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Review on new approach methods to gain insight into the feto-maternal interface physiology

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Non-human animals represent a large and important feature in the history of biomedical research. The validity of their use, in terms of reproducible outcomes and translational confidence to the human situation, as well as ethical concerns surrounding that use, have been and remain controversial topics. Over the last 10 years, the communities developing microphysiological systems (MPS) have produced new approach method (NAMs) such as organoids and organs-on-a-chip. These alternative methodologies have shown indications of greater reliability and translatability than animal use in some areas, represent more humane substitutions for animals in these settings, and – with continued scientific effort – may change the conduct of basic research, clinical studies, safety testing, and drug development. Here, we present an introduction to these more human-relevant methodologies and suggest how a suite of pregnancy associated feto-maternal interface system-oriented NAMs may be integrated as reliable partial-/full animal replacements for investigators, significantly aid animal-/environmental welfare, and improve healthcare outcomes.

KEYWORDS

microphysiology, organ on a chip, pregnancy, parturition, organoids

Introduction

Since at least the fourth century Before Current Era, non-human animals have represented a large and important feature of biomedical research (1). While that history has occasionally been punctuated by the scientific victimization of certain human populations (condemned criminals, victims of war crimes, slaves, and orphans), revulsion by and rejection of such human experimentation by the majority of cultures left non-human animals as the default organisms for studying human physiology and pathophysiology (1, 2). During the more than one-thousand years before general anesthesia, there is no question that the conduct of such studies involved a ghastly experience for the unfortunate non-human. It is unlikely that early investigators accepted the Cartesian view of animals as automatons (3), and more likely that they steeled their reasoning (if not sentiment) to these agonized animal subjects as the best experimental substitute for their own species. Over many centuries, different groups advocated for changes to – even abandonment of – this practice known as vivisection. Arguably, the most important legislation for protecting animals used in laboratory settings originated in the 19th-century (4, 5) and has continued to this date

(5). The recently ratified Food and Drug Administration (FDA) Modernization Act (6) contains an element of animal protection by strengthening consideration of validated non-animal methods in new drug regulatory submissions and by removing the requirement of animal testing for biosimilars. While no legislation has promoted either animal use or animal replacement, it is important to understand the advantages and limitations of each, if the best science is to be pursued.

Animal models

The history of animal use in research extends over more than one and one-half millennia, while microphysiological systems (MPS) are relative infants. Clearly, there has been far more opportunity to scrutinize the value of animal use than that of MPS and our comments herein solely focus on the scientific value of each methodology. Animals are complete biological organisms, while MPS exist as single, or multiple (less than complete organism constituency) organ systems is created in specific formats by its designers. The representation of an entire, very complex biology is the largest advantage of animal use over presently available MPS (7, 8). Conversely, despite the phenotypic resemblance of many animal models to humans, mechanistic differences very often exist at the cellular level. Additionally, animal models differ in endocrine, paracrine, and immunological aspects to humans and there are structural differences of intrauterine organs compared to humans. Non-human primate models are the closest to human pregnancy; however, their use in pregnancy research is limited due to the high cost of conducting experiments to generate statistically and biologically relevant data, scarcity in obtaining the right models, and the duration of pregnancy. This leads to translatability problems, which are their largest disadvantage. In response, breeders of some animals used in laboratories will genetically manipulate their stock to more closely "humanize" them. However, as we learn more about the influence of other factors such as the microbiome and silent genetic mutations, this still may not be enough to "idealize" these animal models. The use of systematic reviews may have some role in mitigating this lack of similarity, by optimizing selection of well-defined non-human experimental subjects. Regardless of these negatives attached to animal models, they are the only complete near-human representations that we have currently. For that reason, they have maintained our attention and attract us to their use.

Non-animal methods/new approach methods

A variety of MPS fill the categories of new approach methods (NAMs). MPS are engineered microenvironments that recapitulate the function of one or more organ systems. Among these physiological and pathophysiological miniaturizations, there are those which are subject to microfluidic flow and certain external forces and others in which those features are absent. A complete review and analysis of these types of systems (9) is beyond the scope of this article, and we will limit our discussion to the general groups of MPS and their most common applications. Our review

is to introduce the concept and utility of NAMs and we restrict this review to describe NAMs developed to study feto-maternal interface (FMi) biology.

Two-dimensional cell culture systems

The present majority of cell culture models are those having two dimensions (2D). They are the flat microscopic worlds familiar to most investigators. On the downside, they do not approximate the three-dimensional (3D) construct that is the human body, they soil and damage their own space, and their predictivity for drug development can be quite poor. Clearly, a methodological upgrade would be valuable and MPS developers have created and continue to create more human-relevant 3D systems.

Three-dimensional organ systems

Spheroids represent the entry level for 3D organ systems. They are cultured as free-floating, spontaneously aggregating (single cell or multicellular mixtures adhering to each other), minimally complex and spherical cellular designs (10). They derive from immortalized cell lines, primary cells, or human tissue fragments and have found use in drug testing, nanoparticle examination, and the study of neurodegenerative diseases and liver physiology/pathophysiology (11). Heterogenous in nature, spheroids contain layers of proliferating and non-proliferating cells; of important note is the often-necrotic core of larger spheroids, which results from diffusion difficulties of nutrients and waste across their structures. However, this is a problem that also provides an opportunity to the study of drug delivery to the similarly necrotic cores of solid tumors (12, 13).

Another tiny workhorse is the organoid, a microengineered, self-assembling cluster of organ-specific cells arising from adult stem cells, embryonic stem cells, or induced pluripotent stem cells (iPSCs) (such as those from skin or blood which have been reprogrammed into an embryonic-like pluripotent state) (14). Unlike the cellular monocultures from which spheroids are derived, the variety of stem cells producing organoids can selfrenew and differentiate into multiple lineages in vitro; this renewal process is responsible for an organoid's greater longevity. While patient-derived tumor organoids and patient-derived spheroids (tumorspheres) have a place in cancer research and precision medicine strategies, the more functionally advanced organoid can be sourced as a living tissue biobank, provide disease-specific models, and (like some spheroids) be useful in drug screening (15-17). The spheroid and organoids are now being developed and getting used for several FMi research (18-27). These approaches and tools will be optimized and available widespread for research use in reproductive biology and medicine.

Organs-on-chips and Tissue-chips are the most sophisticated of the MPS (28, 29). In addition to having 3D tissue architecture and multiple cell type composition, they are subject to biomechanical forces and linked by microfluidics, which permit accurate recapitulation of certain key aspects of human physiology (30). This superiority to 2D models and the less advanced 3D models (especially, viewed in consideration of the reliability and translatability problems associated with animal use), should encourage development of this MPS category as a pathway toward developing species of interest (human) methodologies. The challenges and limitations of such development have been reviewed elsewhere (31) and we will only herein comment that they are not yet engineered to full human physiological complexity. Like animal models, MPS can have reproducibility problems associated with cellular and other system variability. These difficulties might diminish with improved standardization of cellular and technical components. Additionally helpful are resources such as that of the aggregated human and animal exposure data contained within the publicly available University of Pittsburgh Microphysiology Systems Database (MPS Db),¹ which can aid in the selection of the best Organ-on-chip system for a particular study. Validation of Organs-on-chips, like other in vitro methods, is of current developer and regulatory interest, but a recent international roundtable of MPS stakeholders (privileged communication-Second BWF MPS Roundtable) noted that animal studies have not been subject to the same level of validation scrutiny as that proposed for MPS. Such thought prompts continued study of human MPS in various contexts, including its use in certain studies of reproductive physiology/pathophysiology.

Pregnancy (maternal-fetal interface) physiology NAMs

Pregnancy is a unique condition where two independent physiological systems, the fetus and the mother, co-exist for a defined period to maintain pregnancy and aid fetal growth and development (32). This co-existence ends with parturition, a unique synchronized process that terminates all homeostatic states of pregnant uterine tissues (33-37). Unfortunately, an unacceptable and growing number of pregnancies do not end at term with the delivery of a fully developed fetus due to various adverse pregnancy outcomes (APOs). Currently, mechanistic knowledge of complex physiologic interactions between fetal and maternal tissue that maintain pregnancy and potential drug actions to treat APOs is unavailable. Due to this knowledge gap, therapeutic interventions are limited during pregnancy, making women and their fetuses a highly vulnerable population. The impact of maternal exposures like drugs or vaccines (efficacy and toxicology) during pregnancy is difficult to measure as longitudinal sampling of intrauterine tissues and other biological specimens for analysis is impractical. Similarly, various environmental exposures (pollutants and toxicants) disrupting pregnancy homeostasis and contributing to APOs are also difficult to predict. This lack of knowledge creates a major clinical dilemma during pregnancy, and the only option is to deliver the fetus prematurely, contributing to almost 1 million neonatal deaths around the globe annually (38-44).

Pregnancy complications and outcomes like preterm births are difficult to predict due to heterogeneities in the risk factors and biomolecular pathways that are effectors of preterm labor. One of the major risk factors of pregnancy complications are environmental pollutants and toxicants (45-56). An extensive study on pregnant subjects that tested a broad array of 59 chemicals reported that 80% of the chemicals were commonly found in both maternal and fetal samples (57), suggesting a maternal-fetal transport and potential harm to pregnancies (58). Unfortunately, epidemiological association between exposure and APOs have not reduced PTB rates stemming from such exposures. To better understand the impact of hazardous substances on pregnancy risk, a molecular mechanistic knowledge by which toxicants activate pathways causing preterm birth in maternalfetal tissues are needed. Animal models have not been successful in determining pathological events contributing to adverse events during pregnancy, primarily due to the differences in the endocrine and paracrine systems between animal models and humans. Current in vitro toxicity testing methods cannot be used to accurately assess the hazard of tested substances on pregnancy outcomes as homeostatic disturbances resulting in immune imbalances at the feto-maternal interfaces cannot be measured using current approaches. Exposure to environmental toxicants often occurs in the form of mixed chemicals, especially during environmental disasters. However, current hazard assessment is conducted by evaluating single chemical at a time (59). It is well understood that simply combining the effect of individual chemicals cannot accurately determine the effect of mixed chemicals (60, 61), a major gap and challenge that is also recognized and prioritized (62, 63). It was found that, in some cases, when comparing toxic effects of a mixture of chemicals, the combined effects was greater than simply adding up the effects of the individual components (64, 65). These results strongly indicate that testing of actual environmental samples is desirable, rather than assuming that the effects of individual components from a mixed sample can be simply added together.

Therefore, reducing the risk of APOs is a global need (40, 66); however, pregnant women are excluded from most clinical trials and remain therapeutic orphans due to the lack of evidence for proper intervention (67–71). Challenges in conducting research and drug development, toxicology testing in pregnancy include (i) recruitment hurdles due to the paucity of data needed to convince the clinicians, subjects, and regulators of the utility of a drug in pregnancy, (ii) absence of informative biomarkers to assess the fetoplacental response to therapeutics or other exposures, and (iii) lack of suitable preclinical models (*in vitro* or animal models) to address both pharmacokinetics and pharmacodynamics. These limitations lead to the systematic exclusion of pregnant women from clinical trials (68, 72).

Maintenance of human pregnancy and initiation of parturition requires complex coordination of cells among the fetal (placenta and fetal membranes)-maternal (cervix, decidua, and myometrium) organs (F-M). Therefore, a critical challenge to a reproductive scientist is modeling the coordinated inter-tissue and intercellular interactions among various F-M uterine systems, especially at the FMi tissues and their multiple functions that maintain a harmonious state of pregnancy. Current studies using 2D cell cultures (73–75), mixed co-cultures (76–78), transwell co-cultures (79–81), tissue explants (75, 82), organoids (83–85), organ baths (86), and *in vivo*, animal models (87, 88) have provided valuable information on respective organ systems' contributions to pregnancy maintenance and some insights into

¹ https://upddi.pitt.edu/microphysiology-systems-database/

malfunctions associated with adverse pregnancy events (89). However, limitations associated with these models are far more than reliable information needed for regulatory agencies to act on an intervention or a biomarker to reduce the incidences of APOs. The primary limitations of current cell culture models (2D or 3D and transwells) are: (1) these models do not provide intercellular interactions required to understand the homeostasis mechanisms during pregnancy. Transwells are still limited to two cell type models, (2) primary cells currently used do not survive many passages to replicate data, (3) difficulty to control inter individual variability of cells when tissues are collected from different subjects for cell isolation, and (many cell lines are not derived from pregnancies (e.g., choriocarcinoma cell lines used as the placental cell) and do not reflect true physiology expected during pregnancy, and (4) explant cultures can generate plethora of data; however, still cellular manipulations (gene knockout studies, cellular interaction studies, cell type specific response studies, etc.) are difficult.

Improving women's reproductive health and fetal and neonatal outcome require physiologically relevant and experimentally manipulatable models that can generate high throughput data to study the human intrauterine organ system. Recent advances in our group's efforts to create a suite of five pregnancy and women's health-focused human MPS that model "healthy" and "disease" states of intrauterine tissues (90-97) present a unique opportunity to translate them into drug development tools. A biomimetic "organ-on a-chip (OOC)" recapitulates the multi-cellular organ system as seen in vivo using in vitro cell culture systems. These models can better mimic organ systems' structure, functions, and responses, though they do not mean maintaining or growing actual organs on a chip. The purpose of OOC is not to build a whole living organ but to synthesize minimal functional units that recapitulate tissue- and organ-level functions (98). The combination of microfabrication, microfluidics, and iPSC technologies has provided many physiological models that better mimic human anatomy, functions, and responses more accurately, as seen in vivo (9, 99). These MPS can provide compartmentalized chambers that enable culturing and organizing cellular, extracellular matrices (ECMs), and other microenvironmental layers within these compartments while still propagating cellular signals, and sometimes even cells themselves, to propagate between the compartments through interconnected fluid paths. Animal models do not always mimic human pregnancy, particularly parturition (100, 101) and preterm birth induced in many of these models is not a naturally occurring pathobiological process (102). To overcome this, microphysiologic systems (OOC technology platform) has been used to recreate the placenta (103-108), fetal membrane (90, 96, 109-113), cervix (94, 114), vagina (115, 116), and blood-brain barrier models (117-120).

Our MPS models of various intrauterine organ systems can address the limitations of existing models and ultimately accelerate preclinical drug development and provide reliable data for clinical trials. MPS models have been in use in other branches of medicine and are extensively recommended for preclinical trials (121, 122); however, reproductive medicine and the women's health field have not yet advanced in utilizing MPS. Such MPS models can also accelerate the repurposing of already approved drugs during pregnancy. These futuristic approaches using the *non-animal models* (also referred as NAMs) provide the next best opportunity for advancing research and filling knowledge gaps in the perinatal and reproductive health field. The basic design of each MPS model is that they have a (1) cell culture layer having cell culture chambers connected with arrays of microchannels and a (2) reservoir layer having large media reservoirs, where this layer is placed on top of the cell culture layer to support cell culture for 48–72 h without the need for media exchange. All the devices fit within a well of a 6-well plate (Figures 1–4; 92, 94–96, 112, 113) and can be operated with pipetting. The microfabrication of these MPS devices from polydimethylsiloxane (PDMS) has been extensively tested and validated (89, 90, 91, 112, 92, 94–96, 123–127).

Currently available MPS – NAM and their main directions in the field of maternal-fetal physiology

Modeling healthy and disease states using the feto-maternal interface on a chip

We extensively reported on the use of FMi-OOC (Figure 2) for modeling various maternal and fetal exposures and using a four-chamber device. During pregnancy, the fetal membrane (amniochorion) lines the intrauterine cavity and provides mechanical, immune, and endocrine support to the growing fetus (128). The fetal membranes also form one of the FMi by connecting to the maternal decidua. No MPS models have recreated this important FMi until now. The FMi-OOC chip has four concentric circles for cell cultures (96). Each chamber is 250 µm in height, and the width of each chamber was designed to mimic the thickness of each maternal and fetal layer [decidua, chorion, and amnion (mesenchyme and epithelium)] as seen in utero. This design allowed us to test four different cell types in four separate microenvironments (different culture mediums). Type IV collagen filled microchannels modeled the basement membranes and connected the chambers aligned with a media reservoir. This device contains maternal decidua (DEC) in the center chamber (green), connected to fetal chorion trophoblast cells (CTC) (yellow), amnion mesenchymal cells (AMC; pink), and amnion epithelial cells (AECs; purple) in the outer chamber. CTC and AMC layers are grown in a semi-3D suspension with Matrigel and primary amnion membrane-derived collagen to provide biological context (91, 92, 95, 96). This is the first MPS device to model the fetal membrane FMi allowing us to recreate disease phenotypes and assess the role of the *fetal inflammatory* response, a major determinant of adverse pregnancies and fetal morbidities. This device has been utilized to model ascending infection and inflammation [lipopolysaccharide (LPS) (96) and Ureaplasma parvum], identify the biological role of fetal-derived exosomes on labor induction (95), validate the functional role of drug transport proteins within fetal membrane cells (126), environmental toxicology studies (92), and conduct a preclinical trial with pravastatin and rosuvastatin (91). Results from these studies provide cellular and effluent biomarkers of various disease states and drug pharmacokinetics (PK), metabolism, and efficacy. Most importantly, the findings from FMi-OOC studies on infection-associated inflammatory changes leading to preterm



birth and exosome-mediated feto-maternal communication, associated inflammatory changes, and parturition were replicated physiologically using animal models validating the usefulness of MPS devices as an alternative to animal models. To note, physiological validation in animal models has its own limitations, as they do not mimic human pregnancy and or parturition.

Testing cigarette smoke and dioxin exposure induced cellular changes on feto-maternal cells

A two-chamber model of AEC and DEC separated by a semipermeable membrane was used to recreate FMi. Cells were treated with cigarette smoke extract (CSE) or dioxin for 48 h. The same experiments were conducted in transwells for comparison. Compared to transwell cultures, cells in OOC produced better membrane permeability regardless of the side of exposure (fetal vs. maternal). Membrane permeabilization was higher in AECs directly treated with CSE (1.6-fold) compared to similar treatment on the decidual side (1.2-fold). Both CSE and dioxin treatment on the maternal side induced cellular senescence on the fetal side. This effect was minimal in the transwell system. This confirmed the hypothesis that maternal oxidative stress (induced by cigarette smoke) can potentially cause premature aging of the fetal cells.

Testing exposures by the mother or fetus and its pathologic impact on fetal cells

Limitations of the two-chamber model, where we cannot include all the cellular components of the FMi, promoted us to develop the four-chamber device. FMi-OOC was used extensively by our lab to report the impact of cigarette smoke and Cadmium toxicity.

Cigarette smoke causes sterile inflammation, as seen in non-infectious inflammatory conditions in preterm birth

Using FMi-OOC, we recreated an environment induced by cigarette smoke extract (CSE – oxidative stress inducer). Maternal and fetal CSE exposures induced the following: (1) time-dependent propagation of cigarette smoke as measured by nicotine in the media in various cell chambers, (2) cellular transitions [epithelial mesenchymal transition (EMT)] similar to that seen in pathologic preterm birth, (3) senescence of cells induced by oxidative stress, and (4) sterile inflammation characterized by increased proinflammatory cytokines and reduced antiinflammatory cytokines. These changes were similar to our prior reports using the animal models (129), suggesting that MPS models are ideal for replacing animal models in experimental reproductive biology.



Cadmium toxicity and fetal outcome

Exposure to environmental chemicals can prematurely trigger labor-initiating signals at the FMi, leading to spontaneous preterm birth. Using Cadmium as a model toxicant, we tested the effect of maternal exposure to Cadmium (Cd) (92). Cd transport through the FMi and its impact on the cell cycle, cell death, and inflammation were analyzed. Cd treatment resulted in significant cell death and a pro-inflammatory environment in the maternal decidua but had minimal effect on the fetal chorion cells and no effect in the fetal amnion cells compared to controls. The mother primarily mediated the inflammatory response and not the fetus. The maternal response, but lack of fetal response, indicates that Cdmediated adverse effects originate from maternal pathophysiology rather than fetal-derived triggers of preterm labor. This study demonstrates that the FMi-OOC can indeed predict the response of FMi upon exposure to chemicals, opening the possibility for using OOC models for environmental toxin screens (92). Compared with cigarette smoke exposure described above, these data suggest that FMi-OOC can determine the differential effects, i.e., the pathological triggers (pathways and biomarkers) that contribute to APOs are maternal or fetal driven. This knowledge is critical in determining intervention strategies.

Extracellular vesicle (exosomes) mediated communication in response to toxicant exposure

Maternal and fetal exposure to various pollutants and toxicants can lead to premature senescence of the feto-maternal FMi tissues (130–133). These pathological changes prematurely produce exosomes that cargo cellular damage associated inflammatory mediators. These exosomes are communication channels



width and microchannel length. (D) Media reservoir layer that was aligned on top of the cell loading inlets and outlets. (E) Brightfield image showing STB and CTB cell barrier (white arrow) formation covering the microchannels. The cell barriers were confirmed to contain tight junction marker E-cadherin (green) expression confirming their functionality. Additionally, microchannels between the CTB and HUVEC chamber are filled with type I collagen to recreate the placenta stroma. Collagen was stained with Masson trichome for visualization (blue color, right image). (F) Brightfield and fluorescence images documents cell morphology, microvilli expression (black arrow), cytoskeletal markers [cytokeratin-7 (CK-7; red); tight junction marker (E-cadherin; green); and endothelial cell marker (MUC18; green)].

between the mother and the fetus and often cause functional changes, including labor (131, 133). We tested the feto-maternal communication using FMi-OOC where exosomes carrying High mobility group box 1 protein [HMGB1; one of the damages associated molecular pattern marker (DAMP)] related changes were tested. Exosomes with HMGB1 is hypothesized to increase the inflammatory load at the FMi (95). To test this, exosomes from AECs grown under normal conditions were engineered to contain HMGB1 by electroporation (eHMGB1). eHMGB1 propagation through FMi was tested using FMi-OOC. We have reported that eHMGB1 propagated through the fetal cells to the maternal decidua and increased inflammation [receptor expression (RAGE and TLR4) and cytokines]. Furthermore, intra-amniotic injection of eHMGB1 (containing 10 ng) into pregnant CD-1 mice on embryonic day 17 led to PTB. Injecting carboxyfluorescein succinimidyl ester (CFSE)-labeled eHMGB1, we determined in vivo kinetics. We reported that eHMGB1 trafficking resulting in preterm birth in animal models was also associated with increased FMi inflammation, such as that we observed in cells of the FMi-OOC. This study provided in vivo functional validation of FMi-OOC experiments and strengthened the reliability of such devices to test physiologic and pathologic systems (95). Bioengineered microvesicles (particles with higher size than exosomes) are also ideal for studying biochemical gradients and compartmentalized responses in in vitro models. Besides the FMi MPS models described above, other microphysiologic platforms exist, such as the placenta on a chip. This two-chamber device that contained two cells types (one from choriocarcinoma and other HUVEC cells also contained ECM) (105). Variations of these designs have been used by other investigators (104, 134–138). Some of these models are used for environmental toxicology trials.

The MPS based on NAMs have notable strengths. MPS models that can recreate an entire organ can overcome the limitations of many of the current approaches (2D cultures, explant models, transwell models, organoids, and animal models). MPS models require lesser number of cells and media compared to traditional approaches. Cellular migration, propagation of paracrine factors (e.g., extracellular vesicles) can be easily studies using MPS platforms. Creation of a healthy and disease states of an organ can mimic the state of the organ *in vivo* and can be used to determine pathophysiology at different cellular levels in an organ, discover biomarkers of the pathophysiology, and test efficacy of compounds that can reinstate the healthy state of the organ.

With all of this said, these MPS are not without limitations. Although the architecture of the organ can be reproduced in an *in vitro* model, they are difficult to manufacture, and miniaturized cell culture chambers are difficult to operate. PDMS used for many of the currently used MPS models have been reported to absorb certain compounds and this may pose a limitation to study specific secreted products or incorporating these substances in experimental models. Translational research requires high



culture layers. (1) Brightfield microscopy of microchannels connecting the VEC-ECTO chambers; (2) brightfield microscopy image of microchannels connecting the ENDO-DEC chambers; (3) brightfield image showing the microchannels between the cervical epithelial cell chambers and the MES chamber filled with Matrigel; and (4) representative SEM image showing the type IV collagen in the microchannels. (C) Representative fluorescence microscopy images showing intermediate filaments and cell-specific markers [pan-cytokeratin (pan-CK), cytokeratin (CK)-14, CK-18, vimentin, α -smooth muscle actin (α -SMA), mucin (Muc5a), and type I collagen] of the cells cultured in VCD-OOC. (D) The VCD-OOC device with the

integrated media reservoir filled with a dye in each cell culture layer for visualization.

throughput and large volume of data generation and most academic research lab settings may depend on commercial sources for their design and manufacture, prior to conducting any experiment; this can be cost prohibitive. Low number of cells and media used may also pose difficulties in terms of having sufficient materials to conduct several routine lab experiments (e.g., Western blot analysis). However, many labs have now designed ways to overcome many of these limitations and certain technological advancements have helped to reduce the sample volume required to conduct several lab tasks.

A future for NAMs

Reliability, reproducibility, and translatability are the features needed to make NAMs part of the research mainstream. As we have written before, this will require them to perform at a level that satisfies the needs of investigators (139). Reproducing the features and function of complex biological systems is an inescapably daunting task. If the past work of developers is any indicator for future advances in NAMs, then we should anticipate an equally awesome evolutionary process. Along that creative timeline, input

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from other stakeholders is crucial. While regulators will remain as the gatekeepers for releasing end products, conversations among the stakeholder groups most closely associated with the challenges confronting NAMs will have relevance for informing and driving the developmental process. The initial core demographics of academic scientists, industry, government funders, government regulators, venture capital, philanthropy, animal research advocacy, and animal protection, should be joined by patient advocacy groups, specialty clinicians, ethicists, and other stakeholders holding significant interests in the promise that NAMs have for delivering improved healthcare outcomes, increased scientific accuracy, reduced and/or eliminated reliance on animal use, and optimized environmental welfare. NAMs are ideal but still there are limitations to accessing them and conducting experiments using them. For example, microfluidic and other platforms are not accessible to several labs and trained personnel to conduct experiments are also currently lacking. Human primary cells, cell lines or human iPSCs are also needed for use in these experimental models. Accessibility or cost associated in accessing them may also limit several researchers from acquiring these technologies or using them.

Author contributions

RM: Conceptualization, Data curation, Funding acquisition, Writing – original draft, Writing – review and editing. LL: Conceptualization, Supervision, Writing – original draft, Writing – review and editing. LM: Conceptualization, Writing – original draft, Writing – review and editing.

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