1	Supporting Information
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3	Single-Particle Assay of Poly(ADP-ribose) Polymerase-1 Activity with
4	Dark-Field Optical Microscopy
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37	Materials. All oligonucleotides listed in Table S1 were sythesized by Shanghai
38	Sangon biotechnology (Shanghai, China). Bis(p-sulfonatophenyl)phenylphosphine
39	(BSPP) and nicotinamide adenine dinucleotide (NAD ⁺) were purchased from
40	Sigma-Aldrich (Shanghai, China). (3-aminopropyl)-triethoxysilane (APTES) was
41	purchased from Macklin Biochemical (Shanghai, China).
42	Hexadecyltrimethylammonium bromide (CTAB) was obtained from J&K Scientific
43	Ltd. (Beijing, China). Trisodium citrate, chloroauric acid tetrahydrate (HAuCl ₄ ·4H ₂ O)
44	and polyvinylpyrrolidone (PVP) were purchased from Sinopharm Chemical Reagent
45	(Shanghai, China). Gold chloride trihydrate (HAuCl ₄ \cdot 3H ₂ O) was obtained from
46	Energy Chemical (Shanghai, China). Human PARP-1 was obtained from Trevigen
47	(Wuhan, China). Dulbecco's Modified Eagle Medium (DMEM) medium, RPMI-1640
48	medium, fetal bovine serum (FBS) and
48 49	medium, fetal bovine serum (FBS) and 3-(4,5-dimethylthiazol-2-yl)-2-diphenyltetrazolium bromide (MTT) were all from
48 49 50	medium,fetalbovineserum(FBS)and3-(4,5-dimethylthiazol-2-yl)-2-diphenyltetrazoliumbromide(MTT)wereall fromKeyGen Biotech (Nanjing, China). The formation of PAR catalyzed by PARP-1 was
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48 49 50 51 52	medium,fetalbovineserum(FBS)and3-(4,5-dimethylthiazol-2-yl)-2-diphenyltetrazoliumbromide(MTT)were all fromKeyGen Biotech (Nanjing, China). The formation of PAR catalyzed by PARP-1 wasoperated in the reaction buffer solution (R-buffer) composed of 50 mM Tris-HCl, 2mM MgCl2, 50 µM Zn(OAc)2 and 50 mM KCl (pH 7.4). Ultrapure water (18.2
 48 49 50 51 52 53 	medium, fetal bovine serum (FBS) and 3-(4,5-dimethyltazol-2-yl)-2-diphenyltetrazolium bromide (MTT) were all from KeyGen Biotech (Nanjing, China). The formation of PAR catalyzed by PARP-1 was operated in the reaction buffer solution (R-buffer) composed of 50 mM Tris-HCl, 2 mM MgCl ₂ , 50 µM Zn(OAc) ₂ and 50 mM KCl (pH 7.4). Ultrapure water (18.2 MΩ·cm at 25 °C, Thermo Scientific, USA) was used in the whole assay. All other
 48 49 50 51 52 53 54 	medium, fetal bovine serum (FBS) and 3-(4,5-dimethylthiazol-2-yl)-2-diphenyltetrazolium bromide (MTT) were all from KeyGen Biotech (Nanjing, China). The formation of PAR catalyzed by PARP-1 was operated in the reaction buffer solution (R-buffer) composed of 50 mM Tris-HCl, 2 mM MgCl2, 50 µM Zn(OAc)2 and 50 mM KCl (pH 7.4). Ultrapure water (18.2) MQ·cm at 25 °C, Thermo Scientific, USA) was used in the value assay. All other reagents were of analytical grade and needed no further treatment. Mark
 48 49 50 51 52 53 54 55 	medium, fetal bovine serum (FBS) and 3-(4,5-dimethyliazol-2-yl)-2-diphenyltetrazolum bromide (MTT) were all from KeyGen Biotech (Nanjing, China). The formation of PAR catalyzed by PARP-1 was operated in the reaction buffer) composed of 50 mM Tris-TCl, 2 mM MgCl2, 50 µM Zn(OAc)2 and 50 mM KCl (pH 7.4). Ultrapure water (18.2) MQ·cm at 25 °C, Thermo Scientific, USA) was used in the vhole assay. All other reagents were of analytical grade and needed no further treatment. Apparatus. Transmission electron microscopy (TEM) images were carried out with a
 48 49 50 51 52 53 54 55 56 	medium, fetal bovine serum (FBS) and 3-(4,5-dimethyliazol-2-yl)-2-diphenyltetrazolium bromide (MTT) were all from KeyGen Biotech (Nanjing, China). The formation of PAR catalyzed by PARP-1 was operated in the reaction buffer solution (R-buffer) composed of 50 mM Tris-HCl, 2 mM MgCl2, 50 µM Zn(OAc)2 and 50 mM KCl (pH 7.4). Ultrapure water (18.2) MΩ·cm at 25 °C, Thermo Scientific, USA) was used in the whole assay. All other reagents were of analytical grade and needed not further treatment. Apparatus. Transmission electron microscopy (TEM) images were carried out with a Cenai G2 20 (FEI, Czech Republic). Scanning electron microscopy (SEM) images Serue (SEM) images Serue (SEM) images

absorption spectra were recorded with UV-visible spectrometer (Shimadzu UV-2450,

59	Kyoto, Japan). Zeta potentials of each sample were all measured by NanoBrook Omni
60	Zeta potential analyzer (Brookheaven, USA) at 25 °C and measurements were
61	performed at least three times after diluted by deionized (DI) water. An inverted
62	optical microscopy (Eclipse Ti-E, Nikon, Japan) equipped with a dark field condenser
63	(0.8 < NA < 0.95) and a 60X objective lens (NA = 0.7) was used for dark-field
64	spectroscopic experiments. White light source (100 W halogen lamp) through
65	condenser was focused onto the sample and excited the nanoprobes to generate
66	plasmon resonance scattering light. After collected by a 60X objective and captured
67	by a true color digital camera (Nikon digital sight DS-Ri1, Japan), the scattering light
68	was split by a monochromator (Acton, P-2300, Princeton Instruments, USA) equipped
69	with a grating (grating density: 50 g·mm ⁻¹ ; blazed wavelength: 600 nm). Then a
70	spectrometer charge-coupled device (CCD) (PyLoN, Princeton Instruments, USA)
71	was utilized to record the scattering spectra. In order to avoid interference and ensure
72	accuracy, all scattering spectra of the nanoprobes were corrected by subtracting the
73	background spectrum generated by the instrument itself.

Synthesis of Gold Nanoparticles. The synthesis process of AuNPs with size of 13 nm (Au₁₃) was according to trisodium citrate-based classic reduction method. In brief, an aqueous solution of HAuCl₄·4H₂O (1 mM, 50 mL) was poured into the round-bottom flask and brought to a vigorous boil while stirring. Once the gold solution started to reflux vigorously, trisodium citrate solution (38.8 mM, 5 mL) was added rapidly into HAuCl₄ solution. The color of the mixture solution changed from yellow to clear, to black, to purple and finally to wine red throughout the whole 81 reaction process. Fifteen minutes later, the heating source was removed and the 82 obtained solution was allowed to cool to room temperature (RT). Finally, the colloidal 83 solution was stored in a brown glass bottle at 4 °C for further use.

84 The seed-growth method was used for preparation of Au₅₀. Briefly, 1 mL of the above prepared Au₁₃ (used as seed), 25 mL of DI water, 360 µL of 0.2 M 85 NH₂OH·HCl solution and 300 µL of 1 % w/v PVP (44000-54000 MW) were mixed in 86 a 50 mL round-bottom flask in sequence. Then 8 mL of 0.1 wt % HAuCl₄ was 87 injected dropwise into the mixture within 30 min under violent stirring at RT. The 88 89 reaction was stopped after another 30 min. Finally, the obtained Au₅₀ was stored in a brown glass bottle at 4 °C for further use. The concentration of the Au₅₀ solution was 90 estimated to be 0.1 nM, which was calculated by the Lambert-Beer law with the 91 extinction coefficient of $1.5 \times 10^{10} \cdot M^{-1} \cdot cm^{-1}$.¹ 92

Extraction of Cytoplasm and Nucleus of Cells. The extractive cytoplasm and 93 nucleus of two kinds of cancer cells (A2780 and MCF-7) and normal cells (IOSE80) 94 were obtained according to the previously reported method.⁷ Firstly, A2780 cells were 95 washed by $1 \times PBS$ (pH 7.4) one time or treated by ethylenediaminetetraacetic acid 96 (EDTA). Then the obtained cells were centrifuged with discarding supernatant. 97 Subsequently the cytoplasmic protein and nuclear extraction reagent were added 98 separately followed by vigorous shaking. The resulting mixture was centrifuged at 99 12000 rpm at 4 °C for 5 min. Finally, the obtained supernatants of cytoplasm and 100 nuclei were stored respectively in precooling centrifuge tubes at -80 °C for further use. 101 Cytoplasm and nucleus protein from MCF-7 and IOSE80 were also extracted on the 102

103 basis of our previous study.

Preparation of Au₅₀-dsDNA-aptamer. Firstly, excess BSPP was added to 1 mL of 104 105 Au₅₀ solution. And the mixture was incubated at RT overnight under stirring, which replaced the citrate ligands on the surface of Au₅₀ with BSPP ligands and stabilized 106 107 AuNPs at high ionic strength in the process of DNA modification. Then the mixture 108 was centrifuged at 8000 rpm for 15 min to collect the precipitation which was redispersed in 1 mL of DI water. 50 µL of 10 mM BSPP was then added to the 109 solution and shaked uniformly to obtain Au₅₀-BSPP. Secondly, the resulting 110 111 BSPP-protected Au₅₀ was mixed with 30 µL of 1 µM specific-DNA (s-DNA) and 30 µL of 1 µM AS1411 aptamer, followed by gently rocking at RT for 12 h. For 112 113 enhancing the salt resistance of Au₅₀, 1 M NaCl was introduced every 3 h until the 114 final concentration up to 150 mM. After that, 18 µL of 10 µM thiol-polyethylene glycol 800 (PEG-800) was added to the above Au₅₀ solution with incubation for 1 h to 115 block nonspecific binding sites of AuNPs. Thirdly, 30 µL of 1 µM complementary 116 DNA (c-DNA) was added with gentle vortexes for 2 h to realize the modification of 117 activated dsDNA on the Au₅₀ surface. Finally, the Au₅₀-dsDNA-aptamer was collected 118 by centrifugation at 8000 rpm for 15 min. 119

Preparation of Au_8 -aptamer. Firstly, 30 µL of 1 µM AS1411 aptamer was added to 121 1 mL of Au_8 solution, followed by gently rocking at RT for 12 h. Then, the 122 Au₈-aptamer was collected by centrifugation at 13000 rpm for 15 min.



Figure S1. Zeta potentials of individual Au₅₀ (A), Au₅₀-dsDNA (B) and Au₅₀-dsDNA@PAR (C). 5.0 mU PARP-1 was used for forming Au₅₀-dsDNA@PAR. The error bars represent standard deviations of three repetitive measurements.



Figure S2. (A) Dark-field images of Au_{50} -dsDNA without Au_8 . Real-time dark-field images of Au_{50} -dsDNA@PAR incubated with Au_8 for different times: (B) 0, (C) 5, (D) 10, (E) 20 and (F) 30 min. 5.0 mU PARP-1 was used for the optimization.



Figure S3. Dark-field images of control assay without PARP-1: (A) Au_{50} , (B) Au_{50} -dsDNA, (C) Au_{50} -dsDNA + NAD⁺, then incubated with Au_8 . Representative dark-field images of Au_{50} -dsDNA incubated with NAD⁺, Au_8 and PARP-1 with different concentrations: (D) 0.2, (E) 0.3, (F) 1.0, (G) 2.0, (H) 5.0 and (I) 10 mU.



Figure S4. TEM images of MCF-7 cells (A) and IOSE80 cells (B) incubated with 10 pM Au_{50} -dsDNA-aptamer for 5 h and followed incubation with Au_8 -aptamer for 1 h at low magnification.

Name	Sequence and modifications (from 5' - 3')
Specific DNA	HS-(CH ₂) ₆ -CCCGTGCGTGCGCGAGTGAGTTG
Complementary DNA	CAACTCACTCGCGCACGCACGGG
AS1411 aptamer	HS-(CH ₂) ₆ -GGTGGTGGTGGTGGTGGTGGTGG

Method	System	Detection range	LOD	Reference
ELISA	PARP-1 antibody	Semiquantitative	-	2
PAGE	Clickable NAD ⁺	Qualitative	-	3
Fluorescence	NAD ⁺ analogue	Qualitative	-	4
Fluorescence	MnO ₂ quenched PFP	0.03-1.5 U	0.004 U	5
Fluorescence	TOTO-1	0.02-1.5 U	0.02 U	6
UV-Vis	AuNRs aggregation	0.05-1 U	0.006 U	7
Photoelectrochemistry	PFP	0.01-2 U	0.007 U	8
Nanochannel	Diffusion flux	0.05-1.5 U	0.006 U	9
Single nanoparticle				m1 · 1
scattering spectrum	AuNPs	0.2-10 m∪	-	I his work

Table S2. Comparison of analytical performances of various methods fordetermination of PARP-1.

Cancer colle	Found (U) in	Found (U) in
Cancel cells	Nuclei	Cytoplasm
A2780	0.3091	0.1313
MCF-7	0.2336	0.1358

Table S3. Detection of PARP-1 in cancer cells by the clinical ELISA method.

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