



## Spatially reinforced Au nano-cavities as a reaction nano-reservoir for trace analysis of DNA hybridization



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### ABSTRACT

Spatially reinforced Au nano-cavities (SR-nAu) with reduced tip-to-tip displacement create strongly localized surface plasmon resonance by Raman scattering. Within SR-nAu, the nano-cavity structure with high aspect ratio was optimized as the nano-reservoir for the distinction of DNA hybridization responses from specific DNA sequences of Avian influenza virus. In the nano-reservoir, the suspended or immobilized probe, its complementary and noncomplementary target sequences were characterized at very low concentration through enhanced Raman spectra. The presence of Raman shift at  $738\text{ cm}^{-1}$ , the characteristic breathing mode of thymine, and the change of relative peaks intensity at  $1603$  and  $1554\text{ cm}^{-1}$  before and after hybridization were particularly taken as the major indications to validate the match or mismatch state of DNA sequences. Through a labeling-free means, and from the evidence provided by specific complementary and noncomplementary DNA sequences, the recognition capability of nano-reservoir can be down to subnano-molar concentration ( $10^{-10}\text{ M}$ ).

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### 1. Introduction

Accurate detection using biomarkers, e.g., antibody, antigen, DNA sequences, at early stages of a disease is of utmost importance in monitoring infectious and contaminated systems [1]. Specific DNA analyses have become a dominant method over the past decade, along with the unraveling genetic codes of variable biomarkers [2,3]. One of the widely established analytical methods is the use of fluorescence spectroscopy [4]; however, some drawbacks mainly correlated with multiplex techniques and stability of fluorescent dyes are presumably resulted. For increasing demands on the fast-screening technologies, e.g., for the control of influenza virus, the development of labeling-free means, where the recognition readouts can be measured directly from the analyze without additional target manipulation, has been constantly emphasized [5]. The related bio-sensing systems may consist of electrical [6,7], fluorescence, surface enhanced Raman scattering (SERS)-based [8,9], and microgravimetric techniques. The technique of SERS has

received much attention owing to its high enhancement for directly characterizing the specific molecular bonding of target species. With an appropriate SERS-active substrate, it requires only minimal sample preparation without labeling. In recent years, SERS technique has been increasingly and successfully employed in the study of, e.g., biological molecules [10,11], DNA sequences [12], peptides [13], proteins [14]. Furthermore applications are increasingly applied to, e.g., gene profiling [15], clinical diagnostics [16].

Nano-structure created on a noble metal film with an exceptional SERS property is of great interest for analyzing relatively small size and dimension of target species [17]. As the target species is attached, the effect of SERS may originate from a coupled laser wavelength with dramatic amplification of electromagnetic field accumulated with in the nano-structure, in particular for nano-cavity that may generate a localized surface plasmon resonance (LSPR) [18]. Presumably, the recognition of Raman-active species restricted within the nano-cavity with LSPR will be remarkably enhanced. In addition, comparing nano-rod with nano-cavity structure, the latter has the advantage of capturing target species in solution. Smaller or comparable target species is thus anticipated to be trapped into nano-cavity, then its Raman-active signals can be significantly enhanced by LSPR. In particular, a nano-cavity with multiple edges and a high aspect-ratio angle tend to enhance LSPR [19]. As nano-cavities are uniformly arrayed, the tip-to-tip distance between or among nano-cavities will also influence the

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effect of SERS [20]. It has been reported that spatially reinforced Au nano-cavities (*SR-nAu*) with a reduced tip-to-tip displacement are competent to create strong effect of SERS [21]. Based on the electromagnetic theory, a variety of LSPR modes can exist within the nano-cavities of different dimensions, each with its own characteristic electromagnetic field distribution and the resultant effect of SERS [22], which may thereafter provide wide applications in sensing Raman-active species [23].

It is therefore of great interest to apply *SR-nAu* as the nano-reservoir for distinguishing a controllable reaction of DNA hybridization at an extremely low concentration. For example, DNA sequences can be well distinguished by Raman signals of the labels attached onto its backbone structure [24]. Nevertheless, the obtained signals are those of the labels but not of the oligonucleotide [25]. As a consequence, it is very promising to develop a novel technique to characterize the oligonucleotides directly. DNA hybridization is the most common, specific, and spontaneous reaction between complementary DNA sequences, and it also take an important part in most biochemical reaction, followed by polymerase chain reaction (PCR). In this study, target and its complementary DNA sequences under suspension or immobilization condition are anticipated to be annealed and hybridized within the nano-reservoir. To analyze the change from single strands to hybridization responses, annealing temperatures and noncomplementary sequences are taken as the variables and verified. The labeling-free technique for the recognition of DNA hybridization at low concentration is thereafter realized and diagnosed within SERS-active nano-reservoir.

## 2. Materials and methods

### 2.1. Fabrication of SERS-active *SR-nAu*

Au substrate was prepared by thermal evaporation of  $\approx 200$  nm of Au (99.99% purity) onto the polished single crystal silicon (100) wafers (Silicon Sense, Germany) primed with a  $\approx 5$  nm-thick titanium adhesion layer. The as-prepared Au substrate surface (shortened as the *apAu*) was polycrystalline with a grain size of 20–50 nm; its grains predominantly exhibited (111) orientation. The grain size and the root mean squared roughness ( $\approx 0.62$  nm) of the *apAu* were measured by scanning probe microscope (SPA300HV, SEIKO, Japan).

The *SR-nAu* was fabricated using one-step loading-unloading mode of dynamic contact module system (NanoIndenter G200, Agilent Technologies, USA). In the making process, a triangular pyramidal tip of Berkovich diamond with a radius of  $\approx 20$  nm was used under a controlled relative humidity ( $\approx 32\%$ ) and room temperature ( $\approx 24^\circ\text{C}$ ). Before each batch, the nano-indentation tip was cleaned by indenting on the single crystal aluminum. The loading procedure was controlled to have a constant drift rate of 0.05 nm/s for stabilizing the fabrication steps. The applied loading force was in the range of 1500–1700  $\mu\text{N}$  for indentation depths ( $D_v$ ) of  $\approx 90$  nm measured using the vertical displacement. The tip-to-tip displacements ( $D_{t-t}$ ) of  $\approx 300$ ,  $\approx 500$ , and  $\approx 700$  nm, measured using the parallel displacement, were used. The  $D_{t-t}$  and the derived geometrical changes were chosen as the major parameters for comparison.

The field-emission scanning electron microscope (FE-SEM, JSM-7001, JEOL, Japan) was employed to determine the aperture size of cavity ( $D_a$ ) and  $D_{t-t}$ . The measured  $D_v$  was referenced in Supplementary data 1. The approaching velocity and the harmonic displacement of the nano-indentation tip toward the target surface were maintained at 1 nm/s and 1 nm for all the testing surfaces. Depending on the unspecified contact points, surface approaching sensitivity was optimized at 10% for the *apAu* [26].

**Table 1**

Probe (P) and target (T) sequences of H7 and their modifications.

Name	Description	Sequence 5'–3'
H7-T	H7 target sequence	TACTC-AATTT-GACTG-GGTCA-ATTG
H7-P	H7 probe sequence	CAAAT-TGACC-CAGTC-AAATT-GAGTA
HS-H7-P	Thiol-terminated H7-P	HS-C6-CAAAT-TGACC-CAGTC-AAATT-GAGTA
Cy3-H7-T	H7-T with Cy3 labeling	Cy3-TACTC-AATTT-GACTG-GGTCA-ATTG

### 2.2. Suspended molecular probes and DNA sequences

The *SR-nAu* used for distinguishing a controllable reaction at an extremely low concentration is called the nano-reservoir. R6G molecule (Sigma, Germany) is commonly used as a tracer dye in biotechnology applications, e.g., the use of fluorescence microscope. It is used as the index to examine the effect of SERS from the target species-included nano-reservoir. R6G solution was diluted with phosphate-buffered saline (PBS, Sigma, Germany) solution to a concentration of  $10^{-4}$  M. Thereafter the nano-reservoir with the molecular probes-containing solution was covered with glass slide and then immediately measured using Raman spectroscopy [27].

All types of DNA sequences used in this study were obtained from MD-Bio Inc. (Taipei, Taiwan), their names and base compositions were shown in Table 1. The target sequence (denoted as H7-T) was a specific biomarker of Avian influenza viruses (AIV), and its complementary sequence (denoted as H7-P) was designed as a capture site for DNA hybridization. These sequences, including the hybridized H7-T/H7-P (denoted as H7-hybrid) were stored at  $-4^\circ\text{C}$  before use. Then, they were diluted to  $10^{-8}$ ,  $10^{-9}$ ,  $10^{-10}$ , and  $10^{-11}$  M in PBS solution for further uses. A quantity of 5  $\mu\text{l}$  from different dilutions of each DNA sequence was placed on the *apAu* and within the nano-reservoir for subsequent experiments.

### 2.3. Immobilized DNA sequences on *SR-nAu*

Alternatively H7-P was immobilized on the nano-reservoir for further bio-sensing applications. At first, the *apAu* was employed to verify the immobilization process using high resolution X-ray photoelectron spectroscopy (HRXPS, ULVAC Inc., Kanagawa, Japan). The thiol group was modified at the 3' end of H7-P strand and prepared as the thiol-terminated H7-P (denoted as HS-H7-P), as described in Table 1. This sequence was diluted to  $10^{-6}$  M in PBS solution for further uses. HS-H7-P ( $10^{-6}$  M, 10  $\mu\text{l}$ ), dithiothreitol (DTT, 0.1 M, 10  $\mu\text{l}$ ), and triethylamine (TEA, 0.1 M, 1.5  $\mu\text{l}$ ) were mixed to reduce the disulfide bonds at the 5' end of HS-H7-P and generate free thiol groups for getting immobilization on the *apAu* and the nano-reservoir at  $4^\circ\text{C}$  for 24 h. After that, the HS-H7-P immobilized surface was washed by deionized water for 10 min, and then stored for further analyses. The HS-H7-P immobilized *apAu* was labeled with Cyanine dye (Cy3) to verify DNA hybridization. Raman spectra from the HS-H7-P immobilized nano-reservoir were then taken.

### 2.4. Raman spectroscopy and enhancement factor of SERS

A confocal microscopic Raman spectrometer (In Via Raman microscope, RENISHAW, United Kingdom) using 633 nm radiations from the excitation of He–Ne laser was employed. The excitation power of laser was  $\approx 17$  mW and the power on the substrate was  $\approx 8$  mW measured by an illumination meter. The scattering light was collected by a  $50\times$  air objective lens to a CCD detector. A grating of 1800 lines/mm was used to disperse the scattered light. All the reported Raman spectra were resulted from a single 10 s accumulation in a range of 500–3000  $\text{cm}^{-1}$ . Since there was no Raman signal in the range of 2000–3000  $\text{cm}^{-1}$ , the spectra in the range of

500–2000  $\text{cm}^{-1}$  were presented. Before each batch, Raman shifts were calibrated using the signal of 520  $\text{cm}^{-1}$  with the same absolute intensity from a standard silicon wafer.

The enhancement factor (EF) of the measured SERS was estimated according to the standard equation [28]:

$$EF = \frac{I_{\text{SERS}}/N_{\text{SERS}}}{I_{\text{NRS}}/N_{\text{NRS}}}, \quad (1)$$

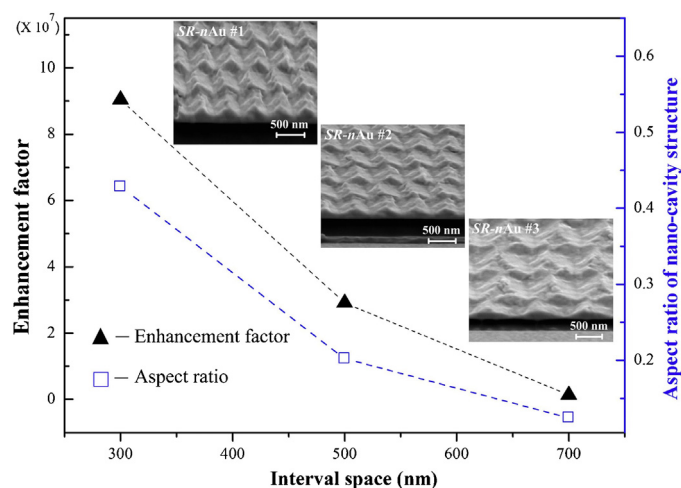
where  $I_{\text{SERS}}$  and  $I_{\text{NRS}}$  are SERS and normal Raman scattering intensity, respectively;  $N_{\text{SERS}}$  and  $N_{\text{NRS}}$  are the numbers of molecules contributing to the inelastic scattering intensity in each cavity structure, which are respectively evaluated by SERS and normal Raman scattering measurements. The calculated value of EF can represent the ability of inducing SERS effect in each cavity structure. Raman intensity was averaged from ten consecutive measurements. Note that the *ap*Au and the bare nano-reservoir are initially Raman-inactive and used as the reference substrate surface for the calculation of EF when Raman-active R6G molecules are adsorbed within the nano-reservoir.

### 3. Results and discussion

#### 3.1. Optimization of SR-nAu as the nano-reservoir

FE-SEM photo-images of SR-nAu#1, #2, and #3 were presented in Supplementary data 2. The  $D_a$  and the aspect ratio,  $D_v/D_a$ , of the inter-cavity structure were manipulated by  $D_{t-t}$  of each indentation step, the cross-section of cavity structures were observed in Fig. 1. According to the inset figure of Fig. 1, the aspect ratio was correlated with the percentage of volume shrinkage attributed to the mechanically made nano-cavity. For SR-nAu#3, each nano-cavity structure was mostly independent;  $D_a$  was measured  $\approx 725 \pm 8.3$  nm. As  $D_{t-t}$  was decreased to  $\approx 500$  nm (i.e., for SR-nAu#2),  $D_a$  was  $445 \pm 8.4$  nm. As  $D_{t-t}$  was further decreased to  $\approx 300$  nm (i.e., for SR-nAu#1),  $D_a$  was  $210 \pm 8.3$  nm. The nano-cavity structure progressively accumulated and became “spatially reinforced”, as  $D_{t-t}$  or  $D_a$  decreased.

The effect of SERS on SR-nAu#1–3 was firstly evaluated using an optimized concentration of  $10^{-4}$  M R6G. In Figure S-2, the major peak of the molecule, designated “ $P_{\text{R6G}}$ ”, was clearly found at 1361  $\text{cm}^{-1}$  (i.e., assigned to  $\nu(\text{C}=\text{C})$ , aromatics). The Raman



**Fig. 1.** Raman-active peaks of R6G molecules at 1360  $\text{cm}^{-1}$  ( $P_{\text{R6G}}$ :  $\nu(\text{C}=\text{C})$ , aromatics) was taken as the index for the calculation of SERS enhancement factors ( $\blacktriangle$ ) and corresponding aspect ratios ( $\bullet$ ) with respect to their  $D_{t-t}$  values within SR-nAu#1–3. The inset figures represented the cross-section of nano-cavity structures for SR-nAu#1–3.

intensity of “ $P_{\text{R6G}}$ ”, denoted as  $I_{\text{peak(R6G)}}$ , showed the enhancement. The  $I_{\text{peak(R6G)}}$  values for SR-nAu#1–3 increased with the decreasing  $D_{t-t}$  or  $D_a$ , which were recorded as intensity indicators for subsequent comparisons.

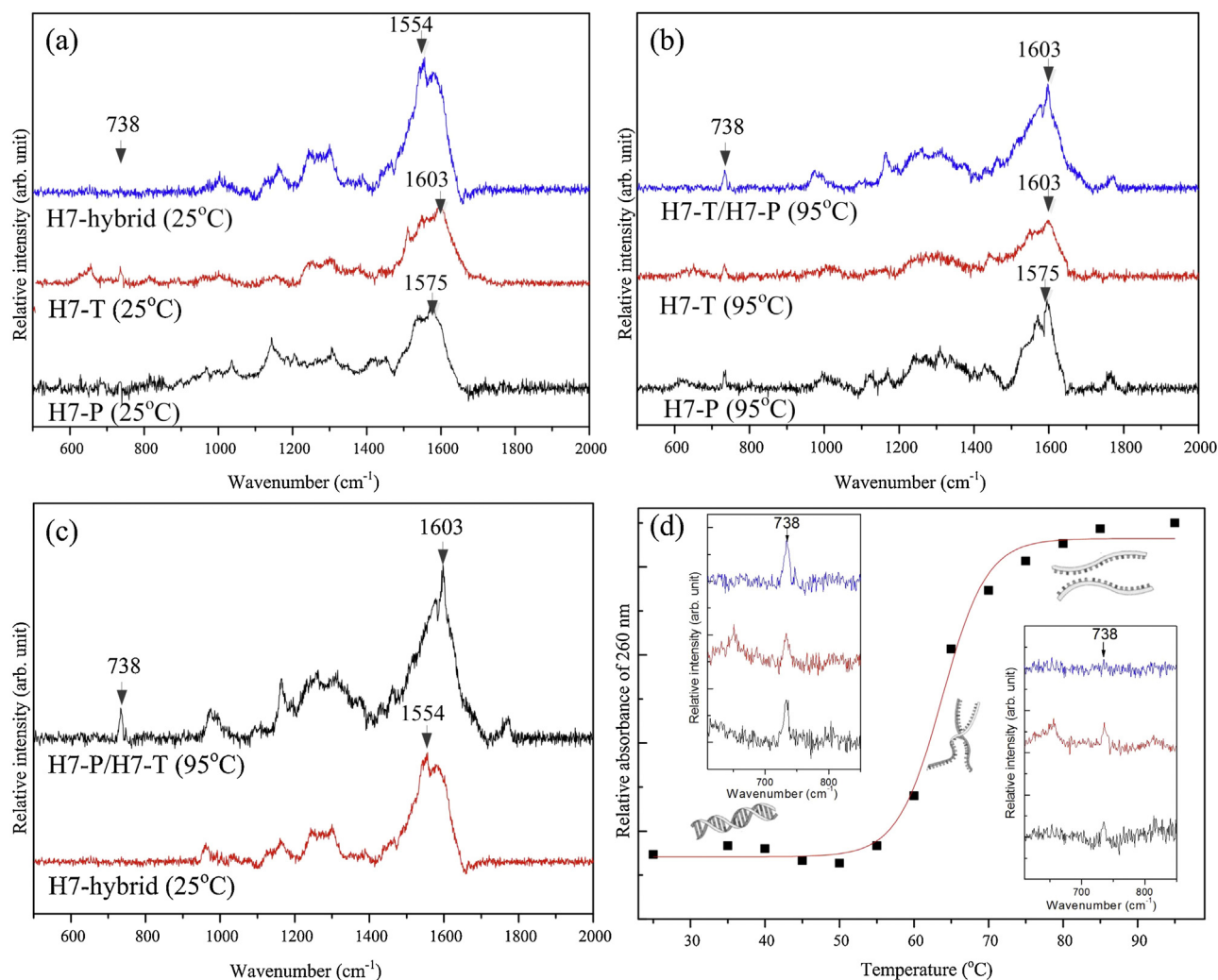
In Fig. 1, according to Eq. (1), EFs of SERS for SR-nAu#1–3 were estimated using  $I_{\text{SERS}} = I_{\text{peak(R6G)}}$  and  $N_{\text{SERS}} = N_{\text{R6G}}$ . Note that the  $N_{\text{NRS}}$  value of R6G is estimated using  $\approx 5 \mu\text{l}$  of  $10^{-2}$  M R6G spread over a diameter of 3 mm, and then the surface is dried off and measured. The average  $N_{\text{NRS}}$  was calculated  $\approx 4.2 \times 10^9$  molecules. The  $I_{\text{NRS}}$  value was measured  $\approx 150$ . The  $I_{\text{SERS}}$  value was divided by the amount of R6G per unit volume to obtain the EF of SERS for one single cavity of SR-nAu. In Fig. 1, EFs of SERS decreased with increasing  $D_{t-t}$  and decreasing  $D_a$ . The optimal EF of SERS for R6G within SR-nAu#1 was thereafter estimated  $\approx 9.0 \times 10^7$ .

R6G molecules were physically adsorbed near SR-nAu,  $I_{\text{peak(R6G)}}$  increased with decreasing  $D_{t-t}$ . Physically adsorbed R6G molecules within SR-nAu#1 were distinguished by a significant enhancement of Raman signals. By comparing the volume of SR-nAu, the increase of EF was most probably contributed by LSPs generated within the nano-cavity structure. The SR-nAu#1 with a chosen  $D_{t-t}$  of  $\approx 300$  nm was competent to adjust and maximize the resonance frequency of LSPs and thereafter to induce the electromagnetic effect within the nano-cavity.

#### 3.2. Recognition of single-strand DNA sequences and hybridization

The SR-nAu#1 was used as the nano-reservoir for the distinction of DNA hybridization. Based on the dimension of measured area (i.e., 1  $\mu\text{m}$ ) by Raman spot, four nano-cavities within SR-nAu#1 were defined as one nano-reservoir for the subsequent process of recognizing DNA sequences. In Fig. 2(a), Raman shifts in the range of 600–1800  $\text{cm}^{-1}$  appeared as fingerprinting information for H7-T, H7-P, and H7-hybrid, as given in Table 2. Note that H7-T, H7-P, or H7-hybrid on the *ap*Au did not acquire Raman signals. However, as shown in Supplementary data 2, SERS spectra of H7-hybrid within the nano-reservoir over a concentration range of  $10^{-8}$ – $10^{-11}$  M were mostly distinguishable. An optimized detection limit for further studies was preferably chosen at  $10^{-10}$  M, which could be 10–100 times lower than the detection limit of surveyed reports [29].

From Fig. 2(a), (b), and Table 2, Raman shifts in the regions of 650–750 and 1560–1610  $\text{cm}^{-1}$  were clearly shown by varying with DNA sequences and annealing temperatures. Raman spectra of H7-P and H7-T showed similar peaks assignment for those at 25 or 95  $^{\circ}\text{C}$ . Relative strong peaks at 1575 (for H7-P) and 1603  $\text{cm}^{-1}$  (for H7-T) were respectively distinguished by the arrangement of nucleic acid. In particular, the presence of Raman shift at 738  $\text{cm}^{-1}$ , the characteristic breathing mode of thymine, was clearly found for the sequences H7-T (25 and 95  $^{\circ}\text{C}$ ), H7-P (25 and 95  $^{\circ}\text{C}$ ), and H7-T/H7-P (95  $^{\circ}\text{C}$ ) and disappeared for H7-hybrid (25  $^{\circ}\text{C}$ ). Relative strong peaks at 1603 (i.e., H7-T,  $\text{NH}_2$  (A, G, C)) and 1554  $\text{cm}^{-1}$  (i.e., H7-hybrid (25  $^{\circ}\text{C}$ ), amide II) were obviously modified after hybridization. In Fig. 2(c), the representing Raman spectra for H7-hybrid (25  $^{\circ}\text{C}$ ) and H7-T/H7-P (95  $^{\circ}\text{C}$ ) were compared, which indicate DNA hybridization response that changes from double to single-strand by annealing temperatures. In Fig. 2(d), H7-T and H7-P were mixed and annealed within the nano-reservoir from 25 to 95  $^{\circ}\text{C}$ . The values of relative absorbance at 260 nm were recorded, while the hybridization responses were also illustrated. The result corresponded to the presence of Raman shift at 738  $\text{cm}^{-1}$ , which indicated the response change from double to single strand by annealing temperatures. Otherwise, the relative intensity of peak at 1554  $\text{cm}^{-1}$  showed similar intensity before and after hybridization, because this peak was contributed by amide II from nucleotides. The amount of amide II would not be affected by hybridization,



**Fig. 2.** SERS spectra of target sequence (H7-T), its complementary sequence (H7-P), and hybridized or mixed H7-T/H7-P were measured at different annealing temperatures: (a) 25 °C and (b) 95 °C, respectively. (c) SERS spectra of H7-hybrid (25 °C) and H7-T/H7-P (95 °C) were particularly compared. Raman shifts were assigned and described in Table 2. (d) The mixed H7-T/H7-P was annealed within the nano-reservoir from 25 °C to 95 °C; relative absorbance of 260 nm along with the increase of temperatures was recorded.

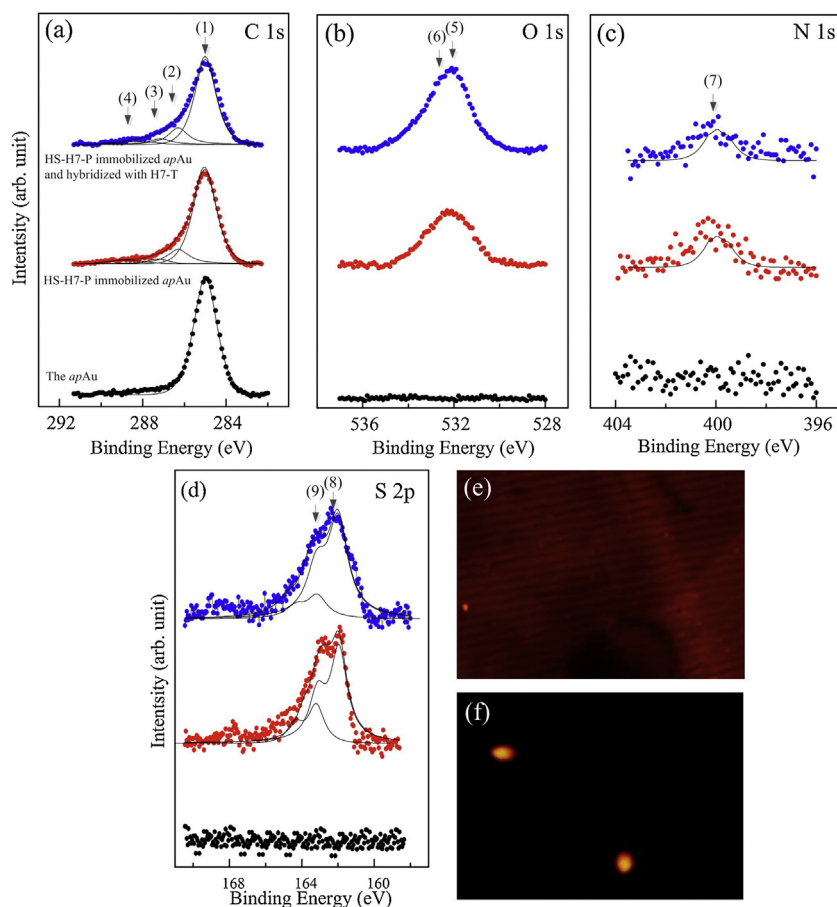
**Table 2**  
Raman shifts and assignments for H7-P, H7-T, H7-hybrid (25 °C), and H7-P/H7-T (95 °C).

Raman shift (cm <sup>-1</sup> )	Assignment	H7-P	H7-T	H7-hybrid (25 °C)	H7-T/H7-P (95 °C)
738	Ring breathing mode of T	✓	✓		✓
961	Deoxyribose	✓	✓	✓	✓
1181	T, C	✓			✓
1252	C, A		✓	✓	✓
1306	A	✓	✓	✓	✓
1554	Amide II	✓	✓	✓	✓
1575	Ring breathing modes G, A	✓	✓	✓	✓
1603	NH <sub>2</sub> (A, G, C)	✓	✓	✓	✓

so it was taken as a reference for the following comparison. The intensity ratio of 1575 cm<sup>-1</sup>/1554 cm<sup>-1</sup> changed from 1.65 to 0.82; this result is reproducible, which indicates that the breathing modes of nucleotides may be changed with the hybridization response. As a result, one may take Raman shift at 738 cm<sup>-1</sup> as the major indication to validate the match or mismatch state of DNA sequences. The nano-reservoir is therefore competent to identify DNA hybridization responses by identifying Raman shift at 738 cm<sup>-1</sup> and relative Raman intensity ratio between 1603 and 1554 cm<sup>-1</sup>.

### 3.3. Recognition of DNA sequences using H7-P-immobilized nano-reservoir

For practical applications, bio-specimen might contain some impurities, which tended to interfere with, e.g., in this study, the characteristic Raman spectra of H7-T, H7-P, and H7-hybrid sequences. It was therefore preferable to immobilize H7-P upon the nano-reservoir firstly as a capture site, followed by washing away the unbound species. A preliminary study on the HS-H7-P immobilized *ap*Au and then hybridized with H7-T was performed. In



**Fig. 3.** XPS spectra of (a) C 1s, (b) O 1s, (c) N 1s, and (d) S 2p were presented. To verify the immobilization of HS-H7-P upon the *apAu* with S–Au bonds, their respective chemical structures with binding energies were given: (1) C–C (285.0 eV), (2) C–O (286.5 eV), (3) C–N (286.7 eV), (4) N–C=O–N (289.0 eV), (5) C=O–N (531.6 eV), (6) C–O–H (532.8 eV), (7) C–N (400.2 eV), (8) S–Au (162.0 eV), and (9) dialkylsulfide (163.2 eV). (e) A fluorescent image of Au-S-H7-P/H7-T-Cy3 after washing (in red) was shown. (f) A fluorescent image of suspending H7-P/H7-T-Cy3 after washing (in dark) was also shown. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of the article.)

**Fig. 3(a)–(d)**, C 1s, O 1s, N 1s, and S 2p HRXPS spectra for the respective surfaces were presented. The results indicated that HS-H7-P was firmly immobilized upon the *apAu* and presumably competent to hybridize with H7-T. The binding energies for peaks (1)–(7) represented the presence of H7-P or H7-hybrid, while those for peaks (8) and (9) showed strong S–Au bond (162.0 eV) on the respective surfaces. In **Fig. 3(e)** and **(f)**, fluorescent images for the washed Au-S-H7-P/H7-T-Cy3 (in red) and H7-P/H7-T-Cy3 (in dark) were presented. Obviously the washing procedure could remove the unbounded species (e.g., the suspending H7-P/H7-T) from Au surface and remain the immobilized H7-hybrid.

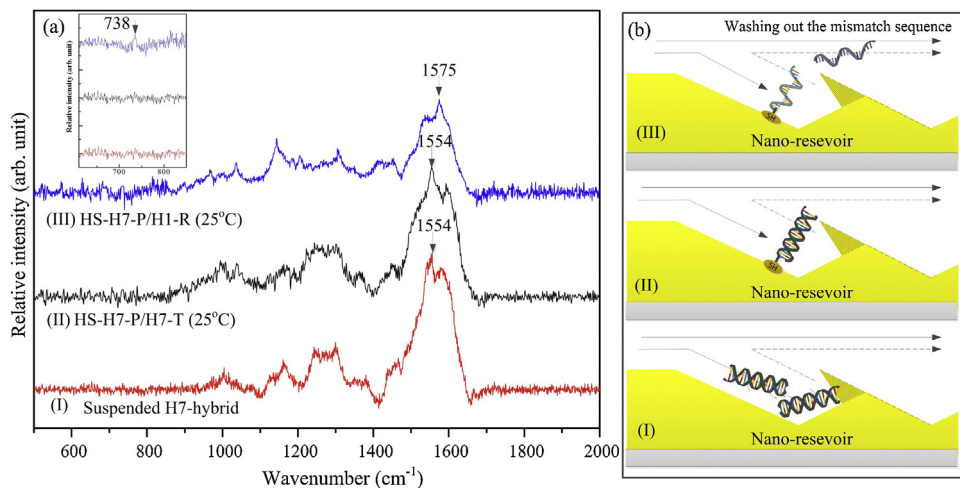
The nano-reservoir was thereafter taken as reaction nano-cavities, in which the suspended H7-hybrid and the immobilized HS-H7-P upon Au, followed by hybridized with H7-T, were studied. The H1-R, which was a mismatch sequence with HS-H7-P, was taken as the reference. Before taking Raman spectra, all the DNA sequences-containing nano-reservoirs were washed. In **Fig. 4(a)**, corresponding to the respective illustrations in **Fig. 4(b)**, Raman shift at  $738\text{ cm}^{-1}$  was checked with respect to the suspended H7-hybrid, HS-H7-P/H7-T ( $25^\circ\text{C}$ ), and HS-H7-P/H1-R ( $25^\circ\text{C}$ ). For the former two cases, H7-T and H7-P were hybridized and Raman shift at  $738\text{ cm}^{-1}$  was disappeared. However, the shift reappeared as the mismatched H1-R sequence was added to HS-H7-P at  $25^\circ\text{C}$ . In addition, the change of major peak intensity from  $1554\text{ cm}^{-1}$  (i.e., with hybridization) to  $1575\text{ cm}^{-1}$  (i.e., without hybridization, HS-H7-P remained upon Au) could be found as the immobilized

HS-H7-P reacted with the respective H7-T and H1-R at  $25^\circ\text{C}$  and then washed. This result indicated that HS-H7-P immobilized nano-reservoir was capable to recognize specific target sequence. It is therefore promising to apply this immobilization method as a labeling-free technique for capturing target sequence and recognizing a specific biomarker correlated with, e.g., a disease.

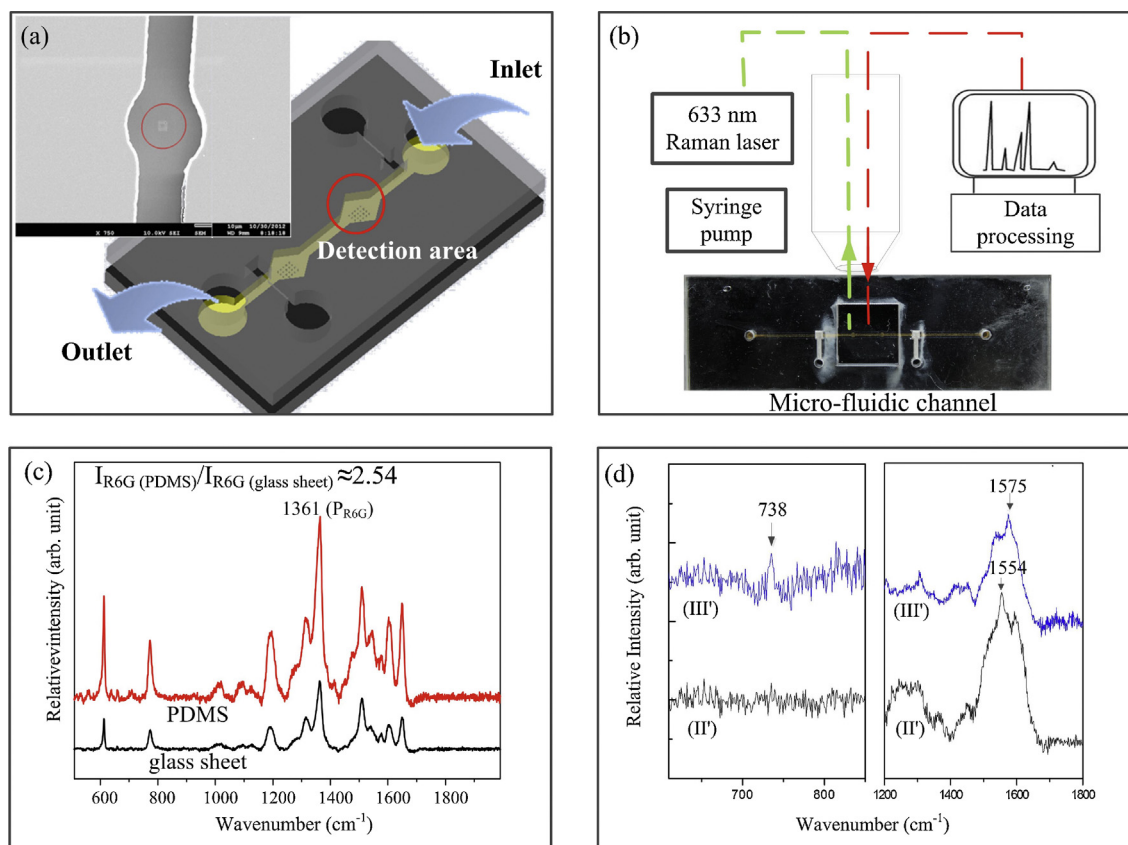
#### 3.4. HS-H7-P immobilized nano-reservoir embedded in a micro-fluidic channel

In **Fig. 5(a)**, FE-SEM micrograph and the layout of the nano-reservoir embedded micro-fluidic channel were presented. In **Fig. 5(b)**, schematic description of a simplified and automated micro-fluidic process was illustrated. Note that the pre-treated bio-specimen is directly transferred from the inlet to the detection area embedded with the nano-reservoir and followed by Raman analysis. After DNA hybridization, PBS solution was entered and washed out the unbounded species.

In **Fig. 5(c)**, R6G molecule was employed to verify the enhancement ability of the micro-fluidic channel. Comparing PDMS with thin glass sheet as the covering materials, relative intensity ratio ( $I_{\text{R6G}}(\text{PDMS})/I_{\text{R6G}}(\text{glass sheet})$ ) by taking  $1361\text{ cm}^{-1}$  ( $P_{\text{R6G}}$ ) as the indication was averaged around 2.54. PDMS was advantageous to be utilized for further application. In **Fig. 5(d)**, typical Raman shifts of HS-H7-P/H7-T ( $25^\circ\text{C}$ ) and HS-H7-P/H1-R ( $25^\circ\text{C}$ ) covered with



**Fig. 4.** (a) SERS spectra of H7 sequences with varied mixing reactions within the nano-reservoir, denoted as (I) suspended H7-hybrid, (II) HS-H7-P/H7-T (25 °C), and (III) HS-H7-P/H1-R (25 °C). (b) The corresponding illustrations of mixing reactions (I)–(III) within the nano-reservoir.



**Fig. 5.** (a) FE-SEM micrograph of the as-designed nano-reservoir embedded micro-fluidic channel was shown. (b) The nano-reservoir embedded micro-fluidic channel integrated with Raman microscopy as a bio-detection platform was illustrated. (c) Enhancement capabilities of the nano-reservoir embedded micro-fluidic channel covered by glass sheet or PDMS with the same thickness (0.2 mm) were compared. (d) SERS spectra with the major indications of HS-H7-P/H7-T (II') and HS-H7-P/H1-R (III') directly analyzed within the nano-reservoir embedded micro-fluidic channel at 25 °C were compared.

PDMS were recorded and denoted as spectra (II') and (III'). The characteristic spectra were comparable with those in Fig. 4(a), directly analyzed within the nano-reservoir. It is therefore validated that SR-nAu#1 is a capable SERS-active substrate to be used as the nano-reservoir for the distinction of specific DNA hybridization into micro-fluidic channel. Furthermore the nano-reservoir immobilized with DNA probe sequence is presumably enabled to

be embedded into a micro-fluidic channel for capturing the target DNA sequence without labeling.

#### 4. Conclusion

A novel DNA diagnostic method based on the reaction nano-reservoir has been developed. In this study, SR-nAu with high

aspect ratio for the nano-reservoir was firstly optimized using R6G molecule as the probe and the EF of SERS was estimated  $\approx 9.0 \times 10^7$ . Specific biomarkers of AIV, H7-T, H7-P, and H7-hybrid, within the nano-reservoir over a concentration range of  $10^{-8}$ – $10^{-11}$  M were mostly distinguishable. Raman shifts for the presence of  $738\text{ cm}^{-1}$  and relative intensities among 554, 1575, and  $1603\text{ cm}^{-1}$  were used as the major indications to validate the match or mismatch state of DNA sequences at 25 and  $95^\circ\text{C}$ . Thereafter the HS-H7-P immobilized nano-reservoir, in comparison with the suspended H7-hybrid, was taken as the reaction nano-cavities, hybridized with H7-T and H1-R (a mismatch sequence with HS-H7-P) at  $25^\circ\text{C}$ , and then washed. Raman shifts for the presence of  $738\text{ cm}^{-1}$  and the change of major peak intensity from 1554 (i.e., with hybridization) to  $1575\text{ cm}^{-1}$  (i.e., without hybridization, HS-H7-P remained upon Au) could be clearly found. Furthermore HS-H7-P immobilized nano-reservoir was embedded into a microfluidic channel and covered with PDMS; the integrated system was competent to capture and distinguish the target DNA sequence. Through labeling-free means, the recognition capability of SERS-active nano-reservoir for DNA sequences could be down to  $10^{-10}$  M.

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### Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.snb.2013.09.120>.

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