Abstract. The aim of the present study was to discuss the design of a microfluidic chip consisting of columns, and its use for the enrichment of nasopharyngeal cancer (NPC) cells. A microfluidic chip experiment was simulated using FLUENT software. Within the microfluidic chip, aptamers were bound to the reaction chamber (consisting of columns) using a biotin-avidin system. Cell suspension was introduced into the reaction chamber to capture NPC cells. NPC cells were subsequently eluted, and the capture rate of the cells was calculated. The modified aptamer-bound microfluidic chip was able to capture NPC cells with a capture rate of ~90%. The modified aptamer-bound microfluidic chip has a wide range of potential applications for the diagnosis of NPC.

Introduction

Nasopharyngeal cancer (NPC) is one of the most common cancer types in South China. In the early stages of the disease, NPC is usually asymptomatic, with the majority of patients being diagnosed in the middle or advanced stages. Radiotherapy is the primary treatment for NPC; however, 40-60% of patients relapse following definitive radiotherapy (1-5). Relapse primarily occurs locally, in the neck region, or a combination of the two. The majority of relapse cases appear between 2 and 3 years after radiotherapy.

Relapse and metastasis of NPC are direct causes of mortality. Radiological examination is the major method of detecting relapse and metastasis. However, in order to be visualized radiologically, lymph node lesions resulting from relapse and metastasis must reach a maximum diameter of 0.8 cm, according to the Union of International Cancer Control (6), at which point the chance of successful treatment begins to decrease (7). Preceding relapse and metastasis, tumour cells gain access to the peripheral blood, where they are referred to as circulating tumour cells (CTCs) (8). Tumour burden is directly associated with the number of CTCs in the blood; therefore, tumour status may be determined by CTC detection (9).

Aptamers are oligonucleotide or peptide molecules, which, through their unique secondary structure, bind to a specific target molecule (10). Aptamers may be screened using the Systematic Evolution of Ligands by Exponential Enrichment technique, and due to ease of synthesis, storage and transportation, have been employed in an extensive number of applications (11). Microfluidic chips contain complex fluids that are manipulated, observed, detected and controlled at the micron level. In the present study, the microfluidic reaction chamber of a chip was modified for the incorporation of an array of bypass columns. This design increased the area/volume ratio of the fluid environment, thus increasing the reaction efficiency of the microfluidic chip (12,13). Aptamers targeting NPC cell markers were prepared in a preliminary study (14) and subsequently affixed to the inner wall of the reaction chamber within the microfluidic chip. This modified aptamer-bound microfluidic chip is capable of capturing NPC cells from peripheral blood, for the non-invasive diagnosis of NPC.

Materials and methods

Modelling of an array of bypass columns in the microfluidic chip. The shape, arrangement and length of each major axis of the bypass column were simulated using Fluent 6.3.26 software (ANSYS, Inc., Canonsburg, PA, USA). Alterations in the magnitude of shear force and flow rate were analysed in the microarray. The columns were designed with a circular, elliptical (ratio of major axis to minor axis, 10:7) or square form, and arranged in either an aligned or staggered fashion. Flow rates between 10 and 2,000 µm/sec were employed. Multiple arrays of bypass columns were simulated using the finite element method (FEM) (15). The microarray with optimal performance was selected for preparation.
Preparation of the microfluidic chip. The polydimethylsiloxane (PDMS) microfluidic chip was manufactured by Suzhou Wenhao Chip Technology Co., Ltd. (Suzhou, China). The SU-8 mould was prepared via a photolithographic process (16,17), while the PDMS array and cover slips were prepared by PDMS demoulding technology. The typical final step in preparing a closed chip for microfluidic experiments was accomplished using the bonding process (18,19). The chip was composed of an inlet, reaction chamber and an outlet; the reaction chamber was designed with an array of circular bypass columns in a staggered arrangement. The diameter of the bypass columns was 40 µm, and the spacing between columns was 100 µm.

Modification of the inner wall of the microfluidic chip. Biotinlabelled aptamers (Sangon Biotech Co., Ltd., Shanghai) were bound to the inner wall of the reaction chamber of the microfluidic chip using a two-step approach, firstly by binding the avidin (Thermo Fisher Scientific, Inc.) through physical absorption to the inner wall of the reaction chamber, followed by the immobilization of the biotinlabelled aptamers to the inner wall of the reaction chamber using the biotin-avidin mechanism. Avidin is a glycoprotein and each avidin molecule consists of four subunits and each subunit can bind to one biotin molecule. Immobilization of the aptamers to the inner wall of the reaction chamber was determined using fluorescence microscopy (magnification, x400).

Aptamer synthesis and buffers. Aptamers were synthesized by Takara Biotechnology Co., Ltd., (Dalian, China). The aptamer sequence is as follows: Fluorescein isothiocyanate-5'-ACC GACCGTGCTGACCTACCCGGCGAGTGAGTTGGTACGTGTACGCTGAGGGGGTACGGTTGTTACGTTCCAGTGATAGCGGCGGCGGT GCG-3'-Biotin. Buffers used in this study included elution buffer (DPBS containing 0.1 mg avidin/ml, 4.5 g NaCl/ml, 0.25% tRNA and 1 mg bovine serum albumin), and capture buffer (DPBS containing 0.1 mg avidin/ml, and 1 mg bovine serum albumin), and capture buffer (binding buffer and Histopaque-1119 from Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) at a 1:1 ratio.

Cell culture. The NPC cell line C666, used to screen aptamer in the present study, human gastric cancer cell line SGC7901, human colorectal cancer cell line HT29, human ovarian cancer cell line SKOV3, human cervical cancer cell line Hela, and normal nasopharyngeal epithelial cell line NP69 were preserved at our laboratory. Keratinocyte-serum-free medium (SMF), 1640 medium, and foetal bovine serum (FBS) were purchased from Gibco (Thermo Fisher Scientific, Inc.). C666, SGC7901, HT29, A549, SKOV3 and HeLa cells were cultured in 1640 medium containing 10% heat-inactivated FBS and 100 U/ml penicillin-streptomycin. NP69 cells were cultured in keratinocyte-SFM. All cells were washed twice with Dulbecco's phosphate-buffered saline (DPBS) and treated with 0.25% trypsin. Following trypsinization, the cells were either harvested or routine passage was performed.

Cell capture experiment. Prior to cell capture, the concentration of the cell suspension was adjusted to 10⁶ cells/ml, and cells were stained with Vybrant DiI or DiD dye (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Staining was performed at 37°C for 5 min according to the manufacturer's protocol. The cells were rinsed once with wash buffer and resuspended in capture buffer. The cell concentration was adjusted to 10⁶ cells/ml, and the solution was kept on ice prior to use.

The microfluidic chip was treated with 1 mg/ml avidin for 1 min and washed three times with binding buffer. After incubating the chip in 30 µM of biotinlabelled aptamers for 1 min, the chip was again washed three times with binding buffer. Finally, 1 ml of cell suspension (C666 and NP69) was prepared using capture buffer, and injected into the microfluidic chip using a micro-pump. To elute the target cells, the chip was washed three times with binding buffer. To verify the capture efficiency of the modified microfluidic chip, a flat microfluidic chip was constructed and the capture of target cells for the two methods was compared by t-test (n=4).

Separation of tumour cells from whole blood. Whole blood samples (7.5 ml) were collected from healthy volunteers and combined with an EDTA anticoagulant. Physical absorption and aptamer immobilization were performed as described above. Avidin were bound to the inner wall of the reaction chamber by physical absorption, followed by the immobilization of the biotinlabelled aptamers to the inner wall of the reaction chamber. To prevent cell loss during trypsinisation, NPC cells were treated with enzyme-free cell disassociation solution, and subsequently added to the whole blood of the healthy volunteers. The NPC cells from the whole blood samples were captured by the microfluidic chip; the final concentrations of NPC cells were 10, 100, 1,000, and 10,000 cells/ml, and each test was performed in triplicate.

Capture of NPC cells. Whole blood samples containing a C666 cell suspension or NP69, SGC7901, HT29, SKOV3 and Hela cells were transported to the reaction chamber via the inlet of the microfluidic device. The outlet of the reaction chamber was connected to a cell harvester. To avoid introducing variation into the whole blood samples, no buffer was added during cell separation. A micro-magnetic stirrer was placed near the inlet and outlet to prevent the cell aggregation and ensure uniform distribution within the sample.

Statistical analysis. The shear force within different column types, fluid flow rate and length of the major axis was analysed by independent sample t-test for two-group comparison, and by analysis of variance (ANOVA) using the least significant difference, Bonferroni and Dunnett's T3 post hoc accordingly, for the comparison of more than two groups. The purity of captured cells and the capture efficiency at different lengths of the major axis and flow rate were determined by Pearson correlation analysis. The difference in capture efficiency between the ordinary and the modified microfluidic chip was analysed by t-test. To confirm the association of input and capture of different types, fluid flow rate and length of the major axis was analysed by Pearson correlation analysis. All data were analyzed by SPSS 20.0 (IBM Corp., Armonk, NY, USA). P<0.05 was considered to indicate a statistically significant.

Results

Effects of different parameters on bypass column efficiency. Different arrays of bypass columns were simulated using Fluent software. The impact of column shape and fluid flow rate on
the flow of NPC cells was examined. The fluid flow rate at the inlet had a considerable impact on the shear force within the micro-chamber. The flow rate was gradually increased, and the maximum shear force in different arrays was compared. As the flow rate near the inlet increased, the maximum shear force also increased (Fig. 1). With a flow rate <400 µm/sec (50, 100, 200 µm/sec), no significant difference was observed in the maximum shear force across different arrays by ANOVA (P>0.05; Fig. 1A). At a flow rate ≥400 µm/sec (400, 600, 800, 1,000, 1,500 and 2,000 µm/sec), the shear force significantly differed between circle staggered and circle in line, circle staggered and ellipse in line, circle staggered and ellipse staggered, circle staggered and square in line, circle staggered and square staggered, square staggered and circle in line, square staggered and ellipse in line, square staggered and ellipse staggered, square staggered and square in line, ellipse staggered and ellipse in line, and circle in line and ellipse in line arrangements (Fig. 1B). Maximum shear force was observed for the circle columns in a staggered arrangement, followed by the square columns in a staggered arrangement. The shear force of the elliptical columns in a staggered arrangement did not differ from that of the elliptical columns in an aligned arrangement (as determined by ANOVA). As determined by t-test (P<0.05), the shear force with flow rate <400 µm/sec was

Figure 1. Effect of flow rate at the inlet on the maximum shear force of the array. The maximum shear forces at numerous flow rates and column shapes were assessed for a staggered arrangement and in-line arrangement of columns (n=3). The maximum shear forces at (A) 50-200 µm/sec and (B) 400-2,000 µm/sec flow rate. (C) The shear forces of flow rate <400 and ≥400 µm/sec. *P<0.05 compared with the circle in line, ellipse in line, ellipse staggered, square staggered and circle staggered.
determined the optimal shear of the microfluidic chip simulations, a range of 20-45 was selected for the subsequent experiments. Separation of target cells by microfluidics. was lower compared with that observed for values >100 (P<0.05), the shear force with a major axis length ≤100 was observed in the maximum shear force across different of the major axis was ≤100. staggered (P<0.05; Fig. 3A). The shear force of square staggered was bigger compared with the circle in line, ellipse in line, ellipse staggered, square staggered and circle staggered (P<0.05; Fig. 3A). The shear force of square staggered was bigger compared with the circle in line and ellipse in line arrangements by ANOVA (P<0.05). When the length of the major axis was ≤100 µm, no significant difference was observed in the maximum shear force across different arrays by ANOVA (P>0.05; Fig. 3B). As determined by t-test (P<0.05), the shear force with a major axis length ≤100 µm was lower compared with that observed for values >100 µm (Fig. 3C). Thus, a major axis length of between 25 and 100 µm was selected for the subsequent experiments.

Separation of target cells by microfluidics. Based on the results of the microfluidic chip simulations, a range of 20-45 µm was selected as the major axis length for the bypass columns; microfluidic chips were prepared with varying lengths from within this range. The effect of major axis length and inlet flow rate on the capture efficiency and purity of target cells was examined. Fig. 4 illustrates that as the major axis length increased, the purity of captured cells also increased, whilst the capture efficiency decreased. Pearson correlation analysis indicated a significant positive correlation between axis length and the purity of the captured cells (R=0.94; P<0.01), a significant negative correlation between axis length and the capture efficiency (R=-0.95; P<0.01), and negative correlation between the capture efficiency and the purity of captured (R=-0.86; P<0.05) at a range of 20-45 µm axis length. With the increase of capture efficiency, the purity of capture decreased therefore, the proper long axis was maximum number of cells obtained from colligating capture purity and capture efficiency (input cells x capture purity x capture efficiency). When the long axis was 45 µm, the target cells were captured the most. Therefore, the length of the major axis was set at 45 µm and the effect of different flow rates on the capture efficiency and purity of the captured cells were also analysed. Fig. 5 illustrates that as the flow rate increased the capture efficiency decreased, whilst the purity of captured cells increased when the long axis was fixed. A negative correlation was indicated between flow rate and capture efficiency (R=-0.97; P<0.01), a positive correlation between flow rate and capture purity (R=0.92; P<0.01), and a negative correlation between capture efficiency and the capture purity (R=-0.92; P<0.01). Therefore, the proper flow rate was the maximum number of cells obtained from colligating capture purity and the capture efficiency (input cells x capture purity x capture efficiency). At flow rates between 200 and 800 µm/sec, the capture efficiency and purity of the captured cells were >70%. The number of captured target cells was the highest at a flow rate of 700 µm/sec; therefore, for subsequent experiments, a major axis length of 45 µm, and a flow rate of 700 µm/sec were selected. Fig. 6 compares the capture efficiency between a conventional flat channel microfluidic chip and modified version, the microfluidic chip reaction chamber with an array of bypass columns defined as the micropillar channel microfluidic chip. The t-test confirmed that the capture efficiency of the modified microfluidic chip was considerably higher compared with that of the conventional one (P<0.01).

To verify the enrichment capacity of microfluidic chip for NPC cells, C666 cells, a nasopharyngeal cancer cell line used to screen aptamer in the present study, were used as target cells, and NP69, SGC7901, HT29, SKOV3 and Hela were selected as controls. C666 cells were enriched using a microfluidic chip bound with biotin-labelled aptamers. For 1 ml of cells, a 1:1,000 ratio of target to control cells was sought, at a final concentration of 10⁶ cells/ml. The two cell types were stained prior to the experimentation; C666 cells were pre-stained with Vybrant DiI dye (red), and NP69, SGC7901, HT29, SKOV3 and Hela were pre-stained with Vybrant DII dye (blue). Fig. 7 represents the cell mixture (target cell, C666; control cells, NP69) prior to (Fig. 7A) and following (Fig. 7B) separation of NPC cells using the microfluidic chip. The modified aptamer-bound microfluidic chip effectively enriched NPC cells with a capture efficiency of 92%, the percentage of tumor cells isolated/total tumor cells present.
Separation of NPC cells from whole blood. The addition of cultured NPC cells into peripheral blood samples was used as a model for CTCs in the peripheral blood of tumour patients. Specifically, varying concentrations of C666 cells were added to 1 ml of peripheral blood, and NPC cells were separated and enriched using the microfluidic chip. The capture efficiency for NPC cells from blood samples was >90%. Fig. 8 indicates that the quantities of captured cells ranged from 10-10,000 with varying concentrations of total NPC cells, and the coefficient of correlation was 0.99 (calculated using linear regression).

Figure 3. Effect of major axis length on the maximum shear force of the array. The maximum shear force at numerous lengths of the major axis and column shapes was assessed for a staggered arrangement and in-line arrangement of columns (n=3). The maximum shear force at (A) 0-75 µm axis length and (B) 150-300 µm axis length. (C) The shear forces axis length of ≤100 and >100 µm. *P<0.05 compared with the circle in line, ellipse in line, ellipse staggered, square staggered and circle staggered.
Discussion

Liquid biopsy for tumours is currently a prevalent topic in the field of tumour diagnosis and treatment. Tumour cells in the peripheral blood of patients can be detected using different techniques (22). Methods commonly employed for the detection of CTCs include: i) Flow cytometry, used for detecting DNA from eight CTCs in breast cancer; ii) immune‑magnetic bead separation; and iii) the CellSearch system (Menarini Silicon Biosystems, Inc., Huntington Valley, PA, USA) developed based on the principle of immune‑magnetic bead‑based selection (23). The CellSearch system is a semi-automatic CTC detector that binds the epithelial cell adhesion molecule of CTCs to facilitate their separation from peripheral blood cells. It is currently used to determine the prognosis of patients with breast cancer. Additional methods include membrane filtration, which separates tumour cells from the peripheral blood on the basis of size and selected cell markers. The aforementioned techniques (except for DNA detection) use high‑affinity antibodies to capture and count CTCs (24‑26). However, the number of available antibodies that are able to bind to tumour cells or cell lines is limited, and off‑target binding remains a major challenge (27). Specific preservation
conditions are required to maintain the functional activity of protein probes, (the principal limitation of antibody-based capture), which restricts the application of these techniques in tumour detection.

Aptamers not only exhibit higher affinity and specificity compared with antibodies, but are also more easily synthesized, preserved, transported and surface-modified. The use of aptamers for capturing CTCs has been reported in recent years, with capture efficiency varying between 70 and 98% (28,29). In the present study, a microfluidic chip was modified by incorporating an array of bypass columns into the reaction chamber to increase the contact area between cells and the inner surface of the chamber. Experiments indicated that the capture efficiency of the modified microfluidic chip (90%) was higher compared with that of a conventional microfluidic chip (78%). Additionally, the modified aptamer-bound microfluidic chip targeting NPC cells that was screened by preliminary research, successfully detected NPC cells in the peripheral blood. The capture efficiency was 90%, higher compared with the reported efficiency in existing studies (30,31).

It was demonstrated that an aptamer-bound microfluidic chip was able to separate NPC cells from whole blood at a higher efficiency than is currently attainable, and neither the aptamer-bound microfluidic chip nor the detected samples required pre-treatment. The method may be performed quickly and with an effective lower limit of 10 cells. The capture efficiency of the aptamer-bound microfluidic chip was determined by preliminary in vitro cell-based experiments, though the true diagnostic value may be verified by clinical patient sample. Additionally, the aptamer of the microfluidic chip may only recognize the corresponding matched epitope, thus may not reliably capture all types of cancer cells. Subsequent research may involve samples from patients with nasopharyngeal cancer with mixed aptamers to capture a greater range of cancer cell types. Despite these limitations, the results of this study implicate far-reaching clinical applications for the modified microfluidic chip.

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Availability of data and materials

The datasets used and/or analysed during the current study are available from the corresponding author by reasonable request.

Authors' contributions

WXC, JGJL, XSZ and WLC were responsible for the conception and design of the present study. XHW, SYQ, YQZ, QMW, JYL, WMX and CX were responsible for the experiment. WXC, WLC and XHW assisted in data analysis and interpretation. All authors were involved in the writing of the manuscript. All authors have read and approved the final version of the manuscript.

Ethical approval and consent to participate

The study was approved by the ethics committee of Jiangxi Cancer Hospital, Nanchang, China. Written informed consent was obtained from all volunteers.

Patient consent for publication

Written informed consent was obtained from all volunteers.

Competing interests

The authors declare that they have no conflict of interest.

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