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Accelerated maturation of branched organoids confined in collagen droplets

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Droplet-based organoid culture offers several advantages over conventional bulk organoid culture, such as improved yield, reproducibility, and throughput. However, organoids grown in droplets typically display only a spherical geometry and lack the intricate structural complexity found in native tissue. By incorporating singularised pancreatic ductal adenocarcinoma cells into collagen droplets, we achieve the growth of branched structures, indicating a more complex interaction with the surrounding hydrogel. A comparison of organoid growth in droplets of different diameters showed that while geometrical confinement improves organoid homogeneity, it also impairs the formation of more complex organoid morphologies. Thus, only in 750 μm diameter collagen droplets did we achieve the consistent growth of highly branched structures with a morphology closely resembling the structural complexity achieved in traditional bulk organoid culture. Moreover, our analysis of organoid morphology and transcriptomic data suggests an accelerated maturation of organoids cultured in collagen droplets, highlighting a shift in developmental timing compared to traditional systems.

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Introduction

Progenitor cells isolated from human adult tissue or stem cells are able to navigate a complex spatiotemporal differentiation pattern that gives rise to organoid structures that capture important properties of native tissue.^{1–4} Many developmental processes result in branched structures as found in several distinct tissues, including the lungs,^{5,6} kidneys,^{7,8} mammary

glands^{9,10} and pancreas.^{11,12} This process of branching morphogenesis also plays a relevant role in pathological conditions such as tumour progression in pancreatic ductal adenocarcinoma (PDAC). Indeed, PDAC organoid growth successfully replicates the various developmental processes and maturation steps characteristic of PDAC tumours *in vivo*.¹³ Furthermore, the ability of branched PDAC organoids to accurately capture intratumoural heterogeneity and drug response patterns highlights their potential for drug screening and personalised medicine.¹⁴ Beyond scalability and parallelisation, drug-screening applications rely heavily on the ability to steer and control such maturation steps.¹⁵ The use of droplet microfluidics^{16,17} to encapsulate cells into small hydrogel droplets or core-shell structures has enhanced the throughput^{18,19} and scalability²⁰ of organoid culture systems. Most of those recent applications focused on establishing spheroid culture systems from cell culture aggregates in matrigel without higher-order structure formation processes.^{21–27} Single-cell-derived organoids from progenitor cells cultured in hydrogel droplets displaying distinct developmental phases remain to be implemented. Confining organoid culture to small droplets alters the mechanical properties of the surrounding hydrogel by the modified boundary condition. Its impact on the developmental processes during organoid growth remains unexplored. Here, we demonstrate that branching organoid structures can be grown in collagen droplets, recapturing faithfully the structures found in bulk assays and even *in vivo*. To this end, we incorporated

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singularised murine pancreatic ductal adenocarcinoma cells into the collagen droplets and tightly controlled the temperature gradient in the microfluidic device, ensuring uniform collagen polymerisation within the hydrogel droplets. While previous studies established the use of 100–300 μm sized collagen droplets for spheroid culture,^{28–30} our results reveal that increasing droplet size to 750 μm enables complex branching morphogenesis and lumen formation, thereby supporting the growth of single cell derived PDAC organoids. We showed that geometrical confinement accelerates organoid maturation, further highlighting the potential of this approach for high throughput and time-sensitive experiments.

Materials and methods

Provenance of cell lines

Primary tumour cells (PDAC) were gifted by the laboratories of Prof. Maximilian Reichert and Prof. Dieter Saur (Technical University of Munich). Cells were authenticated by genotyping. The Phoenix ECO helper-free retrovirus producer cell line was a gift by the laboratory of Prof. Carsten Grashoff (Universität Münster). All cells were tested and confirmed negative for mycoplasma contamination every 6 months.

2D cell culture

Pancreatic ductal adenocarcinoma cells (PDAC) were cultured in high glucose DMEM medium with 1% penicillin/streptomycin (Thermo Fisher Scientific) and 10% fetal bovine serum (FBS) (Sigma). Cells were grown at 37 °C with 5% of CO₂ and 80% humidity. Media was exchanged every 48 hours, and cells were passaged by trypsinisation upon 80–85% of confluence. For 2D analysis, cell culture with 80–85% confluence was used. Phoenix ECO helper-free retrovirus producer cell line (Phoenix ECO cells) was cultured with high glucose DMEM (Thermo Fisher Scientific) supplemented with 10% FBS and 1% penicillin/streptomycin (Thermo Fisher Scientific) in an incubator with 80% humidity and a 5% CO₂ atmosphere, at 37 °C.

Generation of fluorescently labelled cell lines

As already described previously,³¹ for the stable expression of fluorescent markers, retroviral transduction as well as CRISPR/Cas9 transfections were performed with the same primary tumour cells. Retroviral transfection of PDAC cells was implemented during an 8 day protocol using a retroviral plasmid for mCherry-LifeAct (Addgene)³² or Teal-H2B (gifted by the laboratory of Carsten Grashoff). At day 1, Phoenix ECO cells were seeded in a 175 cm² flasks. Having reached a confluence of 50 to 60% (day 2), Phoenix ECO cells were transfected using Mirus TransIT-X2 Dynamic Delivery System (VWR MIRUMIR-6000) as described in the manufacturer's protocol. Media exchange was performed after 24 hours (day 3) of incubation. Simultaneously, PDAC cells were seeded in a 75 cm² flask. Virus was harvested after 48 hours (day 4) from Phoenix ECO cells and sterile filtered (0.45 μm pore size). Supplemented with 7.5

$\mu\text{g ml}^{-1}$ polybrene (Sigma-Aldrich) the virus conditioned media was added to the PDAC cells and incubated for 24 hours. This was repeated on day 5. After 48 hours of incubation with virus conditioned media, media was exchanged for PDAC cells with fresh high glucose DMEM (Thermo Fisher Scientific) supplemented with 10% FBS (Sigma F7524) and 1% penicillin/streptomycin (Thermo Fisher Scientific). After 72 hours cells were passaged as described above. Selection of fluorescently labeled cells was implemented by antibiotic resistance, using puromycin dihydrochloride (Gibco) for mCherry-LifeAct, and Geneticin selective antibiotic (G418 sulfate) (Gibco) for Teal-H2B. The FACS sort was carried out using BD Aria Fusion. A CRISPR/Cas9 system was used to endogenously label E-cadherin in the used primary tumour cells.

Organoids preparation in bulk collagen

For organoids growing in bulk collagen, we used a previously published protocol.^{31,33} Briefly, PDAC cells (9591) with confluence 80–85% were detached using trypsin and prepared to a final concentration of 500 cells per ml of DMEM media. Next, the cell suspension was mixed gently with 1.3 mg ml⁻¹ or 3 mg ml⁻¹ collagen type I solution (rat tail from Corning, NaOH to stabilise pH and neutralizing buffer). The solution was incubated for 1 h at 37 °C to polymerise the collagen fibres, and DMEM media was added after detaching the gels from the dish walls. Media was changed first after 72 hours and then every 48 hours. All conditions were performed simultaneously in 4 technical replicates $n = 4$.

Chip design

Two different chips were used to achieve different droplet diameters for droplet production. To generate large droplets of 750 μm diameter, a microfluidic chip with a channel height of 600 μm was used, whereas for the production of the smaller sized droplets with a diameter of 370 μm , a chip with 200 μm channel height was used. For both chips, the same underlying CorelCAD design was employed (Fig. S1a).

SU8 deposition

To achieve wafers coated with 600 μm thick SU8, we mixed 47.41 g of SU8-3005 (Kayaku Advanced Materials, United States) and 4.66 g SU8 2000 Thinner (Kayaku Advanced Materials, United States). The silicon wafers (Nano Quarz Wafer, Germany) were placed on a heating plate that had been calibrated to ensure a horizontal orientation of the hot plate's surface. After 7 ml of SU8-thinner mixture had been added to the wafers, the heating plate was turned on and heated to 95 °C. The thinner evaporated overnight, and the SU8-coated wafer was used for photolithography.

Chip fabrication

To crosslink the exposed pattern, the SU8-coated wafers were exposed to UV light at an exposure energy of 720 mJ cm⁻² for 200 μm thick wafers and to 311 mJ cm⁻² for 600 μm thick



SU8 wafers. After UV exposure, the wafers were baked at 65 °C for 5 minutes and at 95 °C for 13 minutes (200 µm thickness) and 33 minutes (600 µm thickness) during the post-exposure bake. The wafers were immersed in SU8 developing solution (PGMEA, Sigma-Aldrich) to develop the structures for 15 minutes (200 µm thickness) and 35 minutes (600 µm thickness) and washed with isopropanol. After developing the structures, the samples were baked at 150 °C for 5 minutes during the post-exposure bake. After master fabrication, PDMS was mixed with curing agent at a ratio of 10:1 (Sylgard 182 Silicone Elastomer, Dow Corning, United States) and poured on the master. To remove air bubbles, the master was then put under vacuum for 15 minutes. Then, the PDMS was cured by overnight incubation in the oven at 70 °C. The next day, the PDMS replica was peeled off the mold. The channel inlets and outlets were generated by using a biopsy puncher of 1 mm diameter. The PDMS was bonded to a glass coverslip after treatment with oxygen plasma. After bonding, the microfluidic chip was incubated at 70 °C for at least 1 hour. Eventually, the microfluidic chips were treated with commercially available Aquapel for 5 minutes at room temperature to obtain a hydrophobic channel surface. The chips were then air dried and incubated at 70 °C for 15 minutes.

Characterisation of channel height

To characterise the height of the SU8 layer used to form the 600 µm channels, we sectioned the PDMS mold cast from the master wafer into thin slices (Fig. S2a). We then measured the channel height at multiple positions for each mold replica on the wafer. To ensure uniform droplet size, we compared the average channel height and its variability across replicas (Fig. S2b). Replicas 2 and 3 were selected for experiments as their average height was closest to 600 µm and they displayed minimal variance in channel height.

Cell preparation for organoid seeding on the microfluidic chip

The collagen solution for organoid seeding on the microfluidic chip was prepared as previously described for bulk organoid culture.^{9,33} To achieve the incorporation of single cells into collagen droplets the PDAC cells were mechanically separated after trypsin treatment by aspiration with a syringe needle (28G, Braun). The remaining doublet cells were removed by using a cell strainer (40 µm, Corning). To generate small droplets with single incorporated cells the PDAC cells were diluted to a final concentration of 30 000 cells per ml in 3 mg ml⁻¹ collagen solution, respectively to obtain single incorporated cells in large droplets the cells were diluted to a final concentration of 5000 cells per ml.

Droplet production for organoid culture on the microfluidic chip

For droplet production, fluorinated oil Novec 7500 (Iolitec, Germany) with 3% (w/w) surfactant RAN008 (RAN Biotechnologies, United States) was used to generate the small

droplets. For the production of large droplets, we used fluorinated oil with 5% surfactant. We loaded the oil with surfactant into 5 ml BD plastic syringes to initiate droplet production. The collagen solution containing the cells was loaded into either a 1 ml BD glass syringe or a 5 ml BD plastic syringe, depending on the required droplet amount. The syringes were connected to the microfluidic chip using fluid dispenser tips (PT-050-27, Drifton, Denmark) and polyethylene micro tubing (PE/2, Scientific Commodities, United States). The syringes were operated using programmable or computer-operated syringe pumps (KDE Technologies, Harvard Pumps or Landgraf HLL). Typical operating speeds for droplet production were 2000 µl h⁻¹ and 1000 µl h⁻¹ for the oil and collagen to generate small droplets or 1000 µl h⁻¹ and 1000 µl h⁻¹ for the oil and collagen to generate large droplets. After droplet production, the droplets were collected in a 2 ml Eppendorf tube at 37 °C to initiate collagen polymerisation immediately. After droplet production, the Eppendorf tubes were transferred to an incubator and kept at 37 °C with 5% CO₂ for 1 hour. After collagen polymerisation, the fluorinated oil with surfactant was removed, and the droplets were washed three times with only fluorinated oil. After the last washing step, the fluorinated oil was removed, and fluorinated oil with 4% PFO (v/v) was added to the droplets to break the emulsion. Cell culture medium was added on top of the oil phase and the droplets were left for 5 minutes to transition into the aqueous phase. The droplets were centrifuged using a tabletop centrifuge for 5 minutes to improve droplet recovery. After centrifugation, the pellet formed by the droplets at the interface of the oil and cell culture medium was aspirated with a pipette, and the droplets were transferred into the cell culture medium. The droplets were distributed in 6 well plates. For microscopy imaging, the droplets were transferred to achieve a final concentration of 750 droplets per well, whereas for RNA sample collection, the desired final concentration was 1500 droplets per well.

Fluorescent collagen preparation

Collagen was fluorescently labeled as previously described.⁹ Briefly, collagen was dialyzed at 4 °C to reach pH 7. Subsequently, collagen was conjugated with Atto 488 by overnight incubation at 4 °C. To remove any non-bound dye, dialysis was performed for 8 hours followed by an overnight dialysis step with acid to avoid any premature polymerisation. The fluorescently labeled collagen was stored at 4 °C until further usage.

SiRDNA staining

For nuclei visualisation, PDAC organoids were incubated with SiRDNA (Spirochrome SPY650-DNA SC501) for a minimum of 1 h before imaging at 1–2 µg ml⁻¹.

Microscope analysis

Samples were analysed using scanning confocal microscope Leica Sp5 (PL FLUOTAR 10×/0.30 and HC PL APO 20×/0.70),

Leica Mica (PL FLUOTAR 10 \times /0.32) and epifluorescent microscope Leica Thunder (PL FLUOTAR 10 \times /0.32).

Characterisation of the droplet diameters

Collagen droplets were segmented using Cell Pose 2.0 running on Google Colab.³⁴ The diameter was extracted from the binary images using the scikit image package.

Measuring cellular incorporation frequency into droplets

After droplet generation, droplets in 6 well plates were screened for the amount of encapsulated cells.

Counting the amount of spheroids and branched structures present among organoid phenotypes

Four days after droplet generation, droplets in 6 well plates were screened for the number of structures that developed into spheroids and for the number of structures that displayed an invasive phenotype with the ability to form branches.

Droplet and organoid segmentation with feature extraction

Binary masks of 2D projected images of collagen droplets containing PDAC organoids and bulk-grown organoids were analyzed using the selection tool in ImageJ (ver. 1.54f).³⁵ Python and the scikit-image package were used to analyze morphological features. The compactness of the droplets was defined as $4\pi A/P^2$. Therefore, the projected 2D area, A , and the perimeter P were extracted from the segmented droplets.

Preparation of RNA samples for mRNA sequencing

2D PDAC cells were cultivated until 80–85% of confluence, detached, and collected by spinning down (300 g) to obtain cell pellets. Organoids on days 4, 7, 9, and 13 were subjected to collagenase treatment (2.6 mg ml⁻¹ for 1.3 mg ml⁻¹ collagen and 6 mg ml⁻¹ for 3 mg ml⁻¹ collagen) to remove the organoids from the collagen gel. Next, the solution with organoids was spun down (1200 rpm), and the pellet was gently washed with PBS. Total RNA was extracted from the organoids and 2D samples using RNeasy Mini Kit (Qiagen, Cat. No. 74104) according to the manufacturer's instructions. All steps were performed in RNase free environment. Next, RNA quality was determined using Agilent RNA 6000 Nano Assay. Library preparation was performed on samples with RIN \geq 7 and concentration \geq 30 ng μ l⁻¹. mRNA sequencing was performed by BMKGENE company using Illumina Novaseq X. 6GB of raw data was obtained for each sample. Data was next analyzed using Galaxy Europe server to get normalised read counts.³⁶

Differential gene expression (DEG) analysis

We used the DESeq2 R package from iDEP for differential gene expression analysis.³⁷

Statistical analysis

Experiments were performed two to three times independently from each other with similar outcomes. For each set of experiments, at least 5 replicates were measured. 95% confidence intervals were computed *via* bootstrapping with 1000 bootstrap iterations.

Results and discussion

Generation of collagen droplets using droplet-based microfluidic chips

To generate collagen droplets, we used PDMS microfluidic chips with two inlets: one for the collagen solution mixed with singularised PDAC cells and another one for the fluorinated oil with surfactant (Fig. 1a and S1a). Collagen droplets were generated at the microfluidic chip's cross-section and collected in a heated reservoir at 37 °C. To achieve full collagen polymerisation, the droplets were maintained in emulsion at 37 °C for 1 hour (Fig. 1a). Subsequently, the droplets were recovered from the emulsion and transferred to cell culture medium. The cells were cultured in the collagen droplets for up to a week and developed into branched organoids (Fig. 1a). The microfluidic chip was operated in a cold room to prevent premature collagen polymerisation before droplet formation. We performed experiments to assess the impact of the microfluidic setup and the collecting reservoir temperature on collagen polymerisation (Fig. 1b). Our findings indicate that polymerisation at 37 °C ensures evenly polymerised hydrogel (Fig. 1c–e). Moreover, it is crucial to maintain the temperature of the collagen at 4 °C throughout the seeding process to prevent premature collagen polymerisation (Fig. 1c and d). The collagen polymerises homogeneously only under such tightly controlled temperature gradients (Fig. 1e). To investigate the impact of droplet size on organoid growth behaviour, we fabricated microfluidic devices of two different heights, producing collagen droplets of 373 ± 21 μ m (small droplets, mean \pm s.d.) and 756 ± 30 μ m (large droplets, mean \pm s.d.) in diameter (Fig. S1b). The generation of large droplets was achieved by fabricating a microfluidic chip with 600 μ m high channels. To obtain this channel height 600 μ m thick SU-8 photoresist was deposited onto silicon wafers (Materials and methods). While previous studies also reported the generation of large droplets with diameters ranging from 600–1000 μ m using extrusion dripping techniques,^{38,39} this microfluidic approach offers a reproducible and accessible method for controlled droplet formation. The production rates were 10 Hz for the small droplets and 1 Hz for large droplets. A cell in a large droplet experienced a microenvironment with 8 times the collagen volume compared to a cell in a small droplet. The collagen solution containing the singularised PDAC cells was well-mixed to ensure even cell distribution in the seeding volume. H2B-Teal expressing PDAC cells were encapsulated in hydrogel spheres following a Poisson distribution during droplet formation. Consequently, 37% of large droplets were empty, 36% contained a single cell, and 27% contained two or more cells. For small droplets, 33% were empty, 33% contained



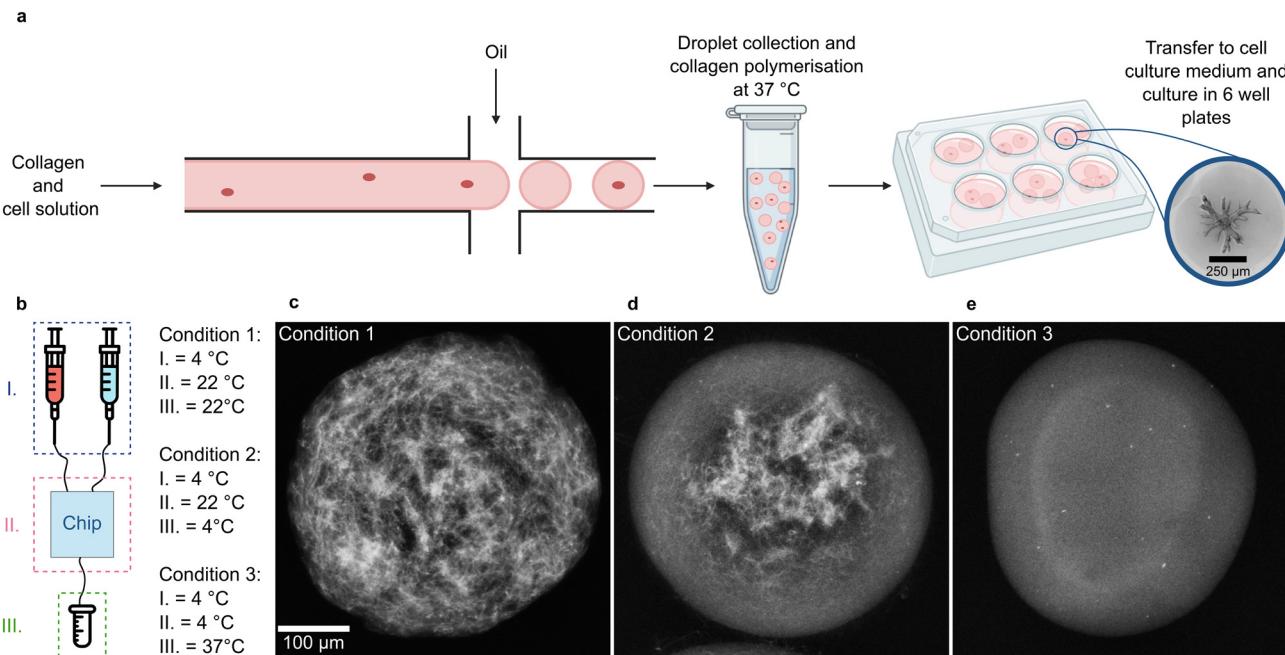


Fig. 1 Microfluidic setup for organoid production. (a) Singularised cells are encapsulated in collagen droplets using droplet-based microfluidics. After breaking the emulsion, the droplets are transferred to cell culture medium, and the organoids are cultured for 8 days. In large droplets, the seeded cells develop into branched organoids. (b) The microfluidic chip was operated with three different temperature gradients throughout the pipeline to establish optimal collagen polymerisation conditions. (c) At room temperature, the collagen droplets polymerise heterogeneously because of collagen fibre aggregation ($n = 2$ independent experiments). (d) The microfluidic chip is operated at room temperature, and the reservoir is cooled during droplet collection. Afterward, polymerisation is induced at 37 °C. The collagen polymerised prematurely as the temperature control was disrupted during droplet production ($n = 2$ independent experiments). (e) Maintaining the whole setup at 4 °C and the collecting reservoir at 37 °C allowed for homogeneous collagen polymerisation throughout the droplet ($n = 2$ independent experiments).

one cell, and 34% contained two or more cells (Fig. S1c). We determined the proportion of droplets containing at least one developed three-dimensional structure on day 4 to quantify organoid growth efficiency in the hydrogel spheres. For large droplets, containing on average 1.1 cells per droplet, $47 \pm 5\%$ (mean \pm s.d.) of droplets contained at least one organoid on day 4. Small droplets, containing on average 0.8 cells per droplet, showed a slightly lower efficiency, with $40 \pm 8\%$ (mean \pm s.d.) of droplets containing at least one organoid on day 4 (Fig. S3). Thus, droplet-based organoid culture yielded approximately one organoid for every second seeded cell, which represents a significantly higher efficiency than compared to bulk, where extreme limiting dilution analysis indicates that only every fifth cell forms an organoid.¹³

Characterisation of organoid growth in small and large collagen droplets

Single PDAC cells, seeded in collagen gels with a volume of approximately 1 ml, develop into mature structures with dimensions reaching several millimeters. By day 5, PDAC organoids typically measure 542 ± 155 μm (mean \pm s.d.) along their major axis.^{9,13} To investigate the impact of volume confinement, we analyzed organoid development in small (≈ 370 μm diameter, 27 nl) and large (≈ 750 μm diameter, 221 nl) collagen droplets. Organoids in small droplets had an increased tendency to grow in a spherical shape. In contrast,

organoids in large droplets tended to evolve into elongated structures with branched morphologies (Fig. 2a). We analyzed segmented 2D projections of the organoids and computed the shape irregularity index over time by comparing the perimeter of the organoid's shape to that of a circle with the same area. We found that organoids grown in small droplets displayed a lower shape irregularity index after 6 days in culture compared to organoids grown in large droplets, supporting the hypothesis that organoids in small droplets rather grew as spheroids (Fig. S4). To evaluate the efficiency of branching morphogenesis, we screened large and small droplets four days after seeding and quantified the proportion of spheroids *versus* branched structures. Given that the first 5 days of bulk organoid culture are characterised as the onset phase, we distinguished between spheroid structures and invasive phenotypes.¹³ Spheroids were defined as structures with a major axis a inferior to 60 μm, a minor axis b greater than $a/2$, or fewer than 2 branches of at least 30 μm in length. Structures that did not meet these criteria were classified as branched, displaying at least 2 branches or an invasive phenotype with the potential for further branching in later developmental stages. Our analysis revealed that in small droplets, only $44 \pm 5\%$ (mean \pm s.d.) of the structures were classified as branched structures, whereas in large droplets, $59 \pm 5\%$ (mean \pm s.d.) did fall under this classification (Fig. 2b). The major axis of organoids in large droplets increased more rapidly than in small droplets



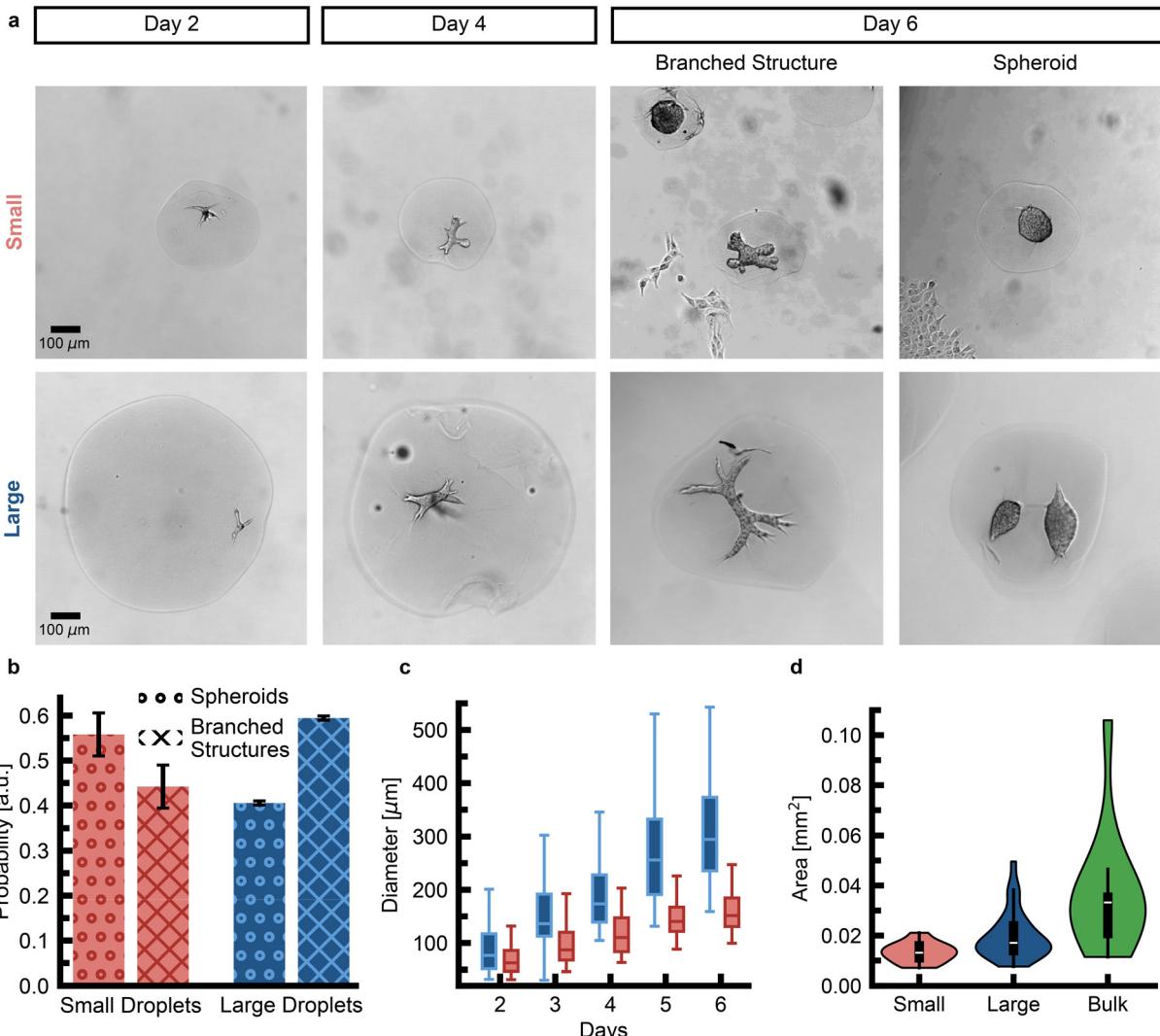


Fig. 2 Characterisation of organoid morphology in small versus large droplets. (a) Morphology of organoids grown in large and small droplets over time ($n_{\text{Large}} \geq 34$ and $n_{\text{Small}} \geq 24$ for each day). (b) Probability of obtaining spheroids versus branched structures in small (red) versus large (blue) droplets ($N = 3$ independent experiments with $n_{\text{Small}} \geq 116$ and $n_{\text{Large}} \geq 153$ structures evaluated for each experiment). Error bars indicate the standard deviation. (c) The major axis measured for organoids grown in small (red) and large (blue) droplets ($n_{\text{Large}} \geq 34$ and $n_{\text{Small}} \geq 24$ for each day). (d) The distribution of the area for 2D projected organoids on day 6 for organoid culture in small (red) and large droplets (blue) and in bulk (green) ($n_{\text{Small}} = 24$, $n_{\text{Large}} = 60$, $n_{\text{Bulk}} = 13$).

(Fig. 2c). By day 5, organoids in small droplets measured $150 \pm 42 \mu\text{m}$ (mean \pm s.d.) along their major axis, while those in large droplets reached a mean major axis of $266 \pm 89 \mu\text{m}$ (mean \pm s.d.). Despite this, even organoids in large droplets only attained half the size of those grown in bulk organoid gels. To assess if droplet-grown organoids displayed a more homogeneous size distribution, we measured the area of the 2D projected organoids, A_{proj} , and computed the polydispersity index $\text{PDI} = \sqrt{(\langle A_{\text{proj}}^2 \rangle - \langle A_{\text{proj}} \rangle^2) / \langle A_{\text{proj}} \rangle}$ as an estimate of the homogeneity of the organoid culture.¹⁹ The polydispersity index for bulk-grown organoids is $\text{PDI} = 63\%$. The one for organoids grown in large droplets decreases to $\text{PDI} = 47\%$. The polydispersity index for organoids grown in small droplets is reduced to $\text{PDI} = 28\%$. Therefore, a

restriction in the available volume of the hydrogel improves the homogeneity of the organoid culture (Fig. 2d), while at the same time reducing morphological complexity.

Characterisation of collagen interaction

The formation of branched structures depends on the cells' ability to interact and plastically deform their surrounding matrix.^{9,13} By pulling on the collagen fibres, the organoid forms a collagen cage that facilitates the development and growth of branched structures. Given the increased branching tendency of organoids in large droplets, we investigated if these organoids also displayed increased interaction with their surrounding matrix. To assess this, we monitored changes in droplet diameter over time. Large droplets underwent visible deformation after

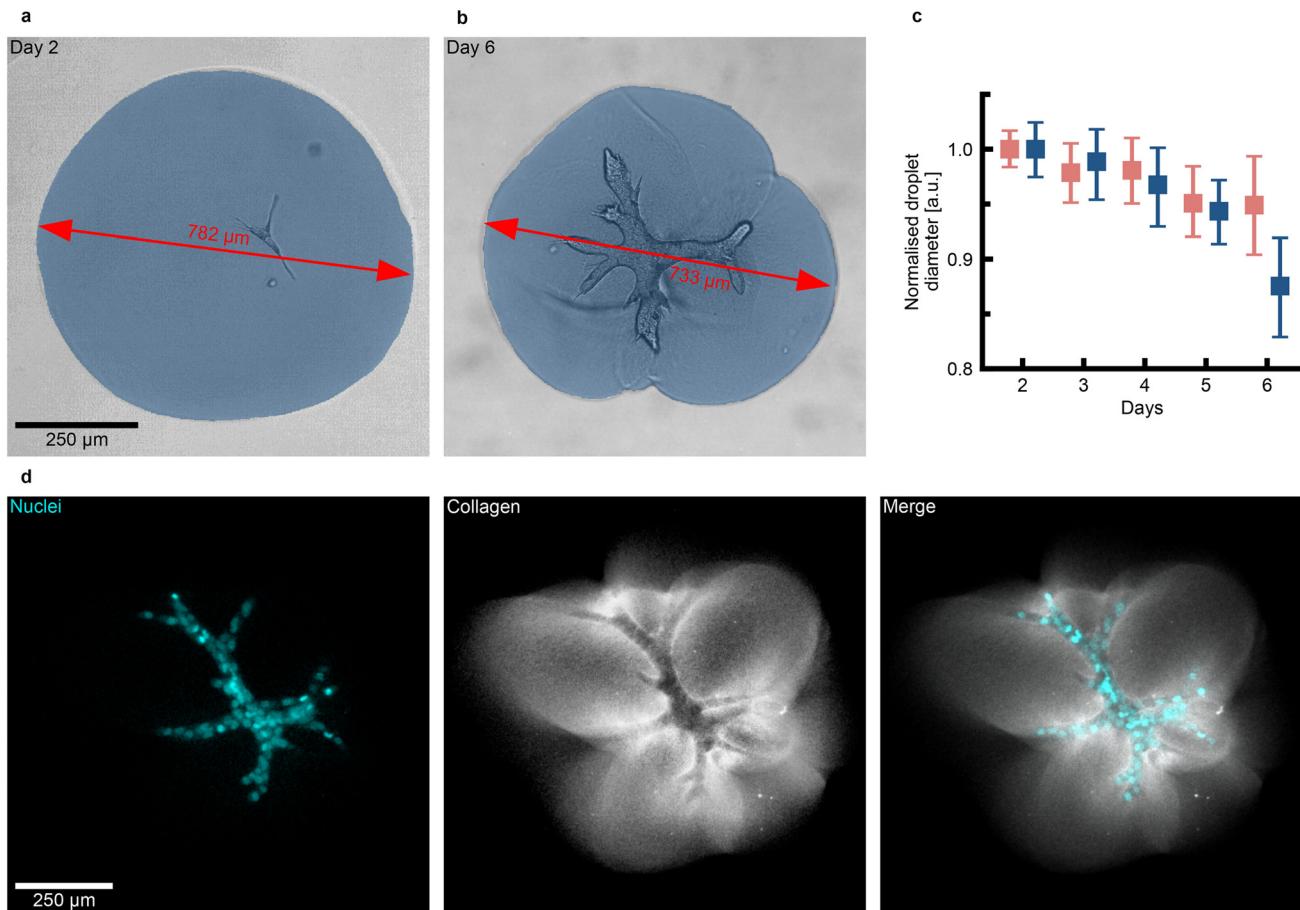


Fig. 3 Organoids contract the collagen droplets. (a) The droplets' surface is homogeneous 2 days after organoid seeding as the growing organoid does not yet exert sufficient mechanical force to deform the collagen matrix ($n = 32$ droplets). The segmented collagen droplet is visualised in blue. (b) The droplet's surface is increasingly irregular 6 days after organoid seeding, as the branched organoid pulls on the surrounding matrix and causes increasing droplet deformation, resulting in a reduced diameter ($n = 29$ droplets). (c) For large droplets, branched structures cause a 12% reduction in droplet diameter because of collagen deformation. In contrast, small droplets only undergo 5% of diameter reduction on day 6 ($n_{\text{Large}} \geq 23$ and $n_{\text{Small}} \geq 23$ for each time point). Error bars indicate the 95% confidence interval. (d) The deformation of a collagen droplet by an organoid cultivated for 5 days leads to the formation of a collagen cage. The collagen deformation is the highest at the tips of the organoid's branches ($n = 2$ independent experiments).

one week in culture (Fig. 3a and b). We measured the 2D-projected shape of the droplets to quantify these deformations and examined whether the continuous pulling exerted by the organoids led to a decrease in droplet size. Our results showed that small and large droplets experienced similar amounts of contraction up to day 5. However, by day 6, large droplets were contracted by 12%, compared to only 5% for small droplets (Fig. 3c). To further investigate how organoids interact with the collagen droplet, we quantified their spatial positioning within the droplets on day 2 (Fig. S5a-c). Organoids cultured in large droplets were located within 40% of the droplet radius. In contrast, in small droplets, organoids were found within 80% of the radius, placing them much closer to the periphery. As the organoids were not consistently located at the centre, we computed the normalised minimal distance from the organoid boundary to the droplet boundary (Fig. S5a). Up to day 5, the normalised minimal distance gradually decreased, and the values measured for organoids grown in small and large droplets were

comparable (Fig. S5d). However, by day 6, the organoids in the large droplets displayed a stronger decrease in the normalised minimal distance compared to the organoids cultured in small droplets, suggesting more extensive growth and mechanical deformation in the large droplets at day 6. This observation was further supported by the analysis of the droplet compactness, which captures the anisotropic deformation of the collagen droplets (Fig. S5e). The compactness for large droplets also shows a stark decrease on day 6, which aligns with the observed droplet contraction. To examine how complex morphologies, such as branched structures, influence mechanical deformation, we used fluorescently labeled collagen and stained PDAC cell nuclei with SiRDNA to visualise the interaction between the branched structures and the collagen matrix. By day 5, branched structures were observed to deform the surrounding collagen. Indeed, the areas with the most deformation coincided with the locations of organoid branches. Additionally, the increased collagen intensity at the organoid boundary suggested the formation of a collagen

cage, which is crucial for the cells to expand into a branched structure (Fig. 3d). Beyond day 6, organoids grown in large droplets continued to contract the collagen (Fig. S6a and b). Simultaneously, we observed that the tips of the organoids appeared increasingly rounded.

Characterisation of organoid morphology

In bulk, branched PDAC organoids exhibit a characteristic lumen formation phase in their final development stage.¹³ Leading up to lumen formation, the organoids undergo increasing epithelialisation, which is marked by a buildup of E-cadherin and F-actin at the future apical sites where the lumen will form (Fig. 4a). Lumen formation sites can be observed in bulk-grown organoids as early as day 11.³¹ In contrast, droplet-grown organoids display a notably faster

progression. On day 4, the droplet-grown organoids still display a mesenchymal phenotype characteristic for bulk-grown organoids on day 7 (Fig. S7a and b). However, by day 6, clear signs of epithelialisation are already evident, with E-cadherin and F-actin accumulation at lumen nucleation sites within the branches, along with some fully opened lumens on day 7 (Fig. 4b and c). This accelerated development in droplet-grown organoids results in lumen formation and maturation occurring six days earlier than in bulk-grown organoids, indicating a more rapid maturation process.

Characterisation of the transcriptome for organoids grown in large droplets *versus* bulk

Our imaging data showed that organoids grown in large droplets developed key morphological features and reached

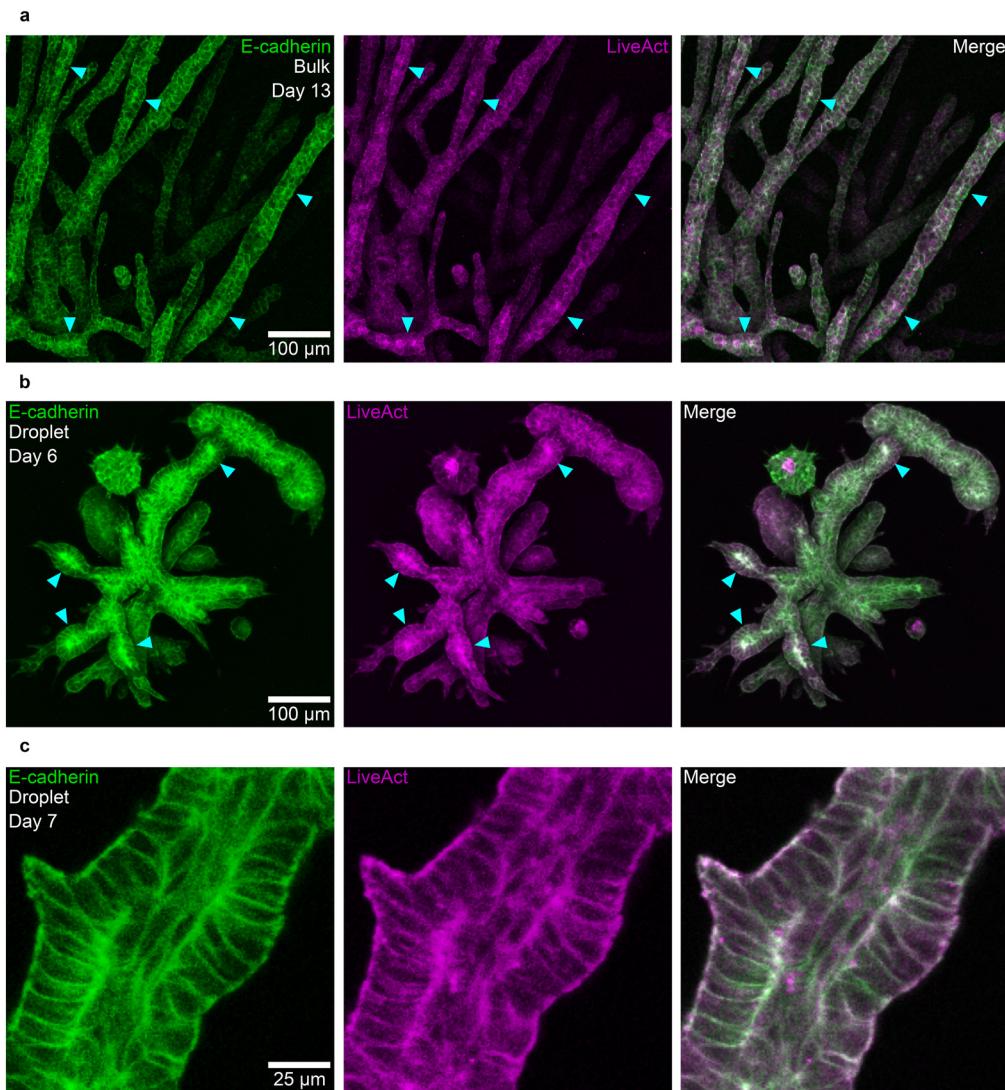


Fig. 4 Branched organoids in large droplets display similar developmental stages to those in bulk. (a) PDAC organoids cultured in bulk display local heterogeneous expression levels of E-cadherin and F-actin that indicate lumen formation sites on day 13 ($n = 3$ independent experiments). (b) PDAC organoids grown in large droplets display local heterogeneous expression levels of E-cadherin and F-actin, indicating lumen formation sites already on day 6 ($n = 3$ independent experiments). (c) A fully formed lumen in a branched organoid grown in a large droplet on day 7 ($n = 3$ independent experiments).



developmental hallmarks five days earlier than those in bulk organoid culture. To rule out the possibility that the higher collagen concentration of 3 mg ml^{-1} used in droplet cultures, compared to the 1.3 mg ml^{-1} used in bulk cultures, were responsible for the accelerated onset of lumen formation, we

analyzed the transcriptomic profiles of organoids grown in 1.3 mg ml^{-1} and 3 mg ml^{-1} bulk collagen gels on day 4 and 7 (Fig. S8a-h). In both cases, the transcriptomic profiles remained highly similar, suggesting that the differences in collagen concentration cannot account for the observed acceleration in

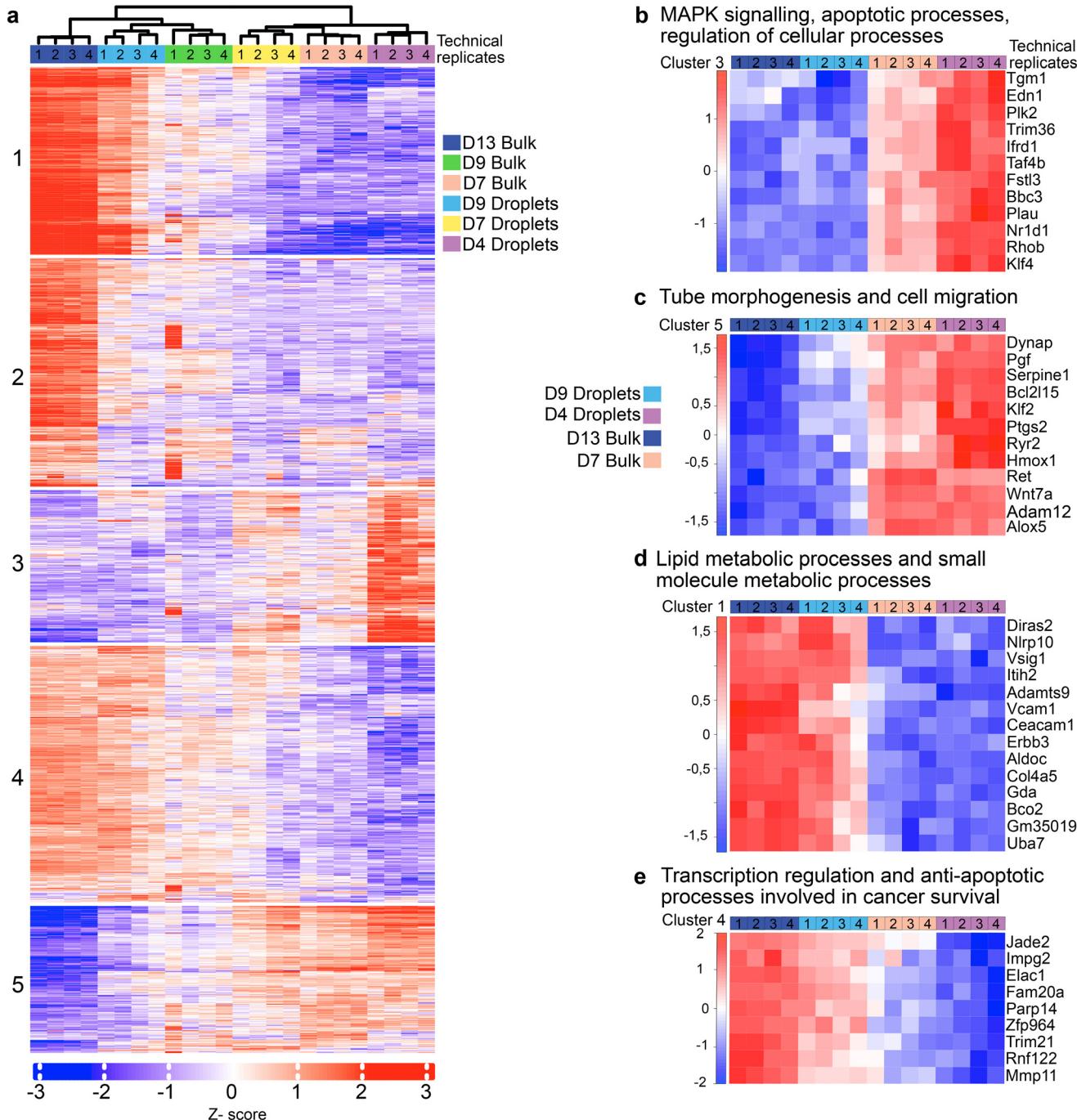


Fig. 5 PDAC organoids expression pattern in different growth conditions. (a) K -means clustering with five clusters ($k = 5$) using 2000 genes with the most variable expression level between different growth conditions: droplets day 4, 7, 9 and bulk day 7, 9, 13 ($n = 4$ individual experiments), FDR = 0.05. (b) GO biological processes enrichment analysis from cluster 3 for selected conditions (droplets day 4, 9 and bulk day 7, 13) highlights genes involved in MAPK signaling and apoptosis-related processes. (c) Gene expression heatmap (cluster 2) for selected genes regulating tube morphogenesis and cell migration. (d) Gene expression heatmap (cluster 1) of a subset of genes involved in lipid metabolism and small molecule metabolism. (e) Heat map for selected genes (cluster 4) involved in transcription regulation and anti-apoptotic pathways facilitating cancer survival.



organoid maturation. Furthermore, our analysis only identified a few significantly differentially expressed genes for organoids grown in small and large droplets on day 4 and day 7 (Fig. S8e and f). In contrast, the comparison of organoids grown in 1.3 mg ml⁻¹ bulk collagen to those cultured in large droplets on day 4 revealed a greater number of significantly differentially expressed genes, than the comparison between bulk-grown organoids on day 7 and droplet-grown organoids on day 4, consistent with the morphological observations of accelerated maturation (Fig. S8g and h). To further explore this, we compared the transcriptomes of organoids grown in bulk culture to those grown in large droplets. As previously reported, bulk organoid culture shows a stage-specific gene expression pattern, with genes being up- or downregulated according to the developmental phase. Early-stage organoids exhibit a mesenchymal-like phenotype, transitioning to a more epithelial-like phenotype at later stages.¹³ To evaluate how the geometrical confinement effects influence these gene expression dynamics, we analyzed the top 200 most variably expressed genes across all samples, using Euclidean clustering to visualise a gene expression heat map. In bulk culture, we observed the expected oscillatory pattern, where certain genes were downregulated at day 7 and upregulated at day 13 as the organoids matured (Fig. 5a). Day 9 represented a transition point, with partially overlapping gene expression profiles between the two states. In droplet-grown organoids, the gene expression profile on day 9 closely resembled the profile on day 13 in bulk culture, consistent with the accelerated maturation timeline observed in their morphology. Given this shift in the timing of gene regulation, we conducted a deeper analysis of the expression patterns across different gene clusters. We identified a switch in the gene expression pattern involved in critical metabolic processes, such as lumen formation and branching morphogenesis, relevant during PDAC organoid maturation (Fig. 5b and c).^{13,31} For later time points (D13 bulk and D9 droplets), we detected a population of overexpressed genes involved in cancer survival and lipid metabolic processes characteristic of enhanced metabolic activity during tumour progression (Fig. 5d).⁴⁰ In droplet-based organoid culture at day 9, genes responsible for cancer survival and anti-apoptotic processes follow the characteristic upregulation pattern observed in bulk organoid culture at day 13, indicating faster maturation (Fig. 5e).

Conclusion

Encapsulating singularised cells into collagen droplets resulted in the formation of individual highly branched organoid structures. Thus, droplet-based organoid culture improved yield and sample homogeneity and achieved a faster maturation of the organoids. A reduction in droplet size, while improving organoid homogeneity, entailed a loss in morphological complexity. Thus, only in large droplets did we obtain organoids with similar structural complexity to organoids grown in bulk culture. Moreover, it demonstrates the feasibility of cultivating droplet-grown organoids with complex morphologies rather

than spherical shapes. Using microscopic and transcriptomic analysis, we confirmed the accelerated maturation of organoids grown in geometrically confined microenvironments, which aligns with previous studies on stem cell aggregates cultured in microwells.⁴¹ Although our study highlights the influence of droplet size, it does not directly assess the effective mechanical stiffness experienced by the organoids. Future work incorporating physical modeling and molecular force reporters could address this gap and help uncover the biological mechanisms underlying the observed acceleration in maturation. While previous studies have emphasised the role of collagen concentration in modulating individual and collective cell behaviour,^{42,43} our findings demonstrate that mechanical boundary conditions are fundamental for steering organoid structure formation processes. This highlights the importance of understanding how altered geometrical and mechanical microenvironments affect maturation timelines. Although the mechanisms behind the observed accelerated maturation remain unclear, our findings underscore the need to account for potential shifts in timescales when using droplet-based organoid cultures. Such shifts may impact critical aspects of dose-response studies, including the timing of drug exposure, sensitivity to treatments, and the reproducibility of results.

Author contributions

I. R. performed experiments with A. P.'s support under the supervision of A. R. B., M. K. R. generated the cell lines used in this study and performed bulk-organoid experiments. I. R. and W. X. and Y. L. conceived the microfluidic chip design and fabricated the masters. A. R. B., and D. W. conceived the project with the support of M. R. Data Analysis was performed by I. R. and A. P. All authors participated in the writing of the manuscript.

Conflicts of interest

The authors declare no competing interests.

Data availability

The datasets generated and analyzed during the current study are available in a Zenodo repository titled “Accelerated Maturation of Branched Organoids Confined in Collagen Droplets” (DOI: <https://doi.org/10.5281/zenodo.14070465>).⁴⁴ The raw data obtained for this study are available from the corresponding author upon reasonable request.

Supplementary information is available: The SI includes the microfluidic chip design and statistical analyses characterising droplet size and organoid growth within collagen droplets. It also provides morphological and transcriptomic data supporting the comparison of bulk organoid cultures with droplet-grown organoids. See DOI: <https://doi.org/10.1039/D5LC00287G>.



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References

- 1 G. Rossi, A. Manfrin and M. P. Lutolf, *Nat. Rev. Genet.*, 2018, **19**, 671–687.
- 2 Z. Zhao, X. Chen, A. M. Dowbaj, A. Sljukic, K. Bratlie and L. Lin, *et al.*, *Nat. Rev. Methods Primers*, 2022, **2**, 1–21.
- 3 S. Y. Lee, I. S. Koo, H. J. Hwang and D. W. Lee, *SLAS Discovery*, 2023, **28**, 119–137.
- 4 S. M. Kang, D. Kim, J. H. Lee, S. Takayama and J. Y. Park, *Adv. Healthcare Mater.*, 2021, **10**, 2001284.
- 5 A. I. Vazquez-Armendariz and S. Herold, *Front. Cell Dev. Biol.*, 2021, **9**, 631579.
- 6 D. Iber, *Curr. Top. Dev. Biol.*, 2021, **143**, 205–237.
- 7 S. I. Mae, M. Ryosaka, S. Sakamoto, K. Matsuse, A. Nozaki and M. Igami, *et al.*, *Cell Rep.*, 2020, **32**, 107963.
- 8 N. Gupta, E. Dilmen and R. Morizane, *J. Mol. Med.*, 2021, **99**, 477–487.
- 9 B. Buchmann, L. K. Engelbrecht, P. Fernandez, F. P. Hutterer, M. K. Raich and C. H. Scheel, *et al.*, *Nat. Commun.*, 2021, **12**, 1–10.
- 10 L. Yuan, S. Xie, H. Bai, X. Liu, P. Cai and J. Lu, *et al.*, *Nat. Methods*, 2023, **20**, 2021–2033.
- 11 L. Flasse, C. Schewin and A. Grapin-Botton, in *Cellular Networks in Development*, Academic Press, 2021, vol. 143, pp. 75–110.
- 12 Z. Ghezelayagh, M. Zabihi, M. K. Ashtiani, Z. Ghezelayagh, F. C. Lynn and Y. Tahamtani, *Cell. Mol. Life Sci.*, 2021, **78**, 7107–7132.
- 13 S. Randriamanantsoa, A. Papargyriou, H. C. Maurer, K. Peschke, M. Schuster and G. Zecchin, *et al.*, *Nat. Commun.*, 2022, **13**, 1–15.
- 14 A. Papargyriou, M. Najajreh, D. P. Cook, C. H. Maurer, S. Bärthel and H. A. Messal, *et al.*, *Nat. Biomed. Eng.*, 2024, **9**(6), 836–864.
- 15 S. W. Lee, M. J. Song, D. H. Woo and G. S. Jeong, *Biomed. Pharmacother.*, 2024, **174**, 116511.
- 16 L. Nan, H. Zhang, D. A. Weitz and H. C. Shum, *Lab Chip*, 2024, **24**, 1135–1153.
- 17 T. Moragues, D. Arguijo, T. Beneyton, C. Modavi, K. Simutis and A. R. Abate, *et al.*, *Nat. Rev. Methods Primers*, 2023, **3**, 1–22.
- 18 R. F. Tomasi, S. Sart, T. Champetier and C. N. Baroud, *Cell Rep.*, 2020, **31**, 107670.
- 19 H. Zhao, Y. Cheng, J. Li, J. Zhou, H. Yang and F. Yu, *et al.*, *Fundam. Res.*, 2024, **4**(6), 1506–1514.
- 20 P. J. Cohen, E. Luquet, J. Pletenka, A. Leonard, E. Warter and B. Gurchenkov, *et al.*, *Biomaterials*, 2023, **295**, 122033.
- 21 H. F. Chan, Y. Zhang and K. W. Leong, *Small*, 2016, **12**, 2720–2730.
- 22 Z. Chen, S. Kheiri, A. Gevorkian, E. W. Young, V. Andre and T. Deisenroth, *et al.*, *Lab Chip*, 2021, **21**, 3952–3962.
- 23 A. Saint-Sardos, S. Sart, K. Lippera, E. Brient-Litzler, S. Michelin and G. Amselem, *et al.*, *Small*, 2020, **16**, 2002303.
- 24 G. Ronteix, S. Jain, C. Angely, M. Cazaux, R. Khazen and P. Bousso, *et al.*, *Nat. Commun.*, 2022, **13**, 1–13.
- 25 Z. Wang, M. Boretto, R. Millen, N. Natesh, E. S. Reckzeh and C. Hsu, *et al.*, *Stem Cell Rep.*, 2022, **17**(9), 1959–1975.
- 26 E. Princee, S. Kheiri, Y. Wang, F. Xu, J. Cruickshank and V. Topolskaia, *et al.*, *Adv. Healthcare Mater.*, 2022, **11**, 2101085.
- 27 M. Trossbach, E. Åkerlund, K. Langer, B. Seashore-Ludlow and H. N. Joensson, *SLAS Technol.*, 2023, **28**, 423–432.
- 28 S. Hong, H. J. Hsu, R. Kaunas and J. Kameoka, *Lab Chip*, 2012, **12**, 3277–3280.
- 29 H. Liu, Y. Wang, H. Wang, M. Zhao, T. Tao, X. Zhang and J. Qin, *Adv. Sci.*, 2020, **7**, 1903739.
- 30 Y. Xiao, Q. Huang, J. W. Collins, J. Brouchon, J. A. Nelson and Z. Niziolek, *et al.*, *Organoids*, 2023, **2**, 204–217.
- 31 S. J. Randriamanantsoa, M. K. Raich, D. Saur, M. Reichert and A. R. Bausch, *iScience*, 2024, **27**, 110299.
- 32 T. Kiyomitsu and I. M. Cheeseman, *Cell*, 2013, **154**, 391.
- 33 J. R. Linnemann, H. Miura, L. K. Meixner, M. Irmler, U. J. Kloos and B. Hirschi, *et al.*, *Development*, 2015, **142**, 3239–3251.
- 34 M. Pachitariu and C. Stringer, *Nat. Methods*, 2022, **19**, 1634–1641.
- 35 J. Schindelin, I. Arganda-Carreras, E. Frise, V. Kaynig, M. Longair and T. Pietzsch, *et al.*, *Nat. Methods*, 2012, **9**, 676–682.
- 36 L. A. L. Abueg, E. Afgan, O. Allart, A. H. Awan, W. A. Bacon and D. Baker, *et al.*, *Nucleic Acids Res.*, 2024, **52**, W83–W94.
- 37 S. X. Ge, E. W. Son and R. Yao, *BMC Bioinf.*, 2018, **19**, 1–24.
- 38 Z. He, J. Wang, B. J. Fike, X. Li, C. Li, B. L. Mendis and P. Li, *Biosens. Bioelectron.*, 2021, **191**, 113458.
- 39 K. R. Jegannathan, E.-S. Chan and P. Ravindra, *J. Mol. Catal. B: Enzym.*, 2009, **58**, 78–83.
- 40 H. Cheng, M. Wang, J. Su, Y. Li, J. Long and J. Chu, *et al.*, *Life*, 2022, **12**, 784.
- 41 N. Brandenberg, S. Hoehnel, F. Kuttler, K. Homicsko, C. Ceroni and T. Ringel, *et al.*, *Nat. Biomed. Eng.*, 2020, **4**, 863–874.



42 S. I. Fraley, P.-h. Wu, L. He, Y. Feng, R. Krisnamurthy, G. D. Longmore and D. Wirtz, *Sci. Rep.*, 2015, **5**, 14580.

43 C. Mark, T. J. Grundy, P. L. Strissel, D. Böhringer, N. Grummel and R. Gerum, *et al.*, *eLife*, 2020, **9**, e51912.

44 I. Ruider, A. Pastucha, M. Raich, W. Xu, Y. Liu, M. Reichert, D. Weitz and A. R. Bausch, Accelerated maturation of branched organoids confined in collagen droplets, *Zenodo*, 2025, DOI: [10.5281/zenodo.14070465](https://doi.org/10.5281/zenodo.14070465).

