CLINICAL RESEARCH ARTICLE

See Editorial on pages 4-5 in this issue.

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Elevation of fast but not slow troponin I in the circulation of patients with Becker and Duchenne muscular dystrophy

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Abstract

Introduction: One of the hallmarks of injured skeletal muscle is the appearance of elevated skeletal muscle proteins in circulation. Human skeletal muscle generally consists of a mosaic of slow (type I) and fast (type IIa, IIx/d) fibers, defined by their myosin isoform expression. Recently, measurement of circulating fiber-type specific isoforms of troponin I has been used as a biomarker to suggest that muscle injury in healthy volunteers (HV) results in the appearance of muscle proteins from fast but not slow fibers. We sought to understand if this is also the case in severe myopathy patients with Becker and Duchenne muscular dystrophy (BMD, DMD).

Methods: An enzyme-linked immunosorbent assay (ELISA) that selectively measures fast and slow skeletal troponin I (TNNI2 and TNNI1) was used to measure a cross-section of patient plasma samples from HV (N = 50), BMD (N = 49), and DMD (N = 132) patients. Creatine kinase (CK) activity was also measured from the same samples for comparison.

Results: TNNI2 was elevated in BMD and DMD and correlated with the injury biomarker, CK. In contrast, TNNI1 levels were indistinguishable from levels in HV. There was an inverse relationship between CK and TNNI2 levels and age, but no relationship for TNNI1.

Discussion: We define a surprising discrepancy between TNNI1 and TNNI2 in patient plasma that may have implications for the interpretation of elevated muscle protein levels in dystrophinopathies.

KEYWORDS

biomarker, creatine kinase, muscle injury, muscular dystrophy, troponin

Abbreviations: BMD, Becker muscular dystrophy; CK, creatine kinase; DMD, Duchenne muscular dystrophy; EDTA, ethylenediaminetetraacetic acid; ELISA, enzyme-linked immunosorbent assay; eMHC, embryonic myosin heavy chain; HRP, horseradish peroxidase; HV, healthy volunteer; IgG, immunoglobulin G; IR, infrared; IRB, Institutional Research Board; MLPA, Multiplex ligation-dependent probe amplification; NADH, nicotinamide adenine dinucleotide; NADP, nicotinamide adenine dinucleotide phosphate; PBS, phosphate-buffered saline; PBS-T, phosphate-buffered saline containing 0.1% Tween-20; REC, Research Ethics Committee; TNNI, troponin inhibitory subunit; U, unit.

1 | INTRODUCTION

Human skeletal muscle consists of both fast (type IIa, IIx) and slow (type I) myofibers defined by the expression of their myosin isoform, a sarcomere contractile protein, required for force development. Human

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skeletal muscle consists of approximately 50% slow and 50% fast fibers,¹ although individual muscles can vary in their slow/fast proportion.² Each fiber type has distinctive properties with respect to contractile speed, substrate utilization, and fatigue resistance, and the body uses these different properties to efficiently coordinate movement.

When healthy muscle is subjected to excessive, unaccustomed exercise, it develops soreness and sustained reductions in strength and range of motion. Proteins also leak from injured muscle fibers into circulation, including creatine kinase (CK), lactate dehydrogenase, and myoglobin. These biomarkers are not unique to either fast or slow fibers and so do not provide detail regarding differences in fiber responses to injury. Troponin I (TNNI) is a component of the troponin complex that controls initiation of contraction of muscle by calcium. It is distinct in that there is a different isoform for each type of striated muscle: TNNI1 in slow skeletal muscle, TNNI2 in fast skeletal muscle, and TNNI3 in cardiac muscle. Selective enzyme-linked immunosorbent assays (ELISAs) have been used to demonstrate that TNNI2 but not TNNI1 is elevated in circulation after injurious exercise,³ even under extreme conditions.⁴

Duschene muscular dystrophy (DMD) and Becker muscular dystrophy (BMD) are caused by an absence (DMD) or truncation (BMD) of the dystrophin protein.⁵ Dystrophin provides a structural link between the actin cytoskeleton and the basement membrane through the dystrophin-glycoprotein complex.⁶ When dystrophin is absent or truncated, contraction of muscle leads to heightened muscle stress and injury with normal use.

While the sensitivity to injury is much higher in DMD muscle than in BMD or healthy muscle, fast fibers still appear to be more susceptible than slow fibers, with young DMD patients exhibiting histological evidence of disruption in fast fibers⁷ and early loss of type IIx fibers.⁸ We sought to explore the relative susceptibility of these fibers to leak muscle contents.

2 | METHODS

2.1 | Patient samples

Healthy volunteer (HV) frozen plasma samples were purchased from BioIVT (Westbury, NY) and are confirmed to have been collected with institutional review board (IRB) approval and with appropriate informed consent. Plasma and serum for affected individuals were received from the Newcastle MRC Centre Biobank for Rare and Neuromuscular Diseases (DMD), and a Becker muscular dystrophy biomarker study at Binghamton University - SUNY (BMD). Upon receipt, all samples were aliquoted into working volumes of 50–100 μ L and stored at –80°C to minimize freeze-thaw damage. For the BMD biomarker study at Binghamton University - SUNY, recruitment was through foundation social media posts, with centralized remote consent of subjects via telephone (IRB approval by Binghamton University). Patients had centrally curated DMD gene testing reports with mutation data consistent with a BMD diagnosis (in-frame mutations, or other mutations consistent with partial dystrophin expression). For patients with a clinical diagnosis but without a prior genetic report, genetic confirmation was made via multiplex ligation-dependent probe amplification (MLPA) genetic analysis (Fullerton Genetics Laboratory, Asheville, NC).

A mobile phlebotomist was sent to the subject's home for sample collection. Red top serum vacutainer tubes, containing silica act clot activator, were used for the blood collection. If a subject required MLPA testing, an ethylenediaminetetraacetic acid (EDTA) tube would be added for those collections, but was not used for any other analysis. After the serum tubes were left to clot for 30 min, they were processed in a centrifuge at 1000-1300 \times g for 10 min. The serum (top layer) fluid was then pipetted from the vacutainer tube and transferred into cryovials and immediately frozen on dry ice for shipment and later storage at -80° C. Serum samples were sent frozen on dry ice to Binghamton University and stored at -80° C. Samples were collected from 2017 to 2019 and analyzed in 2019.

Plasma samples from the Newcastle MRC Centre Biobank were collected from patients attending clinics at The John Walton Muscular Dystrophy Research Centre. DMD donors were genetically diagnosed via the NHS Northern Genetics Service Laboratory prior to providing blood samples to the MRC Centre Biobank for Rare & Neuromuscular Diseases. A multidisciplinary clinical team from the John Walton Muscular Dystrophy Research Centre were responsible for the clinical observations of this cohort. Collection of samples for storage and subsequent release for research was ethically approved by North East - Newcastle & North Tyneside 1 Research Ethics Committee (REC reference: 19/NE/0028). Informed consent was obtained from donors using REC approved consent forms. Purple top BD EDTA Vacutainers were used to collect peripheral blood from the DMD cohort. Blood was drawn into vacutainers, gently inverted 5-10 times to ensure adequate mixing of blood with EDTA and then centrifuged at $1500 \times g$ for 10 min. The upper plasma fraction was transferred via pipette into cryovials and immediately stored at -80°C. Samples were collected over a period of 9 y (2010-2019) and stored at -80°C prior to release to Edgewise Therapeutics for analysis.

2.2 | CK assay

Blood plasma CK activity was assayed using a coupled-reaction kit purchased from Pointe Scientific (Canton, MI). Plasma was diluted 25-fold with phosphate-buffered saline (PBS), of which 2 µL was added to the 384-well plate. The CK assay reagent (70 µL, 4:1 kit Buffer A:Buffer B) was added using the Multidrop Combi (ThermoFisher, Inc., Waltham, MA) and the reaction progress monitored by absorbance at 340 nm for 30 min with the SpectraMax M3 plate reader (Molecular Devices, San Jose, CA) over approximately 20-30 min. Following the termination of the reaction, pathlength correction values were measured with near-infrared (IR) absorbance at 900 and 975 nm. The raw absorbance data were processed in Microsoft Excel to exclude points with $A_{340} > 2.5$ and to correct for pathlength using a system-specific K-Factor of 0.168, which was measured and applied as described.⁹ The corrected absorbance data vs time was fit to a linear model in GraphPad Prism (GraphPad Software, San Diego, CA) to yield reaction slopes, which were compared

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to a standard curve of nicotinamide adenine dinucleotide (NADH; 5–100 μM) to yield enzyme rates in U/L, where U is defined as the amount of enzyme that results in the reduction of 1 $\mu mol \cdot L^{-1} \cdot min^{-1}$ nicotinamide adenine dinucleotide phosphate (NADP).

3 | TNNI ELISAS

Plasma concentrations of TNNI isoforms for slow and fast muscle were measured by capture ELISA. The slow isoform (TNNI1) was measured using a commercially available test kit (LS-F7068, LifeSpan Biosciences, Inc, Seattle, WA) and was performed according to the manufacturer's instructions. The fast isoform (TNNI2) was assayed as described previously.⁴ Briefly, high-binding ELISA plates were coated with α -TNNI2 monoclonal antibody (Clone 7G2, OriGene, Inc., Rock-ville, MD) at a concentration 6.4 µg/mL overnight at 4°C. The wells were blocked with 1% w/v non-fat dry milk in PBS for 30 min at 37°C, followed by incubation for 2 h at 37°C with the samples or recombinant human TNNI2 as a standard curve. The wells were washed with PBS containing 0.1% Tween-20 (PBS-T) and incubated with 1 µg/mL polyclonal α -TNNI2 antibody (PA5-76303, Thermo-Fisher, Inc.) for 90 min at 37°C. After washing with PBS-T, the detection antibody (horseradish peroxidase [HRP]-conjugated goat- α -rabbit IgG, 0.08 µg/mL, Pierce Biosciences) was added for 45 min at 37°C,

TABLE 1 Demographics of muscular dystrophy patients and healthy controls

	Healthy (n = 50)	BMD (n = 49)	DMD (n = 132)
Age (y)	45.5 +/- 15.9	38.4 +/- 18.8	13.0 +/- 5.7
Age range (y)	19-75	6-73	2-33
Non-ambulant	-	17 (35%)	60 (45%)

Note: Data are mean ± SD.



FIGURE 1 Plasma concentrations of CK enzymatic activity (A), TNNI1 (B), and TNNI2 (C) were measured in samples from BMD patients (squares) and DMD patients (triangles), with HV as controls (circles). In each panel, the error bars represent the median +/– the interquartile range. In panels (B) and (C), samples that exhibited no detectable TNNI concentration were assigned values equal to the assay's limit of detection (0.1 ng/mL and 0.001 ng/mL for fast and slow TNNI, respectively) When compared with each other, a significant correlation was found between CK and fast TNNI2 (D), with an R² of 0.67. There was no significant correlation between CK and slow TNNI1 (E) nor between fast TNNI2 and slow TNNI1 (F) In panels (D) – (F), healthy samples are represented as black triangles, BMD as blue diamonds, and DMD red circles. ****: P < .0001. All other comparisons are non-significant

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BARTHEL ET AL.

and the HRP was visualized with Ultra-TMB colorimetric reagent (ThermoFisher) followed by quenching with 2 N H_2SO_4 and measurement of the absorbance at 410 nm. Selectivity of these assays for fast vs slow TNNI has previously been confirmed using human muscle extracts.⁴

4 | RESULTS

Demographics of subjects included in the study are described in Table 1. We observed a wide range of CK activities in all groups with higher average values in BMD patients compared to HV (678 \pm 923 U/L and 355 \pm 1743 U/L, respectively) and greater activity than both BMD and HV in DMD samples (3402 +/- 2969 U/L; *P* < .0001) (Figure 1A).

We observed a similar pattern of elevated TNNI2 concentration (Figure 1B; HV: 6.0 +/– 38.9, BMD: 3.1 +/– 5.7, DMD: 51.0 +/– 71.6 ng/mL), although unlike CK, 83% of HV samples had TNNI2 concentrations below the level of detection for the assay (0.1 ng/mL). In contrast, all samples but one had concentrations of TNNI1 below 0.4 ng/mL (Figure 1C) with no clear relationship between disease group and concentration (HV: 0.036 +/– 0.052, BMD: 0.034 +/– 0.048, DMD: 0.16 +/– 1.13 ng/mL). A significant correlation was present between TNNI2 and CK ($R^{2=}0.68$, P < .001) (Figure 1D) while no

such correlation existed between TNNI1 and CK or TNNI1 and TNNI2 (Figure 1E, F).

CK activity was highest in younger DMD patients with a decline in activity in older patients (Figure 2A, $T_{1/2}$ for the first-order decay fit: 5.6 y). Similar results were obtained for TNNI2 (Figure 2B) but not TNNI1 (Figure 2C). In the BMD patients, CK and TNNI2 levels were higher in younger patients, but the relationship with age was less clear and flattened compared to DMD (Figure 2D, E). As with DMD, TNNI1 levels in BMD were low and had no age relationship (Figure 2F).

Non-ambulant DMD patients had significantly lower CK activity and TNNI2 concentrations than ambulant patients (Figure 3A, B, CK: 798 +/- 955 U/L vs 3276 +/- 1704, TNNI2: 25.2 +/- 32.2 vs 82.6 +/- 86.9) but not lower TNNI1 (Figure 3C). Within the BMD patient population, the changes were not significant at the 95% confidence level (Figure 3D–F).

5 | DISCUSSION

This cross-sectional, retrospective study describes unexpected differences between fast and slow skeletal muscle biomarkers in DMD/BMD patient plasma. Although the data demonstrate a marked difference between fast and slow biomarkers between groups, it should be noted that a limitation of the current study is that these are



FIGURE 2 Concentration of CK enzymatic activity (A), TNNI2 (B), and TNNI1 (C) vs patient age in DMD patient samples. The same comparisons were made for BMD patients in panels (D, E, and F) for CK, TNNI2, and TNNI1, respectively

47



FIGURE 3 Ambulatory status for DMD was compared against plasma concentrations of CK enzymatic activity (A), TNNI2 (B), and TNNI1 (C). The same comparisons were made for BMD (D, E, and F). A patient was defined as "ambulatory" so long as the patient was not described as wholly dependent upon a wheelchair for mobility. Bars represent the mean +/- the standard error for the population. ****: P < .0001, ns: non-significant

not age matched, leaving the possibility of a different conclusion with closer age comparison. Also of note, aggregated CK activity levels in our BMD and DMD samples were lower than those measured in recent published datasets,¹⁰ possibly due to differences between assay systems. Finally, the mean CK of 355 U/L in the HV population was higher than is generally observed in healthy populations. This high value is driven by two individuals with much higher levels than the rest of the population (approximately 1000 and 10 000 U/L; Figure 1A). Unaccustomed exercise in healthy individuals can raise circulating CK above upper limits of normal and beyond 10 000 U/L.¹¹ Therefore, the most likely explanation is that these individuals performed some form of strenuous exercise prior to sampling. The population median (65.0, shown in Figure 1A) is in line with normal ranges for males (22–334¹²).

The findings of differential troponin levels do, however, agree with a previous study that included two myopathy patients,¹³ although the specific disorders and absolute concentration of TNNI isoforms were not provided in that study. Our findings are also similar to studies in HV performing injurious eccentric exercise. In this setting, transient elevations in TNNI2 but not TNNI1 were observed after unilateral arm exercise³ or more extreme whole body exercise regimens⁴ and levels of TNNI2 were correlated with those of CK. Elevations in both TNNI1 and TNNI2 have been recorded in

severe trauma and ischemia,¹³ suggesting that the release of TNNI1 into circulation may require muscle injury beyond that caused by contraction-based injury, even in the context of dystrophin absence or mutation. Similar observations have also been made for cardiac TNNI3, which is only measurable in circulation after a severe ischemic event.¹⁴

Differences in disease susceptibility between fast and slow skeletal muscle fibers has been documented in both clinical and preclinical studies. Histological evidence of increased fast fiber disruption was first documented in DMD patients by examination of the fiber distribution of embryonic myosin heavy chain (eMHC), a contractile protein only expressed in regenerating fibers.⁷ Similar findings have also been documented in preclinical models of DMD, including dystrophic pigs and dogs.^{15,16} Transgenic *mdx* mice that have a higher proportion of slow fibers also have milder disease progression.^{17,18} Our observations provide additional detail regarding this relative susceptibility but also raise a number of questions.

As disease progresses in DMD, fibrosis and fatty tissue slowly replace muscle. Remaining skeletal muscle fibers also exhibit advanced pathophysiology including co-expression of fast and slow myosin along with eMHC, which is observed in mixed, slow, and fast fibers.¹⁹ Rates of muscle turnover (measured with eMHC) are approximately constant with age (~30%),²⁰ suggesting that changes

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in muscle composition do not affect rates of injury. In light of these changes, our results of low TNNI1, independent of age are surprising and suggest that slow or mixed fast/slow fiber injury or turnover occurs differently from fast fibers. One possible explanation is that membrane injury in DMD and BMD is considerably more severe in pure fast fibers compared to slow fibers. Fast myofibers develop greater relative force,²¹ and this would be expected to impart greater physical stresses on the dystrophin-deficient membrane. Structural response to stress may also be different as fast fibers contain smaller integrating z-discs²² that are more susceptibility to disruption.²³ Additional studies will be required to determine why slow muscle fibers resist membrane rupture with contraction injury and may inform novel strategies to protect fast fibers from excessive disruption in BMD and DMD.

Our observations of decreasing CK and TNNI2 with age and ambulatory status in DMD are consistent to those seen with other studies.^{10,24} This trajectory is usually assigned to decreases in muscle volume due to replacement through fibrosis. Our observations of differences between TNNI1 and TNNI2 levels with age suggest that the decreases in CK may be more nuanced, reflecting only the injury status of fast fibers.

CK is the most commonly measured circulating biomarker of muscle injury, and elevated levels are commonly used to diagnose pediatric skeletal muscle myopathies. Our cross-sectional data reveal interesting differences between the distribution of CK and TNNI2. The majority of HV (83%) had TNNI2 levels below the lower level of detection of the ELISA (<0.1 ng/mL), while only 4% of BMD and 6% of DMD patients had non-measurable levels of TNNI2. In contrast, CK activity was measurable in all samples tested with large overlap between HV and BMD activity. This difference is consistent with other cross-sectional examinations of TNNI2 vs CK.¹⁰ A more sensitive TNNI2 assay will be required to properly define basal concentrations in HV and differences between healthy, BMD, and DMD patients. Greater understanding of these differences may provide a useful biomarker tool for the assessment of novel treatment effects in patients with relatively lower CK activity including BMD or older DMD patients.

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CONFLICT OF INTEREST

Benjamin Barthel and Alan Russell are paid employees of Edgewise Therapeutics, Inc. The remaining authors have no conflicts of interest.

ETHICAL PUBLICATION STATEMENT

We confirm that we have read the Journal's position on issues involved in ethical publication and affirm that this report is consistent with those guidelines.

DATA AVAILABILITY STATEMENT

Data available on request from the authors

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