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RECEIVED 07 June 2025 ACCEPTED 03 September 2025 PUBLISHED 29 September 2025

CITATION

Stephanie GA and Vander Roest AS (2025) Cardiac disease mechanobiology: advances using hiPSC-CMs.

Front. Cardiovasc. Med. 12:1642931. doi: 10.3389/fcvm.2025.1642931

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Cardiac disease mechanobiology: advances using hiPSC-CMs

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Cardiovascular diseases (CVDs) remain the leading cause of mortality worldwide. Despite advances in diagnosis and treatment, there is a critical need for sophisticated models that accurately reflect human cardiovascular pathophysiology. This mini review explores recent advancements in cardiac microenvironment engineering for modeling cardiac mechanobiology and investigating genetic and acquired cardiac diseases. Cardiac function relies heavily on mechanical cues, with integrin- and cadherin-based adhesion complexes mediating mechanosensitive signaling that drives disease progression. However, studying these processes in humans remains challenging. Although animal models have been indispensable, they often fail to recapitulate human-specific cardiac features. Human-induced pluripotent stem cells (hiPSCs) have been transformative, enabling patient-specific modeling and the identification of disease-specific phenotypes that are challenging to replicate in traditional animal models. Despite their promise, hiPSC-CMs are constrained by their immature phenotype and heterogeneity, which limits their efficacy in modeling adult cardiac physiology. Emerging in vitro systems, particularly those engineered using biomaterials such as hydrogels, address these limitations by mimicking the mechanical and biochemical environment of native cardiac tissue. We discuss the potential and challenges of these hiPSC-derived cardiomyocytes (hiPSC-CMs) in modeling cardiac mechanotransduction, focusing on the interplay between mechanical stress and cellular maturation, mechanics, and signaling. By integrating advanced biomaterials and genome editing technologies, these in vitro platforms hold the potential to revolutionize cardiac research, offering the prospect of more precise interventions and improved patient outcomes.

KEYWORDS

hiPSC-CMs, mechanobiology, disease modeling, engineered platforms, cardiomyopathies

Introduction

Cardiovascular diseases are characterized by altered mechanical function and remain the leading cause of mortality globally despite progress in diagnosis and treatment (1). The field of molecular genetics has identified key mutations and pathways involved in disease onset and progression, improving diagnosis and informing therapies (2). Additionally, tissue engineering has revolutionized research on cardiac mechanobiology—the process by which cardiac cells sense and respond to mechanical stimuli in vitro (3). By leveraging advances in material science and cell culture techniques, in vitro cardiac tissue models that mimic the native heart environment can elucidate cardiac physiology and pathology.

Animal models have been pivotal in advancing our understanding of cardiovascular diseases but have significant practical and biological limitations. Surgical maneuvers

including transaortic constriction or shunts have been used in both large and small animal models to induce pressure or volume overload, respectively. These procedures trigger dysfunction and cardiac remodeling, highlighting cardiac sensitivity to changes in mechanics in disease (4). Mouse models are commonly used due to their short reproductive cycles, low cost and ease of genetic manipulation. However, species-specific differences in cardiac physiology and genetic composition limit their translational relevance (5-8). For example, two common transgenic models of inherited hypertrophic cardiomyopathy (HCM)-R403Q-MyHC and R92W-TnT-show marked differences in gene expression, mitochondrial function, and redox balance, not only from each other but also from human myectomy tissue, with minimal overlap in dysregulated genes or pathways between species (8). These findings highlight that genotype alone is insufficient for predicting phenotype or therapeutic responses in cardiac disease modeling. Large animal models such as pigs match the physiology, heart size, immune system and cardiac anatomy of humans, but are limited by the difficulty of introducing disease-causing mutations as well as high rates of sudden cardiac death due to tacharrhythmias (9-11). While primary human cardiomyocytes retain relevant, native structural and electrophysiological properties, their limited

availability, poor viability in culture and lack of scalability pose significant challenges for disease modeling. These limitations underscore the need for complementary models that balance physiological relevance with experimental accessibility (Figure 1).

Human induced pluripotent stem cells (hiPSCs) offer a scalable, patient-specific platform that enables mechanistic insights into cardiac pathophysiology (12, 13). Additionally, genome-editing approaches such as CRISPR-Cas9 can be used to repair disease genotypes in patient-derived hiPSCs or introduce relevant mutations to study their pathophysiology (14–17). Improvements in the efficiency of hiPSC production processes, from the harvesting of somatic cells for hiPSC generation, to the differentiation of cardiomyocytes and other cardiac cell types, have driven advancements in drug discovery, disease modeling, and tissue engineering.

Furthermore, engineered 2D and 3D *in vitro* models that mimic the composition and organization of the cardiac extracellular matrix (ECM), have revolutionized cardiac disease modeling, drug screening technologies, and regenerative medicne (18–21). Hydrogels have become increasingly popular as platforms for modeling cardiac diseases *in vitro*, addressing the limitations of traditional tissue culture plastic, which fail to replicate the complex physiological and

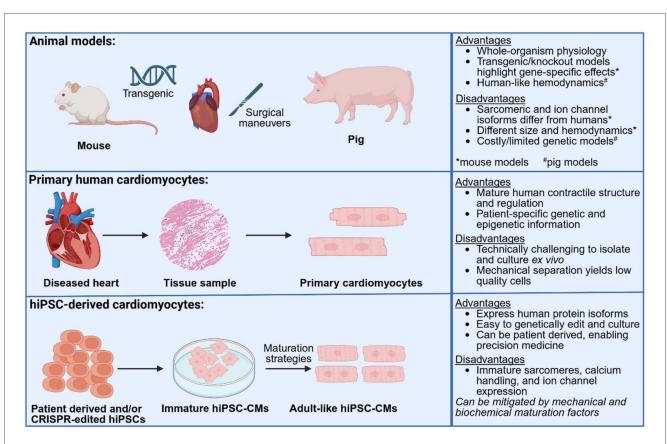


FIGURE 1

Schematic overview of three commonly used platforms: animal models, primary human cardiomyocytes, and hiPSC-CMs. Large (e.g., pig) and small (e.g., mouse) animal models enable whole-organism studies and genetic manipulation, but are limited by species-specific differences and difficulty in modeling human disease heterogeneity. Primary cardiomyocytes from patient tissue retain native structure and genotype but are technically challenging to isolate and culture. hiPSC-CMs offer scalable, patient-specific platforms for modeling cardiac disease and drug response, though they exhibit fetal-like features without appropriate maturation. Each model presents unique advantages and limitations that inform their use in cardiovascular research. Created in BioRender. UIUC, J. (2025) https://BioRender.com/a66j7s6.

dynamic environment of cardiac tissue. Their potential for biofunctionalization, tissue-like elasticity and tunable mechanical properties make hydrogels a suitable platform for modeling cardiac disease, thus facilitating the study of cardiomyocyte function, cell-matrix interactions, cardiac mechanobiology and thus disease progression *in vitro* (22–24). Leveraging these biomaterial platforms allows for the understanding of cardiomyocyte mechanosensing in both healthy and diseased myocardium.

Myocardial mechanobiology: a double-edged sword

The heart undergoes continuous contraction and relaxation cycles and must be able to respond and adapt to varying mechanical stress and hemodynamic load throughout an animal's lifespan. This is enabled by complex remodeling processes that are mediated by both cardiomyocytes and non-myocytes. Cells respond to mechanical cues transduced into biochemical signaling cascades to promote remodeling under physiological and pathological conditions, a process known as mechanotransduction or mechanosensing (25, 26).

Mechanobiology in cardiac development and physiological remodeling

The heart experiences significant changes in load and demand during pre- and post-natal development, increasing in size via hypertrophy and maturation of cardiomyocytes. As the heart grows to meet increased circulatory needs, cardiomyocytes experience continuous mechanical stress from the heart's beating and the surrounding ECM. These biomechanical cues are sensed through mechanotransduction pathways, which coordinate sarcomeric growth and alignment, contributing to the functional organization of the myocardium (27). Under normal conditions, mechanotransduction is a compensatory mechanism that maintains homeostasis as the cell adapts to an altered mechanical environment.

The cytoskeleton plays a crucial role in the mechanotransduction process, linking the cellular and contractile machinery to forcesensing complexes that interface with the ECM. Costameres align with z disks in cardiac muscle and function as attachment sites for transmission of lateral forces to the ECM. They comprise a network of mechanosensitive proteins, including integrins, that coordinate contraction of the heart muscle and are crucial for mechanosensing. Integrins are a family of cell-surface receptors composed of α and β subunits with extracellular domains that bind an ECM ligand while their cytoplasmic domains engage cytoskeletal and force-sensitive proteins like talin, paxillin, and vinculin (28). These domains also recruit intracellular signaling mediators including focal adhesion kinase, IQGAP1, and melusin, initiating cascades such as the Raf-MEK-ERK pathway, which culminates in ERK phosphorylation and cardiomyocyte hypertrophy (29, 30). Beyond signal transduction, integrins contribute to adhesion stability and cytoskeletal organization, influencing cell shape, size, and morphology. The dominant integrins in cardiomyocytes— $\alpha1\beta1$, $\alpha5\beta1$, and $\alpha7\beta1$ —bind to collagen, fibronectin, and laminin, respectively (31). Notably, integrin expression is developmentally regulated; the $\alpha5$ subunit is prevalent in fetal and neonatal cardiomyocytes but is later replaced by $\alpha7$ during maturation. Integrin $\alpha5$ has been shown to be critical for maturation of hiPSC-CMs cultured in 3D hydrogels. Its knockdown impaired epithelial-to-mesenchymal transition, reduced expression of key mesodermal markers (T Brachyury, TBX6), and diminished cardiomyocyte differentiation and contractility. These findings highlight the importance of integrin-linked signaling pathways in shaping hiPSC-CM development and maturation (32).

HiPSC-CMs replated onto collagen I at later stages of differentiation have also been shown to exhibit a shift in integrin isoform expression—particularly upregulating $\alpha 1$ and $\beta 1$ subunits-accompanied by activation of FAK and ERK signaling pathways. This integrin-mediated mechanotransduction cascade drives structural and electrophysiological maturation, including increased expression of adult myosin isoforms and improved calcium handling (33). In parallel, focal adhesion-associated kinases, such as FAK and members of the MAPK family (ERK, JNK, p38), have also been shown to be key modulators of cardiomyocyte specification from pluripotent stem cells, linking external mechanical signals to lineage commitment via phosphorylation of transcriptional and epigenetic regulators. These kinases integrate ECM and growth factor cues to guide mesodermal induction, cardiac progenitor formation, and sarcomeric gene activation, positioning them as essential nodes in both mechanobiological signaling and cardiogenesis (34). Moreover, integrin-linked kinases (ILKs) are critical for cytoskeletal remodeling in reprogramming. iPSC kinome-wide functional analysis suggests a critical role of integrin in the cytoskeletal remodelling process. Specifically, integrin-linked kinases within the integrin-linked kinase (ILK) network play a crucial role in cellular reprogramming through cytoskeletal remodeling (35). These findings highlight the critical role of extracellular matrix composition and the temporal dynamics of cell-matrix interactions in modulating integrin-mediated signaling cascades in development and disease modeling.

Additionally, cell adhesion proteins such as N-cadherin, have been shown to play a critical role in maintaining the structural integrity and function of cardiomyocytes as part of adherens junctions at intercalated discs. N-cadherin binds and stabilizes β -catenin, which is a crucial component of the adherens junction involved in the WNT signaling pathway (36).Standardized hiPSC-CM differentiation protocols primarily target the WNT signaling pathway to direct mesodermal and cardiac lineage specification (37–39).

Mechanobiology in cardiomyopathy and pathological remodeling

Under abnormal conditions, mechanotransduction can drive maladaptive responses and pathological remodeling (40-43).

Recent studies using primary human cardiomyocytes from patients with heart failure have shown that cytoskeletal forces are relayed to the nucleus via desmin and microtubule networks, and that disruption of this architecture can lead to chromatin reorganization, altered gene expression and impaired contractile function (44). Additionally, recent studies leveraging hiPSC-CMs have highlighted the centrality of cytoskeletal components such as filamin C (FLNC), in maintaining these mechanical linkages. FLNC truncation variants disrupt cytoskeletal stiffness, impair cell-ECM adhesion, and induce arrhythmic beating profiles in a gene-dosage-dependent manner. Atomic force microscopy analysis revealed compromised mechanical integrity at both the membrane and cytoplasmic levels, underscoring the importance of costameric and cytoskeletal integrity for proper mechanosignaling (45). Changes in ECM composition and mechanical integrity due to cardiac injuries or genetic cardiomyopathies often result in changes in integrin expression patterns. For instance, integrin expression has been shown to revert to fetal isoforms in murine models, with upregulation of $\alpha 5$, $\alpha 7$, and $\beta 1D$ (46, 47). Additionally, loss of integrin-linked kinase expression in cardiomyocytes has been linked with arrhythmogenic cardiomyopathy in mice (48).

Interestingly, increased β1-integrin expression at the sarcolemma of N-cadherin-deficient murine cardiomyocytes suggests enhanced cell-ECM interactions as an adaptive mechanism for providing additional structural stability in the absence of intact adherens junctions (49). Targeted N-Cadherin deletion in CMs has also been shown to impede cardiac regeneration in neonatal mice, leading to excessive scarring following ischemic injury (50). However, the mechanisms through which these changes drive remodeling at the microscale level are challenging to resolve in vivo. Furthermore, the pathophysiology of many cardiomyopathies manifests differently in animal models. For instance, volume overload caused by extended endurance training can exacerbate the chance of arrhythmia in human patients with ACM-associated mutations, including mutations in the cell-cell junction protein plakophilin 2 (PKP2). However, prolonged volume depletion protected mice with ACM-prone PKP2 mutations from developing arrhythmias, even with prolonged exercise (51). An hiPSC-CM model of arrhythmogenesis showed exaggerated lipogenesis and apoptosis in mutant PKP2 hiPSC-CMs. Furthermore, hiPSC-CMs with a homozygous PKP2 mutation also had calcium-handling deficits potentially associated with arrythmia, a phenotype not observed in ACM mouse models (52). Additionally, mechanosensitive ion channels that are crucial for excitation-contraction recoupling have also been shown to be implicated in pathological remodeling. Recent studies of HCM hiPSC-CMs cultured on stiff polyacrylamide gels have identified hypermetabolic mitochondrial remodeling in HCM due to disrupted calcium handling between the L-type calcium channel and mitochondria (40). These findings further emphasize the need for in vitro platforms that incorporate human-derived cells and allow for spatial and molecular resolution of mechanotransductional remodeling during cardiac pathophysiology.

Engineered *in vitro* systems for cardiac mechanobiology: challenges and opportunities

The application of hiPSC-CMs to in vitro model systems is constrained by their immature, embryonic-like phenotype and intrapopulation heterogeneity, limiting their ability to model adult cardiac physiology (53). For example, hiPSC-CMs have decreased, hierarchical organization of myofibrils, which is crucial in the myocardium, where normal function is highly coupled to tissue organization and mechanics. hiPSC-CM immaturity is further characterized by differences in electrophysiological properties, ion channel expression, and metabolic profiles. This immaturity has been attributed to the lack of in vivo environmental cues that the cells experience during development. For example, hiPSC-derived cardiomyocytes cultured on traditional tissue plastic as monolayers often lack the anisotropic alignment of myofibrils, which is crucial for mechanical transduction and electrical activity (54, 55). The native myocardium has a Young's modulus of 10-18 kPa under physiological conditions, whereas a diseased myocardium can stiffen beyond 50 kPa (56). The Young's modulus of most traditional tissue culture plastics is approximately 1 GPa, about 5-6 orders of magnitude stiffer than even the diseased myocardium. Studies have also shown that both ECM composition and density are crucial in modulating cellular mechanics as well as downstream signaling cascades that promote homeostasis (57, 58). This therefore establishes a need for physiologically relevant materials that allow for resolution of cardiac mechanobiological remodeling under both physiological and pathological conditions. Biomimetic materials have been explored for better replication of the cardiac microenvironment. Key biomaterial features such as stiffness, topography, and viscoelasticity have been shown to critically influence cardiomyocyte alignment, maturation, and force generation. Both synthetic hydrogels such as polyacrylamide gels, and natural hydrogels such as collagen and hyaluronic acid have been explored as alternatives to tissue plastic for in vitro cell culture (59-61).

In addition to bulk material properties, spatial cues have proven to be critical for hiPSC-CM culture. Microfabrication techniques such as microcontact printing (µCP) have been employed to engineer spatially patterned substrates that provide alignment cues and promote myofibril alignment and cardiomyocyte maturation in vitro. Single-cell micropatterning techniques afford precise control over cell shape, polarity, and spatial confinement. Rodshaped micropatterns, designed to mimic the elongated morphology of native cardiomyocytes not only enhance structural organization but also significantly improve electrophysiological maturation, including increased sodium current density, faster action potential upstroke velocity, and improved calcium handling (62). Traction force microscopy analysis of single-cell hiPSC-CMs has also demonstrated that enhanced myofibrillar organization leads to greater sarcomeric activity, improving contractile output as measured by microbead displacement during contractionrelaxation cycles (63). Taken together, these findings show that substrates with physiological stiffness significantly improve contractile activity of patterned hiPSC-CMs, calcium transients,

electrophysiology, mitochondrial organization, and transversetubule formation and facilitate improved disease modeling *in vitro*.

The tunability of hydrogel platforms has also enabled mechanistic insights into pathological remodeling linked to disease-associated changes in tissue stiffness. For example, increasing polyacrylamide substrate stiffness exacerbates Duchenne muscular dystrophy (DMD) hiPSC-CM dysfunction by impairing calcium handling, reducing contractile force, and inducing contraction-dependent telomere shortening. The resulting telomere attrition activates a p53-mediated DNA damage response, linking mechanical stress to progressive cardiomyopathy and identifying telomere maintenance as a potential therapeutic target (64).

While single-cell microcontact printing has been shown to achieve highly consistent single hiPSC-CM geometry, variable contractile function has been reported, potentially due to a lack of connectivity to neighboring cardiomyocytes. Patterns that allow for multi-cellular contact have been used to promote connectivity and enhance mechanical and electrical coupling. For example, innovations in micropatterning, such as the integration of ECM geometry with N-cadherin "end caps" on hydrogels, have been shown to reshape cardiomyocyte morphology with minimal impact on contractile function, suggesting that cell-cell adhesion cues may fine-tune architectural features without substantially altering baseline mechanical output (21). Additionally, developmentally inspired fibronectin micropatterns derived from embryonic heart ECM have been shown to guide cardiomyocyte alignment in a density- and cell-type-dependent manner, mediated in part by N-cadherin-based cell-cell interactions, thus emphasizing the need for spatially and temporally controlled microenvironments that recapitulate both cell-cell and cell-ECM interactions (65).

Importantly, the spatiotemporal dynamics of cell-cell junction assembly have also been shown to be critical for normal hiPSC-CM development. Adherens junctions and desmosomes assemble progressively in response to mechanical and organizational cues, and their maturation is dependent on proper spatial alignment. Disruption of desmosomal proteins in hiPSC-CMs derived from patients with PKP2 mutations impairs junctional integrity, force transmission, and contractile coordination. Notably, these defects can be partially rescued by modulating the mechanical microenvironment to promote cytoskeletal alignment, focal adhesion organization, and enhanced cell-cell contact, reinforcing the concept that junction assembly is tightly regulated by mechanotransduction and cellular architecture (15). Thus, biophysical context, such as cellular alignment and force transmission, modulates the severity of these phenotypes, offering insight into mechanotransduction-dependent pathogenesis.

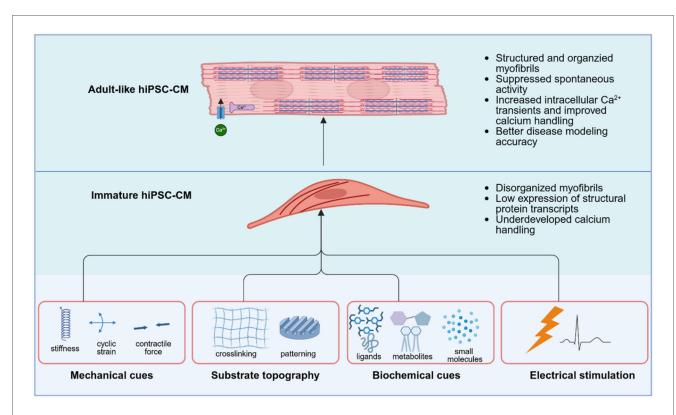
Further, culture substrates that promote robust, mechanically organized cardiac muscle tissue formation yet still allow for contractile analysis in 2D have been reported. Micropatterning of elastic polydimethylsiloxane (PDMS) to anchor thin, purified cardiac muscle strips in 2D arrays showed that myofibrillar alignment and physiological contractions in an optimized media environment drive improved development of contractile function, calcium handling, and electrophysiology (66).

Additionally, recent work has also shown that applying cyclic mechanical stretch to hiPSC-CMs cultured on collagen I-coated

PDMS membranes enhances sarcomeric organization, contractility, and gene expression through nuclear mechanotransduction, reinforcing the importance of mechanical cues in guiding maturation via chromatin remodeling and transcriptional activation (67). Electrical stimulation has also emerged as a crucial tool in promoting the structural and functional maturation of hiPSC-CMs. For example, it has been observed that electrical stimulation applied during differentiation promotes hiPSC-CMs into a specialized conduction system phenotype, marked by increased expression of connexin 40. Additionally, electrically stimulated hiPSC-CMs exhibited faster action potential depolarization, longer intracellular Ca2+ transients, and slower action potential duration at 90% of repolarization (68). In addition to mechanical and electrical stimulation, chemically defined maturation strategies have been developed to emulate the adult cardiac metabolic environment. For example, protocols incorporating reduced glucose, increased fatty acid concentrations (e.g., palmitate, oleate), and galactose have been shown to shift hiPSC-CM metabolism from glycolysis to oxidative phosphorylation, leading to increased mitochondrial content, enhanced membrane potential, and upregulated fatty acid oxidation enzymes and expression of maturation markers (69-71). In conjunction with bioengineering approaches, these hiPSC maturation strategies have improved the electrophysiological, structural, and metabolic fidelity of hiPSC-CMs, advancing their utility in cardiac disease modeling

Extending beyond 2D systems, optimized 3D ECM environments, defined by specific combinations of matrix composition, stiffness, and timing of cell–ECM engagement, significantly enhance murine cardiomyocyte differentiation from embryonic stem cells. Using a statistical modeling framework, fibronectin-based matrices with intermediate stiffness and early exposure timing have been shown to be optimal for promoting cardiac gene expression and sarcomeric organization, reinforcing the importance of ECM mechanobiology in guiding lineage commitment and maturation (72).

Additionally, engineered heart tissues (EHTs), have revolutionized the field of cardiac mechanobiology, have been crucial for modeling the mechanical behavior of the myocardium (73-75). Cell-dense hydrogels, typically a mixture of fibrin and collagen, are cast between two stretching posts. These posts provide mechanical stretching, thus facilitating the modeling of preload and afterload dynamics. EHTs have also been integrated to provide a heart-on-a-chip strategy, wherein engineered, anisotropic ventricular myocardium with muscular thin film are multiplexed to provide real-time, parallel physiological and pharmacological assessment across an ensemble of tissues (76). For example, it has been shown that dynamic loading of wild-type EHTs improves contractile force, enhanced alignment, conduction velocity and contractility. Interestingly, it has also been shown that dynamic loading is necessary to recapitulate ACM due to mutations in the desmoplakin gene (73). Collectively, these platforms have advanced our ability to model biomechanical cues and disease-specific phenotypes in vitro, offering a scalable framework for probing the mechanistic underpinnings of cardiac function and dysfunction with high spatial and physiological fidelity.



Strategies to promote functional maturation of hiPSC-derived cardiomyocytes. Immature hiPSC-CMs exhibit disorganized myofibrils, low expression of structural protein transcripts, and underdeveloped calcium handling, which limit their utility for accurate disease modeling. Multiple bioengineering strategies have been developed to drive their maturation toward an adult-like phenotype, characterized by organized sarcomeres, suppressed spontaneous activity, and enhanced intracellular calcium transients. These strategies include mechanical cues (e.g., substrate stiffness, cyclic strain, and contractile loading), substrate topography (e.g., ECM-crosslinked micropatterning), biochemical modulation and electrical stimulation. Integration of these cues promotes structural, electrophysiological, and metabolic maturation of hiPSC-CMs, thereby improving their fidelity as human cardiac disease models. Created in BioRender. UIUC, J. (2025) https://BioRender.com/a66j7s6.

Beyond disease modeling, hiPSCs' regenerative and pluripotent capacity are also being heavily explored for regenerative therapies. Initial preclinical studies explored various stem cell sources for cardiac repair in animal models. These studies demonstrated that intramyocardial injection of mouse iPSCs following coronary ligation improves cardiac function and attenuates remodeling in immunocompetent mice (77). However, other studies observed teratoma formation in immunodeficient mice, raising concerns about potential risks associated with transplantation of undifferentiated iPSCs (78). To address this, differentiation protocols have been refined to derive committed cardiovascular lineages such as cardiomyocytes, endothelial cells, and smooth muscle cells prior to transplantation, reducing the risk of tumor formation. However, a few challenges remain, such as poor engraftment and limited cell retention at the infarcted site due to rapid clearance. Strategies have shifted towards using tissueengineered constructs, where hiPSC-derived cardiomyocytes are encapsulated within bioactive hydrogels that mimic the native myocardial ECM. For instance, the development of a multicellular cardiac patch consisting of hiPSC-derived cardiomyocytes, smooth muscle cells, endothelial cells and cardiac fibroblasts, significantly enhanced myocardial repair in a porcine model of myocardial infarction (79). Improved left ventricular ejection fraction, vascular

density, enhanced engraftment of mature human cardiomyocytes, and electromechanical integration with host tissue were observed, without the induction of arrhythmias. These findings underscore the promise of hiPSC-derived tissue constructs as next-generation therapeutic platforms for cardiac regeneration.

Conclusions and future perspectives

The use of iPSC-derived cardiomyocytes (CMs) to model cardiac pathophysiology and understand cardiac mechanobiology has ushered in a new era of potential insights into the intricacies of heart disease. These models have allowed researchers to overcome the limitations posed by traditional animal models and primary human cardiac cells, providing a more accurate representation of human-specific cardiac conditions, genetic influences, and patient-specific phenotypes. Despite the ongoing limitations of hiPSCs, advancements in biomaterials, especially hydrogels, alongside genome editing technologies like CRISPR-Cas9, have highlighted their potential in mimicking the physiological and pathological conditions of the human heart. This has allowed the study of mechanobiological signaling pathways and their role in regulating cardiomyocyte maturation, contractility, and disease phenotypes in

vitro. Future research could focus on refining biochemical maturation strategies of hiPSC-CMs as well as enhancing the robustness microengineered cardiac tissues to replicate the dynamics of the native cardiac environment more faithfully.

Author contributions

GS: Writing - original draft. AV: Writing - review & editing.

Funding

The author(s) declare that financial support was received for the research and/or publication of this article. The authors received funding from the NIH (HL153679-05) which supported the preparation of this minireview.

Conflict of interest

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

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The handling editor AMR declared a shared affiliation with the author AVR at the time of review.

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