

Where We Stand: Lung Organotypic Living Systems That Emulate Human-Relevant Host–Environment/Pathogen Interactions

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Jimenez-Valdes RJ, Can UI, Niemeyer BF and Benam KH (2020) Where We Stand: Lung Organotypic Living Systems That Emulate Human-Relevant Host–Environment/Pathogen Interactions. Front. Bioeng. Biotechnol. 8:989. doi: 10.3389/fbioe.2020.00989 Lung disorders such as chronic obstructive pulmonary disease (COPD) and lower respiratory tract infections (LRTIs) are leading causes of death in humans globally. Cigarette smoking is the principal risk factor for the development of COPD, and LRTIs are caused by inhaling respiratory pathogens. Thus, a thorough understanding of host–environment/pathogen interactions is crucial to developing effective preventive and therapeutic modalities against these disorders. While animal models of human pulmonary conditions have been widely utilized, they suffer major drawbacks due to inter-species differences, hindering clinical translation. Here we summarize recent advances in generating complex 3D culture systems that emulate the microarchitecture and pathophysiology of the human lung, and how these platforms have been implemented for studying exposure to environmental factors, airborne pathogens, and therapeutic agents.

Keywords: spheroids, organoids, bio-scaffolds, precision cut lung slices, bronchial biopsies, 3D bioprinting, lungon-a-chip, inhalation models

INTRODUCTION

Lung disorders represent a great socioeconomic challenge and a major burden on health care systems worldwide; chronic obstructive pulmonary disease (COPD) and lower respiratory tract infections (LRTIs) are the third and fourth leading causes of death globally, respectively (World Health Organization [WHO], 2018). Cigarette smoke exposure is the principal risk factor for COPD development and exacerbation, and LRTIs are consequent to inhalation of airborne pathogens. Therefore, a better understanding of the pulmonary system at health and disease and recreation of respiratory host–environment/pathogen interactions are crucial if we are to prevent and treat lung disorders. While animal models of human respiratory conditions have been instrumental in advancing our knowledge, they suffer major drawbacks due to inter-species (e.g., genetic, homeostatic physiology, pathology, and respiratory tree anatomy, airway histology, etc.) differences hindering clinical translation. Rodent animal models, for instance, which have widely been applied to shed light on signaling networks and lung development, are fundamentally different from

humans in terms of lung cellular composition, airway branching, and immune system function (Rackley and Stripp, 2012). Besides, rodents are obligate nose breathers with intricate and highly developed nasal turbinates that lead to different particulate deposition patterns compared to humans during breathing (Hecht, 2005). Due to these factors, rodents at best are passive inhalers and, unlike humans, are unable to actively inhale therapeutic candidates or cigarette smoke.

To circumvent some of these shortcomings with animal models, simple two-dimensional (2D) cell culture models that emulate naturally present lung air-liquid interface (ALI) have been developed to study human lung pathophysiology in vitro (Prunieras et al., 1983). Recently, these systems have been used in combination with cell exposure systems that mimic the natural route of exposure of lung epithelium to environmental factors in order to measure the risks of potentially damaging inhaled nanoparticles (Lenz et al., 2009) or cigarette smoke (Thorne and Adamson, 2013). However, despite their use of primary human lung airway epithelial cells (hAECs) and recapitulation of mucociliated histology, they have several limitations in recreating organ-level functionalities of the human lung. For example, they do not enable (1) real-time analysis of dynamic intercellular (e.g., leukocyte-endothelial cell) interactions under physiological flow, (2) study of inhalation exposure to whole cigarette smoke, as a representative inhaled material, under physiological breathing airflow without disturbing ALI, (3) recreation of blood-like vascular perfusion to continually supply nutrients and growth factors, (4) introduction and perturbation of vascular and airflow shear stress to simulate different pathophysiological conditions, and (5) high-resolution kinetic analysis of biological responses (e.g., the time-course release of secreted factors, like cytokines/chemokines, in response to pathogenic challenge) (Fang and Eglen, 2017; Ainslie et al., 2019). In addition, these platforms are commonly used in the absence of sub-epithelial extracellular matrix (ECM; Yamaya et al., 1992; Gray et al., 1996).

Recent advances in tissue engineering and microfabrication have led to the development and application of more complex 3D culture systems and biomimetic microfluidic platforms to capture the structural and functional complexity of the human lung (Miller and Spence, 2017). In this review, we assess these stateof-the-art complex *in vitro* lung models, and the efforts made to expose these models to environmental factors, pathogens, and therapeutic agents. Finally, we discuss future directions on tackling the challenges of *in vitro* lung models that were observed over the past decade and share our vision on how to further enhance these models for more accurate and better clinical translation.

RECENT ADVANCES IN CREATING COMPLEX 3D IN VITRO MODELS OF HUMAN LUNG

Spheroids and Organoids

Among the simplest of the 3D lung models are spheroids. Spheroids are multicellular sphere-like culture systems that

enable recapitulation of cell-cell interactions; they are either utilized as freely floating cell aggregates or get seeded onto 3D bio-scaffolds (Ainslie et al., 2019). Spheroids utilize tumorderived cells, patient-derived xenograft cells, and immortalized cell lines, and have been applied to study intercellular interactions and response to therapies in the context of cancer (Ekert et al., 2014; Meenach et al., 2016; Klameth et al., 2017; Lewis et al., 2018), lung progenitor cells differentiation (Chimenti et al., 2017), and pulmonary fibrosis (Surolia et al., 2017). However, spheroid culture systems still suffer deficiencies such as a lack of vasculature and ALI, hard to control cell ratios and aggregate size, as well as a failure to mimic organ function (Fang and Eglen, 2017); therefore, spheroids do not represent good in vitro lung models to study host-environment/pathogen interactions. Another similar yet distinct preclinical system is organoids; these are cultured organ-specific cell types, which are derived from a population of stem cells (adult or pluripotent), and are capable of maintaining stem cells during in vitro culture. During formation, organoids develop into 3D tissues that recreate in vivo-observed microanatomy through self-organization (Fang and Eglen, 2017; Ainslie et al., 2019). Compared to spheroids, organoids exhibit long-term viability and rely on internal developmental processes (rather than cell-cell adhesions) to drive tissue/organ-like microarchitecture formation. Human lung organoids have been generated to replicate bronchi/bronchioles (Konishi et al., 2016; Mccauley et al., 2017; Miller et al., 2018), alveoli (Zacharias et al., 2018), and even multi-lineage structures (Dye et al., 2016). While helpful in advancing our understanding of the respiratory system, particularly from a developmental perspective, this models lack several of key organ features such as vascularization, and mechanical forces associated with breathing, and it is difficult to obtain fully differentiated lung cell types (Barkauskas et al., 2017). Lung organoids have been sparsely used for hostenvironment/pathogen exposure studies, since the organoid lumen, which represents the apical surface of the lungs through which a natural exposure would occur in vivo, is difficult to access, thus treatments or stimuli are applied in the environment in which the organoids are embedded. In this scenario, murine organoids have been infected with influenza virus (Quantius et al., 2016), and human organoids have been infected with respiratory syncytial virus (RSV; Sachs et al., 2019). Some attempts have been made to microinject pathogens or their byproducts into organoids using custom made tools, primarily in gastrointestinal organoids (Williamson et al., 2018); however, in lung organoids, this technique has only been applied to propagate Cryptosporidium, a protozoan parasite (Heo et al., 2018).

Precision Cut Lung Slices

To address the disconnection that often exists in spheroids and organoids in including both cellular and organ-level complexities *in vitro*, some groups have turned to an *ex vivo* system called precision cut lung slices (PCLS), which maintain the cellular structure and the biological processes of the lung (Ainslie et al., 2019). Importantly, PCLS generation from healthy explants and diseased tissue can reveal differences in the cellular and molecular interactions within the microenvironment of the lung. Disease modeling of healthy tissue can be achieved *ex vivo*

through mimicking disease characteristics and used for exploring therapeutic treatments (Alsafadi et al., 2020). Precision cut lung slices have been applied for cytotoxicity assessment to lowmolecular-weight (LMW) chemicals (Lauenstein et al., 2014), evaluation of biological responses to inflammatory stimuli such as lipopolysaccharide endotoxin (LPS), macrophage-activating lipopeptide-2 (MALP-2), interferon gamma (IFNgamma) (Henjakovic et al., 2008), and recreation of pathologies such as pulmonary fibrosis, through the use of a combination of profibrotic growth factors and signaling molecules (Alsafadi et al., 2017) and COPD models induced with 1-(3-Hydroxy-5-(thiophen-2-yl)phenyl)-3-(naphthalen-2-yl)urea (FzM1) (Skronska-Wasek et al., 2017). However, PCLS are often cultivated only for short periods of time (they are very challenging to maintain viable and functional for weeks), their use is impacted by the limited availability of whole/resected organ for slicing, and they only represent a brief snapshot of the cell populations in the tissue. Besides, the PCLS cannot fully recapitulate all attributes seen in vivo; for instance, there are major challenges with preserving arteriole morphology and localization of lung airway smooth muscle cells (Sanderson, 2011). Precision cut lung slices can reproduce the initial interactions and inflammatory responses to industrial chemicals (Lauenstein et al., 2014) and pathogens [e.g., influenza viruses (Liu et al., 2015), rhinovirus (Beale et al., 2014), Yersinia pestis (Banerjee et al., 2019), and Coxiella burnetiid (Graham et al., 2016)] in the human respiratory system; however, the extent to which the immune response can replicate the in vivo situation is limited, as demonstrated by Neuhaus et al. (2017), who observed that LPS-induced tumor necrosis factor-alpha (TNF- α) secretion decreased significantly over a period of 15 days, and it is not possible to recruit non-resident immune cells. Moreover, while murine-derived PCLS models have been adapted to study the effects of cigarette smoke (Donovan et al., 2016), the PCLS per se have not been utilized for inhalation exposure studies in vitro, principally because the route of administration represents a challenge (e.g., physiological ALI cannot be established in this model system), and often the entire slice is bathed in the compound or stimulant of interest (Liu et al., 2019). Changing from ALI conditions to liquidsubmerged conditions represents a drawback, as it impacts epithelial cells' glycoprotein secretion profile, tight junction integrity, and permeability (Grainger et al., 2006). In addition, the physio-chemical properties of pulmonary stimuli such as airborne pollutants, gaseous substances, novel medications, and therapeutic agents can alter when in liquid suspension (Upadhyay and Palmberg, 2018). In addition to this, PCLS become static systems, lacking the shear flows associated with air and blood flow in the ALI and vascular endothelium, respectively, which could impact cell-cell/ECM responses and interactions in tissues.

Bronchial Biopses

Bronchial biopsies are samples of airway tissue obtained from the carinae of large and small cartilaginous airways. These samples are often fixed and used to measure airway remodeling by morphometric analyses of airway epithelial mucin stores, measurements of reticular basement membrane thickness, quantification of the number and size of globlet cells, the analysis of the content and density of the smooth muscle of the airways, as well as quantification and characterization of inflammatory cells (Woodruff and Innes, 2006). In addition, some effort have been made to culture the bronchial biopsies for short-term exposure studies in vitro. For instance, this culture model has been used to study the effects of chemotherapeutic treatments on non-small cell lung cancer (Lang et al., 2007), to evaluate cellular responses by exposing asthmatic lung tissue to allergen Dermatophagoides pteronyssinus (Jaffar et al., 1999; Lordan et al., 2001; Vijayanand et al., 2010), and to test the antiviral efficacy of therapeutic drugs against influenza virus infection (Nicholas et al., 2015). The advantages of bronchial biopsies include retaining the three-dimensional structure of the lungs if the extraction procedure is successful, which requires a trained bronchoscopist, allowing sampling from healthy subjects as well as patients with lung diseases, and the minimally invasive nature of the sampling procedure. However, this system has similar drawbacks as the PCLS, and is impacted by scarcity of donor tissue availability, small sample size (usually 1-3 mm) and number of biopses that can be obtained at any given time $(\sim 4-10)$. In addition the *in vitro* viability in often low, therefore exposure studies on bronchial biopsies usually do not last more than a single day.

Cellularized Bio-Scaffolds

Extracellular matrix is the 3D network of extracellular macromolecules that provide structural integrity and biochemical support to tissue-resident cells. Extracellular matrix both serves as a scaffold and modulates cellular responses such as self-renewal, quiescence, migration, proliferation, phenotype maintenance, differentiation, and apoptosis (Akhmanova et al., 2015). Hydrogels can mimic ECM in vitro using natural products (like collagen, hyaluronic acid, chitosan, alginate, or Matrigel) or synthetic polymers [such as polyethylene glycol (PEG) or polyacrylamide (PAA)] (Zhou et al., 2018). Natural hydrogels have been predominantly used to culture lung cells and organoids (Sato et al., 2017; Zacharias et al., 2018); however, artificial matrices have been demonstrated as a viable alternative (Miller et al., 2019), since their components are biologically inert, and can be functionalized by the addition of proteins, peptides, and/or polysaccharides (Akhmanova et al., 2015).

Besides hydrogels, lung ECM can be generated via decellularization of the whole/resected lung through perfusion with detergents, which allows considerable preservation of the micro- and macro-architecture of the organ, and the ECM composition (Gilpin and Wagner, 2018). Extracellular matrix has been recognized as a bioactive medium that modulates cellular responses in its surroundings. By using cellularized bio-scaffolds it will therefore be possible to mimic some pathological alterations, and the consequent functional changes that occur in lung diseases, especially those with a chronic nature (Burgess et al., 2016). Decellularized lung scaffolds have enabled studying cell-ECM interactions in tissues of healthy people and patients with chronic lung diseases, such as lung cancer (Stratmann et al., 2014), idiopathic pulmonary fibrosis

(IPF; Van Der Velden et al., 2018), and scleroderma (Sun et al., 2016). These matrices are instrumental in evaluating lung repair and regeneration and have been recellularized with different types of human cells, including fibroblasts, endothelial cells, epithelial cells, stem cells from multiple organ sources, and organoids (Porzionato et al., 2018; Giobbe et al., 2019). However, it has not been possible to completely recellularize these matrices to allow recreation of lung physiology and function, their application has been partly hindered due to the limited number of whole/resected human lungs, and to our knowledge, their utilization [beyond regenerative medicine and orthotopic transplantation (Guenthart et al., 2019)] for studying host–environment/pathogen interactions has not been demonstrated.

3D Bioprinting

To more faithfully replicate tissue microarchitectures of the human lung, 3D bioprinting has emerged as a great technological platform ideally positioned to create finely defined and controlled biological structures and living systems by utilizing a wide array of materials (natural, synthetic, or even hybrid hydrogels) and cell types. This technology can achieve a high degree of precision in cell positioning and can be scalable for automation (Gungor-Ozkerim et al., 2018). However, 3D bioprinting has been predominantly applied for the development of vascularized tissues (Kolesky et al., 2014), rather than generating functional whole organ or organ-like mimicries. To our knowledge, only a few studies have focused on the lungs. Horvath et al. (2015) biofabricated a three-part mimetic (composed of an endothelial cell layer, basement membrane, and an epithelial cell layer) of airblood barrier analog of the human lung alveoli within a transwell insert. While a great initiative, the study utilized cell lines, rather than primary human-derived cells, and the authors only focused on model validation (viability, cellular proliferation, and barrier function) and no data were presented on the application of the platform for the analysis of host-environment/pathogen interactions. More recently, a new method for 3D bioprinting of tissues by stereolithography was reported, whereby it was possible to make a model inspired by alveolar morphology, capable of withstanding mechanical strain for cyclical ventilation and oxygen transport (Grigoryan et al., 2019). While successful in creating entangled vascular networks of the lung alveoli, the bioprinted tissue lacked alveolar epithelial cells, and the platform was applied neither for toxicological studies nor for inhaled exposure to a respiratory pathogen or therapeutic agents. Altogether, 3D bioprinting offers the possibility of developing biologically inspired lung organotypic models; however, despite the advantages of 3D bioprinting, wide spread utilization of these techniques has been limited by the requirement of expensive and complex technologies as well as a steep learning curve.

Lung-on-a-Chip Models

Organ-on-a-Chips are biomimetic, microfluidic, cell culture devices created with microchip manufacturing methods that contain continuously perfused hollow microchannels inhabited by living tissue cells arranged to simulate organ-level physiology (Bhatia and Ingber, 2014; Benam et al., 2015). By recapitulating the multicellular architectures, tissue–tissue interfaces, chemical gradients, mechanical cues, and vascular perfusion of the body, these devices produce levels of tissue and organ functionality with well-defined structures and highly controlled microenvironments that would not be possible with conventional 2D or 3D culture systems. They also enable high-resolution, real-time imaging and *in vitro* analysis of biochemical, genetic and metabolic activities of living human cells in a functional human tissue and organ context. Adaptation of Organ-on-a-Chip technology by pulmonary scientists has led to the development of model systems that emulate human lung pathophysiology *in vitro*.

Lung-on-a-Chip devices have been developed to model different regions of the respiratory system and recapitulate in vivo-observed multicellular architecture and physicochemical environment of the lung (airway and alveoli). The use of this technology (Alveolus-on-a-Chip) has enabled a better understanding of different aspects of human alveolar pathologies; such as evaluating the influence of breathing-associated mechanical cues on the growth and migration pattern of Non-Small Cell Lung Cancer tumor cells (NSCLC; Hassell et al., 2017), reproducing interleukin-2 (IL-2) induced alveolar edema observed in human cancer patients at similar doses and over the same time frame, and its pharmacological inhibition (Huh et al., 2012), mimicking pulmonary thrombosis by treatment with inflammatory stimuli (TNF- α) and bacterial products (LPS) corroborated in murine models (Jain et al., 2018), studying inhaled exposure of the alveolar epithelium to aerosolized LPS and its immune-modulatory impact (Artzy-Schnirman et al., 2019), and recreating recruitment of circulating leukocytes and inflammation induction after treatment of epithelial cells with TNF-a, Escherichia coli, and silica nanoparticles (Huh et al., 2010). In these microphysiological systems primary, cancerous or immortalized cell line of human alveolar epithelial origin have been co-cultured with human lung microvascular endothelial cells (Huh et al., 2010, 2012; Hassell et al., 2017) or human umbilical vascular endothelial cells (HUVECs; Jain et al., 2018).

In the context of conducting airways, we have developed and applied Lung-on-a-Chip devices (Airway-on-a-Chip) for culture and differentiation of primary human epithelial cells in co-culture with primary human lung microvascular endothelial cells, to model small airways, reproduce the mechanical forces associated with inhalation-exhalation respiration cycles (airflow shear), mimic the shear forces across endothelial cell layers (via vascular flow), and recreate the ALI barrier of human airways (Benam et al., 2015, 2016a,b, 2017, 2018, 2019, 2015; Benam and Ingber, 2016; Niemeyer et al., 2018). This technology has been adapted to allow first-in-kind studies that reproduce and characterize tissue-tissue crosstalk between pulmonary epithelium and airway smooth muscle (Humayun et al., 2018) and enable better understanding infectious disease biogenesis processes through analysis of the production of inflammatory cytokines and immune cell recruitment (neutrophils) following exposure to respiratory fungi (e.g., Aspergillus fumigatus) and the bacteria (e.g., Pseudomonas aeruginosa) (Barkal et al., 2017) by other investigators. Similay utilizing Airway-on-a-Chip we have been able to dissect inflammatory processes involved in the pathobiology of chronic lung conditions such as asthma induced by the exposure of ephitelium to IL-13 and COPD by recreating terminally differentiated mucociliated airway epithelia on-chip, and exposing cells to the viral mimic polyinosinic-polycytidylic acid (poly(I:C)) or to LPS (Benam et al., 2016b), and identify novel biomarkers of human lung diseases and test efficacy of lead therapeutic compounds (Benam et al., 2016b). We developed a Breathing-Smoking Lung-on-a-Chip platform that recreates smoke-induced pathologies in humans (Benam et al., 2016b). The system consisted of a Lung Small Airway-on-a-Chip (Benam et al., 2016b) that reproduces the living bronchiolar tissue for exposure to inhaled whole cigarette smoke (WCS), a "microrespirator" that emulates diaphragm and rib cage function and reproduces inhalation-exhalation airflow at physiological rhythms and patterns, a "biomimetic smoking robot" (Benam et al., 2020) that mimics human mouth and generates fresh WCS



FIGURE 1 | Advantages and disadvantages of complex 3D *in vitro* culture models of the human lung for inhalation exposure studies. Each model presented has unique benefits and challenges. Improved airway modeling can be achieved through incorporating inhalation exposure systems. The digital images of human lung at the center and bronchus near the bottom left were acquired from Shutterstock and Biorender, respectively.

and regulates the passage of inhale/exhale air/smoke, and software that represents the brain of human smoker and controls "smoking topography" and "breathing" behavior. Using this platform, we flowed WCS horizontally across the apical surface of differentiated human bronchiolar epithelia (as occurs in the lungs of human smokers), from normal subjects and COPD patients, at representative smoking topography parameters for puff time, puff volume and interpuff interval under clinically relevant breathing conditions. We found that inhaled smoke exposure transformed ciliary beat frequency (CBF) pattern, range, and variability and induced oxidative stress. Importantly, it was observed that submerging the ciliated airway epithelia with cell culture medium alone was sufficient to mask the impact of inhaled WCS on ciliary micro-pathologies. In addition, we discovered new smoke-mounted biological markers that distinguish COPD epithelia from those from healthy individuals (Benam et al., 2016a).

Recently, Elias-Kirma et al. (2020) developed an interesting model of branching Airway-on-a-Chip for toxicological evaluation of inhaled particulate matter. The authors utilized primary human bronchial epithelial cells for culture, and exposed the epithelium on-chip to anti-vanilloid receptor 1 (VR1) antibody-coated polystyrene microparticles. The microparticles were generated using a commercial aerosolizing machine and brought into contact with the cells through a pinch valve-controlled antistatic tube. While the device design in this study was unique, the exposure did not mimic what naturally occurs during rhythmic breathing cycles and the authors did not study biological impact of exposure to environmental pollutants, tobacco products or therapeutic agents.

These study implies that Lung-on-a-Chip platforms have the potential for recreation of in vivo-like inhaled exposure to pathogens, environmental threats, and drug treatments. However, for such platforms to be more applicable and physiologically relevant, it is important to enhance their multicellularity and complexity, for instance by the addition of parenchymal cells, such as fibroblasts, tissue-resident and circulating immune cells, muscle cells, and/or ECM. One of the drawbacks of microfluidics is that manufacturing is often expensive and time consuming; in addition, instruments such as syringe pumps or air pressure systems are required to control fluid flow within the chips. Furthermore, the devices need to become more user friendly (particularly for biologists with little/no engineering expertise) and be amenable for increased throughput. On the other hand, although microphysiological systems have great flexibility with respect to the experimental design, there are important challenges due to miniaturization, such as reduced media volumes, different media exchange rates and methods, and lower number of cells tested; finally, and an additional challenge in conducting hostpathogen/environment studies is the stimulus dose. Therefore, it is necessary to carefully evaluate these differences when comparing cell behavior and viability in microfluidic devices versus macroscopic cultures, and especially when comparing against in vivo results to validate the systems. Moreover,

from a high-level perspective investigators utilizing Organon-Chip devices must consider as accurate and as feaible as possible validating and qualifying these models against tissues that they are aiming to replicate - i.e., human samples and clinical data. Lastly, we would like to mention that multiple factors have contributed to popularity and quick and widespread adaptation of PDMS for fabrication of Lungon-a-Chip microfluidic devices. These include ability for rapid prototyping and multilayer device fabrication, optical transparency from 240 to 1100 nm (enabling utilization by various optical detection schemes), flexibility (allowing axial stretch), gas-permeablity (so that oxygen can feasibly penetrate the device and reach the cells embedded within the chip), being non-toxic, inexpensive and not breaking (like glass). However, a drawback of PDMS has been its absorption of small, hydrophobic molecules from flowing solution (Toepke and Beebe, 2006). To mitigate this, we apply either of these two approaches: (1) correcting for absorption of small hydrophobic molecules by quantifying the loss (e.g., via mass spectrometry) and (2) saturating the chips prior to cell culture or experimentation to minimize loss.

CONCLUSION AND FUTURE DIRECTIONS

In this article, we reviewed the most recent advances in creating preclinical 3D model systems that reproduce human lung pathophysiology in vitro and discussed their respective merits and drawbacks in the context of host-environment/pathogen interactions (Figure 1). It has been a long road for the transition from 2D to 3D cell culture systems; however, 3D models are gaining popularity in use and adaptation due to their critical advantages over static cell cultures. Spheroids, organoids, and PCLS have proven to be invaluable tools for studying lung development and pathologies, but they lack natural ALI and are unable to reproduce breathing-associated airflow, thus it is not possible to perform inhalation exposure studies using these platforms. Cellularized scaffolds have been widely used to study cell-ECM interactions at health and disease, and for regenerative purposes; however, these have not yet been used to assess host-environment/pathogen interactions. Recently, 3D bioprinting has emerged as a promising technique for developing more intricate lung living systems with great potential for high throughput yield; but numerous technical challenges need to be tackled and complex-system building to integrate multiple organ-specific cells and tissue (beyond generating vascularized constructs) must become a priority. In contrast, microfluidic Lung-on-a-Chip microdevices have emerged as an attractive alternative to mimic lung function and biology as seen in vivo. In the particular case of lung models, and more specifically in the study of the host-environment/pathogen interaction, it is important how the treatments and/or stimuli are delivered, in this sense, the microfluidic platforms have demonstrated to be very robust and physiologically relevant in vitro tools that can incorporate biochemical, physical and mechanical cues. We envision considerable improvement and adaption of Lung-on-a-Chip platforms for the study of hostenvironment/pathogen interactions. The improvements include allowing higher throughput analysis, and integration of ECM and additional cellular complexity. These microdevices will be utilized for the study of nebulized and dry powder treatments and recreation of clinically relevant exposure to airborne respiratory pathogens (instead of lung cells being exposed submerged to the infective agents). We also anticipate the *in vitro* cell culture of cellularized matrices and potentially PCLS within the Lung-on-a-Chip system to provide dynamism and enhanced viability *ex vivo*. In line with this, 3D bioprinting techniques could revolutionize the way microfluidic devices are currently manufactured, allowing for increased production and reproducibility.

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AUTHOR CONTRIBUTIONS

KB conceptualized and critically revised the manuscript. RJ-V, UC, and BN drafted the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: KB is the founder and holds equity in Pneumax, LLC. KB and BN are co-inventors on patent applications describing Lung Chip and related technologies.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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