

Simulation of Single Cell Trapping via Hydrodynamic Manipulation

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Article history

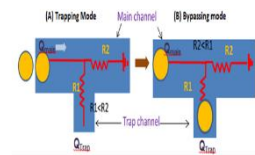
Received :20 February 2014

Received in revised form :

27 May 2014

Accepted :6 June 2014

Graphical abstract



Abstract

Microfluidic devices are important for the single cell analysis such as cell mechanical and electrical characterization. Single cell characterization could be related to many significant applications including early disease diagnosis. However to perform the single cell manipulation, firstly a single cell have to be isolated and a platform for the cell manipulation have to be provided. One of the methods to trap a single cell is by using hydrodynamic trapping in the microfluidic channel. This study provides a finite element model for single cell trapping for a yeast cell model. The objectives of the simulations are to obtain the appropriate channels' geometry and optimized ratio of the fluid's inlet and suction flow rate to trap a single yeast cell. Trap channel was designed to trap a $5\mu\text{m}$ yeast cell with a suction hole placed in the end of the trap channel. Design geometry and the ratio of fluid flow rates for the cell trapping model were studied using the hydrodynamic resistance concept. The analysis was carried out using numerical solutions from the finite element ABAQUS-FEA software. Using the cell trapping model, a single yeast cell able to be trapped into the trap channel with optimized channel's suction hole's geometry and appropriate fluid's inlet and suction flow rate ratio. The appropriate $Q_{\text{Trap}}/Q_{\text{Main}}$ ratio to perform cell trapping using hydrodynamic resistance concept is the ratio value above 1. A $5\mu\text{m}$ yeast cell model able to be trap inside a trap channel with the height, width and length of $7\mu\text{m}$ by manipulating the suction hole's flow rate of 1.5 and $2.0\mu\text{m}$ of height, 7 and $3\mu\text{m}$ of length and width, respectively which situated at the centre edge of the trap channel.

Keywords: Single cell; hydrodynamic resistance; microfluidic; cell trapping

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1.0 INTRODUCTION

Biomedical and biological research nowadays has moved towards a single cell approach. Previous conventional biological studies usually being performed to study a large cell populations and population approach prevent the investigation of an individual cell. The measurement of population based study is the summation of responses that can only reflect the average responses of a cell population. The information inherent to single cells will allow us to resolve heterogeneity and eventually improve understanding of enduring problems in molecular biology, cancer diagnosis, pathology and therapy. The analysis of single cells with a sufficient number of measurement is a need to elucidate heterogeneities which is important to obtain a statistically meaningful data to reveal properties of individual cells and cell differences.

Microfluidic platforms have become an important tool for single cell analysis as they allow constructing fluidic channels with dimensions adapted to a specific cell size and provide fluidic environments for cell analysis with minimal dilution. Microfluidics have advantages to overcome challenges of traditional assays and perform medical diagnosis and handle small sample sizes

and thus minimized the use of valuable reagents in the analysis. The main benefits of micro fabricated systems for cell studies are the capability to design cellular microenvironments, precisely control fluid flows, and to reduce the time and cost of cell culture experiments.

Microfluidic devices can be operated using hydrodynamic forces thus exhibiting numerous advantages such as on-chip cell labeling, short detection time and high reproducibility based on simple and robust experimental procedure. The size-based approach is relatively less invasive because it does not require any chemical or biological interactions between the cells and the fluid. Single cells trap should only allow spatial localization of single cells, but also create reaction chambers, where reactions with stimuli can take place and manipulations could be performed.

The field of Computational Fluid Dynamics (CFD) is improving fast with the ability to achieve approximate, but realistic results of a wide variety of complex and three dimensional viscous flows. The CFD modeling is an invaluable tool that has been applied only relatively recently in the area of microscale cell culture that enables a better understanding of the hydrodynamic environment and the factors that

modulate it. CFD is now enabling us to understand the implications of fluid flow and transport on cell function thus provides important insights into the design and optimization of microfluidic culture chip

This study presents development of the single cell trapping using the microfluidic finite element model using hydrodynamic manipulation techniques. In this paper, we discuss the simulation of the single yeast cell trapping inside the microfluidic channel. The single cell trapping model was studied by performing

W \ U b b Y \ D g \ X Y g \ [b \ [Y c a Y h f m \ c d h] a \ n U h] c b \ U b X \ h \ Y \ f U h] c \ c Z \ Z \ i \] X \ U b X \ g i W h] c b D g \ Z \ c k \ f U h Y g \ h c \ U W \] Y j Y \ g i W W Y g g \ g] b \ [\ Y \ W Y \ \ \ h f U d d]

The paper is divided into five sections with the first section as an introduction and second section explains the hydrodynamic trapping idea and concept of the proposed model. Third section discusses the experimental setup of the simulation, next section involves results and discussion and the last section will be the conclusion.

2.0 THE IDEA AND CONCEPT OF THE MODEL

The concept of hydrodynamic trapping was originally proposed by Tan *et al.* The micro channels are designed such that: (i) when a resistances are than that of the trapping stream and subsequently into the trap; (ii) the bypass channel (main channel in our model) and subsequent trapping site. Darcy-Weisbach equation is used to determine the pressure drop or pressure difference in a channel and solving the continuity and momentum equations for the flow problem.

From Hagen-Poiseuille equation defined as following equation:

$$\Delta P = \frac{128 \mu L Q}{\pi r^4}$$

where ΔP is the pressure drop, μ is the hydrodynamic flow resistance of the rectangular channels, L , H and W are length, height and width of the channel respectively. Considering micro channels as a resistive circuit is analogous to resistance in an electric circuit, this equation is the analogs of V and I , respectively

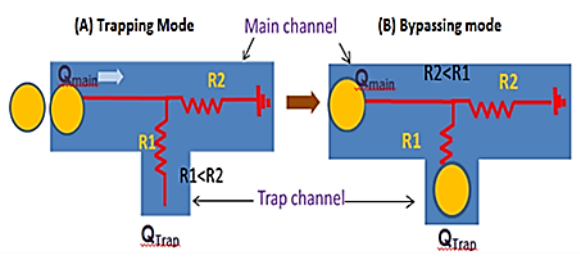


Figure 1 Simple schematic of single cell trapping channel with the hydrodynamic resistance concept

Figure 1 shows the schematic explanation of the hydrodynamic trapping concept with $R1$ and $R2$ representing the flow resistance for trapping channel and main channel,

At intersection (Figure 1), the flow is divided into the main path and the trap path. Yellow circle denotes the yeast cells trapped. The flow rates of the main path (Q_{Main}) and the main path (Q_{Trap}) are distributed depending on the corresponding flow resistances. By using relationships of $A = W \times H$ and $P = 2(W + H)$, the hydrodynamic flow resistance can be formulated as following equation:

where C is a constant that depends on the aspect ratio of the channel. A is the cross-sectional area and P is the perimeter of the channel. The flow rate ratio between trap path and main path is modeled as given in equation (3), approximating that the pressure drop across main path and trap path are the same. For the trap to work, the flow rate along trap path must be greater than that of main path ($Q_{Trap} > Q_{Main}$).

3.0 SIMULATION SETUP

The analysis was carried out using finite element-ABAQUS analysis software which able to perform physical analysis. At first, the simulation analysis was carried out using the parameters in micro dimensions properties. However due to time consumed in simulation to converge is too long (data not shown), the parameters was appropriately scaled into meter dimension with the ratio of 1 m is proportional to the advantage of dimension scaling is that a simulation works could be carried out in a reasonable simulation time. This approach to represents a nano scale model by giving nanometer dimensions to the geometry and using the material property values identical to the real model suffers from two major drawbacks. Firstly, the simulation will face a very small incremental time steps which would make real time simulation prohibitively expensive if not impossible and secondly, using properties with nanometer dimensions cause numerical issues in finite element programs

The single cell trapping system was modeled using Abaqus CAE software. The fluid micro channel was modeled as 3D eulerian explicit EC3DR and a node linear eulerian brick element part assigned with water properties (density, equation of state, viscosity). Figure 2A shows the different parts involved in the model; a eulerian part with the fluid channel and a three dimension (3D) deformable part of the elastic yeast cell model diameter and Figure 2B shows the assembly setup with a yeast cell diameter. The micro channel consists of two channel; the main channel with the width and depth of $10 \mu m$ and total length of $100 \mu m$ and a

channel with $7\mu\text{m}$ of length, width and depth. There is a rectangular suction hole placed at the end of the trap channel. The dimension of the suction hole is a variable ranging from $1.0\mu\text{m}$ to $2.0\mu\text{m}$ with $1.0\mu\text{m}$ was set as the height for the initial simulation analysis.

Results obtained showed that the cell was not trapped inside the trap channel (data not shown). This is due to the suction hole is situated at the bottom of the trap channel. Uneven distribution of fluid velocity produced in the trap channel (low velocity: dark blue, high velocity: grey) caused the pressure drop produced was not enough to capture cell inside the trap channel (Figure 3A).

A very high suction rate (Q_{Main} ratio of 60) is needed to produce wide velocity distribution inside the trap channel. Cell was found able to be trapped when the suction rate is 60 and above. Suction rate required is quite high for the application of micro channels as it may cause deformation of cells. Therefore, another strategy such as increasing the suction hole size have been carried out in the subsequent analysis to produce the appropriate pressure drop with a lower suction rate. The height of suction hole was increased to $1.5\mu\text{m}$ and (Figure 4).

Subsequent simulation is carried out using the design with $Q_{\text{Trap}}/Q_{\text{Main}} > 1$. Results obtained show that cell was able to be trapped into the trap hole using the modified design (Figure 5). The hydrodynamic concept was found able to be applied in the modified design. The verification using the modified model proved that the hydrodynamic concept works accordingly (trapping was successful with $Q_{\text{Trap}}/Q_{\text{Main}} > 1$).

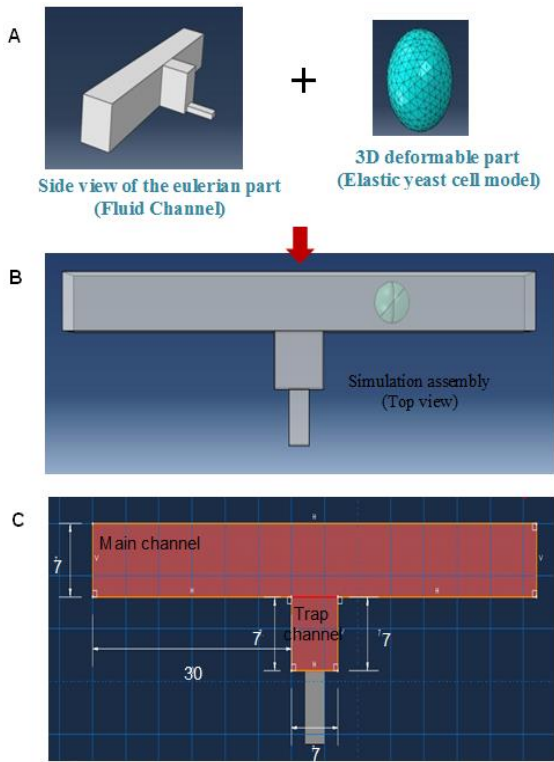


Figure 2 Construction of the finite element model of single cell trapping system. (A) Parts involved in the model. (B) Simulation assembly setup. (C) The dimensions of fluid channel (top view)

A spherical yeast cell ($5\mu\text{m}$ in diameter) was model as an elastic 3D standard solid deformable C3D8R and an 8-noded hexahedron element. The cell was assembled to develop the finite element model for the proposed system (Figure 2). Figure 2C shows the dimensions of the proposed channel. The fluid channel and cell were meshed using hexahedron and tetrahedron respectively. No-inflow and non-reflecting outflow eulerian boundary conditions were applied to the walls of channel. Inflow velocity of 0.3ms^{-1} was applied to the inlet and atmosphere pressure was applied to the outlet of the channel. Various suction velocities ranging from 4 to 40ms^{-1} was applied (depending on the flow rate ratio of main channel and trap channel ($Q_{\text{Trap}}/Q_{\text{Main}}$)) to the suction hole in the trap channel. The interaction between objects and water was modeled using general contact with tangential behaviour and the interaction was modeled using normal behaviour.

4.0 RESULTS AND DISCUSSION

According to the hydrodynamic trapping concept, cell/particle trapping should able to be achieved when $Q_{\text{Trap}}/Q_{\text{Main}}$ ratio is more than 1^9 . To verify whether the concept works in the proposed device, a finite simulation analysis was carried out. Flow rates and hydrodynamic resistance analysis was carried out to the cell trapping model designed with a specific dimensions starting with

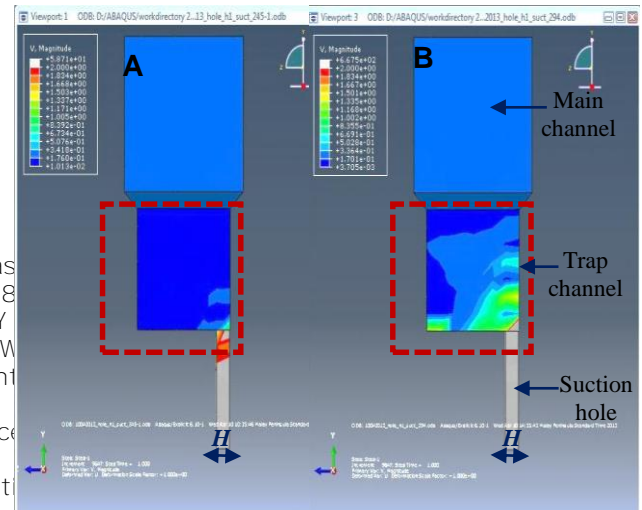


Figure 3 Simulation results show the distribution of fluid velocity inside the trap channel (side view) for $Q_{\text{Trap}}/Q_{\text{Main}}$ ratio of 3 (A) and 60 (B)

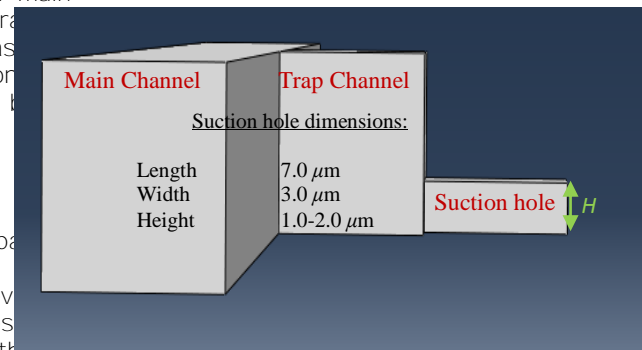


Figure 4 Suction hole's height modification (side view)

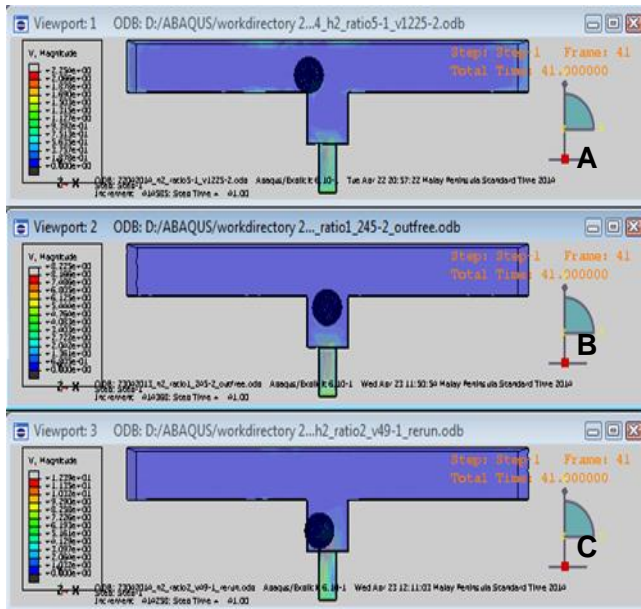


Figure 5 Simulation results (top view) of the modified’s design (suction hole’s height 2 μm) for Q_{Trap}/Q_{Main} ratio of (A) 0.5 (B) 1.01 (C) 2.0

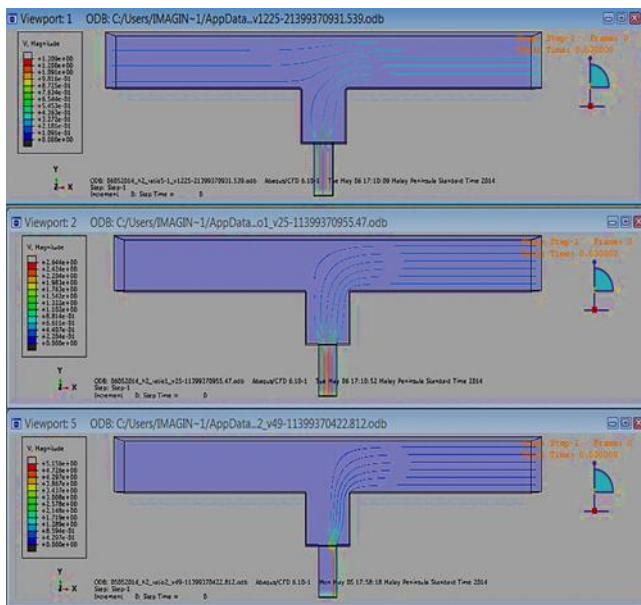


Figure 6 Streamline velocity field of the modified’s design (top view) suction hole’s height of 2 μm for Q_{Trap}/Q_{Main} ratio of (A) 0.5 (B) 1.01 (C) 2.0

Streamline plots of the modified design’s with suction hole’s height of 2 μm were obtained for Q_{Trap}/Q_{Main} ratio of 0.5 to 2.0 are shown in figure 6. The streamline velocity field for Q_{Trap}/Q_{Main} ratio above 1 (Figure 6 B and C) show that the flow diverged from main channel to the trap channel and all the streamlines are directed towards the trap channel. In contrast to Q_{Trap}/Q_{Main} below 1 (Figure A), the velocity streamlines are not fully directed towards the trap channel as portions of the streamlines are directed past through the main channel and towards the trap channel. The streamlines obtained are not fully focusing towards trapping channel and unable to produced not enough force to trap the cell into trapping channel.

Simulation analysis was preceded further with another simulation to study the trapping concept as cell was found able to be trap into the trap hole when Q_{Trap}/Q_{Main} ratio is more than 1 (Figure 8 down images). Both design with suction hole height of 1.5 μm and 2.0 μm are able to obey the hydrodynamic trapping concept as cell was found able to be trap into the trap hole when Q_{Trap}/Q_{Main} ratio is more than 1 (Figure 8 down images). Therefore, subsequent analysis due to the size is not too big and enough to trap single cell and to minimize access stress from executing the trapped cell.

Figure 8 shows the distribution of fluid velocity inside the channels and the trapping of the cell. Colour contours in the result (upper images) show different velocity from low (dark blue) to high (grey). The distribution of fluid velocity from inlet to the outlet are almost the same although the height is different (Figure 8B and 8C upper images). This results show that both design produced almost the same pressure drop for the trapped cell. Therefore, subsequent analysis due to the size is not too big and enough to trap single cell and to minimize access stress from executing the trapped cell.

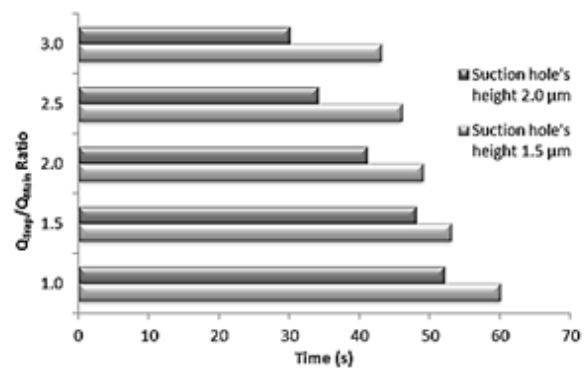


Figure 7 Simulation results of cell trapping time for different Q_{Trap}/Q_{Main} ratio of 1 to 3 for channel suction hole’s height of 1.5 and 2.0 μm

Table 1 Simulation findings for the optimization of suction hole’s height

Ratio of Q_{Trap}/Q_{Main}	Ability to trap cell		
	H : 1.0 m	H : 1.5 m	H : 2.0 m
0.5	x	x	x
1.0	x	yes	yes
1.5	x	yes	yes
2.0	x	yes	yes
2.5	x	yes	yes
3.0	x	yes	yes

Subsequent analysis was carried out using the single cell simulation to study the trapping concept as cell was found able to be trap into the trap hole when Q_{Trap}/Q_{Main} ratio is more than 1 (Figure 8 down images). Both design with suction hole height of 1.5 μm and 2.0 μm are able to obey the hydrodynamic trapping concept as cell was found able to be trap into the trap hole when Q_{Trap}/Q_{Main} ratio is more than 1 (Figure 8 down images). Therefore, subsequent analysis due to the size is not too big and enough to trap single cell and to minimize access stress from executing the trapped cell.

between main channel and trap channel (refer nodes position in chosen is dependent on the type of cell and type of experiment on Figure 9A and 9B). The Q_{Trap}/Q_{Main} ratio value above 1 able to analysis and how will it be performed

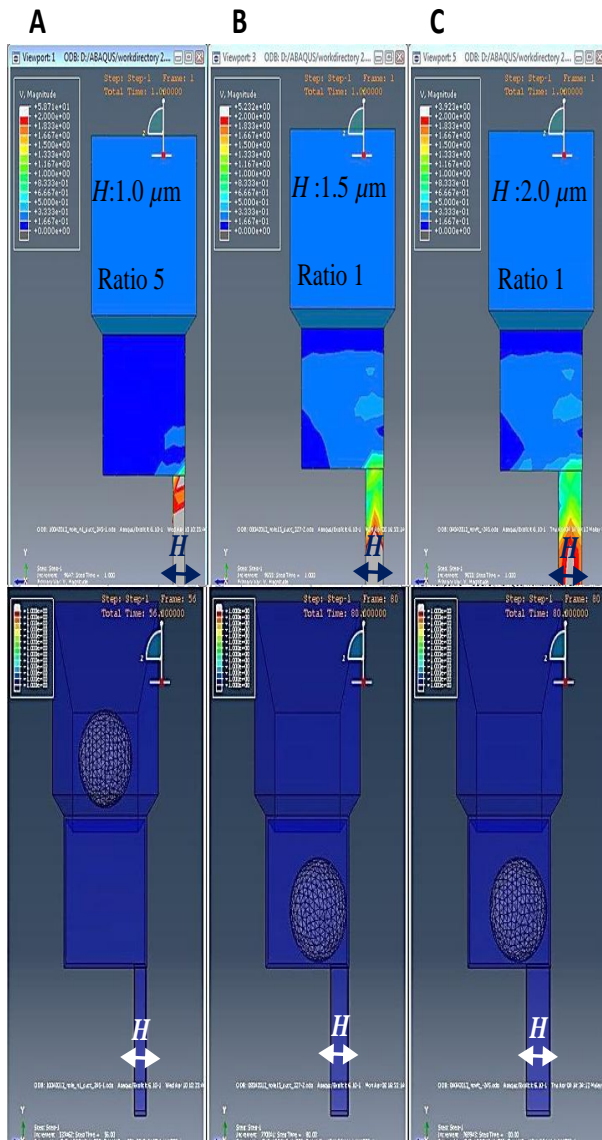


Figure 8 Simulation findings for suction hole's height optimization. Upper figures shows the velocity color contour in the fluid channel (side view) and down figures show the cell trapping result. (A) Suction height 1.0 μm with Q_{Trap}/Q_{Main} ratio = 5. (B) Suction height 1.5 μm with Q_{Trap}/Q_{Main} ratio = 1.01. (C) Suction height 2.0 μm with Q_{Trap}/Q_{Main} ratio = 1.01

and to prevent position variable of the trapped cell. This is important in providing a good platform for cell manipulation in studying the biological, biophysical or biomedical aspect of cells and in also achieving accurate and consistence result. Using hydrodynamic resistance concept is the ratio value above 1. A 5 μm yeast cell model able to be trap inside a trap channel with the height, width and length by manipulating the suction hole of height 7 and 3 μm of length and width, respectively which situated at the center edge of the trap channel. Cell trapping model able to isolate an individual yeast cell inside fluidic environment thus provide a platform to further study the mechanical or biological behaviour of single cell. Single cell manipulation such as chemical and

This study provides a finite element model for single cell trapping specifically trap a 5 μm yeast cell via hydrodynamic resistance manipulation using a suction hole placed in the end of the trap channel. The single cell trapping finite element model was found able to trap a single cell at different flow rates was applied by referring to the hydrodynamic concept. The single cell trapping finite element model was found able to trap a single cell at different flow rate ratio.

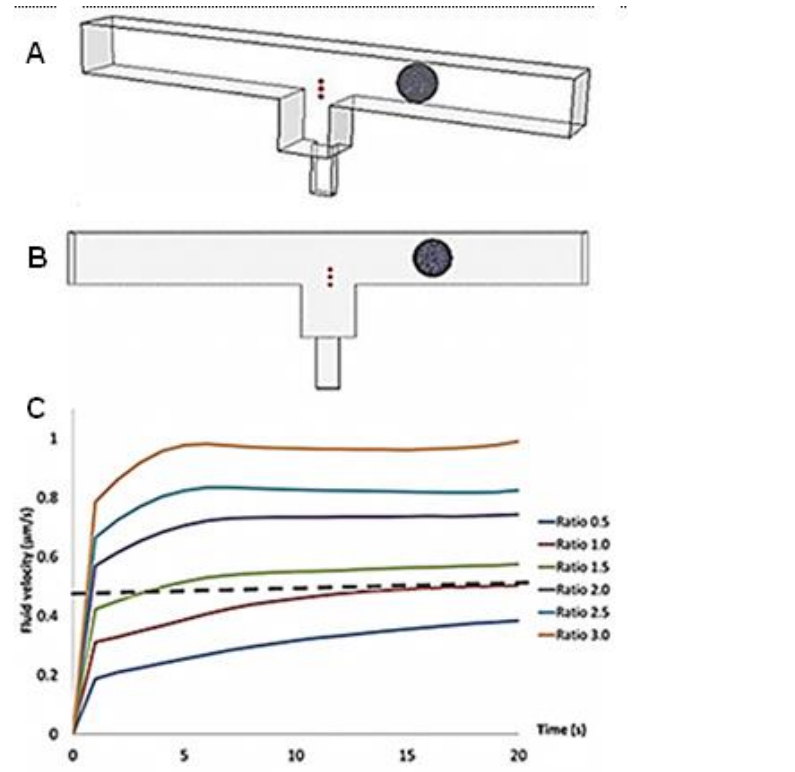


Figure 9 (A) 3D view and (B) Front view of single cell trapping model with the position of 3 nodes (red dots) selected for fluid's velocity data. (C) Graph of velocity of fluid versus time for Q_{Trap}/Q_{Main} ratio from 0.5 to 3.0. Dashed black lines represent the minimum of fluid velocity needed (average velocity of 3 nodes) at the nodes shown in upper figures to trap cell into the trap channel for a fluid's inlet's velocity of 0.3 ms^{-1} .

5.0 CONCLUSION

This study presents the finite element model of single cell trapping inside microfluidic channel. This single cell trapping system able to be constructed using ABAQUS. The single cell trapping model able to obey the hydrodynamic resistance trap concept as the appropriate ratio to perform cell trapping. Using hydrodynamic resistance concept is the ratio value above 1. A 5 μm yeast cell model able to be trap inside a trap channel with the height, width and length by manipulating the suction hole of height 7 and 3 μm of length and width, respectively which situated at the center edge of the trap channel. Cell trapping model able to isolate an individual yeast cell inside fluidic environment thus provide a platform to further study the mechanical or biological behaviour of single cell. Single cell manipulation such as chemical and

biophysical treatments and also mechanical characterization can be performed inside microfluidic channel using this system.

Acknowledgement

We would like to express our appreciation towards Ministry of Higher Education Malaysia (MOHE) grant no. (FR637), (MOHE) grant no. 4L038 (ERGS) and Universiti Teknologi Malaysia, grant no. 57973(NAS), O3H80 (GUP) and O2H34 (GUP) for funding this project and for their endless support.

References

- [1] R. M. Johann. 2006. *Analytical and Bioanalytical Chemistry*. 385: 408.
- [2] M. A. McClain, C. T. Culbertson, S. C. Jacobson, N. L. Allbritton, C. E. Sims, and J. M. Ramsey. 2003. *Analytical Chemistry*. 75(21): 5646–5655, Sep. 2003.
- [3] G. Roman, Y. Chen, P. Viberg, A. Culbertson, and C. Culbertson. 2007. *Analytical and Bioanalytical Chemistry*. 387: 9–12.
- [4] A. K. Price and C. T. Culbertson. 2007. *Analytical Chemistry*. 79: 2614.
- [5] C. E. Sims and N. L. Allbritton. 2007. *Lab on a Chip*. 7: 423.
- [6] S. M. Kim, S. H. Lee, and K. Y. Suh. 2008. *Lab on a Chip*. 8: 1015.
- [7] S. Bhattacharya, T.-C. Chao, and A. Ros. 2011. *Electrophoresis*. 32: 2550.
- [8] M. Huang, S. Fan, W. Xing, and C. Liu. 2010. *Mathematical and Computer Modelling*. 52: 2036.
- [9] W.-H. Tan and S. Takeuchi. 2007. *Proceedings of the National Academy of Sciences of the United States of America*. 104: 1146.
- [10] T. Teshima, H. Ishihara, K. Iwai, A. Adachi, and S. Takeuchi. 2010. *Lab on a Chip*. 10: 244.
- [11] Z. Tang, Y. Akiyama, K. Itoga, J. Kobayashi, M. Yamato, and T. Okano. 2012. *Biomaterials*. 33: 7405.
- [12] I. Kumano, K. Hosoda, H. Suzuki, K. Hirata, and T. Yomo. 2012. *Lab on a Chip*. 12: 3451.
- [13] J. Chung, Y.-J. Kim, and E. Yoon. 2011. *Applied Physics Letters*. 98: 123701.
- [14] G. Thiagarajan, K. Deshmukh, Y. Wang, A. Misra, J. L. Katz, and P. Spencer. 2007. *Journal of Biomedical Materials Research Part A*. 83:332.
- [15] T. Gervais, J. El-Ali, A. Günther, and K. F. Jensen. 2006. *Lab on a Chip*. 6: 500.
- [16] A. K. Bryan, A. Goranov, A. Amon, and S. R. Manalis. 2010. *Proceedings of the National Academy of Sciences of the United States of America*. 107: 999.
- [17] M. R. Ahmad, M. Nakajima, S. Kojima, M. Homma, and T. Fukuda. 2008. *IEEE Transactions on Nanobioscience*. 7: 185.
- [18] A. E. Smith, Z. Zhang, C. R. Thomas, K. E. Moxham, and A. P. Middelberg. 2000. *Proceedings of the National Academy of Sciences of the United States of America*. 97: 9871.
- [19] J. D. Stenson, C. R. Thomas, and P. Hartley. 2009. *Chemical Engineering Science*. 64: 1892–1903.
- [20] J. D. Stenson, P. Hartley, C. Wang, and C. R. Thomas. 2011. *Biotechnology Progress*. 27: 505.
- [21] T. P. Burg, M. Godin, S. M. Knudsen, W. Shen, G. Carlson, J. S. Foster, K. Babcock, and S. R. Manalis. 2007. *Nature*. 446: 7139.
- [22] J. Lee, R. Chunara, W. Shen, K. Payer, K. Babcock, T. P. Burg, and S. R. Manalis. 2011. *Lab on a Chip*. 11: 645.
- [23] K. V. Christ, K. B. Williamson, K. S. Masters, and K. T. Turner. 2010. *Biomedical Microdevices*. 12: 443.
- [24] Christ and K. T. Turner. 2010. *Journal of Adhesion Science and Technology*. 24: 37.