REVIEW



Engineering biomimetic intestinal topological features in 3D tissue models: retrospects and prospects

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Abstract

Conventional 2D intestinal models cannot precisely recapitulate biomimetic features in vitro and thus are unsuitable for various pharmacokinetic applications, development of disease models, and understanding the host-microbiome interactions. Thus, recently, efforts have been directed toward recreating in vitro models with intestine-associated unique 3D crypt-villus (for small intestine) or crypt-lumen (for large intestine) architectures. This review comprehensively delineates the current advancements in this research area in terms of the different microfabrication technologies (photolithography, laser ablation, and 3D bioprinting) employed and the physiological relevance of the obtained models in mimicking the features of native intestinal tissue. A major thrust of the manuscript is also on highlighting the dynamic interplay between intestinal cells (both the stem cells and differentiated ones) and different biophysical, biochemical, and mechanobiological cues along with interaction with other cell types and intestinal microbiome, providing goals for the future developments in this sphere. The article also manifests an outlook toward the application of induced pluripotent stem cells in the context of intestinal tissue models. On a concluding note, challenges and prospects for clinical translation of 3D patterned intestinal tissue models have been discussed.

 $\textbf{Keywords} \ \ Intestine \ tissue \ models \cdot Microfabrication \cdot Biophysicochemical \ and \ biomechanical \ cues \cdot Coculture \cdot Induced \ pluripotent \ stem \ cells$

Abbreviations

2D	Two-dimensional
3D	Three-dimensional
SI	Small intestine

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- LI Large intestine
- CBC Crypt base columnar cells
- TA Transit-amplifying
 M cell Microfold cell
 UV Ultraviolet
- PDMS Polydimethylsiloxane
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PTFE	Polytetrafluoroethylene
PEG	Poly(ethylene glycol)
AA	Acrylic acid
ECM	Extracellular matrix
UV-LIGA	Ultraviolet-lithography, electroplating, and
C V EIGIT	molding
PLA	Poly(lactic acid)
CVD	Chemical vapor deposition
PMMA	Poly(methyl methacrylate)
PLGA	Poly(lactic-co-glycolic acid)
PEVA	Poly-ethylene-co-vinyl-acetate
CAD	Computer-aided design
	÷
PEGDA	Poly(ethylene glycol) diacrylate
VMEPS	Vertically moving extrusion-based printing
HINEC	system
HUVECs	Human umbilical vein endothelial cells
TEER	Transepithelial electrical resistance
MUC17	Mucin 17
RT-PCR	Reverse transcription polymerase chain reaction
FITC	Fluorescein isothiocyanate
PCL	Poly-ε-caprolactone
ZO-1	Zonula occludens-1
P-gp	P-Glycoprotein
ALP	Alkaline phosphatase
CYP3A4	Cytochrome P450 3A4
EdU	5-Ethynyl-2'-deoxyuridine
Olfm4	Olfactomedin 4
CK20	Keratin 20
MUC2	Mucin 2
E-cad	E-cadherin
ISC	Intestinal stem cell
RGD	Arginine-glycine-aspartate
GAGs	Glycosaminoglycans
Wnt	Wingless-related integration site
TGF-β	Transforming growth factor beta
FGF	Fibroblast growth factors
LGR5	Leucine-rich repeat-containing G-protein
	coupled receptor 5
IFN-γ	Interferon gamma
TNF-α	Tumor necrosis factor alpha
YAP	Yes-associated protein 1
BCRP	Breast cancer resistance protein
MRP2	Multidrug resistance protein 2
iPSCs	Induced pluripotent stem cells
STAT1	Signal transducer and activator of transcrip-
ENIC	tion 1
ENS MID 2	Enteric nervous system Magraphaga inflammatory protein 2
MIP-2	Macrophage inflammatory protein 2
IL-10	Interleukin 10
ISEMFs	Intestinal subepithelial myofibroblasts
SCFAs	Short-chain fatty acids
ELCs	Enterocytes-like cells

DELCs	Definite endodermal-like cells
IPLCs	Intestinal progenitor-like cells
HLCs	Hindgut-like cells
EGF	Epidermal growth factor
5-aza	5-Aza-2'-deoxycytidine
BIO	6-Bromoindirubin-3'-oxime
DAPT	N-[(3,5-difluorophenyl)acetyl]-L-alanyl-
	2-phenyl-1,1-dimethylethyl ester-glycine
WRN	Wnt3A,R-spondin,Noggin
PEPT1	Peptide transporter 1
SIOs	Small intestinal organoids
Cos	Colonic organoids
BMPs	Bone morphogenetic proteins
SATB2	Special AT-rich sequence-binding protein 2
HOX	Homeobox
WDR43	WD Repeat Domain 43
TALEN	Transcription activator-like effector nuclease
CFTR	Cystic fibrosis transmembrane conductance
	regulator

Inflammatory bowel disease

Introduction

IBD

The intestine is one of the most vital metabolic organs involved in digestion and absorption of nutrients [1–3]. Its unique microarchitecture accompanied by the peristaltic contractions provides a much required large surface area to aid nutrient absorption [2–4]. Owing to its constant contact with gut microbiome and xenobiotics components, it is at high risk of various pathological disorders [5–7]. Thus, to find a potential treatment for these disorders, a clear understanding of their mechanistic aspects is necessary.

Animal models are still the gold standard clinical models to study intestinal anatomy, physiology, pathologies and xenobiotic-mediated tissue responses [8, 9]. However, the underlying differences with the human intestinal anatomy, gut microbiota, immunological activities, expense of animal maintenance, and ethical considerations, further limit their usage [8, 10]. Therefore, the current thrust of the healthcare and diagnostic sector is on the development of functional in vitro intestinal tissue models.

Up to date, various intestinal models, including 2D/3D or dynamic microfluidic chip-based, have been proposed and developed [1, 11]. However, limited physiological relevance of these models, in terms of topological and microenvironmental features, makes them far from actually recapitulating the complexity of the native intestinal tissue, and thus, are unsuitable for clinical applications [1, 11–14].

In this regard, decellularized intestinal submucosal tissue holds the most promising prospects as it retains, to an extent, native-like architectural and matrix's biochemical characteristics [15, 16]. Owing to these aspects, decellularized



scaffolds have generated a great wave of enthusiasm, mainly in the domain of regenerative medicines and in vitro model development [17, 18]. But their applicability is often limited by a low tissue availability, tedious non-standardized decellularization procedure, damage to the native ECM (microstructural, biochemical, and mechanical), difficult handling, and low experimental reproducibility [15, 17, 18].

Consequently, a current focus of research fraternity is on devising strategies for patterning intestine-like topologies on 2D substrates. These patterned substrates could be employed for developing biomimetic in vitro 3D tissue models. Advanced microfabrication techniques, including soft lithography, laser-induced ablation, and 3D (bio) printing, have paved the way for the same [1, 11]. In fact, cells cultured on these models exhibit more differentiated phenotype, compared to those on other conventional models [19–22]. To the best of our knowledge, to date, there exists no review that critically highlights the patterned 3D intestinal tissue models, their corresponding fabrication technologies, as well as the impact of these topological cues on cellular activity. Thus, in the present work, we have taken the opportunity to discuss these topics comprehensively. Further, we discuss the importance of various biophysical and biochemical aspects of the matrix, external biomechanical cues, and coculture strategies (with other cell types and intestinal microflora), supported by recent development in the domain. Besides, to further extend the scope of the review, aspects of stem cell engineering are also included.

Intestinal epithelium: understanding the biology of crypt-villus architecture

The intestinal epithelium is the innermost lining of the intestine, acting as a physical barrier between the internal tissue and the luminal environment and is responsible for the digestion of the chyme and absorption of the nutrients [5, 23]. The intestinal epithelium is organized into a specialized crypt/villus configuration [3, 24]. Notably, this configuration varies along the length of intestine, from the small intestine (SI) to large intestine (LI).

In the SI, the configuration consists of a villus (finger-like projection of the intestinal wall), surrounded by multiple crypts (invaginations into the intestinal wall, appropriately termed as the crypt of Lieberkuhn) [3, 24]. The villus is the site of intestinal brush-border enzyme-mediated digestion (mainly in the duodenum) and nutrient uptake (in the jejunum and ileum) [3]. The absorbed nutrients are further channelized into the blood mainstream via the underlying blood and lymphatic capillaries. The duodenum zone of SI has the longest villi (> 1 mm length) [24]. On the other hand, the crypt primarily acts as glands that secrete antimicrobial agents and several hormones. They are also characterized

by the presence of an intestinal stem cell (ISC) niche [1]. In contrast, the LI does not have villi-like structures but abounds of cryptic invaginations [1].

The ISC niche is highly dynamic and is responsible for restoring the intestinal epithelium every 4–5 days [25]. It contains active multipotent ISCs (also known as crypt base columnar cells or CBCs), placed alternatively to Paneth cells [24, 26]. Besides, the stem cell niche also has a quiescent stem cell population that plays a crucial role in regenerating the ISC niche in case of any tissue injury [24]. The CBCs undergo constant proliferation and also give rise to transit-amplifying (TA) cells, which further differentiate into secretory (Paneth, goblet, enteroendocrine and tuft cell) and absorptive (enterocytes and M cell) lineages [1, 24, 26]. In contrast, LI does not have Paneth cells, and the ratio of goblet cell to enterocytes is relatively higher than in SI [1].

During the proliferation and differentiation process, the fully mature epithelial cells show a gross upward movement along the crypt-villus axis in the SI and toward the luminal surface in the LI [1]. Besides this, the damaged or apoptotic cells shred off into the intestinal lumen. Notably, unlike other differentiated epithelial cells, Paneth cells move downwards to form the part of stem cell niche [24].

The distribution and specific functions of the different cell types are illustrated in Fig. 1 and Table 1.

Developing crypt-villus architecture: the current state of the art

Over the last decade, enormous efforts have been dedicated to recapitulating the crypt-villus architecture associated with the intestinal tissue in vitro [1, 11]. This section delineates different fabrication strategies employed for developing these topological features and the impact of these topological features on the behavior of cultured intestinal cells.

Fabrication strategies

Replica molding

Replica molding is a strategy that involves the development of master mold containing patterns, followed by multiple imprinting and remolding steps to fabricate scaffolds with biomimetic 3D intestinal topology.

Photolithography has widely been employed in various fields to obtain micropatterns. Typically, it is based on the crosslinking of a photosensitive polymer or photoresist, that is selectively exposed to ultraviolet (UV) light through a photomask with a desired pattern [44]. Conventional photolithography with SU-8 photoresist on silicon wafer followed by PDMS replica molding was employed to fabricate molds for 3D villi-like hydrogel units and the PDMS fluidic



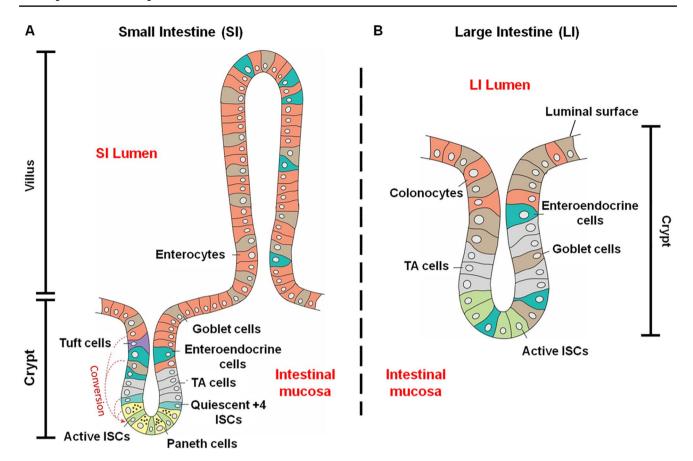


Fig. 1 Distribution of cells across the **a** crypt-villus axis of SI and **b** crypts of LI. ISC niche consisting of actively dividing ISCs (green) and Quiescent+4 ISCs (light blue) is present at the cryptic base. TA cells (gray) further differentiated to form enterocytes (in SI)/colonocytes (in LI) (orange), goblet (brown), tuft (purple), enteroendocrine

(blue), and Paneth cells (yellow). Paneth cells are particularly present in the crypts of SI. Red arrow designates the ability of different cell types to convert into active ISCs. Reprinted with permission from [26]

channel [45]. The resultant alginate/collagen hydrogel units with different aspect ratios were assembled inside the fluidic channel, recreating 3D intestinal environment on-chip for culturing cells under flow conditions. In another study, the sole crypt structures in a collagen scaffold were reproduced with one-layer photolithography, within a modified Transwell (Fig. 2b) [46] or within a customized platform in which a photo-patterned hydrophilic polytetrafluoroethylene (PTFE) membrane [47]. In an effort to recreate human SI epithelium with crypts and villi, two-layer photolithography was used. The strategy involved reproducing the negative of the villi first, and then the negative of the crypts in a single mold. Two steps of polydimethylsiloxane (PDMS) soft lithography followed the preparation of micromolds; after that, crosslinked collagen hydrogel scaffolds (via EDC/ NHS chemistry) were developed, reproducing crypts and villi (Fig. 2a) [48].

Recently, reaction-diffusion mediated photolithography (which involves controlled oxygen gradients within

the polymerization set up) has been developed as a simple, single-step and mold-free procedure to mimic the 3D villus-like geometry using a polymeric formulation containing poly(ethylene glycol) diacrylate (PEGDA), acrylic acid (AA), and Irgacure D-2959 [49]. The dimensions and geometries of the microstructures were established by playing with fabrication parameters such as the oxygen diffusion/ depletion timescales, the exposure dose, and distance to the light source. AA allowed covalent conjugation of ECM proteins (laminin or collagen), whose density could be tuned without affecting the mechanical properties of the hydrogel, making the constructs suitable for cell culture. Inexpensive and rapidly implementable UV-LIGA (lithography, electroplating, and molding) fabrication technology, involving backside exposure-based photolithography with SU-8, was used to produce an array of microneedles, with projective structure and size comparable to those of the mouse duodenal villi [50]. These molds were replicated to develop poly(lactic acid) (PLA)-based 3D scaffolds.

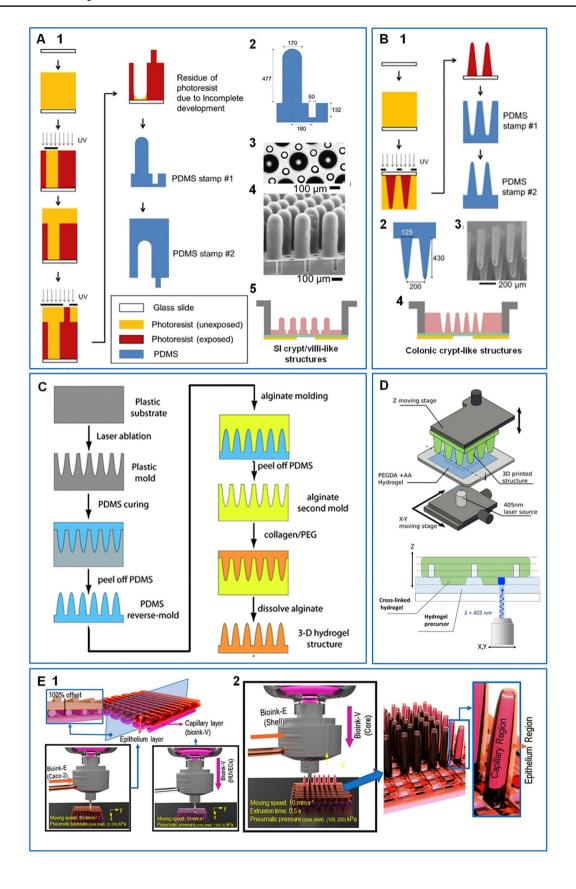


Table 1 Cells of intestinal tissue: distribution, major functions, and key cell markers

Cells	Localization	Major functions	Key cell markers	References
CBC cells	Crypt	Multipotent active ISC population. Bulk of proliferation and maintenance of crypts	LGR5, Olmf4, Ascl2, Smoc2, Sox9	[24, 27]
+4 cells	Crypt	Reserve/quiescent ISC population. Replenishes active stem cell population after injury	Bmi1, Tert1, Hopx and Lrig1	[24, 27]
TA cells	Crypt	Forms the intermediate progenitor pool. Differentiate into intestinal cells of secretory or absorptive lineages	DCAMKL1, Lrig1	[27–30]
Paneth cells	Crypt	Secretion of antimicrobial proteins (lysozyme, α -defensins, C-type lectins, and phospholipase A2). Provides necessary biochemical stimuli (Wnt, EGF, and Notch ligands) as well as essential nutrients for the maintenance of ISCs	Lysozyme, α-defensins	[24, 31, 32]
Tuft cells	Crypt and villus	Involved in chemosensing. Regulate type 2 immune response. Has a specific role in the detection of infection by parasitic helminths	Dclk1,Vil1, Pou2f3, α-gustducin, IL-25, TRPM5, Cox-1, Cox-2	[7, 33]
Enteroendocrine cells Crypt and villus	Crypt and villus	Involved in chemosensing. Regulate appetite, glucose level, intestinal motility, digestion, and even immune response (via secretion of peptide hormones including somatostatin, CCK, GIP, GLP-1, GLP-2, motilin, neurotensin, secretin, 5-HT, etc.)	Chromogranin A	[34–36]
Goblet cells	Crypt and villus	Protects against bacteria by the production of mucus. Modifies MUC2, MUC5AC, MUC5B, MUC1, MUC3, MUC17, Tff3 T cell-mediated immunity (via secretion of resistin-like molecule β). Aid in the restoration of epithelia post-mucosal injury (via secretion of trefoil factor). Involved in antigen uptake and presentation to subepithelial dendritic cells. The produced mucus also lubricates the gastrointestinal tract	MUC2, MUCSAC, MUCSB, MUC1, MUC3, MUC17, TIF3	[4, 32, 36–38]
Enterocytes	Villus	Involved in nutrient uptake, trans- or paracellular trans- portation of macromolecules. Secretion of antimicrobial proteins. Regulate immune response (antigen processing and presentation). Maintains homeostasis between host tissues and intestinal microbial communities (via secretion of IgA). Regulate inflammatory responses (via detoxification of LPS by ALP enzyme)	ALP, maltase glucoamylase, sucrase isomaltase, carbonic anhydrase II	[32, 36, 39–41]
M cells	Peyer's patches and isolated lymphoid follicles	Involved in transepithelial transport of intact microbe and luminal antigens for presentation to the immune cells	Glycoprotein 2, Uromodulin, PrPC, ANXA5	[32, 42, 43]

like domains-protein 1, DCAMKLI doublecortin- and calmodulin kinase-like 1, Wnt wingless-related integration site, EGF epidermal growth factor, DclkI doublecortin-like kinase 1, VilI villin 1, Pou2/3 pou domain, class 2, transcription factor 3, IL-25 interleukin 25, TRPM5 transient receptor potential cation channel, subfamily m, member 5, Cox-I cyclooxygenase 1, Cox-2 cyclooxygenase 2, CCK cholecystokinin, GIP gastric inhibitory polypeptide, GLP-1 glucagon-like peptide 1, GLP-2 glucagon-like peptide 2, 5-HT 5-hydroxytryptamine, MUC2 mucin 2, MUC5AC mucin 3, MUC17 mucin 17, Tff3 trefoil factor 3, LPS lipopolysaccharides, ALP alkaline phosphatase, PrPC LGR5 Leucine-rich-repeat-containing G-protein-coupled receptor-5, Ohnf4 olfactomedin 4, Ascl2 achaete-scute complex homolog 2, Smoc2 secreted modular calcium binding protein-2, Sox9 SRY-box, Bmil B lymphoma Mo-MLV insertion region 1 homolog, Terrl telomerase reverse transcriptase, Hopx homeoboxdomain-only protein, Lrig1 leucine-rich repeats and immunoglobincellular prion protein, ANXA5 annexin A5







∢Fig. 2 Fabrication strategies. **a** Two-layer photolithography to replicate crypt-villi features. 1 The fabrication process of PDMS stamps. 2 Measurement specifications the developed PDMS stamp #1. Values mentioned in µm. 3 Bright-field micrographs of the stamp #1 from the top-view. 4 Scanning electron micrographs of stamp #1 from the side view. 5 Schematic representation of the SI-associated patterns developed on collagen scaffolds in the modified transwell insert. Reprinted with permission from [48]. b Single-layer photolithography to replicate colonic crypt features. 1 The fabrication process of PDMS stamps. 2 Measurement specifications the developed PDMS stamp #2. Values mentioned in um. 3 Scanning electron micrographs of the PDMS stamp #2. 4 Schematic representation of the colon-associated crypt structures developed on collagen scaffolds in the modified transwell insert. Reprinted with permission from [46]. c Fabrication steps to obtain 3D collagen villi-like structures, starting from a plastic substrate created by laser ablation from which the PDMS mold and then alginate mold are replicated. The alginate mold is finally dissolved. Reprinted with permission from [51]. d Schematic representation of SLA strategy for layer-by-layer fabrication of 3D printed PEGDA scaffolds (in the presence of a photoinitiator and AA polymer chains) using a 405 nm laser. Reprinted with permission from [58]. e Extrusion-based bioprinting strategy. Representative schematics of the fabrication process of 1 2D mesh-like crypts and 2 3D finger-like villi. Two bioinks, namely bioink-E (with Caco-2 cells) and bioink-V (with HUVEC) were employed. The villus architecture consisted of inner vascular core (bioink-V) and an exterior shell of epithelial cells (bioink-E), developed using VMEPS with core-shell nozzle. Reprinted with permission from [62]

UV laser ablation was used to develop plastic molds with high-aspect-ratio holes. The features were replicated on PDMS and on calcium alginate that was subsequently dissolved to achieve final 3D collagen/PEG villus-like structures. The use of calcium alginate as a sacrificial mold eliminated the stress associated with the separation of the mold from the hydrogel structure, enabling the fabrication of patterns with high aspect ratio and curvature, without damaging them (Fig. 2c) [51]. Similarly, CO₂ laser was also employed for developing poly(methyl methacrylate) (PMMA) templates and subsequently 3D patterned collagen scaffolds [52, 53]. The CO₂ laser system is often advantageous than UV laser for developing molds with high aspect ratios due to its higher etching capability.

Porous PLGA scaffolds were produced by a modified version of the porogen leaching-thermally induced phase separation method. PLGA was mixed with sodium bicarbonate as a porogen, and the solution was poured into the agarose molds. After PLGA casting, the scaffolds were frozen at -20 °C, submerged in pre-cooled ethanol to extract chloroform and finally immersed in warm distilled water to dissolve the porogen [54]. In another study, the same strategy was employed to develop porous 3D patterned scaffolds with poly-ethylene-co-vinyl-acetate (PEVA) polymer [55].

Although the substrates mimicking crypt-villus geometry have widely been fabricated using the above-mentioned strategies, the obtained microstructures are only an approximation of the irregular and multiscale topography of the

native tissue. To enable the polymeric replica of biomimetic irregular SI geometries, chemical vapor deposition (CVD) of Parylene C on decellularized porcine SI was introduced [56]. After Parylene C was deposited, the porcine tissue was removed by bleach immersion. The side of the material, in contact with the tissue, contained the negative pattern of the native intestine and was used as a mold to prepare a PDMS replica. Resulting PDMS substrates exhibited multiscale resolution features, including crypt-villi structures and the underlying basement membrane [56].

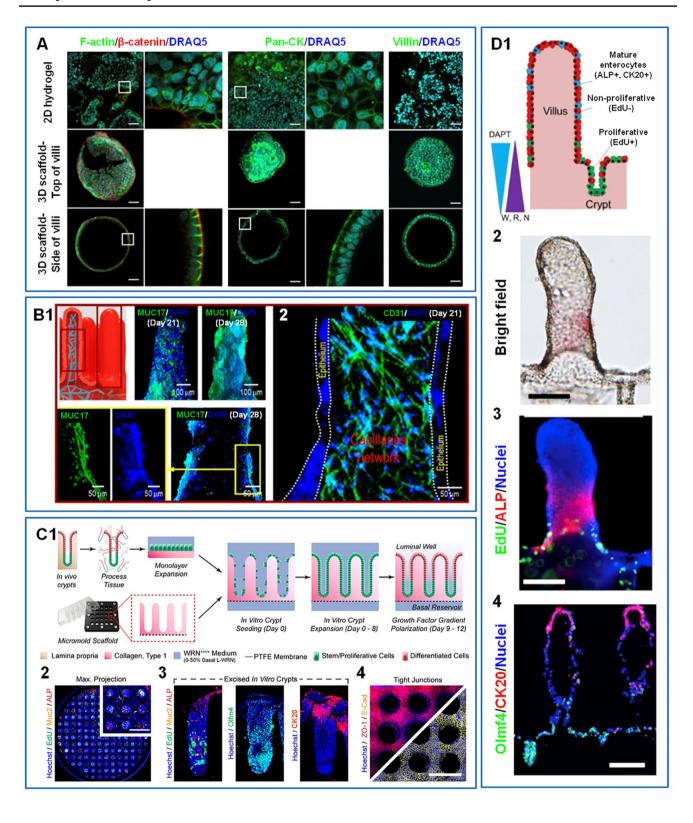
3D bioprinting

3D printing or additive manufacturing involves the fabrication of 3D objects by layer-by-layer deposition of material(s), following a computer-aided design (CAD) model [57]. Recently, stereolithography (SLA), a form of 3D printing, was employed to fabricate artificial scaffolds matching the topography and dimensions of the mouse intestinal epithelium with a microscale resolution (Fig. 2d) [58]. The authors employed the same PEGDA-AA-based polymeric formulation, similar to that used in the other study [49]. The printed scaffolds were further conjugated with ECM proteins (fibronectin or laminin or collagen) to improve its suitability for cell culture. Notably, unlike the previous study which used photolithography to fabricate villi-like structures with PEGDA-AA formulation [49], implementation of 3D printing offered more uniformity in the printed structures as well as allowed to develop scaffolds with both crypts and villi geometries [58].

Further, the advanced versions of 3D printing even allow fabrication of cell-laden 3D constructs, which is otherwise difficult to achieve with traditional ones. They often use a bioink formulation consisting of cells and cell-protecting hydrogels. Moreover, this strategy can enable the printing of multiple cells in the desired region within the scaffold [59, 60]. In this regard, 3D bioprinting was employed to mimic the human intestinal villi with Caco-2 cell-laden collagen bioink, crosslinked with tannic acid, a natural polyphenol. The bioprinting process involved two steps: (i) printing of a flat mesh structure to fabricate the crypt region and (ii) printing in the vertical direction by a vertically moving extrusion-based printing system (VMEPS) to obtain a protruded villus structure [61].

The same group introduced a dual-cell printing system supplemented with a core–shell nozzle to fabricate 3D intestinal villi containing epithelium as well as a blood capillary network (Fig. 2e). They prepared two bioinks: bioink-E with Caco-2 cells, for shell region, and bioink-V with human umbilical vein endothelial cells (HUVECs) for the core region. This time, the process involved three steps: (1) printing of a layer of flat mesh structure using bioink-V; (2) printing of a second layer using bioink-E on the bioink-V





structures with 100% offset; and (3) the villus-like structures were obtained by simultaneously printing bioink-E and bioink-V in the vertical direction using a core–shell nozzle

[62]. The same strategy was further opted to develop 3D intestinal patterns using collagen/decellularized porcine SI submucosa-based bioinks [63].



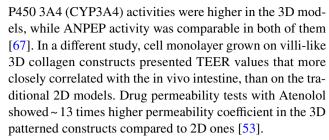
∢Fig. 3 Cell responses on the 3D intestinal models. **a** Representative confocal micrographs of Caco-2 cells cultured on 2D hydrogels and 3D SLA-printed scaffold (at the top and side of a villus) for 21 days. Expression and distribution of F-actin/ Pan-cytokeratin/villin are shown in green, β-catenin in red, and the nucleus is stained blue with DRAQ5 (Scale bar 50 µm). White square mark the area zoomed for better visualization; images provided in the right-hand side of each subset. Reprinted with permission from [58]. b Representative immunofluorescence micrographs demonstrating the presence of 1 MUC17+Caco-2 cells (at the exterior) and 2 CD31+capillary-like network (in the core) in the bioprinted intestinal model. Nucleus was stained with DAPI. Reprinted with permission from [62]. c Generation of colonic crypts. 1 Schematics of the culture of colon-derived crypt cells on micromolded collagen scaffold. Post-seeding on the scaffolds, the crypt cells were expanded till day 8. Cellular differentiation and polarization were induced via exposure to growth factor gradient. Representative images demonstrating spatial expression of intestinal markers in colonic crypts differentiated in vitro in the 2 topview (max projection along the z-axis and summing up the images) and 3 side view. 4 Confocal micrographs showing expression of ZO-1 and E-cad at the luminal surface of the differentiated in vitro crypts. Reprinted with permission from [47]. d Generation of SI-associated crypt-villus pattern in vitro. 1 Illustration showing the effect of combined WRN growth factors and DAPT gradient on cell polarization. 2 Bright-field images of the cell-seeded 3D structure (Scale bar 100 µm). Representative images, demonstrating the distribution of 3 EdU/ALP and 4 Olfm4/CK20 positive cells along with the 3D pattern with nucleus stained in blue (Scale bar 100 µm). Reprinted with permission from [48]

Functional behavior of the cells on 3D intestinal models

Gene expression and barrier functions

Substrate topography plays a fundamental role in determining the physiological function of a variety of cell types including osteoblasts [64], hepatocytes [65], cardiomyocytes [66], and even stem cells [64]. It influences cell–matrix interaction, cell–cell interaction, and biomechanical cues, modulating the intracellular signaling cascade, and subsequently, the cell behavior [52]. Intestinal cells are no different in this regard.

A notable difference between that arises due to the presence of villus-like structures is the available surface area for cellular growth that may influence their physiological behavior. A study demonstrated that cells on both 2D and 3D models grew at a similar rate for the first 12 days, but after 16 days, they had grown twice on patterned 3D substrates. Additionally, a decrease in TEER values (i.e., the integrity of tight junctions formed by cell–cell contacts) and an increase in absorptive permeability of fluorescein could be attributed to an increase in culturable surface area. A reduction in the expression of occludin and p-glycoprotein (P-gp, an efflux transporter protein) was also evident as compared to the 2D model. Metabolic enzymatic activities associated with intestinal epithelium were also evaluated; ALP and cytochrome



3D topological features also affected the differential phenotype in a spatial manner. Poly-ε-caprolactone (PCL)/ PLGA 3D patterned membranes facilitated differentiation of Caco-2 cells, as evidenced by the brush-bordered cell phenotype (at the top of the villus), temporal increase in the activity of (ALP) during differentiation, and uniform zonula occludens-1 (ZO-1) expression. Lower TEER, along with higher uptake of ²⁵ Mg and ⁴⁵Ca, was witnessed on the 3D membranes as compared to the non-patterned 2D ones [68]. In another study, Caco-2 cells seeded onto PEGDA-AA 3D scaffolds showed directional migration from the base to the top of the villus, to fully cover them. After 21 days, Caco-2 cells expressed markers such as villin (at the apical side of the microvilli), ZO-1 (in tight junctions), and β -catenin (in the basolateral cell membranes). The 3D villi-like constructs had lower TEER values and higher permeability coefficient than the conventional Transwell controls [49].

Recently, printed PEGDA-AA-fibronectin hydrogel scaffolds, 2D and 3D, were used to study the impact of 3D topography on Caco-2 differentiation. 2D coverslip (with or without fibronectin coating) was also taken as 2D control. Cells formed a confluent monolayer in all three surfaces; however, there existed significant differences in their phenotypic characteristics. The cells on glass coverslip exhibited flattened morphology without any apicobasal polarization, irrespective of fibronectin coating. On the contrary, elongated morphology and polarized cells were observed on the printed hydrogel matrices, both 2D and 3D. Cells on the 3D scaffolds demonstrated higher cell elongation, better cell polarity, and higher expression of villin and ALP, as compared to 2D hydrogels. The study also revealed that the height of the villus (500 or 1500 µm) did not critically affect cell polarization (Fig. 3a) [58].

As mentioned above, intestinal epithelium harbors multiple cell types. Thus, to better mimic cellular profiles, existing in vivo, Caco-2 and HT29-MTX were cocultured on porous PLGA scaffolds in a 3:1 ratio. A directional cellular migration from the base to the top of the scaffold was evident, with villus tips having better differentiated phenotypic characteristics. Higher ALP secretion and mucus production, along with lower TEER values, were observed in 3D constructs than their 2D counterparts [54]. Integration of native-like vascular network is necessary, yet challenging aspect for developing the intestinal model. In this regard, the coaxially 3D printed models with spatially defined localization



of Caco-2 (at the exterior) and HUVEC (in the core) were shown to support an improved differentiated phenotype of the epithelial cells, in terms of ALP and ANPEP activity, expression of MUC17, and the presence of brush-border epithelium, as compared to the models without HUVEC. Besides this, HUVEC also underwent a transition to form a capillary-like network in coculture models (Fig. 3b) [62].

Intestinal models are also applied for ascertaining the role of the microbial population (both healthy gut microbes and pathogenic ones) in the gut. The conventional 2D models fall short in mimicking the original bio-interactive profiles of the intestinal epithelium with the microbes. A recent study highlighted a lower invasion by pathogenic bacterial strains, *Escherichia coli* O157: H7 and *Salmonella typhimurium* on 3D patterned scaffolds. Such an outcome was linked to higher expression of mucin-related genes, particularly MUC17, in the case 3D substrates than the 2D monolayer cultures [52].

Intestine-associated cellular compartmentalization

Although the studies mentioned in the previous section demonstrate physiologically relevant spatially discrete phenotypic features, they fail to closely recapitulate the aspects of native-like cellular compartmentalization. The intestinal epithelium has a distinct crypt-villus (in SI) or crypt-lumen (in LI) architecture; wherein stem cell niche lies in the crypts, while the differentiated polarized cells are present at the villus/luminal surface [69]. Current efforts to recapitulate such complexity utilize gradient of growth factors and small molecules across the 3D patterned scaffolds, seeded with intestinal crypt cells, which drives migration and differentiation process, creating lineage compartmentalization, similar to the native tissue.

In this regard, collagen-coated microhole array was fabricated for reproducing the cell compartmentalization in crypt structures. Primary mouse colon crypt cells were seeded, and localized gradients of growth factors (Wnt3A, R-spondin, Noggin; together termed as WRN) were applied across the microholes, thereby yielding a stem/proliferative region (5-ethynyl-2'-deoxyuridine (EdU)+) inside the microholes, and non-proliferative and differentiated (ALP+) zones distant from the microholes [20]. Another crypt-lumen model was fabricated to study intestinal cell compartmentalization process, where human intestinal crypt cells were cultured on a micromolded collagen scaffold (with an array of crypt-like invaginations) and subjected to a biochemical gradient by placing WRN-rich medium in the basal reservoir and WRN-deficient medium in the upper (or luminal) reservoir. Polarization of the in vitro tissue was observed with proliferative cells (EdU+, olfactomedin 4 (Olfm4)+) in the crypt region and differentiated cells (EdU+, keratin 20 (CK20)+ and erzin+) at the luminal surface. Interestingly,

mucus-secreting goblet cells (MUC2+) were also observed [46].

Scalability of the intestinal model is one of the most pertinent challenges. In this regard, a scalable platform consisting of ~3875 crypts over a single membrane, where chemical gradients could be simultaneously imposed in distinct regions; this enables a parallelizable in vitro crypt formation system. Here also, crypts were seeded with human colon crypt cells, and exposure to WRN gradients resulted in cellular compartmentalization. Proliferative cells (EdU+, Olfm4+) were restricted to the basal region of the crypts, whereas differentiated colonocytes (ALP+, CK20+) were present at the luminal plane. Proteins responsible for tight and adherens junction maintenance such as ZO-1 and E-cadherin (E-cad) were displayed at the luminal plane. Goblet cells were also present (Fig. 3c) [47].

Besides, in a more recent study, the conventional WRN-gradient (basal to luminal) was found to be insufficient to fully induce cellular compartmentalization on 3D collagen constructs, seeded with human small intestinal crypt cells. Incorporation of the second gradient of a small molecule, DAPT (luminal to basal), overcame this issue. After applying a dual gradient strategy, proliferative (EdU+, Olfm4+) and differentiated (CK20+, ALP+) cells were observed explicitly in crypt and villus regions, respectively. Moreover, EdU+ cells migrated upwards along the crypt-villus axis over time, similar to in vivo (Fig. 3d) [48].

These studies indicate that mimicking the 3D geometry of the intestinal epithelium influences the cellular phenotype. Table 2 summarizes all the studies that have evaluated cellular behavior on a 3D microstructured construct to date.

Modulation of intestinal cell behavior: biophysical and biochemical determinants

Controlling the in vitro behavior of the cells (both the stem cells and differentiated ones) and achieving desirable in vivo-like effects, are some of the biggest challenges faced by the researchers. The recent understanding of existing multifactorial determinants (intrinsic and extrinsic) of cell behavior has led to significant advancements in this field. Principally, the biochemical nature of in vivo ECM is mimicked, by utilizing in vivo-like mechanical and physical cues that have now been established as substantial regulation factors of cell fate and behavior. Moreover, recreating the cellular environment and interactions with neighboring cell population and intestinal microflora has also being explored. Besides, creating various gradients of ECM components, biochemical factors, and gases, similar to the native cryptlumen (in LI) and crypt-villus (in SI) axis are also pertinent options (Fig. 4).



Table 2 Additional studies on in vitro intestinal tissue model with patterned crypt-villus geometry

Refer- ences		[20]	[46]	[70]	[47]	[21]	[51]
	ananysis (days)	4	12 (extendable to able to atleast 32)	4	2	7	7–10
Secretive P_{3D} (v/s P_{2D}) (in cm/s)	Metabolite ana- Iyzed (Values)	1	I	1	1	1	1
Absorptive P_{3D} (v/s P_{2D}) (in	cm/s) Metabolite analyzed (Values)	I	1	1	1	1	1
TEER _{3D} (v/s	$(\ln\Omega\mathrm{cm}^2)$	1	1	I	1	I	ı
Compartmentali- zation		Crypt: EdU+; zones distant from crypts: ALP+, MUC2+gob- let cells were observed near the crypt microholes	Crypt: EdU+, Olfm4+; Luminal plane: CK20+, Erzin+; MUC2+goblet cells were also observed	Crypt base: EdU+,Sox9+; Luminal plane: ALP+	Crypt: EdU+, Olfm4+; Luminal plane: ALP+, CK20+, ZO-1+, E-cad+; MUC2+ goblet cells were also present	1	I
Gene/protein expression or	acuvity in 3D (v/s 2D)	ALP, MUC2 (not compared with 2D)	CK20, Olfm4, Erzin, MUC2 (not compared with 2D)	Sox9-EGFP, ALP, MUC2 (not compared with 2D)	E-cad, ZO-1, CK20, MUC2, ALP, Olmf4 (not compared with 2D)*	CrgA, SI, E-cad (not compared with 2D)	1
Cells used		Murine intestinal crypts	Human colonic crypts	Mouse colonic epithelial cells	Primary human colon epithelial cells	Caco-2 or juvenile mouse small intes- tinal crypt cells	Caco-2
tt.	Crypt-villus	ı	1	1	1	1	1
ypts or villi or both	Crypt	Diameter: 50 µm	Depth: 430 µm; base diameter: 125 µm (at the base); density: 35/ mm²	Depth: 250 µm; diameter: 75 µm	Depth: 419.1 μm; diameter: 65.4 μm (at the base); diameter: 128.1 μm	1	1
Specifications of crypts or	Villus	ı	1	1	1	Height: 340 µm; width: 184.5 µm (at the base) to 49 µm (at the tip); spacing: 69.5 µm	Height: 450– 500 µm; aspect ratio: ~ 5
Material used		Collagen	Collagen (crosslinked with EDC/ NHS)	Collagen I hydrogel (EDC/NHS crosslinking)	Collagen (crosslinked with EDC/ NHS)	PGS (along with a sodium bicarbonate as porogen; crosslinked with MDI)	Collagen or PEGDA
Fabrication methodol-	ogy used	Photolithog- raphy	Photolithog- raphy and replica molding	Photolithog- raphy and replica molding	Photolithog- raphy and replica molding	Laser ablation of PMMA and replica molding	Laser ablation of PMMA and replica molding
3D architectural	reatures	Intestinal crypt configuration				Intestinal villus con- figura- tion	





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	Refer- ences		[67]	[50]	<u>8</u>	[73]	[74]	[63]	[62]
				<u></u>	۲	드		<u>~</u>	<u> </u>
	Time before experimental analysis (days)	(c.fm)	20	I	21	10	7	20–21	14-21
	Secretive P_{3D} (v/s P_{2D}) (in cm/s)	Metabolite ana- lyzed (Values)	Rhodamine-123: 2.9×10 ⁻⁶ (v/s 4.3×10 ⁻⁶ for polycarbonate membrane)	ı	1	ı	ı	1	1
	Absorptive P_{3D} (v/s P_{2D}) (in cm/s)	Metabolite analyzed (Values)	FITC-dextran (4 kDa): 1.5×10 ⁻⁶ (v/s 0.78×10 ⁻⁶ for polycarbonate membrane): fluorescein: 9.1×10 ⁻⁶ (v/s 7.1×10 ⁻⁶ for polycarbonate membrane)	1	Fluorescein: 4.378×10 ⁻⁶ (v/s 8.756×10 ⁻⁶ for non-patterned nembrane and 1×10 ⁻⁶ for polycarbonate membrane)	1	1	FITC-dextran (4 kDa): 1.44×10 ⁻⁶	FITC-dextran (4 kDa): 0.622×10 ⁻⁶ (mesh); 1.889×10 ⁻⁶ (villi)
	TEER _{3D} (v/s TEER _{2D}) (in Ω cm ²)		60–80 (v/s 200 for polycar- bonate mem- brane)	I	85 (v/s 176 for non-patterned membrane and 215 for polycarbonate membrane)	ı	ı	1	1
	Compartmentali- zation		1	ı	Villus tip: More ZO-1 expression, clear microvilli projections; Base: Less defined micro- villi projections with low ZO-1 expression	ı	ı	1	1
	Gene/protein expression or activity in 3D (v/s 2D)		Decreased: P-gp and ZO-1	1	Increased: ALP	ZO-1 (no comparison shown)	1	ZO-1, E-cad, MUC17, ALP, and ANPEP (not compared with 2D)	E-cad, ZO-1, MUC17, ALP, ANPEP (not compared with 2D)*
	Cells used		Caco-2	Caco-2	Caco-2	Caco-2	intestinal epi- thelial cells (IEC6)	Caco-2 and HUVEC	Caco-2 and HUVEC
		Crypt-villus	1	1	1	1	1	1	1
	ots or villi or bo	Crypt	1	1	1	I	ı	Mesh (0/90 struts)	Mesh (0/90 struts)
	Specifications of crypts or villi or both	Vıllus	Height: 400 µm; diameter: 100 µm (at the base); surface area 0.1 mm²	Height: 420 μm; base diameter: 130 μm; density: 32.5/mm²	Height: 500 µm; diameter: 450 µm (at the base) to 400 µm (at the tip); spacing: 80 µm; density: 5/mm²	Height: 1000 µm	Height: 1000 µm; diameter: 700 µm (at the base) to 500 µm (at the tip)	Height: 831.1 µm, diameter: 190.9 µm; aspect ratio: 4.4	Height: 770 µm; diameter: 183 µm; aspect ratio:4.2
	Material used		Collagen (crosslinked with glutaral- dehyde)	PLA	PCL and PLGA	GelMA and PCL-MA (pho- tocrosslinked)	PLGA	Collagen/decel- lularized SI submucosa (crosslinked with tannic acid)	Collagen- Based Bioink (crosslinked with tannic acid)
iable z (continued)	Fabrication methodol- ogy used		Photolithog- raphy and replica molding	Photolithog- raphy and replica molding	Photolithog- raphy and replica molding	Digital light processing stereolithog-raphy	3D printing and replica molding	Vertically moving extrusion- based cell printing	Vertically moving extrusion- based dual cell printing
מטוע ד	3D architectural								



 Table 2 (continued)

1													
3D architectural	Fabrication methodol- ogy used	Material used	Specifications of crypts or villi or both	ots or villi or botl	ч	Cells used	Gene/protein expression or activity in 3D	Compartmentali- zation	TEER _{3D} (v/s TEER _{2D})	Absorptive P_{3D} (v/s P_{2D}) (in cm/s)	Secretive P_{3D} (v/s P_{2D}) (in cm/s)	Time before Referexperimental ences analysis	Refer- ences
			Villus	Crypt	Crypt-villus		(v/s 2D)		$(in \Omega cm^2)$	Metabolite Metabolite and analyzed (Values)	Metabolite analyzed (Values)	(days)	
	Vertically moving extrusion-based cell printing	Collagen (crosslinked with tannic acid)	Height: 708 µm; diameter: 162 µm; aspect ratio: 4.4	Mesh (0/90 struts)	I	Caco-2	MUC17, ALP, and E-cad (not compared with 2D)	. 1	1	. 1	ı	21–30	[61]
Intestinal crypt- villus con- figura- tion	Laser ablation of PMMA and replica molding	PLGA (along with a sodium bicarbonate as porogen)	Неіght: 500 µm	1	1	Caco-2	1	Villus tip: differentiated cells; Base: undifferentiated cells	449.18 (not compared with 2D)	1	1	21	[75]
	Photolithog- raphy and replica molding	Collagen (crosslinked with EDC/ NHS)	Height: 477 μm; diameter: 170 μm	Height: 132 µm; diameter: 60 µm	Total mucosal thickness: 609 µm; crypt.villus ratio: 2; each villus surrounded by 6 crypts	Human small intestinal crypts	CK20, ALP, and Olfm4 (not compared with 2D)	Villus: EdU., ALP+, CK20+; Crypt: EdU+, Olfm4+	1	1	1	9–12	[48]
	High-resolution stereo-lithography 3D printing	PEGDA-AA (containing fibronectin)	Height: 500 µm; diameter: 300 µm (at the base) to 150 µm (at the tip)	Height: 200 µm; diameter: 50 µm	Total scaffold thickness: 750 µm; each villus surrounded by 6 crypts	Caco-2	Pan-cytokeratin, β-catenin, ALP, and villin (not compared with 2D)	Villus tip: More polarized, ALP+, Villin+ cells; Villus base and crypts: Low differentiation observed	1	1	1	9–12	[28]

*Coculture with HUVEC significantly improved the functionality of the tissue constructs

E-cad E-cadherin, ANPEP alanyl aminopeptidase, PGS poly(glycerol sebacate), MDI 4,4'-methylenebis(phenyl isocyanate), CrgA chromogranin A, SI sucrase isomaltase, GelMA gelatin methacrylate, EDC 1-ethyl-3-(3-dimethylaminopropyl)-carbodiimide, NHS N-hydroxysuccinimide, Sox9 SRY-box 9, WGA wheat germ agglutinin, SGLT-1 sodiumglucose transporter 1, Glut-2 glucose transporter 2

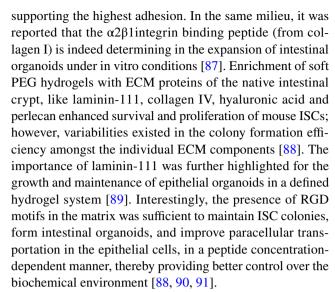


ECM composition

As the biological milieu surrounding cells, the ECM is a fibrous network of crosslinked proteins and glycosaminoglycans (GAGs) which makes up most of the cellular microenvironments [76]. Highly hydrated, it forms a hydrogel secreted by cells and connects the latter within tissues, and organs with each other. It provides structural and spatial organization to tissue components, protects the cells against mechanical pressures, mediates cellular attachment and also controls downstream signaling cascades that further determine their behavior [77, 78]. Abounding of signaling molecules for the regulation of tissue function and homeostasis, the ECM further translates information by its composition and molecular architecture, overall constituting a dynamic system of macromolecules implicated in tissue development, maintenance, and repair [76, 79, 80]. Continuously degraded and remodeled according to tissue needs [77], the ECM also presents a tissue-specific composition that enables tissuespecific functions.

In the intestinal epithelium, the ECM in the crypt base is preponderantly made up of the isoforms of fibronectins, laminins, collagens, and GAGs [81], suggesting isoformdependent regulation of the ISC niche. This is further emphasized by the differential distribution of diverse components along the crypt-villus axis, with laminins α1 and α 2 abundant in the crypt base, laminins α 3 and α 5 present in the villus, fibronectins surrounding the crypt, and tenascin found at the base of villus and in the surrounding of crypts [81–84]. Amongst the collagen superfamily, collagen type IV forms a major component of the intestinal basement membrane and is a key regulator of the mechanical microenvironment of the intestinal crypt cells [81, 84]. Besides the biochemical composition, the ECM also directs cell behavior by its selective entrapment of growth factors and chemical stimuli. In fact, among GAGs, heparan sulfate proteoglycan (also known as perlecan) presents the ability to bind various growth factors such as wingless-related integration site (Wnt), Hedgehog, transforming growth factor beta (TGF-β), and fibroblast growth factors (FGF) proteins, thereby contributing to ISC maintenance [81, 84, 85]. Hence, substrate biochemistry has been used to regulate cell behavior in vitro, often detaining a substantial role in optimizing stem cell microenvironment and their differentiation.

In a study, the authors screened various basement membrane ECM proteins and found that collagen IV readily enabled the attachment of mouse intestinal epithelial cells, while collagen I overlay was essential for preserving the Lgr5+expression (stem cell marker) in the sandwich cultures [86]. Notably, in the case of collagen type IV coating, the cell attachment was mediated by integrin $\alpha 2\beta 1$, and was dependent on the ECM concentration; $100 \mu g/mL$



Further emphasizing on 3D patterned substrates, a synergistic effect of substrate topology and ECM protein on the phenotypic characteristics of cultured Caco-2 cells was reported [92]. For this, crypt-like collagen microwells, coated with either laminin or fibronectin, were used. The study revealed better cell differentiation characteristics, barrier function, ANPEP, and ALP activity, on laminin-coated substrates than on the fibronectin-coated ones. On the other hand, the topological features affected cell adhesion, spreading and barrier functions. In the same backdrop, a recent study highlighted that adding decellularized SI submucosa to collagen type I-based bioink improves cellular activities in bioprinted Caco-2 cells, allowing for the formation of the more biomimicking intestinal epithelium, aided by a bioengineered villus geometry [63].

Altogether, these findings underline the essential role of matrix biochemistry in modulating intestinal cell growth and differentiation, with ECM proteins conditioning in vitro cell adhesion, proliferation and differentiation.

Matrix stiffness

Among the parameters regulating cellular behavior, substrate stiffness represents one of the most critical factors, controlling cell phenotype and determining stem cell fate [93, 94]. Under in vivo conditions, the collagen content in the intestinal submucosa is 2–4 times higher than that in the mucosa region (that contains the epithelium and displays crypt-villus architecture), pointing out the existence of stiffness gradients between these layers [81]. Thus, various studies have attempted to exploit substrate stiffness in optimizing in vitro intestinal epithelium models. However, the consensus is yet to be reached for optimal stiffness due to the cellular response often being a result of both substrates physical and biochemical properties, with different reports depending on the nature of the matrix material. For instance,



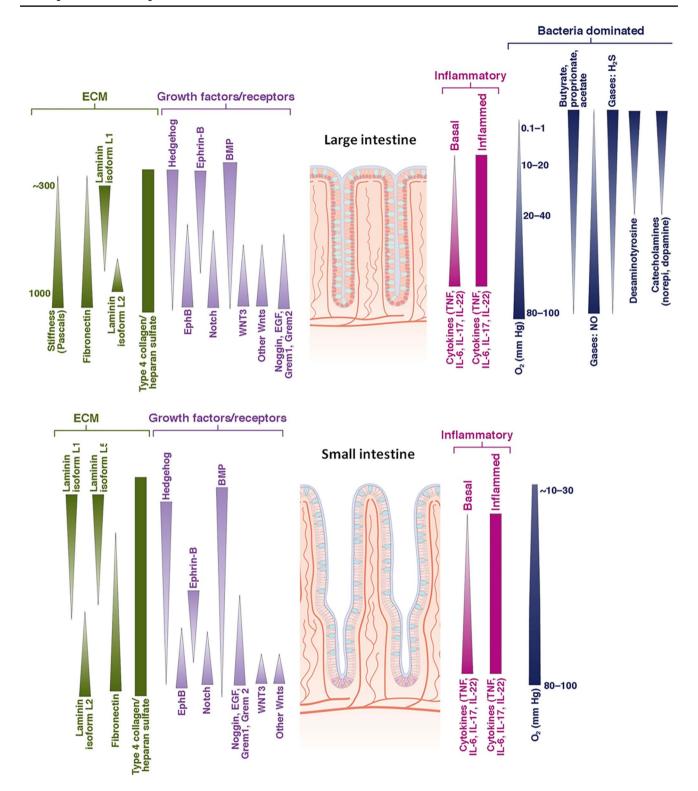


Fig. 4 Illustration of various gradients found along the crypt-lumen (in LI) and crypt-villus (in SI) axis in vivo. Gradients of ECM, growth factors/receptors, inflammatory cytokines, and bacteria-dominated components (gases and small molecules) have been shown with

the shades of green, purple, pink, and blue color. The pointing end and the broad base of the triangle represent lower and higher concentration, respectively. Reprinted with permission from [81]



mouse ISCs embedded in PEG hydrogels (conjugated with 1 mM RGD motifs) of higher stiffness (1.3 kPa) supported higher cell proliferation than those with lower stiffness (300 Pa); this involved the yes-associated protein 1 (YAP), a mechanosensing effector and key mediator in ISC self-renewal [88]. In another study, fibrin gels (supplemented with 10% Matrigel) of stiffness range between 77 ± 25 and 140 ± 47 Pa showed higher and bigger cyst formation from mouse ISCs, 4 days post-cell seeding [89]. Similarly, modulation in the matrix stiffness of protein-engineered scaffolds via tetrakis(hydroxymethyl)phosphonium chloride-based crosslinking affected the yield of mouse organoids; softer scaffolds (180 Pa) showed higher organoids formation than the stiffer ones (1220 Pa), irrespective of the content of adhesion motifs [90].

Besides, efforts have been made to create monolayer cultures of intestinal cells that have physiological relevance. Varying stiffness was achieved by simply modulating the thickness of Matrigel coating in polystyrene substrates; with $2.9 \pm 0.1 \, \mu m$ (Young's modulus of $\approx 3 \, \text{GPa}$) and $\sim 2 \, \text{mm}$ (storage modulus $\approx 50 \text{ Pa}$), thin and thick coatings formed stiff and soft substrates, respectively. Organoid-derived crypts or single cells, when cultured on stiffer substrates underwent spreading and formed epithelial monolayers, whereas those on softer substrates formed organoids [95]. In a separate study, collagen hydrogel substrates with a gradient of stiffness (230 \pm 140 Pa) were developed within Transwell inserts using EDC/NHS-mediated gradient crosslinking. The luminal surface with lower stiffness was used for culturing the cells, while the stiff basal surface prevented gel deformation; this closely mimicked the lamina propria of the intestine. The study revealed that lower TEER values along with a higher expression of breast cancer resistance protein (BCRP) and multidrug resistance protein 2 (MRP2) in the case of gradient crosslinked substrates relative to the conventional thinner and stiffer collagen membrane $(1.50 \pm 0.27 \text{ MPa})$. Moreover, grown primary human small intestinal monolayers enabled prazosin (a substrate of BCRP) transport as in vivo, in contrast with the stiffer substrates [96]. As an extension to this work, the authors demonstrated that the cells on the conventional scaffold lacked various drug-metabolizing enzymes relative to those on the gradient crosslinked hydrogels, additionally not properly responding to known inducers and inhibitors in contrast with the latter substrates [97].

Therefore, substrate stiffness represents an essential factor of cellular microenvironment with potential for directing the behavior of intestinal cells and thus can be leveraged to create more biomimicking in vitro intestinal epithelium models. As to the effect of matrix stiffness in 3D topological constructs, no report is yet available that corroborates

the translation of the above stiffness-related findings in such settings, thus extending the scope of future developments.

Shear stress and peristaltic motion

The intestine constitutes a dynamic organ where cells are continuously exposed to peristalsis, an alternation of contraction-relaxation cycles that enable the progression of the luminal content of the hollow organ and facilitate food digestion; notably, differences exist in the flow/transit time and contraction specification of SI and LI [1, 4]. Such microenvironment has been recapitulated in vitro in microfluidic platforms that recreate in vivo cues to affect culture outcome. For instance, using PDMS and soft lithography, a microfluidic device was designed that permitted liquid flow at a low rate (30 μ L h⁻¹), producing low shear stress (0.02 dyne cm⁻²) and cyclic strain of 10% (0.15 Hz), thus reminding physiological peristaltic movements [13]. When Caco-2 cells were cultured therein, compared to static culture in Transwell inserts, the assembled cues led to the growth of a columnar epithelium with improved polarization, paracellular transport, ANPEP activity, and barrier function that further formed intestinal villi-like folds, much like the whole intestine. Moreover, the cells grown under such complex conditions, unlike those under static conditions, maintained their viability and barrier function even when cocultured with a probiotic bacterium, Lactobacillus rhamnosus GG. Under similar microenvironmental conditions, the reconstitution of proliferative crypts was highlighted. EdU+cells further differentiated into enterocytes, goblet, enteroendocrine, and Paneth cells along the crypt-villus axis [98]. Better CYP3A4 activity was also observed. In a recent study, intestinal morphogenesis was shown to be influenced by fluid-flow rates in a time-dependent manner, the flow regime between 70 and 120 μ L h⁻¹, being most effective [14]. The authors also identified Frizzled-9 receptor as the mediator of this flow-induced spontaneous morphogenesis. By changing the fluid shear stress from ~ 0 to 0.03 dyne cm⁻² (physiological range), it was observed that phenotype of Caco-2 cells (mucus secretion, microvilli formation, expression of tight junctions and CYP3A4) could be modulated, with a shear stress of 0.02–0.01 dyne cm⁻² creating the optimum parameters [99]. Another study also revealed that the response of intestinal epithelium to strain, in terms of proliferation and activation of downstream signaling cascades, was dependent on ECM proteins. Collagen I, collagen IV, and laminin positively modulated cell proliferation under cyclic strain, but not fibronectin [100]. Interestingly, better tissue growth and maturation were observed when human intestinal organoids, exposed to uniaxial strain, were transplanted in mice mesentery, according to Poling et al. [101] who performed morphometric, transcriptomic, and functional studies on the explanted constructs.



In an attempt to integrate fluid-flow dynamics with intestine-associated topological features, recently, a 3D printed bioreactor was developed [55]. The study revealed the formation of site-specific expression profiles of Caco-2 differentiation with elevated apoptosis at the tip of the villus (regions subjected to the highest shear), and less ALP but increased mucus secretion near the base [55]. These results were more similar to in vivo intestine than those obtained in the case of conventional 2D Transwell culture or static structures. Active glucose transportation (at a higher rate) along with decreased TEER values were also evident in the presence of 3-D topography and fluid-induced shear. In the same prospect, the combination of 3D topology and shear yielded synergistic effects, giving Caco-2 cells an adequate stimulus for differentiation and considerably improving their metabolic activities; specifically, CYP3A4 activity was 7 times higher on 3D scaffold under fluidic condition than static 2D conditions [72]. Also, ANPEP activity and cell absorptive permeability were enhanced in the 3D fluidic conditions.

Given all these regulatory effects, mechanical cues constitute a major factor in the design and fabrication of devices and tissue models aiming to recapitulate the intestinal tissue for various pathophysiological and pharmacological studies.

Coculture with other cell types as well as microflora

Another strategy for biomimicking the physiology of the intestinal epithelium is by coculturing epithelial cells with other impactful cells (including microbial cells) that provide structural support and/or participate in regulatory crosstalks with the cells of interest, via generation of gradients of growth factors, gases, and microbial products, thereby regulating the dynamics of ISC niche and maintenance of intestinal health [81]. Thus, many efforts have been directed toward developing suitable in vitro coculture modules.

In a study, Caco-2/HT29-MTX coculture and Caco-2/ HT29-MTX/Raji B triple culture models showed more physiological relevance than Caco-2 monoculture models for studying the intestinal permeability of the drugs [102]. Intestinal muscularis cells were cocultured with intestinal epithelium on collagen-coated PCL tubular scaffolds, showing the presence of 11 different cells types from the mucosa, the muscularis, and the serosa and long-term spontaneous and periodic contractions, thereby recapitulating native-like characteristics, to an extent [103]. Incorporation of human intestinal microvascular epithelial cells with intestinal epithelium in a parallel channel of a microfluidic device (accompanied by flow-induced shear and cyclic strain) enhanced the rate of intestinal morphogenesis compared with those cultured without endothelial cells and showed better resemblance with native human duodenum [104].

One of the most utilized components in coculturing approaches is the addition to epithelial cells of supportive

stromal cells, generally fibroblasts or myofibroblasts [105, 106]. The presence of the NIH-3T3 fibroblasts accelerated the formation of epithelial layer by Caco-2 cells, strengthened its barrier function and improved its absorptive permeability as well as promoted the recovery upon disruption of intestinal barrier [105]. Intestinal subepithelial myofibroblasts (ISEMFs) were shown to support the long-term culture of intestinal organoids, even in the absence of a few critical growth factors required for maintaining cultures [107]. Moreover, supplementation of ISEMFs conditioned medium was sufficient to prevent autolysis of human intestinal organoids in collagen gels [108]. Enteric nervous system (ENS) cells have also been utilized to elucidate their interactions with murine ISCs [109]. The findings showed that ENS cells and myofibroblasts together control the stem cell behavior, particularly inducing their differentiation into enteroendocrine cells at a higher proportion relative to that done by coculture with myofibroblasts alone. Moreover, better barrier function and reduced flux of FITC-dextran was also evident in co- or tri-cultured models as compared to mono-cultured ones. Besides this, all three systems also showed variations in the production of cytokines.

Considering the involvement of gut microbiota in the various metabolic processes as well as pathobiological conditions [110, 111], they have also been introduced in various in vitro intestinal models, thereby aiming toward recreating more in vivo-like microenvironment. Coculture of organoids with Lactobacillus rhamnosus GG resulted in an increased percentage of proliferative cells and improved differentiation into Paneth cells, while Lactobacillus reuteri D8-induced proliferative effects in the organoids and prevented them from the damaged caused by TNF- α [112]. Besides, pathologic bacteria are introduced in intestinal epithelial culture systems to study pathophysiological pathways involved in infections and certain pathological conditions [22, 113]. Here, it is also important to mention that the microbiome products such as polyamines and short-chain fatty acids (SCFAs) also critically determine intestinal dynamics, and thus need consideration. For instance, butyrate, a type of SCFA, is essential for differentiated colonocytes for undergoing healthy mitochondrial respiration and promoting barrier function (at a lower concentration), while it acts as an inhibitor to ISC proliferation [81, 114–116]. Polyamines are essential for the growth and development of intestinal mucosa [81, 117, 118].

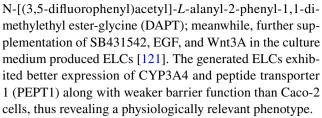
Implementation of coculture strategies on the patterned 3D scaffolds also revealed improvements in the functional aspects of the intestinal epithelium (studies discussed in section "Functional behavior of the cells on 3D intestinal models") [54, 62, 63]. Moreover, treatment of butyrate in 3D colonic model inhibited cryptic proliferation, while inducing differentiation into ALP+ absorptive lineage [20, 46, 47].



Exploring avenues in induced pluripotent stem cells (iPSCs)-derived intestinal cells

Intestinal crypts have gained considerable attention for the development of in vitro intestinal tissue models with biomimetic architecture, cellular distribution, and physiological functionality. However, their clinical utility is limited by lack of ideal sources, tedious isolation protocols, the rapid loss of functionality, and existing ethical issues. Therefore, recent research interventions have been directed toward the use of iPSCs, which overcome many of these limitations. Such a strategy has also been advantageous for developing patient-specific intestinal models (healthy/diseased) and corresponding personalized therapies. Moreover, humanspecific pharmacokinetic features, closely mimicking the native conditions, could be achieved with the derived intestinal cells, which is otherwise difficult with the available human cell line alternatives. The current section highlights the recent developments in stem cell engineering for deriving intestinal cells. 2D cultures and 3D organoid cultures have majorly been employed for stem cell differentiation and the expansion of intestinal cells; thereafter, these differentiated cells could be harvested and used for different applications including in vitro model development.

In a study, human iPSCs were differentiated into enterocytes-like cells (ELCs) following a step-by-step protocol [119]. iPSCs were first committed toward definite endodermal-like cells (DELCs) lineage using Activin A, followed by differentiation into intestinal progenitor-like cells (IPLCs) using FGF2. The developed IPLCs expressed caudal type homeobox 2 (CDX2, a marker for intestinal development and differentiation) and LGR5 (ISC marker). Further, maturation of IPLCs into ELCs occurred in the presence of epidermal growth factor (EGF) under low serum (2%) conditions. The differentiated cells expressed multiple mature enterocytes markers; however, their expression was relatively lower than adult small intestinal tissue. The study also revealed that Wnt3a treatment had negligible, while FGF4 (instead of FGF2) treatment or the concentration of serum had a pronounced effect on differentiation efficiencies. In an extension to this study, authors conducted iPSCs to IPLCs differentiation following the same Activin A/FGF2 protocol; however, for generating ELCs, different cocktail of small molecules (glycogen synthase kinase 3 inhibitor XV, Dorsomorphin, PD98059, 5-aza-2'-deoxycytidine (5-aza), and A-83-01) were added along with EGF treatment [120]. The study revealed that resultant ELCs exhibited pharmacokinetic functions and showed the expression of mature enterocytes; however, the functional phenotype of the ELCs varied significantly among the treatment groups. In the same backdrop, iPSCderived DELCs were efficiently differentiated into IPLCs using a cocktail of 6-Bromoindirubin-3'-oxime (BIO) and



Besides the directed differentiation of iPSCs to specific intestinal cell types, multicellular intestinal tissues have also been developed using organoid culture technology. Sequential treatment of iPSCs with Activin A and a growth factor cocktail (FGF4 and Wnt3A, at high concentration) generated 3D spheroids containing CDX2+hindgut-like cells (HLCs) [122]. The study revealed that individually neither FGF4 nor Wnt3A could cause DELCs to HLCs differentiation. Also, the time of exposure and the growth factor concentration affected the stable expression of CDX2. Further, these HLC spheroids were transferred to Matrigel and cultured for long-term in medium containing another cocktail (EGF, Noggin, and R-spondin-1) to direct differentiation into small intestinal organoids (SIOs). The developed SIOs contained functional enterocytes, goblet, Paneth, and enteroendocrine cells. In a different study, a distinct population of SIOs and colonic organoids (COs) were generated [123]. The authors employed specific treatment of Activin A, FGF4, and CHIR99021 to develop CDX2+HLC spheroids. The spheroids were encapsulated in Matrigel and exposed to either SIO (containing EGF, Noggin) or COs (containing EGF, bone morphogenetic proteins (BMP)) differentiation medium. The resultant SIOs and COs exhibited differential phenotype, similar to the native tissues. The study revealed the regulatory role of BMP signaling in stable expression of special AT-rich sequence-binding protein 2 (SATB2, a definitive marker of the presumptive large intestinal epithelium) and activation posterior homeobox (HOX) genes to direct the differentiation of iPSC-derived HLC spheroids to COs.

Human intestinal organoids were generated from iPSCs following a multistep protocol involving differentiation into DELCs (Activin A, Wnt3A), HLCs (FGF4, CHIR99021), and SIOs (EGF, Noggin, CHIR99021) [124]. The differentiated cells from the organoids were cultured in a microfluidic platform with continuous media perfusion. The study revealed that the presence of physiologically relevant shear stress resulted in intestinal morphogenesis, and showed the presence of Paneth, goblet, enterocytes, enteroendocrine cells as well as LGR5+stem cell and WD Repeat Domain 43 (WDR43)+transit-amplifying cell population. Moreover, the developed intestine-on-chip model showed responsiveness toward exogenous stimuli (IFN- γ and TNF- α), with in vivo–like patterns.

Specific merit of iPSCs is that they also offer an endless opportunity to recapitulate patient-specific pathobiological



Table 3 Additional studies demonstrating generation of intestinal cells from human iPSCs

References		[127]	[128]	[129]
Research highlights		The resultant cells exhibited SI epithelial cells (SIECs)-like functions and expression profile. Differentiated cells predicted drug permeability, better than Caco-2 cells. The authors upbeat high differentiation potential of the reported methods	The study screened different growth factors and small molecules for their effectiveness in differentiating iPSCs to IPLCs and ELCs. The differentiated cells exhibited functions similar to SIECs; with a low expression of colonic epithelial cell (CECs) markers	SI-like function and gene expression profiles were observed. Organoids showed the presence of different intestinal cells (enterocytes, ISCs, goblet, enteroendocrine, Paneth, smooth muscle cells, and fibroblasts)
	COs	1	1	ggin, EGF, – 8059,
	SIOs	1	I	R-spondin1, Noggin, EGF, A-83-01, PD98059, 5-aza, DAPT
	HLCs	I	1	CHIR99021, FGF4
entiation induction into	ELCs	EGF, forskolin, PD98059, 5-aza, A-83-01	LY2090314 Wnt3A, R-spondin3, Noggin, SB202190, Dex, EGF, IGF-1	1
cules for differe	IPLCs	FGF2	LY2090314	1
S. no. Growth factors/small molecules for differentiation induction into	DELCs	Activin A, BMP4, VEGF, FGF2 FGF2	Wnt3A, Activin A	Activin A
S. no.		-	7	ю



Table	Table 3 (continued)						
S. no.	S. no. Growth factors/small molecules for differentiation induction into	ecules for dif	ferentiation induction into			Research highlights	References
	DELCs	IPLCs	ELCs	HLCs	SIOs COs		
4	CHIR99021	, i	Ť.	CHIR99021, FGF4, Noggin	Noggin, R-spondin, EGF, – MMP-8 inhibitor	Direct on-chip differentiation of iPSCs was done in a step-by-step manner. The developed intestinal tubules expressed markers for enterocytes, Paneth, and neuroendocrine cells. They also developed barrier function and showed responsiveness toward pro-inflammatory cytokines	[130]
'n	Activin A, CHIR99021	1	1	Activin A, CHIR99021, FGF4	R-spondin1, Noggin, EGF –	Developed organoids expressed markers of ISC, enterocytes, goblet, and Paneth cells. Retinoic acid treatment for 48 h, promoted iPSC differentiation to intestinal organoids	[131]
9	Activin A	1	1	CHIR99021, FGF4	EGF	Differentiated hindgut cells were cocultured with iPSC-derived vagal neural crest cells in Matrigel, to form intestinal organoids with the enteric nervous system	[132]
r	Activin A	1	1	FGF4, Wnt3A	R-spondin1, Noggin, EGF –	The developed organoids expressed markers of enterocytes, goblet, and Paneth cells. Mesenchymal cell marker expression was also observed. Organoids were maintained for long-term and exhibited drug-metabolizing enzyme activity	[133]



Table 3 (continued)

S. no.	S. no. Growth factors/small molecules for differentiation induction into	ules for diff	ferentiation induction into				Research highlights	References
	DELCs	IPLCs	ELCs	HLCs	SIOs	SOO		
∞	Activin A, Wnt3A, FGF2	ı	ı	FGF4, Wnt3A	R-spondin1, Noggin, EGF	I	The developed organoids expressed markers of enterocytes, goblet cells, enteroendocrine, Paneth, and proliferative cells	[134]
6	Activin A, CHIR99021	1	1	FGF4, CHIR99021		EGF, CHIR99021, LDN193189	Developed COs showed phenotypic characteristics, specific to the adult human colon, respectively. COs were employed for modeling colorectal cancers and drug screening	[135]
10	Activin A, BMP4, FGF2, LYS294002, CHIR99021	1	1	Activin A, FGF2, CHIR99021, retinoic acid	R-spondin1, Noggin, EGF, CHIR99021, Y-27632, prostaglandin E ₂	1	The developed organoids expressed markers of enterocytes, goblet, enteroendocrine, and Paneth cells. Organoids showed responsiveness toward Salmonella enterica serovar Typhimurium by expressing genes associated with infection and inflammation.	[136]

Dex Dexamethasone, VEGF vascular endothelial growth factor, SIECs SI epithelial cells, IGF-1 insulin-like growth factor 1, CECs colonic epithelial cell, MMP-8 matrix metalloproteinase-8



conditions in vitro. For instance, iPSCs from patients with cystic fibrosis (p.F508del mutant) were corrected using transcription activator-like effector nuclease (TALEN)mediated repair and further employed for generating intestinal organoids [125]. Differentiation protocol employed Activin A and Wnt3A for definite endoderm (DE) specification, Wnt3A and CHIR99021 for hindgut specification, and R-spondin1, Noggin, EGF, and Wnt3A for intestinal maturation (post-embedding in Matrigel). Organoids (both corrected and non-corrected), developed using this protocol, were able to grow without a need of mesenchymal niche, which is often present in iPSC-derived organoid culture, thus, facilitating their use in a forskolin-induced swelling assay. Moreover, the organoids from corrected iPSCs showed expression of cystic fibrosis transmembrane conductance regulator (CFTR) and functional recovery of its activity, as compared to non-corrected ones. In a separate study, patient-specific SIOs and COs were generated by differentiation of iPSCs, derived from healthy and inflammatory bowel disease (IBD) patients (adult-onset and very early onset IBD) [126]. The developed organoids were used for in vitro assessment of pro-inflammatory cytokine-induced alterations in the barrier functions of healthy and diseased patients.

Additional studies in this domain are summarized in Table 3.

Conclusion and outlook

The present article depicts superiority of the intestinal tissue models with biomimetic 3D architecture (crypt-villus or crypt-lumen) over the conventional 2D models; in reflecting native-like apicobasolateral cell polarity, functional phenotype (xenobiotic metabolism and TEER values), and cellular compartmentalization (differentiated cells at the villus/luminal end, while less differentiated/stem cell niches in the crypts). However, the future of these patterned intestinal models lies in integrating native-like biochemical cues (such as matrix composition and biochemical gradients), biophysical cues (such as matrix stiffness), mechanobiological aspects (such as shear stress and peristaltic motion), and opting for suitable coculture strategies.

In this regard, microfabrication methodologies (like photolithography and laser ablation) have a limited scope of development as they are majorly employed for creating the primary molds to mirror the 3D intestinal pattern in the bulk hydrogels. Apart from primary mold fabrication, control over the distribution of biochemical and biophysical cues as well as cellular components is quite challenging. Interestingly, 3D printing could prove to be quite handy and advantageous in this context. The recent developments in

tissue engineering and regenerative medicines have already validated that 3D bioprinting provides better spatiotemporal control over the distribution of different cues and cellular components [59, 137, 138]. 3D printing strategy is still in its infancy for the development of 3D intestinal models; in fact, to date, only SLA [58] and VMEPS (a type of extrusion-based 3D printing) [61] have only been employed. However, exploring newer modalities such as scaffold-free bioprinting [139], freeform bioprinting in suspension bath [140], digital light processing (DLP)-based bioprinting (an advanced version of SLA) [141], aspiration-assisted bioprinting [142], and 4D bioprinting [143] could further widen our horizon and ensure a promising future ahead.

Besides, innovative strategies for integrating native-like fluid flow and peristalsis with these 3D patterned models are also needed. Bioreactor [55] and microfluidic [72] technologies have already been explored to address these avenues; however, they still exhibit limited clinical applicability. Notably, working with hydrogel-based 3D tissue constructs is challenging as they are fragile and often undergo disintegration under continuous mechanical stress. A critical consideration should also be paid to the biodegradation susceptibility of these hydrogel-based platforms. Improving the mechanical properties of the hydrogel system (via crosslinking) is a potent strategy, but the impact of these biomechanical cues on intestinal cell behavior should be examined as well. Moreover, current research in this domain is mostly limited to use of either human cancer cell lines or intestinal crypts (derived from rats or mouse), which cannot accurately predict drug pharmacokinetics that occur in the human system. Owing to the source and ethical limitations associated with human ISCs, the use of patient-derived iPSCs, instead, may add precision to these studies. Besides, developing strategies for high-throughput platforms with multifaceted applicability is also needed and demands profound investigation.

On a concluding note, we upbeat that endeavors guided toward integrating biomimetic architectural cues with the necessary biophysicochemical and mechanical cues along with the aspects of stem cell engineering would accord for more physiologically relevant intestinal model and could further open up new vistas in the domain of personalized medicine in the near future. Breakthrough in this field could be witnessed at faster rates via dedicated collaborative efforts from engineers, material scientists, biologists, and clinicians. Policymakers and stakeholders are also an essential part of this collaborative network.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval This article does not contain any studies with human or animal subjects performed by any of the authors.

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