

Microfluidic Droplet-Assisted Fabrication of Vessel-Supported Tumors for Preclinical Drug Discovery

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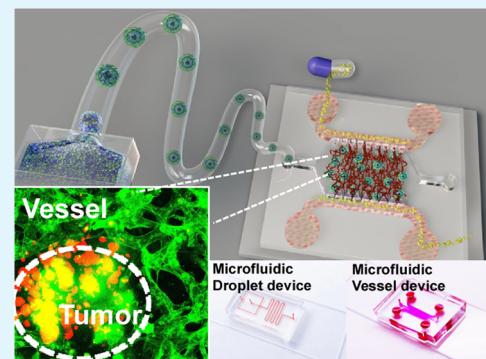
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ABSTRACT: High-fidelity *in vitro* tumor models are important for preclinical drug discovery processes. Currently, the most commonly used model for *in vitro* drug testing remains the two-dimensional (2D) cell monolayer. However, the natural *in vivo* tumor microenvironment (TME) consists of extracellular matrix (ECM), supporting stromal cells and vasculature. They not only participate in the progression of tumors but also hinder drug delivery and effectiveness on tumor cells. Here, we report an integrated engineering system to generate vessel-supported tumors for preclinical drug screening. First, gelatin-methacryloyl (GelMA) hydrogel was selected to mimic tumor extracellular matrix (ECM). HCT-116 tumor cells were encapsulated into individual micro-GelMA beads with microfluidic droplet technique to mimic tumor–ECM interactions *in vitro*. Then, normal human lung fibroblasts were mingled with tumor cells to imitate the tumor–stromal interaction. The tumor cells and fibroblasts reconstituted in the individual GelMA microbead and formed a biomimetic heterotypic tumor model with a core–shell structure. Next, the cell-laden beads were consociated into a functional on-chip vessel network platform to restore the tumor–tumor microenvironment (TME) interaction. Afterward, the anticancer drug paclitaxel was tested on the individual and vessel-supported tumor models. It was demonstrated that the blood vessel-associated TME conferred significant additional drug resistance in the drug screening experiment. The reported system is expected to enable the large-scale fabrication of vessel-supported heterotypic tumor models of various cellular compositions. It is believed to be promising for the large-scale fabrication of biomimetic *in vitro* tumor models and may be valuable for improving the efficiency of preclinical drug discovery processes.

KEYWORDS: microfluidic droplet, core–shell tumor, vessel-on-a-chip, tumor microenvironment, preclinical drug screening



INTRODUCTION

Chemotherapy remains one of the most widely used and effective anticancer treatments.¹ For a new molecular entity, it usually takes 10 years from the drug discovery stage to market approval. Most drug candidates that pass the preclinical development fail in phase I clinical trials.² Particularly, the accurate predictive ability of preclinical physiological models is one of the important factors affecting the efficiency of drug development.^{3–5} In recent years, multicellular tumor spheroid has been increasingly used as *in vitro* model for drug screening assays.^{6–8} Compared with the traditional two-dimensional (2D) cellular monolayer model, the three-dimensional (3D) spheroids can exhibit more similar *in vivo* physiological characteristics, including hypoxic core, pondus hydrogenii gradient distribution, cellular metabolism, and gene expression, thus providing a more realistic drug response.⁹ Accordingly, the manufacturing technology of tumor spheroids has also been continuously improved and developed.^{10–16} Among them, droplet-based techniques stand out for their capability to generate uniform spheroids at a high-throughput manner.^{17–26} As a relatively automated production method, microfluidic droplet technology breaks the device limitations

of traditional well-array plates²⁷ and can mass-produce identical tumor spheroids for drug screening in a time-dependent manner.²⁸ For instance, Kwak et al. reported a microfluidic water-in-oil droplet system for the mass fabrication of uniform tumor spheroids.²⁹ The highest generation rate of the system can reach 1000 droplets/min, and the encapsulated cells in the single droplets can be developed into 3D homogeneous tumor spheroids in 24 h. However, the homotypic tumor spheroids still cannot fully represent the drug response in real tumors. Due to the heterogeneity of tumor tissues, tumor extracellular matrix (ECM), tumor–stromal cells, and tumor–tumor microenvironment (TME) interactions all play important roles in drug resistance.^{30–34} Notably, as the main component of TME, blood vessels have a special vascular transport function and

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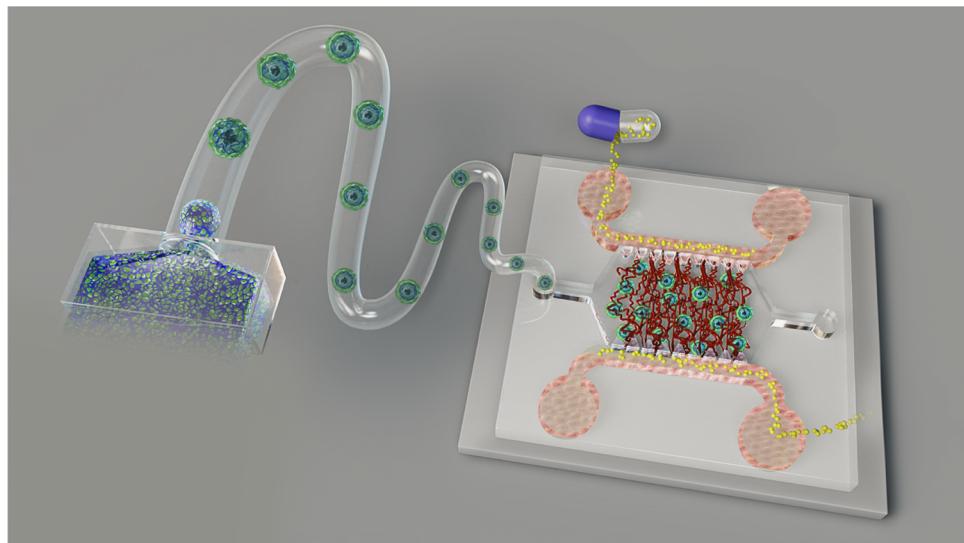


Figure 1. Schematic of the vessel-supported heterotypic tumor model for preclinical drug screening.

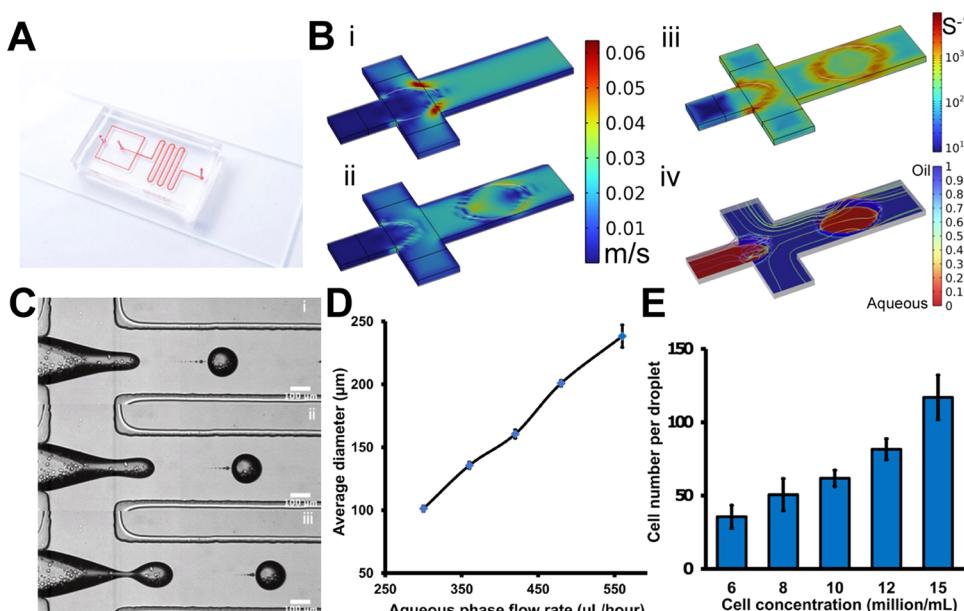


Figure 2. Cell-laden microfluidic droplet generation. (A) Image of the microfluidic droplet generation device. (B) Computational simulation results of microfluidic droplet generation at the flow-focusing zone: the velocity (i, ii), shear rate (iii), and volume fraction (iv). (C) Cell-laden microdroplet generation process. The time interval is 26 μs . (D) Microfluidic droplet size variation according to the aqueous flow rate. (E) Average cell number per individual microdroplet with respect to the original cell concentration. The scale bar is 100 μm .

barrier function, which significantly affect the transport and diffusion of drugs.^{35–40} Therefore, all of these interactions should be considered when constructing tumor models for preclinical drug discovery.

With the continuous development of microfabrication technology, microfluidic tumor–vessel chips not only help to precisely arrange the spatial distribution of cellular components *in vitro* but can also apply biomechanical and biochemical stimuli, including blood flow and growth factor gradient, thereby reproducing the interaction between the heterotypic tumor and TME with high fidelity.^{41–47} For example, Haase et al. reported an on-chip 3D-vascularized tumor model for drug examination in a TME within a large bed of perfusable vasculature.^{48,49} They demonstrated that the dose-dependent effect of anticancer drugs on tumor activity was significantly

affected by the form of vascular infusion administration. Meanwhile, great strides have also been made in the field of biomaterials.^{50,51} As a representative biomaterial, GelMA has been widely used in the regenerative medicine field for 3D cell culture.^{52,53} Various organ models have been successfully established in GelMA, which demonstrate its good biocompatibility.^{5,54,55} Antunes et al. reported an in-air production of tumor spheroid-laden GelMA microgel to expedite drug screening, where GelMA was involved as tumor–ECM.⁵⁶ In addition, GelMA also has the advantages of stability at room temperature, adjustable properties, processable into arbitrary shapes, and instant curing.⁵⁷ Based on this, its combination with different processing techniques makes the spatial distribution of cells in tissues highly controllable, which brings

broad application prospects for tissue engineering (Figure 1).^{58,59}

In this study, inspired by the “bottom-up” engineering concept, a vessel-supported heterotypic tumor model was reported for preclinical drug discovery in a large-scale and controllable manner.⁶⁰ Here, we first mass-produced homotypic GelMA microbeads of tumor cells with a microfluidic droplet generation device to reconstitute tumor–ECM interactions *in vitro*. Then, by mixing the tumor cells with the fibroblasts, heterotypic GelMA microbeads were established to recreate tumor–stromal interactions. After the cellular interactions and self-reassembly, the heterotypic tumor beads presented a core–shell structure with the core as tumor clusters and the shell as the fibroblast monolayer. Next, the developed cell-laden beads were incorporated into an on-chip vessel network platform to reestablish the tumor–TME interaction. As a proof-of-concept application, the antitumor drug, paclitaxel (PAC), was administrated on the individual tumor beads and vessel-supported tumor beads.⁶¹ The drug screening result showed that different tumor components had different responses to drugs. The vigorously proliferating tumor cells were more sensitive to PAC, while the viability of the fibroblasts in the shell layer was not severely affected by PAC. Further, the fibroblast shell had a certain protective effect on the viability of tumor cells in the core area. Afterward, the blood vessel-associated TME further significantly reduced the efficacy of drugs on tumors. By combining these two technologies, microfluidic droplet and on-chip microphysiological vessel platform, we report an integrated engineering system that enables high-fidelity fabrication of vessel-supported biomimetic tumor models for preclinical drug discovery while inheriting the high-throughput properties of droplet technology. It is believed that the reported system can facilitate the drug discovery procedure and contribute to antitumor therapy development.

EXPERIMENTAL SECTION

Microfluidic Device Fabrication. All of the microfluidic devices were designed with AutoCAD and fabricated with standard lithography technology (Figure S7). For the microfluidic droplet generation, the chip is based on a flow-focusing design (Figure 2A).^{62,63} The channel width and height were designed as 400 and 300 μm , respectively. The microfluidic vessel platform design was designed based on the previous literature.^{64,65} Generally, the device consists of a central gel-loading channel and flow channels on the left and right sides. The gel channel and flow channel were separated by a row of trapezoidal micropillars.⁶⁶ The distance between micropillars was 200 μm to provide sufficient surface tension to maintain infused gel solution within the central channel. The device was designed to be 350 μm in height and fabricated with soft lithography. An SU8-2150 (Kayaku Advanced Materials, Inc.) photoresist layer was spin-coated on a 3 in. silicon wafer, followed by UV exposure, post baking and development procedures. All of the fabrication processes were finished in the Center for Photonics and Nanoelectronics (CPN) at Lehigh University. Poly(dimethylsiloxane) (PDMS) (SYLGARD184, Dow Corning) presolution was poured on a solid SU8 mold and cured at 80 °C for 1 h. After being peeled off, the solid PDMS devices were plasma-bonded to glass substrates. All of the devices were autoclaved for sterilization prior to usage.

Computational Simulation. To get the optimized flow condition, finite element method (FEM) was combined with the laminar two-phase level set simulation in COMSOL Multiphysics for the study.⁶⁷ The motion of the continuous and dispersed phase interfaces is governed as described in eq 1

$$\rho \frac{\partial \mathbf{u}}{\partial t} + \rho(\mathbf{u} \cdot \nabla) \mathbf{u} = \nabla \cdot [-p\mathbf{I} + \mu(\nabla \mathbf{u} + (\nabla \mathbf{u})^T)] + \mathbf{F}_{\text{st}}$$

$$\nabla \cdot \mathbf{u} = 0$$

$$\frac{\partial \phi}{\partial t} + \mathbf{u} \cdot \nabla \phi = \gamma \nabla \cdot \left(-\phi(1-\phi) \frac{\nabla \phi}{|\nabla \phi|} + \varepsilon \nabla \phi \right) \quad (1)$$

For simplicity, we assumed the continuous phase as HFE7500 engineered fluid and the dispersed phase as water. In the above equation, \mathbf{u} is the velocity (m/s), ρ denotes the density of the fluid, t is time (s), p is pressure (Pa), μ is the dynamic viscosity ($\text{Pa}\cdot\text{s}$), and \mathbf{F}_{st} is the surface tension force (N/m^3). Then, the effective droplet diameter is calculated as described in eq 2

$$d_{\text{eff}} = 2 \sqrt{\frac{3}{4\pi} \int_{\Omega} (\phi > 0.5) \, d\Omega} \quad (2)$$

Cell Culture. Human umbilical vein endothelial cells (HUVECs) and normal human lung fibroblasts (NHLFs) were purchased from LONZA. HUVECs were cultured in an endothelial cell growth medium supplemented with EGM-2 SingleQuot kit supply and growth factors (LONZA). NHLFs were cultured in fibroblast growth basal medium supplemented with FGM-2 BulletKit (LONZA). HCT-116 cell line was purchased from ATCC. Dulbecco’s modified Eagle’s medium (DMEM, Life Technology) was supplemented with 10% fetal bovine serum (FBS, Invitrogen) and 1% antibiotic and antimycotic (ThermoFisher). All cells were cultured at 37 °C in a humidified 5% CO_2 environment.

Fabrication of Cell-Laden Hydrogel Beads with Microfluidic Droplet Technique. For microfluidic droplet generation, HFE7500 engineered fluid (3 M, Maplewood) was mixed with 2% of 008-FluoroSurfactant (RAN Biotech) as the continuous phase.⁶⁸ The cells were optimized to the desired concentration and resuspended in 5% w/v of GelMA solution supplemented with 0.5% w/v of lithium phenyl-2,4,6-trimethylbenzoylphosphinate (LAP) (CELLINK, Sweden) as the dispersed phase. Both the continuous and dispersed phases were loaded into the microfluidic droplet generation device with two syringe pumps. The generated cell-laden droplets were transferred out of the microfluidic droplet device and collected in a 35 mm Petri dish containing the culture medium. An ultraviolet light (UV) source (405 nm wavelength) was activated for 15 s for photo-cross-linking. The dish was placed on a shaker for half an hour to realize oil–water separation and release the beads into culture medium for the following culture.⁶⁹

On-Chip Vessel Network Construction and Vascularization of the Cell-Laden Beads. For vessel seeding, the HUVECs (8 million/mL) were resuspended in bovine fibrinogen (CAS-9001-32-5, Sigma) solution. After being mixed with thrombin (CAS-9002-044, Sigma-Aldrich), the cell-laden fibrin presolution was injected into the on-chip vessel network platform and cured for 20 min. Hydrostatic pressure was provided to promote vessel network formation (Figure 5A). The left flow channel was applied with a high liquid column (300 μL), and the right flow channel was applied with a low liquid column (100 μL), so that there was a hydrostatic pressure gradient across the central 3D fibrin hydrogel.⁷⁰ The preformed cell-laden beads were mixed in the HUVEC suspension for vascularization. After 3 days of culture, the on-chip vessel system could be constructed, within which the predeveloped cell-laden beads could be integrated. Red blood cells (RBCs) from healthy donors were collected and sent from the University of Maryland School of Medicine. The sample collection process has followed the approved institutional review board protocols with donors providing informed consent.

Immunofluorescence Staining and Imaging. The vessel sample was fixed with 4% of paraformaldehyde in PBS overnight at 4 °C and incubated with a 1:50 diluted Alexa Fluor 488 antihuman CD31 antibody (BioLegend) and 50 $\mu\text{g}/\text{mL}$ of DAPI overnight at 4 °C. The fibroblast-laden beads were fixed and incubated with a human α -smooth muscle actin (α -SMA) Alexa Fluor 594-conjugated antibody (R&D System).⁷¹ The HCT-116-laden beads were fixed and incubated with an antihuman CD326 (EpCAM) antibody (Bio-

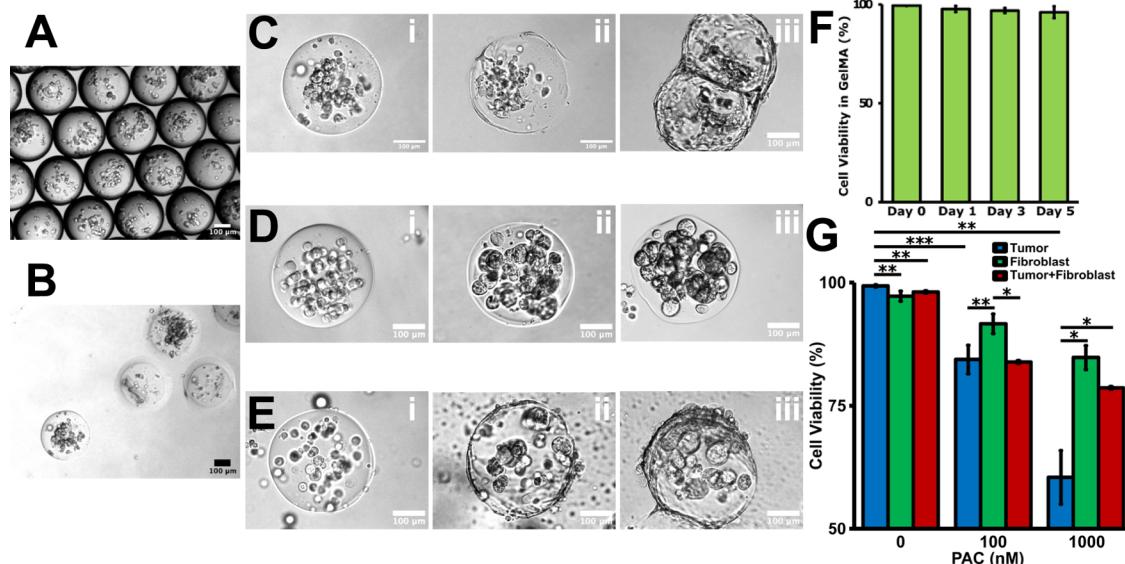


Figure 3. *In vitro* culture of the homotypic and heterotypic cell-laden microbeads. (A) Cell-laden droplet recollection from the microfluidic droplet generation device. (B) The cell-laden microbeads fall to the medium layer during oil/water separation process. (C) Fibroblasts gradually migrated to the bead surface from D0 (i) to D3 (ii) and form a confluent monolayer on D7 (iii). (D) HCT-116 tumor cells just proliferated *in situ* from D0 (i) to D3 (ii), and solid clusters formed inside the GelMA beads on D7 (iii). (E) Coculture of the fibroblast cells and tumor cells was encapsulated into the GelMA beads on D0 (i). The fibroblasts gradually migrated to the bead surface and the tumor cells proliferated *in situ* on D3 (ii). A core-shell tumor model formed on D7 (iii). (F) Viability characterization of the cell-laden GelMA beads during the *in vitro* culture. (G) Drug screening result on the homotypic and heterotypic beads. *P*-values were calculated using a two-sample *t*-test with respect to control. **p* < 0.05; ***p* < 0.01; ****p* < 0.001. The scale bar is 100 μ m.

egend).⁷² The vessel network sample was fixed and incubated with an Alexa Fluor 488 antimouse CD31 (PECAM) antibody (BioLegend).⁷³ All of the fluorescent images and confocal scanning were acquired with a Nikon C2+ laser scanning confocal microscope in the Health Research Hub Center, Lehigh University.

Drug Treatment and Cell Viability Assay. The viability of the cell-laden beads was verified at days 0, 1, 3, and 5 to demonstrate the GelMA biocompatibility. The live/dead staining solution was prepared in complete medium with a concentration of 5 and 10 μ M of calcine AM and propidium iodide (PI, Biotium), respectively, and incubated with the sample for 30 min at 37 °C prior to imaging. The viability was quantified from the fluorescence images and processed with the NIH ImageJ software. The therapeutic drug paclitaxel (PAC, LC Labs) was dissolved in DMSO at a 1 mM stock solution and then diluted into a concentration of 100 and 1000 nM with complete medium.⁷⁴ For the individual bead testing, the PAC solution was deposited to the GelMA beads on day 7. For the vessel-supported condition, PAC was introduced into the microfluidic device from one side channel and flowed through a central vessel network for 3 h on day 3. After 72 h, the viability was examined by the calcine AM (Corning) and propidium iodide (PI, Biotium).

Statistical Analyses. The calcine AM and PI staining fluorescent images of the samples were analyzed to quantify the viability after drug treatment. All of the samples were examined within the same view of the field. Each experimental group was repeated three times. In each experimental group, 20 samples were counted. The error bars of all of the figures represented the standard deviation. The quantified results were analyzed using an independent, two-tailed Student *t*-test, in which the *P*-value <0.05 was considered statistically significant.

RESULTS AND DISCUSSION

Microfluidic Droplet Generation On-Chip. The microfluidic droplet generation device has two inlets for the oil and aqueous phase loadings, respectively. Briefly, the aqueous phase and the oil phase with different flow rates joined at the flow junction area (Figure 2B). Since these two-phase solutions were immiscible with each other, under the action

of viscous shear stress (Figure 2, Biii), the aqueous phase dissociates into a dispersed phase (Figure 2, Biv).²⁵ Specifically, the surfactant-added fluorinated oils are selected to provide sufficient interfacial tension to avoid the coalescence of contacting droplets.²⁴ The size of the droplets depends on the relative flow rates of the oil and aqueous phases (Figure 2C). By fixing the oil flow rate and adjusting the aqueous phase flow rate, droplets of different sizes could be produced (Figure 2D). Here, the diameter of the droplets was set around 250 μ m. Meanwhile, the average cell load per droplet can be controlled by regulating the cell concentration in the original suspension (Figure 2E). To generate uniform cell-laden droplets, the flow rates of aqueous and oil phases were set as 480 and 3360 μ L/h, respectively. The concentration of the original cell suspension was set at 10 million/mL, and the number of cells per drop was about 65. In the end, the generated cell-laden microdroplets were collected in a 35 mm Petri dish and packed into stable arrays for potential gelation operation (Figure 2E).

Fabrication and Characterization of Homotypic and Heterotypic Cell-Laden Beads. GelMA was selected as a tumor-localized matrix scaffold for encapsulating cells into droplets due to its ease of engineering and good biocompatibility.^{75,76} Initially, the cells were resuspended in the GelMA presolution and encapsulated into individual microdroplets. After being collected in the Petri dish, the cell-laden droplets were cross-linked by the 25 mW/cm² UV source into the solid GelMA beads (Figure 3A). Then, a sufficient culture medium was added to the dish. Due to the interfacial tension, the solid beads were encapsulated in the oil layer, floating on top of the medium. While being shaken slightly, the beads were detached from the oil layer and gradually settled into the underlying medium. In this oil–water separation process, the superficial oil shell on the surface of the beads was removed, leaving only

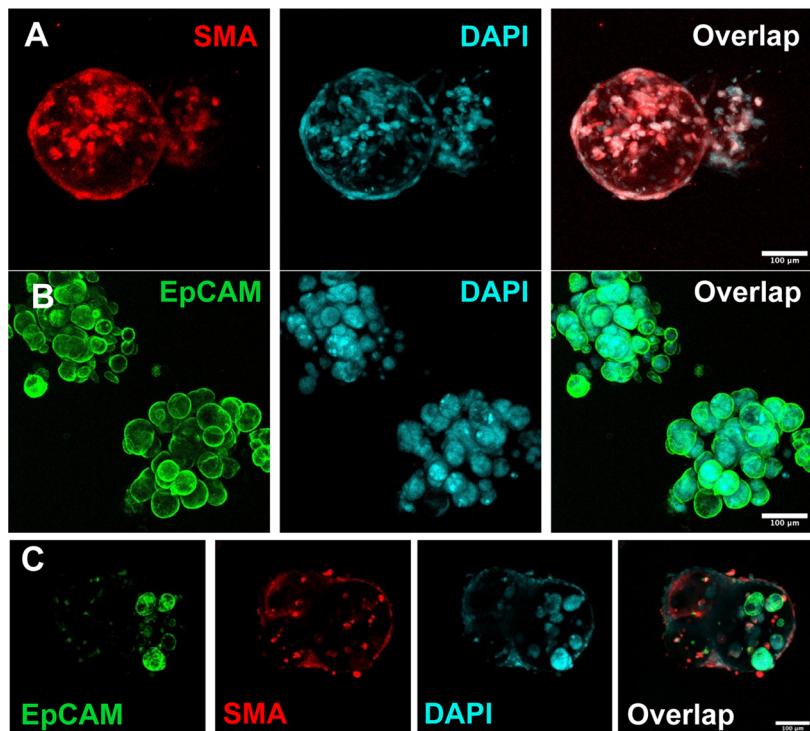


Figure 4. Immunostaining of the homotypic and heterotypic cell-laden beads. (A) The fibroblast-laden beads were characterized by SMA (red), and the nuclei were characterized by DAPI (cyan). (B) The tumor-laden beads were characterized by EPCAM (green), and the nuclei were characterized by DAPI (cyan). (C) The heterotypic beads were characterized by SMA (red) and EPCAM (green), and the nuclei were characterized by DAPI (cyan).

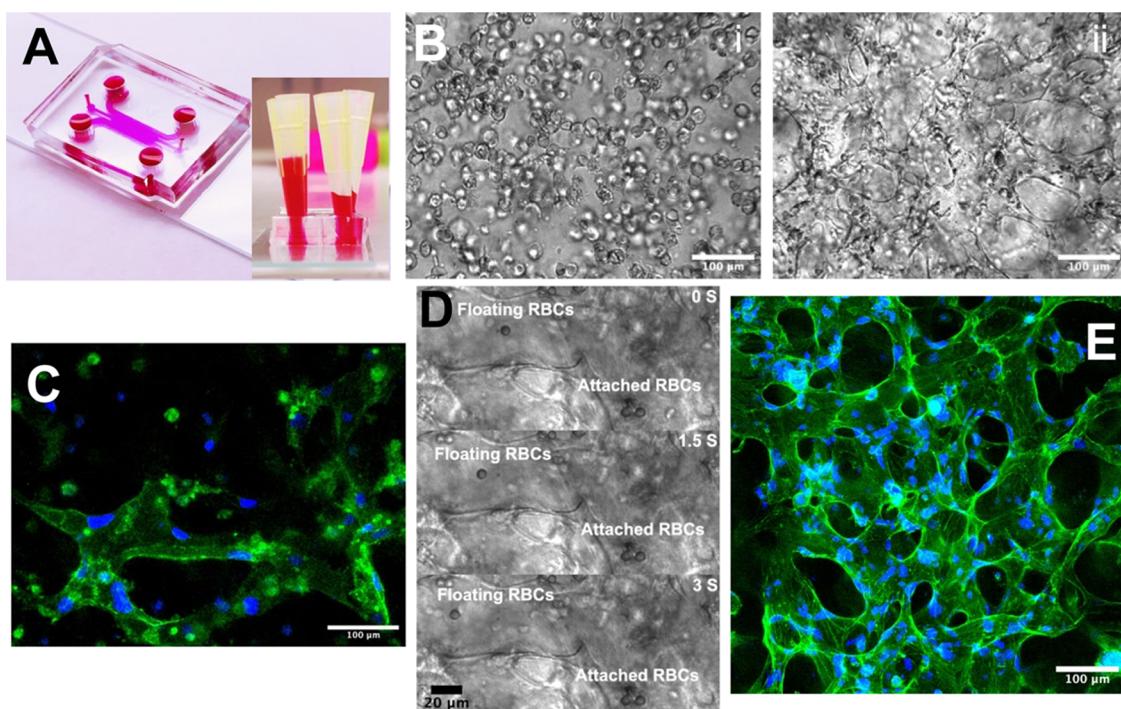


Figure 5. Vessel network-on-a-chip. (A) The microfluidic vessel network-on-a-chip platform. (B) The suspended HUVECs began the self-assembly on day 0 (i), and the perfusable vessel network formed on day 3 (ii). (C) Immunostaining of the vessel tube. CD31 was conjugated with FITC, and cell nucleus was stained with DAPI (blue). (D) RBCs were flowed through the vessel network. (E) Confocal scanning of the 3D vessel network. The F-actin of the vessel network was stained by FITC, and cell nucleus was stained with DAPI (blue).

the beads immersed in the medium for long-term cell culture (**Figure 3B**). For the homotypic cell-laden bead fabrication, the fibroblasts and HCT-116 tumor cells were loaded into the

GelMA beads separately. In a week-long *in vitro* culture, the cell viability was well maintained (**Figure 3F**), which is the basis of the proposed biofabrication system. The cells of two

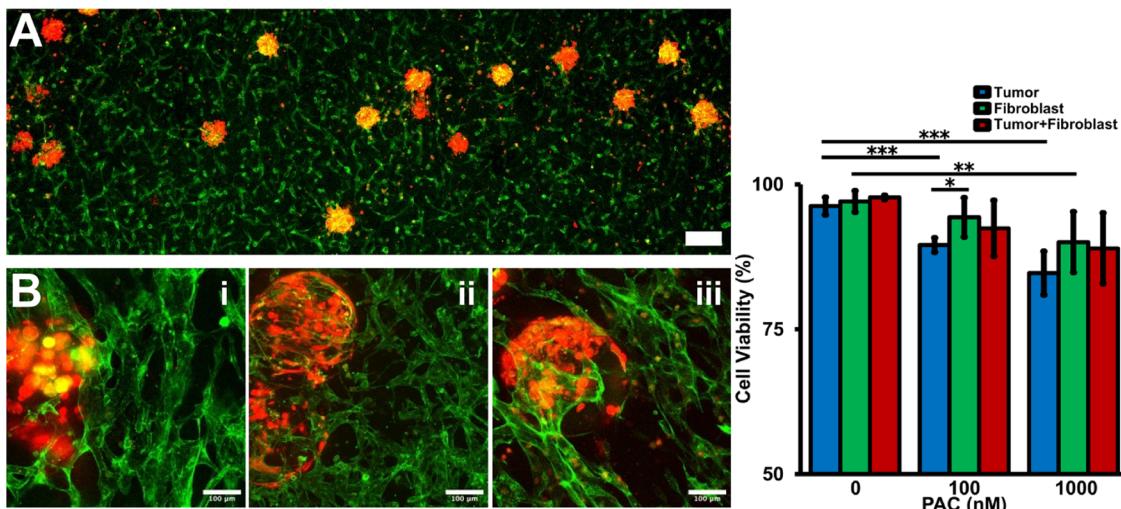


Figure 6. Vessel-supported tumor models for preclinical drug screening. (A) Large-scale integration of the cell-laden beads into the on-chip vessel network platform. (B) The homotypic beads, tumor-laden beads (i), and fibroblast-laden beads (ii) were integrated into the vessel network separately. The heterotypic tumor bead was integrated into the vessel network (iii). (C) The drug screening result on the vessel-supported tumor models. *P*-values were calculated using a two-sample *t*-test with respect to control. **p* < 0.05; ***p* < 0.01; ****p* < 0.001.

different types exhibited different growth behaviors in the GelMA matrix (Figure 3C–E). For the fibroblast-laden beads, the cells migrated to the bead surface and formed a confluent layer. This fibroblast monolayer appeared to have contractile properties as the contour of the bead was slightly compressed. Besides, the cellular layer also exhibited adhesion properties, as the two adjacent beads could be easily bonded together (Figure 3Ciii). α SMA was stained on the selected sample to identify that the fibroblasts were localized on the bead surface (Figure 4A). It is worth mentioning that the morphology of the fibroblasts was related to GelMA composition (Figure S3). In soft (3% w/v) GelMA, the fibroblasts self-assembled into spheroids after overnight culture. In stiff (8% w/v) GelMA, the fibroblasts grew slowly and self-assembled into a network after 1 week of culture.

For the tumor-laden beads, the cells did not show any migration behavior but just proliferated *in situ* into the aggregates. Likewise, EPCAMs were stained on the selected samples to outline the cellular aggregate morphology (Figure 4B). As for the heterotypic cell-laden beads, the fibroblasts were mixed with tumor cells mixed in a 1:3 ratio to better facilitate tumor–vessel interactions.³⁸ Likewise, most of the fibroblasts egressed to the bead surface, with a small amount of them remaining in the core, while the tumor cells proliferated *in situ*. The immunostaining confirmed that a core–shell compartmentalization was formed by the cellular self-assembly, with the fibroblast sheet covering the surface and the tumor clusters in the center (Figure 4C). This separated structure of the two cell types also facilitated the calculation of the viability of the two cells in subsequent drug experiments. If cultured longer, the surface fibroblast sheets could bring adjacent tumor beads together to form macroscopic tumor tissues (Figure S2F). Hence, the tumor–ECM and tumor–stromal interactions were remodeled *in vitro* on the individual microbeads. Beads of other cell compositions and sizes can also be precisely mass-produced as needed based on the capability of the microfluidic droplet technique.⁷⁷

On-Chip Vessel System Construction and Engineering Tumor–TME Interaction. After fabrication and development, the cell-laden beads were integrated into the on-chip

vessel network platform for reorganizing the tumor–TME interaction.⁷⁸ First, the vessel network was formed on the on-chip vessel platform. The high-concentration HUVECs were seeded in the fibrinogen hydrogel. Then, cell culture medium columns of different heights were set at the left and right flow channels of the device to generate interstitial flow (IF). The IF conducted hydraulic pressure across the central gel zone and activated the HUVECs to form the capillary vessel network^{79,80} (Figure 5A). The medium columns were refreshed every 24 h so that there was always hydraulic pressure across the gel zone. On day 0, vacuoles were observed in every single HUVEC, which is often considered a hallmark of vascular tubulogenesis initiation⁸¹ (Figure 5Bi). On day 4, the self-organized vessel network was fully developed on-chip (Figure 5Bii). Biomarker immunostaining shows a strong CD31 signal, indicating tight cell-to-cell junctions (Figure 5C). To better characterize the 3D vessel morphology, filamentous actin (F-actin) of the selected sample was stained and confocal scanned (Figure 5D). As one of the basic properties of blood vessels, the vascular perfusability was demonstrated by flowing suspended RBCs (Supplementary Videos 1 and 2) from the side endothelialized channel through the artificial anastomosis zone and into the central network area (Figure S2). Figure 5E shows a floating RBC flowing through the vessel tube, while some RBCs got attached to the vessel surface. Therefore, a functional vessel network system was established on-chip.

To engineer the tumor–TME interaction, the predeveloped homotypic and heterotypic cell-laden beads were reseeded into the on-chip vessel network platform separately. The beads were cultured 1 week prior to being incorporated into the vessel system. The beads were settled to the bottom of a centrifuge tube by centrifugation and then resuspended into the fibrinogen solution. After being mixed with the thrombin at an appropriate concentration, the solution was injected into the vessel device. On day 3, the functional vessel network formed around the microbeads as well (Figure 6B). The blood vessel with the fibroblasts was slightly smaller in diameter, and the blood vessel with pure tumors was slightly thicker (Figure S6). This may imply that cancer-associated fibroblasts (CAFs)

play an important role that cannot be ignored in real tumor vascularization cases.^{82,83}

In Vitro Antitumor Treatment Testing. To validate the drug resistance brought by the tumor–stromal and tumor–TME interactions, the anticancer drug treatment was administrated on the individual tumor bead model and vessel-supported tumor model, respectively. Paclitaxel is widely employed as an anticancer drug, and the mechanism is to stop the mitotic cells.⁸⁴ By directly administering the beads under static conditions, a concentration dependence of drug efficacy was observed on different cellular components (Figure 3G). The tumor beads were more sensitive to drug than the fibroblasts (Figure S4). The higher the drug concentration, the lower the viability of tumor beads. Although the highest concentration of PAC was the most lethal for tumor cells in 3D tumor beads, the fibroblast beads still maintained about 80% viability. Literatures report that PAC may affect some functions of the stromal cells without significant cell loss.^{85,86} For the heterotypic tumor beads, the viability of the inner tumor clusters was largely preserved compared with the homotypic tumor beads. A possible hypothesis is that the fibroblasts in the shell absorbed part of the drug, reducing the real concentration of the drug that actually reached the tumor cells in the core area.⁸⁷ As for the drug testing on the vessel-supported models, PAC was loaded into the side endothelialized channel and flowed through the artificial anastomosis and went into the capillary vessel networks (Figure S5). Then, the drug diffused across the capillary vessel wall and permeated into the surrounding matrix. The live/dead staining showed that the tumor cell viability was further preserved even at the most lethal drug concentration groups (Figure 6C). In conclusion, the TME composed of different cellular components could bring a non-negligible mitigating effect on the actual killing effect of drugs, and the real drug efficiency decreased.⁸⁸ It is believed that the involvement of heterogeneous TME can import both pharmacodynamic (cell sensitivity-related) resistance and pharmacokinetics (drug delivery-related).⁸⁹ For example, the blood vessels might have transported some of the drugs away before they diffused laterally into the surrounding tissue.⁹⁰ As a result, the drug concentration might not be the same for all of the cells in the vessel network-based TME. In practical drug particle designs, the drug delivery efficiency to overcome the barrier of solid tumor tissues needs extra attention. It may be necessary to define a quantitative model to estimate resistance from different tumor component sources.

CONCLUSIONS

In summary, we report an integrated engineering system to fabricate a vessel-supported heterotypic tumor model for preclinical drug discovery by coupling microfluidic droplet and microfluidic vessel-on-a-chip techniques. After loading into the GelMA microbeads, the tumor cells and the fibroblasts reassembled into a core–shell spatial structure to mimic the tumor–stromal interaction in the ECM. Inspired by the concept of bottom-up engineering, the micro-cell-laden beads were applied as building blocks and consolidated into an on-chip vessel network platform to mimic the tumor–TME interaction. Further, the antitumor drug paclitaxel was tested on the individual tumor model and vessel-supported heterotypic tumor model. The results showed that the vasculature-associated TME had a significantly attenuating effect on the effectiveness of chemotherapy. The reported

system could be beneficial for large-scale preclinical antitumor drug discovery.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsmi.2c23305>.

(Figure S1) Cell viability test in GelMA; (Figure S2) cell-laden bead generation by microfluidic droplet technique; (Figure S3) fibroblast morphologies in the different GelMA constructs; (Figure S4) live/dead staining of the cell-laden beads after the drug testing; (Figure S5) confocal staining of the functional on-chip vessel network; (Figure S6) confocal staining of the vessel-supported tumor-laden beads; and (Figure S7) snapshot of the microfluidic device designs in AutoCAD (PDF)

The suspended RBCs were perfused through the on-chip vessel network (Video 1) (AVI)

Bright-field scanning of the fibroblast-laden beads (Video 2) (AVI)

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Author Contributions

Y.L. conceived and supervised the study. Y.W., Y.Z., Y.Z., and Y. L. designed and performed the experiments. Y.W. collected and analyzed the data. Y.Z. and Y.Z. assisted with data analysis. K.I. assisted with simulation. Y.W. wrote the manuscript. Y.W., Y.Z., Y.Z., K.I., and Y.L. revised the manuscript. All authors discussed the results and approved the submission.

Notes

The authors declare no competing financial interest.

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