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Bioinspired human stomach-on-a-chip with in vivo like function and architecture†

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The lack of biomimetic in vitro models capable of reproducing the complex architecture and the dynamic environment of the gastric mucosa, delay the development of diagnostic and therapeutic tools. Recent advances in microengineering made possible the fabrication of bioinspired microdevices capable of replicating the physiological properties of an organ, inside a microfluidics chip. Herein, a bioinspired stomach-on-a-chip (SoC) device is described, supporting peristalsis-like motion and reconstituting organlevel epithelial architecture and function. The device simulates the upper epithelial interface, representing the three innermost layers of the gastric mucosa, namely the epithelial barrier, the basement membrane and the lamina propria. The dynamic environment imparted by mechanical actuation of the flexible onchip cell culture substrate, was the main driver in the development of epithelial polarization and differentiation traits characteristic of the native gastric mucosa, and allowed partial recapitulation of gastric barrier function. These traits were not affected by the addition of a mesenchymal population to the system, which was able to remodel the surrounding extracellular matrix, nor by the potential epithelialmesenchymal cross-talk. The engineered platform highlights the importance of addressing the mechanical microenvironment of the native organ, to potentiate an organ-level response of the artificial tissue. The proposed SoC represents an appealing tool in personalized medicine, with bio-relevance for the study of gastric diseases and an alternative to current animal models.

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1. Introduction

The stomach plays a central role in human physiology, and gastric dysfunction poses serious health risks with associated non-neglectable mortality and morbidity rates. ¹⁻³ For instance, gastric cancer, represented the fourth leading cause of estimated cancer related deaths in 2020. ⁴ The demand for better study models accurately recapitulating human gastric physiology is high, mainly to improve the way therapies for gastric-associated diseases are currently tested.

Two-dimensional (2D) cell culture models, lacking the complexity of human tissue architecture and physiology, are commonly used for early stages of drug development but, as expected, their simplified nature has a very low predictive power in drug efficacy.^{5,6} Three dimensional (3D) models, engineered with bio-inspired matrices replicating the chemistry, assembly, multicellular nature and architecture of living organs, provide a better simulation of physiological conditions.^{7,8} Similarly, addition of primary and stem cells, was key to the development of increasingly complex *in vitro* gastric models.^{9–12} Notwithstanding, these models were only developed for non-human cells. Recently, a long-lived, airliquid interface gastric mucosoid culture model of human origin, was developed, closely resembling the phenotypical

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features of normal human gastric epithelium. 13 However, it represented the gastric epithelial layer only, limiting the evaluation of physiological responses. We previously reported a 3D two-layered model, mimicking the gastric mucosa and lamina propria.¹⁴ The augmented 3D complexity reflected more closely the architecture of the native organ, while the fibroblastic components provided bioactive signaling to the epithelial layer. Despite notable progress, even the most advanced 3D culture models are limited and still lack the dynamic nature of the gastric mucosa, which is known to be extremely important for normal gastrointestinal physiology. 15,16 In vivo, cells are continuously sensing their vicinity and mechanical cues can, to a large extent, modulate the cellular response at the biochemical level. 17,18 The contribution of biomechanical forces is particularly relevant in tissues such as the gut, where cells are repeatedly exposed to strain. 19-22 Recent developments in soft-lithography, in particular in the fabrication of microactuators, enabled the creation of organ-on-a-chip devices, consisting in microfluidic platforms with capability to replicate a given functional sub-unit of a human organ. ^{21–25} This bioinspired technology enabled the application of precisely defined

mechanotransduction at the cellular and molecular level. 19,26 Recently, Lee and colleagues reported preliminary developments on a stomach-on-a-chip (SoC) model, where gut organoids were encased within a microfluidic device and exposed to cyclic peristalsis-like stretching.²⁷ While presenting a promising model to address the effects of dynamic actuation on a differentiated gastric mucosa, the system used a matrigel-derived supporting gel that poorly recapitulated the human gastric extracellular matrix (ECM) composition and architecture. A more physiologically relevant environment, resembling the native gastric ECM, could be a critical asset when addressing system level response within these devices. Applications focusing on pathological ECM remodeling, as is the case during tumor invasion, could particularly benefit from such a model, since these processes are largely mediated by alterations in ECM components. 28,29

amounts of strain to cell culture substrates, thus allowing to

understand, to a greater level of detail, the implications of

In the current work, we propose a tissue engineering approach to design and fabricate a human-derived biomimetic SoC microdevice, experiencing peristalsis-like movement and intraluminal flow, with architecture and function emulating the native organ. The currently developed model was engineered to represent the three innermost layers of the gastric mucosa, namely the epithelial barrier (characterized by a tight cellular monolayer), the basement membrane (a layer mediating the adhesion and communication between the epithelial layer and the adjacent ECM), and an analogue of the lamina propria (consisting of a collagen type I gel with embedded normal gastric fibroblasts, able to potentially provide paracrine crosstalk between the lamina propria analogue and the epithelial layer and remodel the surrounding ECM). We then characterized the cellular and molecular alterations induced by the dynamic

environment of the SoC, to assess its ability to promote functional and molecular traits characteristic of the native gastric mucosa.

The developed SoC represents a human-derived experimental *in vitro* asset for pre-clinical studies on host-pathogen interactions, drug-drug interactions, drug absorption or targeted delivery of compounds and nanoparticles, among many other possibilities.

2. Results and discussion

2.1. Design and fabrication of a multilayered stomach-on-a-chip (SoC) device

The SoC was designed and fabricated using a rapid prototyping method that we previously described.²⁵

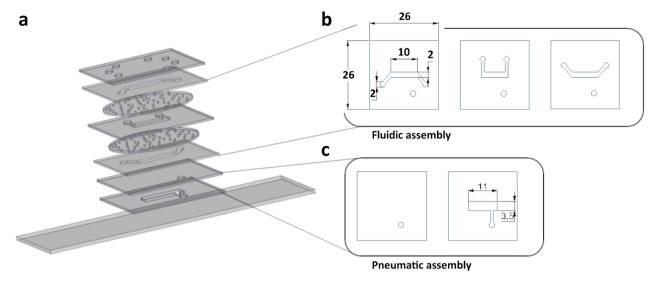
To represent the three innermost layers of the stomach wall (epithelial barrier, basement membrane and supporting lamina propria), the SoC was devised as 9-layered device (Fig. 1a). The top layer was produced from thick PDMS, fabricated by gravity casting. The fluidic network was produced from 500 µm thick PDMS sheets and consisting of 3 fluid routing structures, intercalated by a 16 µm thick polyethylene terephthalate (PET) membrane (Fig. 1a and b). The proposed 3-tiered fluidic structure was designed to enable the replication of an epithelial barrier architecture on the top chamber and importantly, to provide the efficient incorporation of a lamina propria analogue, in the middle culture chamber. This allowed ECM-embedded fibroblasts to be adequately fed by flowing cell culture media above and underneath it. The microactuator was bonded directly below the fluidic network and is composed of a 2-part subassembly, namely a 250 µm thick PDMS membrane juxtaposed on a rectangular vacuum chamber of the same thickness (Fig. 1c). The entire biochip was sealed against a microscope glass slide.

2.2. On-chip emulation of stomach epithelium dynamics and architecture

To model the dynamic stomach microenvironment, peristalsis-like motion was imparted to the cell culture substrate by applying vacuum to the microactuator at cyclical intervals. The stretching pattern was modeled as a sinusoidal wave of 0.15 Hz frequency, which closely resembles the peristalsis pattern of the gut, ^{22,30,31} at a maximum vacuum pressure of –100 mbar. We have previously shown that under these conditions, the integrated microactuator could produce surface stretching of approximately 10%, which is within the physiological range of strain that gut cells experience *in vivo*. ²⁵ Furthermore, as the microactuator was placed directly underneath the cell culture chamber, membrane flexing was effected from below, resulting in a tri-axial displacement of the cell culture membrane.

Cell culture media was perfused over the epithelial layer at a flow rate of 2 μ L min⁻¹. As shear stress is not a dominant force on the stomach lumen, the fluidic routing structures

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d Native mucosa architecture

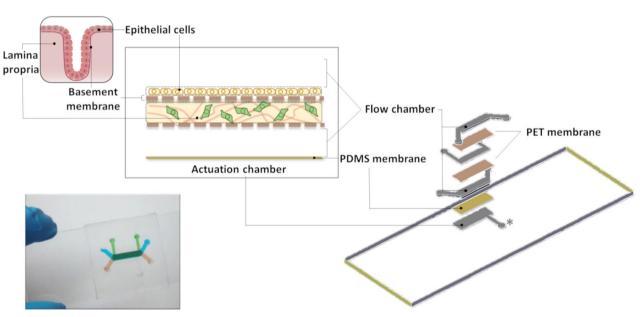


Fig. 1 Graphical representations of the stomach-on-a-chip (SoC). (a) Exploded view of the SoC depicting the 9 intercalating parts composing the fluidic and pneumatic assembly: the topmost layer corresponds to a thick PDMS layer produced by gravity casting to support tube attachment, while the bottommost layer corresponds to a microscope glass slide. (b) Top-down view of the fluidic structures composing the fluidic assembly. Each of the layers is intercalated by a PET membrane. (c) Top down view of the 2-part pneumatic assembly consisting of a thin 250 μm PDMS membrane (left) and a vacuum chamber placed underneath the membrane (right). (d) Schematic of the SoC, highlighting the engineered gastric mucosa with a graphical representation of the 3 innermost layers of the gastric wall, namely the epithelial barrier, the basement membrane and the lamina propria. The exploded view of the biochip, depicts the fluidic and microactuator (*) structures in grey. PET and PDMS membranes intercalate the fluidic structures as depicted. The epithelial and the basement membrane analogues are emulated within the upper chamber. The basement membrane is a thin coat of fibronectin, covering the uppermost PET membrane, on top of which the epithelial layer is seeded. The lamina propria analogue is injected in the middle chamber. Upper and lower fluidic chambers are used to perfuse cell culture media for efficient cell feeding.

were designed to keep shear stress at a low level. Taking in consideration a channel profile, 0.20 cm wide and 0.05 cm high, and a volumetric flow rate of 3.3×10^{-5} cm³ s⁻¹, we estimated the shear stress over the epithelial cell culture to

be about 0.003 dyne per cm². This value is in line with the physiological shear stress experienced by gut cells *in vivo*, which is expected to range between 0.002 and 0.080 dyne per cm². 22,32

Emulation of the stomach epithelium was achieved on the two upper fluidic channels. The top channel encased the basement membrane analogue, consisting of a thin 50 µg mL⁻¹ fibronectin gel coating the PET membrane, on top of which an epithelial monolayer was seeded (Fig. 1d). To mimic the selective permeability of the gastric mucosa it is important that the epithelial layer forms a cohesive, tightly packed monolayer, expressing fully developed cell-to-cell contacts. Here, we used a monolayer culture of MKN74 epithelial gastric cells (Table S1†). This cell line displays moderate differentiation, a typical epithelial cobblestone-like morphology and exhibits several characteristics of its epithelial origin. It displays fully developed and correctly localized tight junctions that mediate paracellular traffic across the epithelial barrier, as shown by staining of tight junction partners ZO-1 and occludin (Fig. S1a†). It also shows functional E-cadherin mediated cell-to-cell contacts, via adherens-junctions, as observed by co-immunofluorescence against E-cadherin and adherens-junction partner α -catenin (Fig. S1b†). This adhesion complex mediates tight monolayer integrity, by establishing cell-to-cell contacts and promotes an indirect link between the actin cytoskeleton of adjacent cells. The ability of cells to form an effective barrier was assessed by measuring the transepithelial electrical resistance (TEER). After day 2 post seeding, the cell line already exhibited TEER values above 100 Ω cm², characteristic of a fully formed barrier, reaching an average of about 157 Ω cm² after 5 days in culture (Fig. S1c†).

Taken together, these observations suggest that the MKN74 cell line was well suited to emulate the gastric barrier in the current SoC model.

To mimic the lamina propria, we used a 3 mg mL⁻¹ collagen type I gel, since it is the most prevalent ECM protein in the human gastric lamina propria.³³ The collagen gel was then embedded with normal gastric NST-20 fibroblasts (Table S1†), potentially introducing physiologically relevant ECM remodeling of the lamina propria analogue. The ECM was directly injected in the central fluidic channel, while the top and bottom perfusion channels acted as a direct feeding layer to the ECM embedded fibroblasts (Fig. 1d).

2.3. On-chip conditions did not affect cellular homeostasis of the epithelial barrier

Although PDMS-based devices are permeable to air, it is unlikely that this exchange is sufficient to ensure an efficient buffering capacity of the atmospheric 5% CO₂, used in conventional cell culture incubators, particularly in pressurized perfusion systems, such as the one used here. To ensure that cells growing under CO₂ depletion and cells growing under conventional cell culture atmosphere had the same basal readings, we compared their phenotype, proliferation, metabolic activity and doubling time, when fed with CO₂-independent culture medium in a CO₂-depleted atmosphere or with RPMI 1640 in the presence of CO₂. No morphological differences were registered between the

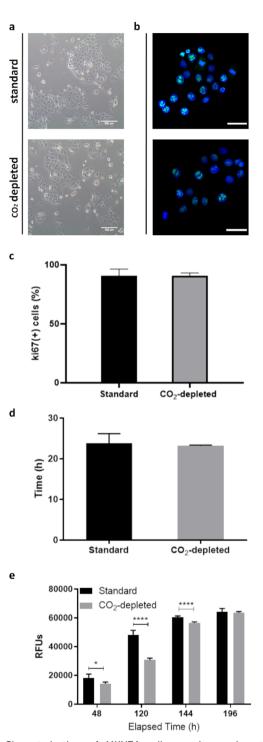


Fig. 2 Characterization of MKN74 cells growing under standard conditions vs. CO_2 -depleted conditions. (a) Morphological study of MKN74 cells growing under standard conditions against cells growing under CO_2 -depleted conditions (scale bars; 200 μ m). (b) Ki67 staining of the same populations (scale bars; 50 μ m). (c) Number of Ki67 positive cells was quantified for both populations and depicted as % of Ki67 positive cells (n=3) (d) corresponding population doubling time (n=6). (e) Comparative study of the metabolic activity of cells growing under normal or CO_2 -depleted conditions (n=6). All graphical representations display average measurement \pm SD.

2a). Furthermore, Ki67 staining and quantification at 48 h of growth showed mostly proliferative populations (Fig. 2b and c), and both exhibited a comparable doubling rate (Fig. 2d). Interestingly, we observed that cells growing under standard conditions, displayed a higher metabolic rate at early time points, compared to those grown in a CO2-depleted atmosphere. However, this difference equalized after ca. 8 days in culture (Fig. 2e). This

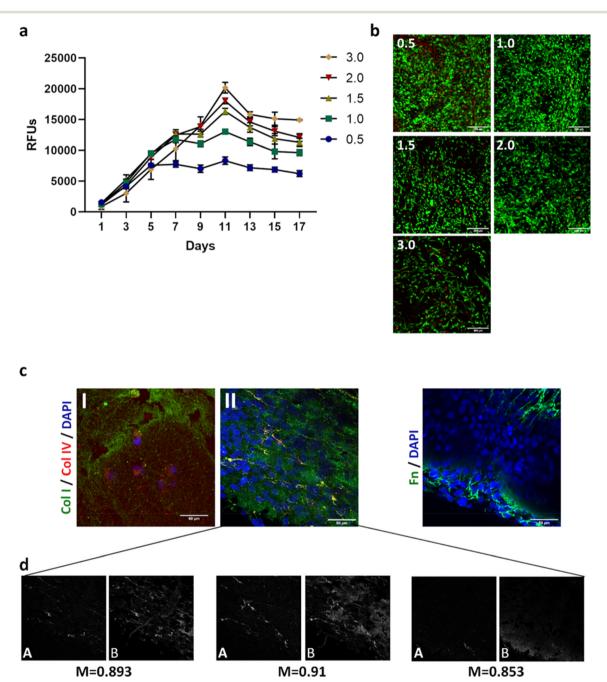


Fig. 3 Characterization of NST-20 normal gastric fibroblasts embedded in a collagen type I hydrogel gel. (a) Metabolic activity study of NST-20 in a range of collagen type I densities, namely 0.5, 1.0, 1.5, 2.0 and 3.0 mg mL⁻¹, over a period of 17 days. (b) Live-dead assay performed for the same populations at day 13 post seeding. Images represent a z-stack projection (z-step: 2 μm) across the entire thickness of the hydrogel. Live cells stained with calcein and depicted in green, whereas dead cells are stained with propidium iodide and seen in red (scale bars: 200 µm). (c) Matrix remodeling as effected by the collagen-embedded NST-20 fibroblasts. Collagen type I staining of the supporting hydrogel can be observed in green (left panel). Collagen type IV stained in red (left panel), 1 day post-seeding (I) and 11 days post-seeding (II) and fibronectin stained in green (right panel), exhibit de novo deposition of ECM components by the fibroblastic population. Nuclei stained with DAPI (blue). (d) Co-localization study for collagen type I and collagen type IV as seen in (c). Panels show collagen type IV (A) and collagen type I (B) and represent the same image source at 3 different z-heights. Mander's coefficient (M) displayed, indicates the fraction of fluorescence signal in A overlapping the fluorescence signal in B and was determined using the Fiji plugin, JACoP on thresholded images. A value of 1.0 represents 100% co-localization of both fluorescence signals.

observation could suggest that cells growing under CO₂-depleted conditions, and fed with buffered medium, may need a longer period to attach and stabilize onto the substrate and start the replicative process, until they reach a similar doubling rate compared to cells growing in standard conditions.

2.4. Impact of embedding stomach fibroblasts within a collagen type I gel

To properly emulate the lamina propria of the stomach wall, it is important that the designed model replicates, at least in part, the cellular milieu of its native counterpart. Here, we used a normal stomach fibroblast cell line (NST-20) embedded in a collagen type I gel. We aimed to potentiate the crucial native ECM remodeling. In this context, it was important to assess if ECM embedded fibroblasts were metabolically stable and growing in optimal conditions within the hydrogel gel.

We first studied the off-chip behavior of gastric NST-20 fibroblasts, seeded at an initial density of 2×10^5 cells per mL, within a collagen type I gel. Observations were carried out over an increasing range of collagen type I densities from 0.5 to 3.0 mg mL⁻¹. Metabolic activity of ECM-embedded fibroblasts was assessed over a period of 17 days and quantified by resazurin assay.

Our observations showed that the metabolic activity increased proportionally with increasing matrix density (Fig. 3a). For 0.5 mg mL⁻¹ gels, metabolic activity peaked at day 5 and reached a stable plateau until the end of the experimental period. For any other gel density, metabolic activity increased steadily, reaching a peak at day 11. The same conditions displayed a small reduction in metabolic activity after day 11, which stabilized around day 15-17 in culture. Our results suggested that fibroblasts growing in a 3.0 mg mL⁻¹ collagen gel needed a longer time to stabilize and start the replicative process (Fig. 3a). This was most likely due to the stiffer environment in which cells were initially seeded. However, once cells stabilized and developed full anchoring, the stiffer conditions seemed more advantageous for cell metabolism. To address the hypothesis that the high metabolic rates observed for higher density gels could result from cells growing selectively on the surface of the hydrogel, and thus not within a truly 3D environment, we performed a live-dead assay at day 13 post-seeding to assess qualitatively the viability of the embedded fibroblasts and imaged the entire hydrogel volume by z-stack imaging (Fig. S2†). Our results showed that cells remained viable within the hydrogel, with no appreciable differences in viability from 1.0 to 3.0 mg mL⁻¹. In contrast, the 0.5 mg mL⁻¹ gel exhibited a high number of dead cells in the core of the collagen discs (Fig. 3b). We have observed that the lower density gels potentiated very fast cell growth and migration and within 3 days, the collagen gel was completely occupied by the embedded fibroblasts. In turn, this could potentiate a higher density of cells in the core of the hydrogel, which could

ultimately lead to a nutrient-depleted region within the core of the gel. Calcein staining also allowed morphological characterization of the stomach fibroblasts within the 3D matrices. Cells displayed a typical elongated spindle shape, characteristic of a mesenchymal phenotype. Protrusions establishing contact between adjacent cells could also be observed, suggesting that cell-to-cell signaling was being promoted within the collagen gel (Fig. 3b). Finally, evidence of ECM remodeling was also observed, as shown by immunocytochemistry against collagen type IV and fibronectin, both components of the native gastric lamina propria. De novo expression of collagen type IV could be observed as early as day 1 post-seeding (Fig. 3c), with extensive deposition of both collagen type IV and fibronectin matrix proteins after 11 days in culture (Fig. 3c). Of note, deposition of collagen type IV seemed to co-localize with thick bundles of collagen type I near the vicinity of fibroblasts, also suggesting de novo deposition of collagen type I (Fig. 3d). These results pointed to a robust remodeling of the lamina propria by the fibroblastic population, thus increasing the biomimetic dimensionality of the lamina propria analogue. Interestingly, remodeling of collagen type IV has been previously associated with malignant transformation in gastric tumors.34 De novo expression of accessory ECM molecules such as collagen type IV within our model, could be effectively explored to elucidate its role in the malignant transformation leading to gastric cancer. Taken together these observations suggested that NST-20 fibroblasts were not adversely affected by the embedding within the collagen type I hydrogel. Under these circumstances, they can reliably be used to mimic the environment of the gastric lamina propria.

It is well understood that fibroblasts anchored to a collagen substrate exert contractile forces that can reshape the collagen fibers into a higher density structure with concomitant loss of volume.35,36 However, integrating onchip a fibroblast-laden collagen hydrogel requires the hydrogel to withstand continued mechanical distension and retain, as much as possible, its shape in order to fill-in the culture chamber completely, thus avoiding disruption of fluid flow. Therefore, we next conducted a gel contraction assay, using the same range of collagen gel densities described above. We monitored fibroblast-laden hydrogel volume contraction every other day, for a period of 15 days. We observed a decrease of gel area proportional to decreasing hydrogel density, with lower density gels contracting significantly more. A 0.5 mg mL⁻¹ gel contracted on average ca. 86% of its original size by day 15 in culture. In sharp contrast, a 3.0 mg mL⁻¹ gel retained 77% of its original size over the same period (Fig. 4a and b).

Taking these observations into consideration, along with the fact that 3.0 mg mL $^{-1}$ hydrogels could sustain mechanical actuation for extended periods of time without loss of integrity (Fig. S3 †), we decided to conduct further experimentation using a fibroblast laden 3.0 mg mL $^{-1}$ collagen type I gel.

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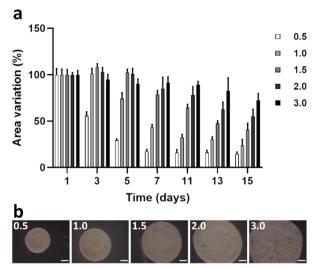


Fig. 4 ECM gel contraction study on fibroblast-laden hydrogels. (a) Contraction tested on a range of collagen type I densities, namely 0.5. 1.0, 1.5, 2.0 and 3.0 mg mL⁻¹ over a period of 15 days. Results graphed as % of area variation to original size (n = 3). (b) Visual representation of the ECM gel contraction assay. Images represent discs at day 15 post-seeding (scale bars: 500 µm).

2.5. A fibronectin thin gel mimicked the basement membrane

In vivo, the basement membrane anchors the epithelial layer, assisting in the maintenance of epithelial integrity and playing a crucial role in tissue compartmentalization. 37,38 Given its preponderant role in tissue maintenance and disease, 39,40 we aimed to reconstitute an anchoring substrate mimicking the in vivo gastric basement membrane. Here, we tested on a single-channel microfluidic chip several coating formulations, consisting of one of the major structural components of the gastric basement membrane, collagen type IV (at 4 mg mL⁻¹) and one accessory molecule, fibronectin (at 50 μg mL⁻¹), 41 that functions as a bridging molecule between the most abundant structural molecules, directly linking the ECM to the epithelial cells. Namely we tested a pure collagen type IV coating, mixtures of collagen type IV and fibronectin at 4:1 and 1:4 ratios and a pure fibronectin coating. Our results showed that MKN74 cells seeded over a thin coating of collagen type IV displayed an atypical phenotype, with a rounded shape. Over time, cells tended to remain isolated and exhibited cellular protrusions (Fig. 5a, panel 1). This suggested that cells were attempting to establish typical epithelial cell-to-cell contacts, but were unable to form a sheet-like epithelial monolayer. In contrast, when a mixture of higher fibronectin concentration was used as surface coating, MKN74 cells started exhibiting the typical epithelial phenotype, with cells displaying a cobblestone-like appearance and occupying the entire culture chamber surface (Fig. 5a, panel 2 and 3). This observation was more prominent when cells were seeded in a gel of pure fibronectin (Fig. 5a, panel 4), where cells displayed a normal phenotype, comparable to that of cells grown routinely in cell cultured-treated plastic vessels. We also quantified the number of cells showing a fully adherent phenotype vs. cells displaying an aberrant loosely adherent phenotype, 4 h postseeding (Fig. 5b). Our results showed that, on average, about 82% of the cells seeded over a fibronectin gel were fully adherent and displayed an epithelial phenotype. In contrast, when seeded in any of the other coating formulations, the percentage of adherent cells was reduced drastically. Populations seeded on pure collagen type IV and a 4:1 mixture of collagen type IV and fibronectin displayed ca. 90% of the population with a non-adherent rounded phenotype. Cells seeded on a 1:4 mixture of collagen type IV and fibronectin performed slightly better, with around 77% of the population remaining non adherent. Increasing the fibronectin proportion did show a gradual, even if not statistically significant, increase in the number of loosely adherent cells, displaying protrusions (Fig. 5b). It was also evident that the pure fibronectin anchoring substrate was well suited for the rapid development of a confluent monolayer under fluid flow conditions. Cells started adhering to the basement membrane as early as 30 min post-seeding forming an 80% confluent monolayer within 24 h (Fig. 5b), similarly to what was observed in Transwell inserts. Taking these observations in consideration, we opted to use a pure thin fibronectin coating as a basement membrane analogue, sacrificing the biomimetics of a mixed gel of collagen type IV and fibronectin, for a more functional adhesion substrate capable of performing well under flow conditions.

2.6. Stomach-on-a-chip device replicates the gastric epithelial barrier

The engineered stomach epithelium was housed within the upper portion of the biochip structure. The perforated membrane allowed communication between the adjacent fluidic structures, while ensuring that cells were kept within their designated confines, as was observed by staining the epithelial cells and the fibroblasts with cell tracker (-Fig. 6a, upper panel). Epithelial cells formed a tightly knit epithelium once confluency was reached (Fig. 6a, lower panel). Importantly, barrier function did not appear to be affected by on-chip culture, since we observed that on-chip, MKN74 cells correctly expressed and localized at the cell membrane, adherens-junctions partner E-cadherin and tight-junctions partner ZO-1 (Fig. 6b). To understand the contribution of the dynamic conditions, towards the establishment of in vitro gastric function, we quantified pepsin activity at day 7 in culture, under static vs. dynamic conditions (n = 3). Comparison with the Transwell static system (S), showed that a regimen of flow alone (F), was not sufficient to induce an increase in pepsin activity. On the other hand, when fluid flow was coupled with biomechanical actuation (F + MA) an increase in pepsin activity was observed (Fig. 6c). Here, we showed that mechanical stretching of the epithelial population induced a significant increase in pepsin activity, suggesting that mechanical actuation might be promoting

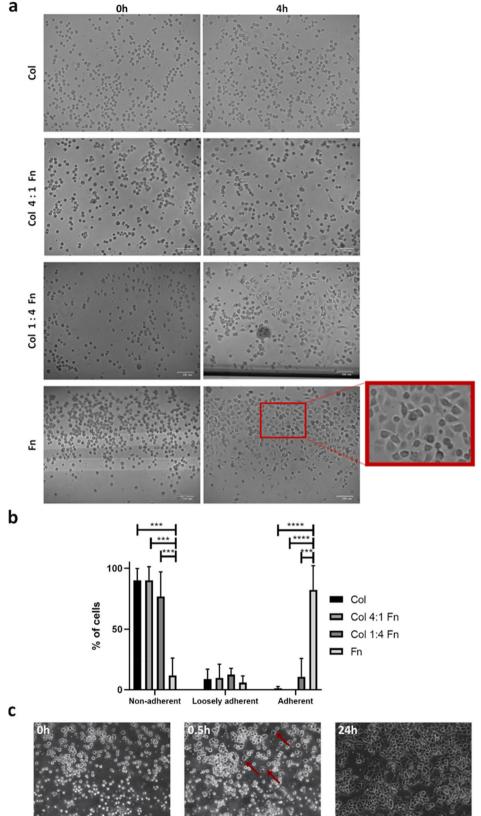


Fig. 5 Selection of a biomimetic basement membrane analogue (a) testing of different coating formulations to mimic the basement membrane. Substrate adhesion was assessed on pure collagen (CoI) type IV, a mixture at different ratios of collagen type IV and fibronectin (Fn), as well as a pure fibronectin coating. The red rectangle on the lower panel depicts a zoomed-in area highlighting the typical cobblestone-like pattern of MKN74 epithelial cells when seeded over a thin fibronectin gel (scale bar: 100 μm). (b) Graphical representation of the number of cells (%) 4 h post-seeding, displaying a rounded non-adherent phenotype, a loosely adherent phenotype or a fully adherent phenotype when cultured over the different substrate coatings. (c) MKN74 cells seeded on-chip, over a fibronectin coating at times 0, 0.5 and 24 h post-seeding (scale bars: 100 µm). MKN74 cells quickly developed stable anchoring when seeded on fibronectin, with cells attaching as fast as 30 min post-seeding (central panel). Red arrows highlight adherent cells.

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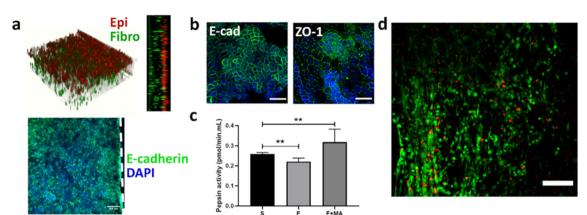


Fig. 6 Representative images of the cellular characterization of the stomach-on-a-chip (SoC). All IF images represent one of at least 3 replicates (a) 3D projection of the SoC biological barrier. On-chip long-term operation generated a fully developed epithelial barrier (stained in red with cell tracker). Below the epithelial layer, fibroblasts could be seen (stained in green with cell tracker) interspersed on a collagen type I matrix. Lower panel shows a typical on-chip, tightly knit epithelial culture at low magnification (scale bar: 200 μ m). Dashed vertical white line represents the channel's wall. (b) Epithelial barrier function was confirmed on-chip by staining MKN74 cells against adherens junction protein E-cadherin and tight junction protein ZO-1, evidencing fully functional cell-to-cell contacts and paracellular selectivity (scale bars: 50 μ m). Nuclei stained with DAPI (blue). (c) Pepsin activity study on static vs. dynamic conditions. Results reflected the conversion of the inactive precursor pepsinogen to the active form pepsin, by autocleavage under acidic conditions. Graph represents mean values \pm SD (n = 3). (d) Live-dead assay of on-chip NST-20 fibroblasts, with live cells stained in green and dead cells stained in red. On-chip viability was within acceptable levels and similar to that registered off-chip using fibroblast-laden collagen hydrogel discs, evidencing the efficiency of the chip design in providing correct feeding to the cells within its core (scale bar: 200 μ m). All images taken after 6 days of culture on-chip.

cytodifferentiation, similar to that observed in the native organ. Finally, as the fibroblast-laden hydrogel is at the inner core of the chip, we also qualitatively assessed the viability of the fibroblastic population growing on-chip by the live-dead assay. No major deviation was observed compared to our previous results obtained on off-chip collagen discs, with most of the fibroblastic population being viable within the hydrogel (Fig. 6d). This confirmed the efficiency of the lower perfusion compartment in providing correct feeding to the embedded fibroblasts.

2.7. Stomach-on-a-chip replicates gastric epithelial barrierfunction and selective permeability

Selective permeability to small molecules and drugs is a characteristic of epithelial interfaces, such as that of the human stomach. Therefore, we compared transport of a small 40 kDa molecule, namely fluorescein isothiocyanate (FITC)-labelled dextran, across the artificial epithelium to assess the impact of dynamic conditions, as well as of the engineered 3D architecture on paracellular transport across the epithelial barrier. Experimental conditions were measured on-chip for epithelial cells with culture media flow only (F), flow and mechanical actuation (F + MA), or the fully assembled SoC containing the epithelial and the fibroblastic populations and subjected to flow and mechanical actuation (SoC + F + MA). All of these conditions were compared to cells growing in static Transwell conditions (S). With this setup, we aimed to assess independently the contribution of each factor, namely the effect of flow, mechanical actuation and the on-chip emulation of the lamina propria, towards the establishment of organ level function within the chip.

Here, a solution of FITC-labelled dextran was either added to the apical side of the epithelium in the Transwell static condition or perfused over the apical side of the SoC at a flow rate of 2 µL min⁻¹. Samples were collected from the basal side chamber every 30 min, for a total period of 180 min. The Transwell inserts were slowly stirred for 5 s after each sample collection, to counteract the static nature of the Transwell system and to avoid the saturation of the epithelial barrier surface with FITC-labelled dextran molecules, which could limit diffusion across the barrier. Our observations showed a marked difference in paracellular permeability for cells growing under static conditions compared to dynamic conditions, with a significant difference already observable at 30 min (Fig. 7a). Flow alone (F) was sufficient to induce a marked decrease in the permeability of the gastric epithelium, in agreement to what we have previously reported, as flow alone was sufficient to induce alterations in the phenotype and molecular signature of these cells on a simplified microfluidic model of the gastric epithelium.²⁵ A similar permeability profile was observed for MKN74 cells under flow and mechanical actuation (F + MA), as well as for the fully assembled SoC (SoC + F + MA). In all conditions tested, the gastric layer was tightly bound and fully formed, as observed by E-cadherin staining (Fig. 7b). Overall, our results suggest that dynamic conditions play a significant role in the establishment of a tightly knit epithelium, displaying highly selective permeability across the epithelial barrier when compared to a static model. This is a relevant observation, as conventional gastric static layouts, used as diffusion models across the epithelial barrier, may be underestimating the contribution of paracellular selectivity across the barrier.

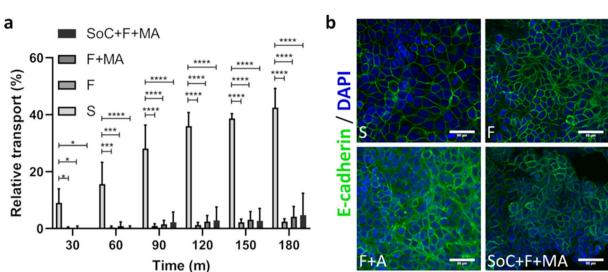


Fig. 7 Gastric permeability study by assessing the capability of a small molecule (FITC-dextran) to cross the epithelial barrier. (a) MKN74 epithelial cells grown without the fibroblastic population, either in Transwell inserts under static conditions (S), on-chip under flow (F) or flow coupled with mechanical actuation (F + MA). The condition SoC + F + MA refers to the fully assembled stomach-on-a-chip (SoC). Results shown as relative transport (%) over time. Graph represents mean values \pm SD (n = 3). (b) Immunocytochemistry against E-cadherin (green), denoting the formation of a cohesive epithelium in all conditions tested (scale bar: 50 μ m). Given the 3D epithelial architecture acquired by the mechanically actuated SoC, portions of the artificial epithelium appear out of focus. Images are representative of the entire surface of the culture epithelium.

Finally, we aimed to characterize the phenotypic and molecular traits developed by the SoC gastric epithelium under dynamic conditions. As reported above, MKN74 cells formed a selective barrier after 2 days in culture when seeded at a density of 3.0×10^5 cells per mL. Taking this in consideration, the MKN74 cell population was seeded onchip and allowed to reach confluency over a period of 3 days, followed by 3 days under flow regime and mechanical stretching of the cell culture substrate. Upon application of mechanical stretching and fluid flow, the SoC epithelium thickened considerably, reaching an average thickness of 38.5 µm and globular-like structures developed across its surface (Fig. 8a, panel 2 and 3). This is in sharp contrast to what was observed in static MKN74 cultures grown in Transwell inserts, where cells formed a flattened monolayer similar to a squamous epithelium (Fig. 8a, panel 1). Accordingly, the bioengineered SoC epithelium acquired a polarized state, with cells displaying a columnar shape with elongated nuclei located at the basal side and E-cadherin expression exclusively across the basolateral region. This phenotype closely resembled a pseudostratified columnar epithelium, similar in many aspects to its in vivo counterpart. Concurrently, expression of mucin-1 could be observed across the apical surface of the globular structures (Fig. 8b). This is in accordance to what can be observed in the native stomach⁴² and as previously reported for other MKN74 3D gastric models.^{25,43,44} Overall, our observations showed that the bioengineered device closely mimicked the gastric architecture and microenvironment by reproducing the three innermost-layers of the gastric epithelium. Furthermore, the peristalsis-like actuation and intra-luminal flow simulation seemed capable of producing physiologically relevant stimuli,

inducing the MKN74 cell line, an otherwise moderately differentiated cell line that typically forms a squamous-like epithelium, to undergo further differentiation and acquire phenotypical characteristics typical of the normal gastric epithelium. Importantly, the introduction mesenchymal population to the stomach-on-a-chip device and the concomitant epithelial-mesenchymal cross-talk, did not affect the development of the differentiated state of the artificial gastric epithelium. The engineered SoC was not only capable of replicating the gastric mucosa architecture, by displaying a degree of differentiation and polarization characteristic of the native organ, but allowed also the partial reproduction of gastric function, as shown by enhanced barrier function. Here, we chose to represent the fibroblastic population, typically found within the gastric lamina propria and essential for tissue homeostasis. However, the lamina propria's cellular and molecular milieu is diverse and other aspects could be easily studied. Components of the immune system could be added to study response to induced inflammation or to tumor initiation. Likewise, an endothelial interface could be established on the lower perfusion channel, thus increasing the complexity of the system to represent the vasculature of the lamina propria, a model that would be well suited to study tumor invasion and intravasation.

3. Concluding remarks

Organ-on-a-chip devices enable a faithful replication of the tissue microenvironment, reproducing both architecture and *in vivo* dynamic conditions in one single device to potentiate the development of organotypic function. In the present



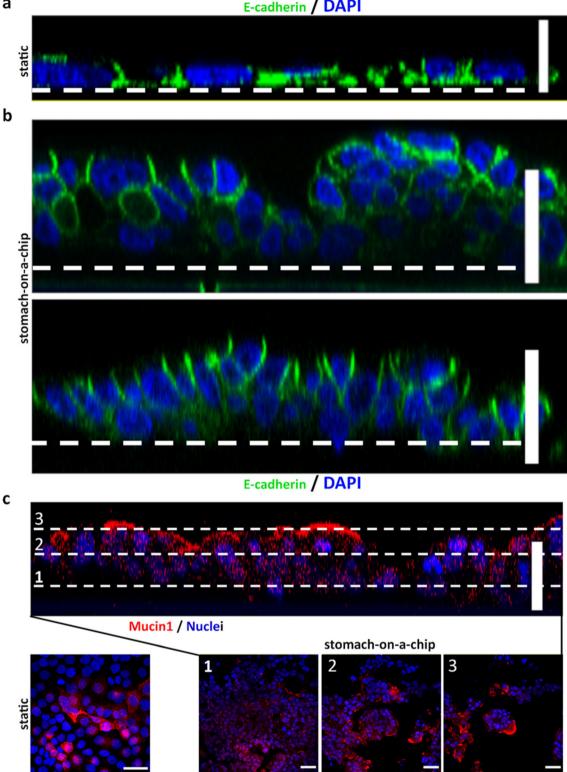


Fig. 8 Comparison study of gastric epithelium under static conditions vs. stomach-on-a-chip (SoC). Cells were analyzed after 6 days, either in static or dynamic conditions. Mechanical actuation was only applied on the last 3 days of the 6-day period. (a) MKN74 cells grown in Transwell inserts under static conditions displayed a flattened, undifferentiated appearance, similar to a squamous epithelium (top panel; scale bar: 20 µm). (b) SoC grown epithelium developed globular-like structures over time (upper panel) and cells acquire an elongated, polarized epithelium with E-cadherin (green) expression exclusively on the basolateral side and nuclei in the basal position, resembling a pseudostratified epithelium (lower panel) (scale bars: 50 µm). (c) Mucin-1 staining of SoC grown epithelium. Orthogonal projection of an on-chip epithelium grown under dynamic conditions (scale bar: 50 µm). Lower panels represent the same microscopic section captured at 3 different z-heights (scale bars: 100 µm), displaying predominant apical expression of mucin-1. Lower left panel shows a MKN74 population grown under static conditions and stained for mucin-1 (scale bar: 50 µm) displaying random cytoplasmatic expression. Nuclei stained with DAPI (blue). All immunofluorescence images are representative of at least 3 independent experiments.

work, we designed and fabricated a stomach-on-a-chip (SoC) device replicating both architectural and functional traits of the native organ. Our model could sustain a long-term 3D co-culture of epithelial and mesenchymal cells, without deleterious effects on cell homeostasis and viability. Importantly, we have demonstrated its capability for eliciting organ-level response, where cells developed phenotypic, molecular and functional traits characteristic of the normal gastric mucosa, that are usually not observable in 2D cultures. These observations outline the relevance of faithfully replicating microenvironment of the gastric mucosa and, particular, the importance of mimicking the dynamic nature of the native organ, in order to engineer a biomimetic analogue. Unlike the organoid-on-a-chip gastric system previously reported by Lee and colleagues, our stratified design allows easy access to both the apical and the basal side of the gastric epithelium, presenting a versatile model that can be used to study or simulate

The presently reported SoC constitutes a valuable tool to study gastric absorption dynamics, as well as host-pathogen interplay and directed-targeting therapies.

both luminal and systemic infection of the gastric

mucosa. A further advantage of the currently proposed

system is the use of a biomimetic supporting extracellular

matrix (ECM), as a lamina propria analogue.

In summary, the low cost of the fabrication method coupled with the robust physiological response that the system elicits, makes the present SoC device a functional microfluidic platform to model the dynamic gastric microenvironment.

4. Experimental

Design

The conceived stomach-on-a-chip (SoC) consisted of 9 intercalated parts (Fig. 1a). From top to bottom, the top 6 layers formed the fluidic portion of the biochip. Layers 7 and 8 composed the microactuator and the bottom layer was a microscope slide. Polydimethylsiloxane (PDMS) microchannels were designed with a 10 mm (length) × 2 mm (width) main rectangular chamber and access holes measuring 2 mm in diameter (Fig. 1b). Each fluidic layer was separated by a 16 µm thick perforated polyethylene terephthalate (PET: it4ip) membrane with pore size of 8 µm and pore density of 6×10^4 cm². The microactuator was a two-part assembly, composed of a 250 µm flexible PDMS membrane on top of a vacuum chamber. The vacuum chamber was 250 μ m-thick and 11 mm (length) \times 3.5 mm (width). The vacuum chamber was larger than the fluidic channels, thus ensuring proper actuation of the entire cell culture surface (Fig. 1c). The whole structure was assembled on top of a microscope glass slide and, for that reason, each structure was centered on a 26 mm square PDMS laminate, to match the width of the microscope slide.

Fabrication

The SoC microdevice was fabricated from pre-cured PDMS laminated sheets and PET porous membranes using a method previously developed by our group.²⁵ Briefly, the device was designed using computer aided design (CAD) software, exported to a cutting plotter (model CAMM-1 GS-24; Roland DG) and machined onto pre-cured 915 mm wide PDMS sheets (MVQ silicones). The top layer was fabricated by gravity casting. Prior to assembly, every material was thoroughly cleaned. Machined PDMS plates were washed with isopropanol (IPA) and air dried with compressed air. PET membranes were immersed in IPA, incubated in an ultra-sound bath for 15 min and dried with pressurized air. Microscope slides were incubated sequentially (5 min by ultrasound) in 2% (v/v) Hellmanex III (Helma Analytics) solution in double distilled (DD) water, followed by acetone and a final rinse in DD water. Clean slides were dried with compressed air and clean materials were either used immediately or stored in a dust-free environment.

Assembly of the chip was performed sequentially from top to bottom (Fig. 1a). PDMS plates and glass slides were bonded by exposure to O2 plasma for 1 min (Zepto Plasma Laboratory Unit, Diener). Structures were then brought into conformal contact, placed between two glass slides and pressure was applied overnight. The outline of the microchannels in each layer was used as a guide to puncture access inlet and outlet holes through the top layer, with a 20 ga luer stub (Instech). To integrate the PET membranes, silanization of the PET surface was required. Following the cleaning procedure, both sides of the membrane were exposed to O2 plasma and immersed in 2% (v/v) bis-amino silane, 1% (v/v) DD water in IPA solution (20 min at 80 °C), followed by thorough cleaning with IPA and cured for 30 min at 70 °C. Membranes were then immersed for 30 min in 70% ethanol (v/v). Prior to bonding, both sides of the PDMS structures were exposed to O2 plasma and the wetted membrane was immediately sandwiched between them. This sub-assembly was further cured for 1 h at 70 °C.

Cell culture

The human-derived gastric cancer cell lines AGS and MKN74 were available at i3S/Ipatimup's cell line repository. For the SoC, the human-derived adenocarcinoma epithelial cell line, MKN74 was used. The human stomach normal fibroblast cell line NST-20 was a kind gift of Dr. Luis Filipe Santos Silva (i3S/Ipatimup, Portugal).

Cell lines were routinely maintained at 37 °C and 5% CO₂, under a humidified atmosphere and fed with RPMI 1640 (Gibco) medium, supplemented with 10% (v/v) fetal bovine serum (FBS; Biowest) and 1% (v/v) penicillin/streptomycin (pen/strep; Biowest). Cells were reseeded when confluency reached about 80% and tested for the presence of mycoplasma when thawed and every month when in culture.

Before on-chip seeding, MKN74 and NST-20 cells were conditioned for at least 48 h in CO_2 -independent cell culture

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medium (Gibco), supplemented with 4 mM of L-glutamine (Gibco), 10% (v/v) FBS (Biowest) and 1% (v/v) pen/strep (Biowest) in a humidified normal atmosphere (*i.e.*, without CO_2 buffering) at 37 °C.

Chip priming was started by sterilization with UV radiation (20 min), followed by flushing with 70% ethanol (20 min). Finally, the devices were rinsed with phosphate buffered saline (PBS; 3 \times 15 min; Fisher BioReagents). Following sterilization, the fluidic portion was incubated with a 50 μg mL⁻¹ fibronectin solution (Sigma) for 2 h at room temperature (RT).

NST-20 fibroblasts were embedded in a collagen type I gel. For that purpose, cells were harvested by trypsinization and suspended in a solution of 3 mg mL $^{-1}$ collagen type I to a final density of 200 000 cells per mL. The resulting suspension was carefully injected into the middle-tier microchannel and incubated for 30 min at 37 °C to allow gelification. The device was then coupled to a piezoelectric controller νia polyether ether ketone (PEEK) tubing (Idex) and CO₂-independent culture medium was perfused at a flow rate of 2 μ L min $^{-1}$. The system was allowed to equilibrate overnight, with the microactuator set to -50 mbar constant vacuum pressure to allow degassing of potential microbubbles present in the culture. 25

To replicate the epithelial barrier, MKN74 cells were collected by trypsinization and resuspended to a density of 3.0×10^5 cells per mL. This suspension was then injected via a 4-way PEEK valve onto the upper fluidic channel. Cell seeding density was controlled under a phase contrast microscope and culture media flow was reinstated in both top and bottom fluidic channels at a flow rate of 2 μ L min⁻¹, 1 h after cell seeding. Cells were allowed a 72 h growth and stabilization period after which mechanical stretching was effected by application of cyclic vacuum. To emulate peristalsis motion experienced by gut cells *in vivo*, the stretching pattern was modelled as a sinusoidal wave of 0.15 Hz frequency and maximum vacuum pressure at -100 mbar.

Cell tracker staining

To monitor cell loading and compartmentalization on the SoC, MKN74 epithelial cells and NST-20 fibroblasts were stained with CellTracker™ green CMFDA and CellTracker™ red CMPTX dyes, respectively. Staining was performed according to the manufacturer's instructions. Briefly, cells were harvested and suspended in CellTracker staining solution (1:1000 dilution) for 30 min at 37 °C. Staining solution was then removed and cell seeding and/or embedding was performed as described above. Epithelial layer was grown to confluency after which the chip was imaged under a confocal microscope (model SP5; Leica Microsystems).

Morphological characterization

For immunofluorescence imaging, cells were washed in PBS with $0.05~{\rm mg~mL^{-1}}$ of ${\rm CaCl_2}$ (3 \times 5 min), fixed

with 4% (v/v) paraformaldehyde suspended in PBS (20 min at RT) and thoroughly washed with PBS (3 × 5 min). Cells were further incubated in 50 mM NH₄Cl (10 followed permeabilization of cellular min), by membranes with 0.2% (v/v) Triton X-100 in PBS (5 min). Blocking was achieved with 5% (v/v) bovine serum (BSA; solution in PBS. albumin Sigma) antibodies used were rabbit anti-E-cadherin (24E10; dilution 1:100; Cell Signaling), rabbit anti-ZO1 (dilution ThermoFisher Scientific), rabbit anti-ocludin (dilution 1:50; Santa Cruz), rabbit anti-collagen I (dilution 1:100; Rockland), mouse anti-collagen IV (dilution 1:100; Dako), and rabbit anti-fibronectin (dilution 1:150; Sigma). Primary antibodies were diluted in 5% (v/v) BSA suspended in PBS and incubated at 4 °C overnight. Secondary antibodies were diluted in 5% (v/v) BSA diluted in PBS and incubation was performed in the dark for 2 h at RT. Antibodies used were antirabbit Alexa Fluor 488 and/or anti-mouse Alexa Fluor Thermo (dilution 1:500; Scientific). Cellular preparations were mounted in Vectashield containing 2-(4-amidinophenyl)-1H-indole-6-carboxamidine (DAPI) for nuclei counterstaining. Image acquisition was performed with a confocal microscope as before.

Characterization of a lamina propria analogue and analysis of ECM deposition

To emulate the lamina propria, a collagen type I (5 mg mL⁻¹; Ibidi) hydrogel embedded with normal stomach fibroblasts (NST-20 cell line) was used. A range of collagen type I densities was tested, namely 0.5, 1.0, 1.5, 2.0 and 3.0 mg mL⁻¹. Cell embedding and gelification of hydrogels performed according to the manufacturer's recommendations. Briefly, NST-20 cells were trypsinized and resuspended to a final cellular density of 2×10^5 cells per mL. After combining the fibroblasts with the collagen type I, 30 µL per gel of the resulting suspension were plated on the bottom of a flat-faced well of a 24-well plate well (Corning). To trigger gelification and ensure cell viability, fibroblast-embedded hydrogels were incubated for 30 min at 37 °C under a humidified, normal (without CO₂) atmosphere. After gelification, 1 mL of CO₂independent medium (supplemented as described above) was added to each well and gel contraction was monitored for 15 days. Discs were photographed every other day under a stereomicroscope (Olympus SZX10) coupled with a camera (Olympus EP50) and disc perimeter was measured using image analysis software (Fiji). The results were plotted as average percent of size difference over time, normalized to initial disc size (day 0). ECM remodeling was studied by immunofluorescence. Fibroblast-embedded discs were collected at day 11 postseeding and stained with primary antibodies against collagen type I, collagen type IV or fibronectin. Image acquisition was done in a confocal microscope as before.

Metabolic activity of collagen-embedded fibroblasts

Metabolic activity was assessed by the resazurin assay. NST-20 cells were embedded in collagen type I discs as described before. Metabolic activity was assessed after the first 24 h in culture and every 48 h for 15 days, for which a stock solution of 0.1 mg mL⁻¹ resazurin (Sigma) was diluted to 20% (v/v; resazurin/cell culture media). Culture media was replaced with 500 μL of this solution and incubated at 37 °C for 2 h. 200 µL of the resulting supernatant were transferred to an opaque 96-well plate with clear bottom (Greiner) and the fluorescence signal was measured (excitation 530 nm/ emission 590 nm) in a multiwell plate reader (model Sinergy MX HM550; Biotek Instruments). Metabolic activity was expressed as average relative fluorescence units (RFUs) ± standard deviation (SD).

Cell viability assay

To assess viability of the NST-20 fibroblasts embedded on a collagen type I gel, a live dead assay was performed. Briefly, a staining solution containing 2 µg mL⁻¹ of calcein and 2.5 µg mL⁻¹ of propidium iodide was prepared. Cells were washed thrice with unsupplemented culture medium, followed by a 20 min incubation at 37 °C with staining solution. Finally, cells were washed twice with unsupplemented culture medium and immediately imaged under a confocal microscope as before. Z-stacks were acquired with a 2 µm z-step and post-processed using image analysis software (Fiji).45

Epithelium characterization

The on-chip formed epithelium was fixed and stained with an antibody against E-cadherin as described above. Z-stacked fluorescence images were acquired with a 2 µm z-step value, using a confocal microscope as before. Z-stacks were postprocessed and epithelium height measured using image analysis software (Fiji). For 3D rendering of the stomach epithelium, z-stacks were processed using Fiji's 3D viewer plugin.

Paracellular permeability

To measure paracellular permeability, transport of fluorescein isothiocyanate (FITC)-labelled dextran (40 kDa; Sigma) across the epithelium was assessed over time. For onchip measurements, cells were grown to confluency and cultured under cell culture media flow (2 µL min⁻¹) with or without exposure to cyclic mechanical actuation for 3 days, at a 0.15 Hz frequency. FITC-dextran solution (1 mg mL⁻¹) was perfused through the top channel and 30 µL samples were collected at the lower channel outlet every 30 min, for a period of 3 h. Samples were analyzed in a multiwell plate reader as before at 488 nm excitation and 520 nm emission. The conditions flow (F) and flow coupled with actuation (F + A) were assessed in a double channel chip, without the addition of the lamina propria analogue. Thus, the contribution of each condition was estimated independently. Static conditions were studied using Transwell inserts (Corning), coated with fibronectin (50 µg mL⁻¹). Cells were allowed to reach confluency and grown for 3 additional days. Permeability was assessed by adding 200 µL of 1 mg mL⁻¹ FITC-dextran solution to the apical side of the Transwell and collecting 50 µL samples from the basal side, while replenishing the same volume with fresh medium. Samples were collected up to 3 h and analyzed in a multiwell plate reader as before at 488 nm excitation and 520 nm emission. In both dynamic and static conditions, permeability was assessed as the relative transport of small molecules across the epithelial barrier and expressed as the fluorescence value collected at the basal channel, normalized to the fluorescence signal of the solution added to the apical side. Results were plotted as average percentage of FITC-dextran release against elapsed time (n = 3).

Pepsin/pepsinogen activity

Gastric epithelial function was measured by quantifying the specific activity of pepsin/pepsinogen. A pepsin/pepsinogen activity assay kit (Biovision) was used. The procedure was performed according to the manufacturer's instructions. Briefly, cells were grown for 3 days post-confluency, either in Transwell inserts (static conditions) or on-chip with culture media flow and in the presence or absence of cyclic mechanical stretching (0.15 Hz frequency and -100 mbar vacuum pressure). Samples (100 µL) were collected from the basal side and 20 µL were added to 80 µL of pepsin assay buffer. A pepsin substrate working solution (pepsin substrate stock solution and pepsin assay buffer at a ratio of 1:10 (v/v), was then added to the previous mixture to start the reaction. Fluorescence was measured in a multiwell plate reader as before at 328 nm excitation and 418 nm emission in kinetic mode for 1 h at 37 °C. Sample pepsin activity was calculated according to the following equation:

Sample pepsin activity = $B/(\Delta T \times P)$

where B is the amount of peptide substrate cleaved, calculated from the standard curve, ΔT is the linear phase reaction time $t_2 - t_1$ (min) and P is the amount of sample (mL). Results are expressed as pmol min⁻¹ mL⁻¹.

Statistical analysis

Data was compiled and analyzed using the software Graphpad Prism (Graphpad software). Data plotted in graphical format is presented as mean value ± SD. Cell culture doubling time and metabolic rates (n = 6) were compared using a two-way ANOVA statistical test. The latter were analyzed in conjunction with a Sidak's post hoc multiple comparison test. Relative transport across the paracellular route and cell adhesion assay, were analyzed using a two-way ANOVA in conjunction with Tukey's multiple comparison test. Pepsin activity mean values were compared using a nonparametric Mann-Whitney U test (n = 3). Statistical significance was defined as p < 0.05 (*), p < 0.01 (***), p < 0.001 (****).

Author contributions

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Daniel A. Ferreira: conceptualization, investigation, writing – original draft, methodology. João P. Conde: writing – review & editing, funding acquisition, supervision. Mario Rothbauer: writing – review & editing, methodology. Peter Ertl: writing – review & editing, funding acquisition, supervision. Pedro L. Granja: writing – review & editing, funding acquisition, project administration, supervision, conceptualization. Carla Oliveira: writing – review & editing, funding acquisition, project administration, supervision, conceptualization.

Conflicts of interest

Authors declare no conflict of interest.

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Notes and references

- 1 R. H. Hunt, M. Camilleri, S. E. Crowe, E. M. El-Omar, J. G. Fox, E. J. Kuipers, P. Malfertheiner, K. E. McColl, D. M. Pritchard, M. Rugge, A. Sonnenberg, K. Sugano and J. Tack, *Gut*, 2015, **64**, 1650–1668.
- 2 M. H. Ross and W. Pawlina, *Histology : a text and atlas*, Lippincott Williams & Wilkins, Philadelphia, Pa., London, 6th edn, 2011.
- 3 P. J. Kumar and M. L. Clark, *Kumar & Clark's clinical medicine*, Saunders Elsevier, Edinburgh, 7th edn, 2009.
- 4 H. Sung, J. Ferlay, R. L. Siegel, M. Laversanne, I. Soerjomataram, A. Jemal and F. Bray, *Ca-Cancer J. Clin.*, 2021, 71, 209–249.
- 5 S. A. Langhans, Front. Pharmacol., 2018, 9, 6.
- 6 J. Arrowsmith and P. Miller, Nat. Rev. Drug Discovery, 2013, 12, 569.
- 7 F. Pampaloni, E. G. Reynaud and E. H. Stelzer, *Nat. Rev. Mol. Cell Biol.*, 2007, 8, 839–845.
- 8 R. Edmondson, J. J. Broglie, A. F. Adcock and L. J. Yang, Assay Drug Dev. Technol., 2014, 12, 207–218.
- 9 N. Barker, M. Huch, P. Kujala, M. van de Wetering, H. J. Snippert, J. H. van Es, T. Sato, D. E. Stange, H. Begthel, M. van den Born, E. Danenberg, S. van den Brink, J. Korving, A. Abo, P. J. Peters, N. Wright, R. Poulsom and H. Clevers, *Cell Stem Cell*, 2010, **6**, 25–36.
- 10 T. Grikscheit, A. Srinivasan and J. P. Vacanti, *J. Pediatr. Surg.*, 2003, 38, 1305–1309.
- 11 T. Sato, R. G. Vries, H. J. Snippert, M. van de Wetering, N. Barker, D. E. Stange, J. H. van Es, A. Abo, P. Kujala, P. J. Peters and H. Clevers, *Nature*, 2009, **459**, 262–265.
- 12 A. L. Speer, F. G. Sala, J. A. Matthews and T. C. Grikscheit, *J. Surg. Res.*, 2011, **171**, 6–14.
- 13 F. Boccellato, S. Woelffling, A. Imai-Matsushima, G. Sanchez, C. Goosmann, M. Schmid, H. Berger, P. Morey, C. Denecke, J. Ordemann and T. F. Meyer, *Gut*, 2019, 68, 400–413.
- 14 B. N. Lourenco, T. Dos Santos, C. Oliveira, C. C. Barrias and P. L. Granja, *Acta Biomater.*, 2018, **82**, 68–78.
- 15 M. Bashashati and R. A. Hejazi, *Dig. Dis. Sci.*, 2018, 63, 3164–3166.
- 16 I. R. Sanderson, Am. J. Clin. Nutr., 1999, 69, 1028S-1034S.
- 17 T. Mammoto, A. Mammoto and D. E. Ingber, *Annu. Rev. Cell Dev. Biol.*, 2013, **29**, 27–61.
- 18 C. T. Lim, A. Bershadsky and M. P. Sheetz, *J. R. Soc., Interface*, 2010, 7, S291–S293.
- 19 O. T. Guenat and F. Berthiaume, Biomicrofluidics, 2018, 12, 042207.
- 20 C. Moraes, M. Likhitpanichkul, C. J. Lam, B. M. Beca, Y. Sun and C. A. Simmons, *Integr. Biol.*, 2013, 5, 673–680.

21 D. Huh, B. D. Matthews, A. Mammoto, M. Montova-Zavala, H. Y. Hsin and D. E. Ingber, Science, 2010, 328, 1662-1668.

- 22 H. J. Kim, D. Huh, G. Hamilton and D. E. Ingber, Lab Chip, 2012, 12, 2165-2174.
- 23 A. O. Stucki, J. D. Stucki, S. R. Hall, M. Felder, Y. Mermoud, R. A. Schmid, T. Geiser and O. T. Guenat, Lab Chip, 2015, 15, 1302-1310.
- 24 M. Wufuer, G. Lee, W. Hur, B. Jeon, B. J. Kim, T. H. Choi and S. Lee, Sci. Rep., 2016, 6, 37471.
- 25 D. A. Ferreira, M. Rothbauer, J. P. Conde, P. Ertl, C. Oliveira and P. L. Granja, Adv. Sci., 2021, 8, 2003273.
- E. Ergir, B. Bachmann, H. Redl, G. Forte and P. Ertl, Front. Physiol., 2018, 9, 1417.
- 27 K. K. Lee, H. A. McCauley, T. R. Broda, M. J. Kofron, J. M. Wells and C. I. Hong, Lab Chip, 2018, 18, 3079-3085.
- 28 X. He, B. Lee and Y. Jiang, Adv. Exp. Med. Biol., 2016, 936, 73-91.
- 29 W. P. Daley, S. B. Peters and M. Larsen, J. Cell Sci., 2008, 121, 255-264.
- 30 C. P. Gayer and M. D. Basson, Cell. Signalling, 2009, 21, 1237-1244.
- 31 M. D. Basson, Digestion, 2003, 68, 217-225.
- 32 T. Ishikawa, T. Sato, G. Mohit, Y. Imai and T. Yamaguchi, J. Theor. Biol., 2011, 279, 63-73.
- 33 A. Gillessen, B. Voss, J. Rauterberg and W. Domschke, Scand. J. Gastroenterol., 1993, 28, 688-689.
- 34 Z. N. Wang and H. M. Xu, World J. Gastroenterol., 2000, 6, 438-439.

- 35 E. Bell, B. Ivarsson and C. Merrill, Proc. Natl. Acad. Sci. U. S. A., 1979, 76, 1274-1278.
- 36 T. Zhang, J. H. Day, X. Su, A. G. Guadarrama, N. K. Sandbo, S. Esnault, L. C. Denlinger, E. Berthier and A. B. Theberge, Front. Bioeng. Biotechnol., 2019, 7, 196.
- 37 A. Pozzi, P. D. Yurchenco and R. V. Iozzo, Matrix Biol., 2017, 57-58, 1-11.
- 38 T. Rozario and D. W. DeSimone, Dev. Biol., 2010, 341, 126-140.
- 39 R. Sekiguchi and K. M. Yamada, Curr. Top. Dev. Biol., 2018, 130, 143-191.
- A. M. Moreira, J. Pereira, S. Melo, M. S. Fernandes, P. Carneiro, R. Seruca and J. Figueiredo, Cells, 2020, 9, 394.
- 41 A. C. Y. Li and R. P. H. Thompson, J. Clin. Pathol., 2003, 56, 885-887.
- 42 D. Boltin and Y. Niv, J. Gastrointest. Dig. Syst., 2013, 3, 15519.
- 43 S. Rocha, J. Carvalho, P. Oliveira, M. Voglstaetter, D. Schvartz, A. R. Thomsen, N. Walter, R. Khanduri, J. C. Sanchez, A. Keller, C. Oliveira and I. Nazarenko, Adv. Sci., 2019, 6, 1800948.
- 44 M. Balmana, S. Mereiter, F. Diniz, T. Feijao, C. C. Barrias and C. A. Reis, Molecules, 2018, 23, 2815.
- 45 J. Schindelin, I. Arganda-Carreras, E. Frise, V. Kaynig, M. Longair, T. Pietzsch, S. Preibisch, C. Rueden, S. Saalfeld, B. Schmid, J. Y. Tinevez, D. J. White, V. Hartenstein, K. Eliceiri, P. Tomancak and A. Cardona, Nat. Methods, 2012, 9, 676-682.