

ON BEHALF OF THE INSPIRE DUCHENNE STUDY TEAM

Update on INSPIRE DUCHENNE: a phase 1/2 study of SGT-003, a next-generation microdystrophin gene therapy for Duchenne muscular dystrophy

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SGT-003 is an investigational product that has not been approved in any region.
No conclusions regarding safety and efficacy can be made.



Disclosures

Clinical trial support:

- Sarepta
- Dyne
- Avidity
- Ultragenyx
- Solid

Advisory boards:

- Armatus
- Encoded
- Insmmed
- Dyne
- Solid

Past royalties:

- Astellas

Duchenne Muscular Dystrophy (Duchenne): Background

Duchenne is an X-linked recessive neuromuscular disorder caused by a lack of functional dystrophin¹



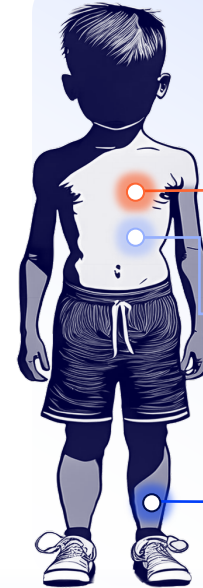
Dystrophin is required for maintaining muscle integrity and function²⁻⁴

- Deterioration of muscle integrity leads to loss of essential membrane proteins and muscle fiber breakdown and leakage, resulting in progressive functional decline



Shortened, functional “microdystrophin” transgenes can be packaged into AAVs to replace dystrophin⁵

- Microdystrophins vary based on their unique compositions⁶



- Decreased heart function
- Cardiomyopathy

HEART FAILURE

- Weak diaphragm

RESPIRATORY FAILURE

- Loss of muscle mass
- Inflammation
- Fibrosis

LOSS OF AMBULATION

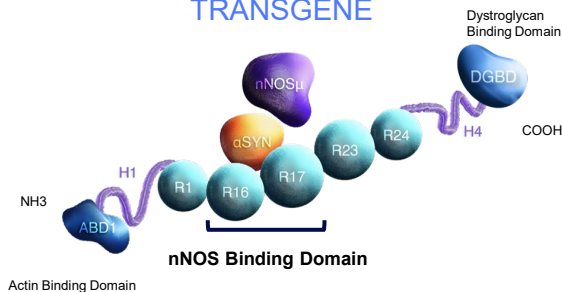
The impact of treatments on muscle integrity is key for patients with Duchenne⁷

AAV=adeno-associated virus.

1. Duan D, et al. *Nat Rev Dis Primers*. 2021;7(1):13. 2. Sheybani A, et al. *Pediatr Res*. 2022;92(6):1613-1620. 3. Voleti S, et al. *Pediatr Cardiol*. 2020;41(6):1173-1179. 4. Wagner KR, et al. *Biomark Med*. 2021;15(15):1389-1396. 5. Crudele JM, et al. *Hum Mol Genet*. 2019;28(R1):R102-R107. 6. Ramos JN, et al. *Mol Ther*. 2019;27(3):623-635. 7. Escobar-Huertas JF, et al. *Cytoskeleton (Hoboken)*. 2024;81(6-7):269-286.

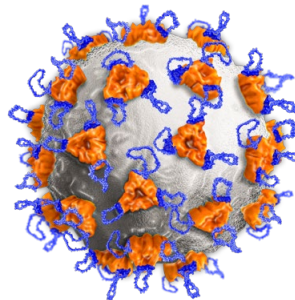
SGT-003: A Next-Generation AAV-Microdystrophin Gene Therapy Candidate^a

SGT-003 MICRODYSTROPHIN TRANSGENE



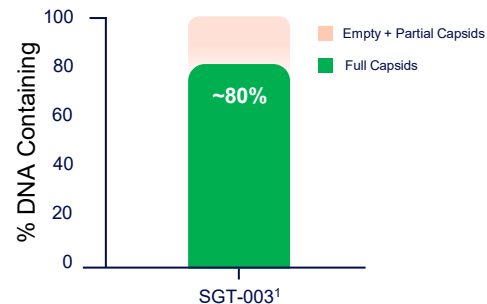
Unique inclusion of nNOS-binding domain designed with the goal of improving blood flow to prevent activity-induced ischemia and associated muscle injury¹

SGT-003 AAV-SLB101 CAPSID



Rationally designed muscle-tropic capsid targeting multiple integrin receptors that are upregulated in dystrophic muscle²

SGT-003 FULL/EMPTY CAPSID RATIO



SGT-003 GMP manufacturing at ~80% full/empty capsid ratio (1000L scale)³

SGT-003's optimized transgene and next-generation capsid were selected to deliver a unique microdystrophin to muscles throughout the body while also reducing liver distribution^{1,2}

^aαSYN=alpha-synuclein; ABD1=actin-binding domain 1; DGBD=dystroglycan-binding domain; H=hinge; nNOS=neuronal nitric oxide synthase; R=spectrin-like repeat.

^aSGT-003 is an investigational product that has not been approved in any region. No conclusions regarding safety and efficacy can be made.

1. Lai Y, et al. *J Clin Invest*. 2009;119(3):624-635. 2. Vu Hong A, *Nat Commun*. 2024;15(1):7965. 3. Data on file. Solid Biosciences.

INSPIRE DUCHENNE: Recently Expanded Study Overview



- Single-dose level, open-label, Phase 1/2 study
- Patients with a confirmed diagnosis of Duchenne
- Actively enrolling: US, Canada, Italy, and the UK
- Prophylactic prednisone regimen as immunomodulation
- Potential for further enrollment of older and non-ambulatory patients (Cohorts 4 and 5)
- NCT06138639

Primary Endpoints:

- Incidence of treatment-emergent adverse events through Day 360
- Change from baseline of microdystrophin protein levels at Day 90

Secondary Endpoints:

- Microdystrophin protein levels and distribution at Days 90 and 360
- TTR, 10MWR, 4SC, NSAA, 6MWT, SV95C at Days 360 and 540

Exploratory Endpoints:

- % predicted FVC, PEF, FEV1; Bayley-4; PODCI

KEY ELIGIBILITY CRITERIA

Age:	Cohort 1: Aged 4 to <7 years Cohort 2: Aged 7 to <12 years Cohort 3: Aged 0 to <4 years
DMD Genetic Variant Exclusions:	Any deletion in exons 1 to 11 and/or 42 to 45, inclusive
Ambulation:	Cohorts 1, 2: Required Cohort 3: N/A
Additional Function:	Cohorts 1, 3: N/A Cohort 2: TTR and 10MWR criteria

Antibodies:	Negative for AAV9 antibodies
Prior Treatments:	No history of gene therapy ≥12-week washout from exon-skipping therapies, vamorolone, and/or givinostat
Steroid Regimen:	Cohorts 1, 2: On a stable dose of daily oral steroids (prednisone/deflazacort) for ≥12 weeks Cohort 3: N/A

4SC=4-stair climb; 6MWT=6-minute walk test; 10MWR=10-meter walk/run; FEV1=forced expiratory volume in 1 second; FVC=forced vital capacity; IV=intravenous; N/A=not applicable; NSAA=North Star Ambulatory Assessment; PEF=peak expiratory flow; SV95C=stride velocity 95th centile; TMA=thrombotic microangiopathy; TTR=time to rise.
Data on file. Solid Biosciences.

15 Participants Enrolled in INSPIRE DUCHENNE

As of a data cutoff of August 12, 2025, 15 participants have received SGT-003 and have follow-up periods ranging up to over 1-year post-dosing

Cohort	Eligible Age Range (years)	Ages at Enrollment (years)	Weights for Dosing (kg)	Participants Enrolled (n)
1	4 to < 7	5 to 6	Up to 27.8	9
2	7 to < 12	7 to 10	Up to 39.7	6
Total	4 to < 12	5 to 10	Up to 39.7	15

No Treatment-Emergent SAEs Reported in INSPIRE DUCHENNE

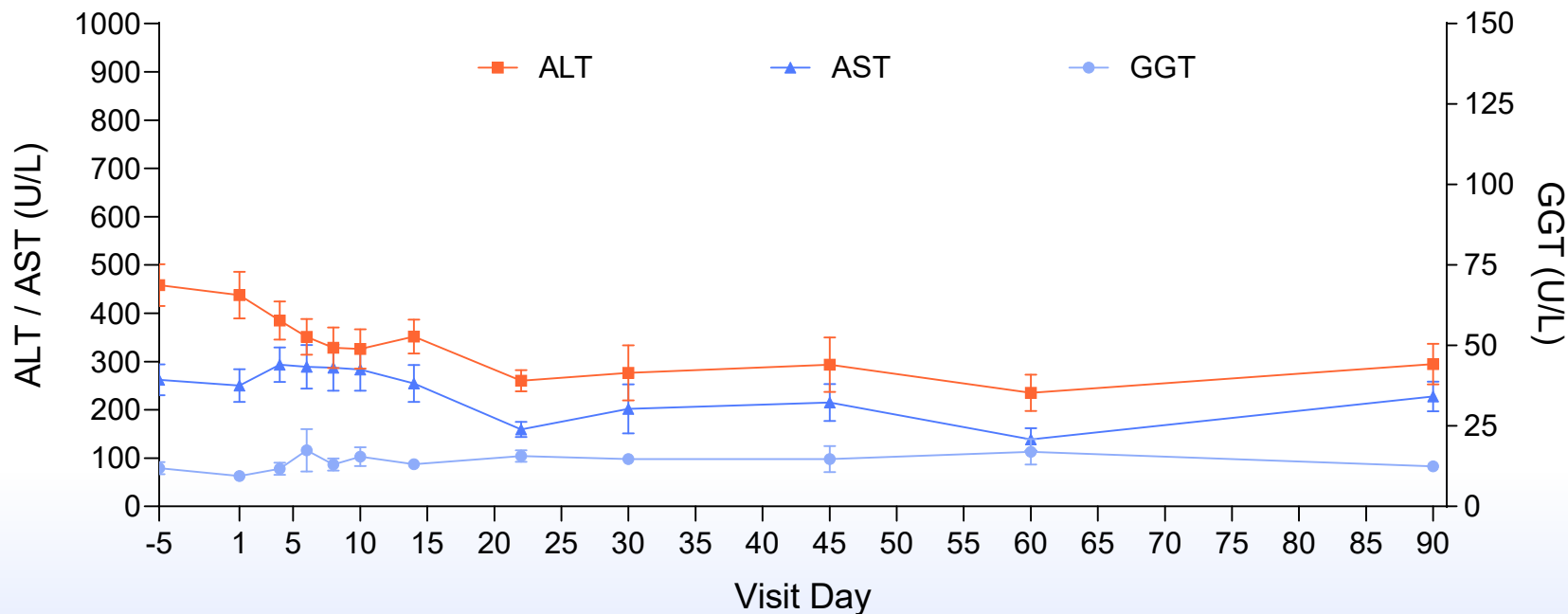
As of a data cutoff of August 12, 2025, mild to moderate AEs observed that resolved without sequelae

SGT-003 Treatment-Emergent Adverse Events (TEAEs) Data cutoff August 12, 2025		Total Participants (N=15)
		n (%)
Serious Adverse Events (SAEs)		0 (0)
Adverse Events of Special Interest (AESIs)	Hepatotoxicity	1 (6.7)*
	Thrombotic Microangiopathy	0 (0)
	Myocarditis	0 (0)
	Myositis	0 (0)
Most Common Adverse Events (AEs)	Nausea	15 (100)
	Vomiting	14 (93.3)
	Thrombocytopenia/Platelet Count Decreased	10 (66.7)
	Decreased Appetite	9 (60.0)
	Headache	6 (40.0)
*n=1 AESI of hepatotoxicity based on laboratory criteria; Grade 1 (mild) hypertransaminasaemia with no clinical symptoms.		

Liver Enzymes Remain Stable in Participants Treated with SGT-003

GGT is monitored as an important biomarker of liver injury

ALT and AST are increased at baseline in Duchenne patients but are also monitored alongside GGT to evaluate changes in liver function



GGT=gamma-glutamyl transferase; ALT: alanine aminotransferase; AST: aspartate aminotransferase
Mean $\pm$ standard error results shown for n=15 participants who reached each time point with a data cutoff of August 12, 2025.
Data on file. Solid Biosciences.

Vector Genome Copies in Day 90 Muscle Biopsies

PCR analysis demonstrated high vector genome copies in muscle

The AAV-SLB101 capsid efficiently transduces muscle



PCR

Microdystrophin protein is expressed in muscle



Western Blot



Mass Spectrometry

Microdystrophin protein is localized throughout the muscle



Immunofluorescence

Vector Genome Copies/Nucleus

Participant	Dose	Copies/Nucleus
1	1.0E14 vg/kg	19.8
2		28.6
3		7.6
Mean		18.7

SGT-003 Microdystrophin Expression in Day 90 Muscle Biopsies

Western blot and mass spectrometry demonstrated high microdystrophin protein levels

The AAV-SLB101 capsid efficiently transduces muscle



PCR

Microdystrophin protein is expressed in muscle



Western
Blot



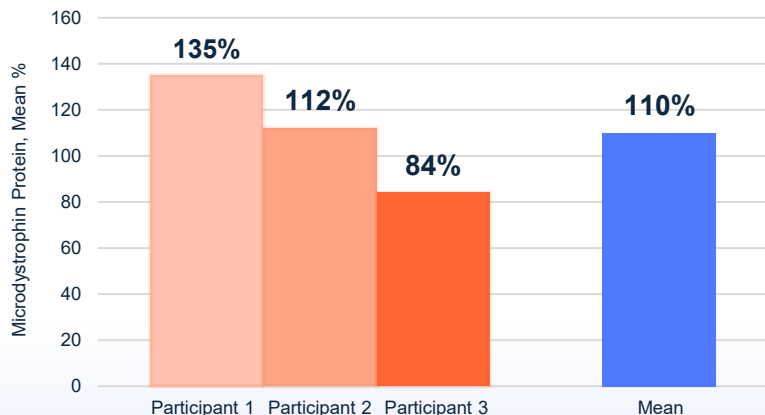
Mass
Spectrometry

Microdystrophin protein is localized throughout the muscle

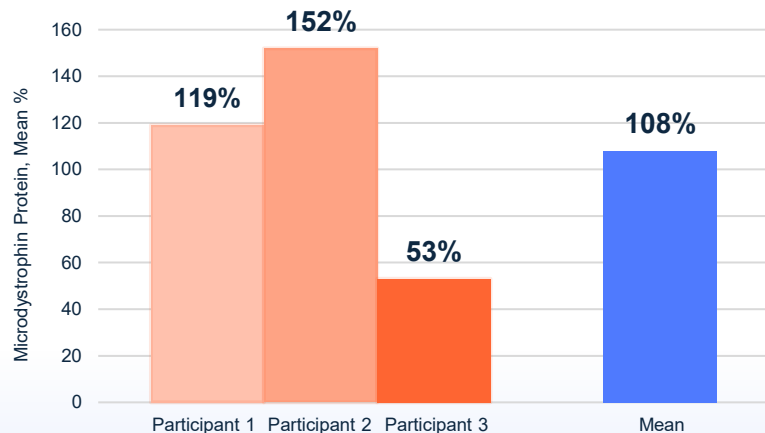


Immunofluorescence

Microdystrophin Expression Measured by Western Blot^a



Microdystrophin Expression Measured by Mass Spectrometry^a



PCR=polymerase chain reaction

^aBaseline Western blot and mass spectrometry were both 0% mean normal dystrophin.

Data on file as of February 11, 2025. Solid Biosciences.

SGT-003 Microdystrophin Protein Distribution in Day 90 Muscle Biopsies

Immunofluorescence demonstrated microdystrophin protein in a high proportion of muscle fibers

The AAV-SLB101 capsid efficiently transduces muscle



PCR

Microdystrophin protein is expressed in muscle



Western Blot



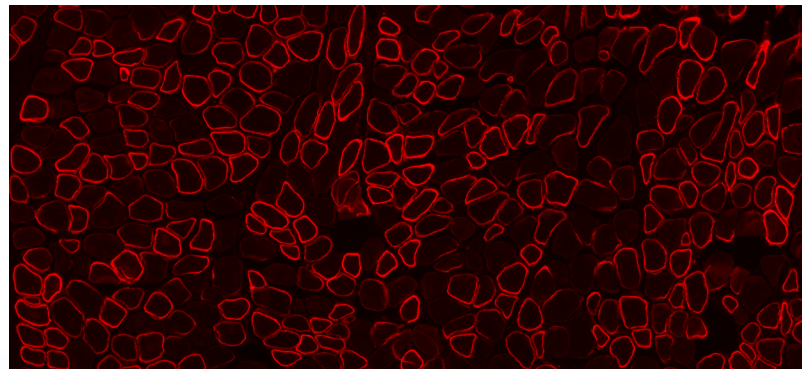
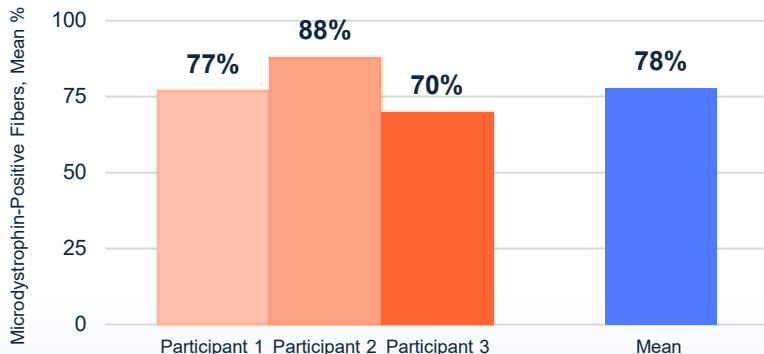
Mass Spectrometry

Microdystrophin protein is localized throughout the muscle



Immunofluorescence

**Microdystrophin-Positive Fibers
Measured by Immunofluorescence^a**



PCR=polymerase chain reaction

^aBaseline mean dystrophin-positive fibers were 1.5% measured by immunofluorescence. Dystrophin-positive fibers are not adjusted for fat and fibrosis; these are absolute numbers.

Participant 2 representative image is shown in the right panel.

Data on file as of February 11, 2025. Solid Biosciences.

Muscle Biopsies Showed Increases in Key Elements of the Dystrophin-Associated Protein Complex

Percent Positive Fibers – Microdystrophin^a

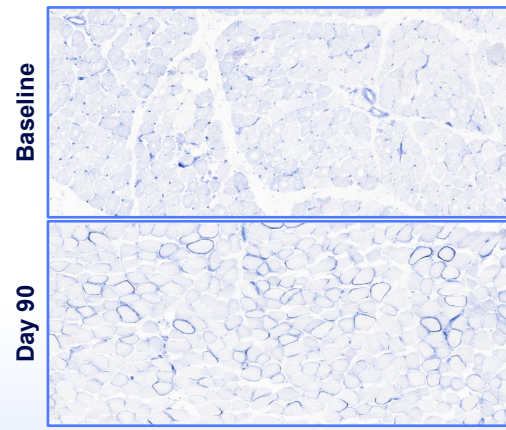
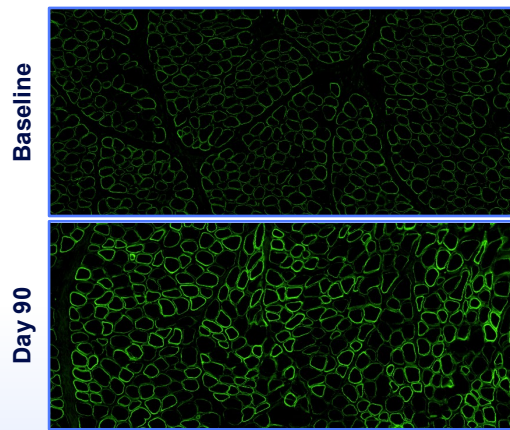
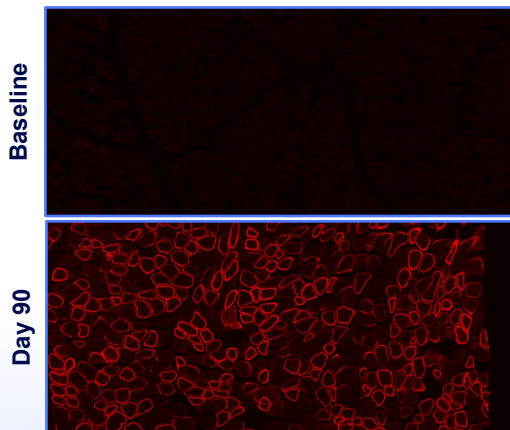
Participant	1	2	3	Mean
Day 90 Values	77%	88%	70%	78%
Baseline Values	0.8%	2.3%	1.3%	1.5%
Change From Baseline (Fold Change)	96x	38x	53x	53x

Percent Positive Fibers – β -sarcoglycan^a

Participant	1	2	3	Mean
Day 90 Values	60%	88%	63%	70%
Baseline Values	0%	2.5%	1.5%	1.3%
Change From Baseline (Fold Change)	∞	34x	41x	52x

Percent Positive Fibers – nNOS activity^a

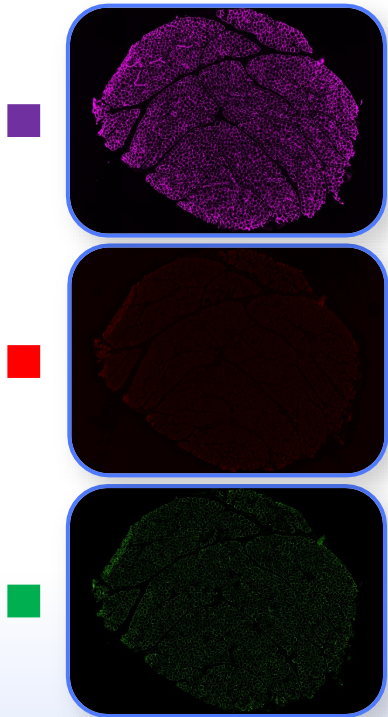
Participant	1	2	3	Mean
Day 90 Values	48%	53%	25%	42%
Baseline Values	0%	1.5%	0.5%	0.7%
Change From Baseline (Fold Change)	∞	34x	49x	62x



^aDystrophin-positive fibers are not adjusted for fat and fibrosis; these are absolute numbers. Participant 2 representative images are shown. Data on file as of February 11, 2025. Solid Biosciences.

Full Slide Scans of Muscle Biopsy Sections Showed Uniform Increases in Key Elements of the Dystrophin-Associated Protein Complex^a

Baseline

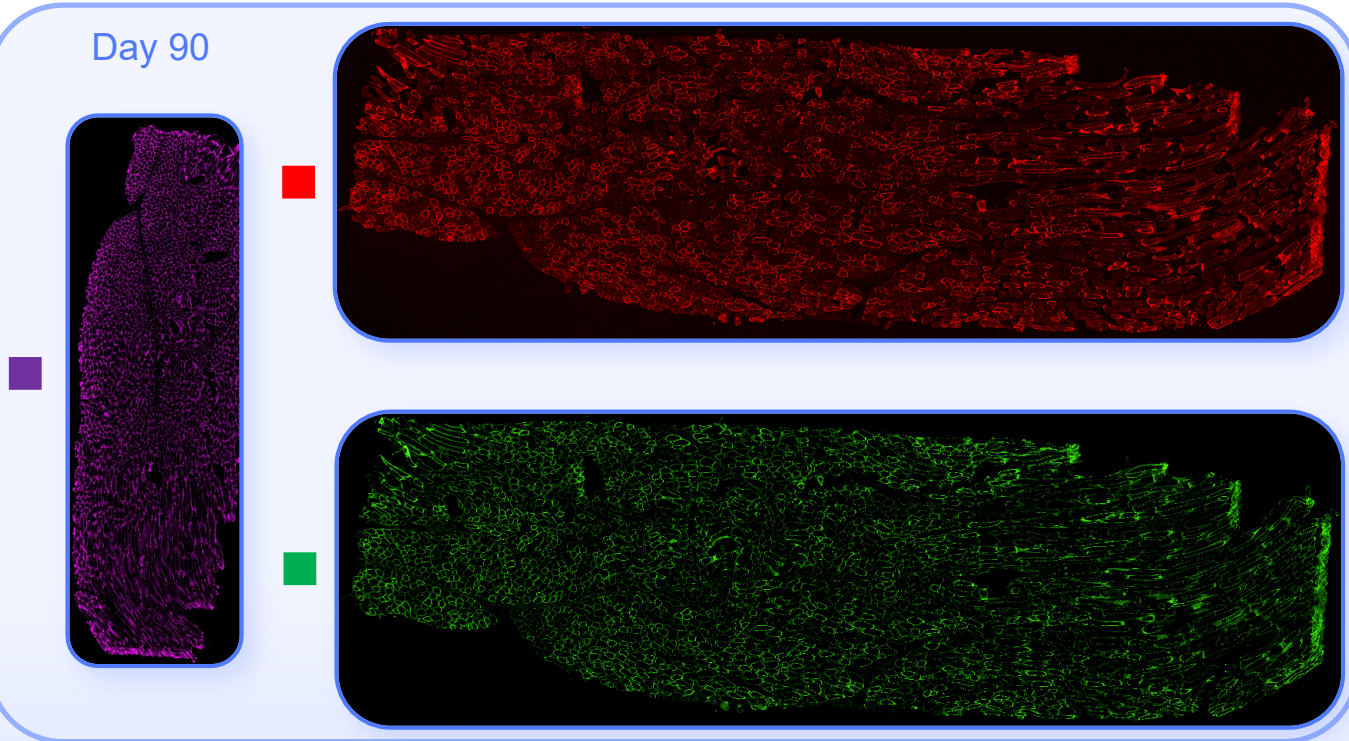


Laminin

Microdystrophin

Beta Sarcoglycan

Day 90

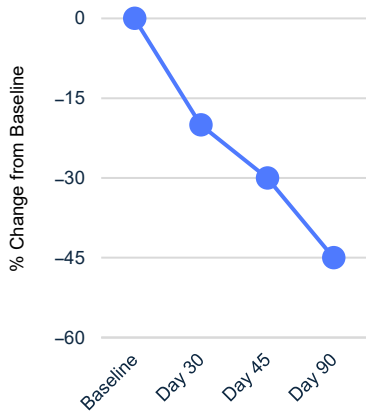


^aParticipant 2 representative images are shown. Laminin staining is used to demarcate muscle membranes.
Data on file. Solid Biosciences, 2025.

Improvements in Markers of Muscle Injury¹

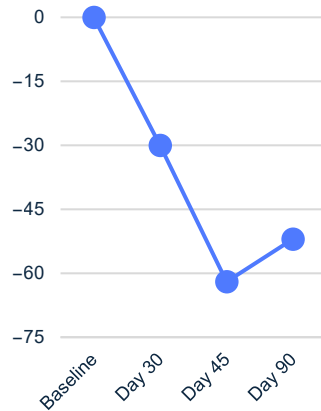
AST, ALT, CK, and LDH are released from muscle into circulation in Duchenne due to tissue damage and muscle injury²⁻⁴

SERUM AST (IU/L)^{1,a}



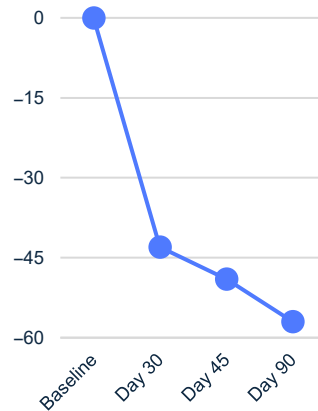
↓ Serum AST (-45%)¹

SERUM ALT (IU/L)^{1,a}



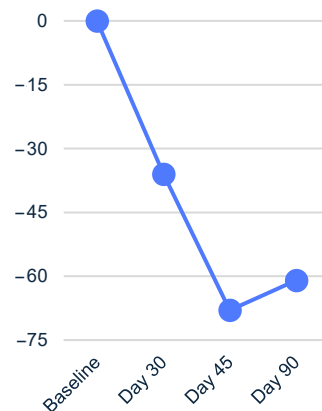
↓ Serum ALT (-54%)¹

SERUM CK (IU/L)^{1,a}



↓ Serum CK (-57%)¹

SERUM LDH (IU/L)^{1,a}



↓ Serum LDH (-60%)¹

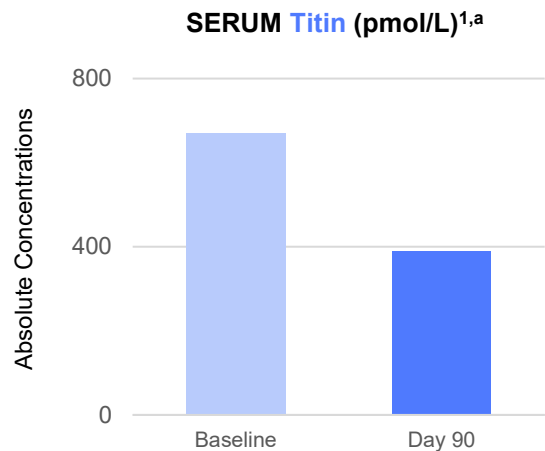
ALT=alanine aminotransferase; AST=aspartate aminotransferase; CK=creatine kinase; LDH=lactate dehydrogenase.

^aMean (n=3) change from baseline results shown.

1. Data on file. Solid Biosciences. 2. Aulbach AD, Amuzie CJ. *A Comprehensive Guide to Toxicology in Nonclinical Drug Development*. 2nd ed. 2017. 3. Kim EY, et al. *Ann Rehabil Med*. 2017;41(2):306-312. 4. Farhana A, Lappin SL. *StatPearls* [Internet]. 2023.

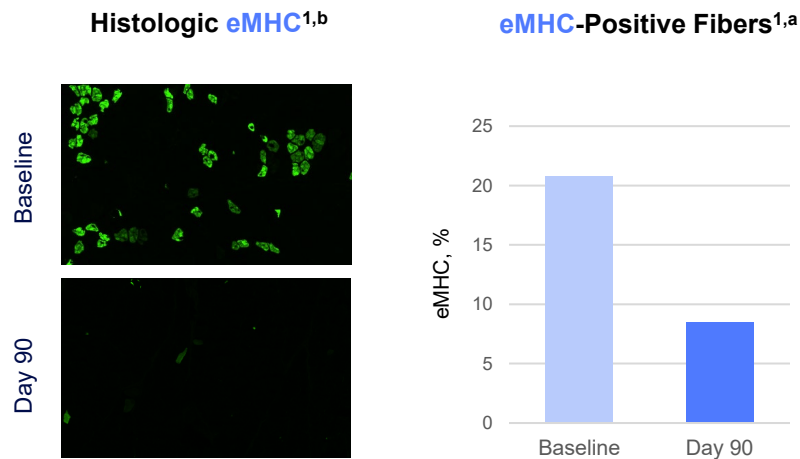
Improvements in Markers of Muscle Breakdown and Dystrophic Regeneration¹

Titin is actively degraded and released into serum and urine when muscle is damaged²



↓ Serum titin (**-42%**)¹

eMHC is expressed in dystrophic muscle fibers that have recently undergone degeneration/regeneration³



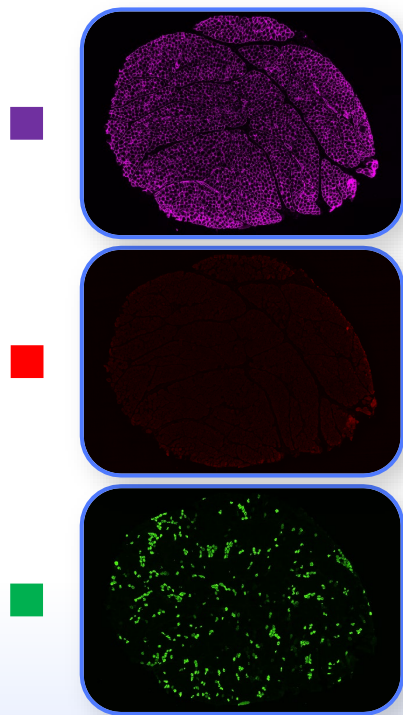
↓ Histologic eMHC (**-59%**)¹

eMHC=embryonic myosin heavy chain.

^aMean (n=3) absolute and percentage change from baseline results shown. ^bParticipant 2 representative images are shown.

Full Slide Scans of Muscle Biopsy Sections Showed Uniform Improvements in eMHC, a Marker of Muscle Breakdown and Dystrophic Regeneration^a

Baseline

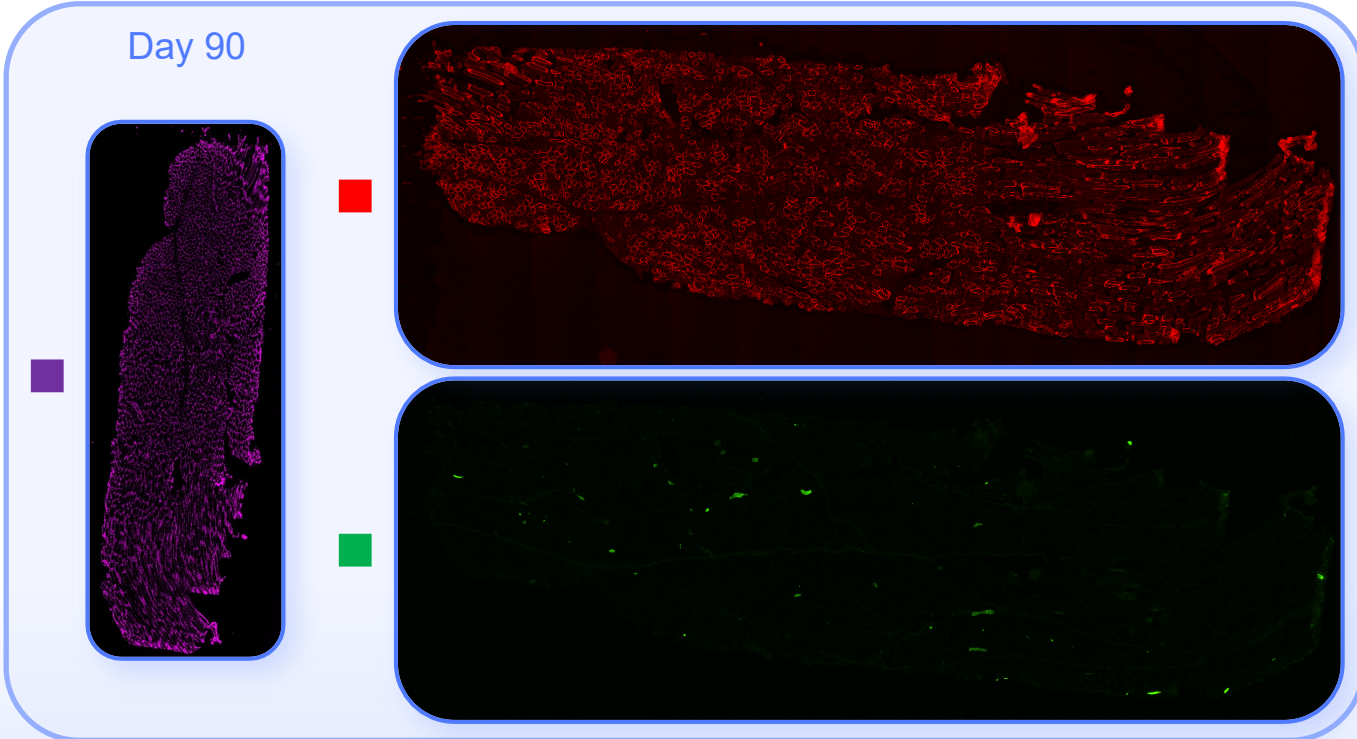


■ Laminin

■ Microdystrophin

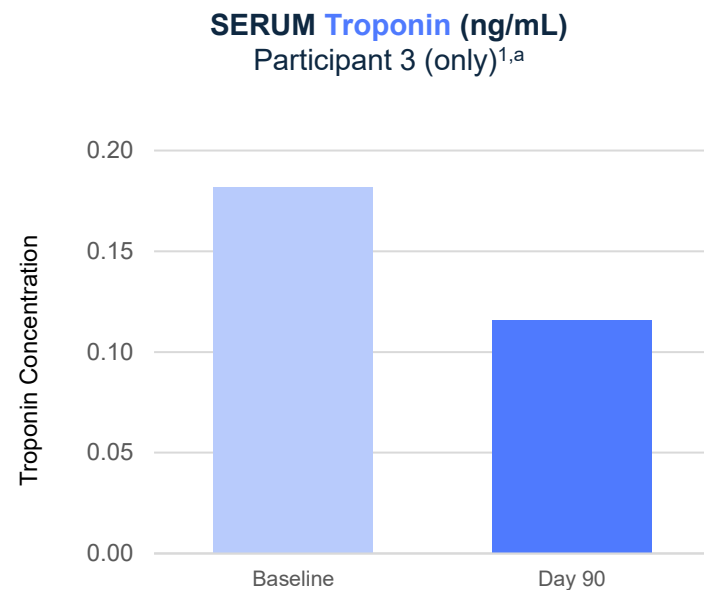
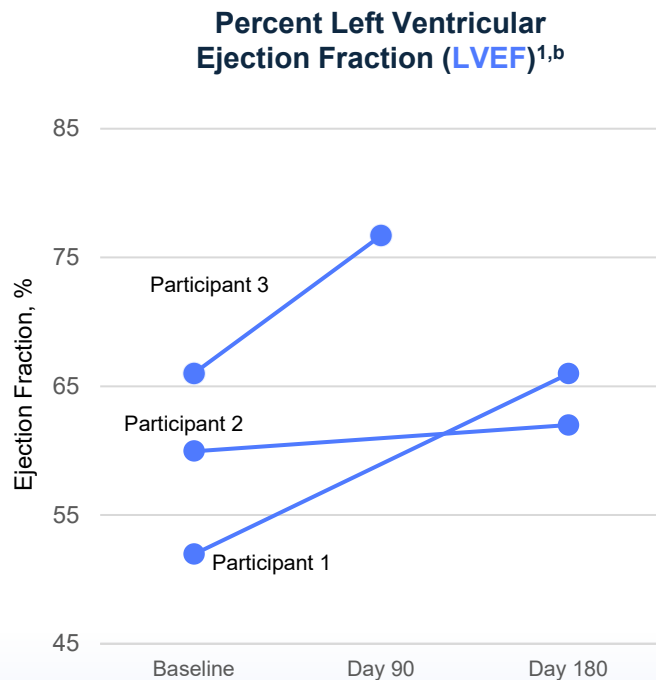
■ eMHC

Day 90



^aParticipant 2 representative images are shown. Laminin staining is used to demarcate muscle membranes
Data on file. Solid Biosciences.

Positive Changes Observed in Cardiac Markers¹



↓ Serum Troponin (**-36%**)^{1,a}

^aSerum troponin data only from Participant 3 at Day 90: Participant 3 had elevated troponin levels at baseline. Troponin levels for Participants 1 and 2 were 0 at baseline. ^bParticipant 3 has yet to reach the Day 180 follow-up as of the data cutoff. All 3 participants demonstrated LVEF above baseline at all follow-up timepoints.

1. Data on file as of February 11, 2025. Solid Biosciences. 2. Voleti S, et al. *Pediatr Cardiol.* 2020;41(6):1173-1179.

INSPIRE DUCHENNE: Current Summary

INITIAL MUSCLE BIOPSY RESULTS FOR THE FIRST 3 PARTICIPANTS REACHING DAY 90

- Mean vector genome copies per nucleus: 18.7
- Mean microdystrophin expression: 110% of normal (Western blot), 108% of normal (mass spectrometry)
- Mean microdystrophin percent-positive fibers: 78%
- Mean β -sarcoglycan percent-positive fibers: 70%
- Mean nNOS-positive fibers: 42%

MUSCLE INTEGRITY BIOMARKER RESULTS FOR THE FIRST 3 PARTICIPANTS REACHING DAY 90

- Consistent improvements across 7 biomarkers

NO SERIOUS ADVERSE EVENTS IN THE 15 PARTICIPANTS TREATED (DATA CUTOFF AUGUST 12, 2025)

- Most common treatment-related adverse events observed were nausea, vomiting, thrombocytopenia/platelet count reduced, decreased appetite, and headache

Acknowledgments

Thank you to the study participants and families!

Thank you to participating clinical sites, investigative teams, and study partners!

Nationwide Children's Hospital (Columbus, OH)

- Kevin Flanigan, MD

Arkansas Children's Hospital (Little Rock, AR)

- Aravindhan Veerapandian, MD

Children's Hospital of The King's Daughters (Norfolk, VA)

- Crystal Proud, MD

Rare Disease Research (Atlanta, GA)

- Renata Shih, MD

Lurie Children's Hospital (Chicago, IL)

- Nancy Kuntz, MD

Washington University at St. Louis (St. Louis, MO)

- Craig Zaidman, MD

University of California, Davis (Sacramento, CA)

- Craig McDonald, MD

University of California, Los Angeles (Los Angeles, CA)

- Perry Shieh, MD, PhD

The Hospital for Sick Children (Toronto, ON)

- Hernan Gonorazky, MD

Gemelli Hospital (Rome, Italy)

- Eugenio Mercuri, MD, PhD

Great Ormond Street Hospital (London, UK)

- Francesco Muntoni, MD

Seattle Children's Hospital (Seattle, WA)

- Alicia Henriquez, MD



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