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WASHINGTON UNIVERSITY IN ST. LOUIS

School of Engineering & Applied Science

Department of Biomedical Engineering

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Elucidation of Emergent Regional Mechanisms of Heart Muscle Dysfunction in the Mouse Model of Duchenne Muscular Dystrophy

by

Ya-Jian Cheng

A dissertation presented to the Graduate School of Arts and Sciences of Washington University in partial fulfillment of the requirements for the degree of Doctor of Philosophy

May 2014

Saint Louis, Missouri

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Table of Contents

List of Figur	es	v
List of Table	s	ix
Acknowledg	ements	x
Abstract of the	he Dissertation	xii
Chapter 1: In	troduction	1
1.1. Du	chenne Muscular Dystrophy	1
1.1.1.	The Gene	1
1.1.2.	The Protein	3
1.1.3.	The Disease Phenotype	5
1.1.4.	Treatment Options	8
1.1.5.	The <i>mdx</i> mouse model	9
1.1.6.	Calcium Overload Hypothesis	9
1.1.7.	Hypothesis	
Chapter 2: M	IRI for Cardiac Sheet Function	
2.1. Ca	rdiac muscle structure	15
2.1.1.	Fiber	
2.1.2.	Sheet	
2.1.3.	Nomenclature of the heart segments	20
2.2. Di	ffusion Tensor MRI	21
2.2.1.	Background	21
2.2.2.	Restricted Diffusion	22
2.1. Me	ethod: DTI for <i>mdx</i> mice cardiac structure	23
2.1.1.	Langendorff perfusion for DTI	25
2.1.2.	Perfusion compatible MR Coil	
2.1.3.	Imaging parameters	
2.1.4.	Post processing, sheet angle calculation (β)	
2.1.1.	Statistical methods	
2.1.2.	Histology	
2.2. Re	sults: Sheet function in 16-month old <i>mdx</i> mouse hearts	
2.2.1.	Morphological observation of β with histology and PAM	

2.2	.2.	16 month old <i>mdx</i> hearts manifest reversible regional diastolic sheet dysfunction	32
2.2	.3.	16 month old <i>mdx</i> and WT hearts exhibit similar fiber organization and function	36
2.2	.4.	3 month old <i>mdx</i> mice exhibit smaller sheet dysfunction, still reversible by low calciur	n
		perfusion	37
2.3.	Dis	cussion of β	38
Chapter	3: Ar	nisotropy information in DTI	40
3.1.	Intr	oduction	40
3.2.	Met	thod: DTI Data Analysis for FA and mode	42
3.3.	Res	ult	42
3.3	.1.	mdx and WT hearts exhibit lower FA & higher mode	42
3.4.	Dis	cussion – abstract shape information	44
Chapter	4: Op	ptical mapping for calcium transients in <i>ex vivo</i> viable <i>mdx</i> hearts	46
4.1.	Ack	nowledgements to the contribution of Dr. Efimov's lab	46
4.2.	Wh	y use optical mapping for calcium?	46
4.3.	Met	thod	47
4.3	.1.	Isolated heart preparation and calcium dye loading	47
4.3	.2.	Optical imaging system	48
4.3	.3.	Optical mapping protocols	49
4.3	.4.	Optical mapping data processing	49
4.4.	Opt	ical Mapping Results	50
4.4	.1.	Optical imaging reveals longer calcium decay time (T50) in 16 month old <i>mdx</i> mice	50
4.4	.2.	3 month old <i>mdx</i> mice exhibit normal calcium transients	51
4.5.	Dis	cussion of regional abnormality at base	52
4.6.	Dis	cussion of the perfusion condition in DTI and optic mapping	53
Chapter	5: At	tophagy and Beneficial Effects of Rapamycin-nanoparticle	55
5.1.	Ack	xnowledgements	55
5.2.	Abs	stract	55
5.3.	Intr	oduction	56
5.4.	Met	thods	58
5.4	.1.	Reagents	58
5.4	.2.	Nanoparticle formulation	58
5.4	.3.	Animal studies	59

5.4.4.	Histology	61
5.5. Res	ults	65
5.5.1.	Rapid enhancement of global physical performance with RNP	65
5.5.2.	Rapid enhancement of cardiac performance with RPN in aged mice	69
5.5.3.	Nanoparticle tissue distribution	69
5.5.4.	Effects on skeletal and cardiac mTOR and autophagic processes	70
5.5.5.	Tissue structural responses	74
5.6. Dis	cussion	75
Chapter 6: Co	onclusion	
6.1. Sur	nmary of major findings	
6.2. Fut	ure works:	83
6.2.1.	Intervention for calcium mishandling and autophagy in DMD	83
6.3. Rer	narks on dystrophin function	84
References		
Appendices		
Principle o	f MRI	
Physics .		
Imaging		
Supplemen	tal tables	
Speculation	n: Might low gravity ameliorate dystrophy?	
C.V. of Ya-Ji	an Cheng	

List of Figures

Figure 7. Photoacoustic microscopic (PAM) image of a unfixed, unstained, and un-dehydrated myocardium, showing sheet structure of a mouse left ventricle. (a), PAM image of the left ventricle free wall (FW) marked in (c) acquired at 532-nm wavelength. (b), Close-up PAM image. The boundaries of the branching sheets are apparent. Red dashed line: long axis. Yellow dashed-dot line: sheet boundary. (c), Photograph of the heart specimen. LV, left ventricle; RV, right ventricle. (Zhang, Cheng et al. 2012). ... 19

Figure 8. Left ventricle (LV) is rounded in shape in short axis view. Right ventricle (RV) has a semilunar shape. Aorta and vena cava are closer to the heart *base* than the *apex*. Apex is pointing to the left in normal human chest. Heart base is physically closer to the head and has larger diameter than apex.20

Figure 12. A & B, representative short-axis view of DTI determined sheet angle (β) in a basal slice of the same WT heart that was sequentially arrested in diastole (A) and systole (B). The posterior-lateral wall of LV possessed two populations of β , exhibiting positive (red) and negative (blue) values. The transmural portion of positive β is increased in systole (compare the portions of red-colored pixels under the dashed lines in A & B). The dotted line in B indicates a 3 population β transition that appears only in systole. C, Photoacoustic microscopy of long-axis view of a freshly half-split WT mouse heart confirmed two populations of β in LV wall. The insert showed H&E stained long-axis WT mouse heart slice with the corresponding region of photoacoustic image squared. LV: left ventricle. RV: right ventricle. PA: papillary muscle.

Figure 13. DTI-determined diastolic and systolic $|\beta|$ in short-axis slices. A, *mdx* hearts exhibited abnormally lower diastolic $|\beta|$ that was recovered to normal value after reducing [Ca²⁺] in the perfusate. For hearts perfusion-arrested in diastole with normal calcium solution, *mdx* hearts (*mdx*-NC) exhibited lower $|\beta|$ than did WT (WT-NC) hearts with the maximal ~10 degree difference occurred at the basal region. In contrast, *mdx* hearts perfused with the low calcium solution (*mdx*-LC) exhibited normal

Figure 18. DTI-determined FA and mode in diastole and systole of WT and mdx mouse heart. $N = 3$	5 for
ach group. *, <i>p</i> < 0.05	44

Figure 21. Rapamycin-loaded nanoparticle treatment improves strength in *mdx* mice......67

Figure 22. Aged <i>mdx</i> mice show muted responses in grip strength after treatment with RNP, but marked improvement in cardiac function commensurate with wild-type animals
Figure 23. Nanoparticles and rapamycin are delivered to cardiac muscle in <i>mdx</i> animals70
Figure 24. <i>mdx</i> animals exhibit a defect in autophagy that is restored by nanoparticle treatment72
Figure 25. Rapamycin nanoparticles alter autophagic biomarkers in heart and skeletal muscle of <i>mdx</i> mice similar to that after steroid therapy74
Figure 26. Intravenously injected nanoparticles do not affect short-term muscle cell destruction or fibrosis
Figure 27. mdx dystrophy pathway proposed by Deconinck et al. (Deconinck and Dan 2007)90
Figure 28. Hypothesized <i>mdx</i> dystrophy pathway

List of Tables

Table 1. Example of sizes of genes and proteins in the human genome	2
Table 2. Post hoc multiple comparisons for diastolic $ \beta $ of 16 month old mdx and WT mice	109
Table 3. Post hoc multiple comparisons for diastolic $ \beta $ of 3 month old mdx and WT mice show significance.	ing no 109

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ABSTRACT OF THE DISSERTATION

Elucidation of Emergent Regional Mechanisms of Heart Muscle Dysfunction in the Mouse

Model of Duchenne Muscular Dystrophy

By

Ya-Jian Cheng

Doctor of Philosophy in Biomedical Engineering

Washington University in St. Louis, 2014

Professor Samuel Wickline, Chair

Cardiac dysfunction is a primary cause of mortality in Duchenne Muscular Dystrophy (DMD), potentially related to elevated cytosolic calcium. However, the regional versus global functional consequences of cellular calcium mishandling have not been defined in the whole heart. Here, we elucidate, for the first time, loci- and age-dependencies between calcium mishandling and myocardial sheet function as a manifestation of dystrophin-deficient cardiomyopathy. We also map calcium transients to illustrate the regional dependence of ion flux disturbances in the dystrophin-deficient (mdx) mouse heart. Furthermore, we elucidate abnormalities in autophagic processes that can be corrected with nanoparticle therapeutics delivering rapamycin to heart tissues to improve ventricular function in affected older mice with incipient cardiomyopathy. We conclude that the rapid reversibility of functional defects by reducing cytosolic calcium or by impacting impaired autophagy points to the significance of regional mechanical factors in the progression of the disease.

Chapter 1: Introduction

1.1. Duchenne Muscular Dystrophy

1.1.1. The Gene

Duchenne muscular dystrophy (DMD) is a fatal X-linked genetic disease that affects 1:3500 of newborn boys who manifest unremitting deterioration of all muscle groups over time. The symptoms appear at the age of 2–6 years. By the age of 10, untreated patients are usually wheelchair-bound, and in their 20s they typically succumb (Fairclough, Wood *et al.* 2013).

The cause of DMD is the near or complete absence of dystrophin protein. The dytrophin gene is located on chromosome X and is 2.3 million base pair (bp) long (Altamirano, Lopez *et al.* 2012), occupying ~0.08% of the human genome of 2.9 billion bp (Alloatti, Gallo *et al.* 1995). It is the longest known human gene (Table 1). The 79 dystrophin exons are spliced into a 14 kbp mRNA¹, which requires 16 hours for processing in cultured myogenic cells (Altamirano, Lopez *et al.* 2012). Although gene therapy has been suggested as a possible cure for the disease, the large size of dystrophin gene exceeds the packaging capacity of currently available viral vectors. Although micro-dystrophin (~30% of original dystrophin cDNA sequence) and Dp71 isoform (~16% length or full dystrophin) were used to tackle the transfection capacity limitation, the effectiveness for this particularly large gene is yet not proven (Nicolas-Metral, Raddatz *et al.* 2001; Hecht, Adams *et al.* 2011; Spurney 2011). Moreover, immune reactions against the transfected gene challenge the clinical application of gene therapy (Berry, Andruszkiewicz et al. 2013).

¹ A gene contains exon(s) and usually intron(s). Exons are encoded into mRNA while introns are removed by RNA splicing.

	Histone H1a	Dystrophin	Titin
DNA length	781 bp	2 M bp	293 k bp ²
mRNA length	781 bp	14 k bp	80 k bp ²
Protein M.W.	22 k Da^3	427 k Da ⁴	3 M Da^5

Table 1. Example of sizes of genes and proteins in the human genome. Histone H1a is small and contains no intron. In the human genome, dystrophin is the largest gene, most of which comprises introns. The titin gene produces the largest single protein in human genome.

Interestingly, both titin, the largest mammalian protein (Han, Grounds *et al.* 2006), and dystrophin, the longest gene in humans, exert a significant mechanical role in the heart. Titin is considered a mechanical spring responsible for diastolic recoil. The size might reflect the mechanical load it bears. Dystrophin is an adaptor between extracellular matrix and cytoskeleton. Its gene length reflects the fact that it produces many isoforms, which require alternative splicing and regulatory functioning of introns. When promoter defects or nonsense mutations, duplications, and deletions lead to *complete absence* of dystrophin, the resultant disease is DMD. When in-frame deletions or other mutations lead to *low expression or low function* of dystrophin protein, the resultant disease is Becker muscular dystrophy (BMD), a milder form of DMD (Bowles, McPhee *et al.* 2012). Figure 1 illustrates the spectrum of mutations and phenotypes for the dystrophin gene.

 ² Franco-Obregon, A. and J. B. Lansman (2002). "Changes in mechanosensitive channel gating following mechanical stimulation in skeletal muscle myotubes from the mdx mouse." <u>The Journal of physiology</u> 539(Pt 2): 391-407.
³ <u>http://www.ncbi.nlm.nih.gov/protein/NP_005316.1</u>

⁴ Monaco, A.P., Neve, R.L., Colletti-Feener, C., Bertelson, C.J., Kurnit, D.M., and Kunkel, L.M. (1986). Isolation of candidate cDNAs for portions of the Duchenne muscular dystrophy gene. Nature 323:646-650. http://www.dmd.nl/isoforms.html

⁵ "ProtParam for human titin". ExPASy Proteomics Server. Swiss Institute of Bioinformatics. http://web.expasy.org/cgi-bin/protparam/protparam1?Q8WZ42@1-34350@



Figure 1. Schematic representation of dystrophin gene taken from He et al (He, Yi et al. 2013) (A), protein isoforms (B), dystrophin mutational spectrum (C). (A) The dystrophin gene is depicted as a 2500 kb arrow pointing from the centromere to the telomere, promoters are shown as rightangled arrows coming off the main arrow with a letter depicting promoter name and dystrophin protein (Dp) in kDa as a number. B, brain promoter (Nudelet al., M. muscle promoter 1989); (Klamut et al., 1990); P, purkinje promoter (Gorecki et al., 1992); R, retina promoter (D'Souza et al., 1995); B=K, brain and kidney (Lidov et al., 1995); S, Schwann cell (Byers et al., 1993); G, general giving rise to both Dp71 (Bar et al., 1990) and Dp45

(Tinsley *et al.*, 1993). (B) Key followed by schematic representations of the various translated products from the dystrophin gene. Key: ABS, actin-binding sites 1 ± 3 , 4 and 5, WW, WW domain; EF, EF hand calcium-binding domain; ZZ, ZZ zinc finger domain; DCC, predicted dimeric coiled coil. Protein schematics are shown aligned under the key with different shaped regions depicting the different domains. Alternatively spliced exons are shown in black and as a square-bracketed 18 below the C terminus of full-length dystrophin, indicating the 18 alternative splice variants in this region (see Fabbrizio *et al.*, 1994, for review).

1.1.2. The Protein

Dystrophin is expressed in muscle, kidney, brain and other human tissues (Chelly, Kaplan et al. 1988). In muscle, the dystrophin protein is a physical adapter between extracellular matrix (ECM) and intracellular cytoskeleton (Figure 2). Based on stoichiometry, Ghate et al have shown that one dystrophin binds directly to 24 actin monomers (Ghate, Baker *et al.* 2011). Dystrophin binds to ECM indirectly through dystroglycans and sarcroglycans. Mutations on

dystroglycans and sarcroglycans both lead to distinct forms of muscular dystrophies (Medvedev, Murro *et al.* 2011). Baron et al have shown that the sarcolemma of the dystrophin-deficient (mdx) mouse muscles is four time less stiff than that of control mice (Baron, Hecht *et al.* 2011), indicating a mechanical role for dystrophin.

The absence of dystrophin results in alteration in membrane structure, which is related to the dystrophin-associated proteins. The dystrophin-associated protein complex is part of a riblike lattice structure called the costamere.(Olamaei, Cheriet *et al.* 2010). These anchor the cytoskeleton to the ECM (Rybakova, Patel *et al.* 2000), and distribute contractile forces generated in the muscle fiber laterally through the sarcolemma to the basal lamina in ECM and thereby maintain uniform sarcomere length along the fiber (Danowski, Imanaka-Yoshida *et al.* 1992). Absence of dystrophin protein leads to complete loss of the dystrophin-associated protein complex and disruption of costamere lattice, resulting in excessive membrane fragility (Rybakova, Patel *et al.* 2000). In addition to these mechanical functions, signaling functions of dystrophin have been theorized as discussed in Chapter 4.



Figure 2. The dystrophin protein in muscle linking the internal cytoskeleton to the extracellular matrix. From (Morelli, Runge *et al.* 2011)

1.1.3. The Disease Phenotype

DMD patients manifest skeletal muscle weakness and fat/fibrosis infiltration in their first decade and succumb to *respiratory* or *cardiac failure* before the third decade (He, Yi *et al.* 2013). DMD skeletal muscle biopsy characteristically reveals necrotic and degenerating muscle fibers, surrounded by macrophages and CD4+ lymphocytes. Smaller immature centrally nucleated fibers reflect ongoing muscle cell regeneration (Deconinck and Dan 2007). Raised levels of muscle enzymes (CK) in DMD patients offer evidence of sarcolemma fragility that occurs after exercise (Stepkowski 2003; O'Brien, Stinson et al. 2008). Subsequent skeletal muscle necrosis and fibrosis result in loss of contractility, eventually in respiratory failure, and ultimately death from muscle weakness. Monitoring of the disease progression with MRI (Magnetic Resonance Imaging) can define regional heterogeneity among various muscle groups, and assess treatment efficacy for limiting fatty infiltration and fibrosis (Kim, Laor *et al.* 2010; Kim, Laor *et al.* 2010).

Histological analysis of DMD hearts demonstrates replacement of myocardium by connective tissue or fat (Schmidt, Burmeister *et al.* 2003). Fibrosis and diastolic dysfunction precede overt dilated cardiomyopathy (DCM), which is subsequently complicated by heart failure and arrhythmia (Judge, Kass *et al.* 2011). In DMD patients, the basal posterior-lateral wall is mostly affected by fibrosis, sparing the apex, RV, and atrium. However, even under these circumstances, heart failure may be induced by stress testing in the face of normal cardiac function at rest (Sethna, Rockoff *et al.* 1988; Schmidt, Burmeister *et al.* 2003).

Cardiac dysfunction accounts for 10~20% mortality in DMD, making it the second leading cause of patient death after respiratory failure (Gulati, Saxena *et al.* 2005). Cardiac dysfunction develops later than does respiratory dysfunction, but advances in respiratory support and patient care have allowed patients to survive into their late 20s and 30s or ordler. (Ishikawa, Bach *et al.* 1999; Ahuja, Kalra *et al.* 2000; Wagner, Lechtzin *et al.* 2007) Accordingly, cardiac complications now are commanding closer scrutiny in the management of advanced disease in DMD patients (Finsterer and Stollberger 2007; McNally 2008; Hermans, Pinto *et al.* 2010; Romfh and McNally 2010).

The early manifestations and functional consequences of DMD cardiomyopathy have been summarized recently by Judge *et al* in (Judge, Kass *et al*. 2011):

The cardiomyopathy that develops in DMD is characterized by normal or thinned left ventricular (LV) wall thickness and progressive decline in ejection fraction or fractional shortening. Variable degrees of LV dilation occur. Abnormal LV *relaxation*⁶ has also been identified as an early manifestation in DMD.

⁶ This abnormal relaxation presages the experimental results in section 2.5 showing that cardiac sheet function is normal in systole but abnormal in diastole.

Notably in DMD, cardiac diastolic dysfunction precedes systolic dysfunction, as observed by Doppler ultrasound (Markham, Michelfelder et al. 2006) and tissue Doppler ultrasound (Bahler, Mohyuddin et al. 2005). DMD cardiomyopathy also exhibits regional abnormalities: MRI scar imaging detects early onset of fibrosis at the epicardium of the more basal segments (Toyota, Fujimoto et al. 2002; Yilmaz, Gdynia et al. 2010). MRI tagging methods have revealed abnormal strain at the base but not the apex in patients with normal ventricular function (Ashford, Liu et al. 2005) as a manifestation of occult cardiac dysfunction. Mertens et al. reported significantly decreased early diastolic myocardial velocities in the anterolateral and inferolateral left ventricular walls with normal global systolic function (Mertens, Ganame et al. 2008). The previous work by Dr. Wickline's group using MR tagging uncovered occult abnormal wall strains occurring regionally at the base of heart in DMD patients (Ashford, Liu et al. 2005) has been supported by subsequent reports by Li et al suggesting that cardiac mechanical strain abnormalities are more prominent in the basal segments of hearts in mdx mice (Li, Liu et al. 2009). These studies indicate that regional myocardial dysfunction precedes global systolic dysfunction and validates the importance of advanced imaging techniques as tools for early delineation of DMD cardiomyopathy.

The observed *regional* dependencies for DMD cardiac mechanical dysfunction remain to be explained, as the mutation is present in *all* cardiomyocytes (Bulfield, Siller *et al.* 1984; van Deutekom, Bremmer-Bout *et al.* 2001; Banks and Chamberlain 2008).

1.1.4. Treatment Options

Unfortunately, there is no effective curative treatment, but palliative therapy has advanced in clinical trials over the last 20 years. To date, the only active pharmacologic interventions proven beneficial for the altering the natural history of the underlying dystrophic process have been the corticosteroids: prednisone and deflazacort (Rinaldi, Mayer *et al.* 2013). Both medications have been shown in clinical trials to lead to modest improvements in muscle strength and function, although the duration of these benefits typically lasts less than 2 years and the mechanism remains undefined (Araujo, Bonuccelli et al. 2013; Hollinger, Gardan-Salmon et al. 2013). Importantly, azathioprine (an immunosuppressant) and prednisone both decrease DMD muscle inflammatory infiltrates equally, but only prednisone improves strength (Griggs, Moxley *et al.* 1991; Kissel, Lynn *et al.* 1993). Animal studies have shown that prednisone enhances strength in dystrophin deficient transgenic mice that lack T-cells and B-cells, indicating prednisone's therapeutic effect does not operate through T-cell/B-cell immune suppressive pathways (Golumbek, Keeling *et al.* 2007).

β-adrenergic receptor blockers, angiotensin-converting enzyme inhibitors (ACEi) and angiotensin II-type 1 receptor blockers (ARBs) are also used for treating DMD cardiomyopathy (Duboc, Meune *et al.* 2007; Ogata, Ishikawa *et al.* 2009; Fayssoil, Nardi *et al.* 2010). Our group has demonstrated a beneficial effect of ACEi on reducing fibrosis and calcium content in Syrian hamster with cardiomyopathy, which harbors a related sarcoglycan deficiency. Although the precise mechanism is unclear, a 10 year long clinical study showed that ACEi reduces DMD mortality (Duboc, Meune *et al.* 2007). A large body of work on ACEi for scar tissue reduction has demonstrated utility for inhibiting focal and global cardiac remodeling after myocardial infarction by interrupting angiotensin signaling in the heart (Iwata, Cowling et al. 2011).

1.1.5. The *mdx* mouse model

Animal models are valuable tools for studying the pathogenesis of disease and for preclinical trials. The *mdx* mouse is the most widely utilized animal model for DMD (Williams and Allen 2007; Banks and Chamberlain 2008). It carries a point mutation leading to complete absence of dystrophin in myocytes. Similar to the later onset of cardiomyopathy in human patients, young *mdx* mice (< 3 months old) develop only a mild phenotype of cardiac disease and exhibit normal resting $[Ca^{2+}]_{I}$ (Williams and Allen 2007; Van Erp, Loch *et al.* 2010). However, both dilated cardiomyopathy and elevated resting $[Ca^{2+}]_{i}$ begin to be observed by about 10 months of age (Quinlan, Hahn *et al.* 2004; Williams and Allen 2007). Since *mdx* mice share the same genetic defect as DMD patients, they represent a standard model for assessing cardiac dysfunction and therapeutics.

1.1.6. Calcium Overload Hypothesis

With respect to pathophysiological features of dystrophin deficiency potentially responsible for mechanical dysfunction, several recent studies in isolated cardiomyocytes of mdx mouse have suggested that intracellular calcium ([Ca²⁺]i) mishandling is a key causative factor (Williams and Allen 2007; Fanchaouy, Polakova *et al.* 2009; Allen, Gervasio *et al.* 2010). Data acquired from mdx mice showed elevated resting [Ca²⁺]i in dystrophin-deficient cardiomyocyte (Pressmar, Brinkmeier *et al.* 1994; Williams and Allen 2007). The calcium overload activates calpains to promote proteolysis leading to cardiomyocyte degeneration (Spencer, Croall *et al.* 1995; Spencer and Tidball 1996; Gailly, De Backer *et al.* 2007). The calcium mishandling might

also affect the cardiac contraction that is controlled by calcium induced calcium release (Bellinger, Reiken *et al.* 2009).

Specifically, (Sapp, Bobet *et al.* 1996) reported that under 1 Hz pulse stimulation, peak force and time-to-peak force were similar in both mdx and wildtype cardiac fibers, but half-relaxation time was significantly prolonged in mdx, indicating normal contraction but abnormal relaxation. However, (Alloatti, Gallo *et al.* 1995; Janssen, Hiranandani *et al.* 2005) both reported depressed contractile force in mdx cardiac fiber, as compared to wildtype. This reported disparity in normal versus abnormal contractile force might be a consequence of a higher stimulation frequency used by Alloatti (4 Hz) and Janssen (2.8Hz). Nicolas et al reported abnormally prolonged relaxation in mdx skeletal muscles, while the active contraction metrics were normal (Nicolas-Metral, Raddatz *et al.* 2001), suggesting a decreased capacity for calcium removal in mdx cells leading to alterations of muscle relaxation that accords with the observation that diastolic dysfunction precedes systolic dysfunction in DMD patients and mdx mice (section 1.1.3).

Several mechanisms for elevated resting $[Ca^{2+}]i$ have been proposed, offering opportunities for developing new therapies. One promising therapeutic approach is to inhibit the calcium influx that occurs through SAC (stretch activated channels) or through transient microruptures of the sarcolemma. It is well recognized that *mdx* cardiomyocytes are sensitive to stretch induced injury that is correlated with increased resting $[Ca^{2+}]i$ (Lynch 2004; Yeung and Allen 2004). Inhibition of SAC or the application of the membrane sealant poloxamer 188 reduces resting $[Ca^{2+}]i$ and prevents injury (Yasuda, Townsend *et al.* 2005; Whitehead, Streamer *et al.* 2006; Ward, Williams *et al.* 2008). In vivo application of poloxamer 188 also protects *mdx* mice from heart failure during dobutamine mediated cardiac challenge (Yasuda, Townsend *et al.* 2005). Inhibiting the leakage of ryanodine receptor (RyR2), a calcium release channel located on the sarcoplasmic reticulum, is another emerging therapeutic approach (Bellinger, Reiken *et al.* 2008; Shannon 2009). Notably, the RyR2 of *mdx* cardiomyocytes appears to be hypernitrosylated, which prevents RyR2 from fully close at resting state (Bellinger, Reiken *et al.* 2009). The resulting sarcoplasmic reticulum calcium leakage through RyR2 is indicated by the presence of calcium sparks (Ward, Schneider *et al.* 2000). Several 1-4-benzothiazepine derivatives, such as S107 and KD201, can stabilize RyR2 to reduce calcium leakage and improve cardiac function (Wehrens, Lehnart *et al.* 2004; Lehnart, Terrenoire *et al.* 2006).

Besides pharmaceutical approaches, transgenic approaches provide potential remediation of calcium mishandling. The root cause of DMD is the lack of dystrophin, making transgene therapy for correction of the dystrophin gene preferable to altering calcium handling genes. However, clinical studies suggest that immunity to transgenic micro-dystrophin in DMD patients represents a significant challenge, because patients that are dystrophin-null recognize transgenic micro-dystrophin as an exogenous antigen (Mendell, Campbell *et al.* 2010). Therefore, transgenic strategies based on endogenous genes, such as sarco/endoplasmic reticulum Ca²⁺-ATPase (SERCA), may offer an immune advantage as compared to dystrophin gene therapies. Indeed, cardiac SERCA2a gene transfection has been shown to correct ECG abnormalities in heart (Shin, Bostick *et al.* 2011); and skeletal SERCA1 gene transfection corrected skeletal muscle calcium reuptake in dystrophic mice (Goonasekera, Lam *et al.* 2011).

The previously reported lack of efficacy of L-type calcium channel blockers against DMD associated muscular dystrophy should not negate the potential value of calcium channel targeted therapy (Phillips and Quinlivan 2008). L-type calcium channels regulate calcium inflow upon depolarization and are closed in the resting state. Since calcium leakage through these channels has not been reported, L-type calcium channels' effect on regulating resting $[Ca^{2+}]i$ should not be expected to be significant.

In summary, calcium targeted therapy holds potential for treating DMD. Because calcium influx cannot be measured in vivo in heart tissue, we propose that development of a surrogate cardiac phenotype that reflected $[Ca^{2+}]i$ would be useful for evaluating the efficacy of treatment. We have previously shown that left ventricular myocardial architecture changes dynamically in cardiac cycle due to calcium regulated contraction-relaxation (Chen, Liu *et al.* 2005).

1.1.7. Hypothesis

How calcium mishandling influences cardiac mechanical dysfunction in whole heart preparations has not been clarified, especially taking into account cellular and mechanical events that are spatio-temporally distributed and regionally heterogeneous (DeAnda, Komeda *et al.* 1998; Balzer, Furber *et al.* 1999). Given the recognized abnormal and regionally heterogeneous wall strains (Figure 3) and fibrosis (Ashford, Liu *et al.* 2005; Yilmaz, Gdynia *et al.* 2008; Yilmaz, Gdynia *et al.* 2010) in DMD hearts of both human patients and *mdx* mice, we hypothesize that the sheet functional consequences of dystrophin deficiency plays out *regionally* in *DMD* hearts over time despite the ubiquitous lack of dystrophin protein in all cardiomyocytes. We hypothesize that elevated resting $[Ca^{2+}]i$ directly affects myofiber and sheet kinetics in *mdx* mouse hearts which can be quantified by diffusion tensor MRI (DTI, Chapter 2).



Figure 3. Cardiac MRI detected abnormal strain at base but not apex of DMD patient hearts, indicating base exhibits more myocardial dysfunction than apex. Figure from (Ashford, Liu *et al.* 2005)

The heart comprises a syncytium of cardiomyocytes that are organized in unique 3D fibersheet architecture that is critical for optimizing a coordinated sequence of excitation, contraction, and relaxation (Streeter and Hanna 1973; Costa, Takayama et al. 1999). Many reports have shown that sheet function produces a sliding motion that is accompanied by dynamic reorientation in systole, contributing to approximately 50% of focal wall thickening (LeGrice, Smaill et al. 1995; LeGrice, Takayama et al. 1995; Costa, Takayama et al. 1999; Dou, Tseng et al. 2003; Chen, Liu et al. 2005). Significant sheet-fiber deformation was also found in early diastole (Ashikaga, Criscione et al. 2004). Because one role of dystrophin is to anchor cardiomyocytes to the extracellular matrix to coordinate contraction-relaxation coupling, we propose that dystrophin deficiency could adversely affect sheet function. Given that ventricular wall strain vary regionally from base to apex, and in light of the expected vulnerability of dystrophin deficient cardiomyocytes to stretch-induced membrane injury, we sought to examine the role of sheet function in the regional mechanical defects reported in the DMD cardiomyopathy (Ashford, Liu et al. 2005; Li, Liu et al. 2009). We investigated the heart sheet structure defect in regions of heart with ultra high-field MRI, because MRI is non-destructive and provides functional assessment according to the methodology described in Chapter 2.

Examination of this regional structural functionality may add to the fundamental understanding of the nature of the heart dysfunction in DMD and lead to improved early monitoring of interventions designed to salvage cardiac structure and function.

Chapter 2: MRI for Cardiac Sheet Function

2.1. Cardiac muscle structure

The normal architecture of the heart muscle remains elusive despite recent advance in high resolution images offered by MRI and photoacoustic imaging (Gilbert, Benson *et al.* 2007). Several models have been proposed to explain cardiac structure without reaching a consensus. Therefore, accurate knowledge about the architecture of heart muscles is critical in understanding normal heart and cardiomyopathy. A comprehensive understanding of myocardial architecture would help us to (1.) explain cardiac electric and mechanical mechanics; (2.) simulate cardiac physiology; (3.) design intervention/monitoring strategy; and (4) understand structure-function relationship of the heart. In the future, we could compare myocardial architecture among fish, amphibian, reptile, bird, and mammal, to understand the evolution of the heart, in which lies the natural history of how the heart adapts the gravity and land environments with high pressurizing ability. A comprehensive review of cardiac architecture during 17th century and year 2002 can be found in a MIT doctorate dissertation (Dou 2002). Below is my review of recent update.

2.1.1. Fiber

Myocardial fibers are cardiac muscle cells, which include sarcolemma and bundles of parallel or sometimes branching myofibrils. *Cell and fiber are terms that may be used interchangeably* (Pollack 1990). The 3D myocardial architecture is critical for optimizing a coordinated sequence of excitation, contraction, and relaxation (Streeter and Hanna 1973; Costa, Takayama *et al.* 1999). The earliest study of fiber helical path was a reprint in Latin by Stensen in 1664. An English review can be found in (Dou 2002).

Based on laborious histology work, Streeter reported a smooth transition of fiber helix angle (α , angle between fibers and local circumferential axis) in the transmural direction (Figure 4) Streeter proposed a wicker basket model to explain how α might reflect the ventricular volume (Figure 4 B). However, the basket model represents the LV incorrectly, because recent studies have found that helix angle *decreases* during longitudinal extension (i.e., diastole) (Chen, Liu *et al.* 2005). In contrast, the wicker basket model predicts *increased* α during longitudinal extension (Figure 4 B). Nevertheless, a smooth transmural transition of α is observed in many mammalian species (Scollan, Holmes *et al.* 1998; Tseng, Wedeen *et al.* 2003; Chen, Liu *et al.* 2005; Helm, Beg *et al.* 2005). Chen 2005 found that $\Delta \alpha$ is decreased by ~7 degree in diastole compared to systole. This small change was not detected by Streeter with his histology method (Streeter and Hanna 1973), probably due to the unavoidable dehydration/deformation artifacts of histology specimen preparation.



Figure 4. **A.** Cartoon of LV fiber orientation from (Streeter and Hanna 1973). Green: epicardial fiber, left handed helix. Blue: midcardial fiber, circumferential. Red: endocardial fiber, right handed helix. **B.** Streeter's proposed wicker basket model of helix angle change between diastole and systole.

2.1.2. Sheet

Robert Grant conceived and illustrated how fibers might aggregate into a laminar sheet structure (Grant 1965) (Figure 5). But the laminar sheet structure was not actually visualized until 1995 by LeGrice using the histology method (LeGrice, Smaill *et al.* 1995). The discovery of the myocardial sheet exemplified how an obvious structure eluded scientists for a long time. All of the popular experimental mammalian hearts exhibit laminar structures, similar to layers of muscle found in a fish fillet (Figure 6). Myocardial sheet is about four layers of fibers aggregating into a laminar structure (simply called a "sheet" in this thesis). Figure 7 shows the sheet architecture in a fresh, unfixed and unstained mouse left ventricle (Zhang, Cheng *et al.* 2012). Figure 7 is acquired with advanced photoacoustic microscopy (PAM) system. The

significances of the PAM image of myocardial sheet are (1) it is the highest resolution of unfixed sheet to the best of my knowledge; (2) the heart is unfixed, meaning there's minimal dehydration artifacts compared to formalin fixed hearts; (3.) it requires no staining nor contrast agent, saving time and artifacts of doing so; (4) it is a three dimensional imaging modality, suitable for investigating the 3D structure like the heart muscle. For technical details, please refer to (Zhang, Cheng *et al.* 2012).



Figure 5. Robert Grant proposed that "planes" can be formed by fibers. "Schema of fiber to fiber relationship which appears to obtain typically in the left ventricle. While fibers are parallel, by their branchings they form 'planes' which appear to rotate." (Grant 1965)



Figure 6. Cooked fish fillet shows sheet structure of skeletal muscle of fish, indicating muscle fibers aggregate into secondary structure in a form of sheets. Cooking helps the separation of sheets, indicating the connective tissues between sheets are more vulnerable to thermal denaturation. The major difference between fish sheet and mammal myocardial sheet is in the fish, fiber is parallel to sheet normal direction. In contrast, in myocardial sheet, fiber is perpendicular to the sheet normal direction.



Figure 7. Photoacoustic microscopic (PAM) image of a unfixed, unstained, and un-dehydrated myocardium, showing sheet structure of a mouse left ventricle. (a), PAM image of the left ventricle free wall (FW) marked in (c) acquired at 532-nm wavelength. (b), Close-up PAM image. The boundaries of the branching sheets are apparent. Red dashed line: long axis. Yellow dashed-dot line: sheet boundary. (c), Photograph of the heart specimen. LV, left ventricle; RV, right ventricle. (Zhang, Cheng et al. 2012).

(C)

Here is a chronicle summary of important discovery on sheet. After LeGrice visualized the sheets with histology (LeGrice, Takayama *et al.* 1995), Costa and Chen inferred that sheet shearing contributes dominantly to heart wall thickening using radiocinegraphy and DTI, respectively (Costa, Takayama *et al.* 1999; Chen, Liu *et al.* 2005). Using diffusion tensor MRI on alive human subjects, Dou reported that sheets change orientation in cardiac cycle (Dou,

Tseng *et al.* 2003). Specifically, sheets align closer to short axis in systole than diastole. Using confocal histology, Pope reported that sheet presents dominantly at mid- and endocardium but not at epicardium (Pope, Sands *et al.* 2008). Using 9.4T MRI with 3 day long scan time, Gilbert confirmed Pope's finding whith the field of view of a whole heart, revealing more mysterious helical nature of sheets in high resolution (25μ m) (Gilbert, Benoist *et al.* 2012). Readers are directed to view the spplemental movie in Gilbert's paper for the 3D whole heart sheet structure.

2.1.3. Nomenclature of the heart segments

Figure 8 illustrates the terminology used to describe the various regions of the heart. This terminology follows AHA consensus (Cerqueira, Weissman *et al.* 2002).



Figure 8. Left ventricle (LV) is rounded in shape in short axis view. Right ventricle (RV) has a semilunar shape. Aorta and vena cava are closer to the heart *base* than the *apex*. Apex is pointing to the left in normal human chest. Heart base is physically closer to the head and has larger diameter than apex.

2.2. Diffusion Tensor MRI

2.2.1. Background

MRI modulates magnetic field to affect nuclear spins to form images (Appendix). Under magnetic gradient, the diffusion of molecules *dephases* the nuclear spins and thus *decreases* MR signal. DTI (Diffusion Tensor MRI) quantifies the ADC (apparent diffusion coefficient) from the signal decay. Specifically, the DTI signal predominantly comes from nuclear spins of water protons. In spin echo imaging, the signal intensity is governed by eq1.

$$S = PD(1 - e^{-TR/T1}) e^{-TE/T2} e^{-bD}$$
(eq1)

S is signal intensity, **PD** is proton density, **T1** and **T2** are relaxation (decay) times of protons within the samples under given **B**₀. **D** is the ADC, which represents the observed Brownian motion of water molecules in tissue. **TR** and **TE** are repetition time and echo time. **b** is the diffusion-weighing factor with units of s/mm^2 . From eq1, one can see higher **b** value decreases **S** (Basser 1995).

TR, **TE** and **b** can be adjusted on the scanner. Longer **TR** and shorter **TE** generate better signal to contrast ratio (SNR) but require longer scan time and faster responding electronics, which are both more expensive. Tradeoff was made between SNR and scan time. We want D for the diffusion information in it, but it is impossible to calculate the D with just one **S** in eq1. Therefore Stejskal and Tanner devised the two-**S** method to calculate the D (Stejskal 1965). See equations below:

$$S_1 = PD\left(1 - e^{-\frac{TR}{T_1}}\right)e^{-\frac{TE}{T_2}}e^{-b_1 D} = S_0 e^{-b_1 D} \qquad \dots \dots \text{measurement 1}$$

$$S_{2} = PD \left(1 - e^{-\frac{TR}{T_{1}}} \right) e^{-\frac{TE}{T_{2}}} e^{-b_{1}D} = S_{0} e^{-b_{2}D} \qquad \dots \dots \text{measurement } 2$$
$$\frac{S_{2}}{S_{1}} = e^{-(b_{2}-b_{1})D} \qquad (\text{eq2})$$

Rearranging eq2, we obtain:

$$D = -\ln\left(\frac{s_2}{s_1}\right) / (b_2 - b_1)$$
(eq3)

That is, we can derive a scalar D by attenuating the signal with two different b values. For example, I have used $b_1 = 0$, and $b_2 = 800$ (s mm⁻¹) in my work.

2.2.2. Restricted Diffusion

Brownian motion of water molecule in an *isotropic* diffusion environment is characterized by a *single* parameter, the diffusion coefficient, D. For example, in threedimensional space, random walkers spread with time t according to a Gaussian distribution would have a mean square displacement, $E(x^2) = 6Dt$. That is, in isotropic diffusion environment, D is the same in all direction. However, complexity in the microscopic structure of myocardium restricts water molecular motion anisotropically and results in non-Gaussian diffusion and hence different values of D in different directions. Therefore, instead of a single scalar D, multiple Ds are required to describe the diffusion anisotropy.

To detect anisotropic diffusion, one could repeat Stejskal's two measurements method (eq3) in 6 non-collinear directions to fill in the following matrix **D**:

$$D = \begin{array}{ccc} Dxx & Dxy & Dxz \\ Dyx & Dyy & Dyz \\ Dzx & Dzy & Dzz \end{array}$$
(eq4)
The D in eq4 is a 3 x 3 matrix, which is also a rank 2 tensor. For example, Dxx is the ADC in x direction. Dxy is ADC in xy direction, which is in the same direction of the vector (x,y,z)=(1,1,0). From the diffusion tensor, **D**, we can assess magnitude of ADC in different directions. Specifically, the ADC is found to be the greatest in fiber direction and smaller in perpendicular directions. Therefore, the tensor not only provides the orientation information (e.g, which direction has the largest ADC) but also relative ratios of ADCs in different directions (i.e., anisotropy of the diffusion environment, Chapter 3). The fact that water diffusion is restricted by the underlying tissue microstructure provides a unique method of assessing myocardium architecture (Safford, Bassingthwaighte *et al.* 1978; Garrido, Wedeen *et al.* 1994; Ennis and Kindlmann 2006).

In summary, DTI is an established method for 3D reconstruction of myocardial microarchitecture that can depict structural changes across the heart cycle in the intact heart (Chen, Liu *et al.* 2005; Sosnovik, Wang *et al.* 2009). The value of DTI is it is non-invasive, non-destructive, 3-dimensional and rapid, compared to traditional histology (Mori and Zhang 2006; Roberts and Schwartz 2007).

2.1. Method: DTI for *mdx* mice cardiac structure

In the heart, the primary eigenvector is considered to be aligned with the local orientation of cardiac myofibers, and thus defines the fiber angle (α) (Hsu, Muzikant *et al.* 1998; Scollan, Holmes *et al.* 1998; Holmes, Scollan *et al.* 2000; Geerts, Bovendeerd *et al.* 2002). The secondary eigenvector is aligned parallel to sheet surfaces (Tseng, Wedeen *et al.* 2003) and thus defines the sheet angle (β , Figure 9 A). This definition system is based on the fact that eigenvalues of a

diffusion tensor are also the apparent diffusion coefficients describing microscopic diffusion movements, which are the largest in the fiber direction and second largest in the sheet direction (Scollan, Holmes *et al.* 1998; Chen, Suzuki *et al.* 2005).



Figure 9. Representative DTI and histology images of sheet architecture in WT mouse hearts. **A**, A local cylindrical coordinate system was used to describe LV myocardial structures. The primary eigenvector (EV1) of the diffusion tensor is characterized by helix angle (α), which is the angle between the circumferential axis and the projection of the first eigenvector onto the circumferential-longitudinal plane; the α shown in A is set as negative. The secondary eigenvector (EV2) is characterized by sheet angle (β), which is the angle between the secondary eigenvector and the radial axis; the β shown in A, toward base and toward endocardium, is set as negative. **B & C**, short-axis views of DTI determined sheet angle (β) in a basal slice of WT heart that was sequentially arrested in diastole (B) and systole (C). The posterior-lateral wall of LV

manifested two populations of β , exhibiting positive (red) and negative (blue) values. D, Histology of sheet organization in a WT heart. Images were acquired from the basal-lateral wall, which is highlighted on the H&E stained image (insert). Two populations of β were observed.(LV, left ventricle; RV, right ventricle; PA, papillary muscle; P, Posterior; L, Left; R, Right; A, Anterior).

Experimental Methods

2.1.1. Langendorff perfusion for DTI

<u>Animals</u>

Adult male *mdx* (C57BL/10ScSn-Dmd*mdx*/J, N=14) and age matched male wild type (WT: C57BL/10SnJ, N=10 and C57BL/6, N=3) mice were purchased from the Jackson Laboratory (Bar Harbor, ME, USA) and the National Institute on Aging (NIA, MD, USA)(18) and housed under a AAALAC-approved facility to 16-month of age. All mice were housed in the same animal facility and cared for under approval by Washington University's animal study committee. All mice were separated into two groups for DTI and optical mapping studies described below.

Isolated Heart Preparation for DTI

Ten *mdx* mice and ten WT mice were used for both diastolic and systolic DTI measurement (Figure 10). Mice were heparinized (1000 U/kg) and anesthetized with isoflurane. Hearts were excised and cannulated for retrograde perfusion using a procedure modified from (Chen, Liu *et al.* 2005). Briefly, hearts were first perfused with Krebs buffer at 60 mmHg constant pressure for approximately 5 min to wash out the residual blood. Five *mdx* hearts and five WT hearts were perfusion-arrested in diastole (perfusion pressure = 60 mmHg) with regular

St. Thomas' cardioplegic solution containing normal $[Ca^{2+}]$ of 1.2 mmol/L (denoted as *mdx*-NC and WT-NC, respectively) under room temperature. DTI of diastolic arrested hearts was performed for 1.5 hours. After that, hearts were reperfused with 37 °C Krebs buffer to resume beating. Upon stabilization of contraction (~6 min, up to 300 beats per min), 2.5 mmol/L Ba²⁺ was then introduced to induce systolic cardiac contracture while Ca²⁺ was lowered to 0.078 mmol/L (Munch, Comer *et al.* 1980; Toyota, Fujimoto *et al.* 2002; Chen, Liu *et al.* 2005). To evaluate the effect of $[Ca^{2+}]$ on diastolic cardiomyocyte architecture in *mdx* mice, another five *mdx* hearts and five WT hearts were perfusion-arrested in diastole with a modified cardioplegic solution containing low $[Ca^{2+}]$ of 0.078 mmol/L (denoted as *mdx*-LC and WT-LC, respectively). The 0.078 mmol/L $[Ca^{2+}]$ was chosen over the calcium-free condition to minimize the risk of complete depletion of intracellular calcium, which can cause calcium overload upon reperfusion with solutions containing normal $[Ca^{2+}]$, thereby exerting unfavorable effects on certain ion channels (Munch, Comer *et al.* 1980; Robinson and Harwood 1991).

The systolic arrest of *mdx*-NC, WT-NC, *mdx*-LC, and WT-LC (defined below) are all achieved with the same modified Tyrode solution containing 2.5 mmol/L Ba²⁺ and 0.078 mmol/L Ca²⁺ (Chen, Suzuki *et al.* 2005). All solutions were equilibrated with 95% O2 plus 5% CO2 (PH 7.4) and contained adenosine (45 μ mol/L) to maximally dilate the coronary vessels (Judd, Rottman *et al.* 1992). Bovine serum albumin (BSA; 61 μ mol/L) was added to the perfusate to minimize interstitial edema (Dunphy, Richter *et al.* 1999; Watts and Maiorano 1999). DTI of systolic contracture hearts were conducted for ~1.5 hours. Upon the completion of MRI, all hearts were perfused with 10% formalin for histological study.



Figure 10. Flow chart for DTI acquisitions for diastolic and systolic arrested hearts. Left, ten mice perfused with normal calcium (NC) concentration for diastolic arrest then high barium for systolic arrest. Right, ten mice perfused with low calcium concentration (LC) for diastolic arrest then high barium for systolic arrest. See text for details.

2.1.2. Perfusion compatible MR Coil

A dedicated MRI-compatible retrograde mouse heart perfusion system was made. This system provides a quantitative and non-destructive assessment of diastolic and systolic myocardial architecture from the same heart. The system includes: (1) a custom-built 15 mmdiameter solenoid coil, which provides high SNR: 65 for non-diffusion weighted images and 40 for diffusion weighted images (See Figure 11 for the coil design.); (2) a Langendorff constantpressure perfusion apparatus. The constant perfusion pressure is controlled by a reservoir with constant height. Perfusate efflux was removed by vacuum suction. The heart was contained in a plastic tube made by syringe, which is omitted in Figure 11 for simplicity but is important to prevent contact between perfusate and the copper coil. A variable capacitor was soldered at the center of the gap of the single turn copper coil for resonance frequency tuning. The distance, **h**, between the cable loop and the copper coil determines the coefficient of coupling, **k**, which determines the mutual inductance, **L**.

$$\mathbf{L} = k_{\sqrt{L_1 L_2}}$$
, k is between 0 (when h is large) and 1 (when d is small) (eq5)

The Figure 11 B is an equivalent circuit of my coil. The resonance frequency, f, of the circuit can be determined by series RLC resonator. Practically, L and C were adjusted to resonate at 499.70 MHz for the 11.7 tesla ultra-high field MR scanner used in this study.

$$f = \frac{1}{\sqrt{LC}}$$
(eq6)



Figure 11. **A.** Cartoon of single turn solenoid coil and perfused mouse heart setup. A plastic tube containing the heart and preventing perfusate from contacting the copper coil is omitted for visual simplicity. Distance between the cable loop and the copper coil is denoted by h. **B.** Equivalent circuit. R is the resistance of loading (perfusate and heart) and copper. L_1 is the inductance of the cable loop. L_2 is the inductance of the copper coil. C is variable capacitor.

2.1.3. Imaging parameters

R

Β.

All MRI experiments were performed on a Varian 11.7 tesla small animal MR scanner (Varian Associates; Palo Alto, CA). Long-axis scout images were acquired as previously described (Chen, Liu *et al.* 2005). A multislice spin-echo sequence with diffusion-sensitizing gradients were used to acquire short-axis, diffusion weighted images. Diffusion-sensitizing gradients were applied in six noncolinear directions. Imaging parameters were as follows: echo time, 46 ms; time interval between diffusion-sensitizing gradients, 30 ms; b-value, 0 and 800 s/mm²; in-plane resolution, $125 \times 125 \,\mu$ m², repetition time, 3 s; and image-acquisition time, 1 h. For diastolic arrested hearts, eight contiguous slices (thickness = 0.7 mm) extending from the

С

base to apex were acquired. For systolic contracture hearts, eight contiguous slices were acquired with slice thickness adjusted to 0.55–0.65 mm based on the fraction of longitudinal shortening. Attention was paid to keep the slice selection and the heart orientation the same between diastole and systole imaging.

2.1.4. Post processing, sheet angle calculation (β)

All MR images were processed using custom-developed software written in Matlab (The MathWorks Inc., M.A., USA) as described previously (Chen, Liu *et al.* 2005). Specifically, fiber helix angle (α) and sheet angle (β) were calculated based on a right-handed helix, the fiber helix angle (α) was calculated as the angle between the circumferential cardiac axis and the projection of the first eigenvector of the diffusion tensor onto the circumferential-longitudinal plane (Scollan, Holmes *et al.* 1998). Sheet angle (β) was determined as the angle between the secondary eigenvector and the radial axis. Sheet inclination toward the base from endocardium to epicardium was set as positive (Figure 9 D) (Scollan, Holmes *et al.* 1998; Tseng, Wedeen *et al.* 2003). Because regional wall thickening is accompanied by a reduced magnitude of the sheet angle, $|\beta|$, therefore $|\beta|$ was used to analyze sheet function (LeGrice, Takayama *et al.* 1995; Chen, Liu *et al.* 2005). LV wall thickness was calculated as the mean distance between epicardial and endocardial borders. The transmural difference of α , defined as the difference in endocardial and epicardial helix angles, i.e., $\Delta \alpha = \alpha_{endocardium}-\alpha_{epicardium}$, was used to quantify transmural changes of fiber orientation.

2.1.1. Statistical methods

Statistical differences were assessed with the analysis of variance (ANOVA) or the multivariate analysis of variance (MANOVA) as appropriate. Bonferroni correction was applied

to post-hoc comparisons. A value of p < 0.05 was considered significant. Data were expressed as mean \pm standard error of the mean (S.E.). All statistics were conducted with the IBM SPSS Statistics program.

2.1.2. Histology

After DTI acquisition, five *mdx* and five wildtype hearts were rapidly fixed in 10% formalin. Fixed hearts were sliced into four blocks with 2 mm thickness from base to apex. Each slice was embedded in paraffin and sectioned at 4 μ m thickness. The tissue sections were stained with picrosirius red for the identification of fibrosis (Van Erp, Irwin *et al.* 2006). The extent of myocardial fibrosis was quantified from digitized histological images using a thresholding algorithm implemented in Matlab. Regions that demonstrate stronger red-to-green ratio in picrosirius red staining were considered to represent interstitial fibrosis. The percentage of tissue fibrosis was calculated as the ratio of pixels representing fibrotic tissue vs. pixels of the entire short-axis slice, and the average value in two adjacent slices was used in analysis.

2.2. Results: Sheet function in 16-month old *mdx* mouse hearts

2.2.1. Morphological observation of β with histology and PAM

DTI, histology, and PAM all show a similar transmural morphological character of sheet, that is, a sharp change of β at midmyocardium of basal posterior mid-wall (Figure 12).



Figure 12. A & B, representative short-axis view of DTI determined sheet angle (β) in a basal slice of the same WT heart that was sequentially arrested in diastole (A) and systole (B). The posterior-lateral wall of LV possessed two populations of β , exhibiting positive (red) and negative (blue) values. The transmural portion of positive β is increased in systole (compare the portions of red-colored pixels under the dashed lines in A & B). The dotted line in B indicates a 3 population β transition that appears only in systole. C, Photoacoustic microscopy of long-axis view of a freshly half-split WT mouse heart confirmed two populations of β in LV wall. The insert showed H&E stained long-axis WT mouse heart slice with the corresponding region of photoacoustic image squared. LV: left ventricle. RV: right ventricle. PA: papillary muscle.

2.2.2. 16 month old *mdx* hearts manifest reversible regional diastolic sheet dysfunction

DTI defined two populations of β with opposite signs in the LV wall of WT and *mdx* mice, as confirmed by several previous reports (LeGrice, Smaill *et al.* 1995; Tseng, Wedeen *et al.* 2003; Chen, Liu *et al.* 2005; Helm, Beg *et al.* 2005; Gilbert, Benson *et al.* 2007; Pope, Sands *et al.* 2008). Under conditions of barium-induced systolic arrest, $|\beta|$ value is decreased in systole (Figure 9 B&C, Figure 13 A&B), agreeing with *in vivo* and *ex vivo* observations (LeGrice, Smaill *et al.* 1995; LeGrice, Takayama *et al.* 1995; Costa, Takayama *et al.* 1999; Chen, Liu *et al.*

2005). At each short-axis imaging slice, the slice-averaged systolic $|\beta|$ was comparable among all groups of hearts (Figure 13 B, p = N.S.). However, the diastolic $|\beta|$ of *mdx* hearts (*mdx*-NC) was ~10° lower than that of WT hearts (WT-NC) at the base when perfused with regular cardioplegic solution (p < 0.05, Figure 13 A). The $|\beta|$ at the apex of *mdx*-NC hearts appeared normal (p = N.S. compared to WT-NC at each corresponding short-axis imaging slice), demonstrating that diastolic sheet dysfunction in *mdx* hearts occurred focally at the base. By reducing the [Ca²⁺] of the cardioplegic solution to 7% of its regular concentration, a complete and rapid restoration of diastolic $|\beta|$ in *mdx* hearts (*mdx*-LC) to normal values was observed. No effect of calcium manipulation on WT cardiac sheet function was observed (WT-LC, p = N.S. compared to WT-NC at each corresponding slice), indicating that the diastolic sheet dysfunction in *mdx* hearts imaging slice), indicating that the diastolic sheet that even at this advanced age, a principle mechanical abnormality in *mdx* mice likely is associated with regionally-dependent incomplete sarcomere relaxation.

At the base, further regional analysis indicated that the maximal decrease of diastolic $|\beta|$ in *mdx*-NC hearts occurred in the endocardium. Compared to WT-NC, *mdx*-NC exhibited 13° lower $|\beta|$ (75° ± 2° vs. 62° ± 3°) in the endocardium (inner 1/3 wall-depth in transmural direction); 10° lower $|\beta|$ (74°± 1° vs. 64° ± 2°) in the midcardium (middle 1/3 wall-depth in transmural direction); and 8° lower $|\beta|$ (68° ± 2° vs. 60° ± 2°) in the epicardium (outter 1/3 wall-depth in transmural direction) (Figure 13 C). Moreover, in the posterior-lateral wall, the diastolic $|\beta|$ was 51° ± 6° in *mdx*-NC (Figure 13 F). This value was approximately 20° lower than that of *mdx*-LC (70°± 4°), as well as that of WT-NC (72° ± 4°) or WT-LC (72° ± 6°) groups (*p* < 0.05 for all comparisons. See post hoc multiple comparison in Table 2 in page 107). Together these

findings indicate a clear regional and transmural dependence of incomplete relaxation, despite the ubiquitous dystrophin mutation.



B. Systolic $|\beta|$ (°)



C. Diastolic Transmural $|\beta|$ (°) Endocardium Midcardium 80 70 t



60

50



E. mdx-LC D. mdx-NC **|β|**(°) <u>90</u>° 30°





Figure 13. DTI-determined diastolic and systolic $|\beta|$ in short-axis slices. A, mdx hearts exhibited abnormally lower diastolic $|\beta|$ that was recovered to normal value after reducing [Ca²⁺] in the perfusate. For hearts perfusion-arrested in diastole with normal calcium solution, mdx hearts (*mdx*-NC) exhibited lower $|\beta|$ than did WT (WT-NC) hearts with the maximal ~10 degree difference occurred at the basal region. In contrast, *mdx* hearts perfused with the low calcium solution (*mdx*-LC) exhibited normal diastolic $|\beta|$. Varying [Ca²⁺] had no effect on diastolic $|\beta|$ in WT hearts (WT-LC). **B**, systolic $|\beta|$ exhibited no difference among four groups of hearts. **C**, diastolic $|\beta|$ in the epicardium, midcardium, and endocardium at the base of WT and *mdx* hearts. In all regions, diastolic $|\beta|$ of *mdx*-LC hearts were comparable to that of WT-LC and WT-NC hearts, but all are higher than that of *mdx*-NC hearts. The largest difference in diastolic $|\beta|$ between mdx-LC and mdx-NC was observed in the endocardium. D & E, representative DTIdetermined diastolic $|\beta|$ maps on short-axis slices at the base of *mdx*-NC and *mdx*-LC hearts. The lower $|\beta|$ in *mdx*-NC heart, indicated by fewer regions with red color, was visually appreciable. F, in the posterior-lateral region (the areas enclosed by dashed line in D & E), the diastolic $|\beta|$ of mdx-NC hearts was approximated 20 degree lower than that of mdx-LC hearts and of all WT hearts. Data represent mean \pm SEM. N = 5 for each group. *, p < 0.05; †, p <0.005 for the indicated short axis location.

2.2.3. 16 month old *mdx* and WT hearts exhibit similar fiber organization and function

Wall thickness and through-wall differences of myofiber helix angle ($\Delta \alpha$) were similar

between mdx and WT mice (Figure 14), suggesting that the major mechanical abnormality was

confined to sheets rather than fibers. The normal wall thickness agrees with former publications

(Li, Liu et al. 2009; Stuckey, Carr et al. 2012)



Figure 14. Averaged wall thicknesses in each short axis slice are similar in all four groups, in both diastole (**A**.) and in systole (**B**.) Averaged through-wall myofiber helix angle differences $(\Delta \alpha)$ in each short axis slice are similar in all four groups, in diastole (**C**.) and in systole (**D**.). Overall, $\Delta \alpha$ is increased in systole, agreeing with a prior publication (Chen, Liu *et al.* 2005). No significance was detected in all comparison in each panel.

2.2.4. 3 month old *mdx* mice exhibit smaller sheet dysfunction, still reversible by low calcium perfusion

The max difference in $|\beta|$ between *mdx*-NC and *mdx*-LC is $6.1^{\circ} \pm 3.6^{\circ}$ (p < 0.66, Table 3 in page 110) and $9.3^{\circ} \pm 2.26^{\circ}$ (p < 0.005, Table 2 in 107), both at base in 3- and 16-month old mice respectively (Figure 15), indicating that this [Ca²⁺]-dependent sheet dysfunction is also loci dependent.



Figure 15. A. $|\beta|$ of 3 month old *mdx* and WT mice under normal calcium (NC) and low calcium (LC) perfusion. B. $|\beta|$ of 16 month old *mdx* and WT mice under NC and LC perfusion. Data represent mean \pm SEM. N = 5 for each group. *, p < 0.05; †, p < 0.005 for the indicated short axis location.

2.3. Discussion of $|\beta|$

This study shows for the first time that mdx hearts exhibit an abnormal myocardial sheet angle magnitude, $|\beta|$, in diastole but not in systole. Agreeably, Goldberg reported that DMD patients have abnormally slower wall thinning but normal wall thickening in echocardiography (Goldberg, Feldman *et al.* 1980) (Figure 16), indicating that dystrophin affects normal diastolic rather than systolic function. Furthermore, regional and intramural gradients of diastolic dysfunction (Figure 13 A & C) are evident, primarily in the more basal segments where abnormal calcium handling also occurs (Chapter 4).



Figure 16. Using echocardiography, Goldberg has shown that DMD patient heart has significantly slower wall thinning but normal wall thickening, indicating that the absence of dystrophin affects diastolic function more than systolic function, agreeing with the DTI measured sheet angle data in Chapter 2. Figure from (Goldberg, Feldman *et al.* 1980).

These co-localized abnormalities in calcium handling and diastolic calcium-dependent sheet dysfunction strongly suggest that features other than the simple absence of dystrophin are critical to the evolution of heart failure in DMD. Indeed, it is well recognized that the mechanical defect in the dystrophin encoding gene does not independently elicit DMD cardiac dysfunctions such as reduced circumferential wall strain (Ashford, Liu *et al.* 2005), which emerge in a regionally heterogeneous fashion over time, and not at all loci where dystrophin is missing. This locidependency agrees with my findings.

Given that systolic $|\beta| < \text{diastolic } |\beta|$ (Figure 13 A & B), the observed lower diastolic $|\beta|$ in *mdx* heart may reflect that cardiomyocytes are not fully relaxed in diastole. We also confirmed that this regional diastolic sheet dysfunction in *mdx* hearts is a likely consequence of cytosolic calcium mishandling, because the regional diastolic sheet dysfunction is quickly reversible upon perfusion with a lower level of calcium (Figure 13 A, *mdx*-NC vs *mdx*-LC).

Chapter 3: Anisotropy information in DTI

3.1. Introduction

In Chapter 2, I have used orientation information (β) derived from DTI to detect the basal myocardial function abnormality. In this Chapter, I sought for the first time to further explore the shape information with fractional anisotropy (FA) and mode of anisotropy (mode) to elucidate regional abnormal myocardial fiber/sheet shape in *mdx* mice hearts. Please refer to Chapter 2 for DTI background and methods. Refer to Advanced Engineering Mathematics by Erwin Kreysiz (9th ed) for tensor matrix properties.

Each diffusion tensor can be decomposed into three eigenvectors and three corresponding eigenvalues. Eigenvectors carry orientation information, while eigenvalues provide anisotropic information for the micro-diffusion environment in the voxel (Bahn 1999). A tensor invariant is a scalar value calculated from eigenvalues. For examples, trace and norm are wildely used tensor invariants in DTI:

 $trace = \lambda_1 + \lambda_2 + \lambda_3 = 3 \times ADC$, where $\lambda_{1\sim 3}$ are eigenvalues

.....eq10

$$norm = \sqrt{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}$$

.....eq11

In an eigenvalue space of a 3-D Cartesians coordinate system, norm is the distance between the origin and the point, $(\lambda_1, \lambda_2, \lambda_3)$. Trace is the sum of the coordinates. The trace is three times of the quantity of the apparent diffusion coefficient (ADC), which is an effective indicator of stroke-sites in brain (van Gelderen, de Vleeschouwer *et al.* 1994; Latt, Nilsson *et al.* 2009). The norm is a measure of total magnitude of diffusion (Grant 1965). Trace and norm are both in the units of mm^2/sec , the same as the unit of eigenvalues.

For example, fractional anisotropy (FA) and mode are two independent scalar values calculated from eigenvalues (Figure 17). FA has been widely used in many DTI measurements (Pierpaoli and Basser 1996), especially in the brain (Papanikolaou, Karampekios et al. 2006). Recently, FA has been applied to delineate the contraction-induced assumption of more isotropy for tissue architecture in skeletal muscle (Galban, Maderwald et al. 2004; Deux, Malzy et al. 2008). Mode was proposed in 2006 as an FA-independent metric for anisotropy (Ennis and Kindlmann 2006). Mode has been used to detect microstructural anatomy of epidermoid cysts and nerve fibers (Jolapara, Kesavadas et al. 2009; Douaud, Jbabdi et al. 2011). However, dynamic changes in myocardial architecture for genetically abnormal dystrophic hearts have not been investigated with the use of FA or mode. Although FA is more widely used in DTI, we propose to include also mode to describe the anisotropy in myocardium, because "No single scalar parameter can be constructed which reflects all of the information about anisotropy which is contained in the eigenvalues" (Bahn 1999). The 3D information acquired in perfused viable mdx hearts should allow delineation of both the locus (regional and intramural) and the cycle dependence (diastole vs. systole) of the principle cardiac defects in dystrophin-deficiency and whether these defects might be reversible (i.e. functional) consequences of calcium mishandling

T.100 -0.87 -0.50 0.00 0.50 0.87 1.00

(please refer to Chapter 4) or be nonreversible consequences of fibrosis (i.e., remodeling).

Figure 17. Demonstration of anisotropic space spanned by FA and Mode. From (Ennis, 2006)

41

3.2. Method: DTI Data Analysis for FA and mode

Please refer to Chapter 2 for isolated Heart Preparation for DTI and MRI procedure. The dataset is the same as acquired in Chapter 2. However, new signal processing method was used to extract information from the dataset. All MR images were processed using custom-developed software written in Matlab. FA and mode are both calculated from eigenvalues and eigenvectors of tensors according to the standard formulas below (Ennis and Kindlmann 2006).

$$FA = \sqrt{\frac{1}{2} \frac{\sqrt{(\lambda_1 - \lambda_{mean})^2 + (\lambda_2 - \lambda_{mean})^2 + (\lambda_1 - \lambda_{mean})^2}}{\sqrt{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}}}$$
$$= \sqrt{\frac{3}{2} \frac{\sqrt{(\lambda_1 - \lambda_2)^2 + (\lambda_2 - \lambda_3)^2 + (\lambda_1 - \lambda_3)^2}}{\sqrt{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}}}$$

, where $\lambda_{1\sim3}$ are eigenvalues and λ_{mean} is the average of $\lambda_{1\sim3}$

.....eq12

$$mode = 3\sqrt{6} \det\left(\frac{A_{aniso}}{|A_{aniso}|}\right), where A_{aniso} = A - \frac{\sqrt{(ev_1 + ev_2 + ev_3)}}{3} \times I_3$$

.....eq13

3.3. Result

3.3.1. *mdx* and WT hearts exhibit lower FA & higher mode

In all hearts, systolic FA was ~40% lower than diastolic FA (p < 0.05 for all comparisons, Figure 18), indicating that a more isotropic diffusion environment was elicited by cardiac contraction (Galban, Maderwald *et al.* 2004; Deux, Malzy *et al.* 2008), as expected from fibers shortening in the long-axis and expanding in the short axis (i.e., becoming rounder, or less oblong). Generally, systolic FA is lower (i.e., spherical instead of a ellipsoidal shape) and mode is higher (i.e., oblong instead of a discoid shape, Fig63) than those of diastole (Fig64). In diastole, *mdx*-NC hearts exhibited abnormally low FA and high mode compared to WT, indicating a regional unrelaxed myocardial fiber/sheet architecture. After perfusing the *mdx* heart with low [Ca²⁺] (*mdx*-LC in Fig64) cardioplegic solution, recovery of diastolic FA and mode to normal values were observed.



Figure 18. DTI-determined FA and mode in diastole and systole of WT and mdx mouse heart. N = 5 for each group. *, p < 0.05. NC, normal calcium. LC, low calcium concentration perfusion. See Methods in Chapter 2.

3.4. Discussion – abstract shape information

The low diastolic FA of mdx hearts, indicating more isotropic fiber/sheets organization (Galban, Maderwald *et al.* 2004; Deux, Malzy *et al.* 2008), also demonstrates this diastolic mechanical defect of mdx hearts. Systolic FA of mdx is lower than that of WT, indicating while the mechanical sheet orientation is not affected by the absence of dystrophin in systole, the microstructural environment of muscular architecture is affected. This requires further investigation.

This study shows for the first time that myocardium microstructure of mdx mice exhibits abnormal anisotropy that is calcium dependent, loci-dependent,⁷ and rapidly reversible. Given that barium-induced cardiac contraction leads to lower FA and higher mode, the observed lower diastolic FA and higher diastolic mode in mdx hearts all reflect that mdx cardiomyocytes are not fully relaxed in diastole.

FA and mode describe the anisotropy of water diffusion in the microscopic environment that could represent several factors, including cell shape/size, extra/intracellular volume ratio, myofiber dispersion, and myocardial sheet extension, all of which broadly impact electromechanical cardiac function. The observation that changing cytosolic $[Ca^{2+}]$ regulates the FA and mode in *mdx* hearts suggests that restoration of cell environment by allowing complete relaxation in diastole can be achieved rapidly by transient adjustments in sarcoplasmic calcium levels. Accordingly, diastolic dysfunction in *mdx* mice may be reversible in part with measures aimed at restoring normal calcium handling, and is not a permanent pathophysiological feature of the disease. Thus, FA and mode are quantitative tissue based metrics to assess microarchitectural alterations in cardiomyopathy.

⁷ Loci-dependent: base is abnormal while apex is normal.

Chapter 4: Optical mapping for calcium transients in *ex vivo* viable *mdx* hearts

4.1. Acknowledgements to the contribution of Dr. Efimov's lab

After I have found the calcium-dependent regional sheet dysfunction described in last Chapter, Samuel Wickline and me proposed a collaboration to Igor Efimov's lab to test the hypothesis of regional abnormal calcium handling in a whole perfused mouse heart. I thank Di Lang, Chaoyi Kang, and Igor Efimov for their generous help on optical mapping for studying the calcium transients of 16 month old (old) and 3 month old (young) *mdx* mice. Di found that the basal calcium transients are abnormally prolonged only when the heart is stretched in old but not young *mdx* mice, for the first time. Combining Di's optical mapping findings with my DTI findings, we were able to infer that *mdx* cardiomyopathy is calcium- and loci- dependent (Cheng, Lang *et al.* 2012).

4.2. Why use optical mapping for calcium?

 Ca^{2+} released by sarcoplasmic reticulum (SR) in cardiac muscle triggers conformation change in troponin, allowing the binding between actin and myosin and the following sliding action, and results in macroscopic contraction of cardiac muscle. The Ca^{2+} is released from the openings of SR Ca^{2+} release channel (ryanodine receptor, RyR). The voltage-gated sarcolemmal L-type Ca^{2+} channels (dihydropyridine receptors, DHPRs) locally increase [Ca²⁺]i and thereby activate the RyRs by Ca^{2+} -induced Ca^{2+} release (CICR). After contraction, sarco/endoplasmic reticulum Ca^{2+} -ATPase (SERCA) uptake the cytosol Ca^{2+} into SR, allowing relaxation (Cheng, Lederer *et al.* 1996). This CICR and Ca^{2+} recycling triggers every heart cycle. How cardiomyocytes handle calcium is critical to both normal and pathological hearts. Chapter 2 suggests mdx mice exhibit calcium-dependent sheet dysfunction at base, leading us to hypothesize that the base handles calcium differently than the apex. We employed optical mapping to investigate calcium handling of mdx mouse hearts for the first time.

Optical mapping of calcium transients in ex vivo, viable, and intact heart (simply "optical mapping" hereafter) uses a camera to capture the time-varying fluorescence signals of calciumsensitive dyes. Efimov's group pioneered the technique two decades ago (Efimov, Huang *et al.* 1994). The optical mapping technique is a powerful tool to investigate cardiac calcium transients in human, dog and mouse (Fedorov, Chang *et al.* 2010; Lang, Glukhov *et al.* 2011; Cheng, Lang *et al.* 2012). However, one needs to either open the chest or prepare an *ex vivo* Langendorff perfused heart to provide access the heart to optical cameras.

For interested readers, the best publication introducing the setup of optical mapping is on *The Journal of Visualized Experiments* (Lang, Sulkin *et al.* 2011), which is also from Efimov's lab. It utilizes online video streaming to explain the complex setup efficiently and clearly.

http://www.jove.com/video/3275/optical-mapping-action-potentials-calcium-transients-mouse

4.3. Method

4.3.1. Isolated heart preparation and calcium dye loading

Seven Langendorff-perfused hearts (Lang, Glukhov *et al.* 2011) from 16 month old mice were separated into two groups (*mdx*, N=4; and WT(C57BL/6), N=3). In brief, after anesthesia and heart excision, a short section of aorta was attached to a 21-gauge cannula. After cannulation, hearts were superfused and retrogradely perfused with oxygenated (95% O_2 , 5% CO_2) constant-

temperature ($37\pm1^{\circ}$ C), modified Tyrode solution (in mmol/L: NaCl 128.2, KCl 4.7, NaH₂PO₄ 1.19, MgCl₂ 1.05, CaCl₂ 1.3, NaHCO₃ 20.0, and glucose 11.1 (pH=7.3)) that was passed through a 5 µm filter (Millipore, Billerica, USA). Perfusion was performed using a peristaltic pump (Peri-Star, WPI, Sarasota, USA) under a constant aortic pressure of 60 – 80 mm Hg.

The isolated heart was pinned at the edge of ventricular apex to the Sylgard bottom of the chamber to prevent stream-induced movement. A small silicon drain tube was inserted into the left ventricle through the pulmonary vein, left atria, and tricuspid valve to prevent solution congestion and subsequent ischemia. A piece of small black tape was used to cover the atria to avoid fluorescence scattering from the atria. The excitation-contraction uncoupler Blebbistatin⁸ (10 μ M, Tocris Bioscience, USA) was used to prevent motion artifact on the action potential duration estimation. The heart was then stained with a calcium indicator (Rhod-2 AM, Invitrogen, Carlsbad, CA, 30 μ L of 1 mg/mL dimethyl sulfoxide, 1:1 mixed with Fluronic F127, Invitrogen, Carlsbad, CA in Tyrode's solution), for 5-7 minutes.

4.3.2. Optical imaging system

The excitation light was generated by a halogen lamp (Newport Oriel Instruments, Stratford, CT; SciMedia, Costa Mesa, CA) and passed through a heat filter, shutter, and excitation band-pass filter (520 ± 45 nm). A flexible light guide directed the band-pass filtered light onto the preparation. The optical mapping apparatus comprised an MiCAM Ultima-L CMOS camera (SciMedia, Costa Mesa, CA) with high spatial (100x100 pixels, 230±20 µm per pixel) and temporal (1,000-3,000 frames/sec) resolution. A band-pass filter (590±15 nm, Thorlabs, Newton, NJ) was fixed in front of the calcium-imaging camera.

⁸ Blebbistatin will stop the muscle contraction, allowing motion free calcium transients recording.

4.3.3. Optical mapping protocols

After isolation and cannulation, motion suppression, and dye staining, prepared hearts were equilibrated for 5-10 min before imaging. The optical mapping signals were recorded sequentially under unloaded and stretched conditions. The unloaded condition was achieved by a pressure releasing silicon tube inserted into the LV chamber of the Langendorff heart preparation mentioned above. The stretched condition was achieved by inflating the LV chamber with Tyrode's solution to 80 mmHg pressure through the silicon tube inserted into the left ventricle. The perfusion pressure was increased from 60 to 100 mmHg to maintain the direction of the retrograde perfusion and the viability of the perfused heart. A steady-state restitution (S1-S1)(Lang, Glukhov *et al.* 2011) pacing protocol was applied under both conditions.

4.3.4. Optical mapping data processing

A custom-designed Matlab program was used to analyze the optical signals, which were filtered with a low-pass Butterworth filter at 256 Hz. Regional calcium dynamics were quantified in WT and *mdx* mouse ventricles (base vs. apex) under both unloaded and stretched conditions. Hand-drawn regions of interest were used to define basal (upper 1/3 area of left ventricle) vs. apical regions (lower 1/3 area of left ventricle). The calcium decay signal is not monoexponential and only the latter half of the decay signal could be fitted exponentially to derive the decay constant. Thus, both T50 and decay constant (Figure 19 A) are commonly used to characterize calcium relaxation kinetics (Giordano, He *et al.* 1997; Vassalle and Lin 2004; Zhu, Altschafl *et al.* 2005; Mukherjee, Apple *et al.* 2006; Taylor, Al-Saadi *et al.* 2006; Williams and Allen 2007; Sahoo and Kim do 2008; Sahoo, Kim *et al.* 2009; Chase and Orchard 2011; Mellor, Wendt *et al.*

2012). Specifically, Ca relaxation time from the calcium transient peak value to 50% relaxation (T50) was calculated to define the early calcium reuptake phase; and Ca decay constant was calculated using an exponential curve fitting program from the 50% calcium peak value to ful relaxation. Taken together, the T50 and the decay constant characterize the whole process of Ca reuptake-dependent relaxation.

4.4. Optical Mapping Results

4.4.1. Optical imaging reveals longer calcium decay time (T50) in 16 month old *mdx* mice

Under unloaded conditions, T50 was comparable between mdx and WT hearts. However, ventricular stretch elicited an approximately 25% increase in T50 (from 40 ± 2 ms to 51 ± 3 ms, p < 0.05, Figure 19) and approximately 50% increase of decay constant (from 25 ± 1 ms to 37 ± 1 ms, p < 0.05) at the base of mdx hearts, confirming abnormal calcium uptake in this region. Ventricular stretch, however, did not affect the T50 nor decay constant at the apex of mdx hearts, nor in the entire left ventricle of WT hearts, indicating a heterogeneous regional dependency on the development of abnormal calcium handling in mdx.



Figure 19. Calcium relaxation for mdx and WT. **A**, representative calcium transients at the base of a WT (blue line) and a mdx (red line) hearts. Signals were averaged from adjacent 25 CCD elements. Definitions of T50 and τ are illustrated. **B & C**, representative T50 and decay constant maps in mdx and a WT hearts under unloaded versus loaded (stretched) condition. **D & E**, under unloaded conditions, both T50 and decay constant were comparable between mdx and WT hearts. Ventricular stretch caused regional increases of T50 and decay constant at the base of mdx hearts. However, WT hearts exhibit similar T50 and decay constant. *, p < 0.05; **, p < 0.01.

4.4.2. 3 month old *mdx* mice exhibit normal calcium transients

Surprisingly, despite the fact that mdx mice lack dystrophin since embryo, the prolonged calcium transient abnormality did not exist in young mice (3 month old) whether stretched or not, while abnormal calcium transient exist at the base of old mdx mice (16 month old), only when stretched. Representative figures are shown in Figure 20.



Figure 20. Calcium transient duration at 80% relaxation is only abnormal (red) in 16 month old *mdx* when stretched; but is normal (blue) when young or unstretched. T50 and decay constant exhibit the same pattern (not shown).

4.5. Discussion of regional abnormality at base

The prior recognition of stretch induced calcium mishandling in isolated mdx cardiomyocytes informs the present work (Williams and Allen 2007; Fanchaouy, Polakova *et al.* 2009). Specifically, mechanical stretch increases opening chance of stretch-activated channels (SAC) (Williams and Allen 2007) and induces transient membrane microruptures in mdx cardiomyocytes (Townsend, Blankinship *et al.* 2007; Townsend, Yasuda *et al.* 2007). The resultant leakage of extracellular calcium into the cytosol then compromises calcium clearance during relaxation and renders mdx cardiomyocytes vulnerable to stretch induced injury (Sapp,

Bobet *et al.* 1996). In this study of intact perfused hearts, we employed a stretch protocol to mimic cardiomyocyte activity under moderate strain regimes by inflating LV pressure to 80 mmHg, which is similar to that used for investigating fibrillation by several groups (Nazir and Lab 1996; Bode, Katchman *et al.* 2000; Zarse, Stellbrink *et al.* 2001; Eijsbouts, Majidi *et al.* 2003; Huang, Tai *et al.* 2003). Under the stretched condition, *mdx* hearts exhibited regional deficiency in calcium uptake, as evidenced by the prolonged T50 and decay time constant, at the base where the sheet functional defect was observed. Thus, our results illustrate a regional correspondence between abnormal diastolic sheet mechanics and disturbed calcium kinetics. Because sheet function is a key determinant of ventricular wall strain (Costa, Takayama *et al.* 1999; Usyk, Mazhari *et al.* 2000), and calcium mishandling in *mdx* cardiomyocytes is associated with cellular injury (Lynch 2004; Williams and Allen 2007; Prosser, Ward *et al.* 2011), the observed stretch-induced regional calcium mishandling in *mdx* hearts may be responsible in part for the local progression of diastolic sheet dysfunction.

4.6. Discussion of the perfusion condition in DTI and optic mapping

Results in Chapter 2 and 3 agree that abnormalities exist at the base of *mdx* hearts. However, Chapter 4 suggests the calcium transient abnormalities only when the left ventricle (LV) is *stretched* by 80 mmHg pressure. Why do I see abnormalities at base in Chapter 2 without stretching? Here is a possible explanation: In the DTI study, the LV was not vented, while in the optical mapping study, the LV was vented with a tube inserted in the LV through left atrium. I did not use the venting tube in DTI study, in order to avoid susceptibility artifacts and sliding motion-artifact of the tube. A venting tube in LV creates two surfaces of water/non-water materials and usually brings microbubbles into the LV. The surfaces and microbubbles both generate unwanted susceptibility artifacts of MRI (Sakurai, Fujita *et al.* 1992). Furthermore, the tube tends to slide by few millimeter during the 1.5 hr long DTI study and generate motion artifacts. However, in optical mapping, all the signals were collected from the surface of a heart. Therefore, a tube in the LV does not affect optical imaging but does prevent the congestion of the perfusate and metabolism acidification in the LV, making a venting tube desirable for optical mapping study. In DTI, the metabolism acidification is less worrisome because the heart was arrested in diastole with the cardioplegic solution under 18 Celsius temperature which minimize the metabolism activity. In conclusion, in DTI study, all the hearts are not vented, therefore, the LV might be stretched by the perfusion pressure delivered through aorta valve incompetence or the Thebesian veins, during the 30~45 min long imaging setup time. In the future, a thin pressure transducer can be used to measure the pressure in the unvented LV for prolonged DTI acquisition to clarify this issue.

Chapter 5: Autophagy and Beneficial Effects of Rapamycin-nanoparticle

5.1. Acknowledgements

Experiment were designed by Bibee Kristine, Jon Marsh, Samuel Wickline and me. Richard Killing and I performed strength measurements. Sharron Yang and I performed autophagy nanoparticle treatments. Huiying Zhang and I performed histology of myocardium. Echocardiography ejection fraction were performed and processed by Dr. Jon Marsh. Western blots were performed by Dr. Kristine Bibee and Dr. James Ching.

5.2. Abstract

Current supportive therapy for DMD patients with corticosteroids results in a modest increase in strength as a consequence of a general reduction in inflammation, albeit with potential untoward long-term side effects and ultimate failure of the agent to maintain strength. Here, we demonstrate that rescuing defective autophagy in *mdx* mice, a model of DMD, with the use of rapamycin-loaded nanoparticles (RNP) induces a reproducible increase in both skeletal muscle strength and cardiac ejection fraction that is not achievable with conventional oral rapamycin delivery. *Rapamycin induces autophagy, or "self-eating", which is a cellular process responsible for recycling metabolites and eliminating toxic byproducts.* The resulted increase in physical performance occurs in both young and adult mice, and surprisingly even in aged wild-type mice. RNP treatment of old mdx mice increases autophagic activity with rapid beneficial consequences for improved LV function after only eight doses.

5.3. Introduction

As discussed earlier, DMD now is treated symptomatically with corticosteroids, resulting in a modest increase in strength albeit with expected and unavoidable side effects (Balaban, Matthews et al. 2005; Markham, Spicer et al. 2005; Moxley, Ashwal et al. 2005; Biggar, Harris et al. 2006). However, the cardiac phenotype has proven difficult to modify, and pursuit of combination therapies that might be steroid sparing remains an active area of research (van Deutekom, Janson et al. 2007; Kinali, Arechavala-Gomeza et al. 2009; Cirak, Arechavala-Gomeza et al. 2011; Goemans, Tulinius et al. 2011).

New pharmaceutical strategies other than steroid are desirable. A recent promising treatment target is the *autophagy pathway*, which is a cellular process responsible for recycling reusable products of metabolism and/or efficient elimination of toxic byproducts (Kundu and Thompson 2008). Indeed, rapamycin, a known inducer of autophagy, was reported by Eghtesad et al. (Eghtesad, Jhunjhunwala et al. 2011) in 6-week-old *mdx* mice to improve diaphragm muscle histopathology. Protein restriction has been shown to activate autophagy in 16-week-old mdx leading to enhanced grip strength (De Palma, Morisi et al. 2012). Recent work by Pauly et al. has demonstrated that autophagy induction in the diaphragm of 6-week-old *mdx* animals by daily intraperitoneal injections of an AMP-activated protein kinase agonist also promotes clearance of defective mitochondria and improves diaphragm strength (Pauly, Daussin et al. 2012). However, the potential benefits of enhancing autophagy in multiple muscle groups with the use of clinically practicable therapeutic agents that might elicit rapid and repeatable responses in *both* global physical performance and cardiac functionality in vivo over a broad range of ages (e.g., ≥ 16 months) have not been examined.

Rapamycin, an immune-suppressing macrolide typically used to prevent organ transplant rejection and as an anti-inflammatory agent to prevent angioplasty restenosis (Vilquin, Asselin et al. 1995; Thomson, Turnquist et al. 2009; Adelman), is known to induce autophagy, thereby inciting a cell survival program that enables useful recycling of amino acids via non-specific degradation of long-lived proteins and dysfunctional organelles (Shigemitsu, Tsujishita et al. 1999). By binding to the mammalian target of rapamycin complex 1 (mTORC1), rapamycin blocks proliferative signaling (Wullschleger, Loewith et al. 2006). Paradoxically, blocking mTORC1 with rapamycin might be expected to exert *deleterious* effects on protein synthesis, muscle fiber regeneration, and cell growth in *mdx* mice, all of which might be expected to maintain strength. However, if autophagy is fundamentally defective in DMD/mdx and can be shown to reverse weakness and cardiac dysfunction, and if a method for effective delivery of clinically approved doses of rapamycin to muscle tissues that might enhance autophagy and improve endurance can be designed, then a compelling mechanistic link between disturbed autophagy and progressive deterioration of global physical strength and contractile performance in vivo would be suggested.

Nanoparticle therapeutics for treating cancer and other pathologies are now undergoing clinical trials in multiple drug combinations (Etheridge, Campbell et al. 2013), but heretofore have not been evaluated in neuromuscular diseases. The potential advantage conferred by such agents is their high local accumulation at sites of disease in concert with the use of lower serum levels of free drug that would limit unwanted off targeted side effects, as has been shown for DoxilTM and AbraxaneTM in cancer for example. Accordingly, we compared both an oral preparation of rapamycin and an intravenous nanoparticle-based formulation of rapamycin (RNP) for ability to improve a clinically significant global performance factor related to endurance and

grip strength, with the use of validated methods that are translationally relevant and highly correlated with mortality in the mdx model (Keeling, Golumbek et al. 2007). Furthermore, we measured autophagic responses to rapamycin nanoparticles in skeletal muscle tissues for correlation with grip strength metrics (designed and performed by Jon Marsh and Kristine Bibee). Finally, we delineated the effects of these agents on cardiac function in aged mdx and wild-type mice.

5.4. Methods

5.4.1. Reagents

All chemicals were purchased from Sigma-Aldrich, Inc. (St. Louis, MO, USA) unless otherwise specified.

5.4.2. Nanoparticle formulation

The perfluorocarbon nanoparticle emulsions without or with rapamycin (at a concentration of 0.275 mol%) were prepared as previously described (Cyrus, Zhang et al. 2008). In general the sizes of the PFC NP range between ~160-240 nm (Wickline, Neubauer et al. 2006; Wickline, Mason et al. 2010). As an example, batch MS1-92b exhibited a diameter of 184.7 nm, polydispersity 0.120, and zeta potential -15.31 mV. We note from prior experience that drug loading has little effect on the size but may affect zeta potential depending on the exact composition and charge of the loaded moiety (Soman, Lanza et al. 2008; Soman, Baldwin et al. 2009). An additional, dual-fluorophore emulsion was formulated to verify delivery of both nanoparticles and rapamycin to skeletal muscle. The emulsion lipid moiety incorporated rhodamine B (Avanti Polar Lipids, Alabaster, AL, USA), and the rapamycin was conjugated with a near-infrared (IR) fluorescent dye (Cy7.5, Lumiprobe Corporation, Hallandale Beach, FL) before formulation into the nanoparticles. Perfluorocarbon nanoparticles are exceedingly stable in blood as they were originally developed as a blood substitute (Flaim 1994). The stability of rapamycin incorporated into nanoparticles has been reported previously (Cyrus, Zhang et al.
2008). Drug release kinetics against an infinite sink of releasing medium in vitro verified retention of 97% of nanoparticle rapamycin over three days.

5.4.3. Animal studies

Strength studies: Male *mdx* mice (strain C57BL/10ScSn-Dmd^{*mdx*/J</sub>) and age-matched controls (strain C57BL/10SnJ) were purchased from The Jackson Laboratory (Bar Harbor, ME, USA). All mice were housed in the same animal facility and cared for under approval by Washington University's animal study committee. Starting at age 14 weeks, mice were injected twice weekly for four weeks with plain nanoparticles (n=18 for *mdx*, n=4 for control) or nanoparticles loaded with rapamycin (n=16 for *mdx*, n=5 for wild-type). Dosage was 1 mL emulsion/kg body mass, injected via lateral tail vein resulting in approximately 0.002 mg rapamycin per animal per treatment. An additional set of *mdx* (n=8) mice was housed in an AAALAC-approved facility until they were 17 months of age, at which point they received 4 weeks of RNP treatment as described above.}

Separate groups were utilized for oral rapamycin therapy. For these animals, rapamycin was solubilized in grain alcohol and was delivered in a mixture of dextrose and water twice weekly for four weeks via micropipette gavage, typically 40–50 μ L per mouse. There were three treatment groups: *mdx* mice given rapamycin (n=6); control mice given rapamycin (n=6); and *mdx* mice given placebo (n=6).

Treatment effectiveness was determined by measuring forelimb grip strength within one week before beginning each treatment regimen, and subsequently within one week after treatment completion. Strength testing was performed in blinded fashion by an experienced technician as previously described by our group (Keeling, Golumbek et al. 2007) and by others who have used this standardized test of physical fitness and endurance. Strength testing was

performed using a grip meter attached to a force transducer (E-DF 002 Digital Force Gauge, Chatillon Force Measurement Systems, Largo, FL, USA). The tests relied on the animal's instinct to grab hold as it was pulled backward. Each mouse was grasped by the tail and encouraged to grab a trapeze bar attached to the force transducer using its forelimbs. The mouse was then pulled away and the peak force was recorded on a digital display. This procedure was repeated five times in a row for each animal. The three highest readings were averaged and normalized by the animal's weight to give the strength score.

Rapamycin nanoparticle drug holiday studies: To determine if the effects of RNP on grip strength were reproducible in individual animals, a cohort (n=7 for *mdx*/no drug, n=8 for *mdx*/rapamycin, n=4 for wild-type/no drug, and n=5 for wild-type/rapamycin) of the originally treated 14-week-old mice was maintained off drug therapy over the span of 18-34 weeks of age to allow washout of the RNP effect. The same one month RNP treatment and grip strength testing regimen was then repeated for 1 month starting at age 34 wks.

Cardiac studies: A cohort of 17-month old *mdx* mice received either RNP (n=9) i.v. via eight tail vein injections (1.5 mL/kg/dose), 2x/week over four weeks to deliver the same cumulative amount of rapamycin as for the other studies above or no treatment (n=7). Wild-type mice of the same age were used as controls (n=6 for RNP, n=8 for no treatment). The mice were lightly anesthetized for cardiac ultrasonography with isoflurane and placed supine for examination with a 16 MHz array (Spark, Ardent Sound, Inc., Mesa, AZ, USA). Prior to imaging, the thorax of each animal was shaved and depilated. Animal body temperature was maintained with a heated water blanket and thermistor-controlled heating lamp. Heated ultrasound coupling gel was liberally applied to the shaved area. Waveforms were sampled at the system's clock frequency (66.67 MHz) using a 12-bit digitizer (GaGe Compuscope 12400, DynamicSignals LLC,

Lockport, IL, USA). Two hundred frames of cardiac data were acquired for each view at a frame rate of 152 Hz, with each frame consisting of 128 A-lines (2048 sampled points/line). Each loop thus consisted of between 8 and 12 heart cycles. Radiofrequency (RF) data were converted to scaled image data using custom plugins written for the open-source image processing software ImageJ (Rasband, W.S., U. S. National Institutes of Health, Bethesda, MD, USA, http://imagej.nih.gov/ij/, 1997-2013). Backscattered ultrasound energy was mapped to grayscale by taking the log of the sum of the squares of the digitized voltage values in a center-weighted moving window. The resulting image loops were scaled to a resolution of 0.05 mm/pixel. Left ventricular ejection fraction (LVEF) was computed using ImageJ by manually tracing the ventricular endocardium border at systole and diastole in the short-axis (SAX) view for each beat and by recording the area region of interest (ROI). Ventricular length was computed from the long-axis (LAX) view by determining the length of a line manually drawn from the apex endocardium to the aortic valve plane. Left ventricular volume at systole (or diastole) was estimated by computing the volume of an ellipsoid having a circular cross-section equal to the mean systolic (or diastolic) ROI area and length equal to the mean systolic (or diastolic) measured ventricular length. LVEF was then computed as the ratio of the difference in volume of this ellipsoid between diastole and systole, and the diastolic volume.

5.4.4. Histology

Fibrosis imaging was performed in the diaphragm by analyzing trichrome stained (Masson), formalin-fixed sections (8-µm thickness) with a stain kit (HT-15-1KT, Sigma-Aldrich) and in the heart with picrosirius red stains per standard protocol (Dolber and Spach 1987). Color light microscope images of collagen were acquired from five randomly selected fields for each

diaphragm section at 200X with an Olympus BX61 microscope and F-View camera (Olympus America Inc., 3500 Corporate Parkway, Center Valley, PA, USA). Similar assessments were performed for basal heart sections from 5 treated and 5 untreated *mdx*. Custom software (written as a plugin for ImageJ) was used to quantify the area of each image associated with collagen-labeled pixels relative to the area of each image associated with tissue of any type.

For fluorescence histologic imaging, 14-week old *mdx* mice were given intravenous injections (1 mL/kg) of dual-labeled fluorescent nanoparticles and sacrificed 4 hours post-injection following systemic saline perfusion to wash out any residual circulating nanoparticles. Nanoparticle and/or rapamycin tissue distribution was demonstrated in the biceps, diaphragm, gluteus, and myocardium of mice injected with fluorophore-loaded emulsions in excised and frozen (OCT medium: Tissue-Tek, Sakura Finetek, Torrance, CA, USA) or paraffin-embedded tissues. Sectioning (8-µm) was performed on 10 consecutive slides from each tissue sample (a total of 60 slides were prepared), and random sections were chosen by a blinded observer for fluorescence microscopy. Adjacent slides were saved for H&E. Multiple images were acquired from each section with an Olympus BX51 fluorescence microscope image system, using an Infinity3 camera (Lumenera Corporation, Ottawa, Ontario, Canada) for optical image recording and digitization, and F-View II B&W CCD camera for fluorescent images.

Tissue nanoparticle biodistribution

¹⁹F perfluorocarbon-core nanoparticles were administered to 14 *mdx* mice (18 weeks old) by tail vein injection; and tissue and organ harvesting conducted at either 2, 12, or 24 hours. Tissues harvested for measurement of nanoparticle concentrations were acquired after in situ

perfusion fixation of the mouse that was initiated by cardiac puncture and flow-through of buffer solution until venous effluent was clear (~10 min). This washout procedure ensured that blood contamination was not a factor in the estimation of tissue nanoparticle content. We also note that the tissue distributions and clearances for similar particles are not affected by drug loading and have been reported before for mice and rabbits (Hu 2007; Soman, Lanza et al. 2008; Soman, Baldwin et al. 2009). The specimens were blotted dry, weighed, fixed in formalin, and then subjected to fluorine spectroscopy with the use of an 11.7T Varian MRI/MRS scanner and a custom-built single-turn solenoid coil. ¹⁹F spectra signal from each specimen and from a standard of perfluoro-15-crown-5 ether were obtained via non-localized 90-FID spectroscopy (2.5-s pulse repetition time TR, and 128, 256, or 512 signal averages NT). The unique and quantitative fluorine signature was expressed as total tissue fluorine content per gram. This methodology fortunately avoids the necessity for extraction of the target compound (perfluorocarbon in this case) with its attendant inefficiencies, since the total amount of ¹⁹F signal in the entire unprocessed tissue can be registered nondestructively. Because there is a direct and linear relationship between the ¹⁹F signal and the nanoparticle concentration, the analytical method yields essentially the nanoparticle concentration per unit tissue mass (Morawski, Winter et al. 2004).

Creatine Kinase

Blood was obtained by left ventricular cardiocentesis, and serum extracted by centrifugation. Serum creatine kinase (CK) levels were measured spectrophotometrically with a Vitros DT60 dry reagent chemistry analyzer (OrthoClinical Diagnostics, Raritan, NJ, USA).

Autophagy and autophagic flux measurements

Muscle tissues were extracted from the *mdx* mice and homogenized in 0.5 mL of RIPA buffer with 1-mM PMSF, Complete Protease Inhibitor Cocktail (Roche, Indianapolis, IN, USA), and Phosphatase Inhibitor Cocktail (Roche) using a glass grinding tube. The tissue homogenates were centrifuged at 10,000xg for 10 min at 4°C and the supernatant was stored at -80°C.

Proteins in tissue homogenates were resolved on a NuPAGE Novex 4-12% Bis-Tris Gel (Invitrogen, Carlsbad, CA, USA), transferred to 0.2 μ M nitrocellulose, blocked in 5.0% milk/TBS-T and incubated with the following primary antibodies: rabbit anti-pS6 (1:1000, Cell Signaling 4857, Boston, MA, USA), rabbit anti-S6 (1:1000, Cell Signaling 2217), rabbit anti-LC3B (1:2000, Sigma-Aldrich), rabbit anti-Beclin1 (1:1000, Cell Signaling), goat anti-actin (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA). Corresponding secondary antibodies were used at 1:5000. Protein bands were visualized using either ECL Western Blotting Substrate (Pierce 32106, Rockford, IL, USA) or Super Signal West Femto Maximum Sensitivity Substrate (Pierce 34095). Western blot image was used to calculate relative protein expression using densitometry with ImageJ. Values were normalized to myosin loading control and reported as fold change compared to the control, non-treated sample.

To illustrate how this approach might apply to actual primary human cells and be potentially generalizable, tumor cells (HIM3 line from the Washington University HAMLET Core) were derived from a biopsy specimen that was procured from a triple negative primary breast tumor and were passaged through immunodeficient animals in a method described previously (Ding, Ellis et al.). Cells were maintained in McCoy's 5A media (Invitrogen, Carlsbad, CA) supplemented with 10% FBS (Sigma, St. Louis, MO) and penicillin/streptomycin (Invitrogen, Carlsbad, CA) in a 37° C humidified tissue culture incubator with 5% carbon dioxide. Lentivirus containing the LC3-GFP fusion was obtained from C. Weihl (Washington University, St. Louis, MO). The cells were transfected using a standard spinfection method with protamine sulfate. After recovery, the cells were sorted on a BD Aria II High Speed Cell Sorter three times with one-week recovery periods between each sort. The stable cell lines were then treated for 24 hours with DMSO (vehicle for rapamycin), rapamycin (100nM), nanoparticles (3.09x10¹⁰ particles/mL), or rapamycin-nanoparticles (3.09x10¹⁰ particles/mL = 100nM rapamycin). After treatment, the cells were washed, fixed with paraformadehyde and imaged with a Zeiss LSM510

microscope. GFP-positive puncta were counted by a blinded observer. Other treated cells were prepared for transmission electron microscopy per standard protocols

For *in vivo* autophagic flux assessment, we utilized a recently developed autophagic flux assay that measures turnover of LC3B-II in mouse skeletal muscle by comparing mice treated with and without a potent blocker of autophagic protein degradation (colchicine) (Ju, Miller et al.). Colchicine was administered over 48 hours in two doses (0.4 mg/kg) delivered intraperitoneally, followed by diaphragm or heart tissue collection. Tissue homogenates were probed as above.

A separate set of experiments using oral prednisolone as the active agent was conducted to delineate potential mechanistic similarities between RNP and steroids with respect to their effects on autophagy for an agent that is known to improve strength experimentally and in patients. Four *mdx* mice were treated with oral prednisolone and four control *mdx* were treated with vehicle for 42 weeks (starting at age 4 weeks), after which muscle tissues [tibialis anterior (TA)] were extracted for autophagy measurements as above.

Statistics

Statistics were generated as noted below for selected ANOVA, t-test, and groupwise comparison procedures using SAS software (SAS Institute, Cary, NC, USA) and the open-source statistical package R (R Foundation for Statistical Computing, Vienna, Austria, <u>http://www.R-project.org</u>).

5.5. Results

5.5.1. Rapid enhancement of global physical performance with RNP

None of the fourteen-week-old mdx and wild-type mice treated with nanoparticles demonstrated apparent adverse effects from the treatment at any time point and all animals

completed the study. Figure 21a shows that grip strength increased significantly when RNP were employed as therapy (30% absolute increase in mean strength of group). Unexpectedly, when nanoparticles (NP) alone were employed as therapy, grip strength exhibited a trend toward a modest strength increase (Figure 21a) of 8%.

To determine whether the grip strength improvement was due to the drug alone, animals were treated with oral rapamycin administered at 10x the equivalent nanoparticle i.v. dose to account for an expected 10% oral absorption rate (Napoli, Wang et al. 1997; Crowe, Bruelisauer et al. 1999; Stepkowski 2003; O'Reilly, McSheehy et al.). As shown in Figure 21a, the oral rapamycin did not improve grip strength. Age-matched wild-type mice exhibited no significant increase in grip strength when treated with rapamycin-loaded nanoparticles or oral rapamycin (Figure 21b).

Figure 21c shows that over this washout interval, strength declined as anticipated. After the repeat dosing of i.v. RNP, grip strength once again improved in the older *mdx* animals (p=0.008, linear contrast model). No significant trend was observed in wild-type mice (Figure 21d). Interestingly, RNP treatment of a cohort of 17-month-old *mdx* (n=8) mice treated for a month in similar fashion revealed a more muted effect on strength (Figure 22a: p = 0.201, two-tailed paired t-test).



Figure 21. Rapamycin-loaded nanoparticle treatment improves strength in mdx mice.

a) Four weeks of treatment with rapamycin-loaded nanoparticles (RNP) delivered intravenously significantly increased mean weight-normalized grip strength in 14-week old *mdx* animals (n=16) vs. groups given plain nanoparticles (NP) IV (n=18), equivalent oral doses of rapamycin (n=6), or oral placebo (n=6) (p=0.012, p=0.027, and p=0.016, respectively, using Fisher's protected least significant difference at 5% significance level). b) No significant difference in mean strength was observed for age-matched wild-type mice given either NP (n=4) or RNP intravenously (n=5) or oral rapamycin (n=6). c) The increase in weight-normalized strength in a subset of *mdx* mice (n=8) treated with IV RNP occurred in both young (14-18 wk) and old (34-38 wk) animals (p=0.008 using a linear contrast model comparing pre- to post-treatment differences at a 5% significance level). Absolute values of the mean change in strength between time points are shown in columns below trend data. d) Wild-type mice (n=5) given a similar longitudinal treatment and drug-holiday exhibited no significant difference between pre- and post-treatment strength (p=0.598). All bar graphs show mean values +/- s.e.m.



Figure 22. Aged *mdx* mice show muted responses in grip strength after treatment with RNP, but marked improvement in cardiac function commensurate with wild-type animals

a) Four weeks of treatment of 17-month-old mdx (n=8) yielded no significant increase in grip strength (p=0.201, two-tailed paired t-test). **b**) LVEF in wild-type mice is greater than mdx before treatment (p=0.016), as expected. **c**) Cardiac function significantly improved after RNP therapy, irrespective of animal type (16% difference for mdx and 9% difference for WT; p=0.024).

5.5.2. Rapid enhancement of cardiac performance with RPN in aged mice

In the cohort of 17 month-old mdx and wild-type mice, one month of biweekly RNP versus no treatment yielded two compelling findings: 1) LVEF in *mdx* was significantly lower than that of wild-type mice before treatment (p=0.016) as expected (Figure 22b); and 2) LV ejection fraction for RNP treated mice was 57 ± 2.9 %, and approximately 16% higher than that of none-treated mdx mice (49 ± 2.5 %; p=0.024).

5.5.3. Nanoparticle tissue distribution

Results of fluorescence microscopy of fluorophore-labeled rapamycin and fluorophorelabeled lipids contained in the outer nanoparticle shell are shown in Figure 23. It is clear that the particles penetrate heart tissue to deliver rapamycin locally (Figure 23). The labeled rapamycin appears to achieve an even greater penetration into tissues than do the nanoparticles, arguing for some degree of local rapamycin release and more complete passive diffusion. We note that the use of the near-IR fluorophore avoids the problem of background autofluorescence, thereby permitting confident assessment of the more diffuse penetration of rapamycin into muscle tissues beyond the loci of nanoparticle deposition. The observation of diffuse rapamycin penetration into tissues beyond the nanoparticle carriers (Figure 23) was further supported by analyzing both blank slides and unstained muscle tissue specimens that were microscopically digitized at the same settings to rule out background signals of any type that might have overlapped with the rapamycin signal, and none were observed (data not shown).



Figure 23. Nanoparticles and rapamycin are delivered to cardiac muscle in *mdx* animals.

Nanoparticle lipid (rhodamine, displayed as blue) and rapamycin (Cy7.5, labeled as red) are colocalized in *mdx* cardiac tissue.

5.5.4. Effects on skeletal and cardiac mTOR and autophagic processes

To further demonstrate that the drug reaches muscle tissue and then acts on a known target, lysates from the muscle were probed for phosphorylated S6. S6 is downstream of mTORC1, so that blocking mTORC1 with rapamycin should cause a decrease in pS6 levels. Indeed, in diaphragm, triceps, biceps, and gluteus, S6 levels were decreased in *mdx* animals treated with rapamycin-loaded nanoparticles when compared to mice treated with nanoparticles alone (Figure 24a). Interestingly, the nanoparticles themselves caused a modest *increase* in pS6 levels when compared with saline treated animals. Thus, the particle itself may induce signaling through mTORC1, but adding rapamycin to the particle then decreases the level of signaling, yet does not cause a return to baseline levels.

Our data confirm that the *mdx* mouse harbors a defect in autophagy in muscle tissues, as there is an increase in p62 protein (Figure 24b) with no difference in p62 transcription (Figure 24c). This expression pattern suggests that p62 is not being consumed in the process of autophagy in the *mdx* animal as it is in the wild-type animals. Further, BNIP3, a protein that is known to trigger autophagy, is found at lower levels in *mdx* animals when compared to wild-type (Figure 24b, c). Additional in vivo confirmation of the effects of rapamycin nanoparticles on autophagy revealed that basal autophagic flux was diminished in *mdx* mouse heart as demonstrated by comparing the change in LC3II protein levels with and without colchicine treatment (Figure 24d).



Figure 24. *mdx* animals exhibit a defect in autophagy that is restored by nanoparticle treatment.

a) RNP cause a decrease in S6 levels in *mdx* animals when compared to NP treatment. However, compared to saline treatment, the NP increased S6 phosphorylation in all muscle groups tested, a trend not observed in the muscles from wild-type animals. **b)** Proteins were extracted from *mdx* and age-matched control animal muscle. Western blot analysis for p62 (n=5 per group) and BNIP3 (n=3 per group) reveal a defect in autophagy as *mdx* animals exhibited higher levels of p62 and lower BNIP3 when compared to wild-type (Student's t-test at 5% significance level). Myosin was used as the loading control. **c)** RNA was isolated from the tibialis anterior muscle of age matched *mdx* (n=5) and control mice (n=3). QPCR was performed and normalized to GAPDH. Graph demonstrates fold change in the following transcripts vs. control: there was no change in p62, BNIP3 or beclin. Beta 2 microglobin is increased as expected, because it is part of MHCI, which is known to be upregulated in *mdx* muscle. Actin levels (normal) serve as an internal control. **d)** Representative Western blots demonstrate that both nanoparticle (NP) and rapamycin-loaded nanoparticles (RNP) increase the levels of LC3B-II in the *mdx* animal, whereas saline treated animals exhibit a low level of LC3B-II expression, even when blocked with colchicine. Scatter plots show results from individual animals.

Additional probing of the tibialis anterior (TA) and heart muscle in the 17 month-old mice treated for 1 month with RNP was performed to assess the autophagic responses. Figure 25a illustrates significant effects of RNP in TA on p62 (decrease), BNIP3 (increase), and LC3II (increase). Figure 25b confirms that RNP also enhanced autophagy in the heart in the same manner. Figure 25c shows autophagic responses to prednisolone in the TA of mdx mice, where changes in p62, BNIP3, and LC3II are all similar in direction and magnitude to those reported above for RNP, further suggesting that autophagy responses to either RNP or steroids in the mdx represents a likely candidate mechanism for the observed augmentation of muscle strength and cardiac function. Taken together, these data point to a rapid and widespread induction of autophagy in diverse muscle tissues after RNP treatment.



Figure 25. Rapamycin nanoparticles alter autophagic biomarkers in heart and skeletal muscle of *mdx* mice similar to that after steroid therapy.

a) Tibialis anterior (TA) muscle in 17-month-old *mdx* mice treated for 1 month with RNP exhibited significant effects of RNP on p62 (decrease), BNIP3 (increase), and LC3II (increase).
b) Cardiac data confirms that RNP also enhance autophagy in the heart in the same manner. c) Young *mdx* mice given 42 weeks of steroid therapy show similar autophagic responses in the TA muscle.

5.5.5. Tissue structural responses

The amount of collagen within skeletal and cardiac muscles (Figure 26a-d) was not different between the treatment groups (p=0.900 for diaphragm using mixed model nested ANOVA, p=0.664 for heart using unpaired two-tailed t-test). Figure 26e demonstrates no significant difference between creatine kinase (CK) levels in saline treated animals when compared to those treated with NP or RNP (p=0.682, one-way ANOVA). From these data, it does not appear that the nanoparticles are exerting their beneficial effect on strength by altering the turnover of myocytes or the replacement of myocytes with collagen in the short term, as might be expected after only 8 doses of therapy.



Figure 26. Intravenously injected nanoparticles do not affect short-term muscle cell destruction or fibrosis.

a) Masson trichrome stains performed on diaphragm sections and visualized with light microscopy (200x) showed increased fibrosis (blue) for *mdx* animals relative to WT. **b**) No significant difference occurred over the short 4 week test interval in *mdx* collagen levels after treatment with either rapamycin-loaded nanoparticles, plain nanoparticles, or saline (mixed model nested ANOVA, p=0.90). c) Picrosirius red stains of *mdx* mice show diffuse heterogeneous fibrosis with d) no significant difference (p=0.664) when treated with RNP. e) Serum creatine kinase levels for *mdx* mice exhibited no significant difference after treatment with either RNP, NP, or saline (ANOVA, p=0.71). All bar graphs show mean values +/- s.d.

5.6. Discussion

The present findings indicate that rapid improvement in physical performance of either skeletal and cardiac muscle can be achieved after only 8 doses of RNP in conjunction with

augmented autophagy. In contrast, oral rapamycin in pharmacological doses exerted no significant effect on strength within the same interval. The observations that the improvement in grip strength is repeatable after a drug holiday in the same mice, together offer further support for the salutary role of RNP on enhancing autophagy, global physical performance, and cardiac function in muscular dystrophy. The muted response to strength testing in aged mdx as compared to the clear enhancement in cardiac function reveals a not unexpected limitation on what might be achievable after initiation of therapy late in the disease process.

Recently published work suggests that autophagy induction may be beneficial to *mdx* animals, whether induced by rapamycin or by dietary protein reduction (Pauly, Daussin et al. 2012). Moreover, a decrease in autophagic flux in vivo has been previously reported for an animal model of Bethlem myopathy and Ullrich congenital muscular dystrophy due to mutations in Col6a-/suggesting disturbed autophagy might other that occur in muscular dystrophies (Grumati, Coletto et al.), yet this mechanism has not been evaluated previously for aged DMD/mdx, as we show here. Furthermore, no prior studies have demonstrated improvements in global physical performance in vivo or cardiac function after rapamycin therapy in aged animals that might be attributable to enhanced autophagy.

The observation of a muted response to short term RNP in the older 17-month-old mice for skeletal muscle performance in the face of a clear improvement in myocardial LVEF and clear induction of autophagy points out certain divergent pathophysiological features of the disease. It is recognized that the heart disease in DMD appears later than the skeletal muscle weakness, but emerges ultimately as a major source of morbidity and mortality. Why heart muscle is relatively more resistant to disease progression than is skeletal muscle despite continuous contractile

activity remains unknown. Interestingly, the disease in the heart evolves *regionally* (e.g., base>apex) despite the ubiquitous loss of the dystrophin (Chapter 2~3).

Although recent retrospective analyses of steroids in DMD are highly suggestive of a beneficial long-term effect on cardiac function (Schram, Fournier et al. 2013), the present data are the first to illustrate such rapid improvement with systemic rapamycin in a prospective trial invoking a unique mechanism of action. Parenthetically, our observation of improved LVEF even in wild type subjects (Figure 22) accords with a recent report by Flynn et al. noting better LVEF in 24 month-old female nondystrophic animalds treated continuously for 3 months with rapamycin in an oral "high delivery" formulation (Flynn, O'Leary et al. 2013). The attenuated response of skeletal muscle strength to RNP in older mdx in our study suggests either that longer term or more aggressive therapy might be required at this stage to enhance strength, or that certain limitations may exist for this (or even any) form of therapy, neither of which is especially surprising. Interestingly, Flynn et al also reported that oral rapamycin increased endurance but not *intensity* of activity in 24 month-old female mice, suggesting a muted effect on strength improvement in older subjects in general, which accords with our observation on grip strength for the 17-month old dystrophic mice (Flynn, O'Leary et al. 2013). Regardless, the ability of RNP to rapidly improve cardiac performance suggests that a single agent might be useful.

The cumulative dose of rapamycin administered to the animals in the rapamycin-loaded nanoparticle treatment group was within the limits of recommended oral doses for patients receiving immunosuppressive therapy when accounting for ~20% oral absorption in humans (Ferron, Conway et al. 1997). The rationale for comparing intravenous nanoparticle rapamycin

therapy with oral rapamycin delivery stems from the conventional practice of oral delivery as standard of care in patients requiring immune suppression, which is known to be effective and well tolerated at the recommended dose levels. We accounted for the well-known and previously reported bioavailability for oral dosing (rough estimates: ~10% in mice (Napoli, Wang et al. 1997; Stepkowski 2003; O'Reilly, McSheehy et al. 2010) by augmenting the doses that were delivered to mice by oral gavage to ensure adequate delivery. Typical clinical maintenance dosing for oral delivery of sirolimus in organ-transplant scenarios for body mass greater than 40 kg is 2 mg/day (Rapamune, Pfizer Inc, Philadelphia, PA). Assuming 20% bioavailability, this equates to approximately 0.4 mg of rapamycin in circulation per day, or approximately 0.006 mg/kg/day for a 70 kg patient. The maximum permissible daily dose of rapamycin is 40 mg, which equates to 0.11 mg/kg/day. Mice treated with RNP i.v. in the current study were given 0.067 mg/kg rapamycin biweekly for a total of 8 times over 4 weeks, or the equivalent of 0.002 mg/kg/day, within the clinical bounds for regular therapeutic use.

The intravenous dose used here was approximately 7-fold less than the calculated absorbed oral dose administered to *mdx* mice by Eghtesad and colleagues to improve muscle histopathology (Eghtesad, Jhunjhunwala et al.). Moreover, Eghtesad also reported that oral or injected rapamycin exerted no affect on the TA muscle, in contrast to the marked upregulation of autophagy reported here in the TA after RNP therapy (see Figure 26). Taken together, these data indicate that at clinically relevant doses of oral rapamycin for *mdx* animals, which scale up to 4–fold greater than the recommended oral dose for patients, no improvement in skeletal muscle strength might be expected. In contrast, a clinically approved cumulative dose of rapamycin administered in NP formulations rapidly elicited marked improvement in both global physical

performance and cardiac function after only eight i.v. treatments, which could represent an advantage for periodic, short-term dosing.

Furthermore, nanoparticles alone did not increase autophagic flux in wild-type mice whereas rapamycin nanoparticles did. While rapamycin is known to promote autophagy via an mTOR dependent mechanism, the mechanism by which nanoparticles influence autophagy is not known. It is intriguing that nanoparticles were unable to enhance autophagy in wild-type mice with presumably normal sarcolemmal structure whereas in *mdx* mice that have defective sarcolemmal integrity due to dystrophin mutations, nanoparticles were effective.

Rapamycin also exhibits mTORC1-independent effects when it binds to FKBP12 on the ryanodine receptor. Previous studies suggest that binding of rapamycin disrupts the ryanodine receptor's functions and therefore would be expected to elicit dysfunctional calcium handling (Ahern, Junankar et al. 1997; Avila, Lee et al. 2003). However, in *mdx* animals, the ryanodine receptor itself appears to be dysfunctional (Bellinger, Reiken et al. 2009). It may be the case that the addition of rapamycin acts to modulate some of these receptors in a favorable manner and effect more normal calcium handling or else acts to decrease the expression of the receptors as has been previously reported (Avila and Dirksen 2005).

The effectiveness of nanoparticles and rapamycin nanoparticles for stimulating autophagy in *mdx* mouse muscle mirrors that of their efficacy in rapidly improving grip strength and cardiac function. Although these data may not establish absolute proof of cause and effect for the relationship between autophagy and strength, they suggest an intriguing candidate mechanism for the decline in strength that might be manageable clinically with an approved drug and a new delivery system.

Although the exact mechanism responsible for rapamycin's entry into muscle tissue remains to be clarified, the effect on autophagy is clear and seems dependent on nanoparticle depot delivery, because oral therapy is ineffective at the doses that were administered. The sustained presence of nanoparticles and rapamycin in skeletal and cardiac tissues is clear from the ¹⁹F spectroscopy and the fluorescence imaging data. Prior pharmacokinetic studies in both rabbits and mice using PFC nanoparticles have confirmed a clearance halftime (or, elimination half-life) of ~3-5 hours in normal mice and rabbits used in various cancer models (Hu 2007; Soman, Baldwin et al. 2009), irrespective of molecular targeting, suggesting that there should be ample opportunity to penetrate into the damaged tissues. We previously have demonstrated rapid permeation of these nanoparticles in other situations where leaky or disrupted vasculature pertains, such as advanced atherosclerosis in cholesterol fed rabbits, and even in atherosclerotic human carotid endarterectomy specimens (Cyrus, Zhang et al. 2008; Zhang, Zhang et al. 2011). It is instructive to note that in dystrophic subjects, recent and prior publications have pointed out the widespread vascular dysfunction and damage that permits egress of circulating components into muscle interstitium (Vidal, Ardite et al. 2012). Dystrophic subjects also harbor defects in vascular endothelium that render lining cells susceptible to apoptosis and impaired angiogenesis (Palladino, Gatto et al. 2013), creating a situation of barrier dysfunction that even applies to the blood-brain barrier (Lien, Mohanta et al. 2012).

Regarding the translational potential of this approach, we note that these beneficial effects on both heart and skeletal muscle were achieved at a cumulative dose of rapamycin that accords with clinically recommended dosing. Although the side effects of rapamycin on glucose tolerance, weight gain, blood pressure, and immune responsiveness are recognized, the drug is generally well tolerated over many years by patients with organ transplantation. The fact that both prednisone and rapamycin appear to induce autophagy as a possible mechanism of action raises the possibility that the potential deleterious consequences of either agent might be mitigated if used in lower doses in combination, or at least if RNP could be employed for a steroid sparing action.

Chapter 6: Conclusion

6.1. Summary of major findings

In this study, we observed that:

- 1. Myocardial sheets can be imaged in fresh/unfixed/unstained heart with photoacoustic image microscopy, indicating that sheet structures are not just dehydration artifacts as criticized in earlier histology studies (Zhang, Cheng *et al.* 2012).
- 2. By using Langendorff perfused heart preparations with DTI, viable myocardial sheet function can be imaged without motion-artifact. We observed that sheet dysfunction is manifest in diastole but not in systole, indicating a relaxation problem with mdx hearts (Chapter 2).
- Fibrosis and sheet dysfunction evolve *regionally* in the *mdx* mouse heart at the base despite the lack of dystrophin in every cardiac cell, indicating that all other observed concomitant pathologies contributing mechanistically to the cardiomyopathy are secondary and developmental rather than primary drivers of the disease process (Chapter 2).
- 4. Sheet dysfunction is *reversible* by lowering the perfusate calcium concentration, i.e., *mdx* heart exhibits *calcium* dependent relaxation-failure (Chapter 2).
- 5. At least two tensor invariants are needed to describe the anisotropy of the diffusion tensor, although most related research has reported only fractional anisotropy for reasons of simplicity. Here we demonstrate how the *mode of anisotropy* and *fractional anisotropy* provide complimentary information regarding myocardial architecture anisotropy in mdx mice (Chapter 3).

- 6. Abnormal calcium transients are *stretch activated* and evolve *regionally* first at the base, indicating that the lack of dystrophin triggers calcium mishandling more quickly at the base but not the apex, especially when stretched (Chapter 4).
- 7. Fibrosis, sheet dysfunction, and calcium mishandling are all more severe in older *mdx* mice (16 month old) than in young mice (3 month old).

6.2. Future works:

6.2.1. Intervention for calcium mishandling and autophagy in DMD

In Chapter 2, we found that mdx sheet dysfunction is regional, progressive and calcium dependent. Therefore, we hypothesized that if we correct the calcium mishandling, we could ameliorate sheet dysfunction. This was carried out in whole hearts by adjusting perfusates with good results, which is compatible with other strategies mentioned above to ameliorate calcium mishandling and systolic dysfunction by genetically engineered approaches. In the case of sheet dysfunction that produces diastolic abnormalities early in the course of heart disease, which itself may predispose to further damage by "stiffening" the heart, even earlier metrics of beneficial interventions might be observed via restoration of sheet mechanics. Although steroids are currently the most effective treatment for DMD patients, their efficacy decreases after prolonged treatment (Markham, Spicer et al. 2005). Therefore, cocktail therapies targeting different pathological mechanisms with the use of steroids, current calcium approaches, ACEi, autophagy, and beta-blockers might be desirable in the hopes that gene therapy might emerge as a first line therapy. The proposed DTI measured sheet angle could offer a surrogate for the efficacy of such interventions.

6.3. Remarks on dystrophin function

Here it is instructive to quote from other author's insights into the progressive regional nature of dystrophin-null cardiomyopathy (Judge, Kass *et al.* 2011):

"One hypothesis regarding the cause of cardiomyopathy for individuals with reduced or absent dystrophin is based simply on a structural deficiency. Since dystrophin is part of the DGC, its primary role may be to hold this complex together. When deficient or absent, the dystrophin-associated glycoprotein complex may deteriorate due to the hemodynamic stress associated with normal left ventricular function. Evidence for this includes the loss of several components of the DGC from the plasmalemma when dystrophin is absent. If this were the only contribution of dystrophin to normal cardiac function, then one would anticipate that cardiac dysfunction would be more *uniform* with regard to its age of onset and severity in DMD. ... Another hypothesis for cardiac dysfunction in DMD and BMD posits that dystrophin and its associated complex of ECM glycoproteins play a *regulatory* role."

Then one might ask, what regulatory function might dystrophin have? And furthermore, why would this be different at the base and apex? In fact, nitric oxide, SMAD, or TGF-beta pathways all are reported to be affected by dystrophin-null conditions (Chao, Silvagno *et al.* 1998; Heydemann, Ceco *et al.* 2009; Goldstein, Kelly *et al.* 2011), but it is still unclear what the role of dystrophin in those pathway might be and whether these problems are primary or secondary to myocyte damage. For example, does dystrophin turn on NO pathway to increase coronary flow in response to certain mechanical input? If dystrophin is a switch, what turns it on in response to what stimulus? The function of dystrophin can be discussed from an evolutionary standpoint. First, dystrophin is an ancient protein that exists even in drosophila and sea urchin (Neuman, Kaban *et al.* 2001; Kucherenko, Marrone *et al.* 2011). Morris 1977 found that "*In Xenopus heart, little or no dystrophin was found on Western blots but the dystrophin-related protein, utrophin, was abundant. … In skeletal muscle, however, Xenopus resembles mammals in expressing dystrophin at the sarcolemma and very little utrophin" (Morris 1997). Based on the fact that*

Xenopus, an amphibian, does not express dystrophin in the heart, one could hypothesize that dystrophin was adapted in mammal hearts for some unknown function that is not required in amphibian hearts. Therefore, a comparative study on heart function and dystrophin distribution in base vs apex of the heart could shed lights on dystrophin's mysterious function. It also would be telling to investigate dystrophin's expression in heart development, especially to focus on different locations exhibiting different stress levels in embryological development; and at different development stages such as before and after the foramen ovale closes. It is possible that cardiac dystrophin's regional expression may change dramatically as a marked redistribution of cardiac stress occurs during and after birth, in light that dystrophin protein was not detected in early embryonic days until prenatal stage in rat (Tanaka and Ozawa 1990).

It also is curious why the apex is spared. Recall the Laplace law (eq41).

$$S = \frac{P}{2} \times \frac{r}{h} \qquad \dots \qquad \text{eq41}$$

Stress in the ventricular wall (S) is proportional to transmural pressure (P) and the ratio between radius (r) to wall thickness (h) (Warren W. Burggren 1997). Indeed, Kim *el al* have reported ~30% higher wall stress at the base of anesthetized canine heart (Kim, Min *et al*. 1985). In 1998, using markers and intraventricular pressure catheters, DeAnda *et al* also estimated up to four times higher wall stress in base compared to apex in awake dogs (DeAnda, Komeda *et al*. 1998), agreeing with the prediction made with Laplace law.

Evolution in some respects is an adaptation to stress. 360 Million years ago, fish evolved into amphibians and faced new challenges in a terrestrial environment, including air exchange, dehydration, and *gravity*. In water, animal's body weight is supported by buoyancy, which is not operative on land. Therefore, gravity might contribute to the fact that the largest extant land

animal, the African bush elephant (4.9 tons, 6m)⁹, is 22 times lighter than the largest marine animal blue whales (110 tons, 25.5m)¹⁰. Humans assume that living on the land is simple, but living in the ocean is challenging. But in evolution, the ability to deal with the land environment is *emergent* as opposed to encoded. Indeed, terrestrial animals likely must "pay" for their body size with certain limiting adaptations. Because area-dependent functions increase less rapidly than volume-dependent functions as an organism's size increase, severe scaling constraints may result as land organisms evolved to larger size (Schmidt-Nielsen 1984). For example, assuming no allometry, body weight is proportional to the body length to the power of **three**, while the muscle cross section area¹¹ is proportional to body length to the power of **two**. Therefore, the stress (force/area) in muscle, in order to bear the body weight of the land animal, is proportional to the "body weight divided by muscle cross section area," which is body length to the power of **one**. Therefore, we might assume that a 180 cm tall man experiences **twice** the stress in the calf as does a 90 cm tall man, all else being equal.

It is logical then that the fundamental function of *dystrophin* is **to protect muscle in high stress environment**. Myocardial sheet/calcium transient abnormalities and fibrosis manifest more at base but not apex in 16 month old *mdx* mice (Chapter 2), consistent with higher stress at the base than apex (DeAnda, Komeda *et al.* 1998). Furthermore, myocardial sheet abnormalities and fibrosis do not appear in 3 month old *mdx* mice, which have 15% smaller heart radius. Potentially, the smaller radius resulting in smaller stress spares the heart from abnormalities at

⁹ African bush elephant. Animal Diversity Web.

http://animaldiversity.ummz.umich.edu/site/accounts/information/Loxodonta_africana.html

¹⁰ Blue Whale. The Marine Mammal Center. <u>http://www.marinemammalcenter.org/education/marine-mammal-information/cetaceans/blue-whale.html#.UjIVGcZ6ZRs</u>

¹¹ Vertebrate striated muscle exerts nearly a constant peak tensile stress, ~200k Pa = 2M dyn/cm², among species. Biewener, A. A. (1989). "Scaling body support in mammals: limb posture and muscle mechanics." <u>Science</u> **245**(4913): 45-48.

earlier ages. Therefore, young mice do not exhibit a cardiac, or even much skeletal dystrophic phenotype, despite the fact that all tissues lack dystrophin before birth.

DMD patients (dystrophin-null) experience the onset of symptoms before two year old and become wheelchair bounded between ages 7 and 13. On the other hand, BMD patients (dystrophin-deficient) have much more benign progress of disease. BMD patients have their onset of symptoms between ages 2 to 21, and become wheelchair bound between ages 22 and 51 (Emery and Skinner 1976). That is, the partially expressed dystrophin in BMD patients is still able to protect the increased stress from the growing body size until age 22 to 51. In a 111 DMD patients study, height was found to be correlated with disease progression (disability accessed by the Vignose scale) with a correlation coefficient of 0.352 (p<0.001), indicating that small stature might protect against the ravages of DMD. A classic paper that has been cited 870 times has reported sarcolemma damage is linearly correlated with stress (force/area cross section). That is, mdx diaphragm muscle shows more damage than does wildtype diaphragm muscle (Petrof, Shrager et al. 1993). Similarly, cardiomyocytes of mdx mice are more vulnerable to mechanical stress induced by isoproterenol and aortic banding (Danialou, Comtois et al. 2001) than are normal myocytes. Indeed, anti human growth hormone (HGH) therapy was found to be effective in DMD patients (Zatz, Betti et al. 1986). Four case reports suggested that patients with deficiency in both HGH and dystrophin show slow or no progress of muscular waste until the administered to solve the short stature (Zatz, Betti et al. 1981; Zatz and Betti 1986; Ghafoor, Mahmood et al. 2003; Calcaterra, Malvezzi et al. 2013). However, an HGH inhibitor, mazindol, fails to inhibit the growth over longer times as suggested by Zatz (Zatz, Rapaport et al. 1988):

"However, the recent report of two new DMD patients with short stature and a benign evolution of the dystrophic process (Passos *et al.*, 1988; Wu *et al.*, 1988) and the observation of wide variability in the progression of the disease in DMD/BMD patients with apparently the same DNA deletions suggests that secondary factors, such as growth hormone (GH) may be implicated in the progression of the disease (Infante and Huszagh, 1988; Malhotra et al., 1988)."

That is, Zatz speculated GH itself might have modified the disease. It appears rather more likely that the $GH \rightarrow height \rightarrow stress$ cascade is the operative factor in modifying the disease.

It is known that, in DMD patients, calf muscles fail to perform normal function first, then the diaphragm fails (Bach 1992; Hamada, Ishikawa *et al.* 2011). The cardiomyopathy emerges last (Fayssoil, Nardi *et al.* 2010). It is natural to ask: among the calf, the diaphragm, and the heart, which muscle bears the most stress (force/area)? Does the failure sequence in DMD muscles correspond to the ranking of stress? To the best of my knowledge, there is no study comparing stresses in these three muscles together in DMD nor in mdx mice. However, the following summarizes the existing data.

Thorpe *et al* found normal young adults (~1.7m ~73kg) have peak stresses of 100 kNm⁻² during high jumps and 150kNm⁻² during 4ms⁻¹ running in calf (gastrocnemius and the soleus) (Thorpe, Li *et al.* 1998). Using the reported height of age 8 DMD patients, ~1.4m (Beenakker, de Vries *et al.* 2002), and assuming no allometry and stress is roughly proportional to the height, we can estimate the high jump and **running** stress in a DMD boy are 82 and **123** kN/m², respectively, which should be the upper limit of stress in DMD patient calf, because they have pseudo-hypertrophic calf and less contractility due to fibrosis.

Stress data in calf muscles in a standing posture in normal humans are scanty. However, Morse *et al* have reported isometric maximum stress of calf muscles in young (131 kN/m², age 25, 1.7m, 79 kg) and elderly (92 kN/m², age 74, 1.7 m, 78 kg) normal people (Morse, Thom *et al*. 2005). Taking elderly subjects stress data and assuming no allometry, a DMD patient of 1.4 m height should have an isometric maximum stress of 92*1.4/1.7=75 kN/m² in calf. Although assuming no allometry may not be true, it is the best approximation devoid of additional calf geometry information. I cannot find any literature addressing the fascicle force and physiological cross-sectional area in calf muscles of DMD patients. However, from the above two literature sources, I estimate that standing calf stress in DMD patient is smaller than **75** kN/m².

mdx mice diaphragm strips were reported to exert maximum tetanic stress of 91~136 kN/m² in ex vivo experiment (Lynch, Rafael et al. 1997; Lynch, Hinkle et al. 2001). In vivo measurement of diaphragm stress were not available in literature, to the best of my knowledge, probably because the thin anatomical geometry of the diaphragm limits the observation opportunity. However, Hughes *et al* used two balloons placed in the stomach and esophagus to measure the transdiaphragmatic pressure in normal adult humans and reported a value of 143 cm H₂O (105 mmHg) (Hughes, Polkey et al. 1999). Meanwhile, Beck reported a maximum transdiaphragmatic pressure of 7.6 cmH₂O (5.6 mmHg) in DMD patients with advanced stage disease (N=8, mean age 25) (Beck, Weinberg et al. 2006). Although the in vivo stress of the diaphragm was not reported, it is not unreasonable to assume that the diaphragm exhibits comparable or higher stress than does heart wall, because the thickness of diaphragm (1.74 mm thick in 10 year old DMD patients; 1.45 mm in normal controls (De Bruin, Ueki et al. 1997)) is >3 times thinner than heart wall (4.8 mm in diastole; 8.7 mm in systole (Goldberg, Feldman et al. 1980)). Diaphragm has comparable maximal transmural pressure as with the heart (105 mmHg vs 120mmHg). Meanwhile, the diaphragm has much larger radius of curvature.

Numerical values of stress in normal adult human heart wall are more abundant. Normal wall stress was reported to be $18 \sim 33 \text{ kN/m}^2$ in systole and $2 \sim 3 \text{ kN/m}^2$ in diastole (Hood, Rackley *et al.* 1968; Grossman, Jones *et al.* 1975). In DMD, one study reported 23 DMD patients with

mean age of 8 have a wall stress of **49** (Markham, Kinnett *et al.* 2008). Unfortunately. this paper ignored the units of wall stress.

In conclusion, values from literature suggest that the calf might be stressed more in 1.4m tall boys than the heart (**75** vs **18~49** kN/m^2). I speculate that the diaphragm might experience even higher stresses than the heart because of the comparable transmural pressures in the face of thinner thickness, and larger curvature radius, but there are no available data reported. After reviewing all of the published values of stresses, it would not be surprising if the stress ranking is actually: calf > diaphragm > heart, which might explain the temporal sequence of failing muscles in DMD patients.



Deconinck proposed a damage pathway initiating with the lack of dystrophin (Figure 27) (Deconinck and Dan 2007). However, this simplistic model fails to incorporate the supposition that it might be stress itself that modifies the degree of damage in different parts of the heart and skeletal muscles at different ages and body sizes.

Figure 27. mdx dystrophy pathway proposed by Deconinck *et al.* (Deconinck and Dan 2007)

Hauser, Whitehead, and Yeung suggested that "NADPH oxidase is a major source of reactive oxygen species in dystrophic muscle and its enhanced activity has a stimulatory effect on stretch-induced Ca^{2+} entry, a key mechanism for muscle damage and functional impairment." (Hauser, Hoger *et al.* 1995; Whitehead, Yeung *et al.* 2010; Yang, Chin *et al.* 2013) That is, these authors have recognized the stretch-induced damage is a factor complementing the lack of dystrophin. Therefore, I propose an updated damage pathway in Figure 28, proposing that the stress in muscle is the activation factor operating on the ubiquitous lack of dystrophin.



Figure 28. Hypothesized *mdx* dystrophy pathway.

To ameliorate DMD, beta blockers, ACE inhibitors, and growth inhibition have all been reported to slow DMD progression (Zatz, Betti *et al.* 1986). In the hypothesis proposed in this Chapter, these therapies work because they all modulate stresses in muscles. However, none of these can have a long-term curative effect, leaving more revolutionary genetic and pharmaceutical strategies desirable.

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Appendices

Principle of MRI

-Physics

In the view of classic physics, hydrogen nuclear spins, and thus creates magnetic moment, which aligns to surrounding magnetic field (B₀) in the same way compass points to the North Pole. ¹² Instead of aligning statically with B₀, the magnetic moment would precess about the B₀ direction. The precession frequency, ω , is named Larmor Frequency and defined by $\omega = \gamma B$, where γ is gyromagetic ratio, depending on nuclear species. While nuclei spin reaches steady state of precession toward B₀, we can excite the steady state temporarily with an added magnetic field (B₁), causing the nuclear spin to deviate from its steady state orientation.¹³ After the B₁ was stopped, nuclear spin will align back to the B₀ direction and release energy in a form of detectable radio frequency magnetic waves (called free induction decay, FID), which can be recorded and signal processed for imaging purposes.

-Imaging

Invented in 1952, MRI allows scientist to acquire FID and forms images to provide structural and functional information. Recall the Larmor Frequency, $\omega = \gamma B$, which is proportional to the magnetic field strength B, allowing us to control the ω by varying the B. For imaging purpose, we are interested in encode spin at different spatial locations with different ω . Therefore, a *gradient* of magnetic field, G_x , was applied along x direction on top of the B₀, making the ω a function of x, that is, $\omega(x)=B_0 + x \cdot G$, therefore, the recorded FID carries spatial information in different frequencies, allowing imaging reconstruction through inverse Fourier transform. This imaging technique is called frequency encoding. Please refer to book in footnote 11 for more information. By turning on-off radio frequency pulses and gradients at specific time and direction/strength of radio frequency pulses and gradients are call **pulse sequence**, which determines the sources of contrast and properties of the images.

¹² The mechanism of nuclear spin is beyond scope, please refer to "Questions and Answers in Magnetic Resonance Imaging", 2000, by Allen D. Elster.

¹³ For detailed physics explanation, see "Spin Dynamics: Basics of Nuclear Magnetic Resonance" by Malcolm H. Levitt (2008)

Supplemental tables

Bonferroni							
			Mean Difference			95% Confide	ance Interval
Dependent Variable	(I) group	(J) group	(I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
magB_slc1	1	2	2.4033	2.01043	1.000	-3.6447	8.4514
		3	1.0321	2.01043	1.000	-5.0159	7.0802
		4	6.7905 [*]	2.01043	.023	.7425	12.8386
	2	1	-2.4033	2.01043	1.000	-8.4514	3.6447
		3	-1.3712	2.01043	1.000	-7.4193	4.6768
		4	4.3872	2.01043	.266	-1.6608	10.4353
	3	1	-1.0321	2.01043	1.000	-7.0802	5.0159
		2	1.3712	2.01043	1.000	-4.6768	7.4193
		4	5.7584	2.01043	.067	2896	11.8065
	4	1	-6.7905 [*]	2.01043	.023	-12.8386	7425
		2	-4.3872	2.01043	.266	-10.4353	1.6608
		3	-5.7584	2.01043	.067	-11.8065	.2896
magB_slc2	1	2	6827	2.36177	1.000	-7.7877	6.4223
		3	-1.3088	2.36177	1.000	-8.4138	5.7962
		4	4.9628	2.36177	.311	-2.1422	12.0677
	2	1	.6827	2.36177	1.000	-6.4223	7.7877
		3	6261	2.36177	1.000	-7.7311	6.4789
		4	5.6455	2.36177	.177	-1.4595	12.7504
	3	1	1.3088	2.36177	1.000	-5.7962	8.4138
		2	.6261	2.36177	1.000	-6.4789	7.7311
		4	6.2715	2.36177	.104	8334	13.3765
	4	1	-4.9628	2.36177	.311	-12.0677	2.1422
		2	-5.6455	2.36177	.177	-12.7504	1.4595
		3	-6.2715	2.36177	.104	-13.3765	.8334
magB_slc3	1	2	9798	2.26177	1.000	-7.7840	5.8244
		3	3786	2.26177	1.000	-7.1828	6.4255
		4	8.3630 [*]	2.26177	.012	1.5588	15.1672
	2	1	.9798	2.26177	1.000	-5.8244	7.7840
		3	.6012	2.26177	1.000	-6.2030	7.4053
		4	9.3428*	2.26177	.005	2.5387	16.1470
	3	1	.3786	2.26177	1.000	-6.4255	7.1828

Table 2.Post Hoc Multiple Comparisons of $|\beta|$ in 16 month old mice

		2	6012	2.26177	1.000	-7.4053	6.2030
		4	8.7416*	2.26177	.008	1.9375	15.5458
	4	1	-8.3630 [*]	2.26177	.012	-15.1672	-1.5588
		2	-9.3428 [*]	2.26177	.005	-16.1470	-2.5387
		3	-8.7416*	2.26177	.008	-15.5458	-1.9375
magB_slc4	1	2	-1.3733	2.27890	1.000	-8.2290	5.4824
		3	3664	2.27890	1.000	-7.2221	6.4893
		4	5.1555	2.27890	.228	-1.7002	12.0112
	2	1	1.3733	2.27890	1.000	-5.4824	8.2290
		3	1.0069	2.27890	1.000	-5.8488	7.8626
		4	6.5288	2.27890	.067	3269	13.3845
	3	1	.3664	2.27890	1.000	-6.4893	7.2221
		2	-1.0069	2.27890	1.000	-7.8626	5.8488
		4	5.5219	2.27890	.166	-1.3338	12.3776
	4	1	-5.1555	2.27890	.228	-12.0112	1.7002
		2	-6.5288	2.27890	.067	-13.3845	.3269
		3	-5.5219	2.27890	.166	-12.3776	1.3338
magB_slc5	1	2	.4927	1.96992	1.000	-5.4334	6.4189
		3	.3064	1.96992	1.000	-5.6198	6.2325
		4	5.6741	1.96992	.065	2520	11.6003
	2	1	4927	1.96992	1.000	-6.4189	5.4334
		3	1863	1.96992	1.000	-6.1125	5.7398
		4	5.1814	1.96992	.109	7448	11.1076
	3	1	3064	1.96992	1.000	-6.2325	5.6198
		2	.1863	1.96992	1.000	-5.7398	6.1125
		4	5.3677	1.96992	.090	5584	11.2939
	4	1	-5.6741	1.96992	.065	-11.6003	.2520
		2	-5.1814	1.96992	.109	-11.1076	.7448
		3	-5.3677	1.96992	.090	-11.2939	.5584
magB_slc6	1	2	1.4707	2.06307	1.000	-4.7357	7.6771
		3	1.4960	2.06307	1.000	-4.7104	7.7025
		4	4.2457	2.06307	.338	-1.9607	10.4521
	2	1	-1.4707	2.06307	1.000	-7.6771	4.7357
		3	.0253	2.06307	1.000	-6.1811	6.2317
		4	2.7750	2.06307	1.000	-3.4314	8.9814
	3	1	-1.4960	2.06307	1.000	-7.7025	4.7104
		2	0253	2.06307	1.000	-6.2317	6.1811
		4	2.7497	2.06307	1.000	-3.4567	8.9561

				·			
	4	1	-4.2457	2.06307	.338	-10.4521	1.9607
		2	-2.7750	2.06307	1.000	-8.9814	3.4314
		3	-2.7497	2.06307	1.000	-8.9561	3.4567
magB_slc7	1	2	2.7587	3.21547	1.000	-6.9145	12.4319
		3	1.1879	3.21547	1.000	-8.4853	10.8611
		4	3.6151	3.21547	1.000	-6.0581	13.2883
	2	1	-2.7587	3.21547	1.000	-12.4319	6.9145
		3	-1.5708	3.21547	1.000	-11.2440	8.1024
		4	.8564	3.21547	1.000	-8.8168	10.5296
	3	1	-1.1879	3.21547	1.000	-10.8611	8.4853
		2	1.5708	3.21547	1.000	-8.1024	11.2440
		4	2.4272	3.21547	1.000	-7.2460	12.1004
	4	1	-3.6151	3.21547	1.000	-13.2883	6.0581
		2	8564	3.21547	1.000	-10.5296	8.8168
		3	-2.4272	3.21547	1.000	-12.1004	7.2460
magB_slc8	1	2	6.8267	5.77138	1.000	-10.5356	24.1889
		3	1.4031	5.77138	1.000	-15.9591	18.7654
		4	8.6303	5.77138	.926	-8.7320	25.9925
	2	1	-6.8267	5.77138	1.000	-24.1889	10.5356
		3	-5.4235	5.77138	1.000	-22.7858	11.9387
		4	1.8036	5.77138	1.000	-15.5586	19.1659
	3	1	-1.4031	5.77138	1.000	-18.7654	15.9591
		2	5.4235	5.77138	1.000	-11.9387	22.7858
		4	7.2271	5.77138	1.000	-10.1351	24.5894
	4	1	-8.6303	5.77138	.926	-25.9925	8.7320
		2	-1.8036	5.77138	1.000	-19.1659	15.5586
		3	-7.2271	5.77138	1.000	-24.5894	10.1351

Table 2. Post hoc multiple comparisons for diastolic $|\beta|$ of 16 month old *mdx* and WT mice.

Group1,2,3, and 4 are WT-LC, WT-NC, *mdx*-LC, and *mdx*-NC respectively. MagB, $|\beta|$. Slc1 is base. Slc8 is apex. *, p < 0.05 level. Bonferroni correction. Most differences exist in the slc3 of group 4, indicating abnormal $|\beta|$ manifest closer to base than apex.

	-		Mean Difference			95% Confide	ence Interval
Dependent Variable	(I) group	(J) group	(I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
magB_slc1	1	2	-1.1000	2.97694	1.000	-10.0556	7.8556
		3	.3200	2.97694	1.000	-8.6356	9.2756
		4	2.4000	2.97694	1.000	-6.5556	11.3556
	2	1	1.1000	2.97694	1.000	-7.8556	10.0556
		3	1.4200	2.97694	1.000	-7.5356	10.3756
		4	3.5000	2.97694	1.000	-5.4556	12.4556
	3	1	3200	2.97694	1.000	-9.2756	8.6356
		2	-1.4200	2.97694	1.000	-10.3756	7.5356
		4	2.0800	2.97694	1.000	-6.8756	11.0356
	4	1	-2.4000	2.97694	1.000	-11.3556	6.5556
		2	-3.5000	2.97694	1.000	-12.4556	5.4556
		3	-2.0800	2.97694	1.000	-11.0356	6.8756
magB_slc2	1	2	6400	3.36391	1.000	-10.7598	9.4798
		3	2.0400	3.36391	1.000	-8.0798	12.1598
		4	3.7400	3.36391	1.000	-6.3798	13.8598
	2	1	.6400	3.36391	1.000	-9.4798	10.7598
		3	2.6800	3.36391	1.000	-7.4398	12.7998
		4	4.3800	3.36391	1.000	-5.7398	14.4998
	3	1	-2.0400	3.36391	1.000	-12.1598	8.0798
		2	-2.6800	3.36391	1.000	-12.7998	7.4398
		4	1.7000	3.36391	1.000	-8.4198	11.8198
	4	1	-3.7400	3.36391	1.000	-13.8598	6.3798
		2	-4.3800	3.36391	1.000	-14.4998	5.7398
		3	-1.7000	3.36391	1.000	-11.8198	8.4198
magB_slc3	1	2	.0600	3.60697	1.000	-10.7910	10.9110
		3	2.1800	3.60697	1.000	-8.6710	13.0310
		4	6.1800	3.60697	.636	-4.6710	17.0310
	2	1	0600	3.60697	1.000	-10.9110	10.7910
		3	2.1200	3.60697	1.000	-8.7310	12.9710
		4	6.1200	3.60697	.655	-4.7310	16.9710
	3	1	-2.1800	3.60697	1.000	-13.0310	8.6710
		2	-2.1200	3.60697	1.000	-12.9710	8.7310

Table 3. Post Hoc Multiple Comparisons of $|\beta|$ in 3 month old mice

Bonferroni

		-					
		4	4.0000	3.60697	1.000	-6.8510	14.8510
	4	1	-6.1800	3.60697	.636	-17.0310	4.6710
		2	-6.1200	3.60697	.655	-16.9710	4.7310
		3	-4.0000	3.60697	1.000	-14.8510	6.8510
magB_slc4	1	2	.1200	3.07448	1.000	-9.1290	9.3690
		3	4.0600	3.07448	1.000	-5.1890	13.3090
		4	7.4600	3.07448	.165	-1.7890	16.7090
	2	1	1200	3.07448	1.000	-9.3690	9.1290
		3	3.9400	3.07448	1.000	-5.3090	13.1890
		4	7.3400	3.07448	.178	-1.9090	16.5890
	3	1	-4.0600	3.07448	1.000	-13.3090	5.1890
		2	-3.9400	3.07448	1.000	-13.1890	5.3090
		4	3.4000	3.07448	1.000	-5.8490	12.6490
	4	1	-7.4600	3.07448	.165	-16.7090	1.7890
		2	-7.3400	3.07448	.178	-16.5890	1.9090
		3	-3.4000	3.07448	1.000	-12.6490	5.8490
magB_slc5	1	2	-1.0200	3.33267	1.000	-11.0458	9.0058
		3	2.5000	3.33267	1.000	-7.5258	12.5258
		4	5.6200	3.33267	.667	-4.4058	15.6458
	2	1	1.0200	3.33267	1.000	-9.0058	11.0458
		3	3.5200	3.33267	1.000	-6.5058	13.5458
		4	6.6400	3.33267	.382	-3.3858	16.6658
	3	1	-2.5000	3.33267	1.000	-12.5258	7.5258
		2	-3.5200	3.33267	1.000	-13.5458	6.5058
		4	3.1200	3.33267	1.000	-6.9058	13.1458
	4	1	-5.6200	3.33267	.667	-15.6458	4.4058
		2	-6.6400	3.33267	.382	-16.6658	3.3858
		3	-3.1200	3.33267	1.000	-13.1458	6.9058
magB_slc6	1	2	3800	3.73493	1.000	-11.6159	10.8559
		3	2.8400	3.73493	1.000	-8.3959	14.0759
		4	4.8800	3.73493	1.000	-6.3559	16.1159
	2	1	.3800	3.73493	1.000	-10.8559	11.6159
		3	3.2200	3.73493	1.000	-8.0159	14.4559
		4	5.2600	3.73493	1.000	-5.9759	16.4959
	3	1	-2.8400	3.73493	1.000	-14.0759	8.3959
		2	-3.2200	3.73493	1.000	-14.4559	8.0159
		4	2.0400	3.73493	1.000	-9.1959	13.2759
	4	1	-4.8800	3.73493	1.000	-16.1159	6.3559

		2	-5.2600	3.73493	1.000	-16.4959	5.9759
		3	-2.0400	3.73493	1.000	-13.2759	9.1959
magB_slc7	1	2	2000	3.96679	1.000	-12.1334	11.7334
		3	1.8000	3.96679	1.000	-10.1334	13.7334
		4	.6800	3.96679	1.000	-11.2534	12.6134
	2	1	.2000	3.96679	1.000	-11.7334	12.1334
		3	2.0000	3.96679	1.000	-9.9334	13.9334
		4	.8800	3.96679	1.000	-11.0534	12.8134
	3	1	-1.8000	3.96679	1.000	-13.7334	10.1334
		2	-2.0000	3.96679	1.000	-13.9334	9.9334
		4	-1.1200	3.96679	1.000	-13.0534	10.8134
	4	1	6800	3.96679	1.000	-12.6134	11.2534
		2	8800	3.96679	1.000	-12.8134	11.0534
		3	1.1200	3.96679	1.000	-10.8134	13.0534

Table 3 Post hoc multiple comparisons for diastolic $|\beta|$ of 3 month old *mdx* and WT mice showing no significance.

Group1,2,3, and 4 are WT-LC, WT-NC, *mdx*-LC, and *mdx*-NC respectively. MagB, $|\beta|$. Slc1 is base. Slc7 is apex. No significance was found among all comparison.

Speculation: Might low gravity ameliorate dystrophy?

In the Chapter 4, we raised the point that higher stress within certain muscles might explain the regional heterogeneity of dystrophy (e.g., severe at heart base but not apex; severe in calf but not forearm). If genetic therapy could not break limitations on gene size and achieve stable transfection, relocating DMD patients to low gravity environments might reduce muscle stress and ameliorate the dystrophy. Furthermore, low gravity affects normal human health due to the redistribution of circulation pressure and decrease muscle/skeleton tension (Smith, McCoy *et al.* 2012). Lower limb muscles atrophy was observed among astronauts to the greatest extent, with the *calf muscles being most affected* and most difficult to target with exercise countermeasures (Lester, Standley *et al.* 2013). Similarly, the most obvious pathological sign of

DMD patients is the pseudo-hypertrophic *calf muscle*, indicating the calf muscle are sensitive to stress, both in space and on the Earth. In fact, DMD patients on the Earth and astronauts in orbits show complimentary loss of muscle fiber types. Specifically, DMD patients lose fast fibers rather than slow fibers (Webster, Silberstein *et al.* 1988), while astronauts lose slow fibers rather than fast fiber (Riley, Bain et al. 2002). I hypothesize that dystrophin is part of stress sensing pathway and signals to reduce astronauts' slow fibers in response to the lower load. When dystrophin is missing, the presumed stress sensing pathway might over-react to the normal stress on the earth and signal to reduce fast fibers in DMD. Since it is unknown how exactly low-gravity changes circulation/muscle/skeleton ¹⁴, how dystrophin-null animals adapt low-gravity environment may shed lights on such mechanism and bring better solutions to counter atrophy in space missions. The hypothetical mechanical sensing and protective mechanism of dystrophin would be critical for extraplanetary exploration, when we would need to fine-tune human body to adapt different gravities.

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¹⁴ "The exact cellular and biochemical events that produce these losses of mass and strength are not as well understood." NASA 2013. Effect of Prolonged Space Flight on Human Skeletal Muscle (Biopsy)

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+HONORS

- · 2013 American Heart Association BCVS Travel Grant
- 2012 American Heart Association Predoctoral Fellowship
- 2012 Washington University Merit Fund of Engineering School
- 2012 Biomedical Engineering Society Design and Research Award
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- 2012 2nd place award in Graduate Research Symposium of Washington University
- 2011 2nd place award in North America Taiwanese Engineering Association Symposium
- 2007 Taiwan Merit Scholar, National Science Council, Taiwan, ROC
- 2006 Outstanding Achievement Award of National Taiwan University Scholarship
- 2006 Award of Taiwan Competition on Biomedical Engineering Creative Design
- 2005 Valedictorian of Class 2005 of Life Science Department, NTU, Taiwan
- 2003 Scholarship for Outstanding Students' Education
- 2001 Silver Medal, International Biology Olympiad, Belgium
- 2001 Presidential Award of National Taiwan University

+RESEARCH

Washington University School of Medicine, St. Louis, MO

2007-2014

Graduate Research Assistant

- Acquire myocardial structure of viable mouse hearts with ultra-high field 11.7 tesla MRI, resulting in a \$52,000 American Heart Association predoctoral fellowship and a \$5,000 merit fund from Washington University
- Designed an *ex vivo* Langendorff mouse heart perfusion system for MRI, resulting in 2012 Design and Research Award of the Biomedical Engineering Society

- Proved dystrophic mice heart is abnormal dominantly at heart base rather than apex, despite mutation is every cell, indicating local heterogeneity modifies cardiac myopathy
- Implement automatic imaging segmentation, signal processing and statistical analysis of ultrasound, MRI, and optic images of heart muscle
- Developed theory of higher stress deteriorates heart muscle regionally at heart base
- Led collaboration among four labs for anti-aging treatments involving drugloaded nanoparticles

National Taiwan University, Taipei, Taiwan

July 2005–June 2007

Graduate Research Assistant

- Designed a microbubble and ultrasound sonification/quantification scheme, resulting in 2006 2nd Place Award in Biomedical Engineering Creative Design Competition in Taiwan
- Quantitative analysis for ultrasound spectrum with MATLAB

Academia Sinica, Institute of Biomedical Science July-August 2004 Taipei, Taiwan

Undergraduate Research Assistant

- Construct DNA clone of cancer proteases
- Imaged euryhaline fish's gill with scanning electron microscope for salt modulation study

+RESEARCH INTRESTS

- Imaging / signal processing
- Compressed sensing in MRI
- Heart muscle form & function
- Diastolic suction mechanism
- Boneless muscle mechanics
- Coronary vasculature

+PRESENTATIONS

•	Oral in American Heart Association, Dallas, TX	November 2013
•	Poster in ISMRM, Salt Lake City, UT	April 2013
•	Poster in FASEB, Boston, MA	April 2013
•	Poster in American Heart Association, Los Angeles, CA	November 2012
•	Oral in Biomedical Engineering Society, Atlanta, GA	October 2012
•	Poster in Washington University Graduate Research Sym	posium, MO
		April 2012
•	Oral in Society for Cardiovascular Magnetic Resonance, (Drlando, FL
		February 2012
•	Poster in American Heart Association, Orlando, FL	November 2011
•	Oral in North America Taiwanese Engineering Association	n Symposium
		April 2011
•	Oral in Acoustical Society of America, San Antonio, TX	-
		October 2009
•	Oral in IEEE International Ultrasonics Symposium, Rome	Italy
		September 2009

+SKILLS

- High field diffusion tensor MRI
- Signal & imaging processing • in MATLAB
- FFT Spectrum analysis
- Image segmentation
- Small animal surgery
- Langendorff perfused mouse heart

- Anatomy / Histology •
- Biomedical Engineering •
- ImageJ / Photoshop •
- SPSS / Linux / LibreOffice •
- SSH •
- Fluent in English, Native in •
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+LEADERSHIP

Washington University Biomedical Engineering Society President	2012
Washington University Taiwanese Graduate Student Association Vice President	2011

+MEMBERSHIPS

Biomedical Engineering Society	2008, 2012
American Heart Association	2010-13
National Society of Professional Engineer	2011
American Association of Anatomists	2012

+PEER-REVIEWED PUBLICATIONS

- Kristin P. Bibee, **Ya-Jian Cheng**, et al. (2014). Rapamycin nanoparticles target defective autophagy in muscular dystrophy to enhance both strength and cardiac function. Federation of American Societies for Experimental Biology, accepted.
- Cheng, Y. J., D. Lang, et al. (2012). Focal but reversible diastolic sheet dysfunction reflects regional calcium mishandling in dystrophic mdx mouse hearts. American journal of physiology. Heart and circulatory physiology 303(5): H559-568
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