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Investigation on the Isolation and Detection of

Circulating Tumor Cells Based on

Microfluidics and Surface-Enhanced Raman Scattering Method

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Abstract

Cancer is a leading cause of death and is an important barrier to increase life expectancy in every country of the world. Besides, two thirds of cancer death occurred in less developed countries because of delayed diagnosis and less accessible treatment. The delayed diagnosis would lead to increased death and disability from cancer. Therefore, it is essential to develop a technology for cancer detection with universality and less cost. The common imaging technologies, such as Magnetic Resonance Imaging (MRI), Computed Tomography (CT) and Positron Emission Tomography (PET), can be used for cancer early screening. After early screening, people who have symptoms and signs consistent with cancer, require further identification of cancer by pathological diagnoses. The traditional invasive biopsy would cause psychological burden to patients and is limited by sample collection in deep tumor and sampling bias. It also has risk of tumor metastasis. In recent decades, liquid biopsy technique has developed quickly and attracted more and more attention. Compared to invasive biopsy, liquid biopsy is noninvasive, cheaper, simple for sample collection with minimum risk. The sensitivity and specificity of liquid biopsy technique has also been improved remarkably due to the continuous innovation in molecular biology technology. Liquid biopsy generally includes the detection of circulating tumor cells (CTCs), circulating tumor-derived exosomes and circulating tumor nucleic acids. Circulating tumor cells are tumor cells that shed from solid tumor and circulate in the blood. Circulating tumor-derived exosomes are vesicles that secret from tumor cells containing some ribonucleic acids (RNA) and protein. Circulating tumor nucleic acids are nucleic acids shedding from tumor cells or released by apoptotic cells. Compared to circulating tumor-derived exosomes and circulating tumor nucleic acids, circulating tumor cells are integrated cells carrying more complete information about tumor. Circulating tumor cell (CTC) detection as a burgeoning detection strategy can identify the tumor lesion in the

early stage, and facilitate the understanding of tumorigenesis, tumor progression, metastasis, and drug-resistance. As so far, many technologies for CTC detection have been developed. Generally, CTC detection can be divided into two stages: first is isolation and enrichment of CTCs and second is downstream analysis of CTCs. There are three main challenges exciting in CTC detection: small number of CTCs, complex blood background, and diversified typing of CTCs. To overcome these difficulties, microfluidic method has been applied to improve CTC isolation combined with Raman fingerprint spectra and surface-enhanced Raman scattering (SERS) method to distinguish CTCs from other blood components. Current microfluidic method cannot separate CTCs completely from blood, which requires further downstream analysis to distinguish CTCs from remaining blood cells. Besides, SERS method might be interfered by complex blood components, which would reduce detection sensitivity of SERS method. Therefore, blood samples require pretreatment to reduce the interference of blood cells for SERS detection and improve repeatability and reliability of detection results. In this thesis, SPION-PEI@Au based SERS biological probe and B-TiO₂ based SERS biological probe were prepared for direct detection of CTCs in the blood. To improve the detection sensitivity and accuracy, microfluidic method was combined for blood pretreatment before SERS detection. The specific research contents mainly include the following three parts:

1. Superparamagnetic iron oxide nanoparticles with poly(ethyleneimine) coated with gold nanoparticles (SPION-PEI@Au) were synthesized according to the previous work of the research group. In this work, SPION-PEI@Au based SERS biological probe was applied on CTC detection of clinical blood samples. The SPION@Au-MBA-rBSA-FA SERS biological probe consisted of four parts: SPION-PEI@Au composite nanoparticles as SERS substrate, 4-mercaptobenzoic acid (MBA) as Raman reporter, reduced bovine serum albumin and folic acid (rBSA-FA) to recognize

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folate receptor (FR) on cancer cell membrane. In this work, 32 clinical blood samples from cancer tumor and 3 clinical blood samples from healthy people were detected directly by SPION@Au-MBA-rBSA-FA SERS biological probe.

- 2. Black TiO₂ (B-TiO₂) nanoparticles were used as SERS substrate. B-TiO₂ has advantages of low cost, high spectral stability and reproducibility, strong anti-interference ability, and selective SERS enhancement to target molecules. The synthesized B-TiO₂ showed good SERS enhancement effect and the LOD of the AR molecule on B-TiO₂ can reach to 5×10^{-8} M. The B-TiO₂-AR-PEG-FA biological probe consisted of four layers. The innermost layer was B-TiO₂ nanoparticles with crystal core and amorphous shell structure. The second layer was alizarin red (AR) molecule which was responsible for providing Raman spectral signal. The third layer was a thin NH₂-PEG2000-COOH layer which was used to improve the dispersion of biological probe and to provide binding sites of folic acid (FA) and thus to increase FA grafting rate. The outermost layer was FA molecule. FA was used to specifically recognize cancer cells by folate receptor (FR) on cancer cell membrane. The research results showed that this B-TiO₂ based SRES biological probe has good specificity and detection accuracy with obvious Raman signal. It can distinguish positive FR-expressing cancer cells (MCF-7) from lower FR-expressing cells (A549 and Raw264.7).
- 3. Because of complex blood components, the sensitivity and stability of SERS biological probe would be attenuated without proper blood pretreatment. To improve the performance of SERS biological probe, microfluidic method was added to isolate CTCs before SERS detection. In this strategy, the relatively low specificity and isolation purity of microfilter could be solved by integrating with highly sensitive and highly specific SERS spectra detection, while the microfilter could reduce the interference

of blood background to SERS detection. Besides, SERS-fluorescence dualmodal in situ imaging method proved that this strategy has high specificity with detection limit of 2 cancer cells per milliliter in rabbit blood. Besides, the operation process was simple and high-speed, with detection time less than 1.5 hours. This strategy has also been applied to CTC detection of clinical blood samples and has detected CTCs from blood successfully.

These results illustrated that in addition to noble metal nanoparticles, semiconductor nanoparticles can also be used for SERS detection with good SERS enhancement effect. Although SERS method has good sensitivity in CTC detection, the performance of SERS method would be affected by complex components of blood and improper blood treatment. Microfluidic method was combined to remove the interference of blood cells and to improve the repeatability and reliability of SERS detection. The combination of microfluidic method and SERS detection method could complement their own shortcomings and thus to improve the detection efficiency. This thesis demonstrated that the combination of microfluidic method and SERS detection method and SERS detection method could open new paths for liquid biopsy.

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List of Abbreviations

3D	Three-Dimensional
AFM	Atomic Force Microscope
Ag	Silver
AgFON	Silver Film Over Nanosphere
ALD	Atomic Layer Deposition
AR	Alizarin Red
Au	Gold
B-TiO ₂	Black Titanium Dioxide
CE	Chemical Enhancement
СК	Cytokeratin
Cu	Copper
COPD	Chronic Obstructive Pulmonary Disease
СТ	Computed Tomography
СТС	Circulating Tumor Cell
CNVs	Copy Number Variations
DEP	Dielectrophoresis
DFF	Dean Flow Fractionation
DLD	Deterministic Lateral Displacement
EE	Electromagnetic Enhancement
EF	Enhancement Factor
EMT	Epithelial-Mesenchymal Transition
EpCAM	Epithelial Cell Adhesion Molecule
FA	Folic Acid
FDA	Food And Drug Administration

FISH	Fluorescence In Situ Hybridization
FMSA	Flexible Micro Spring Array
FR	Folate Receptor
GeIMA	Methacrylated Gelatin
GSSG	Oxidized Glutathione
HB	Herringbone
hMAM	Human Mammaglobin
IF	Immunofluorescence
ІНС	Immunohistochemistry
LOD	Limit Of Detection
LSPRs	Localized Surface Plasmon Resonances
MET	Mesenchymal-Epithelial Transition
MRI	Magnetic Resonance Imaging
NAADP	Nicotinic Acid Adenine Dinucleotide Phosphate
N/C	Nucleus To Cytokeratin Ratio
NIR	Near Infrared
NPs	Nanoparticles
OS	Overall Survival
PBMC	Peripheral Blood Mononuclear Cell
PDMS	Polydimethylsiloxane
PET	Positron Emission Tomography
PFS	Progress-Free Survival
PICT	Photoinduced Charge Transfer
SERS	Surface-Enhanced Raman Scattering
SNVs	Single-Nucleotide Variations

- **SPION** Superparamagnetic Iron Oxide Nanoparticles
- **WBC** White Blood Cells

Chapter 1 Introduction

Cancer is a leading cause of death worldwide with increasing incidence and death cases. In 2021, an estimated 19.29 million new cancer cases occurred and almost 9.96 million cancer cases caused death[1]. There has been consistent growth in global cancer burden, which imposes enormous stress of health, emotion and finance on individual, family, community, and health systems. High proportion of the cancer patients in developing countries including China, are not able to access timely or high-quality treatment and diagnosis. The delay in cancer diagnosis and treatment would increase cancer disability and mortality. There is consistent evidence showing that the early diagnosis with timely treatment could reduce cancer mortality[2-5]. Therefore, early diagnosis and treatment are critical important during comprehensive cancer control. If the tumorigenesis and tumor metastasis can be detected earlier by some reliable detection technologies, the tumors can be strangled in the early stage, thereby improving the treatment and prognosis of cancer. However, early clinical symptoms of tumors are normally atypical, thus, the early diagnosis and treatment of cancer diseases is a worldwide challenging problem. The traditional clinical and imaging diagnostic methods, such as Magnetic Resonance Imaging (MRI), Computed Tomography (CT) and Positron Emission Tomography (PET), are difficult to find early tumor tissue and unable to realize effective diagnosis of early tumors. The inconvenient collection of samples caused the stagnant study of solid tumor. In the background of precision medicine, liquid biopsy technology, with advantages of convenient, minimally invasive, and able to observe tumorigenesis dynamically, has developed rapidly and attracted increasing attention nowadays. Benefit from the continuous innovation of molecular biology technology, the sensitivity and the specificity of liquid biopsy have been improved significantly. Considerable evidence has emerged that cancer cells and related issues would release biomarkers such as circulating tumor cells (CTCs), DNA, RNA, and exosomes into circulating blood. The amount and the type of these biomarkers are closely related to tumor tumorigenesis. Circulating tumor cells are tumor cells shedding from solid tumor and circulating in the blood, which has been used as a biomarker in the detection of peripheral blood of cancer[6]. The comparison between conventional invasive biopsy and CTC detection shows that CTC detection has many advantages apart from high specificity. For examples, samples could be collected in a gentled way, so that better patient compliance can be achieved. More importantly, CTC detection is not limited by tumor heterogeneity, because CTCs could shed from primary tumor as well as metastatic tumor. Therefore, they are carrying more complete information about tumor development. Many tumor challenges, such as accurate auxiliary diagnosis, real-time monitoring, therapeutic evaluation, and prognosis judgment of tumors are expected to be achieved[7]. In addition, CTCs carry a wealth of information about tumorigenesis, tumor progression, metastasis, and drug resistance that can help understand and control tumor diseases. However,

CTC detection is limited by its heterogeneity and extremely small number of CTCs (1-10 CTCs per milliliter) in the blood. Therefore, it is necessary to develop more effective CTC isolation and detection methods.

In this chapter, research background and literature review of circulating tumor cell have been reviewed. Based on the literature review, the limitations of current technology would be summarized and considered for research design. Finally, outlines of this thesis focused on the research of SERS and microfluidics-based CTC detection method.

1.2 Circulating tumor cells

Circulating tumor cells are tumor cells shedding from the solid tumor, intravasating into blood and lymphatic vessels, and translocating to distant tissues via circulation[8]. Figure 1.1 showed the origin of CTCs. During the process of tumor development, CTCs would enter circulation system spontaneously or under the influence of external factors, carrying a large amount of information about tumorigenesis, tumor progression, metastasis, and drug resistance[9]. The survived CTCs would cause tumor metastasis by shedding, invasion, intravasation, circulation, extravasation, and formation of secondary tumor[10]. Current studies show that cancer metastasis causes approximately 90% death of cancer. The detected CTCs in peripheral blood presage the possibility of tumor metastasis. Therefore, CTCs have been considered as effective biopsy method and able to realize accurate auxiliary diagnosis, real-time monitoring, therapeutic evaluation, and prognosis judgment of tumors[7].



Figure 1.1: The schematic of metastatic process including (1) CTC invasion into blood vessels from primary tumor; (2) circulation of CTCs in blood vessel; (3) extravasation after CTC adhesion to blood vessel walls; (4) metastasis and formation of secondary tumor[11].

1.2.1 Biological properties of CTCs

The biological characteristics of CTCs are commonly affected by protein expression, gene mutation, single-nucleotide variations (SNVs), or copy number variations (CNVs) of CTCs. The analysis of the biological characteristics of CTCs can clarify the relationship between CTCs and tumor progression, guide precise anticancer therapy, distinguishing CTCs from other cells and evaluate the targeted drug therapy. The half-life period of CTCs shedding from solid tumor of cancer patients is 1-2.4 h in the circulation system, related to the immune reactions of CTCs involved in the circulation system[12]. CTCs derived from epithelial tumors have the characteristics of epithelial-mesenchymal transition (EMT) and mesenchymal-epithelial transition (MET)[13]. In the EMT process, the surface markers of epithelial cells would be downregulated while the markers of mesothelial cells would be upregulated. Thus, tumor cells would be promoted to break away from intercellular adhesion and acquire variability and invasiveness. In the contrast, the MET process enables CTCs from EMT process to reverse and restore the epithelial phenotype, regain the ability of adhesion, and form metastases[14]. Because CTCs were shed from solid tumor, they have the properties of epithelial cell and mesenchymal cell and can express Epithelial Cell Adhesion Molecule (EpCAM) antigen on their surface, which can be used to distinguish CTCs from blood cells. However, isolation methods only based on EpCAM may limit the detection specificity of CTCs from other types of cancers, so organ-specific cell surface markers or stem cell markers have been used to enrich CTCs. For example, in addition to the commonly used EpCAM, the other biomarkers, such as CK19, HER2, ER, PR, and MUC1, are also used in breast cancer diagnosis[15]; EGR, and PTEN are used for prostate cancer diagnosis[16]; and EpCAM, ASGPR1, N cadherin, Vimentin, Gpc3, and AFP are used for hepatocellular carcinoma diagnosis[17]. The related biomarkers of CTCs have been summarized in Table 1.1.

Biomarkers	Related Cancer Type
EnCAM : Epithelial Cell Adhesion Molecule	Breast Cancer, Prostate Cancer,
	Hepatocellular Carcinoma
CK19: Cytokeratin-19	Breast Cancer
HER2: Human Epidermal Growth Factor Receptor-2	Breast Cancer
ER: Estrogen Receptor	Breast Cancer
PR: Progesterone Receptor	Breast Cancer
MUC1: Mucin-1	Breast Cancer
EGR: Early Growth Response Factor	Prostate Cancer
PTEN: Gene of Phosphate and Tension Homology Deleted	Prostate Cancer
on Chromosome Ten	
ASGPR1: Asialoglycoprotein Receptor-1	Hepatocellular Carcinoma
N cadherin: N Cadherin	Hepatocellular Carcinoma
Vimentin	Hepatocellular Carcinoma
Gpc3: Glypican-3	Hepatocellular Carcinoma
AFP: Alpha Fetoprotein	Hepatocellular Carcinoma

Table 1.1: Biomarkers related with CTCs in different types of cancer

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1.2.2 Physical properties of CTCs

Due to the abnormal metabolism and metabolic disorder of tumor cells after entering circulation system, the change of composition of intracellular substances, gene expression and modification, protein synthesis and aggregation of some polar particulate matters lead to changes of CTCs' physical properties, which are different from normal cells and normal tissue cells.

Size and density

There have been many studies measuring the size of tumor cell and blood cell normally by optical microscopy and flow cytometry. Erythrocytes have a diameter of 7-8.5 μ m with volume of 100.6±4 μ m³ and leukocytes are 5-13 μ m in diameter, including granulocytes with diameter of 7.3-13.2 µm, lymphocytes with diameter of 5.2-10.5 µm, and monocytes with diameter of 9.7-10.5 μ m[18]. In general, the average diameter of leukocytes is less than 10 μ m with wide size distribution. The average diameter of cancer cell lines used to validate various CTC isolation techniques ranged from 15 to 25 μ m, which was significantly larger than that of WBCs. However, cells from cancer cell lines may be larger than CTCs from cancer patients with same cancer type. For examples, CTCs of breast cancer have the average diameter of 13 µm while the diameter of breast cancer cell lines is 15-17 µm; CTCs of prostate cancer have the average diameter of $6.16-9.78 \,\mu\text{m}$, which is much smaller than the diameter of prostate cancer cell lines, $10.84-15.92 \ \mu m$ [19]. The size difference between CTCs and WBCs is less clear than the difference between cancer cell lines and WBCs. The size distribution of CTCs and WBCs is partly overlapped[20]. Besides, CTCs from different melanoma cancer patients have different size distribution. The maximum size of CTCs from two patients is larger than 12 μ m while some of CTCs from one patient are smaller than 10 μ m[21]. Therefore, it is limited to only use cancer cell lines treated blood sample for validation of isolation technology. More realistic samples with variable size distribution of cancer cell lines should be carried out for validation.

CTCs can also be isolated from blood cells by density gradient centrifugation system depending on the density difference of different cells[22]. Erythrocytes and granulocytes have the density higher than separation medium and will precipitate at the bottom of the pipe after centrifugation, while monocytes and lymphocyte have the density smaller than separation medium and will float in the upper layer of separation medium or suspended in medium. CTCs are mainly deposited in the monocyte enrichment layer. After using Ficoll-Paque medium with a density of 1.077 g/mL, CTCs might be detected in the Peripheral Blood Mononuclear Cell (PBMC) layer. However, the density of CTCs is not accurately defined and probably is not uniform. Research shows that some CTCs from prostate and pancreatic cancer appeared much heavier and could not be separated with separation medium with density of 1.077 g/mL[23]. Therefore, the isolation methods based on density should be improved.

Morphology and stiffness

The cell morphology of CTCs and WBCs is different. Some studies have shown that CTCs have lower roundness than WBCs[24]. Most CTCs from breast, colorectal, and prostate cancer have the roundness of 1.5 ± 0.6 , 1.5 ± 1.4 , and 1.5 ± 0.8 , respectively, while the roundness of WBCs is 1.8 ± 1.2 . Apart from cell morphology, the deformability difference between tumor cells and blood cells can also be used for isolation. When cells are encountering forces or restrictions, they will deform to pass through the pores which are even much smaller than cells[25-27]. The cell deformability can be defined by Elastic properties, such as stiffness and Young's modulus, which can be investigated by Atomic Force Microscope (AFM)[28], micropipette aspiration[29], laser optical trapping[30], and microfluidic device[31]. The stiffness of cells is represented by Young's modulus, which can be used to determine cell deformability. Previous studies show that the stiffness of tumor cells ranges from 200 to 2000 Pa with that of neutrophils from 156±87 Pa[32]. However, Lee et al. reported that the stiffness of neutrophils is 1548±871 Pa in center region, 686±801 Pa at the leading edges, and 494±537 Pa at trailing edge, which is similar with the stiffness distribution of tumor cells[33]. Besides, the stiffness of nucleus is higher than that of cytoplasm. Nucleus to cytokeratin ratio (N/C) is defined as the ratio of nuclear area to cytoplasmic area[19]. The N/C ratio of CTCs is higher than 0.8 while the N/C ratio of WBCs is much smaller than 0.8[32]. CTCs from breast, colorectal, and prostate cancer have N/C ratio of 0.8 ± 2.8 , 1.0 ± 3.5 , and 1.0 ± 2.1 , respectively[24]. Therefore, CTCs are concerned to be stiffer than WBCs. In additions, many researchers have studied the relationship between cancer metastatic potential and stiffness and found that CTCs with higher stiffness have lower metastatic potential[34].

Electrical properties

The cell capacitance can reflect the area of plasma membrane resulting from cell transport phenomena and metabolism[35]. Therefore, some researchers use total capacitance to analyze the morphology and physiological states of cells, which has been used to quantify heterogeneity of breast cancer cell lines[36]. Different cell types have significant differences in the membrane capacitance, which has been applied on dielectrophoresis (DEP) method to separate different cell types. Cancer cells have different cell capacitance because of their abnormal transport phenomena and metabolism as well. Besides, under the stimulation of external environment, CTCs induce cell gene expression and the change of membrane skeleton structure and adhesion, which affects the surface charge of cell membrane to resist the shear force of blood fluid. It is reported that breast cancer cells have smaller mean association constant of hydrogen ions, which means that CTCs have more negative charges[37]. The difference in electric properties combined with size difference between CTCs and blood cells can facilitate CTC isolation.

1.3 Clinical application of CTCs

In recent years, clinical trials have demonstrated that CTCs have important value for tumor early diagnosis, prognosis evaluation, therapeutic efficacy, tumor stage and monitoring of tumor recurrence or metastasis.

1.3.1 CTCs and early diagnosis

The theory of CTC detection applied in tumor early diagnosis was based on the research results of animal model[38]. Researchers used mice models to study breast cancer, finding that CTCs could be appeared at early stage of primary tumor formation. In the mice models of pancreatic cancer, CTCs was found in the blood circulation before the formation of solid tumor.

However, there are two key issues remaining in the application of CTC detection in tumor early diagnosis. First, the clinical research found that benign disease patients also have the presentence possibility of CTCs. For examples, Tanaka et al. found that 12% of benign lung disease patients could be detected with CTCs[39]; Pantel et al. have also detected CTCs in the 11% of benign colon diseases patients[40]. Second, the number of CTCs in the blood at the early tumor stage is extremely low. In one milliliter of blood, there are 1-10 CTCs with 10⁶ white blood cells and 10⁹ red blood cells[41]. Besides, due to the rarity of CTCs, no relevance has been found between the number of detected CTCs and tumor volume, tumor classification. Therefore, it is important to develop more sensitive methods for CTC detection.

Although there are several difficulties in CTC detection applied in early tumor diagnosis, it has still been proved that CTC detection is a potential method in

term of early tumor diagnosis. Ilie et al. have detected CTCs in patients with Chronic Obstructive Pulmonary Disease (COPD). Besides, in the follow-up detection within 1-4 years, patients with detected CTCs have been diagnosed with early stage of lung cancer by CT and pathologic biopsy[42]. Therefore, CTC detection has important scientific significance and application value for early diagnosis of tumor.

1.3.2 CTCs and prognosis evaluation

The number of CTCs at different stages of tumor are different which could be considered as an analysis basic of prognosis evaluation. Cristofanilli et al. have verified the hypothesis that the level of circulating tumor cells can predict survival in metastatic breast cancer[43]. In this research, they found that the Progress-Free Survival (PFS) and the Overall Survival (OS) of patients correlated with the number of detected CTCs in circulation. Patients with more than 5 detected CTCs per 7.5 ml peripheral blood have obviously shorter survival and worse prognosis. In the following clinical studies, the corresponding relationship between CTC number and prognosis or treatment evaluation has also been proved to be applicable to prostate cancer, colorectal cancer, and other cancer patients[44, 45].

1.3.3 CTCs and therapeutic efficacy

Nowadays, growing number of tumor clinical research have considered CTCs

as an important value in therapeutic efficacy[46]. As mentioned above, Cristofanilli et al. also applied CTC counting to the monitoring of metastatic breast cancer. In 2006, Budd et al. tested 138 metastatic breast cancer patients with imaging detection and CTC counting before and a median of 10 weeks after the initiation of therapy[47]. They find that patients with more than 5 CTCs in 7.5 mL blood were associated with shorter PFS and OS. Besides, by comparing the use of CTCs to radiology for prediction of OS, they found that assessment of CTCs was an earlier, more reproducible indication of disease status than current imaging methods. CTCs may be a superior surrogate end point, as they are highly reproducible and correlate better with overall survival than do changes determined by traditional radiology.

1.4 CTC detection: isolation and downstream analysis of CTCs

90% of malignant tumor was derived from the epithelial cells where the epithelial tissue cells have extremely different gene expression and biological characteristics compared to other tissue cells, such as blood cells. Therefore, most of CTC detection methods have been developed depending on the difference between epithelial tissue cells and blood cells. However, after decades of development, CTC detection is still challenged because of the extremely small number of CTCs in blood and the differences in size, morphology, molecular marker, gene expression and mutation of CTCs caused by the heterogeneity of CTCs[48, 49]. In general, the CTC detection has two

stages: isolation of CTCs (capture and enrichment), and downstream analysis of CTCs (detection and release)[50].

1.4.1 Isolation technology of CTCs

CTC isolation methods have been divided into two major categories based on biological characteristics and physical characteristics. The former is mainly based on CTCs' biological characteristics that are different from normal blood cells such as expression to epithelial cell adhesion molecule (EpCAM), which has been classified as label-dependent methods. The latter is mainly based on physical differences between CTCs and blood cells such as size, density, deformability, and electric properties, which has been classified as labelindependent methods. The label-dependent methods could also be classified into positive isolation and negative isolation. Positive isolation of CTCs is commonly achieved by EpCAM which is a kind of cell surface marker expressed in CTCs. In negative isolation, WBCs are depleted by antibody against CD45 or other leukocyte antigens. The label-dependent isolation has high specificity but is limited by low surface protein expression level and inherent heterogeneity of tumor cells. In contrast, the label-independent isolation does not depend on cell surface marker expression level. Compared to label-dependent isolation, label-independent isolation can be accomplished in a low cost, simple and fast way. However, it suffers from low specificity because of overlap in size, density, and stiffness between CTCs and WBCs. The

CellSearch system was one of CTC detection technologies based on biological characteristics of CTCs and was the first CTC detection product approved by U.S. Food and Drug Administration (FDA). It could be used for CTC detection and enumeration for metastasis breast, colorectal, and prostate cancer. However, it was limited by its complicated detection process, low purity (< 0.5%) and sensitivity (high false positive and false negative rates)[50]. Therefore, it is urgently needed to develop novel approaches to improve the CTC detection technology, which demands better understanding of biological and physical properties of CTCs. Microfluidics has rapidly developed recently and has become a promising way to tackle the challenges, which has been overviewed in Section 1.5.

1.4.2 Downstream analysis of CTCs

After isolation, there need to be some analysis methods to identify the isolated cells and distinguish CTCs from background cells. Besides, apart from CTC counting for preliminary clinical decision, molecular profiling of clinically relevant CTCs subpopulations is important for the study of tumorigenesis, tumor progression, metastasis, and drug resistance. The recent technological for molecular characterization of CTCs are based on cellular morphology, cellular immunology and cellular molecular.

Compared to the normal blood cells, CTCs have abnormalities of structure,
function, and metabolism, which is the basics of CTC morphologic change[51]. CTCs have some common morphologic features, such as large, irregular, and hyperchromatic nuclei with prominent nucleoli. Besides, the cytoplasm of CTCs is changed due to secretion, condensation, and keratinization. Therefore, cell dyeing can be used to identify CTCs through cellular morphology.

CTCs are shed from solid tumor retaining the properties of epithelial cells and mesenchymal cells. Therefore, they can express gene of epithelial cells and mesenchymal cells, such as Cytokeratin (CK), Vimentin, E-cadherin, and Ncadherin. Besides, the specific tumor can express tissue specific gene, for example, human mammaglobin (hMAM) for breast cancer. CTCs can be identified by immunofluorescence (IF) staining or immunohistochemistry (IHC). The commonly used genes have been mentioned in Section 1.2.1. Fluorescence in situ hybridization (FISH) can also be used to analyze CTCs through fluorescence microscopy to localize fluorescent probes binding specific DNA sequence to parts of the chromosome[52]. In recent years, Surface-Enhanced Raman Scattering (SERS) spectroscopy has been developed in application of CTC analysis. It has been widely applied as optical imaging and detection tool due to the strong signal intensity, excellent photostability, biocompatibility, and especially the multiplexing ability[53]. Through appropriate analysis methods, CTCs can be detected more effectively.

1.5 Microfluidics for CTC detection

Microfluidics is characterized by dramatic reduction in sample and reagent consumption, low fabrication cost, simultaneous analysis of hundreds of samples in short time, and high integration of multiplexing functions such as sample pretreatment, transportation, mixing, reaction, separation, and detection on the same microdevice. It can realize the whole process of sample pretreatment and analysis online[54]. Besides, microfluidic platform featured with chambers, channels, and other nanostructures, such as pillars, tube, rod, and wires, can be used for cell separation because of their biomimetic surface conditions. Because these characteristics are suitable to CTC isolation and analysis as well, many microfluidic devices have been developed for CTC isolation[55]. The microfluidic methods have also been classified into labeldependent methods and label-independent methods.

1.5.1 Label-dependent technologies

Label-dependent technologies, also called affinity-based technologies, depend highly on the expression difference of cell surface markers between CTCs and normal blood cells. Most label-dependent technologies are positive enrichment, that is, capturing CTCs by EpCAM. In addition to positive enrichment, negative CTCs enrichment uses leukocyte-specific surface markers, such as CD45 and CD66b, to deplete leukocytes. Figure 1.2 shows some typical labeldependent microfluidic methods for CTC isolation, and the performance is summarized in Appendix Table 1.

Immunocapture

Immunocapture is the most traditional method among label-dependent technologies for CTC isolation. The surface of microfluidic features or channels can be coated with ligands or capture molecules which can combine with the surface markers of CTCs, as shown in Figure 1.2A[56]. When blood flows through the chip, CTCs will be captured and stay in the channel while other blood cells are taken away. In immunocapture method, varieties of structures are designed to improve the performance of CTC isolation. Micropost array can create turbulent flow to increase the contact chance of CTCs with ligand on the channel. Nagrath et al. developed a microfluidic platform fabricated with EpCAM coated microposts, which has successfully isolate CTCs from the blood[57]. Chaotic mixing is also used to increase the contact area between CTCs and channel surface. For example, the structure of grooved herringbone (HB) pattern is commonly utilized[20]. Immunocapture methods also use nanomaterials and nanostructures with high surface area-tovolume ratio to increase surface coated with antibody, and thereby to improve the capture efficiency. Tseng et al. introduced a NanoVelcro microfluidic device employing silicone nanopillars[58], silicone nanowires[59], and nanofibers coated with antibodies[60].

Hydrogel-based capture

Hydrogel is a kind of polymer material with three-dimensional (3D) network structure formed by chemical or physical crosslinking. It can swell in water instead of dissolving. Compared with other materials, hydrogel has obvious advantages, such as high-water content, soft properties, high permeability, and good biocompatibility and so on, which make it widely used in CTC separation[61-63]. Using the intelligent response of hydrogel to light and heat, researchers fabricated a capture substrate for CTC capture and release. As shown in Figure 1.2B, a near infrared (NIR) light-responsive substrate was designed to capture CTCs by immunocapture and release CTCs by a kind of thermos-responsive hydrogel. This method allows single CTC capture and recovery of captured CTCs[64]. Luan et al. developed a hybrid photonic barcode functionalized with folic acid (FA) to capture and release of CTCs with methacrylated gelatin (GeIMA) gel combined with FA molecules. The soft property of GelMA gel allows better cell viability after CTC isolation[49].

Immunomagnetic capture

The immunomagnetic method use magnetic nanoparticles coated with antibodies, aptamers, or peptides, which can capture CTCs and isolate CTCs from blood cell by a magnetic field. The CellSearch system is based on immunomagnetic method. The method can be classified into positive isolation focusing on capture of CTCs and negative isolation focusing on capture of leukocytes or erythrocyte. In some positive isolation methods, the captured CTCs will be adsorbed in the microchannel, making it difficult for downstream analysis of CTCs. Therefore, Saliba et al. developed a microfluidic device with antibody functionalized superparamagnetic beads which can self-assemble with magnetic traps in the microchannel. The capture efficacy is higher than 94%[65]. However, the self-assembled magnetic columns used in these methods are often unstable due to the large hydrodynamic drag force, which results in low throughput. Figure 1.2C shows a magnetic force gradient-based microfluidic chip which can separate CTCs depending on their expression level of EpCAM. This device can isolate CTCs from 3 mL blood within 1 hour[66]. Among negative isolation methods, magnetic nanobeads coated with anti-CD45 antibodies have been commonly used for separating WBCs from CTCs, thereby enriching CTCs[67]. Although the recovery rate of this negative method is higher than that of positive method, the purity of isolated CTCs is relatively low; therefore, further isolation is required.

Immunofluorescence

Fluorescence in situ hybridization (FISH) is a method that uses specific DNA sequences tagged with fluorophores to detect chromosome or gene status in the nucleus[68]. Conventional FISH technique is time consuming, labor intensive, and expensive[69]. It usually takes 2 or 3 days to complete analysis and the reagent required for assay is expensive such as fluorescently labeled DNA

probes. Therefore, microfluidics-based FISH device has been developed to test CTCs, which can highly reduce time and cost compared to the conventional FISH. After CTCs isolated from blood by microfluidic chip, FISH is used to carry out single-cell molecular diagnosis[70]. Zhao et al. introduced an automatic CTC counting method based on microfluidics and line confocal microscopy. As shown in Figure 1.2D, blood sample was directly labeled with a variety of antibodies combined with different fluorophores and pumped into microchannels linked with line-confocal microscope. Based on the fluorescence signal, CTCs can be counted automatically. Besides, 1 mL blood sample can be analyzed within 30 min due to the high flow rate[71]. The label-dependent technologies can achieve high specific and purity due to specific expression of CTCs. However, label-dependent technologies face the challenges of high cost of antibodies, low flow velocity, and small sample volume.



Figure 1.2: Examples of label-dependent microfluidic methods for CTC isolation. (A) Immunocapture isolation method with aptamer grafted SiNW substrate[56]; (B) Hydrogel-based capture isolation method with NIR-responsive cell-imprinted gelatin[64]; (C) Magnetic force gradient based microfluidic chip[66]; (D) Immunofluorescence isolation method based on microfluidics and line confocal microscopy[71].

1.5.2 Label-independent technologies

Label-independent technologies isolate CTCs from peripheral blood based on the difference of physical properties between CTCs and blood cells. Labelindependent method has advantages of isolating CTCs from blood without any expression of tumor-specific surface markers and maintaining CTCs activity. Figure 1.3 demonstrates some of label-independent microfluidic methods for CTC isolation, and the performance is summarized in Appendix Table 2. Based on the physical differences of size, density, stiffness, morphology and electrical properties, CTCs can be isolated by microfiltering, hydrodynamics, DEP, and acoustophoresis methods.

Mechanical filtering

Mechanical filtering isolation methods use the difference of size and deformability between CTCs and blood cells to isolate CTCs from blood. Almost CTCs have larger cell size and higher stiffness than blood cells. The microfilter commonly consists of pores, cavities, or geometric obstructions fabricated with certain arrangement[72]. When blood flows through the microfilter, the smaller RBCs, WBCs, and platelets will pass the filter while the bigger CTCs are captured. However, CTCs captured by microfilter may be damaged due to high flow rate and filtration pressure. Harouaka et al. designed a new flexible micro spring array (FMSA) device to minimize the damage of cell and maintain cell activity, as well as to increase throughput[73]. The capture efficiency of this device can reach to 90%. However, mechanical filtering isolation method is limited by clogging and adsorption because of the accumulation of cells on the filter, which will cause higher fluid driving forces, damage the captured CTCs, and reduce the isolation performance.

Hydrodynamics method

The hydrodynamics method is mainly depending on the size difference and the different movement response to the hydrodynamic force. There are three main types: Deterministic lateral displacement (DLD), Dean Flow fractionation (DFF), and Microvortex.

DLD is based on the characteristic that CTCs are larger than normal blood cells and is designed with DLD array of which the critical radius is between CTCs and blood cells. CTCs, which are larger than critical radius, will move to one side after collision with arrays, while blood cells which are smaller than critical radius will keep the original path flowing through the array. Loutherback et al. designed a DLD array device that can separate CTCs with the flow rate of 10 mL/min and achieve a capture efficiency higher than 85% without influence on cell viability[74]. DLD method may also suffer from clogging when detecting clinical samples that may be sticky and contain lots of debris.

DFF also uses the size difference between CTCs and blood cells. When fluid flows in a spiral channel, the velocity distribution of fluid is parabolic, with the greatest velocity in the middle of the channel leading to the greatest centrifugal force, thus, flowing to the outer edge of the channel. The fluid near the channel wall has the smallest velocity and the smallest centrifugal force, so it will be pushed by middle fluid. To maintain equilibrium, the Dean flow is formed in the direction perpendicular to the flow of the fluid[75]. Affected by lift force and Dean frag force, lager cells will move to the inner wall and smaller cells will move to outer wall, thereby to separate CTCs from blood cells as shown in Figure 1.3A[76]. A label-free spiral microfluidic device was developed which can separate CTCs from blood with more than 85% recovery rate and 99.99% WBC depletion[77].

Microvortex is caused by inertial force with suddenly expansion-contraction construction. As shown in Figure 1.3B, when cells flow through the expansion-contraction construction, cells larger than critical size will be trapped into vortices. Based on this principle, Renier et al. used microfluidic vortex chip to isolate CTCs from patients with advanced prostate cancer and isolate CTCs with the purity of 1.74-37.59%[78]. However, some microfluidic vortex chip may cause bubbles inside channel. The microfluidic vortex chip can remove large number of WBCs even there is overlapped part between CTCs and normal blood cells[79].

Electrokinetics method

The DEP method utilizes the translational motion of neutral particles in an asymmetric electric field due to dielectric polarization. The DEP force depends on the size and the cell membrane capacitance and suspended media (dielectric constant and conductivity), electric field strength and frequency, and the viscosity of suspended media. The size and cell membrane capacitance of CTCs are different from blood cells. Therefore, compared with other conventional separation methods, DEP method has advantages of higher selectivity, easier control, and higher separation efficiency. Chiu et al. developed an optically DEP method combined with microscopic fluorescent imaging[80]. The purity of isolated CTCs can reach to 100%. Alazzam et al. described the design and microfabrication of CTC isolation microfluidic chip based on DEP. As shown in Figure 1.3C, the device consists of a group of planar interdigitated transducer electrodes. It can precisely detect and count CTCs in the blood[81]. However, DEP method is difficult in manipulation because it requires high voltage and complicate process of resuspending cellular components in an isotonic medium with low conductivity.

Acoustophoresis method

Acoustophoresis is a method used to separate cells based on cellular volume and density[82]. Because cells experience acoustic force with different magnitude depending on size, density, and deformation of cells. Acoustic force will translate cells to the zero periodic pressure variations or maxima pressure node to reach a balance point. Li et al. developed an acoustic-based microfluidic device using tilted-angle standing surface acoustic waves, which can successfully separate CTCs from blood cells with high throughput and recovery rate higher than 83%[83]. Augustsson et al. separated prostate cancer cell lines from WBCs with recovery rate ranging from 93.6 to 97.9% and purity ranging from 79.6 to 99.7% using acoustophoresis method[84]. This research also found that cell viability is almost well kept when using acoustophoresis method. Wu et al. reported a platform integrating acoustics and microfluidics to separate CTCs from blood with cell structure, biological and functional integrity remaining. As shown in Figure 1.3D, CTCs are separated in the PDMS-glass hybrid channel resonator under hydrodynamic and surface acoustic wave force[85]. However, acoustophoresis method may also cause clogging with large volume blood and high acoustic pressure which can lyse cells.

Label-independent methods have advantages of high throughput, low cost, further cell molecular characterization and cell culture. However, overlapping in CTC and WBC sizes reduces the specificity of CTC isolation. The structure of filtering microchip may cause pores clogging. To solve these limitations, multistep technologies have been considered to improve CTC isolation method.



Figure 1.3: Examples of label-independent microfluidic methods for CTC isolation. (A) DLD CTC isolation with spiral microfluidic chip[76]; (B) Microfluidic Vortex HT chip[78]; (C) DEP CTC isolation[81]; (D) Acoustic CTC isolation[85].

1.5.3 Multistep technologies for CTC isolation

Due to the heterogeneity of CTCs, the physical and biological properties of CTCs are variable. It is not enough only to use one property to separate CTCs from blood. Therefore, multistep technologies for CTC isolation have been developed to improve the CTC capture efficiency, targeting two or more properties of CTCs. Figure 1.4 shows some of multistep technologies for CTC isolation and the performance is summarized in Appendix Table 3. A CTC isolation device was developed based on filter microstructure and size-based hydrodynamic method[86]. As shown in Figure 1.4A, CTCs are pre-separated by hydrodynamic force and further isolated by oscillatory flow. Song et al.

combined immunocapture with DLD and enhanced capture efficiency more than 300% and achieved cell viability of 96%[87]. As shown in Figure 1.4B and D, DLD was used to separate CTCs from blood cells and biomarkers coated micropillars were used to capture CTCs. Multivalent aptamer having higher binding efficiency with antigen was used as shown in Figure 1.4D while EpCAM was used as shown in Figure 1.4B. Besides, CTC-iChip was developed using DLD, DFF, and immunomagnetic methods for isolating CTCs [80]. In this design, DLD was used to separate nucleated cells from blood and delete red blood cells, and then inertial force was used to position all the cells in a line and magnetophoresis was used to remove nontarget cells. This device can capture more than 97% CTCs from blood with speed of 8 mL per hour. Ahmed et al. introduced a new device, combining DLD and immunocapture methods, in which pillars were immune-coated[88]. It has over 92% capture efficiency and 82% purity and CTCs were detected from non-metastasis colorectal patients by this method. Jack et al. developed an ultra-specific microfluidic technology based on DFF and immunomagnetic to improve the purity and throughout of CTC isolation[89]. In this design, the blood sample was presorted by inertial force to reduce unwanted cells. CTCs were subsequently labeled by EpCAM-functionalized magnetic beads and then separated by magnetophoresis. Figure 1.4C shows a dual-immunopatterned microfluidic device with double layers coated by anti-EpCAM antibody and anti-63B6 antibody respectively[90]. This device overcame the limitation of low EpCAM expression level of CTCs and achieved 94.47% capture efficiency.

Although the combined methods have better performance for CTC isolation, they require complex design of microfluidic devices, the integration level between different parts of methods is low, the operation is more complicated, and fabrication cost is higher. These shortcomings limit downstream analysis of CTCs and the commercial development of CTC isolation by microfluidics. Therefore, it is necessary to develop a microfluidic device with simple operation, high integration and high performance for isolation and downstream analysis of CTCs.



Figure 1.4: Examples of multi-step methods for CTC isolation. (A) CTC isolation by microfluidic ratchets and hydrodynamic concentrator[86]. (B) Size-dictated immunocapture chip combined immunocapture and DLD methods[88]; (C) Immunocature method using two types of antibody, anti-EpCAM antibody, and anti-63B6 antibody[90]; (D) AP-Octopus-Chip based on the DLD and immunocapture method[87];

1.6 SERS detection method

Surface-enhanced Raman scattering (SERS) spectroscopy has been widely applied as optical imaging and detection tool due to the strong signal intensity, excellent photostability, biocompatibility, and especially the multiplexing ability[53]. The SERS phenomenon was first found by Fleischmann in 1974[91]. It was found that rough metal surface can greatly enhance SERS spectral with enhancement multiple of 10^3 - 10^4 , which is called SERS effect. The discovery of SERS promoted the development of Raman spectral. At present, the total SERS enhancement is considered as the product of the electromagnetic enhancement (EE) and chemical enhancement (CE) mechanisms[92]. The electromagnetic enhancement mechanism is generally received as the dominant contributor to SERS enhancement[93]. The electromagnetic enhancement results from the amplification of the light by excitation of localized surface plasmon resonances (LSPRs). This light concentration occurs preferentially in the gaps, crevices, or sharp features of plasmonic materials, which are traditionally noble and coinage metals with nanoscale features, for examples, silver (Ag), gold (Au), and copper (Cu). The electromagnetic enhancement for SERS is affected by the structure of the supporting plasmonic material, which can reach factors of 10^{10} - 10^{11} [94]. The chemical enhancement mechanism primarily involves charge transfer mechanisms, where the excitation wavelength is resonant with the metalmolecule charge transfer electronic state. Theoretically, the chemical enhancement factors can reach 10^3 . Some studies found that the magnitudes of enhancement through charge transfer transitions are highly molecule specific[95, 96].

1.6.1 Metal-based SERS detection method

Since the electromagnetic enhancement mechanism is the main contributor to

the Raman signal enhancement, noble metal nanoparticles with plasmonic features have extensively been exploited as SERS platforms mainly due to their simplicity of fabrication[97]. The classic SERS substrates of gold (Au), silver (Ag), or copper (Cu), were widely used in SERS enhancement. Au and Ag are the most widely used as SERS substrate because of their stable properties, while Cu is more reactive. Researchers are striving to optimize substrate structure and configuration to maximize enhancement factors by identify new plasmonic materials and vary shapes and structure of nanoparticles[98, 99]. Various methods have been developed for fabrication of different shapes including nanostars[100, 101], nanoflowers[102, 103], nanorods[104, 105], nanotriangles[106, 107], nanocubes[108, 109], and nanocages[110, 111]. Materials with higher roughness and sharpness could achieve better SERS enhancement. Besides, the dense and enhanced plasmonic hot spots in sharp edges and nanogaps drastically increase the surrounding electrical field, enabling even a single molecule level detection[112]. Based on this phenomenon, advanced SERS substrates have been developed by modifying Au and Ag nanoparticles with coatings resulting in structures such as Au particles-coated nanospheres and SiO₂-encapsulated Au particles[113]. In general, the SERS enhancement factor of noble-metal nanomaterials can reach $10^{12}[114].$

1.6.2 Semiconductor-based SERS detection method

Since 1950s, semiconductor have been widely applied in microelectronics. The application on SERS activity of semiconductor materials has been further developed, including ZnO nanocrystals[115], ZnO superstructures[116], TiO₂ nanospheres[117], Cu₂O superparticles[118], W₁₈O₄₉ nanowires[119], Rh₃S₆ microbowls[120], MoO₂ NPs[121], SnO₂ NPs[122], Nb₂O₅ NPs[123], Ta₂O₅ nanorods[124], black TiO₂ nanowires[125] and so on. The enhancement mechanism of the semiconductor SERS effect is mainly attributed to electromagnetic enhancement charge-transfer (EM)and (CT) enhancement[126]. Compared to metal nanomaterials, semiconductor materials have advantages of high spectral stability and reproducibility, strong antiinterference ability, and selective SERS enhancement of target molecules[127]. However, the enhancement factor of semiconductor substrates is relatively low (10^3-10^5) [128]. To overcome this bottleneck, new enhancement mechanism and novel nanomaterials have been developed, such as crystalline semiconductormolecule systems, amorphous semiconductor-molecule systems, and crystalamorphous core-shell semiconductor-molecule systems. Metal oxides are the most used in crystalline semiconductor-molecule systems. For example, Cu₂O superstructure has been developed for SERS enhancement based on the synergistic effect of chemical enhancement and electromagnetic enhancement, with 10^{-9} M and an EF of 10^{5} [118]. People found that amorphous structures could enhance semiconductor-molecule interactions and thereby facilitating interfacial charge-transfer processes, resulting in improved sensitivity of semiconductor-based SERS[129]. The amorphous 2D TiO₂ nanosheets were developed with an ultrahigh EF of 1.86×10^{6} [130]. By integrating the advantages of both crystalline and amorphous structures, Lin et al. designed a kind of TiO₂ NPs with crystal-amorphous core-shell structure which have remarkable SERS activity in visible and near-infrared regions with an EF of 4.3×10^{5} [131]. Table 1.2 summarized the previously published results of SERS-active semiconductor nanomaterials.

 Table 1.2: SERS performance of previously published SERS-active

 semiconductor nanomaterials

Semiconductor	Probe Molecule	Enhancement Factor	Limit Of Detection	Ref.
Cu ₂ o	R6G	8×10 ⁵	6×10 ⁻⁹ M	[118]
Superstructure				
Diamond	MB	3.2×10 ⁵	10-7	[132]
Zno Nanosheets	4-MBA	10 ³ -10 ⁴	1×10 ⁻⁶ M	[133]
Sio ₂ Spheres	CV	104	-	[134]
Ag ₂ s NPs	4-MPY	10 ² -10 ³	-	[135]
Tio ₂ Nanosheets	4-MBA	1.86×10^{6}	-	[130]
Tio ₂ NPs	4-NBT	4.3×10 ⁵	10 ⁻⁶ M	[131]
Tio ₂ Nanowire	R6G	1.2×10^{6}	1×10 ⁻⁷ M	[125]

1.6.3 Application of SERS detection method in biosensing

With the rapid development of novel SERS materials and continuous

improvement of enhancement mechanisms, SERS detection method has been widely applied in biosensing which can be used to detect small molecule, DNA/aptamer, protein/enzyme/peptide/antibody, cellular and in vivo systems.

SERS detection of small molecule

There are several recent research applying SERS for biologically relevant small molecule detection, including the detection of small molecules such as antioxidants (glutathione and glucose), and small molecule markers such as biowarfare agents (anthrax). Zhang et al. designed a silver film over nanosphere (AgFON) substrates for the study of bacillus subtilis spores[136]. As shown in Figure 1.5A, the AgFON substrate can achieve a LOD of $\sim 2.6 \times 10^3$ spores which is lower than the anthrax infectious dose of 10^4 spores, within 11 min procedure. Furthermore, they used atomic layer deposition (ALD) to deposit a sub-1-nm alumina layer on AgFON substrates, which can maintain and stabilize the SERS activity of the underlying silver while present the surface chemistry of the alumina overlayer [137]. In this optimized method, the LOD of bacillus spores reduced to $\sim 1.4 \times 10^3$ within 10s. Compared to original AgFON substrates, the ALD modified AgFON substrates have twice the sensitivity and 6 times shorter data collection time and 7 times temporal stability. SERS method can also be used to detect glutathione due to the C-S stretching band at 660 cm⁻¹ shift. Ozaki et al. used Ag to enhance the Raman spectral of glutathione and achieved a glutathione detection range of 100-800 nM with LOD of 50 Nm[138]. SERS method can also be used for the detection of many other small molecule, such as oxidized glutathione (GSSG)[139], nicotinic acid adenine dinucleotide phosphate (NAADP)[140], glucose[141], lactate[142], lipids[143] and so on.

SERS detection of DNA and aptamer

SERS detection method can also be employed in DNA and aptamer due to its low LODs and good reproducibility. The common SERS detection method of DNA is to functionalize Au or Ag noble metal NPs with reporter molecule and a single stranded piece of DNA. When the single stranded piece of DNA is hybridized with complementary strand of DNA, the SRES signal of the reporter molecule can be observed [144-146]. Natan et al. reviewed the studies of using multiple SERS tags for the labeling or detection of DNA associated with disease[147]. This review has discussed the different schemes of Au and Ag NPs used as SERS substrate for biomolecule measurement. For example, Barhoumi et al. reported a SERS method for single and double-stranded thiolated DNA oligomer detection by binding DNA oligomers to Au nanoshellbased SERS substrates, as showed in Figure 1.5B[148]. They found that the SERS spectral of DNA oligonucleotides are extremely similar and affected by the Stokes modes of adenine. Besides, they introduced a correlation function analysis which can be used to assess reproducibility and quantify the highly complex changes corresponding to modifications in molecular conformation of the adsorbate molecules.

SERS detection of protein

SERS detection is also feasible to detect more complicated and larger molecules, such as protein, enzyme, peptide, antibody[149]. Wei et al. used Au nanoshell substrates to study the SERS spectral of three cysteine-containing aromatic peptides, phenylalanine-cysteine, tyrosine-cysteine, and tryptophancysteine, with excellent reproducibility[150]. They also obtained the relative Raman and SERS cross-sections of the characteristic Stokes modes of the three aromatic amino acids, which can be used to reproduce the Raman spectral of penetratin. Apart from protein identification, the measurements of protein concentration by SERS method have also been studied. For example, Han et al. developed a novel SERS-based method to probe protein concentrations in a solution by collecting the SERS signal of Coomassie Brilliant Blue dye adsorbed non-specifically to silver colloids to monitor the total protein concentration[151]. In this study, the detection range of protein concentration is 10^{-5} - 10^{-9} g/ml with 200 times lower LOD of 1 ng/ml, as showed in Figure 1.5C. This study shows great potential of SERS detection in proteins.

Cellular and in vivo SERS detection

In recent years, applications of SERS detection in cells, tissue and in vivo detection have been widely developed. Stuart et al. first introduced the in vivo

application of SERS which is to measure the glucose concentration of the interstitial fluid. As showed in Figure 1.5D, They functionalized AgFON surfaces with a two-component self-assembled monolayer and subcutaneously implanted it in a rat so that the glucose concentration can be measured by spectroscopically addressing the sensor through an optical window[152]. SERS has also been developed for cancer marker detection in a live cell. Qian et al. reported in vivo tumor targeting and detection based on pegylated gold nanoparticles and SERS[153]. After injecting NPs into the tail vein, SERS spectral obtained from tumor by using targeted nanoparticles can be obviously distinguished from using non-targeted nanoparticles. With a 785 nm laser, SERS tags were over 200 times brighter than NIR-emitting quantum dots. Besides, they could measure SERS spectra at targeted tumor sites up to 2 cm below the skin.



Figure 1.5: Application of SERS detection in biosensing. (A) SERS detection of Bacillus subtilis spores on AgFON substrates; (B) SERS detection of single and double-stranded thiolated DNA oligomers; (C) Concentration-dependent SERS spectra of CBBG from 10⁻⁵-10⁻⁹ g/ml (0-5); (D) Schematic of in vivo glucose measurement by SERS.

1.7 Gold-iron oxide composite nanoparticles

Composite nanoparticles are composed of different functional components, which have attracted more and more interests of scientists. Composite nanoparticles have great potential applications in the areas of electronics, photonics, catalysis, biotechnology, and nanotechnology[154-157]. Abovementioned materials generally have core/shell structure or a binary nanostructure. The successful applications of composite nanoparticles were highly depending on nanostructure, components, stability and dispersion of materials under different conditions. Therefore, researchers were focusing on fabrication of different composite nanoparticles to develop advanced multifunctional materials.

Magnetic composite nanoparticles with specific structure were widely studied in the areas of magnetic fluids, catalysis, data storage, bio-separation, and environmental treatment. To achieve these applications, there were numbers of methods for synthesis of magnetic composites with different components being developed [158-160]. Iron oxide (Fe₃O₄ or γ -Fe₂O₃) nanomaterials have typical structure of cubic inverse spinel. Its inherent magnetic feature, nano size, and surface effect could profit to construct magnetic hybrid nanocomposites, which could highly respond to external magnetic field. They could be applied in the capture of targeted substrates, recyclable nanocatalysis, magnetic-photonic purposes, and magnetic resonance imaging[161-163]. Gold nanoparticles were generally used in optical imaging, hyperthermia, and the detection of DNA and proteins[164-167]. By adjusting proximities and shapes of AuNPs, the plasmonic resonance peak of Au could be shifted to the NIR region[168]. Besides, it has been studied that the gold shell layer could provide a strong plasmonic resonant optical response to the nanoparticle. Therefore, the magnetic composites of Fe_xO_y and AuNPs are promising to be with advantages and properties from both individual Fe_xO_y and AuNPs.

1.7.1 Structure and synthesis of gold-iron oxide composite nanoparticles

The structure of gold-iron oxide composite nanoparticles was generally divided into two types: monodispersed composites and aggregate composites. The monodispersed composites attracted more attentions from scientists because of their better reproducibility and reliable characterizations. In the type of monodispersed composites, there were core/satellite structures, core/shell structures, multi-layer composite structures, nano-dumbbells and nanoflowers as shown in Figure 1.6. The core/satellite structures were generally formed by a single core linked with numerous smaller nanoparticles through covalent bonds or supramolecular interaction. The most common example of core/satellite structure was Fe₃O₄@Au. The Fe₃O₄ cores (50-300 nm) were synthesized by solvothermal reactions and the AuNPs (2-20 nm) were synthesized by Au ion reduction reaction. As shown in Figure 2.6A, The Fe₃O₄ cores were covered by a SiO₂ shell (3-50 nm thickness) by sol-gel reaction and functionalized with ammonium groups so that Fe₃O₄@SiO₂ NPs could be positively charged. After that, the negatively charged AuNPs could be attached onto positively charged Fe₃O₄@ SiO₂ NPs by electrostatic interactions, forming Fe₃O₄@SiO₂@Au core/shell/satellite structure nanocomposites.

Gold-iron oxide composite nanoparticles could be functionalized with antibodies so that they could be used to target specific cancer cells. There were three common ways for functionalization: direct attachment or ligand place exchange of proteins or antibodies through Au-S interactions based on cysteine residual thiol groups[169-171]; coupling reactions by formation of amide of antibodies or respective proteins[172, 173]; supramolecular interactions of biotinylated antibodies with streptavidin-functionalized nanocomposites[174]. Compared to single nanoparticles, the composite nanoparticles have higher surface which could load more ligands and thus improve performance.



Figure 1.6: Schematic representation of gold-iron oxide composite nanoparticles. A) $Fe_xO_y@Au$ core/satellite structures[175], B) $Fe_xO_y@Au$ core/shell structures[176], C) multi-layer $Fe_xO_y@Au$ composites[175], D) Au-Fe_xO_y dumbbell nanocomposites[175], E) gradual change from Au-Fe_xO_y nano-dumbbells to nanoflowers[175].

1.7.2 Applications of gold-iron oxide composite nanoparticles in cell sorting and separation

Gold-iron oxide composite nanoparticles have a wide range of applications including magnetic resonance imaging[177], computed tomography[178],

fluorescent optical imaging[179, 180], magnetic-induced hyperthermia[181], photo-induced hyperthermia[174, 182], drug delivery[179], DNA sensor[183], immunosensor[184, 185], enzyme-based sensor[186], cell sorting[187], and catalysis[188-190]. Among these applications, Magnetic separation and enrichment has become the most commonly used methods for biomarkerpurification, separation and cell sorting. Nash et al. designed a thermally responsive copolymer for coating Fe₃O₄ nanoparticles and model biomarker modified Au nanoparticles [191]. The polymer directed the formation of the Au-Fe₃O₄ aggregates that could be separated efficiently with a magnet. This method could be used for pulling down the biomarker from the human plasma in rapid diagnostic strategies. Wang and Irudayaraj designed two hybrid Fe₃O₄-Aurod-Fe₃O₄ nanodumbbells and Fe₃O₄-Aurod necklace-like structures with tunable optical and magnetic properties [187]. The Fe_3O_4 -Aurod necklace's surface could be conjugated with antibodies relevant to multiple pathogens. This probe could be used for optical detection of specific pathogens. Moreover, due to the magnetization and NIR absorption possessed by this nanocomposite, the target bacteria (E. coli and S. typhimurium) could be separated by the antibody conjugated Fe₃O₄-Aurod necklace under magnetic field, and after absorbing sufficient energy from the NIR laser, the nanocomposite was phototoxic to the specific target pathogen in the mixture.

Liu et al. employed Fe₃O₄@Au core/shell nanocomposites for the application

of purifying CD4+ lymphocytes from the spleen of mice[192]. Streptavidin-FITC was conjugated onto the surface of Fe₃O₄@Au core/shell nanocomposites. For the strong noncovalent binding between streptavidin and biotin, these modified nanocomposites could bind to the biotinlabeled anti-CD4+ antibody, which could also specifically bind to CD4+ lymphocytes. It was noted that Fe₃O₄@Au core/shell nanocomposites could successfully pull down CD4+ T lymphocytes from the whole splenocytes by magnetic separation.

Yu et al. reported the synthesis of poly (diallyldimethylammonium chloride)coated Fe₃O₄ (PDDA-Fe₃O₄) nanoparticles by the coprecipitation method[193]. Negatively charged citrate-capped Au nanoparticles could be effectively selfassembled onto the surface of cationic PDDA-Fe₃O₄nanoparticles. The Au-Fe₃O₄-PDDA aggregated nanocomposites revealed the capability of selectively enriching cysteine-containing peptides (cytochrome c) through the formation of Au-S bonds. The noncysteine-containing peptides could not be extracted by the same nanocomposites.

In recent decades, aromatic boronic acid derivatives have been employed in the construction of receptors for saccharides. Boronic acids form covalent bonds with cis-diol structures, which are present in most sugar moieties, to generate five- or six-membered cyclic esters in nonaqueous or basic aqueous media.

1.8 Black titanium dioxide

TiO₂ is known as the quintessential prototype photocatalyst. However, an obvious drawback of the different pure TiO₂ phases is a wide bandgap of 3-3.4 eV that limits the absorption of solar energy to only a few percent in the UV region. Since photocatalytic absorption is localized at the surface, surface modification appears as the direct route for both engineering the band gap and enhancing the photoactivity[194, 195]. The ideal surface modification would alter the surface of single crystal TiO₂ such that it absorbs the entire spectrum of solar irradiation from the infrared to the UV. Intuitively, the most efficient light-harvesting photocatalyst appears black to naked eye. Black TiO₂ first reported by Chen et al. in 2011 is a nanomaterial that was created by surface modification and distinguished by its black color[196]. This type of black TiO₂ is produced by H₂ reduction of white crystalline TiO₂, exhibiting a narrow band gap and enhanced photoactivity attributed to surface disorder. It has been widely applied in biomedicine. Ren et al. first reported hydrogenated black TiO_2 (H-TiO₂) NPs with near infrared absorption explored as photothermal agent for cancer photothermal therapy to circumvent the obstacle of ultraviolet light excitation[197]. To overcome the low drug loading ability, limited tissue penetration of UV light, and heating effect of 980 nm NIR on normal tissue, Ren et al. designed novel mesoporous silica (mSiO₂) coated black TiO₂ coreshell nanocomposites are designed and constructed as doxorubicin carriers for 808 nm NIR triggered thermal imaging guided photothermal therapy combined chemotherapy of breast cancer. Properties of the nanocomposites such as micro-morphology, size, drug loading ability and release, targeting performance, and therapy efficiency in vitro and in vivo were evaluated[198]. As well as therapy applications, black TiO₂ was studied for tumor diagnosis. Lin et al. successfully designed crystal-amorphous core-shell-structured TiO₂ NPs, the black TiO₂ NPs exhibited remarkable SERS activity in visible and near-infrared regions, and the EF value can be up to 4.3×10^5 . The efficient interfacial PICT can be attributed to the strong synergistic effect of the novel crystal-amorphous core-shell structure of B-TiO₂ NPs. The high-efficiency exciton transition of crystal core provides sufficient photoinduced charges. Furthermore, the interfacial band bending at the crystal-amorphous heterojunction enables effective exciton separation and charge injection, resulting in the enrichment of photoinduced charges in the amorphous shell to facilitate the interfacial PICT between the substrate and target molecules. Low Fermi level and high electronic density of states (DOSs) of the amorphous structure enable strong vibronic coupling in amorphous shell-molecule system. These properties are confirmed by the density function theory (DFT) calculations and Kelvin probe force microscopy (KPFM). In addition, the small band gap, high binding energy, and abundant surface defect states of B-TiO₂ are leveraged to form a stable complex and boost PICT resonance in the substrate-molecule system. Significantly, the remarkable SERS sensitivity

endows B-TiO₂ NPs with the capability of cancer cell diagnosis.

1.9 Research gap

Based on the literature review, larger numbers of technologies have been developed for CTC detection. However, there are some limitations in current technology requiring for improvements. As mentioned above, microfluidic technology could isolate CTCs from blood cells depending on the difference between CTCs and blood cells. The performance of label-dependent methods highly depends on expression level of biomarkers on CTCs and interaction between CTCs and biomarker-coated surface. Besides, excessive binding of CTCs and specific bio-probe might change origin structure of cells which would affect downstream analysis of CTCs and increase detection cost. The label-independent method has the main advantage of isolating CTCs from blood without any expression of tumor-specific markers. However, it is limited by low purity and specificity because of overlap existed in some physical properties between CTCs and blood cells. After isolation, there are still tens of thousands of WBCs remaining, which would disturb the accuracy and sensitivity of CTC detection and require further identification of CTCs. The commonly used way to identify CTCs is fluorescent immunolabeling by characterizing the surface protein expression of CTCs. However, when using multiple fluorophore-conjugated antibodies, the broadness of fluorescence spectrum and the cross-talking of fluorescent tags often confound the results.

Besides, the background noise was magnified in a microfluidic system, which might cause misrecognition of CTCs. SERS method has been widely applied in single cell analysis. However, excessive or unstable noble metal-based SERS biological probes would cause agglomeration and thereby cause hotspot effect leading to false positive. Therefore, it is necessary to develop a novel downstream analysis strategy of CTCs combined with microfluidic method.

1.10 Outline of the thesis

Refer to the research gap, microfluidic method was selected to isolate CTCs and SERS method was used for downstream analysis of CTCs. Three strategies have been adopted for CTC detection in clinical blood samples. In the first strategy, gold-iron oxide composite nanoparticles were selected as SERS SPION@Au-MBA-rBSA-FA SERS biological probes were substrate. reproduced according to the previous work. In this thesis, SPION@Au-MBArBSA-FA SERS biological probes were applied into clinical blood sample detection. 32 blood samples from cancer patients and 3 blood samples from healthy people have been detected. In the second strategy, black B-TiO₂ NPs were used as SERS substrate due to its low cost, high spectral stability and reproducibility, strong anti-interference ability, and selective SERS enhancement of target molecules. The B-TiO₂-AR-PEG-FA biological probe consisted of four layers. The innermost layer was B-TiO2 nanoparticles with crystal core and amorphous shell structure. The second layer was alizarin red

(AR) molecule which was responsible for providing Raman spectral signal. The third layer was a thin NH₂-PEG2000-COOH layer which was used to improve the dispersion of biological probes and to provide binding sites of folic acid (FA) and thus increase grafting rate of FA. The outermost layer was FA molecule. FA was used to specifically recognize cancer cells by folate receptor (FR) on cell membrane. The experiment results showed that this B-TiO₂ based SRES biological probe has good specificity and accuracy with obvious Raman signal. It can distinguish folate receptor-expressing cancer cells (MCF-7) from low folate receptor-expressing cells (A549 and Raw264.7). In the third strategy, microfluidic method was added to isolate CTCs before SERS detection. In this strategy, the relatively low specificity and isolation purity of microfilter could be solved by integrating with high-sensitivity SERS spectra detection, while the microfilter could reduce interference of blood background during SERS detection. Besides, SERS-fluorescence dual-modal in situ imaging method proved that this strategy has high specificity of detection with detection limit of 2 cancer cells per milliliter in rabbit blood. Besides, the operation process was simple and high-speed, with detection time less than 1.5 hours. These results illustrates that both of microfluidic isolation and SERS detection could open new paths for liquid biopsy.

The thesis was divided into six chapter to show the details of research. The first chapter introduced cancer background and gave the literature review of CTC

detection. It has reviewed the importance of CTC detection for cancer, CTC properties and clinical applications of CTCs. CTC detection was divided into two parts: isolation and downstream analysis of CTCs. Microfluidic technologies for CTC isolation have been reviewed, which have been classified into label-dependent method, label-independent method, and integrated method. SERS detection method has been considered for downstream analysis of CTCs. Both metal-based and semiconductor-based SERS substrate has been reviewed. gold-iron oxide composite nanoparticles and black titanium have been introduced as SERS substrates. The second chapter described the methodology used in research: synthesis of SPION-PEI@AuNPs-based SERS biological probe and application on CTC detection, synthesis of B-TiO₂-based SERS biological probe and application on CTC detection, and combination method of microfilter and B-TiO₂-based SERS biological probe on CTC detection. The methodology also contained cell culture, treatment of blood samples and materials and equipment for research. The third to fifth chapters were the experimental results and discussion. The fourth chapter was the characterization and application of SPION@Au-MBA-rBSA-FA SERS biological probes. SPION-PEI@Au were successfully synthesized according to the previous work of the research group. 32 clinical blood samples of cancer tumor and 3 clinical blood samples of healthy people were detected by SPION@Au-MBA-rBSA-FA SERS biological probes. The fifth chapter showed the experiment results of B-TiO₂-based SERS biological probe. The
synthesized B-TiO₂ showed good SERS enhancement effect and the LOD of the AR molecule on B-TiO₂ can reach 5×10^{-8} M. The experiment results showed that this B-TiO₂ based SRES biological probe has good specificity and detection accuracy with obvious Raman signal. It can distinguish folate receptor-expressing cancer cells (MCF-7) from low folate receptor-expressing cells (A549 and Raw264.7). In the sixth chapter, microfluidic method was added to isolate CTCs before SERS detection. In this strategy, the relatively low specificity and isolation purity of microfilter has been solved by integrating with high-sensitivity SERS spectra detection, while the microfilter could reduce the interference of blood background to SERS detection. Besides, SERS-fluorescence dual-modal in situ imaging method proved that this strategy has high specificity of detection with detection limit of 2 cancer cells per milliliter in rabbit blood. Besides, the operation process was simple and high-speed, with detection time less than 1.5 hours. The sixth chapter contained conclusion of this research and perspective of microfluidic method and SERS detection method.

Chapter 2 Methodologies

2.1 Introduction

This chapter has described the methodology of three research. Materials and equipment used for each research have been introduced. The first research was about synthesis of SPION-PEI@AuNPs-based SERS biological probe and application on CTC detection. The specificity in cells and the sensitivity and separation performance in blood of this biological probe have been tested. The traditional way for CTC separation was the combination of density gradient centrifugation and magnetic separation. The second research was about synthesis of B-TiO₂-based SERS biological probe and application on CTC detection. In this section, the SERS enhancement effect of B-TiO₂ was studied. The stability, specificity, and sensitivity of B-TiO₂ based biological probe was tested in the mixed cell suspension. The third research was about the study of combination method of microfilter and B-TiO₂-based SERS biological probe on CTC detection. The performance of B-TiO₂ based biological probe has been tested in the second research. In this section, numeric simulation has been performed to compare the flow dynamics of WBC clearance when flowing through the microfilter with different flow rate at range of 0.1-0.9 mL/min using simplified geometry and boundary conditions by ANSYS Fluent software. The purpose of simulation was to find a proper flow rate to avoid plugging of microfilter and cell damaging. Furthermore, sensitivity, specificity, and capture assay of this method have been tested both in mixed cells and blood.

2.2 Synthesis of SPION-PEI@AuNPs-based SERS biological probe and application on CTC detection

2.2.1 Materials and equipment

Materials: Ethylene glycol ($C_2H_6O_2$), sodium acetate anhydrous ($C_2H_3NaO_2$), sodium borohydride (NaBH₄), folic acid (C₁₉H₁₉N₇O₆), albumin from bovine 1-ethyl-3-[3-(dimethyllamino)propyl] serum (BSA), carbodiimide hydrochloride (EDC·HCl) as well as N-hydroxysuccinimide (NHS) were ordered from Aladdin Reagent Co. Ltd., Shanghai, China. Iron chloride hexahydrate (FeCl₃·6H₂O) was purchased from Alfa Aesar. Tetrachloroauric acid tetrahydrate (HAuCl₄·4H₂O), trisodium citrate (III) dehydrate (C₆H₅Na₃O₇·2H₂O) and Hoechst were purchased from Sinopharm Chemical Reagent Co. Ltd., Shanghai, China. 4-mercaptobenzoic acid (MBA), polyethylenimine (Mw~25000), DMSO and MTT was ordered from Sigma-Aldrich. Lymphocyte isolation was ordered from Slolarbio Life Science Co. Ltd., Beijing, China. Fetal bovine serum (FBS), incomplete DMEM (high glucose) as well as trypsin11 EDTA were ordered from KeyGen BioTech. Anti-CD45 antibody [F10-89-4] (Alexa Fluor® 488) Alexa and Anti-Cytokeratin 8 antibody [EP1628Y] (Alexa Fluor® 647) were purchased from Abcam, Zhejiang, China.

Equipment: The photos of the equipment was shown in Figure 3.1. The

nanoparticles were characterized by transmission electron microscopy (TEM, JEOL. 2100, Tokyo, Japan). The Raman spectra were observed on a confocal microprobe Raman system (Renishaw inVia Reflex, Wolton-under-Edge, U.K.). The laser wavelength was fixed at 785 nm. The time of data acquisition was set to 3.0 s, and the laser power was 280 mW. The SERS spectra were observed from liquid samples with homogeneous SERS hotspot. The zeta distribution of the nanoparticles was measured at room temperature by dynamic light scattering (DLS) using a zeta particle size analyzer (Malvern, England).



Figure 2.1: The photos of the equipment A) Transmission electron microscopy (TEM, JEOL. 2100, Tokyo, Japan); B) Confocal microprobe Raman system (Renishaw inVia Reflex, Wolton-under-Edge, U.K.); C) Zeta particle size analyzer (Malvern, England).

2.2.2 Synthesis of SPION-PEI@AuNPs-based SERS biological probe

Preparation of SPION-PEI

First, 0.68g FeCl₃•6H₂O was weighed by electronic balance and dispersed into 20 mL Ethylene Glycol by ultrasonic vibration method. Second, 1.8g NaAc was added into the above mixed solution. The solution was mixed uniformly by magnetic stirring device. Then, 0.75 g PEI was added into the previous solution. After stirring and heating at 60 °C for 20 minutes, the mixed solution was transferred to reaction kettle and heated at 220 °C for 2 hours by baking oven. After reaction and cooling to room temperature, the reaction product was washed several times by deionized water and alcohol and dispersed in 100 mL deionized water. After centrifugation at 1000 rpm for 5 minutes, the supernatant was collected and reserved at 4 °C.

Preparation of AuNPs

50 mL of 1 mM HAuCl₄ aqueous solution was heated at 165 °C with magnetic stirring. After boiling, 2.256mL of 1.0% Na₃Ct aqueous solution was added into the boiling solution rapidly. After the color of solution changing to purplish red, the solution was further heated for 3 minutes and then cooling at room temperature. After cooling, the AuNPs dispersion was reserved at 4 °C.

Preparation of SPION-PEI@AuNPs

First, 900 µL of SPION-PEI was added dropwise into 3 mL AuNPs solution. Next, the mixed solution was transferred to shaking table and oscillated at 200 rpm for 20 minutes at room temperature. After that, the obtained product was washed repeatedly with deionized water by magnet separation. The final SPION-PEI@AuNPs were dispersed in 4 mL deionized water and kept at 4 °C fridge.

Preparation of rBSA-FA

First, 50 mg FA was weighed and dissolved in 50 mL PBS solution (10mM, pH 7.4). After complete dissolution, 40 mg EDC and 24 mg NHS were added into the primary solution stirring at room temperature for 8 hours. After that, the activated FA was achieved. Second, 0.378 g NaBH₄ and 0.4 g BSA were dissolved in 10 mL and 20 mL deionized water respectively. 260 µL NaBH₄ was added into 20 mL BSA solution and stirred for 1hour to reduce BSA. After that, 5 mL reduced BSA (rBSA) was added into 50 mL activated FA and stirred for 8 hours. The obtained solution was added into ultrafiltration centrifuge tube and centrifuged at 6000 rpm for 30 minutes. The obtained rBSA-FA was dispersed at deionized water and saved at -20°C fridge.

Preparation of SPION-PEI@AuNPs-4MBA-rBSA-FA

The synthesis process of SPION-PEI@AuNPs-4MBA-rBSA-FA was shown in Figure 2.2. 0.000616g MBA was weighed and dissolved in 2 mL ethanol. The above MBA solution was diluted to 2×10^{-5} M. 4 mL MBA solution was added dropwise into 4mL SPION-PEI@AuNPs solution. The primary solution was transferred to shaking table and shock at 200 rpm for 5 minutes at room temperature. After shaking, the product was magnetic washed by PBS solution several times and centrifugated at 1000 rpm for 5 minutes. After centrifugation, the supernatant was collected for further use.



Figure 2.2: Chemistry synthesis process of SERS SPION-PEI@AuNPs-4MBA-rBSA-FA biological probe[199].

2.2.3 Cell culture

MCF-7 is a human breast cancer cell line and A549 is a human non-small cell lung carcinoma cell lines, which are bought from Shanghai Institutes for Biological Sciences (SIBS). Hela is a human cervical cancer cell line supported by General Hospital of Eastern Theater Command. All the cells were cultured in complete culture medium (90% DMEM, 10% FBS) under standard cell culture condition (5% CO₂, 37 °C). For experiments, cells were detached by 0.25% trypsin solution and resuspended in PBS solution.

2.2.4 Specificity of SPION-PEI@AuNPs-based SERS biological probe in cells

To verify the SERS performance of SPION-PEI@AuNPs-based SERS biological probe on the cell membranes and the ability to specifically recognize the FR on the cancer cell membranes, two cancer cell lines of MCF-7 and A549 were selected as models and detected by Raman instrument. First, MCF- 7 and A549 were separately incubated with PBS solution, SPION-PEI@AuNPs-MBA-rBSA-FA biological probe and PEI@AuNPs-MBA biological probe for 1 hour and washed by PBS for three times. After that, the Raman instrument was used to observe the optical image and corresponding SERS mapping image of cells, at 785 nm laser.

2.2.5 Sensitivity and separation of SPION-PEI@AuNPs-Based SERS biological probe in blood

Animal experiments in this study were all carried out according to the protocols approved by the animal care and use committee of the Ningbo University. The blood samples were drawn from rabbit ears and then injected into vacutainer tubes, which contained lithium heparin. MCF-7 cell line was spiked into rabbit blood at 0-500 cells/mL concentration. The blood samples were then separated by density gradient centrifugation. 2 mL blood sample was first diluted by adding 2 mL PBS solution. Next, the blood solution was slowly added to 2 mL lymphocyte separation solution. After centrifugation at 1500 rpm for 25 minutes, the sample was separated into four layers (from top to bottom: plasma, mononuclear cells, lymphocyte separation solution, red blood cells). The layer of mononuclear cells was collected and washed twice with PBS solution by centrifugation at 1000 rpm for 5minutes. After washing, 200 µL SERS biological probes were added into collected cells and incubated for 1 hour. After incubation, the excess biological probes were cleaned away by

centrifugation at 1000 rpm for 5 minutes. Then, the samples were rinsed thrice with 200 μ L PBS solution via magnetic separation and measured by confocal Raman microscopy.

2.2.6 Application in clinical tumor blood detection

To explore the clinical utility of SERS biological probe, blood samples from 32 cancer patients and 3 healthy people were detected. The blood samples were then separated by density gradient centrifugation. The pretreatment of blood was same as Section 2.2.5. 2 mL blood sample was first diluted by adding 2 mL PBS solution. After centrifugation, CTCs were collected by SERS biological probe and separated by magnetic separation. Then, the samples were rinsed thrice with 200 μ L PBS solution and measured by confocal Raman microscopy at 785 nm laser.

2.3 Synthesis of B-TiO₂-based SERS biological probe and application on CTC detection

2.3.1 Materials and equipment

Materials: Titanium(IV) oxide (P25) was purchased from Acros Organics; Sodium borohydride (NaBH₄), Alizarin red, NH₂-PEG2000-COOH (95%), Folic acid (≥98%), N-hydroxysuccinimide (NHS), 1-ethyl-3-(3-(dimethylamino)propyl) carbodiimide (EDC), Fluorescein isothiocyanate (FITC), FITC-PEG2000-FA were purchased from Aladdin, Shanghai, China;

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Tris-HCI buffer (1.0 M, pH 8.5) was purchased from Macklin, Shanghai, China; Dialysis bag (50kD) was purchased from Yuanye, Shanghai, China; FITC linked polyclonal rabbit antibody to FA was purchased from Cloud Clone Corp, USA; Alexa Fluor® 647 Anti-Cytokeratin 8 antibody (CK8) and Alexa Fluor® 488 Anti-CD45 antibody (CD45) were purchased from Abcam, Shanghai, China; Hoechst was purchased from Sinopharm Chemical Reagent Co., Ltd.

Equipment: The photos of the equipment were shown in Figure 3.3. The transmission electron microscopy (TEM) and high-resolution transmission electron microscopy (HRTEM) images of the B-TiO₂ samples were obtained by Talos F200x. The powder X-ray diffraction (XRD) of the B-TiO₂ samples were characterized by the BRUKER D8 ADVANCE DAVINCI diffractometer with Cu K α radiation ($\lambda = 1.54056$ Å). Raman spectra was collected by Renishaw inVia Reflex instrument with 532 nm laser. The fluorescence images were obtained by laser confocal fluorescence microscopy (LEICA, TCS SP5).



Figure 2.3: The photos of the equipment A) Transmission electron microscopy (TEM, JEOL. 2100, Tokyo, Japan); B) High-resolution transmission electron microscopy (HRTEM, Talos F200x); C) X-ray diffraction (XRD, D8 ADVANCE); D) Confocal microprobe Raman system (Renishaw inVia Reflex, Wolton-under-Edge, U.K.); E) Biological laser confocal fluorescence microscopy (LEICA, TCS SP5).

2.3.2 Synthesis of B-TiO₂-based SERS biological probe

Fabrication of B-TiO₂ NPs

B-TiO₂ was fabricated according to the literature [200]. At room temperature with dry condition, the mixture of 1.5 g P25 and 1.5 g NaBH₄ was ground for 30 minutes. Then the mixture was transferred to porcelain boat and placed into tube furnace. The heating process started from room temperature to 350 °C at a heating rate of 10 °C per minute and kept at 350 °C for 4 hours with protection of argon. After natural cooling, the obtained product was washed by water and

ethanol three times and dried at 70 °C. Finally, B-TiO₂ NPs was obtained.

Synthesis of SERS biological probe

The SERS biological probe consisted of four parts as shown in Figure 3.4, a crystal-amorphous core-shell B-TiO₂ as a core structure, Raman reporter AR molecule absorbed on B-TiO₂, a thin NH₂-PEG2000-COOH layer to protect B-TiO₂-AR and to improve the dispersion of biological probe, and FA to specifically recognize folate receptor on cancer cell membrane. First, 5 mL AR solution (10⁻²M) was mixed with 5 mL 1 mg/mL B-TiO₂ by stirring for 4 hours and centrifugating 3 times at 10000 rpm. AR molecule was absorbed onto B-TiO₂ by chemical bonding[201].



Figure 2.4: Chemistry synthesis process of B-TiO₂ based biological probe

The obtained B-TiO₂-AR was dispersed in 50 mL 10 mM Tris-HCI buffer (pH 8.5). Then, B-TiO₂ -AR-PEG was obtained by adding 10 mg NH₂-PEG2000-COOH into the above solution, stirring for 4 hours and centrifugated 3 times at 10000 rpm.

FA was first activated by NHS and EDC in phosphate buffer saline (PBS, pH 7.4). First, 50 mg FA was fully dissolved in 50 mL PBS. Then, 40 mg EDC and 24 mg NHS were added into FA solution stirring for 4 hours at room temperature. After that, the activated FA was mixed with B-TiO₂-AR-PEG solution stirring for 12 hours at room temperature. B-TiO₂-AR-PEG was modified with FA by the amide interaction between the carboxyl group of FA and amino group of NH₂-PEG2000-COOH. Finally, the B-TiO₂-AR-PEG-FA was collected and washed by dialysis bag (50 kD) for 36 hours to remove the excess FA. The obtained B-TiO₂-AR-PEG-FA biological probe was dispersed into PBS solution and saved in 4 °C fridge.

2.3.3 Cell culture

MCF-7 is a human breast cancer cell line. A549 is a human non-small cell lung carcinoma cell line. Raw264.7 is a mouse monocyte macrophage leukemia cell line. These three cell lines are bought from Shanghai Institutes for Biological Sciences (SIBS). All the cells were cultured in complete culture medium (90% DMEM, 10% FBS) under standard cell culture condition (5% CO₂, 37 °C). For experiments, cells were detached by 0.25% trypsin solution and resuspended in PBS solution.

2.3.4 FR expression screening of MCF-7, A549 and Raw264.7 cell lines

The three cell lines were incubated in a 6-well plate with same cell concentration. Each cell line was divided into two wells. The same amount of FITC-PEG2000-FA was added into one well of each cell line respectively and incubated for 30 mins. Then the cells were washed by PBS solution and analyzed by flow cytometry.

2.3.5 FA modification of SERS biological probe

To verify the FA modification, two methods were utilized. For the first one, the conjugation efficiency of FA on the SERS biological probe was calculated by a previous mass balance method[202]. After the activated FA mixed with B-TiO₂-AR-PEG stirring for 12 hours, the resulting B-TiO₂-AR-PEG-FA was obtained by centrifugation at 10000 rpm for 10 minutes. The supernatant was collected, and the residual FA content was determined using the calibration curve of FA standard solutions by measuring the UV-vis absorption. For the second one, a FITC linked polyclonal rabbit antibody to FA was used to simulate the FA receptors on the cell membranes. It was incubated with FA modified SERS biological probe and non-FA modified SERS biological probe for 3 hours, separately. Then, after washing by PBS buffer for three times, the FA modified SERS biological probe and non-FA modified SERS biological probe were observed by fluorescence microscopy gel image system.

2.3.6 Specificity and sensitivity of B-TiO₂-based SERS biological probe in

the mixed cells

To verify the SERS performance of B-TiO₂ SERS biological probe on the cell membranes and the ability to specifically recognize the FR on the cancer cell membranes, two cancer cell lines of MCF-7 and A549 were selected as models and detected by Raman instrument. First, MCF-7 and A549 were separately incubated with PBS solution, B-TiO₂-AR-PEG-FA b biological probe and B-TiO₂-AR-PEG biological probe for 1 hour and washed by PBS for three times. After that, the Raman instrument was used to observe the optical image and corresponding SERS mapping image of cells. Second, after incubated with B-TiO₂-AR-PEG-FA biological probe, MCF-7 and A549 were stained by FITC and Hoechst and observed by confocal fluorescence microscopy. Besides, AR can produce its characteristic fluorescence with its excitation and emission wavelength at 530 nm and 560 nm, respectively.

To verify the ability of B-TiO₂-AR-PEG-FA biological probe to remove interference from FR negative cells, two cell lines of MCF-7 and Raw264.7 were mixed and incubated with B-TiO₂-AR-PEG-FA biological probe for 1 hour and washed by PBS for three times. After that, the mixed cells were stained with Hoechst (blue, MCF-7 positive, Raw264.7 positive), CK8 (red, MCF-7 positive, Raw264.7 negative) and observed by confocal fluorescence microscopy. To test the sensitivity of biological probe, 200 μ L B-TiO₂-AR-PEG-FA biological probe was mixed with 1.0 mL cell suspension of MCF-7 with concentration of 1-100 cells per milliliter. After incubation for 1 hour, the excess biological probes were cleaned away by centrifugation at 1000 rpm for 5 minutes. The precipitate was collected and dispersed evenly in 200 μ L PBS solution for the measurement of confocal Raman microscopy.

2.4 Combination method of microfilter and B-TiO₂-based SERS biological probe on CTC detection

2.4.1 Materials and equipment

Materials: Titanium(IV) oxide (P25) was purchased from Acros Organics; Sodium borohydride (NaBH₄), Alizarin red, NH₂-PEG2000-COOH (95%), Folic acid (≥98%), N-hydroxysuccinimide (NHS), 1-ethyl-3-(3-(dimethylamino)propyl) carbodiimide (EDC), Fluorescein isothiocyanate (FITC), FITC-PEG2000-FA were purchased from Aladdin, Shanghai, China; Tris-HCI buffer (1.0 M, pH 8.5) was purchased from Macklin, Shanghai, China; Dialysis bag (50kD) was purchased from Yuanye, Shanghai, China; FITC linked polyclonal rabbit antibody to FA was purchased from Cloud Clone Corp, USA; Alexa Fluor® 647 Anti-Cytokeratin 8 antibody (CK8) and Alexa Fluor® 488 Anti-CD45 antibody (CD45) were purchased from Abcam, Shanghai, China; Hoechst was purchased from Sinopharm Chemical Reagent Co., Ltd.; Microfilter, PL01 was purchased from Anfang Biotechnology, Guangzhou,

China.

Equipment: The photos of equipment were shown in Figure 3.3. The transmission electron microscopy (TEM) and high-resolution transmission electron microscopy (HRTEM) images of the B-TiO₂ samples were obtained by Talos F200x. The powder X-ray diffraction (XRD) of the B-TiO₂ samples were characterized by the BRUKER D8 ADVANCE DAVINCI diffractometer with Cu K α radiation ($\lambda = 1.54056$ Å). Raman spectra was collected by Renishaw inVia Reflex instrument with 532 nm laser. The fluorescence images were obtained by laser confocal fluorescence microscopy (LEICA, TCS SP5).

2.4.2 Cell culture

MCF-7 is a human breast cancer cell line. A549 is a human non-small cell lung carcinoma cell line. Raw264.7 is a mouse monocyte macrophage leukemia cell line. These three cell lines are bought from Shanghai Institutes for Biological Sciences (SIBS). All the cells were cultured in complete culture medium (90% DMEM, 10% FBS) under standard cell culture condition (5% CO₂, 37 °C). For experiments, cells were detached by 0.25% trypsin solution and resuspended in PBS solution.

2.4.3 Simulation of optimum capture flow rate in microfilter

Numeric simulation was performed to compare the flow dynamics of WBC

clearance when flowing through the microfilter with different flow rate at range of 0.1–0.9 mL/min using simplified geometry and boundary conditions by ANSYS Fluent software. The purpose of simulation was to find a proper flow rate to avoid plugging of microfilter and cell damaging.

Governing equation

First, a 2D numerical model was developed to investigate the WBC clearance when flowing through a hole with depth of 30 μ m, top diameter of 6.5 μ m and bottom diameter of 16 μ m in the microfilter. The photo of microfilter was shown in Figure 2.5. Only one hole was considered for the sake of saving computational cost.



Figure 2.5: Optical image of microfilter.

The volume of fluid (VOF) model in Ansys Fluent 15.0 was used to study the effects of different blood flow rates on the shape and structure of WBCs. In the VOF model, the volume fraction α can be obtained from the continuity equations (1) and (2)

$$\frac{\partial \alpha_d}{\partial t} + \vec{u} \cdot \nabla \alpha_d = 0 \qquad (1)$$
$$\frac{\partial \alpha_c}{\partial t} + \vec{u} \cdot \nabla \alpha_c = 0 \qquad (2)$$

where the subscript c represents the continuous phase (blood), and the subscript d represents the dispersed phase (WBCs). The sum of the volume fractions of the two-phase fluid in each computational cell is 1, i.e., $\alpha_d + \alpha_c = 1$, *t* is the flow time, \vec{u} is the flow velocity which can be obtained from continuity equation and Navier-Stokes equation

$$\nabla \cdot \vec{u} = 0 \qquad (3)$$

$$\frac{\partial \rho \vec{u}}{\partial t} + \nabla (\rho \vec{u} \cdot \vec{u}) = -\nabla p + \nabla \cdot [\mu (\nabla \vec{u} + \nabla \vec{u})] + \rho \vec{g} + \vec{F}s \qquad (4)$$

where p and \overrightarrow{F}_s are the pressure, and surface tension, respectively. In each computational cell, the density ρ and dynamic viscosity μ are calculated by Eq. (5) and (6)

$$\rho = \alpha_c \rho_c + \alpha_d \rho_d \qquad (5)$$
$$\mu = \alpha_c \mu_c + \alpha_d \mu_d \qquad (6)$$

The surface tension $\vec{F}s$ is obtained by the continuous phase surface tension model, as shown in Eq. (7)

$$\vec{F}_{s} = \frac{\rho}{\frac{1}{2}(\rho_{c} + \rho_{d})} \sigma \kappa \nabla \alpha \qquad (7)$$

where σ is the surface tension coefficient, κ is the interface curvature and $\kappa = \nabla \cdot \hat{n}$, \hat{n} is the surface unit normal calculated by $\hat{n} = \vec{n} / |\vec{n}|$, and $\vec{n} = \nabla \alpha$. Wall adhesion is considered by defining a contact angle θ_w at the channel wall, and the surface normal at the reference cell next to the wall is calculated by equation (8)

$$\hat{n} = \hat{n}_{w} \cos \theta_{w} + \hat{t}_{w} \sin \theta_{w} \tag{8}$$

Where \hat{n}_{w} and \hat{t}_{w} are the unit vectors normal and tangential to the wall, respectively. The geometric parameters and material properties in the VOF simulation are shown in Table 2.1 and Table 2.2, respectively. The PRESTO! Interpolation scheme was employed to compute the pressure equation. The convective and diffusion terms were discretized using second-order upwind and central difference schemes, respectively. The momentum equation adopted the second-order upwind scheme, and the volume fraction was solved by Geo Reconstruct method.

Table 2.1: Geometry of microchannel

Upper	Lower	Depth of	Distance from inlet	Distance from outlet
aperture	aperture	hole	to upper pore	to lower pore
6.5 µm	16 µm	30 µm	30 µm	20 µm

Table 2.2: Physical parameters of leukocytes and blood [203]

	Density (kg·m ⁻³)	Viscosity	Interfacial tension (N·m ⁻¹)	Contact angle (°)
		(pa·s)		
Blood	1055.5	0.03	0.027	135
WBCs	1080.0	13		

In the previous simulation, the WBCs are considered as liquid, however the real WBCs are surrounded by a thin membrane. To capture the interaction

physics during the contact between the cell membrane and the solid boundary of the hole in the microfilter, i.e., the wall of the micropore as shown in Figure 2.6, the simulation was also conducted with the effect of carrier flow simplified as an axial pressure at the top. Since the inner structure of the cell was neglected, a homogeneous pressure was applied on the inner boundary of the cell. The solid mechanics calculation was performed for the thin membrane of the cell with a thickness of 0.5 μ m, which has contact with the wall of the micropore. This simulation was conducted using a 3D model by solid mechanics governed by,

$$\nabla \cdot (FS)^T + F_V = 0 \tag{9}$$

Where $F = I + \nabla u$. S is the 2nd Piola-Kirchhoff stress; thus, FS is the 1st Piola-Kirchhoff stress; F_V represents the term of volume force. The contact pressure from the source of contact pair is,

$$T_n = if \ (g_n \le 0, \ -p_n g_n, 0) \tag{10}$$

where $p_n = \frac{E_{char}}{h_{min}}$ is the contact pressure penalty factor, E_{char} is the characteristic stiffness, g_n is the gap distance. The tangential force T_t is computed as

$$T_t = \min\left(\frac{T_{t,crit}}{|T_{t,trial}|}, 1\right) T_{t,trial} \qquad (11)$$

where $T_{t,trial} = -p_t s$, $T_{t,crit} = min(\mu T_n + T_{cohe}, T_{t,max})$ is the critical value of tangential force, T_{cohe} is the cohesion force; p_t is the penalty factor identified as the spring constant, μ is the friction coefficient defined as

$$\mu = \begin{cases} \mu_d + (\mu_s - \mu_d)exp(-\alpha_{def}|v_s|) \\ \mu_s \end{cases}$$
(12)

where μ_d is the dynamic friction coefficient, μ_s is the static friction coefficient, v_s is the slip velocity, and α_{def} is a decay coefficient. The upper equation is the equation of dynamic friction, while the lower represents otherwise.



Figure 2.6: The schematic of the WBC flowing through a hole of the microfilter.

2.4.4 Sensitivity, specificity, and capture assay of microfilter combined SERS spectral detection method in mixed cells

To test the sensitivity and capture assay of this method in mixed cells, the cell suspension solution of MCF-7, A549, raw264.7 and MCF-7 mixed with raw264.7, were added into microfilter device at 1-100 cells/mL concentration. The concentration of cell suspension solution before and after microfiltration was counted by cell counter. The isolation experiment was performed at a

constant pulling speed (0.4 mL/min) controlled by a micro syringe pump. After that, the B-TiO₂ SERS biological probe was added into microfluidic device and incubated with isolated cells for 1 hour. Then, the microfilter was washed carefully by PBS solution. After that, the microfilter was illuminated by 532 nm laser of Raman instrument. The SRES detection results were verified by fluorescence microscope.

2.4.5 Sensitivity, specificity, and capture assay of microfilter combined SERS spectral detection method in blood

To test the sensitivity and capture assay of this method in the blood, MCF-7 cell line was spiked into rabbit blood at 1-100 cells/mL concentration. The MCF-7 spiked blood was diluted by PBS solution with volume ratio of 1:2 and added into microfilter device. The concentration of cell suspension before and after microfiltration was counted by cell counter. The isolation experiment was performed at a constant pulling speed (0.4 mL/min) controlled by a micro syringe pump. After that, the B-TiO₂ SERS biological probe was added into microfilter was washed carefully by PBS solution. After that, the microfilter was illuminated by 532 nm laser of Raman instrument. The SRES detection results were verified by fluorescence microscope.

2.4.6 Application in clinical tumor blood detection

To explore the clinical utility of SERS biological probe, blood samples from 6 breast cancer patients and 2 healthy donors were detected. The blood sample was diluted by PBS solution with volume ratio of 1:2 before microfiltration. The diluted blood sample was added into microfluidic device. The isolation experiment was performed at a constant pulling speed (0.4 mL/min) controlled with a micro syringe pump. After that, the B-TiO₂ SERS biological probe was added into microfluidic device and incubated with isolated cells for 1 hour. After washed by PBS solution, the microfilter was illuminated by 532 nm laser of Raman instrument. The SERS spectral was gathered to analyze the detection result.

2.5 Conclusion

This chapter has introduced methodology used for research. MCF-7, A549 and Raw264.7 cell lines were used for experiment. The performance of both SPION-PEI@Au based SERS biological probe and B-TiO₂ based SERS biological probe have been tested. 32 blood samples from cancer patients and 3 blood samples from healthy people were detected by PION-PEI@Au based SERS biological probe. The sensitivity, specificity, and capture assay of microfilter combined SERS spectral detection method in mixed cells and blood have also been studied respectively. The combined method has also been applied in clinical blood sample detection. The corresponding experimental results would be shown in Chapter 3 to 5.

Chapter 3 Synthesis of SPION-PEI@AuNPs-based SERS biological probe and application on CTC detection

3.1 Introduction

Circulating tumor cells (CTCs) are tumor cells shedding from the primary or metastatic tumor and entering the peripheral blood circulation system. CTCs have incubation period in the blood. Some of CTCs would translate to other tissue and form new tumors[204, 205]. Therefore, CTCs detection have important significance for early diagnosis, prognosis evaluation, therapeutic efficacy, and method selection of cancer treatment. However, the number of CTCs in the blood is extremely small, with only 1-10 CTCs per milliliter, among 1 million white blood cells (WBCs) and 1 billion red blood cells (RBCs)[48]. Because of small amounts of CTCs, there is no effective means to detect CTCs at present. Therefore, developing an ultra-sensitive method for CTC detection is quite necessary.

SERS method is an ultra-sensitive method which can be used for molecular level detection. There are many studies using SERS method to detect CTCs in peripheral blood. In previous work, an improved SERS-active magnetic nanoparticles for CTC detection and analysis has been designed [199]. As shown in Figure 3.1, superparamagnetic iron oxide nanoparticles (SPION) with poly(ethyleneimine) (PEI) on the surface (SPION-PEI), was assembled with AuNPs and then linked with Raman reporter molecule MBA and rBSA-FA sequentially. CTCs could be enriched by SERS biological probe using a magnet and detected by Raman instrument. The SPION-PEI@AuNPs-MBA-rBSA-FA SERS biological probes realized the integration of CTC separation and detection.

This chapter focuses on the clinical application of SPION-PEI@AuNPs-MBArBSA-FA SERS biological probes. The specificity and separation performance of SPION-PEI@AuNPs SERS biological probe has been replicated. After that, the SPION-PEI@AuNPs SERS biological probe has been applied to detect several blood samples from cancer patients. The successful CTC detection in clinical cancer patient bloods demonstrated that SERS detection was worth to be further developed and a promising tool for diagnosis and treatment of cancer.



Figure 3.1: Design of magnetic SERS biological probe and separation mechanism. a) Design of the magnetic SERS biological probe: SPION-PEI@AuNPs-MBA-rBSA-FA; b) Mechanism of the SERS nanoagent for CTC analysis.

3.2 Characterization of SPION-PEI@AuNPs-based SERS biological probe

The SPION-PEI@AuNPs SERS biological probe was composed according to synthetic steps from Figure 3.1a. The SPION-PEI was prepared by solvothermal synthesis method and AuNPs was prepared by sodium citrate reduction of chloroauric acids. Figure 3.2 shows the morphological features of SPION-PEI, AuNPs, SPION-PEI@AuNPs and SPION-PEI@AuNPs-MBA-rBSA-FA. The TEM images showed that the size of AuNPs was 30-40 nm, and the size of SPION-PEI was 200-300 nm (Figure 3.2a and 3.2b). Figure 3.2c showed that SPION-PEI@AuNPs has Core-satellite structures with size of 250-350 nm. After functionalized by rBSA-FA, there was a layer of film on the

nanoparticles as shown in Figure 3.2d. The Zeta potential of AuNPs and SPION-PEI were -47.3 mV and +27.7 mV, respectively (Figure 3.3). AuNPs with negative charge were self-assembled on the SPION-PEI by electrostatic attraction. After electrostatic attraction, the Zeta potential of SPION-PEI@AuNPs was changed to -33.1 mV. The strong repulsive force made SPION-PEI@AuNPs of good dispersibility in water. The SERS spectral of SPION-PEI@AuNPs, SPION-PEI@AuNPs-MBA and SPION-PEI@AuNPs-MBA-rBSA-FA were shown in Figure 3.4. Because the layer of rBSA was relatively thin, the SERS intensity of SPION-PEI@AuNPs-MBA-rBSA-FA showed almost no weakening after covered by rBSA-FA and has good stability. The depth scanning results of SPION-PEI@AuNPs-MBA-rBSA-FA showed that this SERS biological probe has good stability and uniformity in the solution.



Figure 3.2: TEM images of a) AuNPs, b) SPION-PEI, c) SPION-PEI@AuNPs, d) the layer of rBSA-FA.



Figure 3.3: Zeta potential distribution of AuNPs, SPION-PEI, SPION-PEI@AuNPs, and SPION-PEI@AuNPs-MBA-rBSA-FA.



Figure 3.4: a) SERS spectral of SPION-PEI, SPION-PEI@AuNPs-MBA and SPION-PEI@AuNPs-MBA-rBSA-FA; b) SERS spectral of SPION-PEI@AuNPs-MBA-rBSA-FA from randomized 5 points by depth scanning, depth (1-5) -100, -200, -300, -400, -500 μ m. Laser wavelength: 785 nm; lens: 50 × objective.

3.3 Specificity of SPION-PEI@AuNPs-based SERS biological probe in cells

To test the specificity of SPION-PEI@AuNPs-based SERS biological probe, MCF-7 cells and A549 cells were incubated with PBS solution, SPION-PEI@AuNPs-MBA and SPION-PEI@AuNPs-MBA-rBSA-FA, respectively. Figure 3.5 indicated that after functionalized by rBSA-FA, SERS biological probes could target FR positive cells. Because A549 cells were FR negative cells, the FA molecule on the SERS biological probe cannot target A549 cells. Besides, SERS biological probe without FA modification cannot target folate receptor on the cell membrane. On the other hand, it also proved that the SERS biological probe has been successfully modified by FA molecule. As shown in Figure 3.5, there were appreciable SPION-PEI@AuNPs-MBA-rBSA-FA biological probe on the MCF-7 cells while there was few SPION-PEI@AuNPs-MBA-rBSA-FA biological probe on the A549 cells. Figure 4.6a showed the SERS spectral of MCF-7 cells and A549 cells after incubated with SPION-PEI@AuNPs-MBA-rBSA-FA. Besides, the stability of SPION-PEI@AuNPs-MBA-rBSA-FA have been tested by the SERS imaging of single MCF-7 cell and A549 cell as shown in Figure 3.6a. As shown in Figure 3.6b, MCF-7 cells have peak signal at 1076 cm⁻¹ while there was no peak signal for A549 cells. Besides, peak intensity at 1076 cm⁻¹ was increased when there were MCF-7 cells. Therefore, the Raman peak at 1076 cm⁻¹ was utilized for mapping biological probe in the cell surface. The red area showed obvious

SERS intensity at 1076 cm⁻¹ and the dark area showed no SERS intensity at 1076 cm⁻¹. These results showed good specificity and stability of SPION-PEI@AuNPs-MBA-rBSA-FA to FR positive cells, which could profit CTC detection in the clinical blood samples. In addition to the specificity, separation performance has also been repeated. Figure 3.7 showed separated MCF-7 cells.



Figure 3.5: Optical images of MCF-7 and A549 after incubated with PBS solution, SPION-PEI@AuNPs-MBA SERS biological probe and SPION-PEI@AuNPs-MBA-rBSA-FA SERS biological probe. Scale bar: 20 µm.



Figure 3.6: a) SERS imaging of MCF-7 and A549 cell at bright, 1076 cm⁻¹ intensity peak and merge background. Scale bar: 20 μ m. b) SERS spectral of MCF-7 and A549.

3.4 Sensitivity and separation of SPION-PEI@AuNPs-based SERS biological probe in blood

To study the sensitivity of SPION-PEI@AuNPs-based SERS biological probe for CTC detection, different proportions of MCF-7 cells were added into health rabbit blood to imitate peripheral blood environment of cancer patients. Concentration gradient dilution method can accurately control the number of cancer cells added to rabbit blood by diluting cell suspension to concentration of 500, 250, 100, 50, 10, and 5 cells per milliliter. As shown in Figure 3.7, when there were 5 MCF-7 cells added into rabbit blood, MCF-7 cells could be detected and separated.



Figure 3.7: a) Optical image of separated MCF-7 cells from 5-500 cells/mL mixed cell suspension, Scale bar: 20 μm; b) Raman spectral of separated cells.

3.5 Application of SPION-PEI@AuNPs-based SERS biological probe in the clinical blood samples

SPION-PEI@AuNPs-based SERS biological probe has been proved to have good detection efficiency and veracity. It has been applied to CTC detection of clinical blood samples. To test the performance of SPION-PEI@AuNPs-based SERS biological probe in clinical blood detection, 32 blood samples from tumor patients and 3 blood samples from healthy people have been detected. Figure 3.8 showed detection results of healthy people blood samples. It showed that there was no peak enhancement at 1076 cm⁻¹. Figure 3.9 showed SERS detection results of 32 blood samples. Compared to healthy blood results, there were obvious peak enhancement at 1076 cm⁻¹ in almost spectral. However, there were still 5 tumor blood samples detected without 1076 cm⁻¹ peak as shown in red frame, which could be defined as false negative results. Besides, the Raman spectral was unstable due to complex blood background. There were many undesired peaks appeared which would disturb determination of results. Therefore, the stability of SERS biological probe in blood should be improved and it is better to remove the complex background before SERS detection so that the accuracy of results could be improved.



Figure 3.8: SERS detection results of blood from healthy people.


Figure 3.9: SERS detection results of 32 tumor blood samples.

3.6 Chapter conclusion

In summary, SPION-PEI@AuNPs-MBA-rBSA-FA SERS biological probe was successfully reproduced and applied for CTC detection. The SERS-active magnetic nanoparticles were demonstrated to have good specificity and sensitivity to MCF-7 cells. The LOD of CTC detection was measured to be 5 cell per mL. 32 cancer blood samples and 3 blood samples from healthy people were tested. Although the results showed good specificity and separation efficacy, there are still some limitations to be overcome. For examples, density gradient centrifugation as blood pretreatment method may cause loss of targeted cells and redundant SERS biological probes could not be effectively removed, which will cause false negative results and false positive results, respectively. besides, hot-spot effect between noble nanoparticles might also disturb judgment of results.

Chapter 4 Synthesis of B-TiO₂-based SERS biological probe and application on CTC detection

4.1 Introduction

In the third chapter, gold-iron oxide composite nanoparticles were used as SERS substrate for CTC detection. Although it has strong SERS intensity, the hot spot efficacy might cause false positive results. Besides, the toxicity to living organisms and nonselective SERS enhancement of targeted molecules limit their application in CTC detection[117, 206]. It has been reported that low-cost semiconductor nanomaterials can also exhibit high SERS performance with advantages of high spectral stability and reproducibility, strong anti-interference ability, and selective SERS enhancement of target molecules[117]. Lin et al. firstly investigated the crystal-amorphous core-shell heterojunction boosted photoinduced charge transfer (PICT) process for the enhancement of black TiO₂ (B-TiO₂)-molecule system and applied it to the detection of cancer cells[118]. The PICT process is formed between the substrate and the target molecules [130]. The strong vibration coupling in a semiconductor molecule system can effectively increase the molecular polarizability and amplify the Raman scattering cross section[125]. The amorphous structure of TiO₂ improved the PICT efficiency, yield an ultrahigh SERS enhancement factor (EF) and, achieve higher performance of semiconductor-based SERS substrates[207]. Therefore, semiconductor-based SERS substrate shows great potential in CTC analysis.

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In this chapter, black TiO_2 was used as SERS substrate for CTC detection to reduce hot spot efficacy. FA-functionalized SERS biological probe has been designed to distinguish CTCs from the remaining WBCs by targeting the FA biomarkers on the CTC membranes. The SERS biological probe was composed by a crystal-amorphous core-shell B-TiO₂ SERS substrate, Raman reporter alizarin red (AR) molecule, a thin NH2-PEG2000-COOH (PEG) layer to improve the dispersion of biological probe, and FA to specifically recognize folate receptor (FR) on cancer cell membrane (Figure 2.4). Besides, abundant amino group of the PEG layer can provide sufficient binding sites for FA coupling, which is benefit for improving the detection specificity and accuracy of B-TiO₂ based biological probe. SERS-fluorescence bimodal imaging method verified high detection specificity of this strategy. In addition, SERS results demonstrated that FR positive MCF-7 cancer cells could be captured efficiently with reliability and specificity compared to FR negative cancer cells (A549 and Raw264.7 cells).

4.2 Characterization of B-TiO₂-Based SERS biological probe

B-TiO₂ NPs have been successfully fabricated. As shown in Figure 4.1, the color of B-TiO₂ was black while the color of P25 was white. The Low-resolution transmission electron microscopy (TEM) and the High-resolution transmission electron microscopy (HRTEM) were used to identify the

microstructure and morphology of B-TiO₂ NPs. The TEM image showed the particle size of B-TiO₂ NPs was ~25 nm (Figure 4.2a). The HRTEM image showed a crystal-amorphous core-shell structure of B-TiO₂ NPs (Figure 4.2b). As shown in Figure 4.3, there was an obvious oxygen-deficient amorphous shell with 2-3 nm thickness on the surface of crystalline core. The feature of amorphous shell and crystal core structure was further confirmed by the corresponding fast Fourier transform (FFT) image. The diffuse ring for the shell structure and the points pattern for the core structure were consistent with the HRTEM characterization. The lattice plane distance of the crystalline core was 0.35 nm, corresponding to the (101) crystal planes of anatase TiO₂. Meanwhile, X-ray diffraction (XRD) showed that both P25 and B-TiO₂ have the major characteristic diffraction peaks of anatase TiO₂ at 25.4 $^{\circ}$ (101) (Figure 4.4). The XRD image also characterized the crystalline structure of P25 NPs before and after reduction. It showed that the crystalline degree was reduced after the reduction because of the formation of an oxygen-deficient amorphous layer. Besides, the increased half width of the Raman peaks for the P25 at 145, 196, 400, 517, and 642 cm⁻¹ frequencies also indicated the decreased crystallinity in B-TiO₂ NPs (Figure 4.5). At the same concentration of AR, B-TiO₂ has much better SERS performance than P25 (Figure 4.6).



Figure 4.1: Optical photo of P25 and B-TiO₂, scale bar: 2 cm.



Figure 4.2: a) TEM image of B-TiO₂; b) HRTEM image of B-TiO₂.



Figure 4.3: FFT image of B-TiO₂.



Figure 4.4: X-ray diffraction of P25 and B-TiO2 NPs.



Figure 4.5: Raman characterization of P25 and B-TiO2 NPs, respectively.



Figure 4.6: SERS spectra of AR adsorbed on the B-TiO₂ NPs and P25 at 5×10^{-3}

М.

To investigate the SERS performance of B-TiO₂, SERS spectra of AR molecules adsorbed on the B-TiO₂ NPs at different concentrations $(5 \times 10^{-3} - 5 \times 10^{-3})$ 10⁻⁸ M) was measured at 532 nm (Figure 4.7). Figure 4.7 showed the Raman spectra difference between AR and AR absorbed on B-TiO2. Before AR absorbed on B-TiO₂, AR has weak Raman spectral. After AR absorbed on B-TiO₂, the Raman spectral of AR has been strongly enhanced, which indicated that AR has been absorbed on B-TiO₂. When the concentration of AR is reduced to 5×10^{-8} M, the vibration peak could still be distinguished, indicating that the LOD of the AR molecule on B-TiO₂ can reach 5×10^{-8} M. Compared to some semiconductors as shown in Table 1.5, the SERS performance of B-TiO₂ was significantly great. The concentration of AR used to fabricate SERS biological probe was 5×10^{-3} M. To show the binding of AR on B-TiO₂, the conjugation efficiency of AR absorbed on B-TiO₂ was calculated by mass balance method. First, after the centrifugation of B-TiO₂-AR mixture, the supernatant was collected, and the residual AR content was determined using the calibration curve of AR standard solutions by measuring the UV-vis absorption peak at 422 nm (Figure 4.8a and b). To determine the residual AR content more accurately, the supernatant was diluted 3 times before measuring. The conjugation efficiency (CE) of AR on B-TiO₂was calculated as follows:

$$CE_{AR} = \frac{\text{weight of initial AR-weight of residual AR}}{\text{weight of (B-TiO_2-AR)}} \times 100\%$$
(13)

Therefore, the CE of AR in B-TiO₂ was 15.1%, which means the amount of AR

is 15.1 mg per 100 mg B-TiO₂-AR.

To improve the dispersion of B-TiO₂-AR biological probe and avoid nonspecific adsorption, NH₂-PEG2000-COOH was used to enwrap B-TiO₂-AR. The TEM image of B-TiO₂-AR-PEG showed that there was a thin PEG layer on the particles (Figure 4.9). The thickness of PEG layer was about 4-5 nm.

FA was modified to the SERS biological probe by the amide interaction between the carboxyl group of FA and amino group of NH₂-PEG2000-COOH. The conjugation efficiency of FA on the SERS biological probe was 2.1% calculated by mass balance method. After the activated FA mixed with B-TiO₂-AR-PEG stirring for 12 hours, the resulting B-TiO₂-AR-PEG-FA was obtained by centrifugation at 10000 rpm for 10 minutes. the supernatant was collected, and the residual FA content was determined using the calibration curve of FA standard solutions by measuring the UV-vis absorption peak at 350 nm (Figure 4.10). The conjugation efficiency (CE) of FA on B-TiO₂-AR-PEG-FA was calculated as follows:

$$CE_{FA} = \frac{\text{weight of initial FA-weight of residual AR}}{\text{weight of (B-TiO_2-AR-PEG-FA)}}$$
(14)

To further confirm the FA modification, B-TiO₂-AR-PEG biological probe and B-TiO₂-AR-PEG-FA biological probe were incubated with FITC linked polyclonal rabbit antibody to FA, respectively. The fluorescence intensity of B-TiO₂-AR-PEG-FA biological probe (Figure 4.11a) was obviously stronger than B-TiO₂-AR-PEG biological probe (Figure 4.11b). The SERS intensity of B-TiO₂-AR-PEG-FA biological probe was measured as shown in Figure 4.12a. Although the intensity was reduced due to PEG layer, it still has strong signal for satisfying peripheral blood tumor cell detection. The spectra stability of B-TiO₂-AR-PEG-FA biological probe was also verified in Figure 4.12b, which is in favor of CTC detection in complex peripheral blood environment.



Figure 4.7: SERS spectra of AR, B-TiO₂ and B-TiO₂-AR NPs at different concentrations ($5 \times 10^{-3} - 5 \times 10^{-8}$ M).



Figure 4.8: a) UV-vis spectra of AR solution. b) Calibration curve of AR standard solutions at peak 422 nm.



Figure 4.9: TEM image of B-TiO₂-AR-PEG.



Figure 4.10: a) UV-vis spectra of FA solution. b) Calibration curve of FA standard solutions at 350 nm.



Figure 4.11: Fluorescent images of FITC linked polyclonal rabbit antibody to FA. a) FA modified SERS biological probe. b) non-FA modified SERS biological probe. Scale bar: 50µm.



Figure 4.12: a) SERS spectra B-TiO₂-AR-PEG-FA compared to B-TiO₂-AR; b)

The SERS spectra of SERS biological probe at 5 points.

4.3 FR expression screening of MCF-7, A549 and Raw264.7 cell lines

Three cell lines of MCF-7, A549 and Raw264.7 were used to verify the performance of this SERS biological probe. First, the function of FR in the

three types of cell lines was verified by incubating cells with FITC-PEG2000-FA and flow cytometry measurement was carried out. As shown in Figure 4.13, MCF-7 had higher affinity for FITC-PEG2000-FA, while A549 and Raw264.7 showed extremely low affinity. It was demonstrated that MCF-7 was FR positive cell while A549 and Raw264.7 were FR negative cells.



Figure 4.13: Flow cytometry analyses of the function of FR in a) MCF-7, b) A549, c) Raw264.7.

4.4 Specificity and stability of B-TiO₂-based SERS biological probe in the mixed cells

After incubated with B-TiO₂-AR-PEG biological probe and B-TiO₂-AR-PEG-FA biological probe, there were obvious differences between MCF-7 cells and A549 cells (Figure 4.14). For MCF-7cells, there were B-TiO₂-AR-PEG-FA biological probes on the cells while there was few B-TiO₂-AR-PEG biological probe on the cells. For A549 cells, both of B-TiO₂-AR-PEG biological probe and B-TiO₂-AR-PEG-FA biological probe were not observed on the cells. To illustrate detection specificity of B-TiO₂-AR-PEG-FA biological probe, MCF-7 and A549 cells were also stained by FITC and Hoechst, respectively, as shown in Figure 4.15. MCF-7 cells exhibit red fluorescence on the cell membrane, while A549 cells with almost no red fluorescence on the cell membrane, where the red fluorescence was come from AR molecule absorbed on the SERS biological probe.



Figure 4.14: Optical images of MCF-7 and A549 after incubated with PBS solution, B-TiO₂-AR-PEG SERS biological probe and B-TiO₂-AR-PEG-FA SERS biological probe. Scale bar: $20 \mu m$.



Figure 4.15: Fluorescent images of A549 and MCF-7 incubated with SERS biological probe. Scale bar: $20 \ \mu m$.

Besides, the SERS mapping images of MCF-7 and A549 cells after incubated with FA modified SERS biological probe were shown in Figure 4.16a. The typical Raman scattering peak of AR molecule at 1449 cm⁻¹ was more significantly enhanced than that at 1257 cm⁻¹, 1326 cm⁻¹ exhibited in Figure 4.16b, so the Raman peak at 1449 cm⁻¹ was utilized for mapping biological probe in the cell. Three Raman spectra collected from MCF-7 and A549 respectively, were shown in Figure 4.16b. To make sure the repeatability and reliability of results in Figure 4.16b, other 10 measurements for MCF-7 and A549 were provided, respectively as shown in Figure 4.17. The Error Bars in figures and relative standard deviations (RSD) calculated for the SERS intensity at peak 1449 cm⁻¹ demonstrated the good repeatability of the method. There was no SERS signal observed on A549 cells while MCF-7 had obvious SERS signal on the surface. This result demonstrated that FA modified SERS biological probe can effectively target cancer cells with FR expression.



Figure 4.16: a) Optical image and SERS mapping image of MCF-7, and A549 cancer cell at 1449 cm⁻¹. Scale bar: 20 μ m. b) SERS spectral of SERS biological probe with MCF-7, and A549 cancer cell collected from 3 points, respectively.



Figure 4.17: a) SERS spectral of SERS biological probe with MCF-7 collected from 10 points. b) SERS spectral of SERS biological probe with A549 collected from 10 points. c) SERS intensity of SERS biological probe with MCF-7 and A549 at peak 1449 cm⁻¹.

To verify the anti-interference ability of B-TiO₂-AR-PEG-FA biological probe from FR negative cells, MCF-7 and Raw264.7 cell lines were mixed and incubated with B-TiO₂-AR-PEG-FA biological probe for 1 hour and washed by PBS for three times. After stained by CK8 (MCF-7 positive, Raw264.7 negative) and Hoechst (MCF-7 positive, Raw264.7 positive), MCF-7 has red fluorescence on cell membrane, blue fluorescence on nucleus and green fluorescence from AR, while Raw264.7 only has blue fluorescence on nucleus (Figure 4.18). To distinguish the fluorescence from AR and Alexa Fluor 647, the fluorescence of AR was set as green and fluorescence Alexa Fluor 647 was set as red. B-TiO₂-AR-PEG-FA SERS biological probes were obviously observed in MCF-7 cell membrane, while Raw264.7 with only few SERS biological probes. This result showed that B-TiO₂-AR-PEG-FA SERS biological probe exhibited good anti-interference ability for distinguish MEC-7 cells from the FR negative Raw264.7 cell.



Figure 4.18: Fluorescent images of MCF-7 and Raw264.7 incubated with SERS biological probe. Scale bar: $10 \ \mu m$.

4.5 Chapter conclusion

In summary, B-TiO₂-AR-PEG-FA SERS bio-probes were successfully fabricated. Demonstrated by three cell lines of MCF-7 (FR positive), A549 and Raw264.7 (FR negative), SERS biological probes of B-TiO₂-AR-PEG-FA could distinguish FR positive CTCs from FR negative cells by targeting FR on CTC membranes. Besides, through SERS-Fluorescence bimodal imaging method, B-TiO₂-based SERS biological probe has also been verified with high

detection specificity and accuracy. However, the detection results were moderately inaccurate without isolation of CTCs, which should be improved in the future work.

Chapter 5 Combination method of microfilter and B-Tio₂-based SERS biological probe on CTC detection

5.1 Introduction

In the fourth chapter, B-TiO₂ based SERS biological probe could successfully distinguish FR positive CTCs from FR negative cells by targeting FR on CTC membranes in the mixed cells. To improve the detection accuracy and applied it into clinical blood samples, microfluidic method was combined for blood pretreatment.

Microfluidics has been widely used in CTC detection based on the different biological properties and physical properties of CTCs compared with the counterpart of blood cells[208-210]. Microfiltration is one of label-free microfluidic methods which only depends on the size and deformability difference. It comes to the foreground in recent years benefiting from the ability of isolating both epithelial and mesenchymal phenotypes, which are more appropriate for analyses of tumor heterogeneity, and tumor drug resistance[208]. Although microfiltration can isolate CTCs with high capture efficiency and simplicity for operation within a few minutes[211, 212], there are still tens of thousands of WBCs remaining, which would disturb the accuracy and sensitivity of CTC detection. The traditional way to identify CTCs is fluorescent immunolabeling by characterizing the surface protein expression of CTCs. However, when using multiple fluorophore-conjugated antibodies, the broadness of fluorescence spectrum and the cross-talking of fluorescent tags often confound the results[213]. Besides, the background noise was magnified in a microfluidic system, which might cause misrecognition of CTCs[214]. Therefore, it is necessary to develop a novel downstream analysis strategy of CTCs combined with microfluidic method.

In this chapter, the combination of SERS method and microfluidic method could advance both advantages of these two methods as well as overcome their own limitation. A novel strategy for in-situ isolating and detecting CTCs at single-cell resolution via combining microfilter CTC isolation method and SERS biological probe detection method was developed, as shown in Figure 5.1. First, the microfilter efficiently isolated CTCs from peripheral blood based on the size and deformability difference between CTCs and blood cells. Then, a FA-functionalized SERS biological probe has been designed to distinguish CTCs from the remaining WBCs by targeting the FA biomarkers on the CTC membranes. SERS-fluorescence bimodal imaging method verified high detection specificity of this strategy, and the limit of detection (LOD) of CTCs in rabbit blood can reach to 2 cells/mL. In addition, SERS results demonstrated that FR positive MCF-7 cancer cells could be captured efficiently with reliability and specificity compared to FR negative cancer cells (A549 and Raw264.7 cells). The whole processes were simple and efficient, which reduced detection time within 1.5 hours. Importantly, blood samples from

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cancer patients have also been successfully detected based on the microfiltration isolation and B-TiO₂-AR-PEG-FA SERS biological probe detection method.



Figure 5.1: Scheme of CTC detection process.

5.2 Simulation of optimum capture flow rate in microfilter

The pathway of WBC passing through the micropore has been simulated as shown in Figure 5.2. The blue part represented blood and the red part represented WBCs. When the flow rate was 0.1 mL/min, WBC cannot pass through the micropore (Figure 5.2a). When the flow rate reached to 0.2 mL/min and 0.3 mL/min, WBC can pass through the micropore but might be retained within the micropore with tendency of sticking to the micropore wall

(Figure 5.2b and Figure 5.2c). When the flow rate was 0.4 mL/min, the cell can pass through the micropore with a complete structure and intact shape (Figure 5.2d). When the flow rate was 0.5 mL/min and 0.7 mL/min, the structure of WBC was slightly affected by flow condition after passing through the micropore (Figure 5.2e and Figure 5.2g). When the flow rate was 0.6 mL/min, the shape of WBC was slightly deformed (Figure 5.2f). When the flow rate was 0.8 mL/min, the cell was deformed significantly after passing through the micropore (Figure 5.2h), while the cell was completely damaged when the flow rate was 0.9 mL/min (Figure 5.2i). Therefore, the microfilter has the best WBC clearance performance when the flow rate was 0.4 mL/min.



Figure 5.2: The effects of different blood flow rates on WBCs flowing through the pore of the microfilter. The blue part represents blood and the red part represents WBCs. a) u=0.1mL/min, b) u=0.2mL/min, c) u=0.4mL/min, d) u=0.4mL/min, e) u=0.5 mL/min, f) u= 0.6mL/min, g) u=0.7 mL/min, h) u=0.8 mL/min, i) u=0.9 mL/min.

Figure 5.3 shows the variations of volume in blue and contact pressure in red during the period when a WBC flows through the micropore. According to the red curve, the force applied on cell from the source side appears with axial displacement from 2 μ m to 7.5 μ m approximately when there are interactions between the sides of contact pair. In this simulation, the contact of channel and cell is dealt with by setting a penalty function, which cause the deformation of cell described by the volume change shown by the blue curve. The stress on the cell shell from the inner structure is taken into consideration by setting a global equation to restrict the volume of cell to the initial volume. Therefore, the two peaks of volume variation correspond to the local maxima of contact pressure. It should be noted that the trends of the curves are not synchronous, but with a temporal difference since the value of the state variable is adapted in a way that the associated global equation is satisfied. Figure 5.4 shows the variations of volume and contact pressure during the period when a CTC flows through the micropore. As shown in Figure 5.4, the hysteresis of contact pressure is greater than that of WBC shown in Figure 5.3, which might be due to the volume of a CTC is larger than that of a WBC and the reaction of internal pressure is not so sensitive as CTC.



Figure 5.3: The variations in cell volume and maximum contact pressure with regards to the axial displacement of a WBC flowing through a micropore in the microfilter.



Figure 5.4: The variations in cell volume and maximum contact pressure with regards to the axial displacement of a CTC flowing through a micropore in the microfilter.

5.3 Sensitivity, specificity, and capture assay of microfilter combined SERS spectral detection method in blood

To explore the clinical utility of microfilter combined with SERS spectral method, clinical tumor blood model was built to carry out cancer cell capture

and detection studies. The clinical tumor blood model was prepared by mixing 1-100 cells/mL MCF-7 cancer cells into rabbit blood. The WBC clearance could reach 96% by counting the cell concentration before and after filtering. The LOD of this strategy in rabbit blood was 2 cells/mL as shown in Figure 5.5b. Figure 5.5a demonstrated the schematic diagram of the CTC detection process via microfilter and SERS biological probe, which was composed of CTC microfiltering, biological probe targeting, and SERS detection. The SERS spectral of 9 MCF-7 cells after isolation and targeted by SERS biological probe was shown in Figure 5.5d, and SERS signal of the cancer cells exhibited relatively high uniform, illustrating efficient SERS detection of CTC in peripheral blood sample based on effective microfluidic separation technology. Then, the captured cancer cells were further verified by CK8 and Hoechst (Figure 5.5c), Fluorescent image results indicated that the accuracy of this strategy, which showed that microfilter combined with SERS spectral strategy has huge application potential in clinical application.



Figure 5.5: a) Schematic illustration of CTC detection steps. b) LOD of CTC based on the method of combining microfilter and SERS method. c) Fluorescent images of captured cells. Scale bar: 10 μ m. d) SERS spectral of the captured cancer cells. Laser wavelength: 532 nm; power: 0.12 mW; lens: 50× objective.

5.4 Application in clinical tumor blood detection

The strategy of microfluidic isolation process and B-TiO₂-AR-PEG-FA SERS biological probe has been successfully leveraged to detect clinical peripheral blood samples from six breast cancer patients and two health people. The SERS spectra signal acquired from six blood samples of breast cancer patients were unambiguous and reproducible (Figure 5.6), while there was no SERS signal observed in two health people (Figure 5.7). The SERS spectra Cancer patient blood samples and health people blood samples were carefully confirmed by collecting 3 different laser points on the cell examined under a microscope. Besides, the time of completing the whole strategy was less than 1.5 hours. The results illustrated that the CTCs from clinical breast cancer patients can be accurately distinguished via microfilter combined SERS spectral detection method. This novel cell isolation and detection method could be served as an efficient platform utilized in cancer early screening and diagnosis, as well as postoperative evaluation, making great contributions to precision medicine.



Figure 5.6: Schematic illustration of CTC detection based on microfilter and SERS spectra. (b-g) Optical image and SERS spectral of the captured cancer cells from six breast cancer patients. Scale bar: 20 μ m. Laser wavelength: 532 nm; power: 0.12 mW; lens: 50× objective.



Figure 5.7: Optical image and SERS spectral of captured cells from healthy people. Scale bar: 20 μ m. Laser wavelength: 532 nm; power: 0.12 mW; lens: 50× objective.

5.5 Chapter conclusion

In summary, an integrated CTC detection strategy based on microfilter, and SERS detection method was successfully developed for in-situ isolating and recognizing CTCs at single-cell resolution. For CTC isolation, it could reach to 96% WBC clearance by adjusting pulling speed in microfilter and reduce the complexity of blood environment. The low purity and specificity of microfilter were also solved by combining high-sensitivity SERS spectra detection in this study. Demonstrated by three cell lines of MCF-7 (FR positive), A549 and Raw264.7 (FR negative), SERS biological probes of B-TiO₂-AR-PEG-FA could distinguish FR positive CTCs from peripheral blood cells efficiently by targeting FR on CTC membranes and ruling out false positive interference of WBCs with reliability and specificity. Besides, through SERS-Fluorescence bimodal imaging method, B-TiO₂-based SERS biological probe has also been verified with high detection specificity and accuracy. Compared to the traditional fluorescent immunolabeling method, SERS biological probe can distinguish cancer cells from blood cells with high intensity and sensitivity avoiding the influence of background noise. Besides, the use of microfilter improved the efficacy of CTC isolation compared to the traditional density gradient centrifugation method. Benefiting by these advantages, the combination of SERS method and microfilter enhanced the detection efficiency and veracity, which reduced the detection time within 1.5 hours and make the LOD of detection reach to 2 cells/mL. Though the SERS biological probe in this paper was only suitable for FR positive CTCs, it still has great practical values because most of cancer cells are FR overexpressed. These features also facilitated successful CTC detection in several clinical cancer patient bloods which illustrates that the integration of microfluidic isolation and SERS detection was worth to be further developed and a promising tool for diagnosis and treatment of cancer.

Chapter 6 Conclusion and future perspective

6.1 Conclusion

In summary, three strategies have been successfully carried for CTC detection in clinical blood samples. In the first strategy, gold-iron oxide composite nanoparticles were selected as SERS substrate. SPION@Au-MBA-rBSA-FA SERS biological probes were reproduced according to the previous work. In this thesis, SPION@Au-MBA-rBSA-FA SERS biological probes were applied into clinical blood sample detection. 32 blood samples from cancer patients and 3 blood samples from healthy people were detected. Although the results showed good specificity and separation efficacy, there are still some limitations to be overcome. For examples, density gradient centrifugation as blood pretreatment method may cause loss of targeted cells and redundant SERS biological probes could not be effectively removed, which will cause false negative results and false positive results, respectively. besides, hot-spot effect between noble nanoparticles might also disturb judgment of results. In the second strategy, black B-TiO₂ NPs were used as SERS substrate due to its high spectral stability and reproducibility, strong anti-interference ability, and selective SERS enhancement of target molecules. The B-TiO₂-AR-PEG-FA biological probe was composed by a crystal-amorphous core-shell B-TiO₂ SERS substrate, Raman reporter alizarin red (AR) molecule, a thin NH₂-PEG2000-COOH (PEG) layer to improve the dispersion of biological probe, and FA to specifically recognize folate receptor (FR) on cancer cell membrane.

Besides, abundant amino group of the PEG layer can provide sufficient binding sites for FA coupling, which is benefit for improving the detection specificity and accuracy of B-TiO₂ based biological probe. It could accurately target FR positive MCF-7 cancer cells from FR negative cancer cells (A549 and Raw264.7 cells). In the third strategy, microfluidic method was added to isolate CTCs before SERS detection. In this strategy, the low purity and specificity of microfilter could be solved by combining high-sensitivity SERS spectra detection, while the microfilter could reduce interference of blood background during SERS detection. Besides, SERS-fluorescence bimodal imaging method verified high detection specificity of this strategy, and the limit of detection (LOD) of CTCs in rabbit blood can reach to 2 cells/mL. Compared to pretreatment of blood by density gradient centrifugation, the whole processes of combined method was simple and efficient, which reduced detection time within 1.5 hours. Compared to the first strategy, there was less undesired peak in Raman spectral, which made results easier to be determined and with more accuracy. These results illustrates that both of microfluidic isolation and SERS detection could open new paths for liquid biopsy.

6.2 Future perspective

With the rapid development of material science and processing technology, microfluidics has obtained breakthrough progress. Researchers are no longer satisfied with chemical reactions on the microfluidic chip, but also testing various analytes with microfluidics. In recent years, optical and spectroscopic technology has been developed as a high-sensitive, rapid, efficient, and nondestructive detection and imaging method, in which SERS has become one of the most commonly used spectral methods. SERS method has been widely applied in biosensor.

The combination of SERS method and microfluidic method has become a new research trend. SERS method can be used to detect ultra-low concentration samples within microfluidic channel while the multi-channel design of microfluidic chip can facilitate multivariate SERS detection. SERS method can provide excellent analysis method for microflow platform. Besides, the SERS substrate made by microfluidics can solve the limitations of irregular fluctuation of SERS signal in solutions, and thus greatly improving the repeatability and reliability of detection. The combination of SERS and microfluidic chips has provided many new opportunities for the development of both technologies, and their complementarity has also solved their respective problems, laying a solid foundation for multifunctional SERS Labon-Chip systems. There is also a promising of the opportunity in biological, chemical detection, medical diagnosis, and other fields.

Although the microfluidic SERS detection chip has made remarkable research progress, the detection specificity and repeatability, the improvement of chip

multi-functionality and integration are the development directions and research emphases in the future, in particular of the direct detection of actual clinical biological samples. The real clinical samples, such as patient blood, are very complex, containing rich proteins, DNA, and other elements. These biomacromolecules may possess non-specific noise signals during SERS detection, which may result in false positive or false negative results and seriously affect the reliability of detection results. Therefore, for real samples, the specificity of microfluidic SERS detection chip is particularly important. Generally, the specificity of microfluidic SERS detection chip can be solved from two aspects. First, the biological functional modification of SERS substrate or probe can be optimized to improve the selectivity and accuracy of substrate or probe in recognizing the analyte; second, the number of functional units of analyte screening can be increased, such as physical size screening or chemical ligand recognition screening, to eliminate non-specific interference of other substances as much as possible. If microfluidic SERS detection chip can detect real complex biological samples with high accuracy, it will greatly promote clinical diagnosis and treatment. In addition, low-cost automatic microfluidic SERS detection chip is also one of the key objectives. The focus of research is to reduce the production cost of the chip keeping the perfect sensing and analysis functions. Low-cost microfluidic SERS chip has very important social value and practical significance for biochemical detection, such as disease survey and environmental monitoring, in underdeveloped areas.

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Besides, multi-dimensional analysis of CTCs has been a newly emerging research area, instead of restricting to the detection of single gene or protein. As mentioned above, CTCs are heterogeneous in phenotype. Different cellular subpopulations may exist in individual tumors and some of them have shown to be responsible for initiating cancer metastasis and accelerating cancer development. Additionally, during cancer progression and medication, the phenotypic status of CTCs could dynamically evolve. Therefore, it is of great importance to monitor the phenotypic distribution of CTCs at the resolution of individual cells rather than acquiring an average information, which could be more adaptable to predict disease progression and therapeutic efficiency. In addition, the heterogeneity of CTCs is also reflected among different cancerbearing patients. For example, breast cancer has at least three subtypes categorized according to the cancer cell molecular profiles. Each of these subtypes has their specific strategies for treatment and different risks of disease progression. To some extent, the knowledge of phenotype status in cancer cells could determine the therapy and therapeutic outcomes. Consequently, technologies for profiling and categorizing CTCs on the basis of the phenotypic information at single-cell resolution are in great demand.

In the future, SERS and microfluidic chip technology will be further developed. For example, the integration of laser, spectrometer and other devices on the chip will play a greater role, making a great contribution to the improvement of

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on-site real-time detection technology, benefiting the development of life health science, biomedicine and other aspects. It can be envisioned that with the continuous improvement of functions and performance, integrated and automated SERS Lab-on-Chip method will inevitably become a very important technology in the field of biosensor detection for cancer diagnostics related biomedical applications.

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Appendix

	Methods	Isolation	Cancer cell		Cancer cell		CTCs				
Cancer type		marker	lines	Recovery	Capture efficiency	Throughput	Viability	Number of	Sensitivity	Capture	Ref.
								samples	CTCs/ml	efficiency	
Colorectal								13 CRC 25	CRC: 7.2±3.4		
cancer; prostatic	Immunocapture	EpCAM	PC3	74.2-84.4%	-	1 ml/h	96%	DCa	PCa:	-	[215]
carcinoma								PCa	8.3±5.0		
Lung cancer	Immunocapture	EpCAM	A549	~90%	-	1ml/h	Up to 93%	21	-	-	[216]
Breast cancer	Immunomagnetic capture	EpCAM	MCF-7	95.8%	-		93.1 ± 2.6%	10	-	94%	[217]
Lung cancer	Immunomagnetic capture	EpCAM	H-1650	95.7%	-	5-25ml/h	-	5	31–96 CTCs/ml	-	[218]
Breast cancer; non-small-cell lung cancer	Immunocapture	EpCAM	MCF-7, NCI-H1650	76.5%, 82.7%, 83.7% In 1ml whole blood	88%	-	-	-	-	-	[219]
Breast cancer	Hydrogel-based capture	EpCAM	MCF-7	$95 \pm 4\%$	92 ± 6%	1 ml/h	95%	13	3-12 CTCs /ml	-	[63]
Cervical carcinoma	Hydrogel-based capture	Folic Acid	Hela	-	-	-	-	-	-	85%	[64]
Breast cancer	Immunofluorescence	EpCAM	Her2	94%	-	1ml/30min	-	82	90 CTCs /7.5ml	-	[70]

Table 1: Performance of label-dependent microfluidic methods for CTC isolation

Cancer type	Methods	Cancer cell lines	Cancer cell lines				CTCs			
			Recovery	Purity	Throughout	Viability	Number of	Sensitivity	Purity	
							samples	(CTCs/ml)		
Lung cancer,	Vortex	MCF-7	84%	84% 80% 8 ml/min 83.9% ± 4.0% 22 breast cancer Lung cancer 0.5-24.2		Lung cancer 0.5-24.2	-	[220]		
Breast cancer							15 lung cancer	Breast cancer 0.75-23.25 CTCs/ml		
Lung cancer,	DFF	MCF-7, T24	MCF-7: 87.6%	-	350 µL/min	-	5 breast cancer	20-135 CTCs/ml	1CTC/30-	[221]
Breast cancer			T2476.4%				5 lung cancer		100WBCs	
Lung cancer	DFF	MCF-7	-	-	7.5 ml/5min	-	58	Lung cancer 82 CTCs/ml	-	[222]
		NSCLC 1650						Breast cancer 55 CTCs/ml		
Lung cancer	DFF	A549	74.4% capture	-	25 mL/h	-	34		90 % detection rate	[223]
			efficiency							
Breast,	FAST	MCF-7, MDA-	95.9 ±3.1%	-	>3 mL/min	-	142	Breast cancer	-	[224]
stomach, and		MB-231, MDA-						0-540 CTCs/7.5 mL		
lung cancer		MB-436, HCC78,						stomach cancer		
		AGS						2–485 CTCs/7.5 mL		
								lung cancer		
								0-62 CTCs/7.5 mL		
Lung cancer	Filtering	H446, A549, SK-	90%	-	15mL/h	-	45	1.85-68.45 CTCs/mL	-	[225]
		MES-1								
Lung cancer	Vortex	MCF7	56%	35.1% ±	-	-	16		-	[226]

Table 2: Performance of label-independent microfluidic methods for CTC isolation

				7.3%						
Lung cancer	DLD,	Lung cancer cell	86%		-	-	200	I stage	-	[227]
	filtering	line						$5.0\pm5.121CTCs/mL$		
								II stage		
								$8.731 \pm 6.36 CTCs/mL$		
								III stage		
								$16.81\pm9.556\ CTCs/mL$		
Lung cancer	Vortex	A549, H1703,	-	-	-	-	22	1.32 CTCs/mL	48.4% -96.8%	[228]
		H3255							detection rate	

	Method	Isolation marker			Cance	er cell lines	CTCs			
Cancer type			Cancer cell lines	Recovery	Purity	Throughout	Number of	Sensitivity CTCs/ml	Purity	Ref.
					5		samples			
Lung, breast, prostate cancer	DLD and Immunocapture	EpCAM	MCF-7 PC3-9	90% 89.7 ± 4.5%	-	576 ml/h	41 prostate patients	≥0.5 CTC/m	>3.5-log purification 1500 WBCs/ml of whole blood	[229]
Breast cancer	DLD and Immunomagnetic capture	ЕрСАМ	MCF-7	90%	>40%	9.6 mL/min	-	-	-	[230]
Breast, prostate, lung, melanoma cancer	DLD and Immunomagnetic capture	CD16 CD45 CD66b	H1975, H3122, LNCAP, PC3, PC3-9, VCAP, MB231, MCF-7, SKBR3		-	15×10^6 cells/s	38 breast, prostate, lung, melanoma	84.0% 96.4% 68.5% 63.6% of patients	-	[231]
Bladder cancer	PFF and Crossflow	-	UC13	~98%	89%	10 ⁴ cells /h	-	-	-	[232]
Breast cancer	Hydrodynamic and Microsieve	EpCAM	MCF-7	95%	-	1ml/12.5min	26 pancreatic patients	80% of patients	-	[233]

Table 3: Performance of multi-step methods for CTC isolation