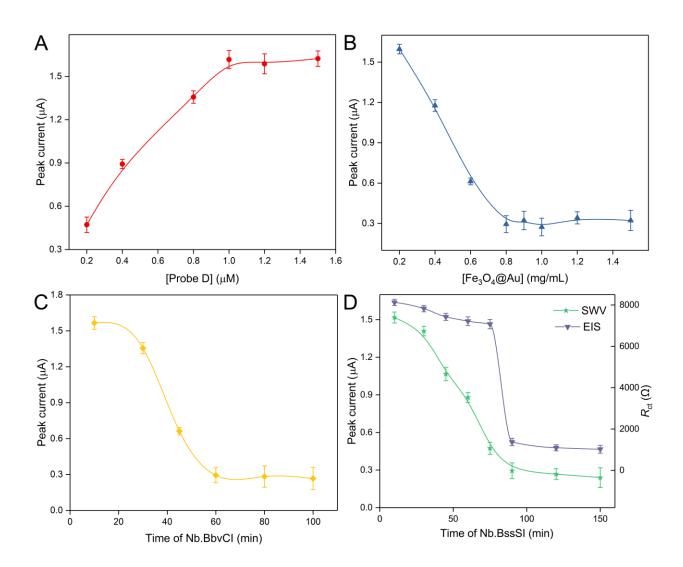
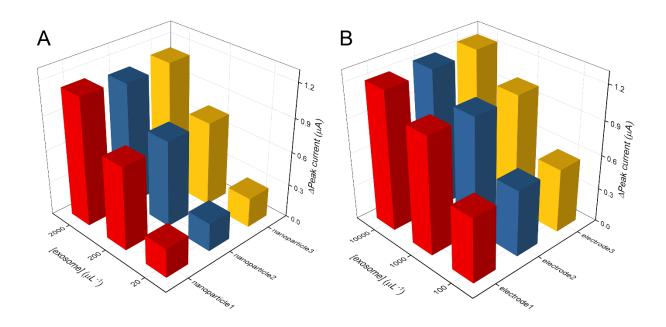


Figure S2. SEM image of Fe<sub>3</sub>O<sub>4</sub>@ Au nanoparticles.



**Figure S3.** Optimization of (A) concentration of Probe D, (B) concentration of Fe<sub>3</sub>O<sub>4</sub>@Au nanoparticles, (C) Nb.BbvCI cleavage time, (D) Nb.BssSI cleavage time.



**Figure S4.** Reproducibility investigation: SWV peak intensities for the detection of exosomes using (A) three sets of Fe<sub>3</sub>O<sub>4</sub>@Au nanoparticles; (B) three independent gold electrodes.

 $\textbf{Table S1.} \ Comparison \ of \ analytical \ performances \ of \ the \ assays \ of \ tumor \ exosomes.$ 

Technique	Strategy	Detection range (/µL)	LOD (/μL)	Ref.
colorimetry	graphitic carbon nitride nanosheets with enhanced peroxidase-like activity	$1.9 \times 10^6$ to $3.38 \times 10^7$	1.352×10 <sup>6</sup>	2
fluorescence	polydiacetylene liposome	$10^5$ to $10^7$	$3 \times 10^{5}$	3
lateral flow assay	double gold nanoparticles conjugates	$7.4 \times 10^3$ to $5.55 \times 10^4$	$1.3 \times 10^{3}$	4
SWASV	quantum dots based immunoassay	$10^2 \text{ to } 10^7$	$10^{2}$	5
ECL	aptamer-binding DNA walking machine	$2 \times 10^2$ to $7.5 \times 10^4$	60	6
SERS	gold nanostar@Raman reporter@nanoshell structures	40 to $4 \times 10^7$	27	7
DPV	DNA walking and exonuclease III-assisted digestion	$50 \text{ to } 10^7$	13	8
giant magnetoresistan ce sensor	2D magnetic MoS <sub>2</sub> -Fe <sub>3</sub> O <sub>4</sub> hybrid nanostructures	$10 \text{ to } 10^6$	10	9
SWV	multipedal DNA walker	10 to $2 \times 10^3$	6	this method

**Table S2.** DNA sequences used in this study.

Name	Sequence (5'-3')		
Probe A	CACCCCACCTCGCTCCCGTGACACTAATGCTATTTTATACGC▼T		
	GAGGATTTT-(CH <sub>2</sub> ) <sub>6</sub> -SH		
Probe B	CACCCCACCTCGCTCCCGTGACACTAATGCTATTTTCTGCAGTT		
	ATTCCGGCTAGT		
Probe C	T <u>CCTCAGC</u> GTAT		
Probe D	MB-ACTAGCCGGGACTC ▼ TCGTGCTTTT-(CH <sub>2</sub> ) <sub>6</sub> -SH		
Probe E	G <u>CACGAG</u> AGTCTAACTGCAG		
Probe B*	TCAAGTCGACTCAAGGTCTGAGTCAACTCCAATTTTCTGCAGTT		
	ATTCCGGCTAGT		