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Toward precision oncology: SERS microfluidic systems for multiplex biomarker analysis in liquid biopsy

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Liquid biopsy-based diagnosis in precision oncology exhibits significant advantages over the traditional tissue biopsies by offering dynamic assessment of tumour heterogeneity, minimally invasive procedures for frequent sampling, and cost-effective tests. Implementation of liquid biopsy-based diagnosis for precision oncology could be the key to provide a confident cancer screening with tailored risk assessment, patient stratification, and real-time monitoring of cancer therapies. To achieve precision oncology with liquid biopsy, the simultaneous analysis of multiple circulating tumour biomarkers is a powerful strategy to establish an accurate signature for each individual patient. Among various developed approaches for tumour biomarker detection, microfluidic devices integrated with surface enhanced Raman scattering (SERS) are emerging as one of the powerful techniques to support precision oncology based on its potential to provide multiplexing and high sensitivity. Particularly, the microfluidic devices provide miniaturised channels for parallel reactions and SERS has the extremely narrow spectra for intrinsic multiplexing. This mini review will focus on recently reported SERS microfluidic techniques, which are capable of simultaneous detection of multiple cancer biomarkers in liquid biopsy and have the promise to be integrated into precision oncology.

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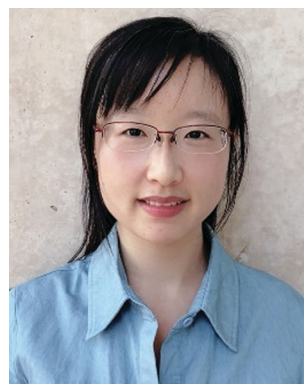
1. Introduction

The emergence of precision oncology in the recent era has radically rejuvenated the field of oncology by introducing the concept of personalized treatment, which analyzes an individual's genetic, epigenetic, or phenotypic characteristics to



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maximize treatment efficiency and minimize side effects. Precision oncology is therefore classified as the cornerstone of next-generation cancer management.¹ Although showing promises in providing personalized treatment, the diagnosis and routine monitoring in precision oncology remain mostly invasive and costly by relying on tissue biopsy to date.^{2,3} In this context, liquid biopsy has emerged as a promising alternative that involves analysis of tumour-derived components in body fluids (e.g., blood, urine, saliva). Due to the advantages of minimally invasive procedures and precise monitoring of tumour biomarkers in real-time, liquid biopsy has shown growing potential to become the next gold standard for cancer diagnostics and therapy monitoring.^{4,5} The recent surge in research for developing liquid biopsy technologies has enabled the discovery of many liquid biopsy-based cancer biomarkers,

such as circulating tumour cells (CTCs), circulating tumour DNA and RNA (ctDNA and ctRNA), soluble proteins, and extracellular vesicles (EVs).⁶ These liquid biopsy cancer biomarkers are found to carry real-time cancer signatures that are displayed on the actual tumour sites.⁷ The analysis of these biomarkers in liquid biopsy could therefore revolutionize the early diagnosis and routine monitoring of cancer resulting in a significant improvement in patient outcome for precision oncology. Despite their immense potential, liquid biopsy technologies have not been established in the clinic yet mostly due to their low sensitivity in accurately identifying the cancer signatures and limited high throughput screening capabilities.⁸ However, these challenges can be addressed with the development of highly sensitive and multiplexed platforms that can specifically identify the biomarkers in trace concentrations in liquid biopsy.⁹

Microfluidic devices have been gaining an increasing attention in the field of biosensing because of their advantageous features, flexible scale up, and mass production at relative low cost. Microfluidic devices are usually derived from micro or nanoscale materials ranging from a few micrometres to nanometres.¹⁰ The integration of 3D nanostructures into microfluidic devices can provide a larger surface area in a miniaturized system and ultra-sensitivity to detect trace concentration of biomolecules.^{11,12} The highly controllable micro/nano environment of such devices is further beneficial as it enables rapid fluid processing and efficient strategies that are required for high throughput biosensing. For instance, it can operate with a low volume of samples, incorporate more than one function in the same device, and enable the detection of multiple targets simultaneously.^{13,14} However, microfluidic devices rely on suitable readouts to detect biomarkers that are captured on the device. Fluorescence-based readouts have been widely used with microfluidic devices for molecular diagnostics. Nevertheless, fluorescence-based methods are limited by the overlapping spectra of the target molecules and photobleaching which significantly limits their sensitivity and multiplexing capability.¹⁵



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Recent years have seen an upsurge in developing surface-enhanced Raman scattering (SERS) based biosensing techniques due to their advantages over fluorescence-based methods. SERS utilizes plasmonic substrates to substantially increase the light scattering resulting in structural characteristics of the molecule bound to or near the plasmonic surfaces.¹⁶ In SERS, there are two main types of surface enhancement mechanisms, *i.e.*, long-range electromagnetic, and short-range chemical enhancement. The electromagnetic enhancement technique is generated by localized surface plasmon resonance which intensifies the scattering intensity of analytes using hot spots, whereas the chemical technique involves bonding interaction of adsorbate and substrate to enhance the cross-section scattering of the analytes.¹⁷ SERS-based detection and characterization approaches have been recognized for their high sensitivity, efficient molecular fingerprint properties, and rapid sample preparation protocols.¹⁸ Significantly, SERS provides narrow spectral width with higher multiplexing capabilities, which makes it a superior readout candidate for biosensing.¹⁹

A multiplex system is particularly important for precision oncology because of the generation of a patient-specific multimolecular profile that can significantly facilitate accurate diagnosis and tailoring of therapy.²⁰ The combination of microfluidics systems with SERS could be an ideal platform that can allow accurate identification of the multiple biomolecules on-chip and significantly enhance the detection capabilities in complex samples with ultra-sensitivity.²¹ As such, the combination of microfluidics and SERS has the potential to address limitations associated with sensitivity and multiplexity and could become the gold standard liquid biopsy technique for applications in precision oncology.

In this mini review, we focus on SERS microfluidic platforms that were evaluated to detect multiple biomarkers in liquid biopsy and could be potentially adapted in precision oncology. As shown in (Fig. 1), we grouped the techniques in different classes based on their three multiplexing types: (i) spatial multiplexing, (ii) barcode-enabled multiplexing, and (iii) label-free multiplexing, and discussed the methods to evaluate their potential for clinical applications.

2. Spatial multiplexing

Spatial multiplexing is a simple form of multiplexing and achieved by creating spatially separated zones for parallel

reactions on the microfluidic system. By mapping the microfluidic system, the spatial information together with the Raman signal enable the biomarker identification. Typically, micro/nanofabrication methods (*e.g.*, lithography) are used to prepare separated compartments (*e.g.*, well structures) or parallel microfluidic channels. The micro/nanofabrication methods allow to integrate micro/nanofabricated structures such as patterning with metallic nanoparticles, which can be used to enhance the Raman signal and support trace biomarker detection in liquid biopsy.¹⁰ Furthermore, droplet microfluidics has also been explored for encapsulation of biomarkers and Raman reporter into individual droplets. The design and fabrication of separate zones for multiplexing and sample delivery to each zone need to be considered carefully. Furthermore, multiple separate zones increase the sample consumption and can also cause additional sample handling errors.

Separated compartments were used by Wang and co-workers for highly sensitive 9-plex detection of miRNA using a microfluidic device with 9 spatially separated zones (Fig. 2a).¹¹ After isothermal amplification of the miRNA, the amplified products were flown through the device. The device had 9 zones that consisted of silver nanoparticles (AgNPs) that were patterned on a silicon substrate. The AgNPs were functionalized with complementary DNA/Raman reporter hairpin probes. After binding of the amplified miRNA to the hairpin, enzymatic hydrolysis resulted in an opening of the hairpin structure that moved the Raman reporter away from the signal enhancing AgNPs, leading to a strong signal decrease. This “signal-on”/“signal-off” detection scheme was highly sensitive and enabled trace miRNA detection of as low as 1 aM in 9 min. This strategy was evaluated for molecular typing of breast cancer, where an 85% concordance with RT-qPCR was found.

Using parallel microfluidic channels, micro RNA (miRNA)²⁰ and soluble proteins^{22,23} were detected. 4 breast cancer-associated miRNAs (*i.e.*, miR-126, miR-191, miR-155, and miR-21) and tumour suppressing miR-125, were detected by Han and colleagues. Each channel comprised of a reaction and detection zone (Fig. 2b).²⁰ After labelled DNA probes were immobilized in the reaction zones, the sample was added for DNA-miRNA complex formation by complementary hybridization. Subsequently, the complexes were desorbed and passed to the Raman detection chamber. The small features of the microfluidic device and dense immobilization of the DNA probes enabled highly efficient miRNA capture from as little as 2 μ L of serum

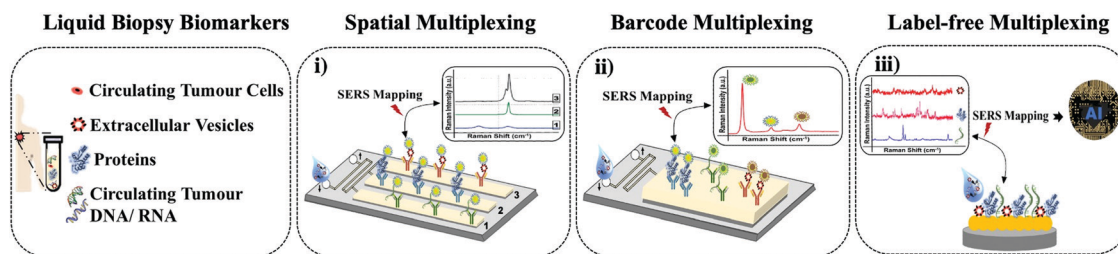


Fig. 1 Schematic of multiplexing in SERS microfluidic systems. Multiplex biomarker analysis can be achieved by (i) spatially separating microfluidic compartments, (ii) using SERS nanoparticle barcodes, or (iii) intrinsic Raman fingerprinting supported by artificial intelligence (AI).



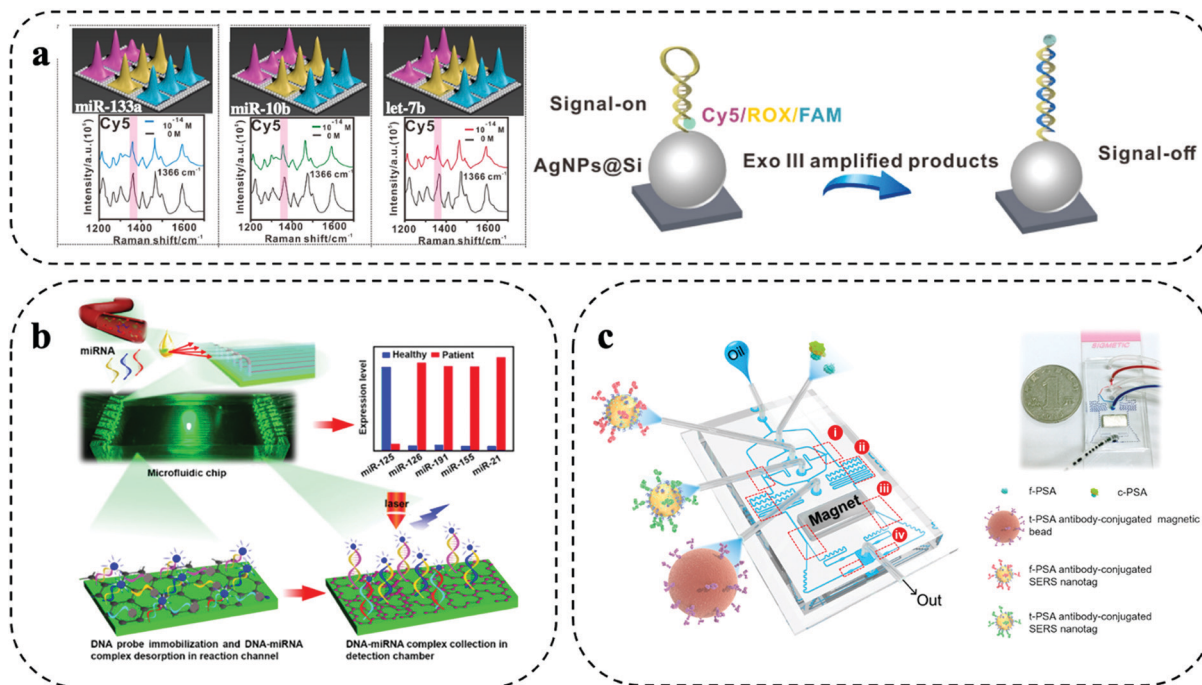


Fig. 2 Spatial multiplexing on SERS microfluidic systems. (a) Separated compartments for 9-plex detection of miRNAs using isothermal amplification. Modified from ref. 11 with permission. Copyright 2021, Elsevier (b) parallel microfluidic channels for 5-plex miRNA detection in a device modified with graphene oxide and poly-L-lysine. Adapted from ref. 20 with permission. Copyright 2021, American Chemical Society (c) droplet microfluidic encapsulation to detect parallelly prostate-specific antigen (f-PSA) and total PSA (t-PSA) in prostate cancer patients. Reproduced with permission from ref. 22. Copyright 2018, Elsevier.

and provided a detection limit of 0.146 aM without requiring RNA amplification. Evaluated on 7 breast cancer patient samples and 3 healthy controls, increased levels of miR-126, miR-191, miR-155, and miR-21 were found in patients, while miR-125 was downregulated compared to the healthy samples.

Similarly, three parallel microfluidic channels were used by Cui *et al.* for 3-plex detection of soluble proteins CA153, CA125, and CEA that are indicated in breast cancer.²⁴ To achieve highly sensitive biomarker detection, the microfluidic channels were patterned with AgNPs that were conjugated to Raman reporters and biomarker-specific antibodies. The AgNPs enhanced the Raman signal by approx. two orders of magnitude compared to the same strategy without AgNPs. Controlling the process to achieve a uniform patterning of the microfluidic channels with metallic nanoparticles is important for Raman signal enhancement and was investigated in detail by Zhang and colleagues.²⁵ They found that a uniform patterning could be easily achieved through a spiral microfluidic geometry. High Raman signal enhancement can also be achieved using bimetallic substrates such as Au–Ag alloys, providing SERS enhancement factor of 10^7 . Such strong enhancement factors are ideally suited for trace biomarker analysis in parallel microfluidic channels. Waluk *et al.* exploited microfluidic channels coated with a bimetallic Ag–Au surface for trace detection of cytokines (*i.e.*, IL-6, IL-8, and IL-18) that are implicated in breast cancer.²¹ The three cytokines were captured in three separated microfluidic channels. The strong and homogenous signal enhancement also reduced Raman signal variation and

contributed to the good assay reproducibility and picogram level detection sensitivity.

Using biomarker encapsulation by droplet microfluidics, Choo *et al.* developed a screening approach for prostate cancer based on the simultaneous detection of free prostate-specific antigen and total prostate specific antigen.²² The microfluidic device integrated a fully automated sandwich immunoassay including droplet generation and immunoreaction, formation of magnetic immunocomplexes, magnetic isolation of immunocomplexes, and SERS detection (Fig. 2c). The simultaneous detection of free and total prostate-specific antigen was achieved by splitting the sample into two droplet streams for separate incubation with SERS nanoparticles conjugated with antibodies against either free or total prostate-specific antigen. The immune-complex was then completed by incubation with antibody-functionalized magnetic beads to pull-down the antigen-SERS nanotag complex for read out. The automated process was completed within 10 min and enabled sensitive biomarker detection in the range of 0.05 to 100 ng mL⁻¹. SERS droplet microfluidics has also been applied to the study prostate cancer cells.²⁶

3. Barcode-enabled multiplexing

Barcode-enabled multiplexing refers to *in situ* detection of multiple biomarkers in non-confined space within the microfluidic system. Typically, the biomarkers are captured by



antibodies immobilized on the microfluidic system, which is followed by a labelling step with a cocktail of target antibodies and Raman reporters co-conjugated SERS nanotags. By varying Raman reporters and target antibody, simultaneous capture and quantitative detection of multiple biomarkers can be achieved.^{10,19} Two main considerations are needed in the selection of Raman reporters. Firstly, the Raman reporters are expected to have separate characteristic peaks without significant spectral overlap. For example, Raman reporter MPY and TFMBA have unique peaks at 1005 cm⁻¹ and 1375 cm⁻¹, respectively. As these two peaks are far away from each other, no mutual interferences can be observed due to spectra overlap. Secondly, the Raman reporters are required to have strong signals to indicate the low abundance of biomarkers. Due to the diverse enhancement capability, the Raman reporters having higher signals have the benefit of sensitively sensing targets for early detection. A novel method was reported by Li *et al.* for effective biomarker monitoring during nasopharyngeal carcinoma (NPC) using SERS sensing. They developed unique SERS reporters using metal carbonyls for enhancing the sensitivity and provided highly multiplex information during the sample testing.²⁷ SERS nanotags with the use of different plasmonic nanostructures including gold nanorod (AuNRs), gold@silver (Au@Ag) nanorods, iron oxide-gold (IO-Au) core shell, gold nanoparticles (AuNPs), gold-silver (Au-Ag) alloy nanoboxes were reported for high-throughput assays with highly selective and sensitive detection. The barcode-enabled multiplexing offers a comprehensive target detection capability, which makes it well suited for wide range of clinical applications from early cancer diagnosis to profiling cancer heterogeneity at a single cell level to monitoring targeted drug therapies for precision oncology. Limitations of barcode-enabled multiplexing are the higher spectral complexity and potential Raman peak overlap. Furthermore, despite the claimed high multiplexing capability of Raman detection, the demonstration of simultaneous measurement of more than 6 biomarkers is rare in liquid biopsy. To deploy the rich Raman spectra for higher multiplexing detection, spectral decoding or deconvolution can be used to reduce signal overlap for identification. In addition, artificial intelligence (AI) based analysis is also increasingly used for Raman data processing.^{28,29} Another strategy to allow higher multiplexing is the use of novel Raman reporters that have lesser Raman peaks to avoid mutual interferences. For instance, research efforts are geared toward the synthesis of novel Raman reporters with few peaks (*e.g.*, one) located beyond 2000 cm⁻¹. For instance, Luo and colleagues, and Liang *et al.* explored the silent region with of the Raman spectrum ranging from ~2050–2150 cm⁻¹ as an extension strategy of the available Raman reporters, which has the promise of improving multi-biomarker detection.^{30,31}

Xiong and colleagues reported a SERS integrated microfluidic chip (MiChip) based on bio-conjugated magnetic nanochains (Magchains) for simultaneous multiplexing of three clinical serum protein biomarkers (*i.e.*, CEA, AFP, PSA) expressed in prostate, colorectal and liver cancer (Fig. 3a).¹³ The integrated MiChip featured various compartments such as fluid ports for sample inlet and waste outlet, mixing chamber, detection

chamber, and two pneumatic microvalves for controlling fluid flow in/out of mixing chamber. For the MiChip assay, the Magchains are conjugated with capture antibodies and SERS nanoprobe, the nanoprobe were prepared using AuNRs synthesized with Raman reporters and specific detection antibodies. The MiChip assay involved the collection of unprocessed liquid specimen, antibody conjugated Magchains and SERS nanoprobe in the mixing chamber followed by mixing driven by Magchains (*i.e.*, synchronised rotation of Magchains in a spinning magnetic field generated by four orthogonal electromagnetic coils). Such a design resulted in the recognition of antigen-antibody and led to the formation of sandwich immune complex of MagChains, target biomarkers and SERS probes. The immune complex was collected in the detection chamber for Raman spectroscopic analysis. For the multiplexed biomarker detection, the specific peaks at 1341 cm⁻¹, 1071 cm⁻¹ and 885 cm⁻¹ of three SERS nanoprobe allowed simultaneous detection of their encoded biomarkers. By measuring the same panel of biomarkers in serum samples, MiChip was able to analyze complex clinical specimens for profiling cancer biomarkers.

Amplification-free SERS biochip integrated with microfluidic channels was demonstrated by Wu *et al.* for simultaneous identification and profiling of multiple point DNA mutations (*i.e.*, KRAS, BRAF, PI3KCA genes) in melanoma and colorectal cancer cell lines (Fig. 3b).³² This barcoding-based *in situ* multiplexing approach was demonstrated on an Au@Ag nanorods SERS substrate assembled in individual microfluidic channel. Three kinds of hair-pin structured nucleic acid – molecular beacon probes were designed with three specific Raman reporter and were covalently linked to the Au@Ag nanorods substrate. The hybridization of target DNA caused a structural change of the molecular beacon probe, and the Raman reporter labelled at the free end were detached leading to low SERS intensity. The intensities of different Raman reporters varied corresponding to the target DNA hybridization. With SERS spectral encoding, signal changes were monitored for simultaneous detection of three DNA mutations in cancer patients. Supervised training algorithm based on classical least square (CLS)-linear discriminant analysis (LDA) was imposed to decode SERS spectra clustering of mutation assay for various cancer cells. SERS biochip demonstrated more than 90% accuracy from analyzing tumour DNAs from cancer patient samples showing its potential for clinical applications.

SERS assay based extracellular vesicles (EVs) phenotype analyser chip (EPAC) was reported by Wang *et al.* for monitoring treatment response in melanoma patients receiving targeted therapy (*i.e.*, BRAF V600E inhibitors) (Fig. 3c).³³ The EPAC based microfluidic system utilized alternating current electrohydrodynamics (ac-EHD) for stimulating the fluid flow within the electrode channels that produced nanomixing strategy for enhancing the target-sensor interaction and to minimize the non-specific bindings. The multiplex assay targets four biomarkers in EVs such as MCSP, MCAM, LNGFR and ErbB3 using uniquely labelled SERS nanotags (*i.e.*, AuNPs conjugated with different Raman reporters and target specific antibodies). EPAC



for breast, lung, and ovarian cancer.³⁵ This ac-EHD based approach provided an enhanced target-sensor interaction and minimized the non-specific binding by using four different SERS barcodes for spectral encoding of multiple cancer protein biomarkers.

Zhang *et al.* demonstrated a SERS immunoassay approach for monitoring epithelial–mesenchymal transition (EMT), a mechanism involved in cancer metastasis, through simultaneous detection of two soluble cadherins biomarkers in breast cancer.³⁶ To enable the parallel cadherin detection, the sample was added on a microchip functionalized with *anti-E-cadherin* and *anti-N-cadherin* antibodies to specifically capture sE-cadherin and sN-cadherin. The captured targets were labelled with duplex SERS barcodes for detection of the targets. Specifically, the duplex SERS barcodes used MBA reporter and DTNB reporter that have characteristic Raman peaks at 1080 cm⁻¹ and 1332 cm⁻¹ to indicate E-cadherin and N-cadherin, respectively. This approach was developed on a microfluidic chip integrated with ac-EHD induced nanomixing for increasing the binding kinetics of the target and different SERS barcode nanotags. Kamińska *et al.* developed a sandwich type triplex SERS immunoassay on a microfluidic device embedded with Ag–Au bimetallic SERS-active substrate for simultaneous detection and monitoring the levels of immune activation of three cytokine interleukins biomarkers majorly responsible for breast cancer initiation and progression.²¹ The use of individual SERS barcoding for each target provided distinct signatures for detection and quantification of multiple biomarkers. SERS enabled microfluidic chip functionalized with modified graphene oxide (GO) platform was reported by Reza *et al.* for the simultaneous detection of three soluble immune checkpoint proteins biomarkers.³⁷ Recombinant nanoyeast single chain variable fragments (NY-scFv) based affinity was introduced as an alternative to monoclonal antibody for capture and different SERS nanotags for multiple target protein detection.

Zhang and colleagues reported an *in situ* approach for CTC isolation combined with spectrally orthogonal SERS aptamer nanovectors for profiling multiple cell membrane protein biomarkers from breast cancer cells.³⁸ The isolated CTCs from multiple breast cancer cells were simultaneously detected by three different DNA aptamers conjugated with the SERS nanovectors by the different target specific SERS intensities. An integrated Raman-Active Nanoprobe (RAN) system combined with microfluidic chip was developed by Cho *et al.*, for simultaneous multiplex detection, isolation, and analysis of circulating cancer stem cells (CCSC) and various subtypes of CTCs in breast cancer cells.³⁹ The simultaneous detection of targets (CD133, EpCAM, EGFR, HER2 and MUC1) was achieved by five different combinations of RANs, prepared by conjugating AuNPs with different Raman reporters and different detection antibodies along with biotinylated double stranded DNA (dsDNA). The isolation was performed using the microfluidic chip system functionalized with avidin–biotin and the restriction enzyme digestion of the dsDNA linker from the RANs. SERS immunoassay approach using a microtube based on magnetic beads and SERS tags was developed by Cheng and co-workers for

determination of the ratio of free to total (f/t) prostate specific antigen for accurate diagnosis of prostate cancer.⁴⁰ The initial part of the assay involved the fabrication of detection probes or SERS nanotags. Briefly, two Raman reporters (*i.e.*, malachite green isothiocyanate (MGITC) and X-rhodamine-5-(and-6)-isothiocyanate (XRITC)) were immobilized on AuNPs, followed by conjugation of two antibodies (f-PSA and c-PSA) using SH-PEG-COOH on the surface of AuNPs. The capture substrate was prepared using carboxylic acid-functionalized magnetic beads to allow EDC/NHS assisted antibody immobilization. Ultimately, this *in situ* approach utilized a microtube for immune complex formation that were captured using a magnetic stand and later transferred capillary tube for SERS detection.¹⁶

As shown in Fig. 4a, Wilson Jr and co-workers demonstrated a combined microfluidic platform-based immunomagnetic enrichment with SERS detection system using magnetic multi-colour SERS nanotags for simultaneous and quantitative detection of CTC using four surface protein biomarkers (*i.e.*, EpCAM, HER2, CD44, IGF1R) on a single breast cancer cell for analyzing CTC subpopulation in clinical sample and monitoring cancer metastasis.⁴¹ For this combined assay, SERS nanotags developed using IO–Au core-shell nanoparticles served as dual function purpose, (i) isolation of CTC bound to the magnetic nanoparticle with the permanent external magnets under the device (*i.e.*, immunomagnetic separation), (ii) subsequent detection of CTCs *via* absorbed Raman reporter under SERS spectroscopy. The on-chip multiplexing assay was performed by synthesizing IO–Au core-shell nanoparticles SERS nanotags with four organic dyes, which included, QSY21, BHQ3, QXL680 and IR740 conjugated with different Raman reporters and target antibody for targeting four different biomarkers in each individual cell. Although Raman reporter displayed different peaks, identification of one distinct peak for each tag when three or more colours needed simultaneously made it a difficult task. Signal deconvolution based on CLS regression method was performed for analysing the levels of each target surface protein biomarkers.

A dual signal enhancement approach using SERS-based droplet microfluidics was reported by Sun *et al.* for simultaneous *in situ* detection of two cytokine biomarkers (*i.e.*, VEGF, IL-8) secreted by a single-cell for studying cell–cell heterogeneity (Fig. 4b).⁴² This assay was built on a droplet based- cross-typed microfluidic SERS platform. The device involved the formation of water-in-oil microdroplets by cutting aqueous phase with oil phase for encapsulation of a single cell. The detection of cytokine biomarkers in the single cell was based on magnetic-field amplified SERS sandwich immunoassay. The immune particles complexes for the detection were prepared by conjugating monoclonal antibodies to silver nanoparticles and polyclonal antibodies to magnetic beads along with two different synthesized Raman active reporters for individual target spectral encoding of two cytokine biomarkers. The dual signal enhancement was obtained with the higher plasmon effect produced by adjacent AgNPs and magnetic field-induced aggregate amplification effect by the magnetic beads for analysing multiple cytokines from single cell. This provided the



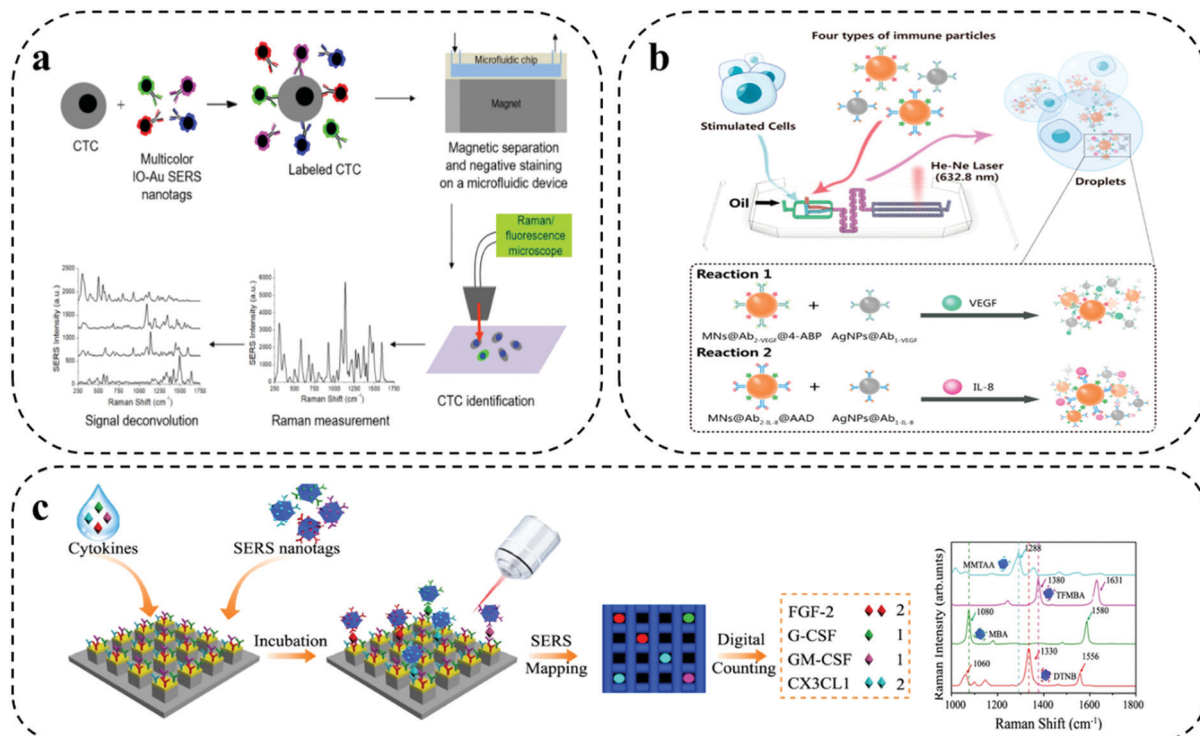


Fig. 4 Barcode-enabled multiplexing on SERS microfluidic systems. (a) IO–Au core–shell nanoparticles based magnetic microfluidic chip for detection and quantification of four surface protein biomarker in individual CTC. Reproduced with permission from ref. 41. Copyright 2020, American Chemical Society. (b) SERS–Droplet microfluidic based dual signal enhancement for identification of two cytokines from a single cell. Adapted from ref. 42 with permission. Copyright 2019, American Chemical Society. (c) Single molecule capture nanopillar platform using SERS nanoboxes for detection of four cytokine biomarkers. Adapted from ref. 43 with permission. Copyright 2021, Springer Nature.

assay with 75 times enhanced SERS signal with a limit of detection at 1.0 fg mL^{-1} in one droplet.

As shown in Fig. 4c, Li and colleagues developed a digital nanopillar platform integrated with SERS barcode detection system for multiplex quantification of single cytokine molecule (*i.e.*, FGF-2, G-CSF, GM-CSF, CX3CL1) to track immune toxicities in melanoma patients receiving immune checkpoint blockade therapy.⁴³ Single cytokine molecule counting of four targets on nanopillar platform was based on three key concepts: (i) Rayleigh criterion separation for fabricating discrete single cytokine nanopillar chip with discretely separated compartments; (ii) probability-driven Poisson distribution for control of target concentration; and (iii) single particle active SERS nanotags for confocal SERS mapping. The gold topped nanopillars were functionalized with target specific capture antibodies for single cytokine capture and SERS nanotags were used for target recognition *via* SERS spectroscopy. The preparation of SERS nanotags involved the conjugation of different Raman reporters and four target antibodies on the Au–Ag alloy nanoboxes. The highly sensitive detection of four cytokines enabled the successful detection of immune toxicities in treated melanoma patients before the onset of symptoms. Furthermore, statistical analysis was performed with the inclusion of four cytokine biomarkers in LDA for wider and accurate patient sample analysis. The outcome from the LDA analysis showed a clear discrimination for the patients with severe cytokine level

and was able to recognize minor differences in grade 1 immune toxicities patients.

4. Label-free multiplexing

Different from the previously discussed barcode-enabled multiplexing strategy, label-free multiplexing approach utilizes the distinct biomolecule-specific Raman signals for establishing the patient signature for personalized identification.⁴⁴ Due to the diverse components and molecular structures of biomolecules, their vibrational and rotational information constitute the unique Raman fingerprints for differentiation.⁴⁴ The direct utilization of the intrinsic Raman spectra from biomolecules for multiplexed analysis features streamlined operation. However, the relatively weak Raman peaks from biomolecules necessitates the employment of high performance plasmonic nanostructures to provide remarkable Raman enhancement.¹⁰ Typically, by integrating highly efficient plasmonic nanostructures with microfluidic chips, the label-free multiplexing analysis can be conducted by two approaches: (1) generating specific biomolecule Raman spectra with the assistance of capture probes to purify the targets; and (2) producing the ensemble spectra as a signature from a mixture, followed by performing multivariate statistical analysis for classification. Due to complex Raman spectra and interpretations, identifying biomarkers based on their intrinsic Raman



fingerprint becomes increasingly difficult. Furthermore, the typical low Raman active cross-section of biomolecules (*e.g.*, protein) results in the weak identifiable signals provides lower sensitivity. With the advantage of revealing subtle distinctions, deep learning and artificial neural networks are seeing a growing impact in the effective processing of big sensing data for complex spectra allowing classification and identification of specific targets.^{28,29} Particularly, with artificial intelligence (AI) for data analysis, the collection and construction of a SERS fingerprint database can have an increased opportunity for future clinical application.⁴⁵ For instance, Shi and co-workers set up a label free SERS database of DNA and developed AI-based discrimination of tumor suppressor genes. On the AgNP-deposited silicon wafer SERS chip, they first collected three kinds of tumor suppressor genes with different base lengths (*i.e.*, p16, p21, and p53 fragments). The collected SERS spectra were then processed to extract independent parameters (*i.e.*, peak number and peak height) as the training data. A deep neural network (DNN) model was constructed with an input layer, multiple hidden layers, and an output layer to train the input data. Particularly, the DNN model involved a back-propagation algorithm and internal adjustable parameters to compute the data in each layer and in turn trained the machine. Verifying the DNN model on new data, both the single DNA targets and DNA mixtures showed recognition with an accuracy rate of 90.28%.⁴⁵

In the first strategy of generating certain biomarker Raman signals for cancer detection, Sim *et al.* developed a head-flocked

gold nanopillar chip to profile three miRNA biomarkers (*i.e.*, miR-10b, miR-21, and miR-373) (Fig. 5a).⁴⁶ Specifically, a gold nanopillar chip was fabricated by depositing gold *via* an electron beam evaporation onto a mask less etched Si nanopillar structure. After functionalized with capture DNA probes, the gold nanopillar chip was able to selectively capture the target miRNAs. Importantly, the gold nanopillars can spontaneously form the “flock” structures with the liquid flowing through the chip to induce the elastocapillary force, which was defined as the “head-flocking” effect. Sufficient hot spots generated by the head-flocking effect greatly contributed to the successful enhancement of three miRNA Raman signals. This label-free microfluidic SERS assay achieved ultra-high sensitivity down to fM level for three miRNA targets and quantified miRNAs directly from simulated patient serum, indicating the potential for future cancer diagnosis.

As for the second strategy of directly interrogating the sample, the ensemble Raman spectra reflects a panel of targets, which is difficult to clearly assign the peaks to specific biomarkers. Nevertheless, with the help of chemometric analysis, the Raman signature from the entire biomolecules is able to distinguish cancer samples from health ones. For instance, Jalali and co-workers reported the label-free Raman measurement of glioma extracellular vesicles (EVs) through integrating the plasmonic nanobowtie structures with a microfluidic device (Fig. 5b).⁴⁷ By leveraging the locally enhanced electromagnetic field provided *via* the nanobowtie-shaped antennas and the

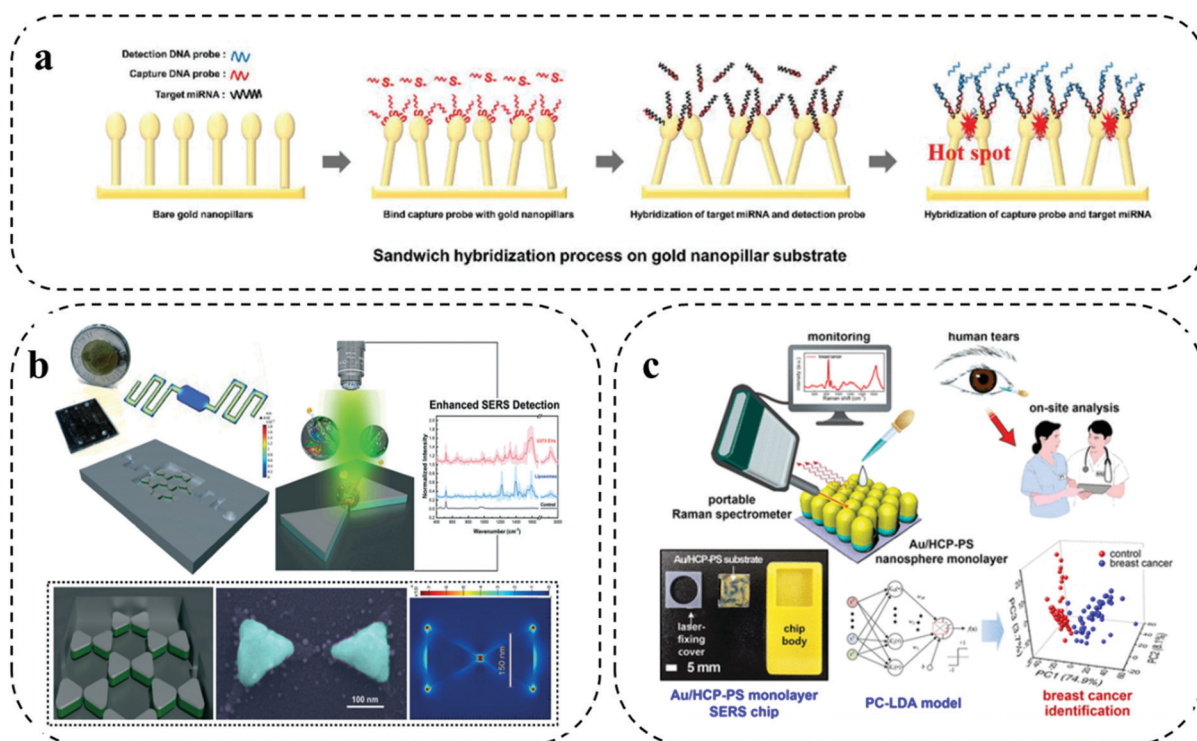


Fig. 5 Label-free multiplexing on SERS microfluidic systems. (a) Gold nanopillar chip for specifically profiling three miRNA biomarkers. Reproduced with permission from ref. 46. Copyright 2019, Royal Society of Chemistry. (b) Nanobowtie structure integrated microfluidic device to enhance EV Raman signals. Adapted from ref. 47 with permission. Copyright 2021, Royal Society of Chemistry (c) Au/HPC-PS SERS chip for generating protein Raman signatures from human tears. Reproduced with permission from ref. 48. Copyright 2020, American Chemical Society.



controllable delivery of EVs *via* the microfluidic flow manipulation, the device can obtain uniform and reproducible Raman spectra. Significantly, the subsequent principle component analysis (PCA) of the Raman spectra unambiguously differentiated the clusters of cancerous cell lines (*i.e.*, U373 and U87) from non-cancerous cell line (*i.e.*, NHA), suggesting the potential of this label-free SERS microfluidic platform in future cancer diagnosis.

Recently, Kim *et al.* demonstrated a label-free SERS microchip for on-site breast cancer detection using human tears (Fig. 5c).⁴⁸ The SERS microchip was fabricated by depositing a gold layer on a free-standing hexagonal-close-packed polystyrene monolayer (*i.e.*, Au/HPC-PS SERS chip). Using human tears as a non-invasive source, the Au/HPC-PS SERS microchip showed the feasibility of generating fingerprint Raman signals from breast cancer patients and healthy controls, which was identified by the principle component linear discriminant analysis (PC-LDA). Furthermore, the leave-one-out cross-validation (LOOCV)-assisted PC-LDA analysis showed that the Au/HPC-PS SERS microchip label-free tear detection can achieve a clinical sensitivity and specificity of 92% and 100%, respectively. Having demonstrated the possibility of this technique in a small cohort of patient samples, a further verification of the performance in a larger patient cohort can help the assay transit into clinical use.

Using highly active silver nanoparticles and a droplet-based device, Hassoun and colleagues designed a microfluidic chip to diagnose different types of leukemia.⁴⁹ In this system, tiny nanoliter droplets were firstly generated by a mixture of silver nanoparticles, KCl, and leukemia cell lysate, in which KCl was used as aggregation agent to create hot spots to enhance Raman signals. Within the confined microfluidic channel, the samples in the droplets were well mixed and allowed the acquisition of SERS spectra using a short integration time (*i.e.*, 1 s). The generated reproducible spectra were mainly associated to the protein and lipids adsorbed to silver nanoparticle, which enabled a robust classification (*i.e.*, accuracy above 99%) of three leukemia cell lines after applying to PCA. This work provided a good example of generating reproducible SERS spectra from nanoparticle colloids by combining with microfluidic devices.

5. Outlook

SERS microfluidic platforms have shown their potential to create multimolecular profiles in liquid biopsy. Table 1 provides an overview of three different multiplexing approaches. Current state of the arts such as polymerase chain reaction

Table 1 Overview of spatial multiplexing, barcode-enabled multiplexing, and label-free multiplexing

Multiplexing system	Biomarkers	Type of cancer	Detection method – Raman reporters (cm ⁻¹)	Advantages	Disadvantages	Ref.		
Spatial multiplexing	miRNA	Breast	FAM – 1332 Cy5 – 1366 ROX – 1503	High sensitivity	High sample volume	11		
	Proteins	Breast	DTNB – 1333 4MBA – 1080 2NAT – 1378				Design complications	24
	Proteins Cytokines	Prostate Breast, Lung	MGITC – 1612 FC – 1178 p-MBA – 1078 DTNB – 1326				Handling errors	22 21
Barcode-enabled multiplexing	Proteins	Prostate, Colorectal, Hepatocellular	NTP – 1341 BTP – 1071 TFTP – 885	High sensitivity	Spectral overlapping	13		
	EVs	Melanoma	MBA – 1075 TFMBA – 1375 DTNB – 1335 MPY – 1000				33	
	DNA	Melanoma Colorectal	Cy5 – 1589 FAM – 1325 ROX – 1493				32	
	CTCs	Melanoma Breast	DTNB – 1337 MBA – 1075 TFMBA – 1175				High multiplexing	15
	CCSC & CTCs	Breast	NBA – 469 QNT – 985 NPT – 1051 MDP – 1096 TP – 1142				39	
Label-free multiplexing	miRNA	Breast	Nil	Label-free Intrinsic spectrum of samples	Lower sensitivity Spectral complexity	41 42 43		
	EVs	Glioblastoma	Nil					
	Human tears	Breast	Nil					



(PCR), enzyme linked immunosorbent assay (ELISA), immunohistochemistry, flow cytometry, cytology, fluorescence *in situ* hybridisation (FISH), or fluorescence microscopy have limitations for biomarker analysis.^{50,51} (For instance, technologies such as ELISA can have long analysis time and low sensitivity, fluorescence signals can suffer from photobleaching *etc.*). Microfluidic based SERS multiplexing platforms may overcome these drawbacks when applied with appropriate SERS strategies and microfluidic substrates.¹⁸ Yet, to translate these promising platforms from proof-of-concept to the clinical environment, future efforts should gear around strategies that make these platforms amendable for routine diagnosis and monitoring. This will require technology innovation from both the 'fluidic' side and SERS side to suit the clinical needs. From the 'fluidic' side, there were only a few demonstrations of fully automated sample-to-answer platforms. Sample preparation remained to be one of the major hurdles for integration into SERS microfluidic platforms. The automation of the platforms would further be advantageous as it improves assay performance (*e.g.*, reproducibility), which is a key factor for clinical implementation. Microfluidic platforms, and to a lesser degree nanofluidic system, can be produced at relative low cost and in large quantities. Thus, the development of disposable sample-to-answer chips seems feasible.

As for the SERS side, several improvements can be made to go together with the microfluidic systems for the advanced and integrated chips. (1) Handheld Raman spectrometers that implement rapid and *in situ* signal readout should be designed to apply on microfluidic devices. The benefit of using handheld Raman spectrometer in chemical analysis has been emerging in the past decades. A further extension of the portable Raman spectrometer into microfluidic systems should open a new area for oncology analysis by allowing routine and point-of-care biomarker profiling. (2) The generation of reproducible and robust SERS signals is necessary to enhance the inter/intra-laboratory comparison and essential for the broad clinical applications. Particularly, the inconsistent signal is an issue for the Raman intensity-based biomarker quantification, which can witness large signal fluctuations due to the uncontrollable hot spots generated from nanoparticle aggregation. To solve this conundrum, the synthesis of highly uniform and stable plasmonic nanostructures as well as the use of an internal standard Raman reporter as a reference can be two promising strategies. (3) A larger panel of biomarkers is expected to achieve simultaneous analysis as the clinical need of a unique patient signature through screening a constellation of biomolecules. Though SERS has the advantage of high multiplexing capability (*e.g.*, 31 plex), the current Raman barcoded detection confines in a relatively low target number (*e.g.*, 4 or 5). To expand the multi-biomarker detection, the utilization of novel Raman reporters (*e.g.*, alkyne molecules) that have minimal spectral overlap should be explored. As another option, the employment of advanced AI technique to learn and deconvolve the overlapped Raman spectra into identifiable codes can be a prospective approach.

Together, future research toward disposable micro/nano-fluidics and combined with handheld, reproducible, and

higher multiplexing SERS will be critical for the development of fully automated microfluidic SERS platform, which enables sample-to-answer analysis of multiple biomarkers. Such a next generation of microfluidic SERS is envisaged to deliver the detailed molecular profile to support accurate decision making in precision oncology.

6. Conclusion

Precision oncology is challenging the idea of "one solution for all" in terms of cancer treatment. Similarly, one marker-based analysis has also been proven not to be enough for diagnosis and monitoring of a complex disease like cancer. An ultrasensitive multiplexed platform could therefore be the central point of precision oncology for accurate diagnosis and monitoring of cancer at the very early stage. In this mini review, we evaluated SERS enabled microfluidic devices in terms of their sensitivity and multiplexing capability. We found that spatial multiplexing is unable to provide a true multiplexed platform. Label free multiplexing started attracting attention and showing promises supported by artificial intelligence which could overcome the limitations associated with the labelling. However, it is still a long way for label free multiplexing to be able to have a place in clinical settings. We conclude that the integration of SERS barcoded systems with spatial multiplexing could be highly powerful to provide a true multiplexed platform at this stage. Moreover, the barcode enabled multiplexed platforms with controllable and rapid fluid flow, 3D nanostructures as the sensor surface as well as the super plasmonic SERS nanoparticles can achieve the required sensitivity for detecting biomarkers at trace concentrations. Finally, we believe that SERS-microfluidic platforms with the support from AI could deliver the ultimate platform for precision oncology to significantly increase the patient survival rate.

Abbreviations

3D	3 Dimensional
ac-EHD	Alternating current electrohydrodynamics
AFP	Alpha-fetoprotein
AgNPs	Silver nanoparticles
aM	Attomolar
Au-Ag	Gold-silver
AuNPs	Gold nanoparticles
AuNRs	Gold nanorods
BTP	4-Bromothiophenol
CCSC	Circulating cancer stem cells
CEA	Carcinoembryonic antigen
CLS	Classical least square
CTCs	Circulating tumour cells
ctDNA	Circulating tumour DNA
ctRNA	Circulating tumour RNA
CX3CL1	Chemokine (C-X3-C motif) ligand
Cy5	Cyanine 5
DNN	Deep neural network



DTNB	5,5'-Dithiobis (2-nitrobenzoic acid)
EDC/NHS	1-Ethyl-3-(3-dimethylaminopropyl)carbodiimide/ N-hydroxysuccinimide
EGFR	Epidermal growth factor receptor
EMT	Epithelial–mesenchymal transition
EPAC	Extracellular vesicles (EVs) phenotype analyser chip
EPCAM	Epithelial cellular adhesion molecule
ELISA	Enzyme linked immunosorbent assay
EVs	Extracellular vesicles
FAM	5(6)-Carboxyfluorescein
fM	Femtomolar
FC	Fuchsin
FGF-2	Fibroblast growth factor receptor
FISH	Fluorescence <i>in situ</i> hybridisation
G-CSF	Granulocyte colony stimulating factor
GM-CSF	Granulocyte macrophage colony stimulating factor
GO	Graphene oxide
HER2	Human epidermal growth factor receptor 2
IGF1R	Insulin-like growth factor 1 receptor
IL	Interleukin
IO–Au	Iron oxide-gold
LDA	Linear discriminant analysis
LNGFR	Low affinity nerve growth factor receptor
4MBA	4-Mercaptobenzoic acid
<i>p</i> -MBA	<i>p</i> -Mercaptobenzoic acid
MCAM	Melanoma cell adhesion molecule
MCSP	Melanoma-associated chondroitin sulphate proteoglycan
MGITC	Malachite green isothiocyanate
miRNA	MicroRNA
MPD	4-Mercaptopyridine
MPY	4-Mercaptopyridine
MUC1	Mucin 1
2NAT	2-Naphthalenethiol
NBA	Nile blue A
NPC	Nasopharyngeal carcinoma
NPT	1-Naphthalenethiol
NTP	4-Nitrothiophenol
NY-scFv	Nanoyeast single chain variable fragments
PCA	Principle component analysis
PC-LDA	Principle component linear discriminant analysis
PCR	Polymerase chain reaction
PSA	Prostate-specific antigen
QNT	2-Quinolinethiol
RAN	Raman-active nanoprobe
ROX	X-Rhodamine
SERS	Surface enhanced Raman spectroscopy
SH-PEG-COOH	Thiol-polyethylene glycol-carboxyl
TFMBA	2,3,5,6-Tetrafluoro-MBA
TFTP	2,3,5,6-Tetrafluorothiophenol
TP	Thiophenol
XRITC	X-Rhodamine-5-(and-6)-isothiocyanate

VEGF Vascular endothelial growth factor

Conflicts of interest

There are no conflicts to declare.

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References

- 1 R. Mirnezami, J. Nicholson and A. Darzi, *N. Engl. J. Med.*, 2012, **366**, 489–491.
- 2 V. Prasad, T. Fojo and M. Brada, *Lancet Oncol.*, 2016, **17**, e81–e86.
- 3 F. Di Nicolantonio, P. P. Vitiello, S. Marsoni, S. Siena, J. Tabernero, L. Trusolino, R. Bernards and A. Bardelli, *Nat. Rev. Clin. Oncol.*, 2021, 1–20.
- 4 J. Kaiser, *Science*, 2018, **359**, 259.
- 5 G. De Rubis, S. R. Krishnan and M. Bebawy, *Trends Pharmacol. Sci.*, 2019, **40**, 172–186.
- 6 N. S. Yee, *J. Clin. Med.*, 2020, **9**, 2556.
- 7 M. R. Speicher and K. Pantel, *Nat. Biotechnol.*, 2014, **32**, 441–443.
- 8 C. Alix-Panabières and K. Pantel, *Nat. Biomed. Eng.*, 2017, **1**, 1–3.
- 9 F. Tian, C. Liu, L. Lin, Q. Chen and J. Sun, *TrAC, Trends Anal. Chem.*, 2019, **117**, 128–145.
- 10 J. Li, A. Wuethrich, S. Dey, R. E. Lane, A. A. Sina, J. Wang, Y. Wang, S. Puttick, K. M. Koo and M. Trau, *Adv. Funct. Mater.*, 2020, **30**, 1909306.
- 11 S. Meng, R. Chen, J. Xie, J. Li, J. Cheng, Y. Xu, H. Cao, X. Wu, Q. Zhang and H. Wang, *Biosens. Bioelectron.*, 2021, **190**, 113470.
- 12 C. Wang, Y. Zhang, W. Tang, C. Wang, Y. Han, L. Qiang, J. Gao, H. Liu and L. Han, *Anal. Chim. Acta*, 2021, 338791.
- 13 Q. Xiong, C. Y. Lim, J. Ren, J. Zhou, K. Pu, M. B. Chan-Park, H. Mao, Y. C. Lam and H. Duan, *Nat. Commun.*, 2018, **9**, 1–11.
- 14 C. Liu, Q. Feng and J. Sun, *Adv. Mater.*, 2019, **31**, 1804788.
- 15 K. K. Reza, S. Dey, A. Wuethrich, J. Wang, A. Behren, F. Antaw, Y. Wang, A. A. I. Sina and M. Trau, *ACS Nano*, 2021, **15**, 11231–11243.
- 16 Z. Wang, S. Zong, Y. Wang, N. Li, L. Li, J. Lu, Z. Wang, B. Chen and Y. Cui, *Nanoscale*, 2018, **10**, 9053–9062.
- 17 A. I. Pérez-Jiménez, D. Lyu, Z. Lu, G. Liu and B. Ren, *Chem. Sci.*, 2020, **11**, 4563–4577.



- 18 J. Wu, X. Zhou, P. Li, X. Lin, J. Wang, Z. Hu, P. Zhang, D. Chen, H. Cai and R. Niessner, *Anal. Chem.*, 2021, **93**, 8799–8809.
- 19 J. Wang, K. M. Koo, Y. Wang and M. Trau, *Adv. Sci.*, 2019, **6**, 1900730.
- 20 Y. Chu, Y. Gao, W. Tang, L. Qiang, Y. Han, J. Gao, Y. Zhang, H. Liu and L. Han, *Anal. Chem.*, 2021, **93**, 5129–5136.
- 21 A. Kamińska, K. Winkler, A. Kowalska, E. Witkowska, T. Szymborski, A. Janeczek and J. Waluk, *Sci. Rep.*, 2017, **7**, 1–11.
- 22 R. Gao, Z. Cheng, X. Wang, L. Yu, Z. Guo, G. Zhao and J. Choo, *Biosens. Bioelectron.*, 2018, **119**, 126–133.
- 23 L. Wu, Z. Wang, Y. Zhang, J. Fei, H. Chen, S. Zong and Y. Cui, *Nano Res.*, 2017, **10**, 584–594.
- 24 Z. Zheng, L. Wu, L. Li, S. Zong, Z. Wang and Y. Cui, *Talanta*, 2018, **188**, 507–515.
- 25 Y. Nie, C. Jin and J. X. Zhang, *ACS Sens.*, 2021, **6**, 2584–2592.
- 26 M. R. Willner, K. S. McMillan, D. Graham, P. J. Vikesland and M. Zagnoni, *Anal. Chem.*, 2018, **90**, 12004–12010.
- 27 D. Lin, C.-L. Hsieh, K.-C. Hsu, P.-H. Liao, S. Qiu, T. Gong, K.-T. Yong, S. Feng and K. V. Kong, *Nat. Commun.*, 2021, **12**, 1–16.
- 28 F. Cui, Y. Yue, Y. Zhang, Z. Zhang and H. S. Zhou, *ACS Sens.*, 2020, **5**, 3346–3364.
- 29 L. Guerrini, E. Garcia-Rico, A. O’Loughlen, V. Giannini and R. A. Alvarez-Puebla, *Cancers*, 2021, **13**, 2179.
- 30 D. Liang, Q. Jin, N. Yan, J. Feng, J. Wang and X. Tang, *Adv. Biosyst.*, 2018, **2**, 1800100.
- 31 S. Tian, H. Li, Z. Li, H. Tang, M. Yin, Y. Chen, S. Wang, Y. Gao, X. Yang and F. Meng, *Nat. Commun.*, 2020, **11**, 1–9.
- 32 L. Wu, A. Teixeira, A. Garrido-Maestu, L. Muínelo-Romay, L. Lima, L. L. Santos, M. Prado and L. Diéguez, *Biosens. Bioelectron.*, 2020, **165**, 112392.
- 33 J. Wang, A. Wuethrich, A. A. I. Sina, R. E. Lane, L. L. Lin, Y. Wang, J. Cebon, A. Behren and M. Trau, *Sci. Adv.*, 2020, **6**, eaax3223.
- 34 A. Rotem, O. Ram, N. Shores, R. A. Sperling, A. Goren, D. A. Weitz and B. E. Bernstein, *Nat. Biotechnol.*, 2015, **33**, 1165–1172.
- 35 K. Kamil Reza, J. Wang, R. Vaidyanathan, S. Dey, Y. Wang and M. Trau, *Small*, 2017, **13**, 1602902.
- 36 Z. Zhang, J. Wang, K. B. Shanmugasundaram, B. Yeo, A. Möller, A. Wuethrich, L. L. Lin and M. Trau, *Small*, 2020, **16**, 1905614.
- 37 K. K. Reza, A. A. I. Sina, A. Wuethrich, Y. S. Grewal, C. B. Howard, D. Korbie and M. Trau, *Biosens. Bioelectron.*, 2019, **126**, 178–186.
- 38 Y. Zhang, Z. Wang, L. Wu, S. Zong, B. Yun and Y. Cui, *Small*, 2018, **14**, 1704433.
- 39 H.-Y. Cho, M. K. Hossain, J.-H. Lee, J. Han, H. J. Lee, K.-J. Kim, J.-H. Kim, K.-B. Lee and J.-W. Choi, *Biosens. Bioelectron.*, 2018, **102**, 372–382.
- 40 Z. Cheng, N. Choi, R. Wang, S. Lee, K. C. Moon, S.-Y. Yoon, L. Chen and J. Choo, *ACS Nano*, 2017, **11**, 4926–4933.
- 41 R. E. Wilson Jr, R. O’Connor, C. E. Gallops, E. A. Kwizera, B. Noroozi, B. I. Morshed, Y. Wang and X. Huang, *ACS Appl. Mater. Interfaces*, 2020, **12**, 47220–47232.
- 42 D. Sun, F. Cao, W. Xu, Q. Chen, W. Shi and S. Xu, *Anal. Chem.*, 2019, **91**, 2551–2558.
- 43 J. Li, A. Wuethrich, A. A. Sina, H.-H. Cheng, Y. Wang, A. Behren, P. N. Mainwaring and M. Trau, *Nat. Commun.*, 2021, **12**, 1–12.
- 44 L. A. Lane, X. Qian and S. Nie, *Chem. Rev.*, 2015, **115**, 10489–10529.
- 45 H. Shi, H. Wang, X. Meng, R. Chen, Y. Zhang, Y. Su and Y. He, *Anal. Chem.*, 2018, **90**, 14216–14221.
- 46 W. H. Kim, J. U. Lee, S. Song, S. Kim, Y. J. Choi and S. J. Sim, *Analyst*, 2019, **144**, 1768–1776.
- 47 M. Jalali, I. I. Hosseini, T. AbdelFatah, L. Montermini, S. W. Hogiu, J. Rak and S. Mahshid, *Lab Chip*, 2021, **21**, 855–866.
- 48 S. Kim, T. G. Kim, S. H. Lee, W. Kim, A. Bang, S. W. Moon, J. Song, J.-H. Shin, J. S. Yu and S. Choi, *ACS Appl. Mater. Interfaces*, 2020, **12**, 7897–7904.
- 49 M. Hassoun, J. Rüger, T. Kirchberger-Tolstik, I. W. Schie, T. Henkel, K. Weber, D. Cialla-May, C. Krafft and J. Popp, *Anal. Bioanal. Chem.*, 2018, **410**, 999–1006.
- 50 J. Perumal, Y. Wang, A. B. E. Attia, U. Dinish and M. Olivo, *Nanoscale*, 2021, **13**, 553–580.
- 51 S. Abalde-Cela, L. Wu, A. Teixeira, K. Oliveira and L. Diéguez, *Adv. Opt. Mater.*, 2021, 2001171.

