



Surface plasmon resonance assay for exosomes based on aptamer recognition and polydopamine-functionalized gold nanoparticles for signal amplification

Guofu Liao¹ · Xiaofeng Liu¹ · Xiaohai Yang¹ · Qing Wang¹ · Xiuhua Geng¹ · Liyuan Zou¹ · Yaqin Liu¹ · Shaoyuan Li¹ · Yan Zheng¹ · Kemin Wang¹

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Abstract

A novel surface plasmon resonance (SPR) strategy is introduced for the specific determination of exosomes based on aptamer recognition and polydopamine-functionalized gold nanoparticle (Au@PDA NP)-assisted signal amplification. Exosomes derived from hepatic carcinoma SMMC-7721 were selected as the model target. SMMC-7721 exosomes can be specifically captured by the aptamer ZY-sls that was complementary to the DNA tetrahedron probes (DTPs), and then the CD63 aptamer-linked Au@PDA NPs recognized SMMC-7721 exosomes for signal amplification. The DTPs were modified on a Au film for preventing Au deposition on the surface during the introduction of HAuCl₄, and PDA coated on the AuNPs was used to reduce HAuCl₄ in situ without any reductant assistance. It results in a further enhanced SPR signal. The assay can clearly distinguish SMMC-7721 exosomes from others (HepG2 exosomes, Bel-7404 exosomes, L02 exosomes, MCF-7 exosomes, and SW480 exosomes, respectively). SMMC-7721 exosomes are specifically determined as low as 5.6×10^5 particles mL⁻¹. The method has successfully achieved specific determination of SMMC-7721 exosomes even in 50% of human serum without any pretreatment.

Keywords SMMC-7721 exosomes · HAuCl₄ · DNA tetrahedron probes · Specificity · Au deposition

Introduction

Exosomes, nanoscale membrane-encapsulated extracellular vesicles with size ranging from 30 to 150 nm, are secreted by many mammalian cell [1, 2] and present in many different human biofluids [3]. In addition, exosomes carrying some biomolecules (such as RNA/DNA and proteins) can involve

the exchange of information between cells and affect the physiological functions of cells [1, 4, 5]. They are well considered as potential and non-invasive biomarkers for significant disease diagnosis [1, 5]. Therefore, it is very necessary to develop an effective method for determining related cancerous exosomes.

There are some researches focusing on the determination of exosomes, such as electrochemistry [6, 7], microfluidics [8, 9], fluorescence [10], surface-enhanced Raman scattering (SERS) [11, 12], and surface plasmon resonance (SPR) [13–16]. Most of these methods had to depend on antibodies so that they have some downsides. For example, preparation of antibody is tedious, expensive, and time-consuming. Besides, given that antibody is easy to denature towards extreme chemical environment, it is difficult for most of these methods to standardize. Aptamer, in addition to having an antibody-like function, has several advantages, such as ease of modification, excellent specificity, and high affinity [17–22]. Hence, aptamer has high potential as a replacement for antibody in the specific determination of cancerous exosomes. Wang et al. developed a portable electrochemical

Guofu Liao and Xiaofeng Liu contributed equally to this work.

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✉ Qing Wang
wwq99@hnu.edu.cn

✉ Kemin Wang
kminwang@hnu.edu.cn

¹ State Key Laboratory of Chemo/Biosensing and Chemometrics, College of Chemistry and Chemical Engineering, Key Laboratory for Bio-Nanotechnology and Molecular Engineering of Hunan Province, Hunan University, Changsha 410082, China

aptasensor for specific determination of hepatocellular exosomes [18]. Huang et al. introduced an aptasensor based on rolling circle amplification–assisted fluorescence signal amplification for specific determination of leukemia-derived exosomes [19]. Zhang et al. also reported electrogenerated chemiluminescence biosensors for MCF-7 exosome–specific determination based on Ti3C2 MXene nanosheets and two aptamers (EpCAM aptamer and CD63 aptamer) [20]. To our knowledge, the application of aptamers on SPR strategy to determine exosomes has so far been relatively rare. Our group has developed gold nanoparticle (AuNP)–amplified SPR biosensor using CD63 aptamer as recognition element for sensitive determination of MCF-7 exosomes [21]. Since CD63 is usually present on most exosomes, the SPR biosensor has poor specific determination of MCF-7 exosomes. Therefore, it is necessary to develop a novel method for specific determination of cancerous exosomes.

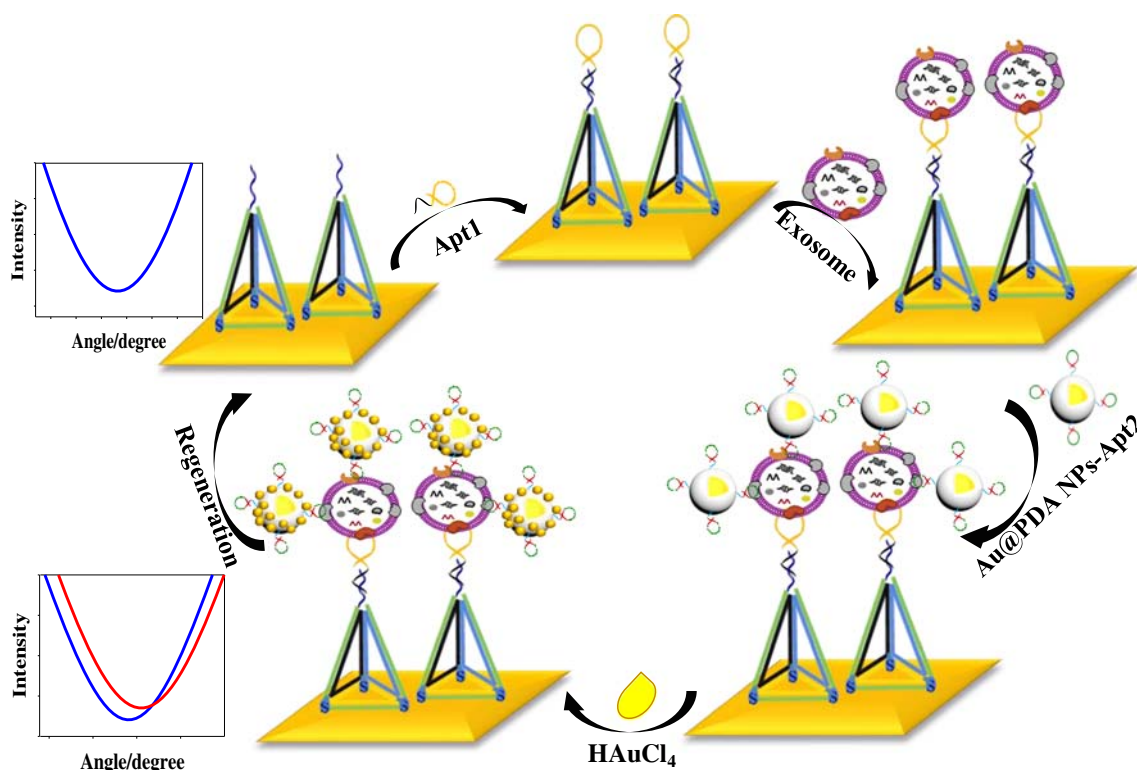
Herein, we first found that aptamer ZY-sl (Apt1), which was generated by the screening of human hepatic carcinoma (SMMC-7721) cells [23], can specifically bind SMMC-7721 exosomes using flow cytometry. Based on the finding, a novel SPR strategy was proposed for the specific determination of SMMC-7721 exosomes based on two aptamers (Apt1 and CD63 aptamer (Apt2)). The strategy is based on polydopamine-functionalized AuNP (Au@PDA NP)–assisted signal amplification, which consists of three main processes (Scheme 1). Firstly, the Apt1 that is complementary to DNA tetrahedron probes (DTPs) with efficient blocking ability and

anti-gold deposition [24–26] can capture specifically SMMC-7721 exosomes. Then, Apt2-linked Au@PDA NPs (Au@PDA NP-Apt2) recognize SMMC-7721 exosomes captured by Apt1 to assist signal amplification. Subsequently, PDA on the surface of AuNPs can reduce HAuCl_4 in situ without any reductant assistance when incubated with HAuCl_4 [27, 28], resulting in further enhancement of the SPR signal. This work can also broaden the potential for specific determination of other cancerous exosomes by replacing other specific aptamers.

Experimental section

Preparation and characterization of Au@PDA NP-Apt2

Au@PDA NP-Apt2 was prepared as shown in Scheme S-1 in the Electronic Supporting Material (ESM). Typically, Au@PDA NPs were formed by in situ polymerizations of dopamine (DA) on the surface of AuNPs under alkaline conditions [29, 30]. Next, the preparation of Au@PDA NP-Apt2 was achieved by covalent bond cross-linking between the $-\text{NH}_2$ on the PDA and the $-\text{COOH}$ on Apt2 [27, 31]. The detail experimental procedures are listed in the ESM. Then, Au@PDA NPs and Au@PDA NP-Apt2 were characterized by UV-vis spectroscopy (Shimadzu, Japan), Zetasizer Nano ZS90 (Malvern Instruments, U.K.), Fourier transform-infrared



Scheme 1 The principle of the SPR assay for determination of exosomes based on aptamer and Au@PDA NPs

(FT-IR) spectra (Brooke group), and transmission electron microscopy (TEM) (JEOL, Japan).

Exosome determination

SMMC-7721 exosomes were derived from human hepatic carcinoma (SMMC-7721) cell line that was purchased from Xiangya School of Medicine (Central South University, China). The specific determination of SMMC-7721 exosomes was conducted on a EC-SPR1010 device (Changchun Dingcheng Technology, China). The SPR chip was prepared following our previous work [21]. Firstly, DNA tetrahedral probes (DTPs) were assembled by referring to traditional simple denaturation and annealing processes [12, 13] (shown in ESM). Figure S-2 indicates that the DTPs are formed. Next, the DTPs (5.0 μM) were first incubated on Au film surface for 3 h at 25 °C. Figure S-3 suggests that DTPs are successfully modified on Au film. Then, the DTP-modified Au film surface was incubated with Apt1 (65 μL , 1.2 μM) for 40 min at 25 °C. Unbound Apt1 was repeatedly washed away with phosphate-buffered saline (10 mM $\text{KH}_2\text{PO}_4/\text{Na}_2\text{HPO}_4$, 137 mM NaCl, 2.7 mM KCl, pH 7.4), and the SPR spectrum was recorded. Next, SMMC-7721 exosomes were respectively introduced and incubated for 40 min at 25 °C, then rinsed thoroughly with phosphate-buffered saline and the SPR spectrum was recorded. Then, Au@PDA NP-Apt2 (65 μL , 4.2 nM) was introduced and incubated for 40 min at 25 °C, and the SPR spectrum was recorded after rinsing with phosphate-buffered saline thoroughly. Subsequently, HAuCl_4 (65 μL , 0.24 mM) was injected and reacted for 20 min at 25 °C, and then the SPR spectrum was recorded after rinsing repeatedly with phosphate-buffered saline. Finally, the regeneration solution (65 μL , 0.1% SDS, 10 mM NaOH) was added for about 5 min to regenerate the SPR chip.

Analysis of real samples

SMMC-7721 exosome in normal human serum was also determined using this assay. The human serum was collected from healthy volunteers of the Hospital of Hunan University. The samples were first obtained through adding different concentrations of SMMC-7721 exosomes in 50% human serum. Then, the above samples were respectively injected and incubated with DTP-Au film for 40 min. After rinsing thoroughly with phosphate-buffered saline, SPR spectrum was recorded. Subsequently, Au@PDA NP-Apt2 (65 μL , 4.2 nM) were introduced and incubated for 40 min. SPR spectrum was recorded after being washed thoroughly with phosphate-buffered saline. Next, HAuCl_4 (65 μL , 0.24 mM) was injected and reacted for 20 min. Afterward, SPR spectrum was recorded after rinsing repeatedly with phosphate-buffered saline. The concentration of the SMMC-7721 exosome in human serum

was determined according to the change of resonance angle. The SPR chip was regenerated using 0.1% SDS/10 mM NaOH for the determination of other samples.

Results and discussion

Choice of materials

Given that SMMC-7721 exosome is one of the markers of human therapeutic carcinoma [32], it is selected as the model target. The SMMC-7721 exosomes were characterized using nanoparticle tracking analysis (NTA), dynamic light scattering (DLS), TEM, and scanning electron microscopy (SEM), respectively. Figure S-4 indicates that exosomes can be successfully obtained.

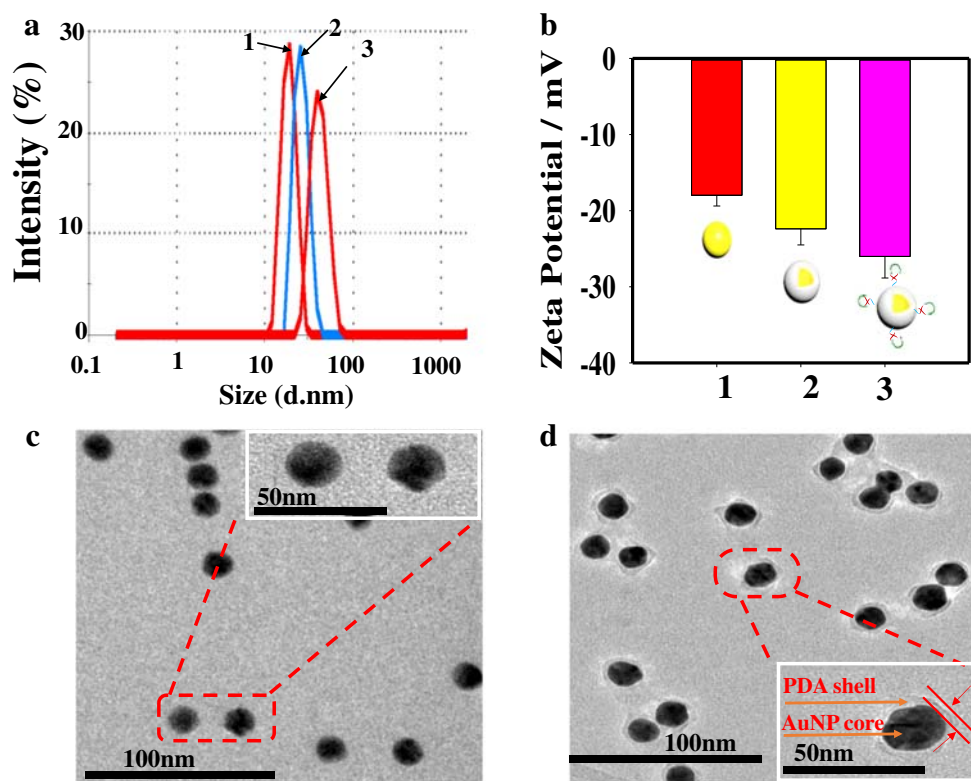
In our previous work [24], it was reported that DTP-Au film not only demonstrated superior antifouling property against protein and cell but also avoided Au deposition on the surface in the process of the catalytic growth of gold nanoparticles. Thus, DTP-Au film is used here for avoiding the Au deposition on the surface of the surface during the reduction of HAuCl_4 by PDA.

Since the good optical properties of AuNPs can enhance the SPR signal and PDA can reduce HAuCl_4 to form small AuNPs [27], Au@PDA NPs are selected here for signal amplification.

Characterization of Au@PDA NP-Apt2

The preparation of Au@PDA NP-Apt2 was confirmed by DLS, zeta potential, UV-vis absorption spectrum, and FT-IR spectra. As shown in Fig. 1(A), as the PDA and Apt2 are modified on the AuNPs respectively, the hydrodynamic size of the nanoparticles increases. As shown in Fig. 1(B), the zeta potential value decreases significantly, which is attributed to the shell of PDA and Apt2 attachment with a large amount of negative charge. Similarly, it also shows the maximum UV-vis absorption peak red shifts from 518 to 538 nm as PDA and Apt2 are modified on the AuNPs (Fig. S-5A). FT-IR profile is also used for the characterization of Au@PDA NP-Apt2. As shown in Fig. S-5B, the newly discovered absorption bands imply the presence of Au@PDA NP-Apt2 (detailed datums are showed in ESM). Simultaneously, the Au@PDA NPs also are confirmed by TEM images. Comparing with AuNPs (15.4 \pm 4.5 nm) (Fig. 1(C)), the core-shell structure of Au@PDA NPs (24.2 \pm 3.3 nm) is clearly visible under TEM image (Fig. 1(D)). In addition, the PDA layer (8.8 \pm 1.2 nm) is clearly displayed on the periphery of the AuNPs. The results show that the Au@PDA NPs and Au@PDA NP-Apt2 can be obtained.

Fig. 1 DLS profile (A) and zeta potential (B) of AuNPs (1); Au@PDA NPs (2) and Au@PDA NP-Apt2 (3). TEM images of AuNPs (C) and Au@PDA NPs (D). Inset: Magnified TEM image of the area marked



Investigation of the resistance of Au deposition using DTP-Au film

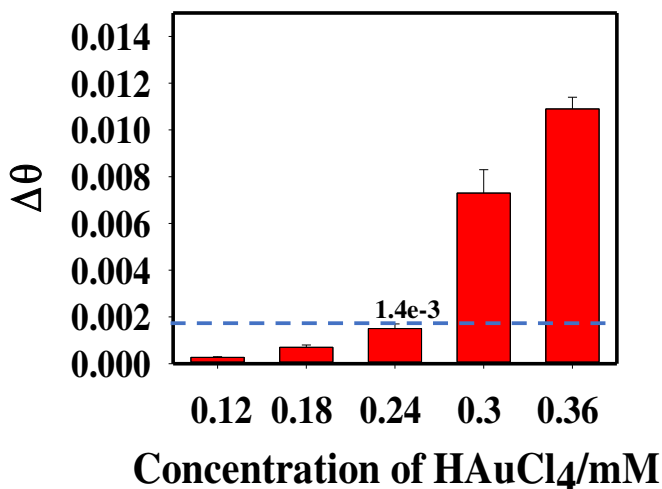
If HAuCl_4 solution is incubated on the Au film, obvious background signal can be observed due to Au deposition on the Au film (Fig. S-6). Thus, the effect of DTP-Au film on the resistance of Au deposition was explored using the EC-SPR1010 device. After different concentrations of HAuCl_4 solution were added and incubated for 20 min, the DTP-Au film was washed thoroughly with phosphate-buffered saline. Then the SPR response was recorded. Thus, the influence from the increase of the concentration of HAuCl_4 solution can be

Fig. 2 Investigation of the resistance of Au deposition using DTP-Au film. The blue dotted line indicated that the $\Delta\theta$ was 0.0015°

ignored. As shown in Fig. 2, the resonance angle change ($\Delta\theta$) increased with the increase of HAuCl_4 concentration. $\Delta\theta$ states a background signal which is caused by the Au deposition on the chip as different concentrations of HAuCl_4 solution are added. Given that the $\Delta\theta (> 0.0015^\circ)$ is acted as a signal [29], 0.24 mM HAuCl_4 is selected for the following experiment.

Effect of HAuCl_4 on Au@PDA NP-Apt2

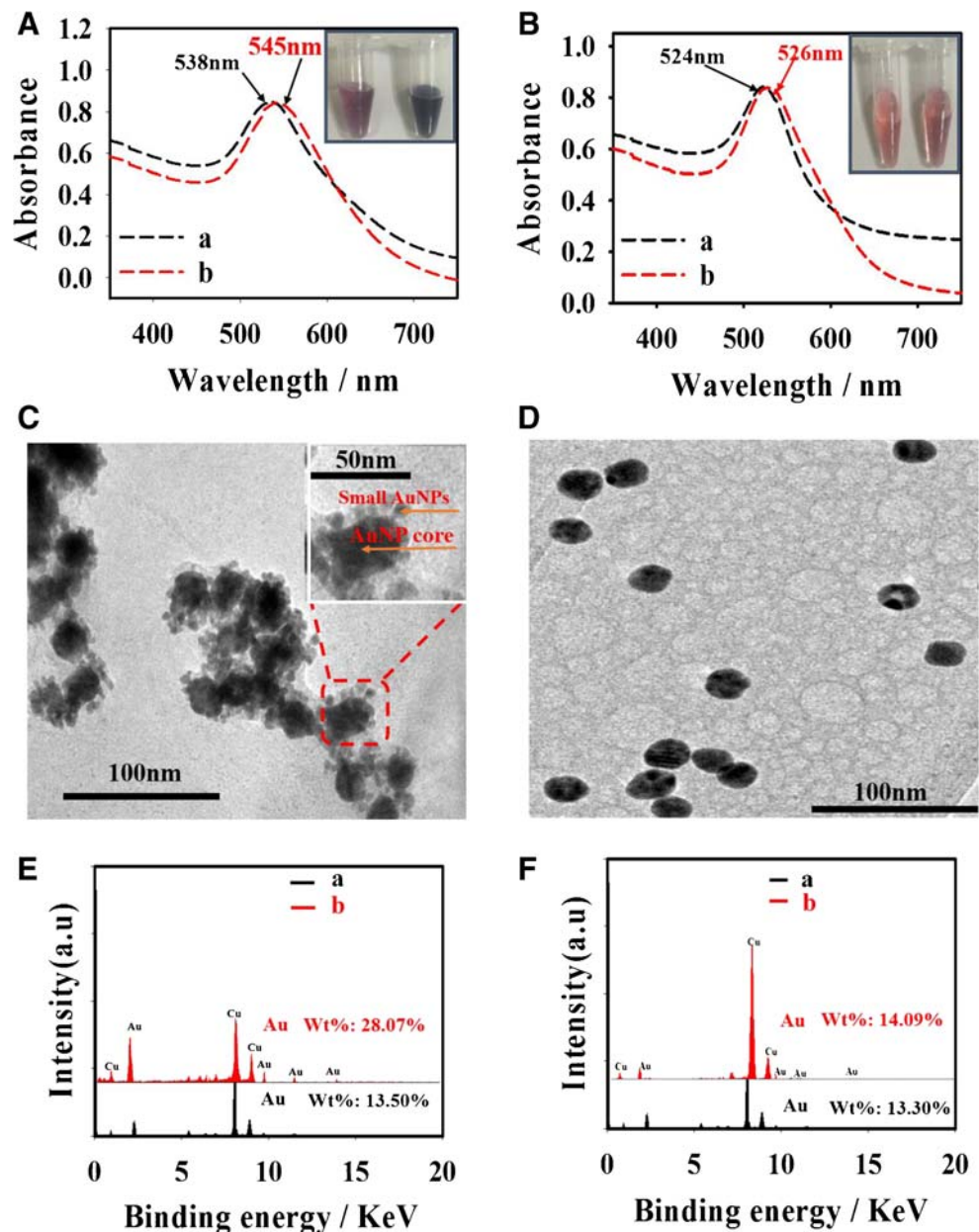
After DTP-Au film had good anti-gold deposition effect on HAuCl_4 (0.24 mM), the effect of HAuCl_4 (0.24 mM) on



Au@PDA NP-Apt2 was also investigated. Figure 3 shows that the maximum UV-vis absorption peak moves from 538 nm to 545 nm as Au@PDA NP-Apt2 was incubated with 0.24 mM H₂AuCl₄. Meanwhile, the color of the solution that incubated with 0.24 mM H₂AuCl₄ changes from dark purple to deep blue. For the AuNP-Apt2 without PDA functionalization (Fig. S-1), the shift in the UV-vis spectrum of AuNP-Apt2 is observed to be negligible after incubation with 0.24 mM H₂AuCl₄ (Fig. 3(B)). Meantime, the color of the solution hardly changes after incubation with 0.24 mM H₂AuCl₄. It is supposed that the size of Au@PDA NP-Apt2 is changed by the reduction of H₂AuCl₄ (0.24 mM) by PDA. Besides, the results are also further confirmed by TEM images. It is clearly

observed that many small nanoparticles were coated on the Au@PDA NPs after incubating with H₂AuCl₄ (Fig. 3(C)), while almost no change is observed for AuNP-Apt2 (Fig. 3(D)). It also suggests that H₂AuCl₄ is reduced by PDA on the Au@PDA NP surface. As shown in Fig. 3, the weight content of Au is increased from 13.5 to 28.07% by an energy dispersive spectrometer (EDS) profile (JEOL, Japan) (Fig. 3(E)), while a negligible change in the weight content of Au was also obtained by EDS profile (Fig. 3(F)). The results also demonstrate that the small AuNPs are deposited on the Au@PDA NP-Apt2 due to the reduction of H₂AuCl₄ by PDA. Therefore, it suggests that Au@PDA NP-Apt2 can provide a feasible platform for assisted SPR signal amplification.

Fig. 3 Effect of H₂AuCl₄ (0.24 mM) on Au@PDA NP-Apt2. The change of UV-vis absorption spectra of Au@PDA NP-Apt2 (A) and AuNP-Apt2 (B) after incubating with H₂AuCl₄. Inset: Photographs of nanoparticles before (left) and after (right) incubated with H₂AuCl₄. TEM images of Au@PDA NP-Apt2 (C) and AuNP-Apt2 (D) after incubating with H₂AuCl₄. EDS profiles of Au@PDA NP-Apt2 (E) and AuNP-Apt2 (F) after incubating with H₂AuCl₄. Wt% represents the weight content of Au



Investigation of the feasibility

The specific recognition of the SMMC-7721 exosomes by the aptamer ZY-sl_s (Apt1) is a prerequisite in the successful construction of the SPR assay. This was first investigated using flow cytometry (Scheme S-2). As shown in Fig. 4(A), the comparison between the fluorescence intensities shows that the experiment group is approximately 80-fold higher than the control group. It provides solid evidence of the specific recognition between the Apt1 and SMMC-7721 exosomes. Simultaneously, the specific recognition between the Apt1 and SMMC-7721 exosomes was further investigated using EC-SPR1010 device. As shown in Fig. 4(B), in experiment group I, SMMC-7721 exosomes are incubated with Apt1 assembled on DTP-Au film to obtain a significant $\Delta\theta$ ($0.0492^\circ \pm 0.0032$). However, control group II is also performed using random sequence1 instead of Apt1, and a low $\Delta\theta$ is observed in comparison with the experiment group I. In addition, HepG2 exosomes (III), Bel-7404 exosomes (IV), and L02 exosomes (V) were used for the control exosomes. The results show that $\Delta\theta$ induced by these control exosomes is negligible. Therefore, the results suggest that the SPR assay for specific determination of SMMC-7721 exosomes can be constructed using Apt1 as the recognition probe.

The feasibility of the assay was then studied. As shown in Fig. 4(C), when Apt1 was incubated with DTPs modified on the Au film, a tiny $\Delta\theta$ (0.0063°) is observed (curve 1 to curve

2). It demonstrates that the Apt1 is assembled on DTP-Au film through hybridization. Next, when SMMC-7721 exosomes were reacted with the Apt1 assembled on DTP-Au film, the SPR spectrum shifts from curve 2 to curve 3 and a clear $\Delta\theta$ (0.0492°) is obtained. It indicates that the SMMC-7721 exosomes are specifically captured by Apt1 assembled on DTP-Au film. Then, Au@PDA NP-Apt2 was injected; the SPR spectrum moves from curve 3 to curve 4 and a $\Delta\theta$ (0.0790°) is observed. It suggested that Au@PDA NP-Apt2 can enhance the SPR signal, implying that the Au@PDA NP-Apt2 are attached on the surface of SMMC-7721 exosomes through the recognition between Apt2 and SMMC-7721 exosomes. Furthermore, the binding between Au@PDA NP-Apt2 and exosomes is characterized using TEM. As shown in Fig. S-7 of the ESM, Au@PDA NPs can attach on the surface of exosomes via Apt2 recognition. Subsequently, HAuCl₄ (0.24 mM) were injected; the SPR spectrum moves from curve 4 to curve 5 and a $\Delta\theta$ (0.0644°) is observed. It implies that Au@PDA NP-Apt2 can reduce HAuCl₄ in situ to further significantly enhance the SPR signals. The results are also shown in Fig. 4(D); compared with Au@PDA NPs (curve a) and AuNP-Apt2 (curve b), a significant $\Delta\theta$ is obtained for Au@PDA NP-Apt2 after treatment with HAuCl₄ (curve c). Finally, after regeneration solution was added, the SPR spectrum moves from curve 5 to curve 6, and then $\Delta\theta$ substantially coincides with the curve 6 to curve 1, indicating that the chip can achieve regeneration. Hence, it is concluded that the assay

Fig. 4 The feasibility analysis of the method. Flow cytometry (A) and SPR (B) investigation of the specific recognition between Apt1 and SMMC-7721 exosomes. SPR response ($\Delta\theta$) of Apt1 recognizing SMMC-7721 exosomes (I); random sequence1 recognizing SMMC-7721 exosomes (II); Apt1 recognizing HepG2 exosomes (III); Apt1 recognizing Bel-7404 exosomes (IV); and Apt1 recognizing L02 exosomes (V). (C) SPR spectra of DTP-Au film (1); hybridization with Apt1 (2); recognizing SMMC-7721 exosomes (3); reaction with Au@PDA NP-Apt2 (4); reaction with HAuCl₄ (0.24 mM) (5); and regeneration (6). (D) Determination of target exosome using Au@PDA NPs (a), AuNP-Apt2 (b), and Au@PDA NP-Apt2 (c) as signal amplification. The final concentration of all exosomes was 6.2×10^6 particles mL⁻¹

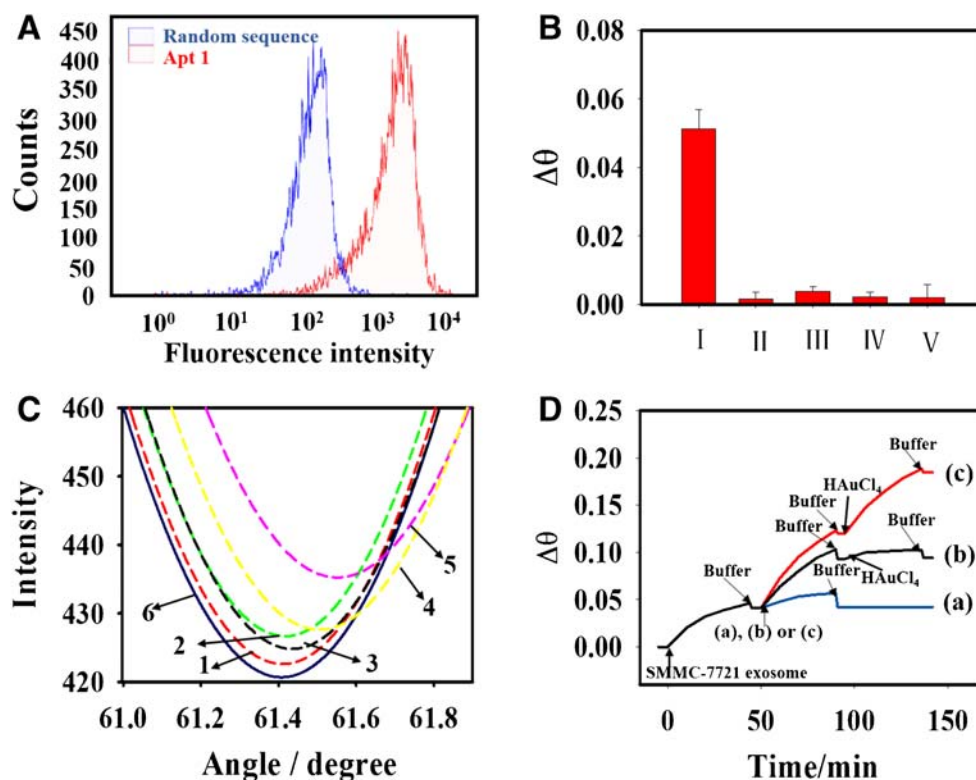
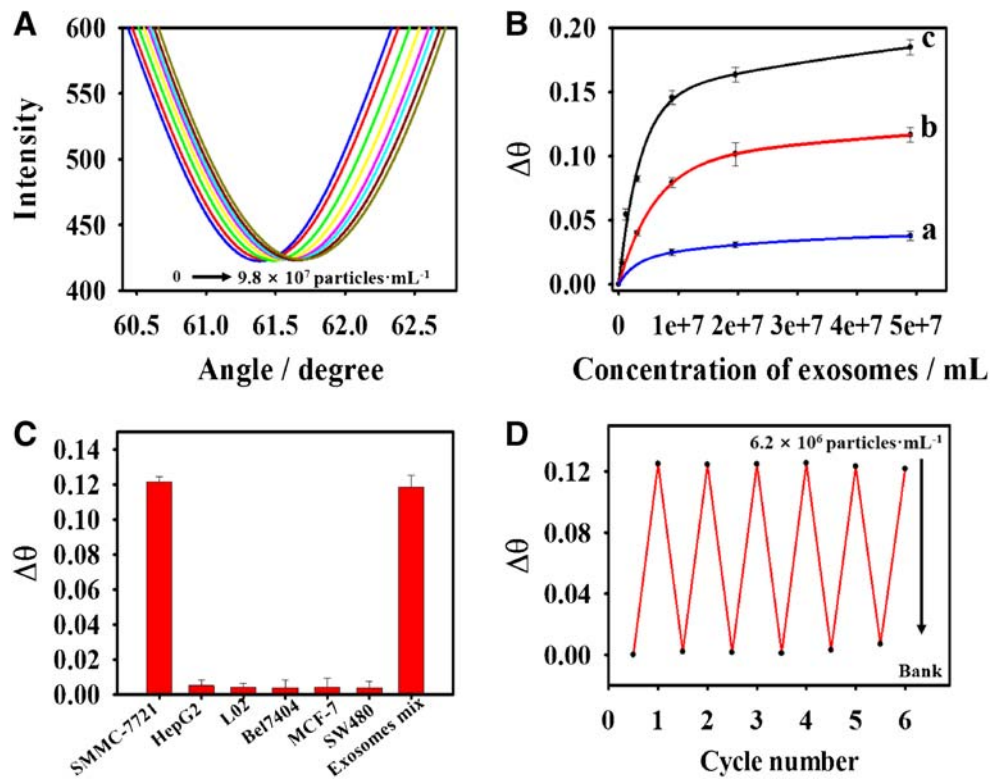


Fig. 5 Exosome determination using the SPR assay. (A) SPR spectra of different concentrations of SMMC-7721 exosomes. (B) The relationship between the $\Delta\theta$ and the concentration of SMMC-7721 exosomes by (a) direct determination, (b) Au@PDA NP-Apt2-amplified SPR assay, (c) Au@PDA NP-Apt2- and HAuCl₄-amplified SPR assay. (C) The selective investigation of the SPR assay. The final concentration of all exosomes was 6.2×10^6 particles mL⁻¹. (D) Reproducibility investigation of the SPR assay



is feasible for the specific determination of SMMC-7721 exosomes under the Au@PDA NP-Apt2-assisted amplification of the SPR signal.

Exosome determination

The sensitivity of the method was significantly affected by the concentration of Apt1 and the reaction time of SMMC-7721 exosomes. Thus, the effects of the concentration of Apt1 and the reaction time of SMMC-7721 exosomes were tested. The results indicate that the optimal conditions for the SPR assay are 1.2 μ M for Apt1 (Fig. S-8A) and 40 min for SMMC-7721 exosome reaction (Fig. S-8B) Next, the SPR biosensor is used to determine SMMC-7721 exosomes under optimal conditions. As shown in Fig. 5(A), the SPR resonance angle gradually increases as the concentration of SMMC-7721 exosomes increased (from 0 to 7.9×10^7 particles mL⁻¹). Figure 5 (B) illustrates the comparison between the $\Delta\theta$ response and the concentration of SMMC-7721 exosomes for

the three processes (direct determination of SMMC-7721 exosomes (curve a); Au@PDA NP-Apt2-amplified SPR assay (curve b); Au@PDA NP-Apt2- and HAuCl₄-amplified SPR assay (curve c)). The results show that the determination limit of direct assay, Au@PDA NP-Apt2-amplified assay, and Au@PDA NP-Apt2- and HAuCl₄-amplified assay is 7.0×10^7 particles mL⁻¹, 6.2×10^6 particles mL⁻¹, and 5.6×10^5 particles mL⁻¹, respectively. It further indicates that the method significantly improves the sensitivity of SMMC-7721 exosomes with the assistance of Au@PDA NP-Apt2 and HAuCl₄. Compared with the existing method, the SPR assay based on aptamer and Au@PDA NPs is sensitive and specific for determining target exosomes in complex samples without any pretreatment (shown in Table S-2). Although many methods can sensitively determine the exosomes, most of these methods use antibody as the recognition probe, which was easy to denature and difficult to standardize. Here, the method for the determination of exosomes based on economical and stable aptamers probe has the same performance as

Table 1 Analysis of human serum samples spiked with SMMC-7721 exosomes

No	Add exosomes (particles mL ⁻¹)	Found exosomes (particles mL ⁻¹)	Recovery (%)	RSD (n = 3, %)
1	5.60×10^5	5.34×10^5	95.3	2.1
2	1.96×10^6	2.03×10^6	103.6	3.7
3	6.20×10^6	6.13×10^6	98.9	2.5
4	9.80×10^6	9.65×10^6	98.5	4.2

the existing methods. This work broadens the application of aptamer based SPR technology.

Next, the specificity of the SPR assay was investigated by using HepG2 exosomes, Bel-7404 exosomes, L02 exosomes, MCF-7 exosomes, and SW480 exosomes under the same condition. As shown in Fig. 5(C), the $\Delta\theta$ exhibits negligible response for HepG2 exosomes, Bel-7404 exosomes, L02 exosomes, MCF-7 exosomes, and SW480 exosomes. However, in the case of SMMC-7721 exosomes or SMMC-7721 exosomes with a mix of other exosomes, it causes a significant change of $\Delta\theta$ ($0.1204^\circ \pm 0.078$). The selectivity of the SPR assay was further investigated to specifically determine SMMC-7721 exosomes in a mixture of SMMC-7721 exosomes and other exosomes (HepG2 exosomes, Bel-7404 exosomes, L02 exosomes, MCF-7 exosomes, and SW480 exosomes, respectively). As shown in Fig. S-9, the $\Delta\theta$ response caused by SMMC-7721 exosomes is similar to that aroused using the mixing of SMMC-7721 exosomes with other exosomes. The results demonstrate that SPR assay show excellent selectivity for SMMC-7721 exosomes.

The regeneration of the SPR assay was studied. As shown in Fig. 5(D), as the SPR chip is dealt with regeneration solution after each determination of target exosomes (6.2×10^6 particles mL^{-1}), and 6 cycles of regeneration are observed without significant signal attenuation. The relative standard deviation (RSD) of 4.3% was also obtained. The results suggest that the assay has good reproducibility.

Determination of SMMC-7721 exosomes in serum

Given that the assay has excellent specificity for SMMC-7721 exosomes (Fig. 5(C)), it can avoid the concern of high levels of biological noise in the absence of a series of human serum pretreatments [19]. Here, for investigating the application of this assay, SMMC-7721 exosomes in 50% human serum are determined without any pretreatment. Figure S-10 illustrates the comparison of the $\Delta\theta$ obtained with determining in 50% human serum and in PBS buffer. It shows that $\Delta\theta$ caused by the same concentration of exosomes in 50% human serum and PBS buffer is almost the same. Besides, the SMMC-7721 exosomes in 50% human serum samples were also determined using the standard addition method. As shown in Table 1, the recoveries of SMMC-7721 exosomes in 50% human serum range from 95.3 to 103.6%, and the RSD is less than 5%. The results indicate that the method has the potential for the specific determination of SMMC-7721 exosomes in liquid biopsy.

Conclusions

A novel SPR assay based on aptamers and Au@PDA NPs was developed for the specific determination of SMMC-

7721 exosomes. This assay shows excellent specificity and stability for the determination of the SMMC-7721 exosome and can be used in 50% human serum analysis. It has the potential for the determination of SMMC-7721 exosomes in liquid biopsy. Theoretically, this assay can potentially be used for determining the other target exosome, as long as an appropriate aptamer can be replaced. However, it has complex operation steps and is limited to use in the laboratory. Therefore, it is required to simplify the operation steps in the future.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no competing interests.

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