

Cite this: *Analyst*, 2020, **145**, 4173

# A novel liquid biopsy-based approach for highly specific cancer diagnostics: mitigating false responses in assaying patient plasma-derived circulating microRNAs through combined SERS and plasmon-enhanced fluorescence analyses†

Adrianna N. Masterson,<sup>‡a</sup> Thakshila Liyanage,<sup>‡a</sup> Claire Berman,<sup>a</sup>  
Hristos Kaimakliotis,<sup>b</sup> Merrell Johnson<sup>ib</sup> <sup>c</sup> and Rajesh Sardar<sup>ib</sup> \*<sup>§a</sup>

Studies have shown that microRNAs, which are small noncoding RNAs, hold tremendous promise as next-generation circulating biomarkers for early cancer detection via liquid biopsies. A novel, solid-state nanoplasmonic sensor capable of assaying circulating microRNAs through a combined surface-enhanced Raman scattering (SERS) and plasmon-enhanced fluorescence (PEF) approach has been developed. Here, the unique localized surface plasmon resonance properties of chemically-synthesized gold triangular nanoprisms (Au TNPs) are utilized to create large SERS and PEF enhancements. With careful modification to the surface of Au TNPs, this sensing approach is capable of quantifying circulating microRNAs at femto-gram/microliter concentrations. Uniquely, the multimodal analytical methods mitigate both false positive and false negative responses and demonstrate the high stability of our sensors within bodily fluids. As a proof of concept, microRNA-10b and microRNA-96 were directly assayed from the plasma of six bladder cancer patients. Results show potential for a highly specific liquid biopsy method that could be used in point-of-care clinical diagnostics to increase early cancer detection or any other diseases including SARS-CoV-2 in which RNAs can be used as biomarkers.

Received 16th March 2020,

Accepted 21st May 2020

DOI: 10.1039/d0an00538j

rsc.li/analyst

## Introduction

Liquid biopsies involve analyzing circulating biomolecules in bodily fluids provide many advantages over traditional tissue biopsies for detecting cancers at an early stage as they are non-invasive, less expensive, and simpler.<sup>1</sup> Small non-coding RNAs,

known as microRNAs, have been found to regulate a large number of human genes and play a significant role in various cancer developments.<sup>2–7</sup> Many studies have demonstrated that the detection of circulating microRNAs at an early stage, via liquid biopsies, could allow better treatment in an effort to prevent cancer progression and metastasis, increasing the chance of patient survival. The current “gold-standard” microRNA assays (*i.e.*, quantitative reverse-transcription polymerase chain reaction (qRT-PCR)-based technologies) are capable of measuring microRNA levels reproducibly from bodily fluids in real-life clinical samples. However, this technology has several drawbacks including: (1) the treatment of biological fluid, (2) complicated labor-intensive RNA extraction, (3) required labeling and amplification prior to analysis, (4) complementary DNA conversion steps, and (5) the large volume of patient samples needed.<sup>8</sup> Together, these traits hinder translation into clinical point-of-care (POC) diagnostics.

In contrast, solid-state, label-free biosensing offers many unique advantages over qRT-PCR techniques, including simple sensor fabrication, high sensitivity, easy to miniaturization (of particular importance when adapting technologies to POC diagnostics), and elimination of receptor tagging with

<sup>a</sup>Department of Chemistry & Chemical Biology, Indiana University-Purdue University Indianapolis, 402 N. Blackford Street, Indianapolis, Indiana 46202, USA.

E-mail: rsardar@iupui.edu

<sup>b</sup>Department of Urology, Indiana University School of Medicine, 535 N. Barnhill Dr., Indianapolis, Indiana 46202, USA

<sup>c</sup>Department of Physics, Purdue University Fort Wayne, 2101 E. Coliseum Blvd., Fort Wayne, Indiana 46805, USA

† Electronic supplementary information (ESI) available: Detailed experimental procedures for Au TNPs synthesis, various spectroscopy and microscopy characterizations, and additional extinction and Raman spectra, and raw Raman data. See DOI: 10.1039/d0an00538j

‡ These authors contributed equally to this work.

§ Integrated Nanosystems Development Institute, Indiana University-Purdue University Indianapolis, 723 W. Michigan Street, Indianapolis, Indiana 46202, United States.

dyes or other specialized reagents ("label-free"). Among advantages, the latter is extremely important as it avoids undesirable interactions between labels and analytes, leading to more reliable and reproducible results. Current solid-state, label-free biosensing techniques such as microarray,<sup>9</sup> electrochemical,<sup>10,11</sup> localized surface plasmon resonance (LSPR),<sup>12–14</sup> surface-enhanced Raman scattering (SERS),<sup>15,16</sup> and photonic microring resonator-based<sup>17</sup> microRNA sensors provide good sensitivity. These single-mode detection methods rely on a specific sensing mechanism. For example, LSPR-based methods detect changes in local refractive index of the nanostructures while electrochemical-based measurements detect changes in current. Unfortunately, such changes can also result from non-specific adsorption of endogenous biomolecules from bodily fluids onto the sensor, as well as structural changes that can be induced following placement within bodily fluids during the measurements. As a result, these techniques suffer from low specificity, particularly when analyzing bodily fluids, and can in turn produce false test results. Further, aside from detecting/analyzing the specific signal change that occurs from the analyte, no additional confirmation of analyte attachment to the sensor can be obtained. Among these techniques, SERS-based biomolecular assays are capable of using Raman signal to provide rich structural information on the analytes. However, current solid-state SERS-based microRNA assays still show many problems, including low sensitivity, poor specificity, and inability to analyze real-life patient samples.<sup>15,16,18–21</sup>

High specificity is the most crucial factor in creating assays that can avoid false-positive and/or false-negative test results for the POC diagnosis.<sup>22,23</sup> To overcome the current technological bottleneck in real-life patient sample analysis, we present, for the first time, the design of a solid-state nanoplasmonic sensor. This novel sensing approach is capable of assaying oncogenic microRNAs directly in human plasma with high sensitivity and specificity. Here, the unique LSPR properties<sup>24,25</sup> of noble metal nanoparticles are utilized for the detection of microRNAs *via* a combined SERS<sup>26,27</sup> and plasmon-enhanced fluorescence (PEF)<sup>28,29</sup> approach. Importantly, the same nanoplasmonic sensor can be used in this multimodal detection method; no changes to structural parameters are required. To demonstrate the potential translational aspects of our dual detection technique in liquid biopsies for POC diagnosis, we successfully assayed two oncogenic microRNAs (microRNA-10b and microRNA-96). Analysis was performed directly in six bladder cancer patient plasma samples at sub-femtogram/microliter ( $\text{fg } \mu\text{L}^{-1}$ ) concentrations.

## Experimental section

### Materials

Chloro(triethylphosphine) gold(i) ( $\text{Et}_3\text{PAuCl}$ , 97%) was purchased from Gelest Inc. Poly-(methylhydrosiloxane) (PMHS,  $M_n = 1700\text{--}3300$ ), triethylamine (TEA, 98%), ACS grade acetonitrile ( $\text{CH}_3\text{CN}$ , 99.9%), and 3-mercapto-1-propanol (3-MP,

95%) were purchased from Sigma-Aldrich. Thiol modified 5'-SH-( $\text{CH}_2$ )<sub>3</sub>-ssDNAs, and various microRNAs were purchased from Integrated DNA Technologies (IDT). RNase H enzyme and RBS detergent solution were purchased from Thermo Scientific. (3-Mercaptopropyl)-trimethoxysilane (MPTMS, 94%) was purchased from Alfa Aesar. All the chemicals were used without any further purifications. RNase free sterile water was obtained from Baxter Healthcare Corporation. The glass cover slips and the tris(2-carboxyethyl)phosphine (TCEP) solution were purchased from Fisher Scientific. Bladder cancer patient plasma samples were obtained from the Indiana University medical school and used as received. All water was purified using a Thermo Scientific Barnstead Nanopure system. Thiol modified -ssDNAs, microRNAs were stored at  $-80^\circ\text{C}$ . PBS buffer ( $\text{pH} = 7.2$ ) was prepared using RNase-free sterile water. All experiments were performed in accordance with the Guidelines of the United States and approved by the ethics committee at Indiana University. Informed consents were obtained from human participants of this study.

### Fabrication of nanoplasmonic sensors

Gold triangular nanoprisms (Au TNPs) were synthesized using our well-established method.<sup>12,14,30</sup>  $\text{Et}_3\text{PAu(I)Cl}$  (10 mg) was dissolved in 20 mL of acetonitrile. After five minutes of stirring at room temperature, 0.019 mL (0.136 mmol) TEA was added, and the mixture was heated to  $40^\circ\text{C}$ . Then, 0.3 mL (2.75 mmol) of PMHS was added, and the reaction ran until the desired LSPR peak of 800 nm was achieved. Au TNPs were then attached onto silanized glass coverslips as we reported previously.<sup>13</sup> Separately, a mixture of  $5\text{ } \mu\text{M}$  -C3-ssDNA-10b/96 with 0.1 M TCEP was prepared and allowed to react for 1 hour to break disulfide bonds. The solution was then diluted with PBS buffer, and the resulting solution was used to incubate Au TNPs for overnight. The addition of -ssDNA-10/96 solutions induced ligand exchange of Au TNPs by removing TEA and PMHS that were present on the surface during the colloidal synthesis. Next, -ssDNA-10b/96-functionalized Au TNPs were washed thoroughly with RNase free water, and both UV-Vis spectrum and Raman spectrum were acquired. Finally, 10 mL, 1.0 mM 3-MP solution was added to ssDNA-10b/96-functionalized Au TNPs to fabricate nanoplasmonic sensors. Various concentrations (100 nM–100 fM) of microRNA-10b or microRNA-96 in 10 mL PBS buffer were used to incubate nanoplasmonic sensors for 2 hours. MicroRNA bound nanoplasmonic sensors were washed thoroughly with RNase free water to remove any loosely bound microRNAs to the sensor. For the reversibility/regeneration tests, microRNA-bound nanoplasmonic sensors were incubated in 15 units RNase H enzyme in RNase free water for 2 hours.

### UV-Vis and SERS measurements, and fluorescence confocal microscopy imaging

UV-Vis spectra were collected from 1100–300 nm on a Varian Cary 50 UV/Vis spectrophotometer, and all sensors were analyzed in RNase free water. SERS spectra were collected of the sensor on clean aluminum foil using a Foster-FORAM 785 HP

Raman spectrometer with a 785 nm diode laser excitation source. Samples were placed on top of the aluminum foil when analyzed due to the requirement of a clean surface to analyze. The background scan was taken of a polystyrene bead to calibrate the laser. For all SERS measurements, two randomly selected spots were analyzed for each sensor and for three-separate sensors (a total of six measurements). Using a Fluoview FV1000 MPE laser scanning biological microscope, 100 nM Alexa-488 dye was used in the analysis of the sensors. A laser power of 40%, a 60 $\times$  water objective, and gain of 1 $\times$  were used for the image acquisition. The diameter of the laser spot was 5- $\mu$ m.

### Scanning electron and atomic force microscopy analyses

The chemically synthesized Au TNPs attached onto the silanized glass coverslips were characterized by scanning electron microscopy (SEM) and atomic force microscopy (AFM). SEM micrographs were obtained using a Hitachi S-4700 Field Emission Scanning Electron Microscope (FESEM) at 20 kV accelerating voltage, and AFM images were obtained using a Bioscope AFM instrument.

### Construction of calibration curves

All SERS spectra were baseline corrected using OMNIC software. The Fit Peaks tool was used in Origin 2018b to find peak intensity with Lorentz fitting. Peak intensity at a specific wave-number was graphed as a function of concentration. Linear fitting was used to find the *R*-squared value.

### Measuring false responses

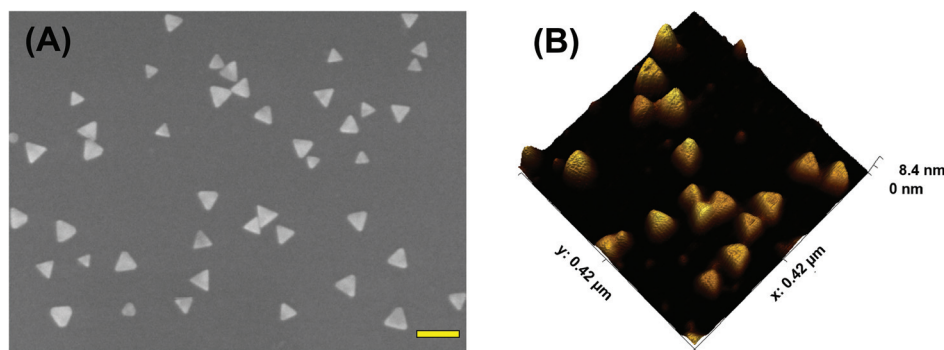
In order to demonstrate the efficiency of our solid-state nanoplasmonic-based sensors mitigating false responses, we analyzed the patient samples that measured the highest and lowest concentrations. Nanoplasmonic-based sensors, which were used to quantify microRNA-10b/96 in patient plasma again re-incubated in 15 units RNase H enzyme in RNase free water for 2 h. Next, both SERS measurement fluorescence confocal measurements were performed. Again, the same sensors were incubated in a 100 nM solution of microRNA-FAM. After

the sensors were thoroughly washed with RNase free water, SERS and fluorescence analyses were conducted.

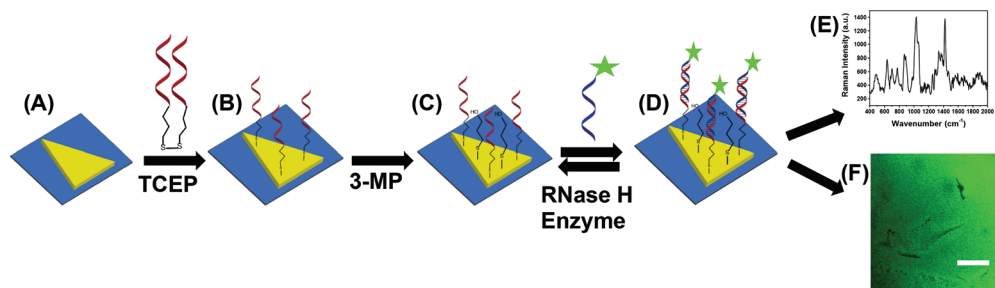
## Results and discussion

To acquire both SERS and PEF signals from the same nanoplasmonic sensor, we used chemically-synthesized gold triangular nanoprisms (Au TNPs) to generate a high intensity LSPR response.<sup>31</sup> This intensified LSPR arises due to strong electromagnetic field (EM) enhancement that occurs at the sharp tips and edges of Au TNPs.<sup>32–35</sup> This leads to the formation of hot-spots, which are ideal for LSPR-based SERS and PEF mechanisms.<sup>26,29</sup> For the first time, we have utilized PEF to unequivocally confirm microRNA attachment onto the nanoplasmonic sensor by utilizing fluorescence confocal imaging. Together, these methods fully eliminate the possibility of false assay responses. Further, in addition to their unique LSPR properties, the atomically flat Au TNPs allow -ssDNAs to form a tightly-packed self-assembled monolayer (SAM) on their surface, allowing for high reproducibility throughout the sensor. This is important in the context of making good interactions and forming a -ssDNA/microRNA duplex. Moreover, tightly-packed SAMs avoid non-specific adsorption of biomolecules onto the Au surface. Au is also extremely stable in human biofluids, therefore sensors are expected to maintain excellent structural integrity. Taking into consideration all their unique structural properties, we chose to Au TNPs with a 42 and 8 nm edge length and thickness, respectively, for the fabrication of our nanoplasmonic sensors, as shown in Fig. 1.

The fabrication of nanoplasmonic sensors is illustrated in Fig. 2, where: (A) Au TNPs were chemically-attached onto silanized glass substrate; (B) TNPs were functionalized with -ssDNA-(10b/96) *via* the Au-S bond, and in order to increase their specificity in SERS, they were further functionalized with 3-mercapto-1-propanol (3-MP) (Fig. 2C) so that the -ssDNA-10b/96 would appear in a “standup” position. This standup formation creates an ideal distance between the fluorophore and Au TNPs to maximize PEF enhancement by reducing fluorescence quenching through the Förster reso-



**Fig. 1** Structural characterizations of Au TNPs. (A) A representative scanning electron microscopy image of Au TNPs attached onto a silanized glass substrate. Scale bar is 100 nm. (B) An atomic force micrograph of Au TNPs.



**Fig. 2** Design of a nanoplasmonic sensor for the detection of microRNAs using combined surface-enhanced Raman spectroscopy (SERS) and fluorescence confocal microscopy techniques. (A) Freshly synthesized Au TNPs were attached onto silanized glass coverslip. (B) Tris(2-carboxyethyl) phosphine (TCEP) addition to DNA-disulfide solution breaks disulfide bonds and subsequent incubation in the solution resulted in the attachment of -ssDNAs on the surface of TNPs. Generally, -ssDNAs were randomly organized on the flat metal surface (e.g., here TNPs).<sup>10</sup> (C) -ssDNA-attached Au TNPs were further functionalized with 3-mercaptopropanol (3-MP) to allow -ssDNAs to “stand up” to achieve the most favorable -ssDNA/microRNA hybridization. The functionalization steps until (C) produced nanoplasmonic sensors, where microRNA-10b/96-FAM can selectively bind (D) to be analyzed by both SERS spectroscopy (E) and fluorescence confocal microscopy (F). Nanoplasmonic sensors can also be regenerated by incubating microRNA-bound sensors in a RNase H enzyme solution; image not to scale.

nance energy transfer (FRET) process. In addition, stand-up -ssDNAs provides adequate space for the microRNAs to form -ssDNA/microRNA duplexes. Finally, 3-MP also improves specificity, as reported in the literature.<sup>10</sup> The functionalizations of Au TNPs with -ssDNA-(10b/96) and 3-MP represent our nanoplasmonic sensors. Fig. 2D shows attachment of a fully complementary microRNA tagged with a 6-fluorescein (FAM), allowing complete -ssDNA/microRNA-FAM hybridization. Most importantly, as shown in Fig. 2D, the nanoplasmonic sensors are fully reversible and regenerative to allow use for multiple analyses. This is critical in the context of determining sensor stability and showing that they maintain complete structural integrity during assays.

Attachment of microRNA-FAM to the nanoplasmonic sensor can be detected/confirmed utilizing both SERS (Fig. 2E) and fluorescence confocal microscopy (Fig. 2F) techniques. It is important to highlight the additional advantages of selecting this particular dimension Au TNP for a SERS-based microRNA assay: (1) As shown in Fig. S1,† the LSPR peak position of nanoplasmonic sensors is  $\sim 860$  nm. The LSPR peak of TNPs and the wavelength of incident laser light controls the intensity of hot-spots. Ideally, for the EM-field-driven SERS enhancement, the LSPR wavelength of nanostructures in the sensors should be longer than the wavelength of laser light; (2) Use of a low energy laser (e.g., 785 nm) that fits the excitation requirements for nanostructures is important for biosensing applications in order to avoid degradation of biomolecules, particularly microRNAs.

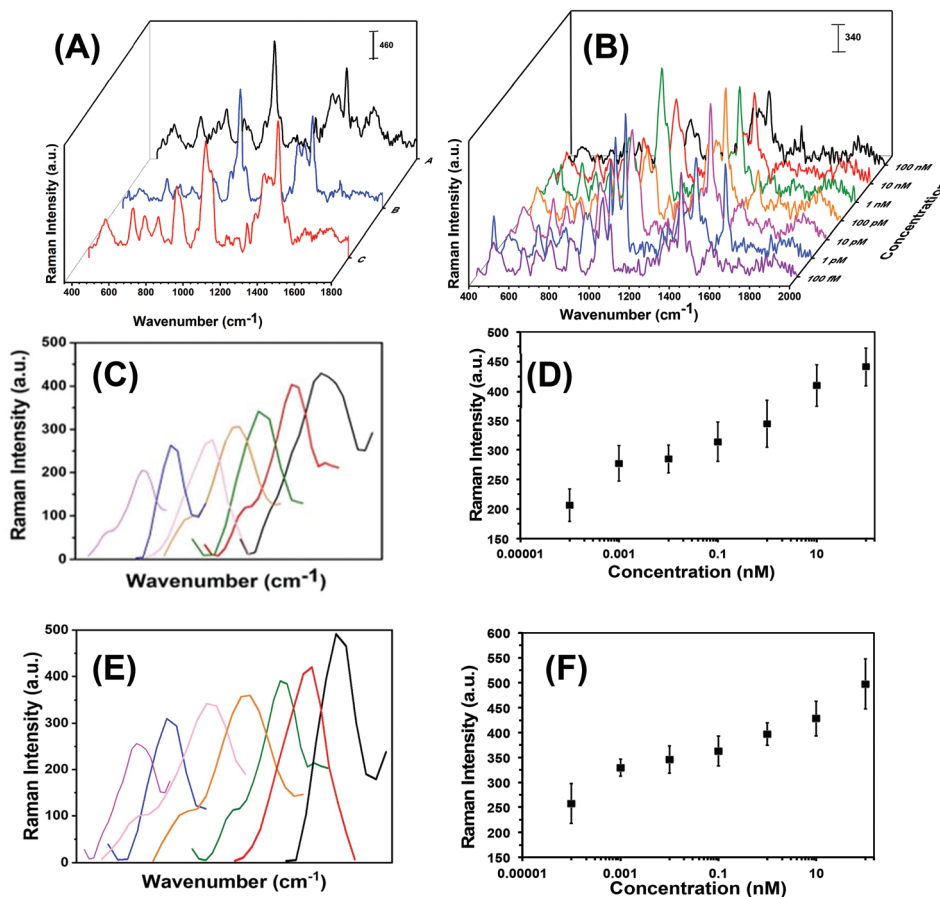
Each step in the fabrication of these nanoplasmonic sensors was characterized through SERS spectroscopy (see Fig. 3). Attachment of -ssDNA-(10b/96) *via* Au-S bonds results in the appearance of a C-S Raman stretch at  $635\text{ cm}^{-1}$ . Furthermore, several new Raman stretches that are characteristic of DNAs (see Fig. 3A and Fig. S2†) such as guanine (G) ring breathing at  $688\text{ cm}^{-1}$ , adenine ring breathing at  $733\text{ cm}^{-1}$ , cytosine ring breathing at  $790\text{ cm}^{-1}$ , and the phosphodiester group of the nucleic acids at  $1289\text{ cm}^{-1}$  are also

observed.<sup>18,21,36–38</sup> The C-H Raman stretch at  $1400\text{ cm}^{-1}$  further confirms the functionalization of Au TNPs with 3-MP (Fig. 3A-blue).<sup>39</sup> Incubation of microRNA-(10b/96)-FAM to the nanoplasmonic sensor displays a prominent Raman peak at  $1062\text{ cm}^{-1}$  of the additional  $\text{PO}_2^-$  symmetrical stretch.<sup>38</sup> Furthermore, the Raman stretch at  $1255\text{ cm}^{-1}$  corresponds to guanine C8-H bending<sup>18,21,36,37</sup> and the low intensity Raman stretch at  $1645\text{ cm}^{-1}$  is unique to the uracil C=O stretch.<sup>18,21,36,37</sup> Fig. 3B illustrates SERS spectra of the microRNA-10b varying concentrations from 100 nanomolar (nM) to 100 femtomolar (fM).

The average edge-length of our chemically synthesized Au TNPs is 42 nm that provides a top surface area of  $764\text{ nm}^2$ . Considering all the -ssDNA probes were attached onto the top surface and thiolate has a  $0.25\text{ nm}^2$  footprint, there would be approximately 3000 -ssDNA-(10b/96) attached per TNP. In the nanoplasmonic sensors fabrication, we used a high concentration of 3-MP in comparison to -ssDNA probes. Therefore, the -ssDNA grafting density is overestimated because a relatively large percentage of TNP surface would be occupied by 3-MP molecules. Further experimental characterization is required to quantitatively determine the grafting density of -ssDNA probe that is currently under our investigation.

In literature, researchers have used a variety of SERS stretches to develop the calibration plots for microRNA assays, but have done so without providing any detailed rationale for their selections.<sup>16,19,40</sup> Theoretical calculations show that the C8-H bending Raman stretch of guanine at  $1255\text{ cm}^{-1}$  could be ideal to develop the calibration plot,<sup>41</sup> thus we selected this SERS stretch for our studies. As shown in Fig. 3C and E for microRNA-10b and microRNA-96, respectively, a continuous increase in C8-H stretch intensity is displayed when corresponding nanoplasmonic sensors were incubated in various concentrations of microRNA solution. The SERS intensity of the C8-H bending mode *versus* the concentration of microRNAs is appeared to be linear over the entire concentration range (Fig. 3D and F, Tables S3 and S4†). The limit of





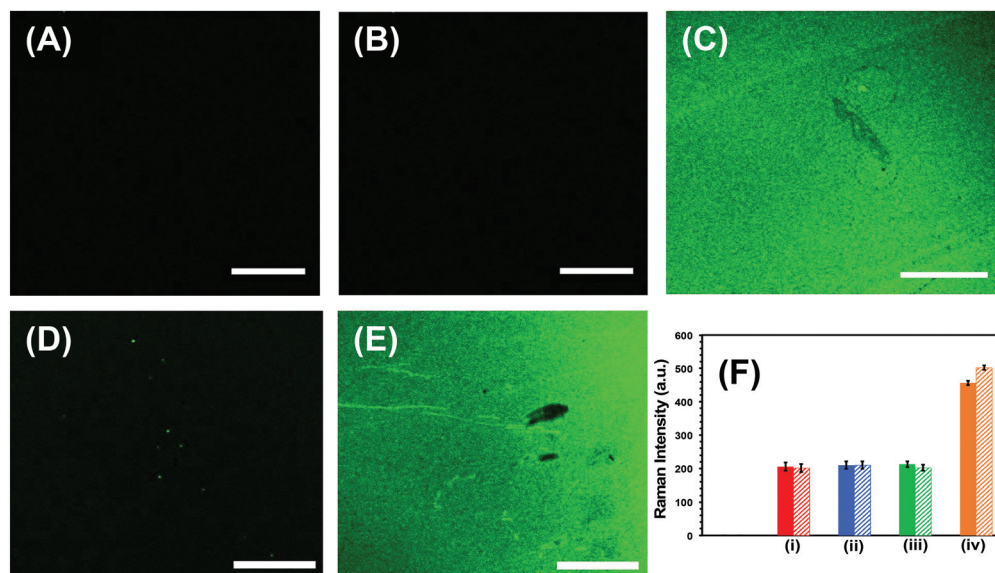
**Fig. 3** The SERS-based microRNA assay using nanoplasmonic sensors. (A) SERS spectra at different stages: Functionalization of AuNPs with -ssDNA-10b, A; after attachment of 3-MP on TNPs, B; and after incubation of nanoplasmonic sensors in a microRNA-10b solution, C. (B) SERS spectra of nanoplasmonic sensors upon incubation in different concentrations of microRNA-10b solution. (C) Expanded SERS spectra of the C8-H bending stretch of guanine at  $1255\text{ cm}^{-1}$  in various microRNA-10b concentrations (light purple: 100 fM, blue: 1 pM, pink: 10 pM, orange: 100 pM, green: 1 nM, red: 10 nM, black: 100 nM). (D) Intensity of the C8-H bending stretch at  $1255\text{ cm}^{-1}$  as a function of microRNA-10b concentration (logarithmic scale).  $Y = 15.94\ln(X) + 361.88$ ,  $R^2 = 0.97$ . (E) Expanded SERS spectra of the C8-H bending stretch of guanine at  $1255\text{ cm}^{-1}$  in various microRNA-96 concentrations (light purple: 100 fM, blue: 1 pM, pink: 10 pM, orange: 100 pM, green: 1 nM, red: 10 nM, black: 100 nM). (F) Intensity of the C8-H bending stretch at  $1255\text{ cm}^{-1}$  as a function of microRNA-96 concentration (logarithmic scale).  $Y = 15.04\ln(X) + 408.73$ ,  $R^2 = 0.95$ . In (C) and (E), the individual concentrations of the microRNA calibration curve are plotted in order to better visualize of the increase in intensity. Each spectrum has the same wavenumber of  $1255\text{ cm}^{-1}$  and does not shift in wavenumber as seen in the figure.

detections (LODs) for microRNA-10b and microRNA-96 calibration was calculated using a published method that follows:<sup>40,42</sup>

$$\text{LOD} = 10^{3m/k} \text{ fM}$$

Here,  $m$  is the relative standard deviation of the blank and  $k$  is the slope of the SERS intensity *versus* microRNA on the concentration curve. Our LODs for microRNA-10b and microRNA-96 are calculated at 1.13 pM and 0.030 pM, respectively. Importantly, this represents a 10-fold improved LODs in comparison to previously reported SERS-based microRNA assays.<sup>38</sup> We believe that the high SERS sensitivity of our nanoplasmonic sensors is due the intensified hotspots at the sharp tips and edges of the Au TNPs, as previously shown through discrete dipole approximation calculations.<sup>32–34</sup>

Successful implementation of the developed biosensors in liquid biopsy-based cancer diagnostics requires unprecedentedly high specificity, specifically to avoid false test results.<sup>1</sup> First, the specificity, which is the ability to unambiguously identify the analyte of our nanoplasmonic sensors, was investigated through fluorescence confocal imaging of labeled target microRNAs. We used microRNA-10b as a model system to study the specificity of the nanoplasmonic sensors. Fig. 4A confirms that our nanoplasmonic sensors do not display any fluorescence (in the absence of microRNAs). Similarly, when Au TNPs were functionalized only with 3-MP (without -ssDNA-10b) and then incubated in a microRNA-10b-FAM solution, no fluorescence signals are detected (data not shown). Further, when the 3-MP functionalized Au TNPs were incubated in a -ssDNA-10b solution, no fluorescence is detected (Fig. 4B). However, when the nanoplasmonic sensors, which

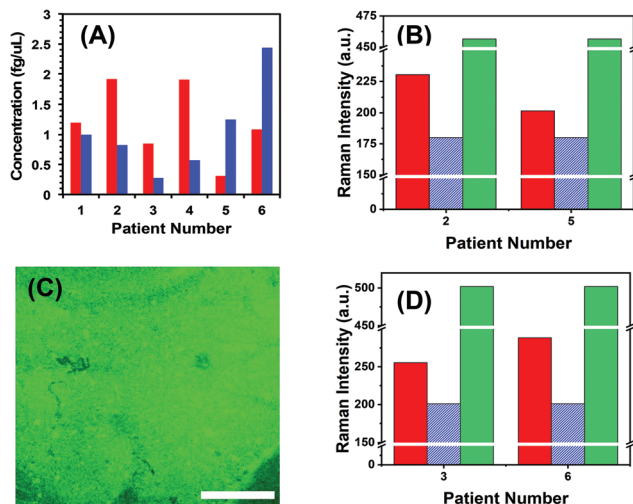


**Fig. 4** Specificity tests of nanoplasmonic sensors for the microRNA detection. Fluorescence confocal microscopy images of (A) glass substrate attached Au TNPs, (B) mixed -ssDNA-10b and 3-MP functionalized Au TNPs (nanoplasmonic sensor), (C) after incubation of nanoplasmonic sensors in a 100 nM microRNA-10b-FAM solution, (D) After treatment of nanoplasmonic sensors with RNase H enzyme to regenerate the sensor, (E) after re-incubation of nanoplasmonic sensors in a 100 nM microRNA-10b-FAM solution. Scale bars are 50  $\mu\text{m}$ . (F) Normalized Intensity of the C8–H bending stretch at 1255  $\text{cm}^{-1}$  for two different microRNAs (solid bars, microRNA-10b; dashed bars, microRNA-96). Black bars represent glass substrate attached AuTNPs (not visible). (i) Red bars represent mixed -ssDNA-10b and 3-MP functionalized Au TNPs or -ssDNA-96 and 3-MP functionalized Au TNPs. (ii) Blue bars represent incubation of nanoplasmonic sensor in a control patient (without the history of cancer) plasma. Green bars represent incubation of nanoplasmonic sensors in a mixture of microRNA-96, 145, 143, and 490-5p solution for the -ssDNA-10b functionalized Au TNPs, or a mixture of microRNA-10b, 145, 143, 490-5p solution for the -ssDNA-96 functionalized Au TNPs. These microRNAs are non-complementary to either -ssDNA-10b or -ssDNA-96. Orange bars represent incubation of nanoplasmonic sensors either in 100 nM microRNA-10b or microRNA-96 solution.

were constructed to detect microRNA-10b, were incubated in a microRNA-10b-FAM solution followed by copious rinsing, confocal imaging shows bright green fluorescence (Fig. 4C). Additionally, a control experiment was performed by incubating a MPTMS functionalized glass coverslip in -ssDNA and microRNA directly without the presence of Au TNPs. The obtained SERS and fluorescence signals showed no results (data not shown). These experimental data suggest the formation of a -ssDNA-10b/microRNA-10b-FAM duplex. Metallic nanostructures are capable of enhancing fluorescence signal through the PEF mechanism, in which LSPR properties of the nanostructure increase the radiative decay rate of the fluorophore.<sup>43</sup> To reduce fluorescence quenching due to FRET processes between the nanostructure and fluorophore, it is important that the fluorophore resides within a particular distance from the metal surface. Recent studies showed that a distance of 7.0 to 16.2 nm, corresponding to 18 to 45 nucleotides, between Au nanostructures and fluorophores provides the strongest fluorescence enhancement.<sup>28</sup> Importantly, microRNA-10b contains 23 nucleotides and thus the spacing between FAM molecules and Au TNPs is within the reported distance for highest fluorescence enhancement. To further confirm our confocal-based imaging results, which demonstrate that nanoplasmonic sensors unambiguously detect microRNAs with high specificity, we incubated microRNA-10b-FAM-attached nanoplasmonic sensors in a RNase H enzyme

solution. This solution is expected to break the -ssDNA/microRNA duplex, removing the attached microRNA, to regenerate the original sensor. As predicted, following RNase H incubation, the fluorescence signal completely disappears, as shown in Fig. 4D. As illustrated in Fig. 4E, the fluorescence signal reappears when sensors were again incubated in then microRNA-10b-FAM solution. These results show the fully regenerative capabilities of the developed nanoplasmonic sensors. Sensor regeneration was further confirmed by the decrease or increase of the 1255  $\text{cm}^{-1}$  peak of the C8–H bending mode in the SERS spectra after -ssDNA-10b attachment, 3-MP attachment, microRNA-10b-FAM attachment, incubation in RNase H enzyme, and re-incubation in microRNA-10b-FAM (Fig. S3†).

Next, we investigated the specificity of our label-free assay for microRNA-10b and microRNA-96 detection using the SERS technique, as shown in Fig. 4F and Fig. S4.† Here, the nanoplasmonic sensors, which are capable of detecting microRNA-10b, were incubated in a mixture of microRNA-96, microRNA-145, microRNA-143, and microRNA-490-5p. The levels of these microRNAs are commonly altered in patients with a history of bladder cancer.<sup>13</sup> SERS analysis shows that the intensity of the C8–H bending mode at 1255  $\text{cm}^{-1}$  does not increase when compared to the original nanoplasmonic sensors. This is because microRNA-96, microRNA-145, microRNA-143, and microRNA-490-5p are non-complementary



**Fig. 5** Nanoplasmonic sensor-based liquid biopsy for cancer detection. (A) SERS-based microRNA quantification in six different bladder cancer-related patient plasma samples. microRNA-10b and microRNA-96 are shown in red and blue bars, respectively. (B) Examination of false test resulting by measuring SERS intensities for two different patients containing the highest (patient 2) and the lowest (patient 5) microRNA-10b levels. Red solid bars represent the Raman intensity of C8–H bending stretch at  $1255\text{ cm}^{-1}$ , blue dashed bars after regeneration of nanoplasmonic sensors through RNase H treatment, and green dotted bars are after incubation of nanoplasmonic sensors in 100 nM microRNA-10b-FAM solution. (C) A representative fluorescence confocal image of the nanoplasmonic sensor after incubating in 100 nM microRNA-10b-FAM solution. (D) Examination of false test resulting by measuring SERS intensities for two different patients containing the lowest (patient 3) and the highest (patient 6) microRNA-96 levels. The bars are color coordinated as shown in panel (B). Scale bars is  $50\text{ }\mu\text{m}$ .

to -ssDNA-10b. However, when the same nanoplasmonic sensor was incubated in the complimentary microRNA-10b solution, an increase in SERS intensity of the C8–H bending mode is observed. Thus, by measuring the Raman intensity in the presence of either non-complimentary/complementary microRNAs, we can unequivocally conclude the high specificity of our nanoplasmonic sensors towards targeted microRNAs. Taken together, our multimodal fluorescence and SERS analyses not only confirm the high specificity of the nanoplasmonic sensors towards the microRNA assay, but also demonstrate the reusability of the newly developed label-free technique. Sensor regeneration is a particularly an important aspect as the ability for repeated measurements not only provides unmatched accuracy, but also it is commercially important in low- and middle-income countries.

To demonstrate the translational aspects of our newly developed technology for clinical POC cancer diagnostics, we successfully assayed two oncogenic microRNAs (microRNA-10b and microRNA-96) directly in plasma from metastatic bladder cancer patients. Fig. 5A demonstrates microRNA-10b and microRNA-96 levels in six patients. It should be noted that all patients were categorized in the metastatic stage; however, the level of microRNA varies from one to another. To mitigate false positive and false negative test results, we re-analyzed the two

nanoplasmonic sensors that measured the highest (patient 2 shows a high microRNA-10b level that could arise from false positive response) and the lowest (patient 5 shows a low microRNA-10b level that could lead to false negative response) microRNA-10b concentrations. It is expected that both sensors that were incubated in the patient plasma containing -ssDNA/microRNA-10b duplexes. As shown in Fig. 5B, SERS intensities decrease following RNase H enzyme treatment. Re-incubation of the regenerated sensors into microRNA-10b-FAM results in nearly the same SERS intensity values, as shown in Fig. 4F. Finally, we also characterized the nanoplasmonic sensor containing microRNA-10b-FAM by fluorescence confocal microscopy, which reveals excellent specificity (Fig. 5C). A similar SERS study was also conducted for microRNA-96 involving patient 3 and 6 (see Fig. 5D) as described for microRNA-10b.

To the best of our knowledge, this is the first example where a SERS-based technique has been successfully implemented in the microRNA assay of clinically relevant samples without any sample processing. Experimental data confirm that the developed nanoplasmonic sensors did not compromise sensitivity or specificity aspects upon incubation in patient plasma, and the concentrations that were determined are true values.

## Conclusion

In conclusion, we have designed and fabricated SERS-based, highly sensitive, specific, and regenerative nanoplasmonic sensors for the first time. These sensors were capable of quantifying oncogenic microRNAs at sub-fg  $\mu\text{L}^{-1}$  concentrations directly from patient plasma, thus obviating the complications associated with current gold-standard qRT-PCR-based technologies. The unique LSPR properties of Au TNPs were utilized to achieve a combined dual-detection SERS and fluorescence analysis that allows mitigation of false test results. The newly developed, label-free microRNA assay technique has unmatched potential to expand SERS-based biomarker quantification in the early detection of cancers through liquid biopsies. Importantly, this holds promise in advancing POC clinical diagnostics, which could be performed using a handheld/portable Raman instrument. Taken together, we believe our multimodal, innovative detection approach, which has been validated using plasma from bladder cancer patients samples, should also be applicable to other cancers (e.g., colon, breast, prostate, and pancreatic) as well as various diseases (e.g., cardiovascular, Alzheimer, and infectious) that involve circulating nuclei acids (e.g., DNAs, long non-coding RNAs, and microRNAs) as biomarkers.<sup>8</sup>

## Conflicts of interest

There is no conflicts to declare.



## Acknowledgements

This work is supported by the National Science Foundation, award number CBET-1604617. We thank Dr Daniel Minner for valuable comments and suggestions. We would like to thank Indiana Clinical and Translational Sciences Institute, which was supported, in parts, by Award Number UL1TR002529 from the National Institutes of Health, National Center for Advancing Translational Sciences, Clinical and Translational Sciences Award.

## References

- 1 J. Das and S. O. Kelley, *Angew. Chem.*, 2020, **59**, 2554–2564.
- 2 D. P. Bartel, *Cell*, 2004, **116**, 281–297.
- 3 A. Esquela-Kerscher and F. J. Slack, *Nat. Rev. Cancer*, 2006, **6**, 259–269.
- 4 D. J. Lockhart and E. A. Winzeler, *Nature*, 2000, **405**, 827–836.
- 5 J. Lu, G. Getz, E. A. Miska, E. Alvarez-Saavedra, J. Lamb, D. Peck, A. Sweet-Cordero, B. L. Ebert, R. H. Mak, A. A. Ferrando, J. R. Downing, T. Jacks, H. R. Horvitz and T. R. Golub, *Nature*, 2005, **435**, 834–838.
- 6 F. Petrocca and J. Lieberman, *RNA Biol.*, 2009, **6**, 335–340.
- 7 J. Wang, J. Chen and S. Sen, *J. Cell. Physiol.*, 2016, **231**, 25–30.
- 8 Z. Jin, D. Geißler, X. Qiu, K. D. Wegner and N. Hildebrandt, *Angew. Chem., Int. Ed.*, 2015, **54**, 10024–10029.
- 9 S. Fang, H. J. Lee, A. W. Wark and R. M. Corn, *J. Am. Chem. Soc.*, 2006, **128**, 14044–14046.
- 10 B. N. Johnson and R. Mutharasan, *Anal. Chem.*, 2012, **84**, 10426–10436.
- 11 M. Wanunu, T. Dadosh, V. Ray, J. Jin, L. McReynolds and M. Drndić, *Nat. Nanotechnol.*, 2010, **5**, 807–814.
- 12 G. K. Joshi, S. Deitz-McElyea, T. Liyanage, K. Lawrence, S. Mali, R. Sardar and M. Korc, *ACS Nano*, 2015, **9**, 11075–11089.
- 13 T. Liyanage, A. N. Masterson, H. H. Oyem, H. Kaimakliotis, H. Nguyen and R. Sardar, *Anal. Chem.*, 2019, **91**, 1894–1903.
- 14 G. K. Joshi, S. Deitz-McElyea, M. Johnson, S. Mali, M. Korc and R. Sardar, *Nano Lett.*, 2014, **14**, 6955–6963.
- 15 D. Graham, B. J. Mallinder and W. E. Smith, *Angew. Chem., Int. Ed.*, 2000, **39**, 1061–1063.
- 16 Y. C. Cao, R. Jin and C. A. Mirkin, *Science*, 2002, **297**, 1536–1540.
- 17 A. J. Qavi and R. C. Bailey, *Angew. Chem., Int. Ed.*, 2010, **49**, 4608–4611.
- 18 J. L. Abell, J. M. Garren, J. D. Driskell, R. A. Tripp and Y. Zhao, *J. Am. Chem. Soc.*, 2012, **134**, 12889–12892.
- 19 J. D. Driskell and R. A. Tripp, *Chem. Commun.*, 2010, **46**, 3298–3300.
- 20 L. Qi, M. Xiao, X. Wang, C. Wang, L. Wang, S. Song, X. Qu, L. Li, J. Shi and H. Pei, *Anal. Chem.*, 2017, **89**, 9850–9856.
- 21 S. Tian, O. Neumann, M. J. McClain, X. Yang, L. Zhou, C. Zhang, P. Nordlander and N. J. Halas, *Nano Lett.*, 2017, **17**, 5071–5077.
- 22 P. S. Gaikwad and R. Banerjee, *Analyst*, 2018, **143**, 1326–1348.
- 23 S. K. Vashist, *Biosensors*, 2017, **7**, 62.
- 24 K. M. Mayer and J. H. Hafner, *Chem. Rev.*, 2011, **111**, 3828–3857.
- 25 J. N. Anker, W. P. Hall, O. Lyandres, N. C. Shah, J. Zhao and R. P. Van Duyne, *Nat. Mater.*, 2008, **7**, 442–453.
- 26 P. L. Stiles, J. A. Dieringer, N. C. Shah and R. P. V. Duyne, *Annu. Rev. Anal. Chem.*, 2008, **1**, 601–626.
- 27 J. R. Lombardi and R. L. Birke, *J. Chem. Phys.*, 2012, **136**, 144704.
- 28 Z. Mei and L. Tang, *Anal. Chem.*, 2017, **89**, 633–639.
- 29 J.-F. Li, C.-Y. Li and R. F. Aroca, *Chem. Soc. Rev.*, 2017, **46**, 3962–3979.
- 30 G. K. Joshi, K. A. Smith, M. A. Johnson and R. Sardar, *J. Phys. Chem. C*, 2013, **117**, 26228–26237.
- 31 T. Liyanage, M. Nagaraju, M. A. Johnson, B. B. Muhoberac and R. Sardar, *Nano Lett.*, 2019, **20**, 192–200.
- 32 L. Scarabelli, M. Coronado-Puchau, J. J. Giner-Casares, J. Langer and L. M. Liz-Marzán, *ACS Nano*, 2014, **8**, 5833–5842.
- 33 T. Liyanage, A. Rael, S. Shaffer, S. Zaidi, J. V. Goodpaster and R. Sardar, *Analyst*, 2018, **143**, 2012–2022.
- 34 G. K. Joshi, S. L. White, M. A. Johnson, R. Sardar and P. K. Jain, *J. Phys. Chem. C*, 2016, **120**, 24973–24981.
- 35 G. K. Joshi, K. N. Blodgett, B. B. Muhoberac, M. A. Johnson, K. A. Smith and R. Sardar, *Nano Lett.*, 2014, **14**, 532–540.
- 36 J. D. Driskell and R. A. Tripp, *Chem. Commun.*, 2010, **46**, 3298–3300.
- 37 F. Madzharova, Z. Heiner, M. Guehlke and J. Kneipp, *J. Phys. Chem. C*, 2016, **120**, 15415–15423.
- 38 L. Qi, M. Xiao, X. Wang, C. Wang, L. Wang, S. Song, X. Qu, L. Li, J. Shi and H. Pei, *Anal. Chem.*, 2017, **89**, 9850–9856.
- 39 B. L. Lambert, S. Gronert, F. H. Shurvell and A. D. Lightner, *Organic Structural Spectroscopy*, 2011.
- 40 Z. Wang, S. Ye, N. Zhang, X. Liu and M. Wang, *Anal. Chem.*, 2019, **91**, 5043–5050.
- 41 B. Giese and D. McNaughton, *Phys. Chem. Chem. Phys.*, 2002, **4**, 5161–5170.
- 42 Y. He, X. Yang, R. Yuan and Y. Chai, *Anal. Chem.*, 2017, **89**, 8538–8544.
- 43 *Principles of fluorescence spectroscopy*, ed. J. Lakowicz, Springer, 3rd edn, 2006.