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### **Bioreactor Engineering of Stem Cell Environments**

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### Abstract

Stem cells hold promise to revolutionize modern medicine by development of new therapies, disease models and drug screening systems. Standard cell culture systems have limited biological relevance because they do not recapitulate the complex 3-dimensional interactions and biophysical cues that characterize the *in vivo* environment. In this review, we discuss the current advances in engineering stem cell environments using novel biomaterials and bioreactor technologies. We also reflect on the challenges the field is currently facing with regard to translation of stem cell based therapies into the clinic.

### Keywords

stem cells; bioengineering; bioreactors; biomaterials; in vitro models; models of disease; regenerative medicine; translation; high-throughput screening; personalized medicine

### 1. Introduction

Stem cells provide enormous opportunities for improving human medicine, through the development of tissue replacement therapies, human *in vitro* models of disease, screening of therapeutic and toxic effects of chemical libraries, and "personalized" medicine. Furthermore, recent advances in stem cell biology, biomaterials, genetic engineering and biomedical engineering have allowed unprecedented ability to create controlled environments and ask specific biological questions. The progression from historical culture plates with animal cells and immortalized cell lines towards embryonic stem cells (ES) and induced pluripotent stem cells (iPS) in 3-dimensional (3D) bioreactors is truly paving the way for new applications in tissue engineering and regenerative medicine, the study of disease, and drug screening (Figure 1). Here we review advances in engineering stem cell environments using dynamic bioreactor systems, and discuss the importance of these novel tools to stem cell research as well as the applications of stem cells in pre-clinical and clinical settings.

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### 2. Limitations of current stem cell research models

Ever since the time of Galen, the famous physician who reportedly dissected pigs and goats, researchers have sought experimental models of human biology. More recently, the Petri dish, invented at the end of the 19<sup>th</sup> century, has proven invaluable for experiments in cellular biology. And in fact, standard Petri dish cultures are still widely used: adherent cells are grown on synthetic surfaces (i.e. tissue culture plastic), basement membrane or extracellular matrix protein coatings (i.e. laminin, vitronectin, collagen), or feeder cells (i.e. mouse embryonic fibroblasts), and are bathed in culture medium containing appropriate nutrients and signaling molecules. Changing of cell culture medium is conducted batchwise, resulting in the variation of medium composition over time.

In Petri dishes, the cells are essentially cultured in two dimensions. Stem cells generally grow in dense colonies with defined borders, which expand in size and merge with other colonies in the culture dish (Takahashi et al. 2007; Thomson et al. 1998). At confluence, cells are passaged for further expansion, or subjected to differentiation protocols. While this culture format recapitulates some aspects of tissues that are essentially two-dimensional (2D), such as skin or bladder, it falls short of providing environments experienced by most cells in the organism. In particular, Petri dish culture lacks the 3D cell-cell and cell-matrix interactions, provision of spatial and temporal gradients of biochemical and physical signals, and systemic regulation including cross-talk between different organ systems (Kaplan et al. 2005; Vunjak-Novakovic et al. 2005). Findings obtained in Petri dish cultures are therefore not always predictable of whole tissues and organs, and are difficult to translate into the *in vivo* settings of pre-clinical studies in animals, and clinical trials in human subjects.

In contrast to the controlled environments of cell culture systems, animal models allow assessment of stem cell developmental potential within whole organisms, and are therefore invaluable for studies of development, disease pathogenesis and toxicity testing (Cheshier et al. 1999; Sacco et al. 2010; Wobus and Loser 2011). After the discovery of mouse ES cells and the completion of human genome sequencing, creation of mice with specific gene knockouts and gene reporters has enabled the study of gene function during development, and cell lineage tracking experiments (Lloyd 2011). Furthermore, specific rodent strains with compromised immune systems have been developed that allow us to study the function of human cells *in vivo* without immune rejection (i.e. humanized mice) (Shultz et al. 2011).

However, despite these advantages, animal models present several limitations when used in disease modeling and toxicological studies. First, very few animal models faithfully reproduce human pathophysiology. Therefore it is important that all disease models - whether surgically or pharmacologically induced or genetic, are clearly defined with regards to the pathology that is being modeled, and to how it relates to the human condition. Second, there are important interspecies differences in pharmaco-toxicological effects between experimental animals and humans (Wobus and Loser 2011), which are only exacerbated when human cells are transplanted into immune-suppressed hosts, potentially also affecting physiological healing responses (Goldring et al. 2011). In this respect, progress in preparation of iPSc from large animals, such as pigs, would advance transplantation studies (Montserrat et al. 2011). Finally, for studies of transplanted cells, *in vivo* models offer less control over the cell microenvironment, and are challenging for on-line monitoring of the outcomes, compared to the *in vitro* systems, which are better defined and better controlled.

A critical application highlighting the importance of developing better *in vitro* systems to model human biology and physiology is that of drug development. A number of high-profile drugs have been recently withdrawn from the market, most commonly due to the cardiotoxicity, neurotoxicity and hepatotoxicity that were not observed until clinical trials

(Report 2011). These negative side effects were not detected because of the limited functional capacity and genetic diversity of current research models, resulting in drugs that pass animal studies but fail in human studies. Part of the problem is that, in the conventional target-centric drug discovery model, compounds are typically not tested in a patient population until Phase II clinical trials have been carried out, thus contributing to a high drug attrition rate (Kola and Landis 2004, Kola 2008).

Further complicating the issue is that even compounds that have been found to be safe and efficacious in animal models can fail spectacularly in clinical trials, adding expense and risk. The cost of bringing a new drug to the clinic has been estimated at \$1.2 billion, up from a 2003 average of \$802 million (DiMasi et al. 2003), and pharmaceutical companies face their largest losses when drugs fail at the late stage of trials, such as Phase III and the post-marketing stage (Report 2011). In addition, some potential targets may be excluded from further development due to false negative results during *in vitro* testing.

The shift from using animal cells and immortalized cell lines to human stem cells and their differentiated progeny, and from simple Petri dish culture to more sophisticated environments capturing critical aspects of the *in vivo* system, could be hugely beneficial for drug development. Bioreactor-based screening technologies could improve prediction of *in vivo* outcomes in pre-clinical and clinical studies. Tissue models using cells from multiple patients could allow study of large and diverse genetic and disease pools, help overcome many of the current limitations of clinical trials (Li 2004). and help advance the state of the art from that of treatments designed for an "average" patient to a personalized medicine approach (Pouton and Haynes 2005, 2007; Rolletschek and Wobus 2009).

### 3. Stem cell cultivation in scaffold-bioreactor systems

Our understanding of stem cell biology has been greatly expanded by the development of novel culture environments using biomaterials and bioreactors that recapitulate selected aspects of native tissue (Gerecht et al. 2007; Gilbert et al. 2010; McBeath et al. 2004; Steiner et al. 2010). The importance of inductive signals present in 3D tissue settings has been recognized long ago, and simple techniques such as cultivation of mesenchymal stem cell (MSC) pellets, human embryonic stem cell (hESC) aggregates (embryoid bodies, or EBs) or neurospheres have been used to differentiate cells (Itskovitz-Eldor et al. 2000; Johnstone et al. 1998; Reynolds and Weiss 1992). However, these cultures present limitations in terms of size (mostly <0.5 mm), control over cell populations and biophysical signals (mechanical, electrical, hydrodynamic), and the lack of extracellular matrix.

In the 1990s, it was proposed that 3D human tissue substitutes could be engineered *in vitro* by cultivation of cells on scaffolding materials, which act as temporary biodegradable templates for tissue development, in bioreactor systems providing environmental control and biochemical and biophysical signaling (Langer and Vacanti 1993) (see Figure 2). This approach was dubbed the "biomimetic paradigm" because it was based upon the idea that these scaffolds in combination with bioreactors would recapitulate the relevant developmental and morphological *in vivo* events *in vitro*. During native tissue development, cells are exposed to a myriad of signals including multiple cell types, extracellular matrix, cytokines, and physical factors, all of which are dynamic in nature. And in fact, since this time, by following the biomimetic paradigm, many biomaterial scaffolds and *in vitro* systems have been designed to recapitulate key aspects of this dynamic environment and support cell growth, matrix synthesis and tissue maturation.

Developments in the field of biomaterials have made it possible to create 3D scaffolds with defined structural features (shape, porosity, pore size, surface roughness), mechanical properties, composition and degradation rate (Kraehenbuehl et al. 2011; Lutolf and Hubbel

2005). A variety of native, synthetic and composite materials is now available in geometries including fibers, meshes, sheets, sponges, hydrogels and combinations (Bhumiratana et al. 2011; Gerecht et al. 2007; Hofmann et al. 2007; Tran et al. 2010). Biodegradable materials allow encapsulation and sustained release of inductive biochemical signals (Ferreira et al. 2007; Son et al. 2011; Wang et al. 2009), and tethering of specific attachment groups (Hanjaya-Putra et al. 2011), to further tailor the cellular microenvironment and direct the cell function.

In order to induce cell growth in the third dimension and to support tissue development, it is critical to provide mass transport to and from all cells using dynamic culture systems such as bioreactors. In static cultures, mass transport is based on diffusion, and generally limits tissue development to thicknesses less than 0.2 mm due to drops in oxygen tension and increased concentrations of toxic metabolites (e.g., acidification). In bioreactors, stirring, perfusion, and dynamic loading have been applied to provide convective transport and allow tissue development on a millimeter to centimeter scale (Chahine et al. 2009; Grayson et al. 2010; Grayson et al. 2011; Hofmann et al. 2007; Marolt et al. 2006).

The direction and rate of medium flow and the viscosity of medium can be tailored to achieve mechanical stimulation by shear stress (Grayson et al. 2010; Grayson et al. 2011; Sikavitsas et al. 2003). Other examples of biophysical stimuli provided by bioreactors include compressive and tensile loading, electrical stimulation, medium flow resulting from stirring or perfusion (Baker et al. 2011; Bian et al. 2012; Maidhof et al. 2011), alone or in combinations. Finally, bioreactors allow online control and monitoring of temperature, pH, oxygen, medium concentrations of nutrients in order to meet the requirements of tissue under study.

### 3.1. Case study 1: Large-scale bioreactor cultivation of pluripotent stem cells

To realize the promise of pluripotent cells for drug discovery and cell therapy, scale up and automation of the manual culture process are necessary, as they assure robust and reproducible production of large numbers of relevant cells. In recent years, suspensionculture bioreactors have been investigated for the production of human pluripotent stem cells and their differentiated derivatives in clinically-relevant amounts, which range from several tens of millions to a few billion cells (Kehoe et al. 2010). These bioreactors offer advantages over monolayer culture, including control of the microenvironment, higher cell densities per volume, low contamination risk, application of scaffolds for encapsulation of anchorage-dependent cells, and tailoring of the hydrodynamic environment to study the effects on self-renewal and differentiation of stem cells.

Agitation is essential for ensuring that all cells are exposed to the culture medium of the same composition, keeping the aggregates or microcarriers suspended, and controlling the aggregate size (Kehoe et al. 2010). In static culture, cell aggregates can form large agglomerates with substantial necrotic areas in their centers, resulting from limited nutrient and oxygen transport (Gerecht-Nir et al. 2004). In stirred suspension bioreactors, impeller-associated hydrodynamic shear can regulate the size of ESC aggregates, prevent agglomeration and increase the final cell numbers. However, high levels of shear stress (>15 dynes/cm<sup>2</sup>) may cause damage to the cells. For hESCs sensitive to shear stress, slow turning lateral vessels and high-aspect rotating vessel bioreactors have been tested, as they contain a low shear environment (Gerecht-Nir et al. 2004). Hydrodynamic shear forces can also be applied in a controlled fashion to guide cell differentiation into specific cell lineages (Sargent et al. 2010).

In initial studies, growth and differentiation of mouse embryonic stem cells (mESCs) were evaluated in suspension bioreactor cultures of cell aggregates, microcarriers or hydrogels

(Kehoe et al. 2010). The microcarrier and monolayer cultures showed similar cell doubling times (14–17 hours), and similar expression of pluripotency markers after 15 days of culture (Fok and Zandstra 2005). mESC aggregates were directly induced into EBs capable of multilineage differentiation by changing the medium composition. Stir-related shear stress reduced colony agglomeration, and regulated the size of aggregates. Uniform growth of mESC aggregates was achieved when the agitation rate was increased from 60 to 100 rpm.

Studies of mESCs in orbital suspension cultures showed that the hydrodynamic environment effectively regulated not only the size and yield of EBs from mESCs, but also the organization and phenotypes of differentiating cells (Sargent et al. 2010). To prevent agglomeration, mESCs have been cultured in hydrogel microcapsules in stirred suspension bioreactors, and differentiated into hematopoietic progenitors (Dang et al. 2004). More recently, gelatine microcarriers were used for mESC differentiation into the chondrogenic and osteogenic lineages. Differentiated cells formed mineralized tissue *in vitro* and *in vivo* in both ectopic and orthotopic locations. Interestingly, cultivation of mESC aggregates in the absence of microcarriers resulted in the formation of muscle tissue after ectopic transplantation (Alfred et al. 2011).

In one of the first studies of hESCs, differentiating EBs in rotating bioreactors that are characterized by laminar flow pattern with mild mixing, reached three times higher cell densities  $(36 \times 10^6 \text{ cells/ml})$  than the static dish cultures  $(13 \times 10^6 \text{ cells/ml})$  after 4 weeks of culture (Gerecht-Nir et al. 2004). The cells remained viable and gave rise to lineages of all three germ layers, suggesting that the bioreactor microenvironment did not alter early developmental events, and could be used to scale-up production of differentiated cells. Subsequent studies included the scale-up of mesoderm and cardiac cell differentiation from hESCs, using microprinting to prepare cell aggregates of uniform size that were then cultured in suspension bioreactors (Niebruegge et al. 2009).

Overall, the culture in suspension systems with differing agitation forces and vessel geometries (rotating dish, roller bottle, spinner bioreactors) significantly improved cell growth compared to static culture. In addition, cultivation of uniform size hESC aggregates under hypoxic conditions in spinner bioreactors further improved the yield of differentiating cells (reaching densities of  $6 \times 10^5$  cells/ml after 16 days of culture) (Niebruegge et al. 2009).

In an alternate approach, encapsulation of mESCs and hESCs in poly-L-lysine coated alginate capsules, and liquefaction of the capsule cores was used to prepare cell aggregates of uniform sizes (Jing et al. 2010). Subsequent cultivation in suspension bioreactors allowed directed cell differentiation into cardiac lineage with higher efficiency than in static culture. In contrast, cultivation of hESCs on microcarriers in suspension bioreactors allowed monolayer-like cell growth, important for directed differentiation rate (to 45 rpm) resulted in up to 45-fold increase in the number of hESCs (~ $1.8 \times 10^6$  cells/ml) expressing pluripotent markers after 8 days of culture. Subsequent change from expansion to differentiation medium resulted in a population expressing definitive endoderm markers. Efficiency of differentiated cells production per unit of surface area was similar to that of static cultures, but the use of microcarriers bioreactors allowed more surface per unit volume.

These studies illustrate diverse effects of scaffolds and bioreactors on mESC and hESC fate, and the potential to develop bioprocesses where cell expansion, differentiation and storage could be integrated within a single culture system (Nie et al. 2009). It has been more challenging to scale up continuous expansion of undifferentiated hESCs compared to

mESCs, due to their propensity to differentiate in the 3D environment, and low survival after dissociation into single cells. Biomaterials supporting undifferentiated hESC growth, such as hyaluronic acid hydrogel (Gerecht et al. 2007) and molecules enhancing cell survival, such as Rho-associated kinase (ROCK) inhibitor (Watanabe et al. 2007), could be implemented in such culture protocols.

In one example, Krawetz and colleagues reported continuous expansion of hESCs in stirred suspension bioreactors using ROCK inhibitor and rapamycin to promote cell survival and undifferentiated growth (Krawetz et al. 2010). Components tested to promote suspension cultures of undifferentiated hESCs included neural sphere culture medium, serum replacements, beta-D xylopiranose, extracellular matrix components (fibronectin, laminin), activin A, and basic fibroblast growth factor (Steiner et al. 2010).

### 3.2. Case study 2: Engineering of functional bone tissue from human stem cells

The shortage of donor tissues and organs suitable for transplantation, and the inherent limitations of artificial prosthetic devices make the development of stem-cell based biological tissue substitutes one of the most active areas of regenerative medicine research. One example is the development of functional bone tissue substitutes for surgical treatment of large bone injuries (Fröhlich et al. 2008).

Many studies have focused on cell lines and animal cells as models for bone tissue regeneration, and the signaling factors and molecular pathways involved in this process. However the development of therapies requires that the findings are tested with human cells. The treatment of smaller defects, such as augmentation of the jaw bones prior to dental implant placement, has been enhanced using culture-expanded primary cells and biomaterials (Schimming and Schmelzeisen 2004). Regeneration of extensive defects (on the centimeter-scale) necessitates large quantities of functional reparative cells (~  $10^8$  per cm<sup>3</sup> of bone), within a template with capacity to guide and undergo regeneration. Stem cells have the ability to self-renew, thereby allowing expansion to large numbers, and to give rise to specialized lineage progenitors, and therefore represent an excellent cell source for bone tissue engineering.

Adult human Mesenchymal Stem Cells (hMSCs), hESCs and iPSCs have all been shown to give rise to bone-forming cells. Research has mostly been performed using hMSCs from bone marrow, as these cells give rise to bone and cartilage during development, and have been shown to promote bone healing clinically in autologous transplants (Salter et al. 2012).

Bioreactor cultivation of human stem cells in an osteogenic scaffold supported cell survival, differentiation, maturation and deposition of bone matrix, while restricting the development of unwanted lineages (i.e. adipose), and facilitating a continued remodeling and vascularization following transplantation (Marolt et al. 2012). The design of scaffold-bioreactor systems is guided by the size and shape of the defect to be repaired and the need for establishing certain mechanical properties and ability for integration. A stepwise approach is often used to select and optimize culture parameters (Grayson et al. 2010; Grayson et al. 2008, 2011; Marolt et al. 2006). For instance, soluble factors to achieve osteogenesis of bone marrow stromal cells (BMSCs) have been extensively studied, and frequently involve a serum-based medium supplemented with a cocktail of osteogenic factors (Hofmann et al. 2007; Marolt et al. 2006; Sikavitsas et al. 2003). A variety of native and synthetic scaffolds has been found to influence bone formation via size, pore architecture (Hofmann et al. 2007), surface roughness (Dalby et al. 2007), cell attachment sites (Shin et al. 2004) and the matierial's biomechanics (Engler et al. 2006).

Among the bioreactors for bone tissue engineering, those providing medium perfusion through the forming tissue have been demonstrated to best allow the scale-up of bone culture to clinical sizes (several centimeters) (Grayson et al. 2010; Martin and Vermette 2005, Salter et al. 2012). The interstitial flow through the scaffold is critical for cell survival and metabolic activity (Grayson et al. 2010; Grayson et al. 2008, 2011). Perfusion also causes shear stress on the cultured cells, stimulating bone development (Grayson et al. 2011; Sikavitsas et al. 2003).

Cultivation of bone substitutes under perfusion conditions requires a mechanical integrity and porosity of the scaffolds to allow fluid flow, with tight control over the perfusion rate and regime (continuous, intermittent etc), oxygen tension, and additional biophysical signals (e.g., mechanical loading). Our group has achieved reproducible formation of compact bone grafts (4 mm diameter  $\times$  4 mm thick discs) from bone marrow derived hMSCs cultured on decellularized bone scaffolds (Grayson et al. 2008; Grayson et al. 2011). Increasing the medium flow velocity from 80 to 1800 µm/s, corresponding to the initial shear stresses of 0.6÷20 mPa, significantly affected bone formation, with the flow velocity between 400÷800 µm/s yielding the best overall outcomes (cell numbers, cell connectivity and bone matrix formation).

This simple and robust scaffold-bioreactor model was easily translated to adult human adipose-derived stem cells (Fröhlich et al. 2010) and hESC-derived mesenchymal progenitors (Marolt et al. 2012), as well as to the cultivation of bone on synthetic silk-hydroxyapatite scaffolds (Bhumiratana et al. 2011). Composite scaffolds containing 4.6% hydroxyapatite resulted in the formation of bone constructs with the Young's equilibrium modulus of 1.5 MPa. A similar perfusion culture approach was used to engineer a 1.5 cm condyle-shaped bone substitute, using anatomically shaped scaffolds and bioreactors, demonstrating the feasibility to prepare clinically relevant bone substitutes (Grayson et al. 2010).

The ability of cultured bone substitutes to survive *in vivo* is limited by the rate of vascularization, and many investigations are currently focusing on vascularized bone substitutes. Strategies include co-cultivation of various adult stem cells and adult vascular (progenitor) cells, as well as induction of pluripotent stem cells into bone-forming and vascular cells. The challenges include designing scaffold-bioreactor environments that would support development of specific tissues in specific scaffold regions (Correia et al. 2011). For engineering of vascularized bone from iPSCs (which can be used as a single source to derive cells of both bone and vascular lineages), the challenge is in efficient differentiation into the specific respective lineages, and in the restriction of unwanted developmental potential of the cells.

Differentiation of iPSCs generally results in mixed cell populations, potentially harboring residual pluripotent cells that could lead to development of tumors or unwanted tissues. Our strategy has been to first derive and expand specific lineage progenitors (for 5 passages), and then to perform the bone substitute engineering (5-week bioreactor cultivation). We have recently shown that ESCs can be induced into mesenchymal-like progenitors with osteogenic potential similar to that of bone marrow derived hMSCs. Importantly, cultivation in decellularized bone scaffolds in perfusion bioreactors resulted in development of homogenous bone tissue (cylindrical plugs, 4 mm in diameter and thickness) without the presence of other lineages, that remained stable and matured *in vivo*, and did not result in uncontrolled cell growth during 8 weeks following transplantation (Marolt et al. 2012).

# 4. Miniature bioreactors for precise, systematic studies of stem cell environments

Many aspects of the native tissue and stem cell niche develop under tight spatial and temporal control, with synergistic effects between various factors. These types of systems typically require tedious combinatorial experimentation, providing strong rationale for tight control over parameter variation and better mimicry of the complex *in vivo* tissue milieu in a high-throughput and scalable manner. Miniaturized versions of 3D bioreactor systems, known as micro-bioreactors, represent an important step towards accurate, multifactorial control of cultured cells and tissues. Because they reduce the amounts of cells and materials required for experimentation, they increase the number of experimental conditions and replicates that may be studied in parallel, making them ideally suited for high-throughput, combinatorial studies (Beebe et al. 2002; Squires and Quake 2005; Toner and Irimia 2005).

Furthermore, the short transport distances in micro-bioreactors allow for even more precise, *"in vivo*-like" control of certain environmental cues. For example, flow in micro-bioreactors is laminar: inertial forces are dominated by viscous forces, and transport occurs by molecular diffusion or by convective regime of well-defined hydrodynamic profile. Laminar flow allows for precise calculation of mass transport parameters and the establishment of flow profiles as a function of the system geometry, flow rate and fluid properties. Since the laminar streamlines remain constant over time and mixing occurs by diffusion, precise dynamic perturbations localized in space and time may be applied (Aasen et al. 2008; Chung et al. 2009; Cimetta et al. 2010; Cimetta et al. 2009; Diao et al. 2006; Jeon et al. 2000; Toetsch et al. 2009). Small transport distances are also key for enabling fast responses to environmental stimuli and their manipulation and control (Cimetta et al. 2009). In summary, micro-bioreactor systems are ideally suited for stem cell research due to their unique characteristics:

- *Down-scalable/parallelizable.* Miniaturization leads to significant reductions in the required amounts of cells and materials for each experimental condition, thus allowing cutting costs and facilitating the increase in test numbers necessary for high-throughput, multi-factorial, quantitative studies. When experiments are massively parallelized, small but significant differences may be detected in biological responses that might otherwise go unnoticed in standard cell culture. In addition, parallelization also enables the screening of large numbers of variables, thus further reducing variability between separate experimental runs. Finally, significant multi-factorial characteristics may be attained by experimental designs in which cell density, cell types, substrate mechanics and biophysical stimuli may be independently and/or simultaneously controlled.
- <u>*"In-vivo-like" fidelity.*</u> Tight spatial and temporal control of the delivery of regulatory factors result in a closer resemblance to the *in vivo* environments. The precise control over transport phenomena and delivery of exogenous stimuli allows the creation of compartmentalized concentration patterns (such as gradients) and of the physiological patterns of electrical and mechanical stimulation. In addition, it is possible to incorporate multiple tissue types on a single platform, either as 2D or pseudo-3D culture.

Key to micro-bioreactor fabrication are micro fabrication techniques, which have been used for applications ranging from micropatterning and microelectronics, to fabrication of micro electro-mechanical systems (MEMS) (McCreedy 2000; McDonald et al. 2000; Richards-Grayson et al. 2004). Typically, polymer-based micromachining is used for obtaining molds with high aspect-ratio structures, while photoresist-based photolithography allows fabrication of microscaled features with ~1 µm precision. Polydimethylsiloxane (PDMS)

micromolding enables the production of polymeric devices from these molds with high accuracy and yield (Kane et al. 1999; Whitesides et al. 2001; Xia and Whitesides 1998). Many types of micro-bioreactor systems have employed these techniques to facilitate drug discovery (Dittrich and Manz 2006; Wen and Yang 2008), stem cell and medical research (Beebe et al. 2002).

### 4.1. Case study 3: Micro-bioreactors for high-throughput screening of environmental factors

One set of challenges in miniaturizing stem cell culture systems is to adapt the complex microenvironment that regulates stem cell fates to down-scaled *in vitro* models. Niche factors are multiple and diverse: for example, biochemical signaling cues and soluble factors from neighboring cells or elsewhere, must be recapitulated along with their associated mass transport characteristics. Moreover, the physiological or pathological cell milieu includes factors such as extracellular matrix elasticity and geometry, mechanical and electrical signals. For the niche effects to be investigated in a combinatorial fashion with physiological relevance, the key relevant factors must all be recapitulated on a microscale with high accuracy.

Certain niche factors, such as microflows are ideally suited for "in-vivo-like" control of transport and convective flow profiles. Figallo et al, for example, have developed a simple device comprised of an array of culture wells to enable systematic and precise variation of mass transport and hydrodynamic shear (Figallo et al. 2007). Using this microfluidic platform, hESCs were systematically studied for their cardiovascular differentiation potential, and it was demonstrated that higher percentages of cells expressing vascular-specific markers was inversely proportional to cell density (increasing from 2% to 23% as cell density decreased from  $1 \times 10^4$  to  $2 \times 10^2$  cells/cm2). This technology is compliant with standard imaging formats, and quantitative image processing.

Cimetta et al have developed a microfluidic device generating stable concentration gradients for long-term cell culture, and applied this method for studying Wnt3a regulation of - catenin signaling (Cimetta et al. 2010). More recently, they designed a new microfluidic bioreactor providing space-resolved gradients of multiple molecular factors in 3D cell culture settings apt for studies of complex signaling pathways in human pluripotent stem cells. A single microbioreactor yields up to 120 data points, corresponding to 15 replicates of a gradient with 8 concentration levels (Cimetta et al. 2012).

Other groups have focused on other factors in the stem cell niche, such as biochemical and extracellular matrix components, in a parallelized, high-throughput manner. Gobaa et al, for example, have developed a micro-platform that simultaneously probes the role of biochemical and biophysical niche factors on stem cell fate (Figure 3). The device is comprised as an array of soft hydrogel microwells with tunable stiffness (shear modulus of 1–50 kPa) which can be independently functionalized with combinations of proteins spotted by robotic technology (Gobaa et al. 2011). The authors addressed three representative stem cell questions to demonstrate the utility of their device: (*i*) the influence of cell density on adipogenic differentiation of hMSCs, (*ii*) the effects of substrate stiffness on osteogenic differentiation and (*iii*) the self-renewal of non-adherent mouse neural stem cells (NSCs). With this high-throughput hydrogel microwells system, features such as cell density, substrate mechanics and protein incorporation could be tested, and more than 2,000 experiments with several adherent and non-adherent cell types can be performed on a single glass slide. Using their method, they showed it is possible to probe the effects of key micro-environmental perturbations on the fate of single cells in high throughput.

One of the challenges in high-throughput screening of soluble factors is to create devices that take advantage of microfluidic phenomena and exhibit the combinatorial diversities achieved by microarrays. Robotic fluid handling and microspotting techniques, for example, facilitate high throughput studies of the influence of different immobilized factors in microarrys, but require time-dependent stimulation to be applied simultaneously. Microfluidic platforms that facilitate studies of diffusible growth factors with spatial and temporal control have historically been difficult to scale up into larger, individually addressable culture chambers. Gomez-Sjoberg et al address these issues with their versatile, fully automated microfluidic cell culture system that creates arbitrary media formulations in independent cell culture chambers (Gomez-Sjoberg et al. 2007). Their microfluidic device is comprised of 96 rectangular culture chambers customizable in terms of cell seeding density, medium composition, and feeding schedule, and each chamber is imaged with time-lapsed microscopy, in long-term (ie > 7 days) experiments.

Certain biophysical cues, such as microflows, are facile to incorporate into microbioreactors, in contrast to other biophysical cues (e.g. mechanical stretch, electrical stimulation etc) that may require additional components and innovative solutions. Kensah and colleagues have constructed a novel scaled-down bioreactor for culturing cardiomyocytes that incorporates the application of mechanical stretch (Kensah et al. 2011). When exposed to cyclic longitudinal stretch during culture via a linear motor, cardiac tissues exhibited hypertrophy, and increased systolic force (~1.42 mN vs. ~0.96 mN in controls). Tandon and colleagues have microscale cell culture system with an interdigitated microarray of excimer-laser-ablated indium tin oxide electrodes for electrical stimulation of cultured cells (Tandon et al. 2010). They constructed systems in a range of geometries for cultivations of cardiomyocytes and human stem cells. Over 6 days of culture, electrical stimulation (2 ms duration, 1 Hz, 180 µm wide electrodes, 200 µm spacing), enhanced proliferation, elongation and alignment, and numbers of gap junctions in both cell types.

Another set of challenges with high throughput screening systems is to automate the cell culture and associated assessments. Lecault et al, for example, have developed a microfluidic platform containing thousands of nanoliter-scale chambers with the capability to perform live-cell imaging, *in situ* immunostaining, recovery of viable cells, and automated medium exchange (Lecault et al. 2011). Ye et al have developed a microfluidic platform which is able to rapidly extract multi-parametric measurements of plasma membrane permeability, nuclear size, mitochondrial transmembrane potential and intracellular redox states in anti-cancer drug-induced apoptosis of human liver carcinoma cells (Ye et al. 2007).

Parallelizable, high-throughput platforms of this kind will likely help characterize stem cells in terms of their self-renewal and differentiation, proliferative capacity and migration in response to combinations of stimuli. Going forward, better control over stem cell niche conditions such as cell density, biophysical cues, while allowing for increased multiplexing abilities, individual addressability, and online assay methods will undoubtedly contribute to higher quality, quantitative data relevant to toxicology, developmental biology, and beyond.

### 4.2. Case study 4: "Body on a chip" devices for drug efficacy/toxicity studies

One of the limitations of single cell cultures (whether of conventional type or bioreactorbased) is that, at best, they only capture the behavior of isolated, single tissues. A particularly important need for *in vitro* modeling of multi-organ systems is in the prediction of toxicity: *in vitro* approaches have hitherto only met with limited success, mainly because of the complexity of the *in vivo* toxic responses. In the animal or human body, there are multiple organs that are interconnected by the systemic circulation, leading to multi-organ interactions and multi-organ metabolism. In the body, cross-talk occurs through soluble

agents: metabolites generated in the liver, for example, can influence the gene expression in other cell types and vice versa (Bhatia et al. 1999; Khetani et al. 2004; Takayama G et al. 2007). Drug screening based on simple culture systems is able to predict many cases of primary drug toxicity, but remains limited in predicting indirect toxicity (eg cardiotoxic effects of breakdown products of a drug metabolized in the liver) and of the pharmacokinetic bioavailability of drugs. Affordable *in vitro* models that mimic the *in vivo* multi-organ human system (so-called "Body-on-a-Chip" systems) are therefore of prime importance for drug discovery and chemical screening. Microscale systems are particularly well suited to the task because of their ability to recapitulate the cellular microenvironment.

Research towards developing "Body-on-a-Chip" systems has been spurred by the success of the development of microscale "Organs-on-Chips" that recapitulate the structural tissue arrangements and functional complexity for each component organ, and efforts in bioprinting, which may be employed for direct 3D cell writing with precise control (Chang et al. 2008). A variety of microscale organomimetic cell culture systems have been developed, such as liver, blood vessels, lung, kidney or brain on a chip, (Figure 3, reviewed in Huh et al. 2012). Notable examples include microscale cultures of cardiomyocytes from human stem cells (Kattman et al. 2006; Zhang et al. 2009), and human liver constructs that retain species-specific drug responses such as induction of P450s, drug metabolism, drug-induced hepatotoxicity and drug-drug interactions (Khetani and Bhatia 2008; Wang et al. 2010). These constructs compartmentalize into functional zones upon exposure to oxygen tension mimicking that along the hepatic sinusoid (Allen and Bhatia 2003; Allen et al. 2005). Even a human gut-on-chip device with intestinal flora and cyclic strain that mimics peristalsis has been developed (Kim et al. 2012).

Several groups have tackled the challenge of culturing multiple organs in a single culture platform, for studying systemic effects of drugs, or cell-cell communication via metabolites. Li et al, for example, have developed a simplified "well within a well" system in which various cell types are cultured in isolation from one another but connected via medium (Li 2008; Li et al. 2004), thus reproducing some organ interactions. When the chemotherapeutic agent tamoxifen was tested in this system against 6 cell types (liver, kidney, lung, central nervous system, blood vessels, and breast cancer) at increasing doses, cell survival values were obtained for each cell type, although rigorous characterization of multi-organ interactions were not performed.

Ma et al have developed a microfluidic device for testing metabolism-dependent drug toxicity. This device also contains two separate chamber arrays, a PDMS quartz layer containing human liver microsomes, and another for cell culture (Ma et al. 2009). Drug-containing solutions can be introduced continuously into the top layer, diffuse through the liver microsomes, and drugs and liver metabolites diffuse to the cell culture chamber. When tested with doses of Acetaminophen above saturation of the detoxification pathway that converts Acetaminophen to non-toxic conjugates, cytotoxic effects were increased, and subsequently reduced when administered below saturation doses. When tested for drug interactions with Acetaminophen, experimental results again aligned with expectations of cytotoxicity. Although the multi-tissue *in vitro* systems attempt to go beyond merely testing direct effects of drugs, towards reproducing some metabolism-dependent actions of the drug, one potentially important aspect which is lacking in these systems is their ability to mimic the dynamics of multi-organ interactions, and especially the physiologically realistic exchange of metabolites.

Several groups have attempted to address these issues. Shuler and colleagues have developed an integrated microscale cell culture analogue (called the  $\mu$ CCA) with several interconnected compartments linked by circulating culture medium (serving as a blood

surrogate) for liver, lung, and fat cells (Ghanem and Shuler 2000, 2000; Sin et al. 2004; Viravaidya and Shuler 2004; Viravaidya et al. 2004). Interestingly, the first prototype of their device was made using two milk dilution bottles and a spinner flask (Sweeney et al. 1995)! The system was designed using simple physiologically based pharmacokinetic models, and used to analyze metabolic products of xenobiotics (in particular, naphthalene) and multidrug resistance with chemotherapy (Tatosian and Shuler 2009), using µL volumes. Two main problems with the device are the fluid links between the compartments that do not model certain nonlinear parameters such as organ volume, and the compartmentalized nature of tissues connected via systemic circulation, which includes some clearing of wastes.

Ahluwalia and colleagues have attempted to better mimic physiological cell-cell interactions. The chambers in their "multicompartment connected culture bioreactor" (MCB) system differ in dimensions, fabrication method, design principles, and applications, in order to mimic physiological cell-cell interactions with much more fidelity than previous systems (Vozzi et al. 2009; Vozzi et al. 2011). In particular, their device was designed with the goal to mathematically correlate nonlinear quantities (organ volume, blood flow, blood retention time, and metabolic rate) and conserve not only kinetic but also metabolic, volumetric, and exchange rate relationships between cells. The system was tested with murine hepatocytes and human umbilical vein endothelial cells, while monitoring albumin, urea, lactate and viability in a connected culture and monoculture in the bioreactor. When the two cell types were connected, increases in endothelial cell viability, hepatic glucose synthesis and albumin and urea production were observed, along with the decrease in lactate production. Although they only studied two cell types for a short period of time, their results showed potential for enhancing cell function, in particular using heterotypic signals from endothelial cells in cultures of hepatocytes.

Zhang, Zhao and colleagues have attempted to link together multiple cell types while maintaining compartmentalization of tissues within a single device (Zhang et al. 2009). Culture medium was supplied through channels to liver, kidney, and adipose chambers and optimized for the four tissue types by supplementation od localized controlled release of organ-specific growth factors (e.g. TGF- 1). Such a technique could, in principle, be applied to deliver a particular biochemical component to only specific cell types without compromising the functions of other cell types and potentially extend culture times.

Going forward, challenges with "human-on-chip" technologies revolve around achieving better predictability of *in vivo* conditions through more authentic representation of cellular behavior, and to construct systems in which conditions are parallelized, in order to achieve higher numbers of test conditions (currently this is an area of future work for the field). By producing consistent, realistic results, these devices will be more trusted as models capable of predicting hitherto unknown reactions and pathways in drug metabolism, for healthy and diseased tissues, and for personalized medicine applications. Ongoing developments of optical and electrical techniques compatible with multiplexing, could further facilitate these efforts. Finally, as the *in vitro* results become more predictable, systems that are easy to operate, more portable, and are less dependent on large equipment will become more and more important.

Translation of stem cell and drug discovery into clinical practice will continue to critically depend on preclinical studies of drug toxicity and disease. As engineered human tissues are being considered for *in vitro* applications in drug toxicity and disease models, advances in connecting different systems to model cross-talk and physiological interaction between organ systems will become of increasing importance. At this point, multi-tissue platforms that can truly simulate human multi-tissue interactions with spatial-temporal control represent a formidable goal. Fortunately, as tissue engineering approaches continue to make

strides towards authentically reproducing the actual milieu of development, regeneration, and disease, the goal of truly developing integrated human physiology *in vitro* via controllable, modular, dynamic, interconnected human tissues, is becoming more attainable. As body-on-chip devices improve the links between *in vitro* and *in vivo* studies, we may hope that the demand for animal studies will be reduced, clinical trials will become more effective, and we may begin to enable a more personalized approach to evaluation of drug regimens.

### 4.3. Case study 5: Integration of advanced models with novel stem cell sources for studying human disease

Integration of advanced microbioreactors and bioengineered environments with novel stem cell sources can open new perspectives in understanding the fundamental biology of disease, towards more personalized approaches to studying the mechanisms of disease, and to develop treatment therapies (Bhadriraju and Chen 2002). *In vitro* models of primary cells or established cell lines are widely used as cellular screening models in toxicology (Rolletschek et al. 2004). Human cardiomyocytes derived from hESCs were used for more reliable cardiac safety pharmacology assays (Braam et al. 2010; Davis et al. 2011). In this study, patch clamp analyses and multichannel electrode arrays (MEA, increasing the numbers of test conditions and/or replicates of the system) allowed generation of field potential duration (FPD) values following exposure to different drugs. A dose-response effect on the prolongation of the FPD values was recorded and comparable to the known drug effects on QT intervals (15÷50% increases in QT time were measured for various drugs at therapeutic concentrations, and was further prolonged at higher concentrations). This method proved to be reliable for preclinical evaluation of new drugs and thus might be used in replacement of the limited, currently used assays.

Since the onset of hESC biology, researchers started focusing on modeling human disease "in a dish" and trying to elucidate their mechanisms. Successful attempts saw the production of hESC lines for cystic fibrosis (Mateizel et al. 2006; Pickering et al. 2005), Huntington's disease (Bradley et al. 2011; Mateizel et al. 2006), and Fragile X syndrome (Eiges et al. 2007). However, the methods used to obtain genetically modified hESC are still inefficient (Giudice and Trounson 2008) thus failing to provide a solid source for predictive disease models that can be experimentally manipulated in vitro. iPS technology can offer an alternative after proving the feasibility of deriving patient-specific pluripotent stem cells, which can be used for a multiplicity of purposes, spanning from drug screenings, to cell replacement therapies and tissue engineering, and to disease modeling studies (Colman 2008).

Recently, two independent groups described the derivation of iPS cells from terminally differentiated mouse cells in stirred suspension cultures (Fluri et al. 2012; Shafa et al. 2012). These reprogrammed cells successfully responded to the "score-criteria" for iPSCs; the implementation of these novel strategies would thus enable standardized, scalable production and differentiation of cells to relevant phenotypes.

A series of diseases have already been successfully modeled *in vitro* via reprogramming of patient cells to iPS cells followed by differentiation into disease-relevant cells and tissues: neurodegenerative diseases, haematopoietic disorders, metabolic conditions and cardiovascular pathologies (Chamberlain et al. 2008; Ebert et al. 2009; Nishikawa et al. 2008; Park et al. 2008; Unternaehrer and Daley 2011). Patient-specific iPS cells were derived from Rett syndrome patients (Ananiev et al. 2011), successfully differentiated into neurons and used to better interpret the disease. Also, iPSCs were generated from patients with Amyotrophic lateral sclerosis (ALS), and were successfully directed to differentiate into motor neurons, the cell type destroyed in ALS (Dimos et al. 2008). Lee and Studer (Lee

and Studer 2011) modeled familial dysautonomia (FD) and demonstrated a partial rescue of few disease-specific phenotypes in FD-iPS-derived via kinetin treatment. These FD-iPSCs could be a fundamental tool for high throughput screens of candidate drugs to reverse *in vitro* phenotypes and for mechanistic studies aimed at understanding disease pathogenesis.

It can be argued that the dynamics of disease progression in diseases such as Alzheimer's or Parkinson's develop in dramatically different scales in the patient and "in the dish". However, researchers are now starting to adopt strategies to accelerate the onset of the pathological phenotypes in culture. Along this line, primary fibroblasts from patients with Parkinson's (Soldner et al. 2009) and familial Alzheimer's disease were successfully reprogrammed, differentiated into neurons and thoroughly characterized (Israel et al. 2012). In this case, iPSCs were successfully used to observe early onstage of Alzheimer's-related phenotypes, even though it can take decades for overt disease to manifest in patients.

### 5. Current challenges

Advanced tissue culture platforms, recapitulating the *in vivo* environment in a controllable manner, have provided great tools to aid researchers in the transition from cell studies to animal models, human clinical trials and into the clinic. Advanced 3D and micro-bioreactor systems each offer unique advantages and present unique limitations associated with their respective size, providing unique engineering tradeoffs and design considerations for *in vitro* control of the stem cell niche (Figure 4). The challenges that remain to be resolved as the field progresses may be categorized into two main themes: (*i*) to establish conditions that are predictive of cell behavior *in vivo*, and (*ii*) to provide bioreactors beyond the laboratory bench.

### 5.1. Producing conditions more predictive of cell behavior in vivo

Although bioreactors do outperform standard culture systems in their ability to provide matrix, molecular, and biophysical factors to cultured cells, current systems still lag behind the nature's ability to deliver highly coordinated sequences of regulatory factors at the level of the cell, which are necessary to regulate cell function in a developing and adult organism.

For reasons that are still not fully understood, researchers have achieved more success with engineering mature tissues of certain types (such as cardiac and osteochondral) than other types (such as pancreatic and hepatic). Even with certain tissue types that grow reasonably well in bioreactor culture (such as bone and cartilage), challenges remain with respect to culturing multiple cell types on a single platform, whether in a 3D or micro-bioreactor configuration.

This issue is further complicated when considering *in vitro* models of disease: although many disease models have been developed using iPSCs, reproducible and consistent results between clones and lines have not yet been achieved. As for hESC lines, challenges remain in generating consistent total numbers of specific cell types, but the issue is further complicated by analysis of whether these cells undergo longitudinal changes that define the disease process. And although it may be argued that it is disease progression that may provide the most reliable assay with regard to compounds that could slow or prevent disease-specific cell death, until appropriately mature phenotypes are achieved consistently in culture, the field will face challenges with adapting engineered tissues to drug screening, disease models or implantable tissue grafts.

Another related challenge is that some known biophysical cues, such as molecular gradients, although relatively easy to apply in microfluidic conditions, are difficult to produce in the larger-scale 3D culture bioreactors, without applying significant shear stresses or using

excessive amounts of expensive culture medium or scarce cells. Others, such as mechanical stimuli, extracellular matrix components, or even long-term cell culture conditions, may be easier to recapitulate in a 3D setting than within a micro-bioreactor.

Perhaps some of these challenges will be ameliorated through future high-throughput combinatorial studies performed in micro-bioreactors, which may shed more light on how tissues emerge from coordinated sequences of cell proliferation, differentiation, and functional assembly that are orchestrated by factors originating from the surrounding cells, matrix, and the external environment. In addition, future research on regeneration of adult tissues will likely help understand how cells respond to the entire milieu of injury or disease, and incorporate the most critical biophysical and/or molecular cues into the repertoire of current bioreactor systems. Finally, improvements in data acquisition and analysis techniques will aid the field in keeping up with an ever-expanding data set and with incorporating knowledge gleaned from basic research into even more advanced cell culture platforms.

### 5.2. Providing bioreactors beyond the laboratory bench

Challenges also remain in the area of bioreactor fabrication, operation, and medical regulation (Figure 4). Many of the bioreactor systems implemented up to date are at the development stage, and have been tested in stem cell research studies but not in basic biological studies. Closer collaboration between fields is expected to widen the opportunities for applying and optimizing bioreactor systems, and help disseminate user-friendly devices that can be easily and robustly operated. In the meantime, because bioreactors are often custom-designed, and because cells are highly sensitive to changes in their environment, even minute changes in molecular and biophysical cues may have implications on the reproducibility of the bioreactor regulation of cell survival, behavior, and differentiation.

A significant challenge is in the scale-up, standardization, and automation of cell culture protocols and bioreactor platforms. Automation will need to be applied wherever possible throughout the production pipeline, in order to derive the appropriate numbers of required cells, during culture when cells are matured (ie changing media etc), and coupled with quality control.

Furthermore, as bioreactor technology systems support engineering 3D tissue grafts for implantation, there will be further challenges to face related to the complexity of regulatory issues, and the design constraints of surgical techniques and the operating theater. For tissue-engineered constructs to be viable and clinically useful, for example, bioreactor design and protocols will need to be compatible with the transportation of biologically stable tissue engineered implants. In the meantime, *in vitro* applications for engineered tissues, such as experimental models of development and disease and chemical screening – will likely accelerate clinical acceptance of engineered tissues for implantation.

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### Figure 1. Development of culture systems

The progression from traditional cultures with animal cells and cell lines towards scaffoldbioreactor systems with human adult, embryonic and iPS cells. The new tissue engineering technologies are paving the way to the new generation of *in vitro* disease models, drug screening systems, and tissue-engineered implantable grafts. Tandon et al.



#### Figure 2. 3D tissue engineering

The so-called "biomimetic paradigm of tissue engineering" includes the cultivation of stem cells on scaffolds within bioreactors that provide *in-vivo*-like environments and produce differentiated, mature tissues. Engineering of anatomically-shaped bone, corresponding to case study #2 in **Section 3.2**, is included in the lower panel.

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#### Figure 3. Miniature bioreactors

Miniature bioreactors have facilitated the precise, systematic studies of stem cell environments via (A) lab-on-chip devices mimicking organs and organ interactions (outlined in **Section 4.2**), and (B) high-throughput screening of environmental parameters (outlined in **Section 4.1**).

	Micro-bioreactors	3-D tissue bioreactors
Applications	• Drug discovery • Study mechanisms of disease • Reduce need for animal testing	<ul> <li>Restore structure and function of damaged tissues</li> <li>Study tissue development</li> </ul>
Advantages	<ul> <li>Increase throughput of chemical and microenvironmental screening</li> <li>Reduce need for animal testing</li> <li>Predict drug interactions</li> </ul>	<ul> <li>Produce autologous transplants</li> <li>Improve integration with host tissue</li> <li>Reduce transplantation needs</li> </ul>
Limitations	<ul> <li>Incomplete recapitulation of 3-D tissue Limited maturity of the cells</li> <li>Single medium limits co-culture ability</li> <li>Labor intensive operation protocols</li> <li>Difficulties with long-term cell culture</li> </ul>	<ul> <li>Large cell numbers (i.e. billions)</li> <li>Labor intensive cell culture protocols</li> <li>Regulatory acceptance</li> <li>Limited maturity of engineered tissues</li> <li>Post-implantation vascularization</li> </ul>

### Figure 4. Advanced culture platforms

Micro-bioreactors and 3-D scaffold/bioreactor systems offer unique advantages and limitations associated with their respective size scale. These characteristics provide engineering tradeoffs and design considerations for *in vitro* control of the stem cell niche.