Supporting Information

An Integrated Platform for Capture of Circulating Tumor Cells and

In Situ SERS Profiling of Membrane Protein through Rationally

Spatial Organization of Multi-functional Cyclic RGD Nanopatterns

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1. Experimental Section

Regent and Chemicals. All the reagents and materials were purchased from commercial suppliers, and are of analytical grade. All aqueous solutions were prepared with ultrapure water obtained with a Milli-Q System with a resistivity of 18 MQ·cm. Gold(III) chloride trihydrate (HAuCl₄•3H₂O, 99%), 1-ethyl-3-(3-dimethylaminoprop-yl)-carbodiimide (EDC), N-hydroxysuccinimide (NHS), hexafluorophosphate (HBTU), Dimethylformamide (DMF), H₂O₂, Quercetin, 2,2-diphenyl-1-picrylhydrazyl radical, dicyclohexylcarrbodiimide (DCC) and all the amino acid (99%) were purchased from Sigma-Aldrich. α -Tocopherol was purchased from Calbiochem. KCl, CaCl₂, MgCl₂, FeCl₃, ZnSO₄, CuSO₄, glucose were obtained from Sinopharm Chemical Reagent Co. Ltd. HER2 was obtained from Abcam Co., Ltd (Cambridge, MA). Phosphate buffer solution (PBS) was prepared using K₂HPO₄ and KH₂PO₄, which was deaerated by purging with nitrogen gas. The galvinoxyl radical, trolox, and ascorbic acid were obtained from Acros Organics. Methanol was of HPLC grade and used directly, whereas ethyl acetate and acetic acid were of analytical grade and purified by standard techniques. The peptide, DNEYFYV(H2)-NH-C=O-CRGDKC(RGD)-acetyl (denoted as H2-RGD) were obtained by Shanghai Biotech Bioscience and Technology Co., Ltd., China. All experiments were carried out at room temperature, unless stated otherwise. TMS was synthesized according to previous report.9b

Instruments and Characterization. All electrochemical experiments were conducted with a computer-controlled CHI 660 (CHI, Shanghai) electrochemistry workstation. Working electrodes were gold-coated glass chips (5 mm \times 20 mm size), used in conjunction with a Pt auxiliary electrode, and a saturated Ag/AgCl reference electrode. The electrodes for electrochemical experiments were purchased from Shanghai Chenhua Co., Ltd., China. Electrochemical activation of TMS/MGF was realized by potential scanning from -0.1 to 0.9 V with a scan rate of 100 mV s⁻¹ in 0.1 M H₂SO₄. SERS spectra were obtained using a confocal microscope Raman system (DXR ThermoFisher Confocal Raman Spectra) equipped with a Renishaw CCD camera. The microscope attachment is based on a system with a 50× long-working length objective, so that the objective will not be in contact with the electrolyte. A holographic notch filter was used to filter the exciting line, and two selective holographic gratings (1200 g/mm, 2400 g/mm) were employed, depending

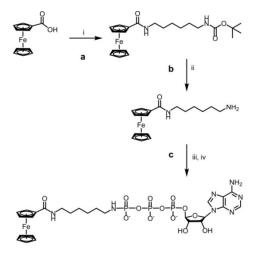
on the spectral resolution required. The exciting wavelength was 633 nm. ¹H, ¹³C, ³¹P NMR experiments were performed on a Bruker Avance 500 MHz spectrometer and chemical shifts were referenced to the residue DMSO (2.50 ppm for ¹H and 39.52 ppm for ¹³C) and H₂O (4.79 ppm). Mass spectrometry was carried out using a Perkin Elmer-Sciex API 365 instrument. SEM images were obtained using a Hitachi S4800 microscope at the accelerating voltage of 5 kV.

Preparation of Au/ITO. The Au-ITOs were prepared on glass chips (5 mm \times 20 mm size) by thermal evaporation of a 500 Å gold layer (99.99%) with a 50 Å chromium as the adhesion layer. The chips were carefully pre-cleaned by repeated sonication in Millipore H₂O and ethanol for at least three times before preparation of the mesoporous gold film (MGF).

Preparation of MGF. Our approach is based on an electrochemical method, which has been practically used for the general preparation of continuous metallic films.⁸ In a typical synthesis, 10 mg of polystyrene-b-poly(oxyethylene) (PS-b-PEO) was dissolved in 3 mL THF completely at 40 $^{\circ}$ C and then 1.5 mL ethanol was added to the mixed solution. An aqueous solution of HAuCl₄ (5 mM) was added slowly to the clear PS-b-PEO solution and spherical micelles were formed by the presence of water. Gentle stirring for 30 min at room temperature is used to make sure that the dissolved Au species were well incorporated into the exterior PEO region of the micelles. Finally, a transparent bright-yellow colored electrolyte (pH 2.5) was obtained and directly used for electrodeposition. Electrochemical deposition from the precursor solutions was carried out by using an electrochemical machine (CHI 660D electrochemical station, CH Instrument, Shanghai) with a conventional three-electrode system, including a platinum wire as a counter electrode, an Ag/AgCl as a reference electrode and various conductive substrate as working electrode. The typical conducting substrate used is Au–ITO with a representative size of 0.45 cm² (0.3 cm \times 1.5 cm), which was fabricated by a dicing cutter. The optimal electrodeposition of Au was carried out at a constant potential of - 0.5 V (versus Ag/AgCl) for 1000 s without stirring at room temperature. During the electrodeposition, a stable current was detected for the Au reduction. After the Au deposition, the micelles (used as soft-templates) were thoroughly removed by ultraviolet-ozone cleaner or low-powered O2 plasma treatment. Calcination of the films in air, which has been commonly used for complete removal of organic templates in mesoporous metal oxide films, led to removal of the templates from the films, but the mesoporous structures collapsed through the grain growth of Au.

Synthesis of Fc-ATP

Fc-ATP was prepared according to previous report.⁹ The synthesis route was illustrated as shown in Scheme S1.



Scheme S1. Preparation routes of the Fc-ATP (Compound 3). Reagents and conditions: (i) TEA and HBTU; (ii) TFA and DCM, (iii) ATP, dicyclohexylcarrbodiimide (DCC), and DMF; (iv) compound 2 and methanol.

Preparation of TMS/MGF. TMS/MGF were constructed by immersing freshly TMS solution in EtOH and 2-(2-hydroxyethoxy) ethylethanethioate (DEG) in 10^{-4} mol L⁻¹ total mixed thiol solution (1 mol % TMS and 99 mol % DEG) for 48 h at room temperature.

Detection limit. This Detection limit value is calculated according to LOD = 3 N/S, where N represents the standard deviation of the I_{320}/I_{1690} values obtained at the H2-cRGD/TMS/MGF within 20 measurements in the absence of HER2, and S represents the slope of the linear equation for HER2.

Cell Culture. Human breast cancer cells (SK-BR-3, MCF-7), human ovarian cancer cells (SK-OV-3) and Hela cells were purchased from the Chinese Academy of Medical Sciences& Perking Union Medical college (Beijing, China). SK-OV-3 represents one type of ovarian adenocarcinoma cells with the overexpressed HER2, Hela cells and MCF-7 cells are HER2-negative cells.^{5b,S1} They were cultured in a flask containing complete medium supplemented with 10% heat-inactivated fetal bovine serum and 1% streptomycin/penicillin. The flask was placed in a humidified atmosphere with 5% CO₂ at 37 °C in a cell culture incubator. The media was replaced once every three days. Cells were cultured to 80-90% confluence before harvest. During harvest, the cells were washed twice with PBS buffer followed by trypsinization with 1 mL of trypsin-EDTA solution (0.25 w/v % trypsin, 2.5 g L⁻¹ EDTA) at 37 °C for 5 min to detach the cells from the flask. The trypsin was neutralized by adding 5 mL of fresh supplemented RPMI-1640 medium, and the harvested cells in medium suspension was transferred into a centrifuge tube and centrifuged at 1000 rpm for 5 min. The supernatant was discarded and the cells were re-suspended in fresh medium. White blood cells (WBCs) were obtained from lysed human blood.^{3c} They were stained with 1,1-dioctadecy 1-3,3,3,3-tetramethylindocarbocyanine and kept at a concentration of 10^6 cells/mL in PBS, respectively.

Cells were cultured on gold-coated glass slides in a 12-well plate under normal culture conditions. After a 48 h incubation period, medium was removed. Then cells were fixed with 4% polyfluoroalkoxy for 15 min and incubated with 0.01% Rhodamine for 5 min, followed by two washes with PBS. The mesoporous gold-coated ITO were viewed by confocal fluorescence microscopy. Then, the culture medium was removed, and cells was washed three times with PBS to remove unbonded samples. Finally, the dish was put on the microscope stage for cell imaging and spectroscopic detection. Confocal fluorescence images were taken using a laser scanning confocal microscope LEICA TCS SP8.

Isolation of CTCs from Cancer Patients' Whole Blood. The EDTA anticoagulated whole blood samples (4 cancer patients, including 3 breast cancer, 1 gastric cancer and 3 healthy individuals) were obtained from Zhongshan Hospital of Fudan University and used without any pretreatment. Before samples processing, the H2-cRGD/TMS/MGF was blocked by 5% BSA and 0.2% Tween-

20, to avoid the nonspecific binding of the blood cells onto the skeleton. Then 1 mL of whole blood samples were incubated with H2-cRGD/TMS/MGF for 30 min at 4 °C and washed three times with PBS to remove unbonded samples. Next, the H2-cRGD/TMS/MGF was incubated with 0.01% Rhodamine for 5 min and followed by two washes with PBS to remove the extra fluorescent dyes. Finally, the H2-cRGD/TMS/MGF electrode was observed under fluorescent microscope.

2. Pore Depth distributions of MGF

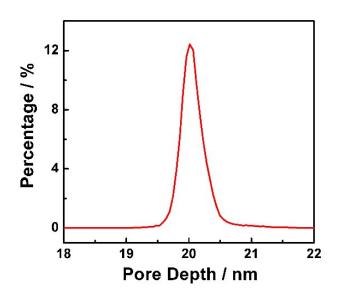


Fig. S1 (a) Pore Depth distributions of MGF (n = 200) from AFM image as shown in Fig. 1e.

3. Characterization of TMS

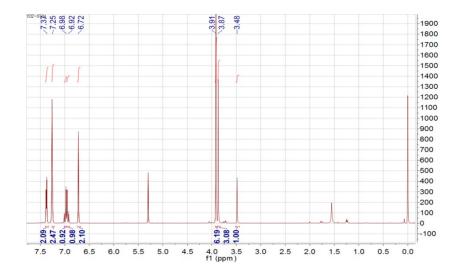


Fig. S2 ¹H NMR spectrum (500 MHz) of TMS molecule in CDCl₃.

¹H NMR (500 MHz, CDCl₃): δ =3.48 (s, 1 H; SH), 3.87 (s, 3 H; OCH₃), 3.91 (s, 6 H; OCH₃), 6.72 (s, 2 H; H2', H6'), 6.92 (d, *J*=16.0 Hz, 1 H; H8), 6.98 (d, *J*=16.0 Hz, 1 H; H7), 7.25 (d, *J*=8.0 Hz, 2 H; H3, H5), 7.37 ppm (d, *J*=8.0 Hz, 2 H; H2, H6); HRMS (ESI): m/z calc.C₁₇H₁₈O₃S for [M+H]+: 303.1; found: 303.1.

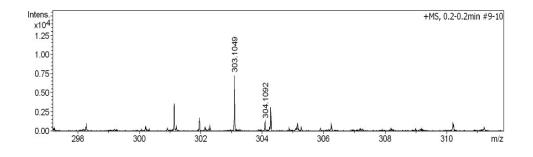


Fig. S3 HRMS (ESI) for TMS.

4. Characterization of Fc-ATP

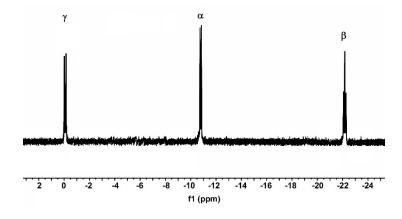


Fig. S4 31 P NMR spectrum (400 MHz) of Fc-ATP in D₂O.

³¹P{¹H}-NMR (δ , D₂O): -0.07(γ) d, J = 21.1 Hz; -10.76(α) d, J = 19.9 Hz; -22.14(β) t, J = 19.9 Hz. [H4·M](ESI⁺) m/z: calc. for C₂₇H₃₉FeN₇O₁₃P₃: 818.1; found: 818.2.

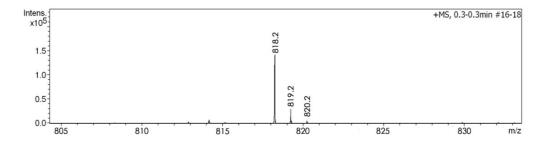


Fig. S5 [H4·M] (ESI+) Mass Spectrum for Fc-ATP molecule.

5. Electrochemical characterization

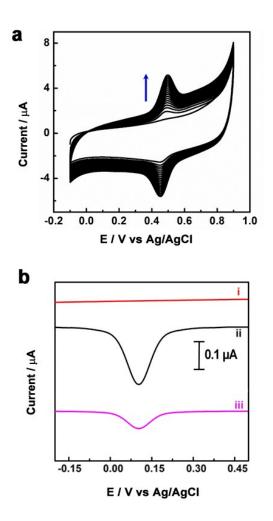


Fig. S6 (a) CVs obtained at TMS/MGF electrode in 0.5 M H_2SO_4 . Scan rate: 100 mV s⁻¹; (b) DPVs obtained at (i) MGF/Au, (ii) TMS/MGF after the first electrochemical activation, and (iii) formed H2-cRGD/TMS/MGF in 10 mM PBS (pH 7.4) after the second electrochemical activation.

6. SERS spectra characterization

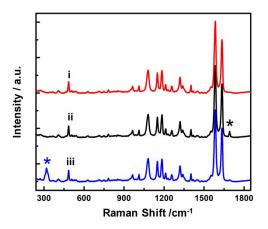


Fig. S7 SERS spectra obtained at (i) electrochemically activated TMS/MGF, (ii) H2-cRGD/TMS/MGF, (iii) H2-cRGD/TMS/MGF in 10 mM PBS (pH 7.4) in the presence of 50 nM Fc-ATP and 40 fg mL⁻¹ HER2.

7. Concentration optimization of Fc-ATP

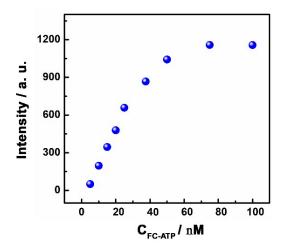


Fig. S8 Responses in peak intensity at 1690 cm⁻¹ (I_{1690}) in Raman spectra with the addition of different concentrations of Fc-ATP and 40 fg mL⁻¹ HER2.

8. Response time for detection of HER2

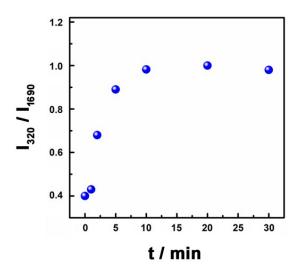


Fig. S9 I_{320}/I_{1690} values obtained at H2-cRGD/TMS/MGF substrate in 10 mM PBS (pH 7.4) containing 50 nM Fc-ATP and 40 fg mL⁻¹ HER2.

9. Fluorescence microscope images of SK-OV-3, MCF-7 and Hela cells captured at H2-cRGD/TMS/MGF platform.

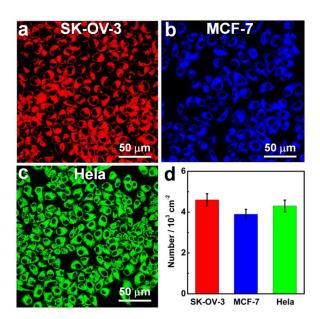


Fig. S10 (a-c) Fluorescence microscope images of SK-OV-3, MCF-7 and Hela cells captured by H2-cRGD/TMS/MGF. (d) Cell capture efficiencies for SK-OV-3, MCF-7 and Hela cells at H2-cRGD/TMS/MGF. Error bars equal to the standard deviation.

10. Fluorescence microscope images of White blood cells

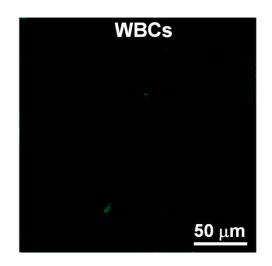


Fig. S11 Fluorescence microscope images of White blood cells captured by H2-cRGD/TMS/MGF.

11. Selectivity

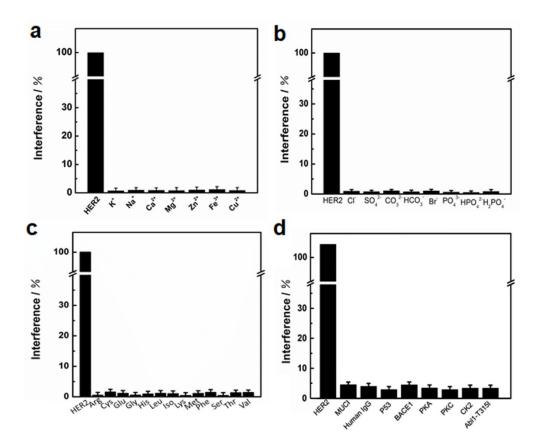


Fig. S12 Interference on peak intensity ratio (I_{320}/I_{1690}) obtained at 320 cm⁻¹and 1690 cm⁻¹ in addition of (a) cations: (i) 10 fg mL⁻¹ HER2, (ii) K⁺ (150 mM), (iii) Na⁺ (150 mM), (iv) Ca²⁺ (2.5 mM), (v) Mg²⁺ (2.5 mM), (vi) Zn²⁺ (1 mM), (vii) Fe³⁺ (10 µM), (viii) Cu²⁺ (10 µM); (b) Anions: (i) 10 fg mL⁻¹ HER2 (ii) Cl⁻, (iii) SO₄²⁻, (iv) CO₃²⁻, (v) HCO₃⁻, (vi) Br⁻, (vii) PO₄³⁻, (viii) HPO₄²⁻, (ix) H₂PO₄⁻ (from ii to ix concentrations are 100 µM); (c) amino acids: (i) 10 fg mL⁻¹ HER2, (ii) Arg, (iii) Cys, (iv) Glu, (v) Gly, (vi) His, (vii) Leu, (viii) Iso, (ix) Lys, (x) Met, (xi) Phe, (xii) Ser, (xii) Thr, (xiv) Val (10 µM from ii to xiv) and (d) proteins: (i) 10 fg mL⁻¹ HER2, (ii) MUC1 (1 ng mL⁻¹), (iii) human IgG (10 µg mL⁻¹), (iv) p53 (0.1 ng mL⁻¹), (v) BACE1 (1 U mL⁻¹), (vi) PKA (1 U mL⁻¹), (vii) PKC (1 U mL⁻¹), (viii) Abl1-T3151 (1 ng mL⁻¹). Herein, the interference % was estimated according to (I_{320}/I_{1690})^{*}/(I_{320}/I_{1690}), in which (I_{320}/I_{1690})^{*} means the I_{320}/I_{1690} after addition of various interferences.

12. Anti-biofouling performance of H2-cRGD/TMS/MGF platform

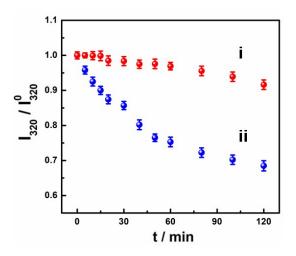


Fig. S13 I_{320}/I_{320}^0 values obtained at (i) H2-cRGD/TMS/MGF platform and (ii) H2-cRGD/TMS/ Au in SK-BR-3 cell lysate containing 40 fg mL⁻¹ HER2 under different time.

13. Comparison of capture efficiency in PBS and whole blood

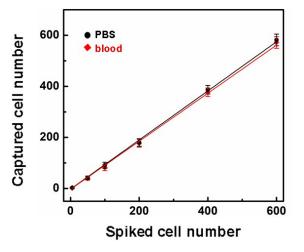


Fig. S14 Comparison of capture efficiency for SK-BR-3 cells in 10 mM PBS buffer (pH 7.4) and whole blood using H2-cRGD/TMS/MGF platform.

14. ELASA kit detection of HER2 activity from different passages of SK-BR-3 cells

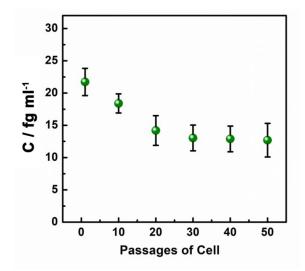


Fig. 15 HER2 activity detected by ELISA kit for cell extracts collected from different passages of SK-BR-3 cells.

15. SERS images of SK-OV-3, MCF-7 and Hela cells captured by H2cRGD/TMS/MGF platform

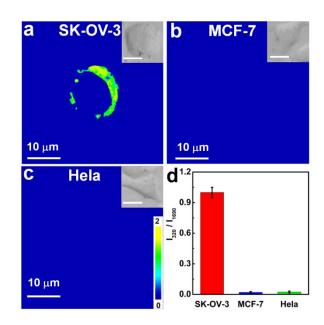


Fig. S16 (a-c) The SERS images of SK-OV-3, MCF-7 and Hela cells captured at H2cRGD/TMS/MGF. Inset: their corresponding dark field images of cells, the scale bar is 5 μ m. (d) The averaged I₃₂₀/I₁₆₉₀ value for SK-BR-3, MCF-7 and Hela cells captured by H2cRGD/TMS/MGF. Error bars equal to the standard deviation.

Sample no.	Туре	Number of CTCs	$C_{\text{HER2-b}} / \text{ng ml}^{-1}$	ELISA kit / ng ml ⁻¹
			(n = 3)	(n = 3)
1	Normal	0	0	-
2	Normal	0	0	-
3	Normal	0	0	-
4	Breast	100	22.18 ± 1.52	23.62 ± 4.85
5	Breast	255	35.75 ± 2.31	38.05 ± 4.23
6	Breast	280	27.09 ± 3.25	29.10 ± 3.86
7	Gastric	457	48.05 ± 2.84	50.00 ± 6.96

Table S1. Comparison of SERS detection at H2-cRGD/TMS/MGF with ELISA Kit

 for detection of HER2 in human serum.

* The level of HER2 in whole blood was estimated through the following equation:

 $C_{\text{HER2-b}} = C_{\text{HER2-c}} \times N$

in which $C_{\text{HER2-b}}$ represents the total HER2 activity in the blood sample, $C_{\text{HER2-c}}$ is the HER2 activity obtained from averaged values for 10 measurements at one CTC. N is the number of CTCs captured at H2-cRGD/TMS/MGF. n = 3 represents three blood samples from the same individual.

References

S1. D. W. Rusnak, K. J. Alligood, R. J. Mullin, G. M. Spehar, C. Arenas-Elliott, A. M. Martin, Y. Degenhardt, S. K. Rudolph, T. F., Jr. Haws, B. L. Hudson-Curtis, T. M. Gilmer, *Cell Proliferat*. 2007, 40, 580.