

ELL-MET

Investigating the Impact of Increased Afterload on Hypertrophic Cardiomyopathy

Maya Roseboro^{1,2}, Miranda Wang^{2,3,4}, Xining Gao^{2,3,4}, Christopher Chen^{2,3}

Isaac Bear Early College High School, 630 MacMillan Avenue, Wilmington, NC 28403, USA¹; Department of Biomedical Engineering, Boston, MA 02215²; Wyss Institute for Biologically Inspired Engineering, Boston, MA 02215²; Wyss Institute for Biologically Inspired Engineering, Boston University, Boston, MA 02215²; Wyss Institute for Biologically Inspired Engineering, Boston University, Boston, MA 02215²; Wyss Institute for Biologically Inspired Engineering, Boston University, Boston Univers and Science, Massachusetts Institute of Technology, Cambridge, MA 02139, USA⁴

Background

Hypertrophic cardiomyopathy (HCM) is the most common inherited heart condition¹. HCM is caused by mutations in genes encoding sarcomere proteins and leads to pathologic left ventricular hypertrophy and fibrosis. At the cell level, these mutations are linked to increased contraction, increased energy usage, and decreased relaxation³. The mechanisms linking sarcomere mutations to the extensive myocardial remodeling observed in HCM are not well understood. This gap in knowledge has limited the discovery of druggable targets for therapeutic development. HCM bears striking resemblance to pressure-overload hypertrophy induced by hypertension, and the sarcomere modulates activity of several mechanosensitive signaling pathways. Therefore, we hypothesize that mechanical stress may play an important role in the development of symptomatic HCM. The goal of our research is to determine the impact of biomechanical loading on HCM progression.

Increased Afterload Exacerbates Hypercontractility and in HCM Microtissues





Fig 1. A. Explants and H&E staining of section from myocardium from a heart from a healthy donor and from a patient with HCM². B. Sarcomere proteins, of which ~60% of HCM patients have a mutation in either β -myosin heavy chain (*MYH7*) or myosin binding protein C (*MYBPC3*)². C. Clinical outcomes of HCM patients who suffer from ventricular hypertrophy.

HCM hiPSC-CMs Exhibit Lower Tolerance for Substrate Stiffness



Methods

Myosin Activity Alone Does Not Drive Fibrotic Response to Increased Afterload







Fig. 5. A. Representative images and contractile stress of healthy cardiac microtissues cultured on soft or stiff pillars, or cultured on stiff pillars with OM. B. Maximum intensity z-projections of microtissues stained for collagen type I and αSMA. Collagen and αSMA intensity were quantified and compared between conditions.

Conclusions

In this work, we built on our previously developed model of HCM to show that HCM cardiomyocytes are hypersensitive to changes in afterload. We found that microtissues formed from hiPSC-CMs housing an HCM-causing mutation in MYBPC3 exhibit hypercontractility when exposed to increased afterload; however, there was no observed statistical difference between passive tension generated between HCM and healthy cell conditions. We found through single-cell traction force microscopy that HCM cells appear less tolerant to seeding on high substrate stiffnesses, showing peak traction force at a lower substrate stiffness than healthy cells. However, further experimentation will be needed to validate this conclusion. Based on our finding that increased afterload exacerbates hypercontractility in HCM microtissues, and building from previous data showing that when exposed to high afterloads HCM microtissues exhibit elevated levels of collagen type I and α SMA, we sought to understand whether increased myosin activity drives the HCM fibrotic response in high mechanical stress conditions. We found that increasing myosin activity with omecamtiv mecarbil augmented contractile stress but did not elevate markers of fibrotic activation in our tissues. Therefore, we posit that increased myosin activity alone is not sufficient to provoke fibrotic activation. Thus, a mechanism other than myosin activity, such as metabolic state, may contribute to stress observed in HCM cells. We hope that this work can provide insight into the impact of mechanical loading on HCM and contribute to the development of new HCM druggable targets.

Fig 2. A. using hiPSC-CMs with a mutation in *MYBPC3* or a healthy WT control and healthy vCFs, we created 3D cardiac microtissues to study HCM *in vitro*. The platform can be tuned to modulate applied afterload via casting with PDMS of different base to crosslinker ratios. All tissues were cultured for 7 days before analysis. B. Healthy or mutant hiPSC-CMs were seeded onto polyacrylamide gels containing fluorescent beads with elastic moduli ranging from 1-22kPa. Videos of cell contraction were collected and bead displacement was used to calculate single cell contraction stress.

References and Acknowledgements

1. Marian & Braunwald. Circulation Research 2017 2. Teekakirikul et al. Journal of Cell Biology 2012 3. Seidman & Seidman. Circulation Research 2011 Figures created with BioRender.com

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