



Lymph Nodes-On-Chip: Promising Immune Platforms for Pharmacological and Toxicological Applications

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Specialty section:

This article was submitted to
Translational Pharmacology,
a section of the journal
Frontiers in Pharmacology

Received: 18 May 2021

Accepted: 04 August 2021

Published: 16 August 2021

Citation:

Shanti A, Hallfors N, Petroianu GA, Planelles L and Stefanini C (2021) Lymph Nodes-On-Chip: Promising Immune Platforms for Pharmacological and Toxicological Applications. *Front. Pharmacol.* 12:711307. doi: 10.3389/fphar.2021.711307

Organs-on-chip are gaining increasing attention as promising platforms for drug screening and testing applications. However, lymph nodes-on-chip options remain limited although the lymph node is one of the main determinants of the immunotoxicity of newly developed pharmacological drugs. In this review, we describe existing biomimetic lymph nodes-on-chip, their design, and their physiological relevance to pharmacology and shed the light on future directions associated with lymph node-on-chip design and implementation in drug discovery and development.

Keywords: organs-on-chip, lymph node, immunity, pharmacology, toxicology, lymph node-on-chip

INTRODUCTION

Organs-on-chip have recently emerged as powerful screening tools in the field of pharmacology and toxicology (Maharjan et al., 2020). Such chips enable more accurate evaluation of the safety and efficacy of investigational drugs compared to traditional tissue culture plates and animal models (Low et al., 2020). This is particularly important since the pharmaceutical industry faces high attrition rates for novel drug candidates (Roberts et al., 2014). It is estimated that only 1 out of 10,000 new chemical entities gains FDA approval and transitions it into the market (Nam et al., 2015). Additionally, the overall failure rate in drug development is reported to be over 96%, including a 90% failure rate during clinical development (Hingorani et al., 2019). These numbers, indeed, pinpoint that the current drug development practice suffers from major drawbacks, among which, is the lack of physiologically relevant platforms for drug screening during preclinical evaluation phases (Liu et al., 2017). Organs-on-chip, therefore, provide promising alternatives to conventional preclinical drug screening platforms.

Organs-on-chip are perfused microfluidic chips composed of micro-sized channels wherein cells of a single type or more are allowed to grow and expand in an *in vivo* like microenvironment. They are not intended to construct whole living organs but instead to build the functional units of the organs that are capable of replicating specific aspects of human physiology *in vitro* (Richardson et al., 2020). In particular, they tend to replicate 1) the 3D microarchitecture of the organ, that is, the spatial distribution of its different cellular types; 2) the biochemical microenvironment of the organ including its extracellular matrix as well as its chemokine, growth factor and nutrient gradients, if any; 3) mechanical microenvironment of the organ including mechanical compressions, cyclic strains and shear stresses and 4) particular function of the organ, for example, filtration, respiration or digestion (Shanti

et al., 2018). Once an organ-on-chip is established, it can then be used to investigate organ physiology both in health and in disease.

Organs-on-chip can be fabricated using various engineering techniques such as micromolding, microetching, soft lithography and photo-polymerization (Shanti et al., 2018). These techniques have enabled the development of extremely intricate structures that closely resemble those *in vivo* and these include the villi of the small intestine and the alveoli of the lungs (Shim et al., 2017; Stucki et al., 2018).

Over the past decade, many research groups have developed different organs-on-chip and demonstrated their ability to be used in pharmacological, toxicological and drug testing applications. For instance, Hassell and others developed a lung-on-chip platform that replicates lung's microenvironment-specific cancer growth and lung's response to tyrosine kinase inhibitor therapy (Hassell et al., 2017). In addition, Vernetti et al. developed a liver-on-chip platform containing primary human hepatocytes along with human endothelial cells, immune cells and stellate cells and assessed the toxicity of various molecules (troglitazone, nimesulide, caffeine, trovafloxacin, levofloxacin, LPS and methotrexate) on the developed liver model (Vernetti et al., 2016). Moreover, Jalili-Firoozinezhad et al. developed a gut-on-chip platform lined by human intestinal epithelial cells and vascular endothelial cells, utilized it to model radiation injury and then deployed it to assess the efficacy of a radiation countermeasure drug (prolylhydroxylase inhibitor) (Jalili-Firoozinezhad et al., 2018). Furthermore, Qian et al. developed a heart-on-chip platform capable of recording cardiac tissue adhesion, electrophysiology, and contractility and used it to study the effect of norepinephrine on the heart (Qian et al., 2017). Still, many other organ-on-chip platforms remain and have been utilized to investigate different processes such as mechanism of action of drugs, pharmacokinetics and pharmacodynamics, efficacy, toxicity and dose response (Richardson et al., 2020).

Despite the development of many organs-on-chip and their utilization in pharmacological and toxicological contexts, only recently has the need for developing a lymph node-on-chip been recognized. This comes after acknowledging the fact that the immune outcome from the lymph node is a key determinant of not only the response to viruses, bacteria, and other foreign particles but also of the immunotoxicity of drugs, i.e. toxicity of the drugs to the immune system (Galarza et al., 2020). Lymph nodes-on-chip can facilitate investigations into the mechanisms of interaction between immune cells and drug candidates thus, unveiling significant knowledge to cut down the high cost of drug development as well as to reduce the high attrition rate.

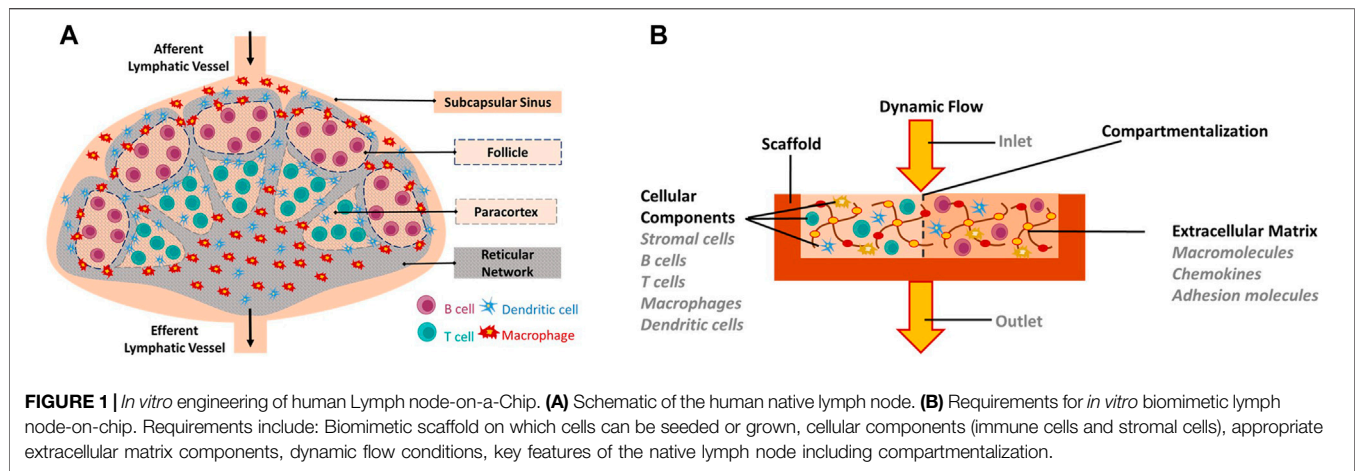
In this review, we describe the human native lymph node, summarize existing biomimetic lymph nodes-on-chip, their design, and their physiological relevance to pharmacology and shed the light on future directions associated with lymph node-on-chip design and implementation in drug discovery and development.

HUMAN NATIVE LYMPH NODE

Lymph nodes are vital organs of the immune system where efficient and protective immune responses are generated and maintained (Gasteiger et al., 2016; Novkovic et al., 2020). Lymph nodes collect and filter the fluid that drains from peripheral tissues, known as the lymphatic fluid, before it is eventually returned into blood circulation (Fullerton and Gilroy, 2016). This process is essential to achieve an efficient adaptive immune response and to protect the body against foreign bodies such as bacteria and viruses.

Lymph nodes provide unique structural microenvironments to support efficient gathering of immunogenic material from peripheral tissues (**Figure 1**) (Sainte-Marie, 2010; Schudel et al., 2019; Shanti et al., 2020). They are compartmentalized into distinct cellular micro-domains populated by a single type of lymphocytes, either B lymphocytes or T lymphocytes (Crivellato et al., 2004; Cupedo et al., 2012). B lymphocytes populate an area known as the primary follicle within the outer cortex of the lymph node whereas T lymphocytes populate an area known as the paracortex. When B lymphocytes residing in the follicles are activated by an antigen, they initiate the formation of germinal centers, which are dynamic microenvironments where B lymphocytes proliferate, differentiate and generate high affinity antibodies necessary for an effective immune response (Domeier et al., 2017).

Not all molecules entering the lymph node have free access to the distinct cellular compartments (follicle and paracortex). Instead, when the lymphatic fluid enters the lymph node through the afferent lymphatic vessel, it is directed to an area known as the subcapsular sinus (SCS) rather than being directly directed to the cellular regions (von Andrian and Mempel, 2003). SCS contains macrophages that sample the lymphatic fluid and remove any microorganisms and debris found in it. In addition, macrophages process antigenic material found in the lymphatic fluid and present it to B and T lymphocytes residing in their specific compartments (von Andrian and Mempel, 2003). As the lymphatic fluid flows through the SCS, a small fraction of it is diverted laterally into a conduit system formed by reticular collagen fibrils that penetrate the B follicle and reach deep into the paracortex (Jafarnejad et al., 2015; Shanti et al., 2020). The conduit system delivers antigens into the follicle and paracortex (Gretz et al., 2000; Liao and von der Weid, 2015; Hughes et al., 2016). In addition, it constitutes a scaffold on which immune cells can migrate and acts as a filtration system (Katakai, 2004). It has been shown that the conduit system only allows small molecules, whose diameter is between 7.2 and 10.7 nm to pass through and penetrate into the follicle or paracortex (Gretz et al., 2000). In their specific compartments, T and B lymphocytes scan the incoming molecules and/or cells for their cognate antigens. If such antigens are not found, cells receive homeostatic survival signals (Cupedo et al., 2012). Conversely, if their cognate antigens are found, cells are activated and induced to proliferate and differentiate: T-cells into cytotoxic ($CD8^+$) effector T-cells, helper ($CD4^+$) effector T-cells or memory cells and B-cells into plasma cells or memory cells (Shanti et al., 2018). Most activated cells then leave the lymph node and head towards



the site of infection to eliminate the foreign antigens (Pape et al., 2007). Remaining lymphatic fluid and soluble molecules, which are not directed into the conduit system, are transported through a number of small trabecular sinuses and medullary sinuses, after which they leave the LN through efferent lymphatic vessels.

The main extracellular component of the lymph nodes is collagen Type I (Kaldjian et al., 2001; Wiig et al., 2010). Furthermore, the organization of the lymph node into distinct cellular regions is controlled by various chemokines and adhesion molecules such as Chemokine (C-C motif) ligand 19 (CCL19), Chemokine (C-C motif) ligand 21 (CCL21), C-X-C Motif Chemokine Ligand 12 (CXCL12) and the selectively chemotactic ligand for B cells CXCL13 (B-cell-attracting chemokine 1) expressed on stromal cells (Cyster, 1999; Katakai, 2004; Lian and Luster, 2015). Besides the secretion of chemokines and adhesion molecules, stromal cells provide functional support to the lymph node (Katakai, 2004). In particular, fibroblastic reticular cells produce various extracellular matrix components and interweave them to make reticular fibers, which in turn, provide mechanical strength to the lymph node, allow migration of immune cells from one area to another and form the conduit system necessary for the filtration of the lymphatic fluid (Gretz et al., 1996, 1997; Mcknight and Gordon, 1998). Dynamic migration processes in the lymph node enable cellular communication between antigen presenting cells (APC), B cells and T cells, which is critical in the generation of the adaptive immune response (Allen et al., 2007).

TECHNOLOGIES FOR GENERATING LYMPH NODES-ON-CHIP

The design and development of an *in vitro* lymph node model has several aspects to be considered and these include: the development of a biomimetic scaffold on which immune cells can be seeded or grown, the generation of a microenvironment with appropriate extracellular matrix components, the production of dynamic flow conditions and the replication of key functional features of the native lymph node (Figure 1). Therefore, researchers have employed various technologies for reproducing the lymph node *in-vitro*. To achieve a biomimetic

scaffold on which cells can be seeded or grown, many relied on soft lithography of SU-8 to create molded polydimethylsiloxane (PDMS) microfluidic devices (Nandagopal et al., 2011; Mitra et al., 2013; Moura Rosa et al., 2016; Parlato et al., 2017; Ross and Pompano, 2018; Shim et al., 2019). These devices can be single a layer bonded to glass, or multiple layers, bonded together to form a central channel. In addition to silicone, some designs utilized other materials such as gelatin, polysulfone, or agarose to form fluidic pathways (Giese et al., 2006, Giese et al., 2010; Haessler et al., 2009). Microfluidics, in turn, enabled the execution of tasks not possible in larger-scale environments. It allowed the establishment of chemical gradients for chemotaxis studies, enabled the selective capture of small populations of cells for time-lapse microscopy and allowed the measurement of tissue diffusion mechanics (Haessler et al., 2009; Zaretsky et al., 2012; Mitra et al., 2013; Moura Rosa et al., 2016; Ross and Pompano, 2018). Still, the ability to select semi-permeable channel material allowed for broader diffusion and a chemical gradient establishment in the surrounding medium (Haessler et al., 2009).

To achieve a microenvironment that mimics the extracellular matrix of the lymph node, three dimensional hydrogels have been used. These create more biologically relevant conditions *in-vitro*, with the choice of gel material having direct chemical and functional outcomes on cell behavior. The ideal biomaterials should have similar stiffness than the target tissue, degradability, and equivalent ability to support the binding of soluble factors, cellular adhesion, cellular communication, and cellular movement. Extracellular matrix is commonly replicated using hydrogels based on poly-ethylene glycol (PEG), agarose, or collagen I, the last being the main component of the extracellular matrix in the naïve lymph node. Depending on the selection, the hydrogel can be improved by adding other molecules. For example, PEG has been functionalized with maleimide (maleic acid and imide) and incorporated with collagen peptides to improve its binding and adhesion properties (Graney et al., 2020). Fibronectin has been also widely used to recreate the extracellular matrix due to its ability to bind multiple proteins such as proteoglycans, growth factors, integrins and chemokines (Pelletier et al., 2000; Woods et al., 2000; Martino and Hubbell, 2010; Rana et al., 2015).

To achieve dynamic conditions, lymph nodes on-chip require active fluidic perfusion. Thus, many researchers utilized peristaltic pumps to flow cell medium and suspended biological material into their developed models (Giese et al., 2006, 2010; Moura Rosa et al., 2016; Sardi et al., 2016; Shim et al., 2019). Active pumping often induces cellular cross-talk by the transport of chemokines or cell markers (Parlato et al., 2017; Polini et al., 2019; Shim et al., 2019).

Another important aspect in the design of a biomimetic lymph node is its ability to support the co-culture and spatial distribution of the different cell types that populate this organ (mainly immune T and B cells, DCs, and stromal cells) in a similar way to how they are organized in the human model. This represents a big challenge that, if achieved, considerably increases the complexity of the design, the experimental conditions and the data analysis. Most of the studies, so far, have used only one or two immune cell types in their devices. The source of the cellular components is also an important point to consider. Some devices are validated using immortalized human cell lines as they are easy to obtain, manipulate and expand, are cost effective and allow experimental reproducibility (Moura Rosa et al., 2016; Birmingham et al., 2020; Shanti et al., 2020; Hallfors et al., 2021). Cell lines are well established models that can be easily modified to target and study molecules of interest. The use of cell lines to mimic *in vitro* what is occurring *in vivo* in the lymph node has nevertheless clear limitations as immortalized cells fail to fully reproduce morphological and functional characteristics of primary cells. For example, cell lines can't be used to fully replicate humoral immunity in which naïve B cells need to proliferate, differentiate and become antibody secreting cells. These limitations can be circumvented by the use of primary immune cells commonly purified from human blood and also tissue slices from human lymph nodes (Lin and Butcher, 2006; Higbee et al., 2009; Giese et al., 2010; Dauner et al., 2017; Radke et al., 2017; Goyal et al., 2019; Kraus et al., 2019). These attractive models, which, in fact, are still under development, attempt to be closer to the physiological behavior of the human lymph node. However, they are also challenging in several aspects such as limitation in the amount of material, short lifespan, specific requirements for cell culture, and donor variability. Special attention should be given to human leukocyte antigen (HLA) -mismatch when using co-cultures of cells from different donors, as it could lead to cell activation and misinterpretation of the results.

We will describe next the most comprehensive Lymph node on chip platforms found in the literature and the technology behind.

Bioreactor Human Artificial Lymph Node

An attempt to model the human lymph node is a commercial perfusion bioreactor system named the Human Artificial Lymph Node reactor (HuALN[®], ProBioGen AG, Berlin, Germany) or HIRISTMIII bioreactor (Giese et al., 2006). It is made of polysulfone (PS), has two culture compartments separated by a double layer of gas-permeable membranes and, two fluidic systems, one for culture medium and one for suspended cells. The HuALN[®] enables long term culture of cells embedded in agarose

matrix and the establishment of well-defined gradients. Authors showed this model is beneficial to study immune reactions and drug responses because they were able to detect micro-organoid structures (by histology and *in situ* imaging), to analyze genetic profiles and, to quantify cytokine and immunoglobulin secretion in the collected cell culture medium (Giese et al., 2010; Radke et al., 2017; Kraus et al., 2019). This model has been used to test glycoprotein vaccines and to study the immunogenicity of protein aggregates of two antibodies, bevacizumab and adalimumab, exposed at different stress conditions (Radke et al., 2017; Kraus et al., 2019).

Multicompartment Lymph Node-on-a-Chip Model

Recently, our research group at the Healthcare Engineering Innovation Center in Khalifa University, UAE has also developed a novel microfluidic platform replicating multiple key features of the lymph node aimed at facilitating biological investigations of immune cellular kinetics, cell-cell interactions, cell-drug interactions and sampling (Shanti et al., 2020; Hallfors et al., 2021). The developed lymph node-on-chip recreates 1) the spatial microenvironment of the native lymph node, 2) the compartmentalization of immune cells within distinct regions 3) the extracellular microenvironment of the native lymph node 4) the fluid flow within the native lymph node and 5) several functional features of the native lymph node. It is a multi-compartmentalized bioreactor consisting of an elliptical body with multiple compartments resembling the different cellular regions of the native lymph node, an inlet aperture resembling the afferent lymphatic vessel and an outlet aperture resembling the efferent lymphatic vessel. The cellular regions are filled with 3D hydrogels encapsulating immune cells. These hydrogels are collagen based and have morphology, porosity, stiffness, and permeability comparable to that of the native human lymph node and sustain cellular viability for duration sufficient for immunotoxicity studies. To demonstrate its feasibility for pharmacological and toxicological applications, we utilized the lymph node-on-chip to assess the effect of a pharmaceutical drug, namely hydroxychloroquine, on the motility of immune cells and their production of reactive oxygen species. We showed that hydroxychloroquine reduced T cell velocity, promoted persistent rotational motion and increased levels of reactive oxygen species. These results highlight the potential of our lymph-node-on-a-chip to be used in pharmacological applications. They also demonstrate that the lymph node-on-a-chip is an attractive alternative to further dissect and understand the mechanisms of action of drugs delivered to cells. The LN-on-chip is intended for identifying comparative behaviour of immune cells in response to drugs and for recognizing patterns of action.

Modular Immune *In vitro* Construct

Another promising immune platform, Modular Immune *In vitro* Construct (MIMIC), was developed by Warren et al. for evaluating the immune response to vaccines (Warren et al., 2005, 2007). The platform consists of a peripheral tissue equivalent mimicking the skin from which the vaccine is

TABLE 1 | Summary of existing attempts to develop *in vitro* lymph nodes on-chip.

| Replicated feature of lymph node | Lymph node Scaffolding material | ECM material | Microfluidics | Cellular component | Application | References |
|---|--|--|---------------------|---|--|---|
| Chemotaxis and Chemokine Diffusion | Agarose | -Collagen hydrogel -Matrigel | Yes, active pumping | Mouse bone-marrow derived dendritic cells | Dendritic cell chemotaxis Recreation of chemokine gradients (CCL21 and CCL19) | Haessler et al. (2009), Haessler et al. (2011) |
| | Photolithography-patterned PDMS | -Heparan Sulfate Proteoglycans No hydrogel Fibronectin coating of surface | Yes, active pumping | Mouse bone-marrow derived dendritic cells | Dendritic cell chemotaxis. Recreation of single and competing chemokine gradients (CCL21, CCL19 CXCL12) | Ricart et al. (2011) |
| | Photolithography-patterned PDMS | No hydrogel Fibronectin coating of surface | Yes, active pumping | Human T cells from blood | -Blood derived T cells migration towards controlled gradients of CCL19 and CXCL12 | Lin and Butcher, (2006) |
| | Photolithography-patterned PDMS | No hydrogel | Yes, active pumping | Mouse lymph node slices | Effective diffusion of cytokines in live mouse lymph node slices (TNF α , IL2 and IFN γ) | Ross and Pompano, (2018) |
| Subcapsular Sinus Dynamics | Adhesive microchannel affixed to a PDMS block previously cured in a polystyrene tissue culture plate | No hydrogel | Yes, active pumping | Thp1 human monocytic cell line LS174T human colon cancer cell line | Effect of subcapsular sinus biophysical (flow and structure) and biochemical (adhesion molecule expression) remodeling on cellular adhesion | Birmingham et al. (2020) |
| | | Portion of channel was functionalized by Fc specific anti-IgG Plates were blocked with BSA | | PANC-1 human pancreatic cell line | | |
| Immune cellular interactions | Photolithography-patterned PDMS | No hydrogel | Yes, active pumping | Mouse primary T and B cells | Assay molecular events during lymphocyte activation Interaction between OT-I CD8 T cells and SIINFEKL-loaded MHCII-eGFP B cells | Dura et al. (2015) |
| | | Surface blocking using bovine serum albumin or pluronic F127 | | | Activation dynamics of CD8 OT-1 T cells and TRP1 transnuclear T cells | |
| | Photolithography-patterned PDMS | Collagen and fibronectin | Yes, active pumping | -MutuDC: Mouse dendritic cell line -MF2.2D, OVAII: Mouse CD4 T cell line -RF33.70/OVAI: Mouse CD8 T cell line | Dynamic interaction of flowing lymphocytes with adherent dendritic cells Effects of low and high shear stress variations on adhesion | Moura Rosa et al. (2016) |
| Germinal center formation and antibody production | Not applicable | -Gelatin activated with silicate nanoparticles -Functionalized poly-ethylene glycol (PEG) that incorporates adhesive peptides | No | -Mouse primary B cells -Fibroblastic cell line expressing CD40L and BAFF molecules | Immuno-engineered organoids to mimic germinal center formation, class switching and antibody production | Purwada et al. (2015), Purwada et al. (2017); Purwada et al. (2019); Béguelin et al., (2017); Purwada and Singh (2017); Graney et al. (2020) |
| | Photolithography-patterned PDMS | - Gel composed of Matrigel and collagen I | Yes, active pumping | - Human T and B cells purified from blood | Mimic germinal center formation, class switching and antibody production | Goyal et al. (2019) |
| Lymph Node Malignancies | Photolithography-patterned PDMS | Thiol-modified Hyaluronic acid, gelatin and PEG | Yes, active pumping | -MLMVECs mouse lung microvascular endothelial cell line -A20 mouse B-cell lymphoma cell line. Cells used from | Model that replicates a blood vessel within a tumor and allows the study of complex interactions between immune cells and endothelial cells in the | Mannino et al. (2017) |

(Continued on following page)

TABLE 1 | (Continued) Summary of existing attempts to develop *in vitro* lymph nodes on-chip.

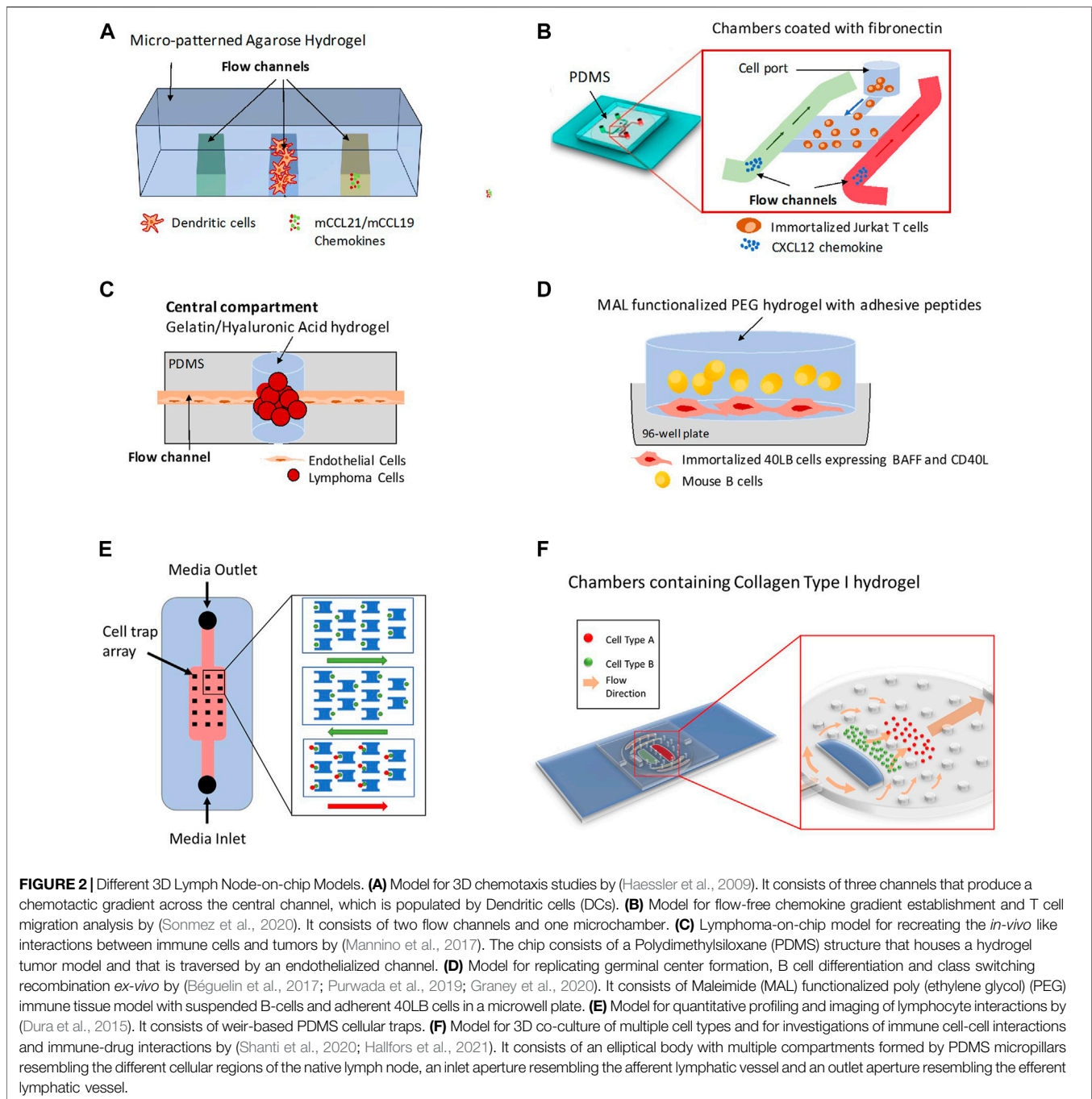
| Replicated feature of lymph node | Lymph node Scaffolding material | ECM material | Microfluidics | Cellular component | Application | References |
|----------------------------------|---------------------------------|--------------|-------------------------------------|--|---|---|
| | | | | explanted A20 tumors in mice -Mouse primary CD3 ⁺ T cells and CD11b ⁺ myeloid cells from spleen | lymphoma microenvironment | |
| | Photolithography-patterned PDMS | No hydrogel | Yes, active pumping | -Mouse lymph node slices -Murine 4T1 breast tumors slices | Model that replicates communication between tumor and lymph node Analysis of protein release and capture between two organs | Shim et al. (2019) |
| Chemical Stimulation | Photolithography-patterned PDMS | Agarose | Yes, active pumping | Mouse lymph node slices | Recreate stimulus-response behavior of the lymph node | Ross et al. (2017) |
| Response to Drugs and vaccines | Photolithography-patterned PDMS | Collagen | Yes, active pumping | -Jurkat human T cell line -Raji human B cell line -Thp1 human monocytic cell line | Model that replicates lymph node architecture, ECM components and flow Analysis of cell interactions and drug effects on cell dynamics | Shanti et al. (2020); Hallfors et al. (2021) |
| | -Polysulfone | Agarose | Yes active pumping (medium and gas) | -Human T and B cells purified from blood | HIRIS™III bioreactor that mimics long term interactions (14 days) between suspended lymphocytes and adherent dendritic cells | Giese et al. (2006); Giese et al. (2010); Sardi et al. (2016); Radke et al. (2017); Kraus et al. (2019) |
| | -Polycarbonate base plate | | | -Human dendritic cells derived from blood monocytes -Mesenchymal stromal cells (MSCs) | Used to test glycoprotein vaccines and the immunogenicity of protein aggregates | |
| | Transwell | - | No | - Human T and B cells purified from blood | Immune response to vaccines including seasonal trivalent influenza vaccine and Tetanus vaccine | Warren et al. (2005); Warren et al. (2007); Higbee et al. (2009); Drake et al. (2012); Dauner et al. (2017) |

introduced, a lymphoid tissue equivalent mimicking the lymph node where interactions between immune cells and vaccine components take place and a functional assay module that allows the assessment of the output of the immune response. Researchers utilized MIMIC platform for evaluating immune response to several vaccines including seasonal trivalent influenza vaccine and Tetanus vaccine (Byers et al., 2009; Higbee et al., 2009; Drake et al., 2012; Dauner et al., 2017). In their study with the influenza vaccine, Dauner et al. used blood derived immune cells from young and relatively old human donors and demonstrated the ability of MIMIC platform to recapitulate differential age-associated response to the vaccine. In particular, they showed that immune cells from elderly population exhibited reduced antibody IgG production and reduced CD154 + IFN γ + IL-2+TNF α + CD4⁺ T activation upon stimulation with the vaccine compared to young

population. Authors concluded that MIMIC platform is an attractive tool to reduce the cost and developmental time of new vaccines and therapeutics.

SELECTIVE FEATURES RECREATED BY LYMPH NODE ON CHIP MODELS

Up to the current time, lymph node on-chip models have only recapitulated selective features of the human lymph node. In the following sections, we describe attempts to recreate aspects of the lymph nodes on-chip and shed the light on future directions associated with lymph node-on-chip design and implementation in drug discovery and development. There are other organ-on-chip models that are immune related and that hold promising potential for investigations related to the immune system



(Stachowiak and Irvine, 2008; Jones et al., 2012; Dura et al., 2015; Sun et al., 2019; Fathi et al., 2020; Bachmann et al., 2021) but we will focus on models that mimic the lymph node physiology. **Table 1** summarizes existing attempts to develop *in vitro* lymph nodes-on-chip. **Figure 2** shows some lymph node on-chip models.

Chemotaxis and Chemokine Diffusion

Initial studies using 3D microfluidic platforms that partially replicated the human lymph node were focused on

characterizing cellular migration. Chemokines and chemokine receptors are key molecules expressed by multiple cell types in the lymph node such as T lymphocytes, B lymphocytes, and DCs. These molecules not only support cellular organization but also guide immune cells during the immune response to allow them to move between the outer zone where antigens are acquired and the areas where B and T cells are activated.

In 2009, Haessler et al. described an agarose-based microfluidic platform with a gradient buffer for 3D chemotaxis studies (Haessler et al., 2009). This platform had a

central cell culture compartment and two side flow channels. DCs were embedded in the central region in a collagen-matrigel-proteoglycans mixture and chemokines and/or media were flowed along the side channels of the device to recreate, in a controlled manner, well defined chemokine gradients. Using this device, authors found that the bone marrow derived murine DCs preferentially migrate towards Chemokine (C-C motif) ligand 21 (CCL21) versus Chemokine (C-C motif) ligand 19 (CCL19) when they are exposed to equivalent and opposite gradients (Haessler et al., 2009). CCL21 and CCL19 are the two endogenous ligands for the chemokine receptor CCR7.

The integration of different chemotactic signals by DCs was also addressed by Ricart et al. through another microfluidic device that similarly allowed the recreation of single and competing chemokine gradients (Ricart et al., 2011). It also enabled the analysis of the effects of actin and myosin inhibitors, as well as pertussis toxin on DC chemotactic migration. This device, described by the authors previously to evaluate neutrophil migration, used fibronectin to favor cell adhesion (Jannat et al., 2010). Authors found that CCL19 signal was more potent than CCL21 or C-X-C motif chemokine 12 (CXCL12) (ligand for receptors CXCR4 and CXCR7, later renamed ACKR3 implying it belongs to the atypical chemokine receptors).

Unlike Haessler et al. and Ricart et al. who studied migration of DCs in microfluidic platforms, Lin et al. were among the first to study blood derived T cells migration towards controlled gradients of CCL19 and CXCL12 using a microfluidic device coated with fibronectin (Lin and Butcher, 2006). Similarly, but more recently, Sonmez et al. developed another PDMS-based microfluidic system able to establish flow-free chemokine gradient patterns and allow T cell migration analysis by time-lapse imaging (Sonmez et al., 2020). Their device had two flow channels and one microchamber for cell culture with the chambers coated with fibronectin to mimic extracellular matrix. Using this system, they generated flow-free concentration gradient by the diffusion of the chemotactic response of Jurkat T cells to CXCL12.

In a different setting, Ross et al. reported a three-layer device to quantify the effective diffusion of cytokines (TNF α , IL2 and IFN γ) in live mouse lymph node slices placed in the cell chamber (Ross and Pompano, 2018). Diffusion was monitored through imaging by time lapse experiments. This model might be applied to study drug delivery and diffusion through the lymph node extracellular matrix.

Subcapsular Sinus Dynamics

In a different attempt, Birmingham et al. developed a novel microfluidic device to investigate the effect of subcapsular sinus remodeling (seen in inflammation) both biophysically (flow and structure) and biochemically (adhesion molecule expression) on human monocytic and metastatic cellular adhesion (Birmingham et al., 2020). The microfluidic device consists of a divergent adhesive microchannel cut from a 125 μ m thick double sided adhesive tape and affixed to a PDMS block previously cured in a polystyrene tissue culture plate. The cells incorporated in the model include THP-1 human monocytic cell line as well as LS174T human colon cancer cell line

and PANC-1 pancreatic cell line. The authors showed that decreased wall shear stress usually seen in flow regimes associated with inflamed lymph nodes influences the trajectories and extent of E-selectin mediated cancer cell adhesion. Interestingly, they showed that adhesion of LS174T cells to E-selectin is enhanced by co-perfusion with THP-1 cells. Furthermore, authors demonstrated that adhesion molecules, intercellular adhesion molecule (ICAM) and vascular cell adhesion molecule (VCAM) may enable cancer cell homing to the lymph nodes through the modulation of the extent and dynamics of cellular adhesion. Taken together, these results emphasize that lymph node remodeling plays a significant role in cancer invasion and lymphatic metastasis.

Immune Cellular Interactions

Microfluidic devices have been also used to understand the cellular interactions and the physiological environment in the lymph node. Dura et al. developed a unique microfluidic device that trapped cells at a pair level and at a single cell level and allowed quantitative profiling and imaging of lymphocyte interactions (Dura et al., 2015). The device comprised an array of weir-based PDMS traps in a flow channel with each trap consisting of a front-side two-cell capture cup and back side single-cell capture cup. Furthermore, the device was treated with 7.5% bovine serum albumin or 10% pluronic F127. The device was utilized to assay the early molecular events during lymphocyte activation, to study the interaction between OT-I CD8 T cells and SIINFEKL-loaded MHCII-eGFP B cells and to characterize early activation dynamics of CD8 OT-1 T cells and TRP1 transnuclear T cells.

In a different attempt, Rosa et al., developed a microfluidic platform to study, in real-time, the dynamic interaction of flowing lymphocytes with adherent DCs (Moura Rosa et al., 2016). The platform, a PDMS based biochip placed on a glass substrate, had one main flow channel, two inlets and two outlets. The channel was coated with collagen or fibronectin to support cell adhesion. This device allowed the study of T cell-DC interactions and antigen-specific attachment of T cells to DCs and to modify in a controlled manner different parameters such as velocity, shear stress, deformation rates and migratory motility.

Germinal Center Formation and Antibody Production

Currently, there are no experimental models that fully recapitulate the human humoral response in an *ex-vivo* lymph node microenvironment, probably due to the complexity of B cell differentiation and affinity maturation process in the germinal center reaction. The laboratory of Ankur Singh which has recently moved to Georgia Institute of Technology in the United States, has extensive work in the field; his group has generated mouse 3D synthetic immune organoids capable of regulating B cell development kinetics and of undergoing germinal center reactions that lead to the differentiation of antigen specific and high affinity germinal center B cells (Purwada et al., 2015, 2019; Purwada and Singh, 2017). Immune organoids are grown in 96-well plates and consist of

mouse B primary cells embedded in a functionalized polymer; a Gelatin matrix activated with silicate nanoparticles or a four-armed poly (ethylene glycol) macromer that has maleimide groups at each terminus (PEG-4MAL) and that incorporates several adhesive peptides including Collagen-1 peptides. 3T3 (“3-days transfer, inoculum 3×10^5 cells”) fibroblasts transduced with CD154, also called CD40 ligand (CD40L) and with B cell activating factor (BAFF) are included in the models to provide CD40L and BAFF signals which are secreted *in vivo* by stromal follicular dendritic cells (FDCs; (Purwada et al., 2019; Graney et al., 2020). Using their models, Singh et al. showed that B cells require “Enhancer of Zeste Homolog 2” (EZH2), a histone-lysine N-methyltransferase enzyme to form germinal centers and characterized the response against membrane bound versus soluble *Klebsiella pneumoniae* bacterial antigens (Béguelin et al., 2017; Graney et al., 2020). In addition, they demonstrated the role of integrin $\alpha 4 \beta 1$ and integrin $\alpha \beta 3$ -binding ligands in inducing GL7+ (GC-like) and GL7- (non-GC-like) phenotype in differentiating B cells (Purwada et al., 2017). This attractive *ex-vivo* model could be improved by adding a microfluidic technology that would increase its potential applications.

Similarly, the laboratory of Donald Ingber at Harvard University, which has extensive work on Organs-on-chip, developed a PDMS-based microfluidic platform that supports the formation of lymph node follicles and germinal centers (Goyal et al., 2019). The platform consisted of two channels separated by a porous membrane whereby the lower channel hosted primary human B and T cells cultured within a gel composed of Matrigel and collagen type I and the upper channel served as a perfusion channel, continuously supplying cells with oxygen and nutrients. Ingber et al. showed that the formed lymphoid follicles and germinal centers support the differentiation of plasma cells when activated with IL-4 and CD40 agonistic antibody or when activated with bacteria (inactivated *S. aureus* Cowan I). In addition, they demonstrated that the introduction of a quadrivalent split virion influenza vaccine into the lymph node platform resulted in plasma cell formation, viral strain-specific anti-hemagglutinin immunoglobulin G (IgG) production, and cytokine secretion similar to that seen in serum of vaccinated humans.

Lymph Node Malignancies

Some 3D models have focused on lymphoid malignancies with the aim of investigating the complex crosstalk between endothelial cells and immune tumor cells. Mannino et al reported a Lymphoma-on-a-chip model in which tumor cells were encapsulated in a hyaluronic acid-based hydrogel and traversed by a vascularized, perfusable, round microchannel (Mannino et al, 2017). This model would be useful for drug delivery studies as it allows the establishment of well-defined diffusion conditions and the evaluation of drug effects on endothelial cell permeability.

Shim et al. added a different degree of complexity to their developed microfluidic LN models by co-culturing lymph node tissue slices with tumor tissue slices in the same device to replicate communication between both organs on chip (Shim et al., 2019).

Culture media circulated through both *ex-vivo* tissue samples and allowed the exchange of soluble factors. Authors found that the model replicated certain features of immune suppression mediated by the tumor, such as T cell activation (Shim et al., 2019).

There is extensive and growing literature describing cancer on chip models that recreate the tumor microenvironment and its interaction with the immune system, recently reviewed in (Maulana et al., 2021). These models represent a novel and attractive approach, closer to the *in vivo* situation than the classical ones, to address important issues in the field such as the tumor immune infiltrate, the immunosuppression mediated by the tumor microenvironment and the check point inhibitors, the drug resistance and the identification and evaluation of therapeutic drugs. Despite its relevance, we don't describe these models here as it is out of the scope of the review, which is focused on lymph node on chips devices.

Chemical Stimulation

To recreate stimulus-response behavior of the lymph node, Ross et al. developed a microfluidic device that hosts murine lymph node tissue slices and stimulated the different regions of the slices with a model therapeutic agent, namely, glucose-conjugated albumin (Ross et al., 2017). The microfluidic device consisted of a 3 layer PDMS in which the first layer constituted microchannels layer, the second layer constituted ports layer and the third layer constituted perfusion chamber. The results of the study demonstrated that the retention of the therapeutic agent was greater in the B-cell zone than it was in the T-cell zone.

LYMPH NODE ON CHIP APPLICATIONS

Despite the fact that most of the developed *in vitro* lymph node models have not been directly employed in pharmacological and toxicological applications, they do hold a huge potential to revolutionize drug discovery and drug development. In particular, the technology they are based upon, i.e. microfluidic technology, has enabled the recreation of intricate anatomical features of organs that closely resemble those *in vivo* in a manner that has not been possible otherwise. In addition, it allowed the design of “plug and play” components that are biologically relevant, simple and easy to use (Donoghue et al., 2021). On the other hand, 3D hydrogels with various extracellular matrix components, coupled with microfluidics, enabled more accurate representations of the cell-cell and cell-environment interactions compared to those in 2D culture plates. In addition, the ability to perfuse microfluidic chips allowed the shift from simple static systems into more dynamic ones in which cell medium and suspended biological material can be flowed, captured and recirculated. All these features make lymph nodes on-chip models, promising alternatives to traditional tissue culture plates. In addition they represent improved and attractive platforms for various applications in immunology, pathology, pharmacology and toxicology, as we will next describe. Although the review here focuses on models

that mimic the lymph node physiology, there are other organ-on-chip models that are immune related and that hold promising potential for investigations related to the immune system (Stachowiak and Irvine, 2008; Jones et al., 2012; Dura et al., 2015; Sun et al., 2019; Fathi et al., 2020; Bachmann et al., 2021).

Immunotoxicity Testing

Testing of potential adverse effects of human pharmaceuticals on immune cells is becoming a standard practice in drug development. As the initial accumulation of immune cells and drugs is often at the lymph node, *in vitro* systems mimicking both the anatomy and physiology of *in vivo* lymph node, will allow mechanistic studies that investigate drug efficacy, efficiency, and toxicity. Cell movement and viability can be easily tracked in real time as seen in some of the existing models described here and other parameters including immune cell proliferation, activation or the expression of soluble factors in the media can also be monitored. Alterations in these, along with antigen presentation, T cell activation, B cell differentiation and antibody production can be used as indicative factors of the immune cellular response to drugs. Fortunately, *in vitro* biomimetic lymph nodes-on-chip are very useful not only for evaluating the immunotoxicity of new immunotherapies, but also for addressing the toxicity generated by drugs already in the clinic.

Immune Cellular Modulation

Lymph nodes-on-chip are of great benefit to medical fields that use drugs to target the immune system with the aim of modifying it and making it useful in the management of different disease conditions. In fact, immunotherapy has been used for long time in the treatment of allergy, to reduce the rejection of transplanted organs and to dampen autoimmunity (Boardman and Levings, 2019). Additionally, it has become a standard treatment for a variety of cancers. Cancer immunotherapy has been shown to induce durable responses in multiple solid and hematologic malignancies and the involvement of the draining lymph node is well documented (Fransen et al., 2018; van Pul et al., 2021). Moreover, it has been shown that immunotherapies also lead to unique toxicity profiles that are related to specific mechanisms of action (Kennedy and Salama, 2020). All these phenomena can be thoroughly investigated in physiologically relevant lymph nodes-on-chip platforms.

Disease Modelling and Investigation

Lymph nodes-on-chip are attractive platforms for investigating diseases in which the immune system is involved such as cancer and inflammatory diseases. Normal and diseased cells or tissues can be loaded into biomimetic lymph node-on-chip platforms and their behavior can be easily monitored. Mechanisms of action of diseases as well as risk factors can be explored and novel pathways or elements can be identified. In addition, lymph nodes-on-chip can be connected to other organs-on-chip thus facilitating studies aimed at investigating the relationship between lymph nodes and other organs. For instance, people with inflammatory bowel disease have been shown to be at an increased risk of developing autoimmune diseases and inflammatory diseases (Halling et al., 2017). Such interesting links could be explored in multi-organ-on-chip platforms

involving the lymph nodes which are known to host key immune cells. Such investigations provide a more systematic and holistic view of physiology, both in health and in disease.

Vaccine Development

One of the major drawbacks associated with vaccine development is the lack of reliable and sensitive platforms to evaluate vaccine formulations in a physiologically relevant preclinical setting (Drake et al., 2012). Thus, lymph node-on-chip platforms are quite promising and advantageous systems for developing novel vaccines. Primary immune cells obtained from human donors can be loaded into the biomimetic *in vitro* lymph node systems along with novel vaccine formulations and their behavior and response accurately monitored. In specific, humoral immunity, antibody production and B cell-T cell interactions in response to vaccinations can be quantified.

Personalized Medicine

While lymph node-on-chip models enable and accelerate drug and vaccine development processes, pending clinical acceptance, they also potentiate as platforms for personalized medicine. This is especially vital since genetic uniqueness might result in drug responses that differ across different populations. Patient immune cells can be obtained through blood sampling or minimally invasive liquid biopsy and be placed into developed lymph node-on-chip systems for evaluation. The possibility of including patient-derived immune cells in the device and replicating *ex-vivo* the cellular responses to different drugs opens new perspectives particularly in the precise and effective treatment of different diseases. For instance, resistance to cancer therapy remains one of the main challenges in the treatment of cancer (Mansoori et al., 2017). This phenomena can be addressed using lymph node-on-chip technology which allows for the direct incorporation of patient-derived cells in a physiologically relevant microenvironment and study of their specific cellular response to cancer therapy.

FUTURE DIRECTIONS

Future lymph node on-chip designs will need to incorporate multiple design elements from foundational work into a complete, multifunctional lymph node. Microfluidic pathways, 3D cellular matrix, co-culture of multiple cell types, chemotaxis and cellular communication, all supported within a biologically sustainable microenvironment will make future lymph node devices more biologically relevant than ever before. Such lymph node models will allow investigations into different immune cellular events and their association with drug discovery and vaccine development. For example, they could be used for examining effects of pharmaceutical drugs on immune cell motility, antigen presentation, T cell activation, B cell differentiation and antibody production in a more realistic 3D scenario.

However, thorough validation of lymph nodes-on chip against both animal and clinical trial results is required to determine the reliability of these models as predictive tools and to define the extent to which these models represent human-relevant physiology (Hwang et al., 2021). Much remains to be done in this field and

there are many opportunities to discover the possibilities lymph nodes on-chip can offer (Hwang et al., 2021). Yet, to ensure widespread adoption, it is necessary that the developed lymph node-on-chip platforms are easy to use, relatively inexpensive, and highly reproducible (Donoghue et al., 2021).

AUTHOR CONTRIBUTIONS

AS, LP: conceptualization and writing. AS, LP, NH: visualization. CS, GP: review and editing. CS: funding

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acquisition. All authors contributed to the article and approved the submitted version.

FUNDING

This work was funded by ADEK Award for Research Excellence (AARE) 2017, project name “AARE17-261, Biomimetic Lymph Node for Pharmaceutical Research” and Khalifa University of Science and Technology under Award No. RC2-2018-022 (HEIC).

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