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## Two-Dimensional Culture Systems to Enable Mechanics-Based Assays for Stem Cell-Derived Cardiomyocytes

J. Notbohm<sup>1,2,\*</sup>, B.N. Napiwocki<sup>2,3</sup>, W.J. deLange<sup>4</sup>, A. Stempien<sup>2,3</sup>, A. Saraswathibhatla<sup>1</sup>, R.J. Craven<sup>2,3</sup>, M.R. Salick<sup>3,5</sup>, J.C. Ralphe<sup>4</sup>, W.C. Crone<sup>1,2,3,5,\*</sup>

<sup>1</sup>Department of Engineering Physics, University of Wisconsin-Madison, Madison WI, USA

<sup>2</sup>Department of Biomedical Engineering, University of Wisconsin-Madison, Madison WI, USA

<sup>3</sup>Wisconsin Institute for Discovery, University of Wisconsin-Madison, Madison WI, USA

<sup>4</sup>Department of Pediatrics, University of Wisconsin School of Medicine and Public Health, Madison, WI, USA

<sup>5</sup>Department of Materials Science and Engineering, University of Wisconsin-Madison, Madison WI, USA

### Abstract

Well-controlled 2D cell culture systems advance basic investigations in cell biology and provide innovative platforms for drug development, toxicity testing, and diagnostic assays. These cell culture systems have become more advanced in order to provide and to quantify the appropriate biomechanical and biochemical cues that mimic the milieu of conditions present *in vivo*. Here we present an innovative 2D cell culture system to investigate human stem cell-derived cardiomyocytes, the muscle cells of the heart responsible for pumping blood throughout the body. We designed our 2D cell culture platform to control intracellular features to produce adult-like cardiomyocyte organization with connectivity and anisotropic conduction comparable to the native heart, and combined it with optical microscopy to quantify cell-cell and cell-substrate mechanical interactions. We show the measurement of forces and displacements that occur within individual cells, between neighboring cells, and between cells and their surrounding matrix. This system has broad potential to expand our understanding of tissue physiology, with particular advantages for the study of the mechanically active heart. Furthermore, this technique should prove valuable in screening potential drugs for efficacy and testing for toxicity.

### Keywords

Cardiomyocyte; Cell culture systems; Micropatterning; Traction force microscopy; Digital image correlation

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\*Corresponding authors jknotbohm@wisc.edu and crone@engr.wisc.edu.

## INTRODUCTION

It is estimated that more than 92 million American adults have at least one form of cardiovascular disease (CVD) [1]. Fortunately, the mortality of CVD has decreased over the last several decades due to improvements in evidence-based medical therapies and a reduction in the prevalence of risk factors [2]. In spite of these promising developments, CVD continues to be the leading cause of death in the United States. Annually, in the United States alone, there are an estimated 610,000 new myocardial infarctions (MI), commonly referred to as a heart attack, and 325,000 recurrent MIs causing approximately 114,000 deaths [3,4].

One of the reasons for the extremely high mortality rate following an MI is the human heart's inability to regenerate. Unlike cells present in several other human tissues [5] or the hearts of some other species [6], the proliferation rate of human cardiomyocytes is extremely low. As few as 1% of human cardiomyocytes replicate annually in adults at 25 years of age, further declining to 0.45% by the age of 75 [7]. Carbon-dating studies have shown that less than 50% of cardiomyocytes are regenerated throughout an entire typical lifespan [7]. In the absence of ability to regenerate lost cardiomyocytes, the myocardial response to infarction triggers a complement cascade that produces a large region of collagen-dense scar tissue to replace the damaged myocardium [8]. This scar tissue is profoundly different from the surrounding myocardium in both mechanical [9] and electrophysiological [10] properties. This heterogeneity and loss of functional muscle contributes to the often relentless progressive heart failure following a MI.

Early stage research is investigating the introduction of stem cells and cardiac progenitor cells to the heart to aid in repair after myocardial infarction [11]. There are also a number of ways in which stem cell-derived cardiomyocytes are already being used in disease modeling, drug screening, toxicity testing, and myocardial tissue repair [12,13]. As this new field continues to develop, more robust and reproducible approaches are needed to promote the physiologic maturation of stem-cell derived CMs and to reduce batch-to-batch variability. Additionally, more information is needed pertaining to how these cells function mechanically, and how their local environment influences their biophysical properties.

Numerous investigators have explored a variety of engineered highly-controlled 2D and 3D constructs that generate reproducible *in vitro* culture platforms to investigate basic cell-cell and cell-extracellular matrix (ECM) interactions [14–24]. 3D culture methods allow cell-cell interactions to occur in a setting that is often more relevant to native biology, which is particularly important when observing phenomena such as organogenesis. Contraction in 3D tissues can be quantified by using force transducers, post deflection, and optical tracking, for example [25–29]. Whereas these methods quantify contraction of the entire tissue, they preclude detailed assessment of single cell function, and they are of relatively low throughput. Advances in lightsheet microscopy [30,31], expansion microscopy [32], and single-cell sequencing [33,34] have begun to address these challenges, but these are still low-throughput in comparison to 2D culture systems.

2D culture methods continue to be a mainstay in the field because they can provide a highly controlled and easily manipulable environment, scalable features with high precision and fidelity, and ease of use for imaging and analysis. 2D systems have advanced basic investigations in cell biology, providing details of cellular architecture, molecular structures, protein interactions, and signal transduction pathways. 2D systems have also provided tools for disease modeling, and high-throughput platforms for drug development and toxicity testing. However, 2D systems do have drawbacks due to limited morphogenesis and differences in cell migration patterns compared to 3D cultures and the *in vivo* environment. A major limitation to traditional 2D cardiomyocyte cultures has been the disorganization of the cells' contractile apparatus, which is in marked contrast to the highly aligned and organized features of *in vivo* cardiac tissue. For detailed contractile physiology studies, a consistently organized, integrated multicellular system is needed. Here we present techniques for studying stem cell derived cardiomyocytes in a 2D cell culture platforms that recapitulated *in vivo* cellular alignment as well as other key markers of maturity. When combined with optical microscopy, this platform enables mechanics experiments to quantify cell-cell and cell-substrate mechanical interactions.

## BACKGROUND

### The Cardiomyocyte

Blood flow in the body results from collective force generated by each cardiomyocyte (CM) of the heart. CM contraction is a complex, choreographed event that involves an action potential change in voltage across the cell membrane, calcium ( $\text{Ca}^{2+}$ ) release within the cell, and mechanical contraction of the cell. The contraction is produced by a highly organized apparatus, formed by numerous cytoskeletal structural, motor, and regulatory proteins that enable the cell to produce force in response to high  $\text{Ca}^{2+}$  concentration within the cytoplasm inside the cell. The individual repeating units of the contractile apparatus are the sarcomeres, which align end-to-end to produce long myofibrils that span the length of the CM and, when provided the appropriate environmental cues, align laterally adjacent myofibrils. Through numerous connections with neighboring cells, the summation of sarcomere contractile events contributes to the overall contraction and relaxation of the cardiac tissue within a chamber of the heart so that blood can be pumped to distant tissues. Inefficient and aberrant contraction in heart disease can be caused by abnormalities in myofibrillar structure, inefficiencies in contractile force generation and relaxation, and disruption of the normal temporal and spatial variation of  $\text{Ca}^{2+}$  flux during excitation-contraction [35–39].

### Human Stem Cell-Derived Cardiomyocytes

Scientific study of human cardiac physiology would ideally be conducted using well characterized human cells and tissues in which as many variables as possible can be controlled. In many studies of the human heart, animal models are required because of limited availability of human cells and tissue. Unfortunately, animal models for the human heart inherently have significant drawbacks due to fundamental physiologic differences between species [40,41]. For example, the hearts of small laboratory rodents have different action potentials, roughly five times the resting heart rate of humans, and their force-frequency relation is opposite that of human hearts [42]. Large animal models match human

heart characteristics more closely, but they are substantially more expensive than small animal models [43]. Access to human heart tissue for research purposes is exceedingly challenging. Additionally, human heart tissue available for research is often obtained only at a significant state of advanced disease. Appropriate healthy controls are frequently not available, because the demand for heart transplants currently exceeds supply. The emergence of stem cell technologies and the ability to produce large numbers of human stem cell-derived CMs provides a promising new avenue for cardiac research. Stem cell-derived CMs have the potential to support groundbreaking discoveries, particularly if human adult-like CMs *in vitro* can be produced reliably and cultured to maturity.

Human embryonic stem cells (hESCs), which are pluripotent cells capable of long-term proliferation, were first successfully isolated from *in vitro* fertilization embryos in 1998 at the University of Wisconsin–Madison [44]. This discovery has led to further breakthroughs in the fields of embryology, cellular biology, tissue engineering, and regenerative medicine. Following this extremely promising discovery, researchers focused their efforts on developing methods for producing specific cell types through a wide variety of methods. As a result, we are now capable of producing a wide array of specialized cell types, including cardiomyocytes [45]. Additionally, the ability to generate induced pluripotent stem cells (iPSCs) from adult cells from any individual [46,47] provides added opportunities for creating disease models, improved drug screens, and clinical therapies that evade autoimmune rejection.

Recent investigations have developed increasingly effective means of producing cardiomyocytes from pluripotent stem cells, which have increased production efficiencies of up to 98% [48,49]. Although these cells have committed to the cardiac lineage, they are immature and only begin to show some markers of maturation after many months in standard 2D culture [50].

## 2D Patterned Substrates

One of the most attractive features of 2D systems is the relative ease with which images can be captured of the cells and their substrates through bright field and fluorescent microscopy. Cells cultured on a micropatterned substrate can be treated with immunostaining techniques to visualize cellular components of interest. Bright field observations and live imaging techniques can be used in concert with image tracking or digital image correlation software. With the incorporation of tracking particles within the substrate, techniques such as traction force microscopy can be employed. As their role is to produce force, cardiomyocytes in particular lend themselves to biomechanics analysis.

Muscle cells respond to certain micropatterns in 2D culture systems with improved cell-cell junctions, increased contractile force, and greater sarcomere organization [51–53]. These micropatterns also influence cell shape and cell attachment [54,55]. The majority of studies involving micropatterned CMs have used rat neonatal ventricular CMs. Deriving human CMs from pluripotent stem cells is a more recent achievement, so researchers have only begun to examine how engineered substrate conditions influence behavior of human CMs. For example, the human CM excitation-contraction coupling and sarcomere structure have been improved by culturing cell populations of CMs onto mesh-like micropatterns for tissue

engineering applications [14]. Culturing human CMs in micropatterned arrays also allows for high-throughput screening [56]. Micropatterning has also been used to demonstrate that cell shape and substrate stiffness impact shortening of sarcomeres and the translation of sarcomere activity to contractility [57].

Although glass and tissue culture plastic are commonly used substrates in cell culture, a wide range of polymeric materials can be employed to more closely mimic the mechanical properties of the *in vivo* environment. In addition to the type and pattern of proteins available to the cells for attachment, the mechanical stiffness of the system is important and allows the cell to display more mature behavior [58]. Cardiac tissue *in vivo* has a Young's modulus of 4 to 50 kPa, with the lowest values being that of healthy myocardium [59]. The desired stiffness can be created with substrates composed of polydimethylsiloxane (PDMS) with tunable stiffness [60–63], polyacrylamide (PA), which can be made with a range of moduli spanning from <1 kPa to >100 kPa [64–66], or alginate, which has also been shown to have a high level of mechanical property controllability through its ionic and covalent crosslinking [67] and can be blended at different ratios with gelatin to modify its mechanical properties for CM culture [68].

In this work we focus on 2D cell culture systems and use micropatterning techniques to create ECM patterns on substrates ranging from glass [69] to polymeric materials such as PA and PDMS in order to examine how patterning impacts the organization and alignment of a CM's contractile apparatus and its ability to enhance maturation of the CMs. When direct microcontact printing is not feasible, we use sacrificial films [60] to transfer ECM patterns to substrates. Importantly, the polymeric substrates can be easily modified to incorporate fluorescent particles that enable techniques such as traction force microscopy to be utilized [70,71,62].

## MATERIALS AND METHODS

### Stem Cell Differentiation

In the research reported here, immature CMs were differentiated from pluripotent stem cells using a modified version of the small molecule Wnt-agonist method [46]. Briefly, on day –3 the cTnT-GFP stem cells were seeded onto Matrigel coated (BD Biosciences) 12-well plates at a density of 400,000 cells per well and fed with E8 medium supplemented with 5  $\mu$ M ROCK inhibitor (Tocris). On day –1, the cells were again fed with E8 medium. On day 0 (exactly 72 hours after initial seeding), cells were treated with RPMI (Life Technologies) supplemented with B27-insulin (Life Technologies), 12  $\mu$ M of the Wnt agonist CHIR 99021 (Tocris) and 1  $\mu$ g/ml of insulin (Sigma). On day 1, cells were fed with RPMI supplemented with B27-insulin. On day 3 for hESCs and when confluent for iPSCs, the cells were treated with RPMI supplemented with B27-insulin and 10  $\mu$ M of the Wnt inhibitor IWP4 (SemGent). On day 5, cells were fed with RPMI containing B27-insulin and then treated with RPMI containing B27-complete on days 7, 9, and every other day until day 15.

On day 15 the media was switched to lactate media to begin CM purification[72]. The lactate media consisted of RPMI-glucose (Life Technologies), B27-complete (Life Technologies) and 5mM lactate (Sigma). The cells were exposed to lactate for 10 days and

then the media was changed back to RPMI containing B27-complete on day 25. Spent medium was replaced with fresh RPMI+B27 every three days thereafter. Purified stem cell-derived CM cultures were washed with PBS and a 10–15 minute exposure to TrypLE (Life Technologies) was then conducted to dissociate and singularize CMs before seeding them onto micropatterned substrates in a 10% serum-containing media.

### Engineered Substrate System

The results reported below were produced on polymeric substrates of PDMS or PA. PDMS of desired stiffness was created by blending Sylgard 184 with Sylgard 527 (Dow Corning) [73]. Sylgard 184 was made by mixing ten parts base to one part curing agent, and Sylgard 527 was prepared by mixing equal weights of part A and part B. Both Sylgard 184 and 527 were mixed for 5 min with a glass stir rod. After each blend was properly mixed, they were then combined in different mass ratios of Sylgard 184:527 and again mixed for 5 min. The mass ratio of 1:10 produced a substrate with Young's modulus of approximately 10 kPa [60]. Once mixed, the PDMS was poured into 100 mm diameter petri dishes and cured overnight at 60°C. The following day the samples were cut into the correct shape with a razor blade and UV sterilized prior to patterning with extracellular matrix proteins.

PA is used for the Traction Force Microscopy results presented below. A solution of 3% acrylamide and 0.1% bisacrylamide was prepared to give a Young's modulus of 1 kPa upon polymerization. Simply mixing fluorescent particles into the prepolymerized solution results in the particles being dispersed throughout the bulk of the substrate. As tractions applied by the cells to the substrate occur only on the top surface of the PA gel, more accuracy is obtained by embedding the particles only at the top surface. For this, we centrifuge the gels upside down during polymerization. The centrifugation brings the fluorescent particles to the top surface of the gel, thereby allowing for imaging with a common wide field fluorescent microscope. We use a high concentration (0.07% volume/volume) of 0.5  $\mu\text{m}$  particles (Fluospheres, Life Technologies), which together produce high contrast bright and dark spots within the image. To avoid leaking of the prepolymerized gel solution during centrifugation, we sometimes use a 2-step method as follows. A 100  $\mu\text{m}$  thick polyacrylamide gel is fully polymerized with no particles. Then, a second gel  $\sim 10$   $\mu\text{m}$  thick with particles is polymerized on the first and centrifuged upside down during polymerization. Although not used in this study, other authors report the fluorescent particles can also be incorporated in PDMS [74] [75].

To accomplish the micropatterning of cells in lanes on soft substrates, reusable microcontact printing stamps were fabricated on a silicon wafer (FlowJEM, Toronto, ON, Canada) with the specific micropatterns designed for study. The PDMS (Sylgard 184) was poured on top of the patterned silicon wafer and cured at 60°C overnight to produce a reverse replica of the silicon wafer. After removal from the silicon wafer, the PDMS was cut into individual stamps which were coated with ECM proteins overnight (83  $\mu\text{g}/\text{mL}$  Matrigel). Using microcontact printing, the ECM-coated PDMS stamps were used to transfer lanes of ECM to the substrate. Fig. 1 illustrates how the micropatterning can be used to transfer Matrigel onto various substrates, so that experiments can be conducted on glass or on a substrate with a more physiologically relevant stiffness.

For patterning on compliant substrates, such as in Fig. 1b, stamping of ECM proteins was conducted on a sacrificial polyvinyl alcohol (PVA) film which was then brought into conformal contact with the substrate [76]. To make a sacrificial film, 0.5 g of PVA (Sigma Aldrich) was dissolved in 10 mL of deionized water and dried overnight in a petri dish. Once dry, the PVA film was removed from the petri dish and cut into rectangles slightly larger than the PDMS stamps. The PDMS stamps were coated with Matrigel overnight and then placed on top of the PVA films. A glass slide and 50 g weight was added to ensure even pressure distribution and better pattern transfer from the PDMS stamp to the PVA film. After one hour, the PVA film was removed from the PDMS stamp and then brought into conformal contact with the soft substrate for 30 min. Afterward, the substrate was washed to dissolve the PVA film, leaving behind the patterned proteins which were then seeded with cells. The remaining areas of the substrate without ECM proteins were then backfilled with Pluronic F127 (Sigma Aldrich) to prevent cell outgrowth. Cells must be seeded onto the patterns with a density high enough to ensure complete pattern coverage by the cells, and low enough to allow a moderate amount of normal growth of each cell. In these experiments, cells were seeded onto patterns in the range of 2000–3000 cells per mm<sup>2</sup> of ECM area available. This is higher than the 1400–1500 cells/mm<sup>2</sup> in monolayer control samples, even though the ultimate seeding density after attachment is equivalent by a count of nuclei.

### Immunofluorescent Staining

For immunofluorescent staining, cells were cultured in the 2D substrate systems, stained at various time points, and then imaged. Staining began with a PBS wash and then exposure to 4% paraformaldehyde (PFA) (Electron Microscopy Sciences) for 15-min at room temperature. The cells were washed again with PBS and then the cell membranes were permeabilized with 0.1% Triton (Sigma) for 6 min at room temperature. After another PBS wash they were treated for 30 min with a blocking solution consisting of PBS, 2% FBS, 0.1% Triton, 11.2 mg/mL glycine, and 50 mg/mL BSA. The following primary antibodies were left on the samples overnight at 4°C: rabbit anti- $\alpha$ -actinin (1:200 dilution, Abcam) and mouse anti-N-cadherin (1:250 dilution, BD Biosciences). The following day, cells were washed once with PBS for 5 minutes, and treated for 1 hour with blocking solution containing the secondary antibody, Alexa Fluor 488 goat anti-mouse and/or Alexa Fluor 647 goat anti-rabbit (1:1000 dilution). When desired, cells were washed once more, and phalloidin conjugated to tetramethylrhodamine B isothiocyanate (TRITC) (Sigma) was applied at a 50 $\mu$ g/mL concentration to label actin filaments, and/or DAPI was applied at a 1:10,000 dilution to label nuclei. Finally, cells were washed with PBS and transferred to coverslips and mounted with ProLong Gold Antifade (Life Technologies). Samples were imaged on a Nikon A1RSi Confocal Microscope. Widefield images were also acquired with an attached Photometrics CoolSNAP HQ2 camera. Nis-D Elements – Advanced Research v. 3.22 software was used for image acquisition and analysis. Objectives for this microscope included a CFI Plan Apochromat Lambda 40 $\times$  NA 0.95 air objective and a CFI Plan Apochromat Lambda 60 $\times$  NA 1.4 oil immersion objective.

To visualize the transfer of Matrigel from the patterned PDMS stamp to the substrate, the primary antibody rabbit anti-laminin (1:500, Sigma-Aldrich) was incubated with the substrate overnight at 4 °C. The next day the substrate was washed with PBS for 5 min and

then treated for 1 h with blocking solution containing the secondary antibody, Cy3-conjugated mouse anti-rabbit (1:1000 dilution, Abcam).

### Surface Feature Tracking and Calcium Imaging

Contraction and relaxation were quantified using surface feature tracking of stem cell-derived CMs grown on micropatterned lanes. Using the IonOptix myocyte calcium and contractility system (Westwood, MA), with the micropatterned slide containing the stem cell-derived CMs placed in a custom open perfusion chamber, detailed cell shortening data was collected. Additionally, the relationship between the calcium transient and contraction was interrogated by introducing the cells to calcium ratiometric dye Fura-2-AM (Invitrogen). Using the IonOptix calcium imaging system with its HyperSwitch in conjunction with the perfusion chamber placed on the flat stage of an appropriate microscope (Nikon TI-U inverted microscope with 40X fluorescence objective), intracellular calcium release and reuptake and cell shortening were simultaneously recorded.

### Digital Image Correlation

Certainly when observing a collection of CMs, but even within an individual CM, the deformation taking place is not uniform across the field of view. The details of the full-field behavior are often critical to developing a deeper understanding of the biomechanics at play. To measure full-field displacements, we used digital image correlation (DIC) [77]. DIC requires an image having a random high-contrast pattern, which in this case was a phase contrast image of the cells obtained with a Nikon Eclipse Ti microscope with a Plan Fluor 10× NA 0.3 objective and Nikon DS-QiMc camera. In our preliminary analyses, we found the results to be the same with two different free software packages, Ncorr ([https://github.com/justinblaber/ncorr\\_2D\\_matlab](https://github.com/justinblaber/ncorr_2D_matlab)) [78] and Fast Iterative Digital Image Correlation (FIDIC, <https://github.com/FranckLab/FIDIC>) [79]. Results presented here used FIDIC.

### Traction Force Microscopy

In addition to measuring displacements and strains of the cells, we measured forces applied by the cells to the substrate using traction force microscopy (TFM), which computes the traction (force per area) at the interface between the cells and the compliant substrate [70,71,80–83]. This required markers to be embedded in to the substrate of our cell culture platform. It is crucial to achieve a high contrast image of a single plane of particles at the top surface of the substrate. As cell-induced deformations cause particle displacements throughout the substrate, the exact location of the particles must be known for accurate calculations. Often measurements are assumed to be taken from particles at the top surface of the substrate nearest the cells. Unfortunately, this assumption is often incorrect, resulting in an image that shows beads for various positions through the bulk of the substrate. To address this issue, we make PA substrates with particles located at the cell-substrate interface as described above.

The cells and fluorescent particles were imaged with a widefield fluorescent microscope (Nikon Ti-E) with a 10× NA 0.3 objective during an experiment. After the experiment, the cells were released from the substrate with trypsin, which eliminated the tractions applied by the cells to the substrate and allowed the substrate to recover to a stress-free state. A final

image of the particles was acquired in that stress-free state; this final image was used as a reference to compute cell-induced displacements of the particles using DIC. After displacements were measured, the tractions were computed by applying equations of force equilibrium to the substrate [70,71,80–83]. There are multiple different ways to implement the equations; we used an approach based on the Fourier transform [71] with corrections for a substrate of finite thickness [82,83].

## RESULTS AND DISCUSSION

### Cell Culture System

Fig. 2a illustrates a pattern designed to investigate the influence of the pattern aspect ratio on CM morphology. Adult human CMs are rod-shaped with a minimum length-to-width ratio of 7:1 and lengths in the 100  $\mu\text{m}$  range [84]. This characteristic elongated shape of mature CMs produces a highly organized internal cytoskeletal network consisting of myofibrils running parallel to the length of the cell body. In these experiments immature cardiomyocytes differentiated from induced pluripotent stem cells (iPSCs) (19–9-11 cell line, WiCell, Madison, WI) were seeded onto PDMS substrate patterned with Matrigel. By fixing and staining the cells after they have grown in the pattern for a period of time, the internal structure of the cell was explored with immunofluorescence microscopy. As Fig. 2c and d show, the internal sarcomere structure of the iPSC-CMs is significantly more organized along the length of the 11:1 feature compared to the disordered central region of the 1:1 feature after just 6 days of being cultured on the patterns.

Although the aspect ratio of the pattern has interesting implications, we found that the pattern width is more important than aspect ratio in controlling the internal organization of the CMs within a multicellular 2D construct [69]. Fig. 2e shows the cell alignment and sarcomere organization within hESC-CMs (H9 hESC line containing a cardiac troponin T GFP promoter [85]) that have been cultured for 18 days in a 20 $\mu\text{m}$  wide Matrigel lane on a PDMS substrate with Young's modulus of 5 kPa (measured by tensile testing). Rather than the internal disarray observed in CMs grown in unpatterned monolayer conditions, the lane guides the cells to create an organized internal structure similar to that of adult CMs *in vivo*. As micropatterning provides control over these intracellular features, it reduces some of the heterogeneity common in biology, thereby providing a means to design well-controlled, repeatable experiments.

Experiments using varying pattern shapes have shown that patterns with widths less than 100  $\mu\text{m}$  induce an increase in nuclear alignment for CMs on substrates of different stiffnesses. Patterns with widths of  $\sim 40$   $\mu\text{m}$  were found to be ideal for producing connected, highly structured, one-cell-wide lanes of hESC-CMs [69]. Using our culture system, we find that CMs cultured in lanes of various width on PDMS substrates display a similar trend in organization to that found on glass (Fig. 3a-d).

We explored more than seven orders of magnitude in Young's modulus with these techniques, from glass to low stiffness PDMS. Fig. 3e-h illustrates the results seen in stem cell-derived CMs under this range of conditions. The actin stain shows not only elements of the sarcomere organization but also stress fibers, which are least prevalent in cells cultured

on the 5 kPa substrate. Analysis of nuclear alignment with the DAPI stain was used to determine the directionality of the cells populating lanes. The cells were defined as aligned if their major axis was within  $20^\circ$  of the lane direction. As Fig. 3d shows, alignment improves with decreasing lane width and is relatively unaffected by substrate stiffness.

### Contraction Displacement and Calcium Imaging

We simultaneously measured contraction amplitude, and intracellular calcium transients in micropatterned hESC-CMs that were cultured in  $30\ \mu\text{m}$  lanes on 5 kPa PDMS for 6 days prior to testing. A major advantage of using our micropatterned lanes rather than conventional monolayer culture of hESC-CMs is that the micropatterning produced organization in the CMs with most sarcomeres aligned parallel to the long axis of the cell, allowing us to assess contraction along a known direction. Contraction was assessed by surface feature tracking, while calcium transients were assessed by ratiometric Fura2 fluorescence using an IonOptix myocyte calcium and contractility system. During testing, micropatterned hESC-CMs were subjected to electrical pacing at 1Hz by field stimulation, allowing us to gather detailed kinetic data of contraction (CM shortening), relaxation (CM re-lengthening), calcium release, and calcium sequestration both in the absence and presence of  $1\ \mu\text{M}$  isoproterenol (Fig. 4).  $\beta$ -adrenergic stimulation with isoproterenol treatment of the micropatterned hESC-CMs increased both contraction amplitude and calcium transient amplitudes while also increasing the rates of contraction, relaxation, calcium release, and sequestration (Fig. 4). This combination of outcomes indicates a mature functional status of the CMs in these culture conditions [86,50].

### Contractile Force

To quantify the forces of CM contraction, we used traction force microscopy. Polyacrylamide substrates embedded with fluorescent particles at the top were generated using the 2-step protocol described in the methods. Matrigel was micropatterned onto the polyacrylamide substrate, and cells were seeded on top (in this case, H9 hESC-CMs). Representative tractions applied by CMs to the substrate are shown in Fig. 5 for unconnected lanes which do not contract in synchrony with neighboring lanes.

Whereas Fig. 5 shows traction at the interface between the cells and substrate, the contraction within a layer or lane of cells can be quantified as well, using an additional analysis that applies the principle of force equilibrium to the layer of cells [83,87–89]. The analysis begins with a micropatterned lane of cells having width  $w$  (Fig. 6a). Cell contraction generates an intracellular force at location  $x$ ,  $F(x)$ , which is balanced by a nearby force in the opposite direction  $F(x+dx)$  and the traction  $T(x)$  having units of force per area. Force equilibrium gives  $F(x) = F(x+dx) + T(x)wdx$ , which can be rewritten  $dF/dx + wT = 0$ . From this relationship, integrating the traction gives the contractile force scaled by lane width,

$$f \equiv F/w = \int T(x)dx.$$

Integration begins at the endpoint of a cell lane, where force is known to be zero. As the force  $f$  is a direct measure of contractile activity within the cell, it is a more physiologically relevant metric than traction. Thus, the micropatterning both produces organized sarcomeres (Fig. 3) and enables mechanics-based quantification of CM contraction. The data show that traction and force increase and decrease periodically over time with repeating magnitude (Fig. 6c-e), indicating collective and consistent cell beating.

### CM Contraction on Patterned Substrates

We designed more complex patterns for multicellular studies to produce alignment of the internal myofibrils while providing for cell-cell connections that span a large area. The resulting connected mesh of CMs displayed synchronized spontaneous contractions. We focus here on a  $15^\circ$  chevron pattern with representative data shown in Fig. 7. In this data set, CMs were cultured for 16 days in the pattern prior to imaging. Bright field images were captured over the course of several spontaneous contractions (representative images are shown in Fig. 7a, g, and j). Displacements were computed by correlating images in the relaxed and contracted states. In contrast to an unpatterned monolayer, the contraction is anisotropic and aligned primarily in the vertical direction (Fig. 7c), which matches the orientation of the chevron patterns and corresponds to the general alignment of the cells in the primary direction of the chevron patterns at  $\pm 15^\circ$  from vertical. Minimal displacements occur in the horizontal direction (Fig. 7b).

To further quantify the contraction, the strain tensor is computed by taking the gradient of the displacement data; subsequently, the principal strains, defined as the eigenvalues of the strain tensor, are computed. Here, we show the sum and difference of the principal strains (Fig. 7e-f), which correspond to changes in local area and shape, respectively (see supplemental data for video). In Fig. 7e, blue (negative) corresponds to a decrease in area, indicating contraction in the plane of the sample. Given the short time between the images, constant volume of the cells can be assumed such that the out of plane strain is  $-(e_1 + e_2)$ . In Fig. 7f, the magnitude of the shape change is illustrated by the difference in principal strains,  $e_1 - e_2$ .

Because the soft substrate is attached to a rigid glass slide below, global macroscopic area and shape changes are not possible. The chevron mesh pattern not only promotes sarcomere alignment through the pattern edges, but also provides the cells with a template for creating a globally self-accommodating structure that permits large coordinated contractions along the chevron orientations of  $\pm 15^\circ$ . The large contractile strains typically reported for CMs in tissue can be observed at the local level and sustained in the chevron mesh pattern of this 2D platform. In contrast when a  $90^\circ$  bridge pattern design was used to connect lanes, CMs were torn apart at junctions by the contractions (data not shown). The primary axis of contraction in the plane of the sample is indicated by the lines in Fig. 7d, which gives the orientations of the second principal strain.

Data in Fig. 7a-f are for CMs on a compliant substrate having Young's modulus of 5 kPa. Results for cells on substrates of Young's modulus 10 kPa and 50 kPa are shown in Fig. 7g-i and Fig. 7j-l, respectively. The data show that contraction decreases nonlinearly as substrate stiffness increases. For example, the root-mean-squares of the differences in principal

strains,  $e_1 - e_2$ , are 10.9%, 7.7%, and 4.0% for cells on 5 kPa, 10 kPa, and 50 kPa substrates, respectively. The strains diminish with increasing substrate stiffness, because the cardiomyocytes are increasingly constrained in their ability to contract by the underlying substrate. Interestingly, the decrease in contractile strains are not proportional to the increase in substrate stiffness, indicating that the CMs produce different contractile force on substrates of different stiffness.

## CONCLUSION

Here we have shown a 2D cell culture platform designed to produce organized collections of cardiomyocytes *in vitro*, thereby enhancing key maturity markers. We quantified cell alignment and contraction using fluorescent dyes, image tracking, digital image correlation, and traction force microscopy. We showed that lanes with widths less than 100  $\mu\text{m}$  induce an increase in nuclear alignment for CMs on substrates of Young's modulus spanning seven orders of magnitude. Calcium transients and contraction amplitude were measured for CMs in lanes with exposure to 1Hz electrical pacing, with and without the presence of isoproterenol. The mature functional status of the CMs exposed to isoproterenol were illustrated by the increases in contraction amplitude, calcium transient amplitudes, rates of contraction, relaxation, calcium release, and sequestration. We also presented data for a connected pattern design (15° chevron) that maintains the feature width restrictions of lanes while producing cell-cell connections that span a large area, recapitulating adult-like cardiomyocyte organization within heart tissue. The connected CMs in the chevron design displayed synchronized spontaneous contractions, with contractile strains of the magnitude typically reported for CMs *in vivo*. Furthermore, we show that contraction decreases nonlinearly as substrate stiffness increases over range of 5 kPa to 50 kPa which spans the range of moduli reported for healthy and diseased heart tissue. Taken together, these findings demonstrate the value of this platform to enhance maturation in stem cell derived cardiomyocytes and study cardiomyocyte behavior. As mechanical contraction is one of the most critical functions of CMs *in vivo*, we expect the methods described here will be useful for drug development, toxicity testing, and diagnostic assays.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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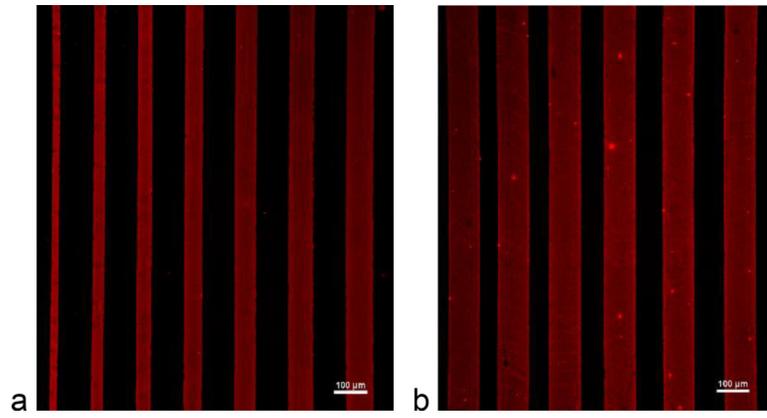
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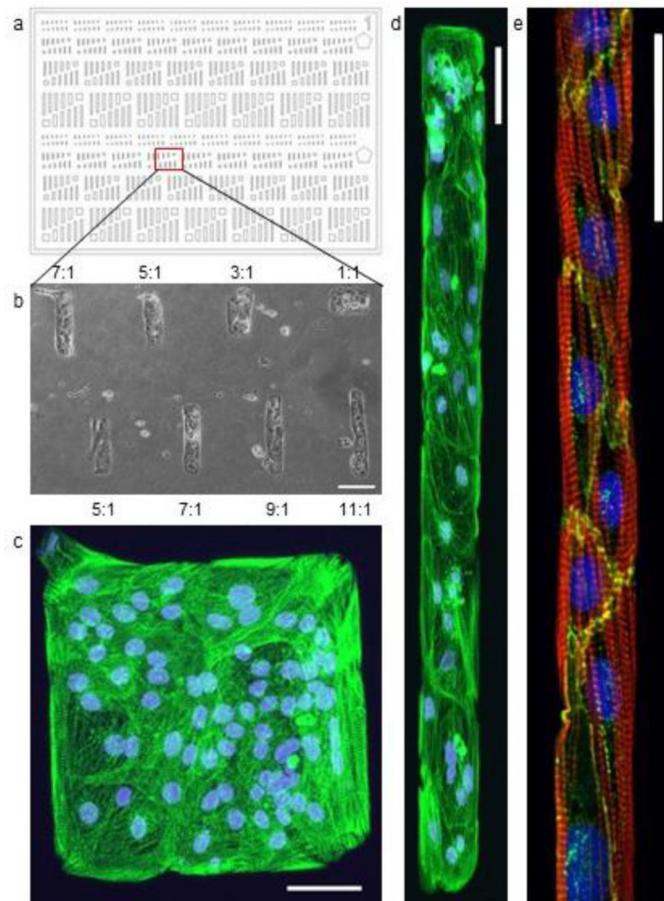
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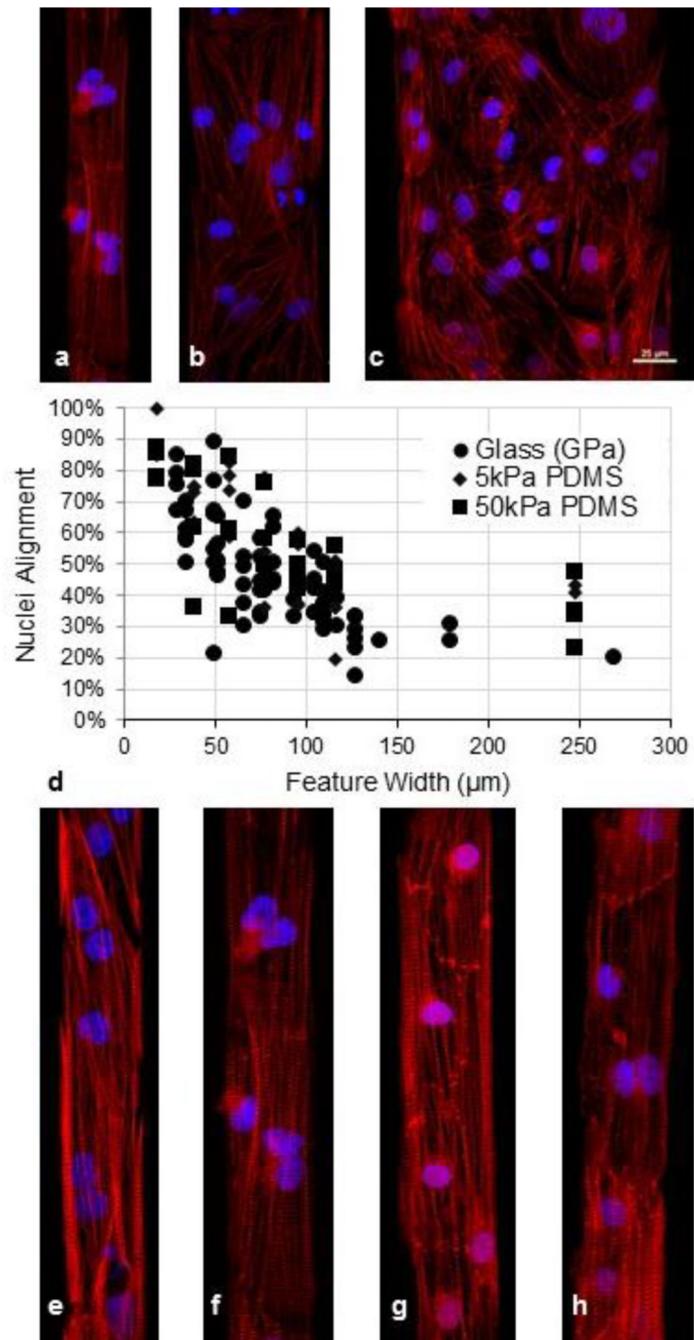


**Fig. 1.** Immunofluorescence images of different patterns of Matrigel (red), visualized with an anti-laminin antibody, on (a) glass and (b) 5 kPa PDMS substrates. Scale bar = 100  $\mu\text{m}$

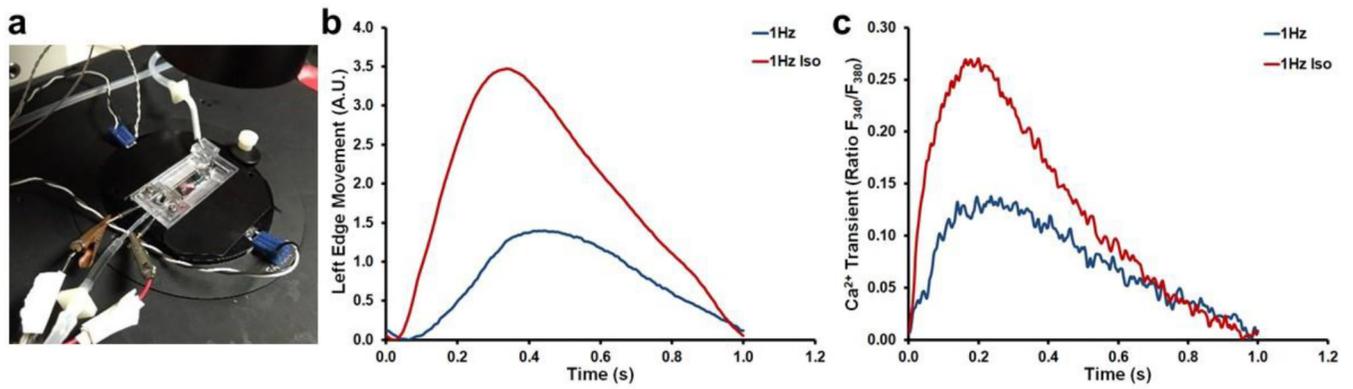


**Fig. 2.**

(a) Pattern consisting of rectangles with a range of aspect ratios between 1:1 and 11:1. (b) A bright field image of 19–9–11 stem cell derived cardiomyocytes cultured on a portion of the pattern created with a Matrigel coated stamp using this pattern design. Aspect ratio is noted for each feature detailed above and below the image. Scale bar: 100  $\mu\text{m}$ . (c, d) Fluorescent microscopy image of a 1:1 and a 9:1 feature. The nuclei are stained with DAPI (blue) and the protein  $\alpha$ -actinin (green) highlights the sarcomere structure within the cells. (a-d) Scale bar: 50  $\mu\text{m}$ . (e) Micropatterned lane with hESC-CMs cultured in Matrigel lanes on a 5kPa stiffness PDMS substrate for 18 days. Cell alignment and sarcomere organization shown in lanes of 20 $\mu\text{m}$  width. The nucleus is stained with DAPI (blue), the protein actin (red) highlights the internal organization of the cell, and the protein N-Cadherin (green) highlights the cell-cell interfaces. Scale bar 50  $\mu\text{m}$ .

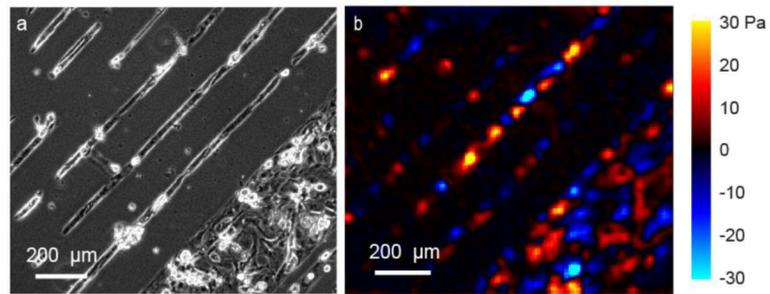


**Fig. 3.** (a-c) Sarcomere organization in hESC-CMs seeded on different lane widths of Matrigel on 1.72 MPa PDMS, including (a) 37  $\mu\text{m}$ , (b) 75  $\mu\text{m}$ , and (c) 150  $\mu\text{m}$ . Red = actin, blue = DAPI. Scale bar is 25  $\mu\text{m}$ . (d) Nuclei alignment for hESC-CMs grown on a range of lane widths spanning 30 to 270  $\mu\text{m}$  using glass and PDMS substrates. (e-f) hESC-CMs grown on substrates of different Young's moduli, each having a lane width of 37  $\mu\text{m}$ : (e) Glass, (f) 1.72 MPa PDMS, (g) 50 kPa PDMS, and (h) 5 kPa PDMS. Red = actin, blue = DAPI



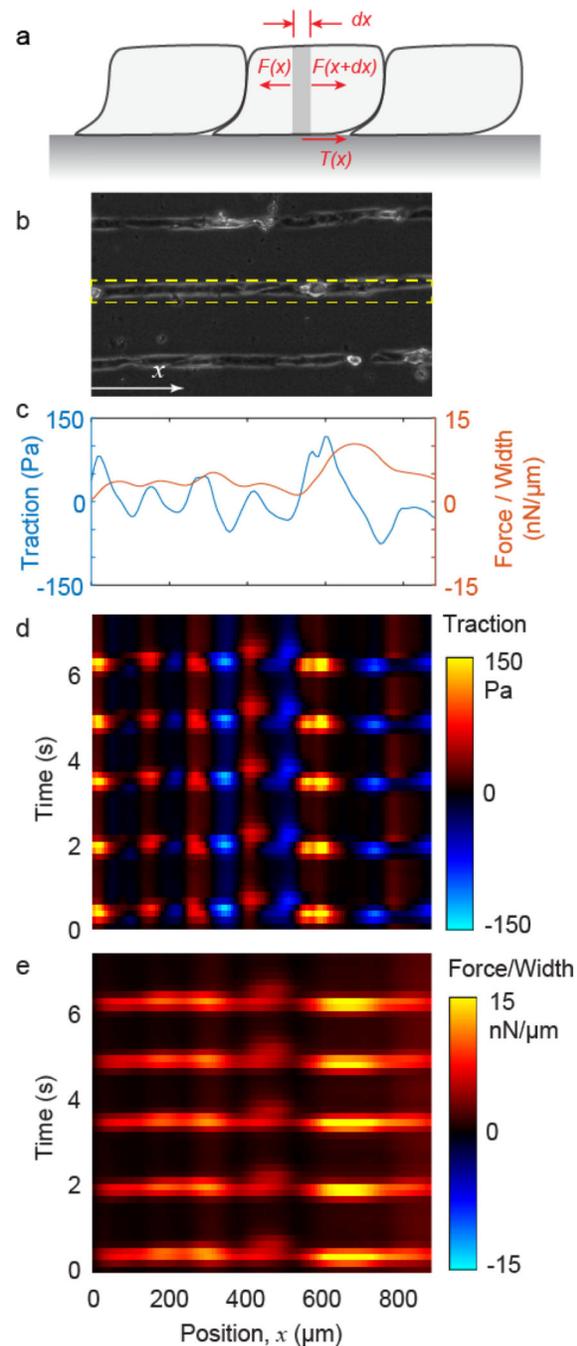
**Fig. 4.**

(a) IonOptix myocyte calcium and contractility system with custom chamber.  $\beta$ -adrenergic stimulation of micropatterned hESC-CMs. hESC-CM (b) contraction amplitude and (c) rate of  $\text{Ca}^{2+}$  transient when paced under different conditions in the presence and absence of isoproterenol. Fura-2-AM ratiometric fluorescent dye (Abcam). Exposure to isoproterenol at  $1 \mu\text{M}$  increased  $\text{Ca}^{2+}$ -transient amplitude, contraction amplitude, the rate of contraction, and the rate of relaxation with 1 Hz pacing.

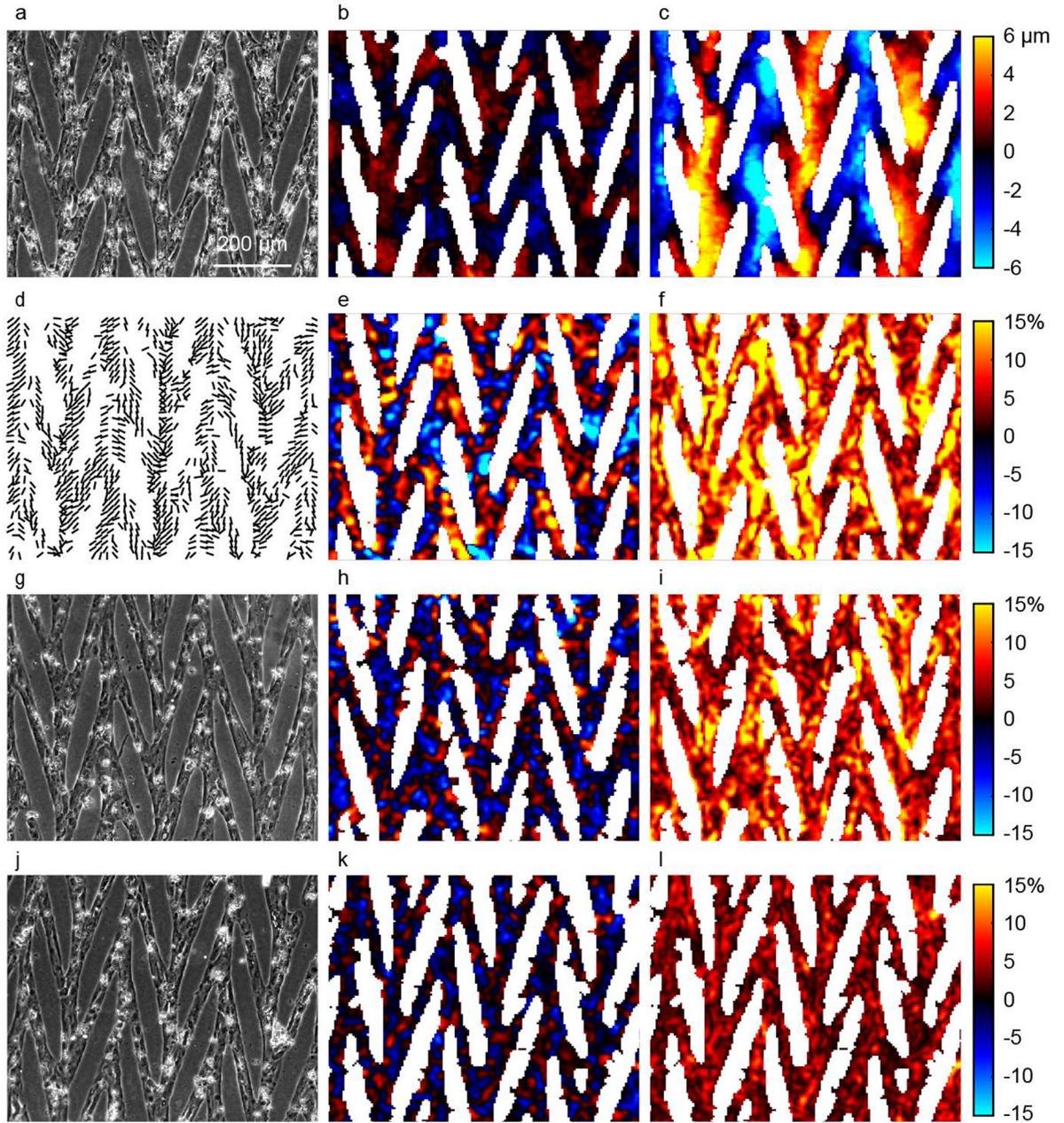


**Fig. 5.**

TFM analysis of micropatterned hESC-CMs seeded within lanes on a 5 kPa polyacrylamide substrate at one moment in time. (a) Phase contrast image of CMs. Narrow lanes are 30  $\mu\text{m}$  in width; wide lane is 350  $\mu\text{m}$ . (b) Traction map showing forces applied by cells to the substrate in direction of lanes. Positive (warm colors) is up and right; negative (cool colors) is down and left



**Fig. 6.** CM contraction over time. (a) Forces within a lane of cells. (b) Cardiomyocytes patterned into lanes on a polyacrylamide substrate. Traction is computed for the lane shown by the dashed rectangle. (c) Traction (averaged over lane width) and contractile force (normalized by lane width) is plotted against position  $x$  for a single point in time. (d, e) Traction and force are calculated for all time points and plotted over position and time as a kymograph. The periodic bands of bright signal in force indicate periodic cell contraction, i.e., beating.



**Fig. 7.** Displacements and strains of a CM array computed with DIC. (a,g,j) Bright field images of CMs micropatterned into a chevron pattern. (b, c) Displacements in horizontal (b) and vertical (c) directions,  $u_h$  and  $u_v$ . Red (positive) indicates rightward and upward. Color scale bars for panels (b) and (c) are the same. (d) Lines show orientation of second principal strain, indicating the primary axis of contraction. Lines are shown only in locations for which the magnitude of second principal strain is greater than 2%. (e,h,k) Sum of principal strains,  $e_1 + e_2$ , gives a measure of relative change in area at each location within the CM array. (f,i,l) Difference of principal strains,  $e_1 - e_2$ , quantifies a relative magnitude of shape change at each location within the CM array. (a-f) CMs on a compliant substrate with

Young's modulus of 5 kPa, in comparison to substrates of 10kPa (g-h) and 50 kPa (j-l).  
Color scale bars for panels (e-f, h-i, and k-l) are the same.

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