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Published in:
PLoS ONE

Document Version:
Publisher's PDF, also known as Version of record

Queen's University Belfast - Research Portal:
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Download date:22. Oct. 2023
Spheroid Formation of Hepatocarcinoma Cells in Microwells: Experiments and Monte Carlo Simulations

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Abstract

The formation of spherical aggregates during the growth of cell population has long been observed under various conditions. We observed the formation of such aggregates during proliferation of Huh-7.5 cells, a human hepatocarcinoma cell line, in a microfabricated low-adhesion microwell system (SpheroFilm; formed of mass-producible silicone elastomer) on the length scales up to 500 μm. The cell proliferation was also tracked with immunofluorescence staining of F-actin and cell proliferation marker Ki-67. Meanwhile, our complementary 3D Monte Carlo simulations, taking cell diffusion and division, cell-cell and cell-scaffold adhesion, and gravity into account, illustrate the role of these factors in the formation of spheroids. Taken together, our experimental and simulation results provide an integrative view of the process of spheroid formation for Huh-7.5 cells.

Introduction

The study of cell culture in three-dimensional (3D) scaffolds is of considerable intrinsic interest and is also important in the context of numerous applications including, e.g., tissue engineering, disease modeling and drug screening platforms [1–3]. The structure and size of the corresponding scaffolds vary in a broad range from two-dimensional (2D) arrays of sub-millimeter wells to complex 3D structures aiming at mimicking specific organs [2, 3]. Chemically, the scaffolds are often fabricated by using natural hydrogels [2], synthetic polymers [1], or combination of such materials [4]. Cells growing in scaffolds typically aggregate. The shape and morphology of aggregates may be different, depending on various factors including the cell type, design of a scaffold and the corresponding fabrication material [1].

Cellular spheroids represent the most common shape of cell assembly [5, 6]. Aggregates of this shape were created, e.g., by concave microwell method [7], hanging drop method [5, 8], or rotating-wall vessel technique [9, 10]. The size (diameter) of spheroids may reach ~1 cm as observed in
experiments with human colon adenocarcinoma cells [9] and rat hepatocytes [11] (the latter cells displayed liver-like morphology or, more specifically, a compact structure with tight cell-cell junctions, smooth and rough endoplasmic reticulum and bile canaliculi lined with the microvilli).

Often, the size is smaller. For example, the size of spheroids composed of mammary epithelial cells was reported to be ~100 μm (these spheroids can produce and secrete milk proteins upon hormonal stimulation) [5], while in the case of hepatocytes the size was ~200 μm [7].

The growth of cell cultures in scaffolds is of interest also in the context of theoretical biology and statistical physics (for general introduction into this area, see reviews [12–16]). The corresponding models are usually based on the mean-field (MF) kinetic equations or Monte Carlo (MC) simulations. The MF approach is convenient in the situations where the geometry is simple. Such models were used to scrutinize the limitations in the nutrient supply and oxygen transport in porous scaffolds on the coarse-grained level without or with explicit description of single pores (see e.g. references [4, 17, 18] and [18, 19], respectively, and references therein).

MC simulations, based often on the lattice approximation and describing evolution of an ensemble of individual cells, are efficient in the situations with complex geometry and/or in the cases when the focus is on aggregation of cells (as in our present study). The available generic 2D and 3D MC simulations have been focused on the growth and differentiation of stem cells [20], cell seeding [21], and formation of cell sheets [4]. Related theoretical studies concern stem-cell niches [22–25] and scaffold-less biofabrication [26].

Herein, we report the results of our study of culturing Huh-7.5 cells in microfabricated low-adhesion microwells. These cells belonging to a human hepatocarcinoma cell line are widely used as a liver cell model for the exploration of HCV infection [27]. Earlier, we observed the formation of Huh-7.5 cell spheroids in PEG-based hydrogels [28] and multilayer cell sheets in a biofunctionalized 3D scaffold [4, 29]. Our present work is focused on the same cells and has three novel ingredients.

First, we use a recently designed microwell platform for direct observation of the proliferation of cells. Its advantages include: (i) The microwell has a total depth that is two times of its diameter, and walls formed of triangular flat fragments are used to separate adjacent wells. So in contrast to conventional microfabricated semi-circular wells, this mechanical stress (shear force)-free design prevents the cells from slipping during medium exchange, and the method of fluid delivery is diffusion based. (ii) Compared to the hanging drop method [5, 8], the microwell system enables flexible medium exchange during incubation, and due to the small radius of curvature of individual wells (smaller than the size of a water drop), we are able to achieve fairly similar distribution of cell aggregates in different wells. (iii) Another feature distinguishing it from the conventional plastic round bottom wells is that the base (fabricated from silicone elastomer) is easily oxygen-permeable. The latter allows us to reduce hypoxia of cells in the centers of aggregates that is inevitable in conventional plastic scaffolds.

Second, the use of microwell platform described above allowed us to observe explicitly in detail the formation of spherical Huh-7.5 aggregates on the length scale up to 500 μm, with both light microscopy and confocal fluorescence microscopy.

Third, our experimental results are complemented by 3D Monte Carlo simulations to illustrate the role of various factors (e.g., cell diffusion, cell-surface adhesion, and gravity) in the process of cell aggregation and spheroid formation.

**Materials and Methods**

**Pretreatment of 3D SpheroFilm™ for cell culture**

The 3D SpheroFilm™ microwell was obtained from Incyto Co. (Chonan, Korea). The inner diameter of the hemispheres in the microwell is 500 μm, and the total well depth is 1000 μm.
The microwell was made of mass-producible silicone elastomer. For proper cell culture, the microwell was cut to 1-cm² squares, and was placed at the bottom of wells in a 24 well plate. 100% ethanol was added into the plate and repeatedly pipetted to remove the air bubbles from the wells. Once ethanol was removed, the wells were washed three times with phosphate-buffered saline (PBS), and then incubated with cell culture medium for at least 24 hours. The cell culture medium was removed prior to cell seeding.

**Cell culture and cell seeding in 3D SpheroFilm**

Human hepatocarcinoma Huh-7.5 cells, purchased from Apath (NY, USA), were maintained in Dulbecco’s Modified Eagle’s Medium (DMEM) supplemented with 10% fetal bovine serum (FBS), 100 U/ml penicillin and 100 µg/ml streptomycin (Life Technologies) in a humidified atmosphere with 5% CO2 at 37°C. Medium was changed every three days. Cells were detached with 0.25% trypsin-EDTA solution (Life Technologies) from the tissue culture plate, counted, and adjusted to 0.05, 0.1 and 0.2 × 10⁶ cell/ml. 1 ml cell suspension was seeded in each piece of microwell square, cut and placed in 24 well plate. After 10 minutes of cell seeding, suspending cells were removed by aspiration, and the remaining cells were washed and incubated in fresh growth media until time of assay.

**Immunofluorescence staining and imaging**

Cells in the microwells were collected at various stages for immunocytochemistry. Cells were washed twice with PBS, fixed with 4% paraformaldehyde (PFA) for 10 minutes, permeabilized with 0.1% Triton X-100 in PBS for 30 minutes, washed again with PBS and incubated in blocking buffer (3% bovine serum albumin (BSA) in PBS) for 1 hour. Cells were stained with mouse primary antibody against Ki-67 (Life Technologies) by overnight incubation at 4°C, and then washed three times with PBS to remove unbound primary antibody. The cells were then incubated with anti-mouse secondary antibody conjugated with Alexa Fluor® 488 (Life Technologies). Meanwhile, filamentous actin (F-actin) was stained with Alexa Fluor® 555 labelled phalloidin (Life Technologies) for 2 hours at room temperature (protected from light). After two washes with PBS, the nuclei were stained with 10 µg/ml DAPI (Life Technologies) for 30 minutes.
minutes. Fluorescent cell images were taken on a LSM 710 confocal microscope with ZEN program (Carl Zeiss).

Results and Discussion

Cell growth and spheroid formation

The shape and dimensions of 3D SpheroFilm™ microwells are schematically illustrated in Fig 1. Huh-7.5 cells were seeded in the microwells at density of $0.05 \times 10^6$ cell/ml. These cells underwent several stages of morphological changes in the following 10 days (Fig 2). Initially they were attached to the bottom of the hemispheres and randomly dispersed. Later on (Fig 2, day 1), multiple cells started to form clusters due to the seemingly random cell migration and cell-cell adhesion. After day 1, several small cell clusters as well as a number of individual cells gradually merged into large cell aggregates at the center of each well of the microwell surface (Fig 2, day 4). These pre-mature spheroids kept growing bigger and denser, and finally became mature with significant thickness and clear 3D structure at day 7 and onward (Fig 2, day 7 and day 10). When Huh-7.5 cells were seeded at higher densities ($0.1$ and $0.2 \times 10^6$ cell/ml, respectively), similar stages of morphological changes can be observed with earlier cell aggregation and denser spheroids formation (Fig 2).

The cell growth patterns were further examined by fluorescent staining of Ki-67 and F-actin. Ki-67 is used as a marker of proliferating cells [30–32]. The expression of Ki-67 is known to be upregulated in the G1, S, G2 and M phases of cell cycle, and reaches maximum in G2 and M phases [33, 34]. In spheroid cell culture (Fig 3), Huh-7.5 cells showed intense Ki-67 staining (green fluorescence) at day 1 after cell seeding, but the green fluorescence was markedly reduced at day 4 compared to day 1. The expression of Ki-67 at day 7 and 10 was similar to day 4. This observation indicates that the rate of cell proliferation was fastest at day 1, and it slowed down at day 4 and the rate was somewhat held constant till day 10. Another interesting observation is the spatial distribution of Ki-67 expression across the spheroid. At day 1, Ki-67 staining is evenly distributed across the cell aggregates; however, at days 4, 7 and 10, strong Ki-67 cells were found scattered on the periphery of spheroids (see white arrows), meaning that cell proliferation was globally suppressed after day 4, but some cells on the periphery still
maintained their proliferation activity. Interestingly, although cells in the spheroids showed suppressed growth rate, the cells growing outside the microwells (those attached to the flat surface) demonstrated consistent high expression of Ki-67 (Fig 4).

The expression of F-actin displayed similar trend with Ki-67 (Figs 3 and 4). Concerning this aspect, we note that Chang and Hughes-Fulford earlier observed remarkable difference between actin cytoskeleton of cells in monolayers and spheroids [35]. In particular, F-actin stress fibers were found in cells of monolayer culture, while cells in spheroids featured cortical actin that clearly distributed at the outline of the cells [35]. They also reported the upregulation of structural genes including cytoskeletal molecules in monolayers. Thus, it seems to be reasonable to attribute the strong F-actin staining of cells in the microwells at day 1 (Fig 3) as well as cells outside microwells at all the time points (Fig 4) to the cell-substrate interaction and monolayer morphology, whereas when there is more cell-cell interaction in cell spheroid (Fig 3), the F-actin expression is weaker.

Monte Carlo simulations

In reality, the proliferation of cells is usually accompanied by their aggregation due to cell-cell adhesion. In microwells, the aggregation may take place near the microwell walls due to adhesion of cells to the wall surface, and it may result in the formation of cell layers attached to the walls. The size and structure of cell aggregates depend on various factors including the adhesion strength, balance between the rates of cell division and diffusion, structure of the
microwell, cell-cell communication, and limitations in the nutrient or oxygen supply, etc. Our
present 3D lattice MC simulations are focused on the proliferation of cells in a single well of
the microwell system (Fig 1). The emphasis is on adhesion, diffusion and the likely role of grav-
ity in this process. The corresponding parameters are varied in a wide range in order to provide
a general view on the pattern formation in the system under consideration. Some other general
factors are ignored in our simulations (which is inevitable due to the complexity of the system).
In particular, the cell-cell communication is not taken into account primarily because at the
moment we have no data specifying this factor. Concerning this aspect, we may note that the
use of the Ki-67 marker indicates that in our experiments the cell growth occurs mainly at the
periphery of aggregates. It may be related to cell-cell communication and also to spatial con-
straints on the cell division. Our simulations take the latter factor into account, and accordingly
the results are expected to be robust even if the former factor is ignored. The limitations of the
nutrient or oxygen supply are not taken into account either. Such limitations are expected to be
significant roughly on the aggregate length scale larger than 200–400 μm

Fig 4. Ki-67 and F-actin staining of Huh-7.5 cells on flat surface outside the microwells. Scale bar is
100 μm.
doi:10.1371/journal.pone.0161915.g004

To mimic the hemispherical well, we use a 60 × 60 × 80 slab of a cubic lattice. This slab is
cut down to a hemisphere in the lower part (at z < 30, where z is the vertical coordinate mea-
sured in the lattice spacing units) and a cylinder in the upper part (at 30 ≤ z ≤ 80). Each lattice
site can be either vacant or occupied by one cell. Cells can diffuse and divide. As usual in the
lattice models, the diffusion of cells is realized via jumps of monomers to vacant nearest-neigh-
bour (nn) sites. Diffusion (sedimentation) of aggregates is not taken into account partly due to
the lack of simple suitable algorithms and partly because in our case (during the cell
proliferation) this process is expected to be important only at the initial stage of the whole process (when the aggregates are small and their number is large) while we are more interested in the situation when the aggregates are relatively large.

Division of a cell is also considered to be possible if it has a vacant nearest neighbour (nn) site. After division, the cell remains on the same site and another (newly born) cell occupies a vacant nn site. Both for diffusion and division, we use the so-called no-flux boundary conditions, i.e., a jump or division is possible if an nn vacant site belongs to the array of sites representing the well.

To mimic the cell-cell and cell-surface adhesion, we introduce the effective attractive dimensionless (i.e., normalized to \(k_B T\)) interaction, \(\epsilon_{\text{cc}} < 0\), between two cells located in nn sites and the effective attractive dimensionless cell-surface interaction, \(\epsilon_{\text{cs}} \leq 0\), for cells contacting the boundary representing the wall of microwells. In this case, the aggregation is known to occur at \(|\epsilon_{\text{cc}}| > 0.89\) [37]. In our calculations, we set \(\epsilon_{\text{cc}} = -1.2\). The value of the other interaction, \(\epsilon_{\text{cs}}\), is varied to illustrate various situations.

To specify diffusion of monomers in the lattice models, one can use various prescriptions, e.g., those corresponding to the initial-state, Metropolis, or Kawasaki dynamics [38]. We employ the initial-state dynamics which reasonably predict rapid diffusion of single cell and slow diffusion inside aggregates. In this case, the jump rate is reduced by \(\exp(n n_{\text{cc}})\) if a cell has \(n\) neighbours. If a cell contacts the boundary, the jump rate is reduced by the additional factor, \(\exp(n m_{\text{cc}})\), where \(m\) is the number of the corresponding contacts. To characterize the relative rates of division and diffusion, we use the dimensionless parameter \(p_{\text{div}}\). The rates of division and diffusion are considered to be proportional to \(p_{\text{div}}\) and \(1 - p_{\text{div}}\), respectively.

With the specification above, our MC algorithm consists of sequential trials to realize diffusion or division events. A site is chosen at random. If the site is vacant, the trial ends. Otherwise, a cell located in the site tries to diffuse if \(\rho > p_{\text{div}}\) or to divide if \(\rho < p_{\text{div}}\), where \(\rho (0 \leq \rho \leq 1)\) is a random number. In both cases, one of the nn site is selected at random and if the latter site is vacant, the division is performed with unit probability, while the diffusion jump is realized with the probability \(\exp(n n_{\text{cc}})\) or \(\exp(n m_{\text{cc}})\) depending on the local arrangement. After each trial, the time is incremented by \([\ln(\rho)] / N\), where \(N\) is the number of lattice sites, and \(\rho (0 \leq \rho \leq 1)\) is another random number. With the latter prescription, the unit time is, as usual, identified with MC step (MCS). On average, one MCS corresponds to \(N\) MC trials. All the MC runs were started with five cells located on the lattice at random (for the representative results with rapid diffusion, the details of the initial arrangement of cells are insignificant). To characterize the cell concentration, we use the average occupation of sites, \(\bar{\rho}\). The MC runs were performed up to reaching \(\bar{\rho} = 0.4\).

In reality, the diffusion of cells is relatively rapid, and accordingly the value of \(p_{\text{div}}\) should be small. To get an integrated view on the process, it is instructive, however, to show typical patterns and kinetics at various rates of diffusion. Following this line, we first show patterns for \(p_{\text{div}} = 1, 10^{-2},\) and \(10^{-4}\) in the absence of the cell-surface interaction, i.e., \(\epsilon_{\text{cs}} = 0\) (Figs 5–7). If the diffusion is negligible (\(p_{\text{div}} = 1\)), the growth takes place around the seed cells and there are only a few aggregates (Fig 5). In the case of slow diffusion (\(p_{\text{div}} = 10^{-2}\)), the aggregates are numerous and their size is small (Fig 6). In the case of relatively rapid diffusion (\(p_{\text{div}} = 10^{-4}\)), the number of aggregates becomes smaller and their size is larger (Fig 7).

Secondly, using \(p_{\text{div}} = 10^{-4}\) and \(\epsilon_{\text{cc}} = \epsilon_{\text{cs}} = -1.2\), we show the likely role of the cell-surface adhesion (Fig 8). In this case, the cell-surface adhesion is appreciable and the cells are located primarily near the walls of microwells.

In the simulations described above, the aggregates are located either at random (Figs 5–7) or near the walls (Fig 8). In our experiments, however, the cells aggregate primarily at the bottom of microwells near the center (Fig 2). This feature can be reproduced by taking the gravity into account. This factor is usually ignored in MC simulations of cell proliferation, and it was
ignored in our simulations above as well. To mimic the effect of gravity, we reduce the probability of jumps upwards by a factor of \( \exp(\frac{\epsilon_g}{C_{15}}) \), where \( \epsilon_g < 0 \) is the corresponding effective energy. The results of MC simulations with \( p_{\text{div}} = 10^{-4} \), \( \epsilon_{cc} = -1.2 \), \( \epsilon_{cs} = 0 \), and \( \epsilon_g = -0.03 \) show that due to the gravity the proliferation takes place near the bottom of a microwell (Fig 9).

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**Fig 5. Monte Carlo simulation of cell proliferation and spheroid formation in the microwells.** Cross sections of a microwell along the main axis at \( \beta = 0.2 \) (a) and 0.4 (b) in the absence of cell diffusion and cell-surface adhesion \( (p_{\text{div}} = 1, \epsilon_{cs} = 0) \). The well and cells are represented by black and red circles, respectively.

doi:10.1371/journal.pone.0161915.g005

**Fig 6. Monte Carlo simulation of cell proliferation and spheroid formation in the microwells.** Cross sections of a microwell along the main axis at \( \beta = 0.2 \) (a) and 0.4 (b) with slow cell diffusion \( (p_{\text{div}} = 10^{-2}) \) and no cell-surface adhesion \( (\epsilon_{cs} = 0) \).

doi:10.1371/journal.pone.0161915.g006
Figs 5–9 exhibit typical cell growth and aggregation patterns obtained in simulations. The corresponding kinetics of cell proliferation are shown in Fig 10. In the absence of diffusion, the kinetics are slow. With diffusion, the kinetics become faster. The growth of the cell population is predicted to be exponential initially and then becomes close to linear.

**Fig 7. Monte Carlo simulation of cell proliferation and spheroid formation in the microwells.** Cross sections of a microwell along the main axis at \( \beta = 0.2 \) (a) and 0.4 (b) with relatively rapid cell diffusion (\( \rho_{\text{div}} = 10^{-4} \)) and no cell-surface adhesion (\( \kappa_{\text{cs}} = 0 \)).

doi:10.1371/journal.pone.0161915.g007

**Fig 8. Monte Carlo simulation of cell proliferation and spheroid formation in the microwells.** Cross sections of a microwell along the main axis at \( \beta = 0.2 \) (a) and 0.4 (b) in the presence of adhesion (\( \kappa_{\text{cs}} = -1.2 \)) of cells to the bottom and side area of the microwell. The adhesion to the top boundary is neglected because in reality the corresponding plane crosses the solution. The diffusion of cells is relatively rapid (\( \rho_{\text{div}} = 10^{-4} \)). The wall and cells are represented by black and red circles, respectively.

doi:10.1371/journal.pone.0161915.g008
In this work, using a new microwell system for cell proliferation and MC simulations, we have demonstrated that the combination of experimental observation and modeling is a powerful tool for the study of cell spheroid formation. The results suggest that in the 3D cell culture platform used in our study, the cell spheroids would form on the basis of strong cell-cell interaction and weak cell-surface adhesion, and growth of spheroids mainly depends on the cell

Fig 9. Monte Carlo simulation of cell proliferation and spheroid formation in the microwells. Cross sections of a well along the main axis at $y = 0.2$ (a) and $0.4$ (b) in the absence of cell-surface adhesion ($\kappa_{cs} = 0$). Cell diffusion is relatively rapid ($\rho_{div} = 10^{-4}$). In addition, the gravity is taken into account ($\kappa_g = -0.03$).

doi:10.1371/journal.pone.0161915.g009

Conclusion

In this work, using a new microwell system for cell proliferation and MC simulations, we have demonstrated that the combination of experimental observation and modeling is a powerful tool for the study of cell spheroid formation. The results suggest that in the 3D cell culture platform used in our study, the cell spheroids would form on the basis of strong cell-cell interaction and weak cell-surface adhesion, and growth of spheroids mainly depends on the cell

Fig 10. Kinetics of cell growth in the Monte Carlo simulations. Lines 1–5 represent the cell growth kinetics of MC runs shown in Figs 5–9, respectively.

doi:10.1371/journal.pone.0161915.g010
proliferation on their periphery. The likely role of the gravity in the spheroid formation was demonstrated as well. We believe that this strategy could potentially be applied to simulate even more complicated cell systems in the future. Furthermore, the capability to direct specific cells to the formation of aggregates of different types on diverse platforms, including cell spheroids in the microwells (as demonstrated in this study), cell spheroids in hydrogels of different stiffness [28], and multilayer cell sheets in biofunctionalized 3D microscaffold [4], could have significant implications for the construction and development of in vitro tissue models.

Acknowledgments

This work was supported by National Research Foundation of Singapore.

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