

# Safety, $\beta$ -Sarcoglycan Expression, and Functional Outcomes From Systemic Gene Transfer of rAAVrh74.MHCK7.hSGCB in Limb-Girdle Muscular Dystrophy Type 2E/R4

Louise R. Rodino-Klapac,<sup>1</sup> Eric R. Pozsgai,<sup>1,2</sup> Sarah Lewis,<sup>1,2</sup> Danielle A. Griffin,<sup>1,2</sup> Aaron S. Meadows,<sup>1,3</sup> Kelly J. Lehman,<sup>2</sup> Kathleen Church,<sup>2</sup> Natalie F. Reash,<sup>2</sup> Megan A. Iammarino,<sup>2</sup> Linda P. Lowes,<sup>2</sup> Erica Koenig,<sup>1</sup> Sarah Neuhaus,<sup>1</sup> Xiaoxi Li,<sup>1</sup> Jerry R. Mendell<sup>2,4</sup>

<sup>1</sup>Sarepta Therapeutics, Inc., Cambridge, MA, USA; <sup>2</sup>Center for Gene Therapy, The Research Institute at Nationwide Children's Hospital, Columbus, OH, USA; <sup>3</sup>Wexner Medical Center, The Ohio State University, Columbus, OH, USA;

<sup>4</sup>Department of Pediatrics and Neurology, The Ohio State University, Columbus, OH, USA

Presented at the 2021 Muscular Dystrophy Association Virtual Clinical & Scientific Conference, March 15–18, 2021



# Disclosures

- LRR-K, ERP, SL, DAG, ASM, EK, SN, and XL are employees of Sarepta Therapeutics, Inc., and may have stock options
- JRM received financial support from Sarepta Therapeutics, Inc., for the travel and accommodation costs of study participants
- KL, KC, NR, and MI have no conflicts to disclose
- LL received fees from Sarepta for licensure of the natural history data set
- This study (NCT03652259) was funded by Sarepta Therapeutics, Inc.
- rAAVrh74.MHCK7.hSGCB (SRP-9003) is an investigational therapy and has not been reviewed or approved by the FDA
- Medical writing support was provided by Paraskevi Briassouli, PhD, of Eloquent Scientific Solutions, and funded by Sarepta Therapeutics, Inc.

# SRP-9003: Investigational gene therapy for limb-girdle muscular dystrophy type 2E/R4 (LGMD2E/R4)

- LGMD2E/R4 is caused by mutations in the *SGCB* gene<sup>1,2</sup>
- Adeno-associated virus (AAV)–mediated gene transfer therapy to express full-length  $\beta$ -sarcoglycan (SGCB) has potential to treat LGMD2E/R4

## SRP-9003 construct: Self-complementary AAV vector

### rAAVrh74



*Efficient skeletal and cardiac muscle transduction<sup>3</sup>*

### MHCK7 PROMOTER



*Optimized for expression in skeletal and cardiac muscle<sup>4-6</sup>*

### hSGCB cDNA



*Full-length human SGCB gene transfer*

**Objective: To report the interim findings of an ongoing Phase 1/2 clinical gene transfer trial delivering SRP-9003 to patients with LGMD2E/R4 (NCT03652259)**

# Study design: First-in-human, open-label, Phase 1/2 study

3-year study: Open label, single dose

Day 0      Day 60      Year 2      Up to 3 years

Cohort 1: SRP-9003  $1.85 \times 10^{13}$  vg/kg (n=3)<sup>a</sup>

Day -1: Prednisone 1 mg/kg daily for 30 days, taper

Cohort 2: SRP-9003  $7.41 \times 10^{13}$  vg/kg (n=3)<sup>b</sup>

Day -1: Prednisone 1 mg/kg daily for 60 days, taper

Open label

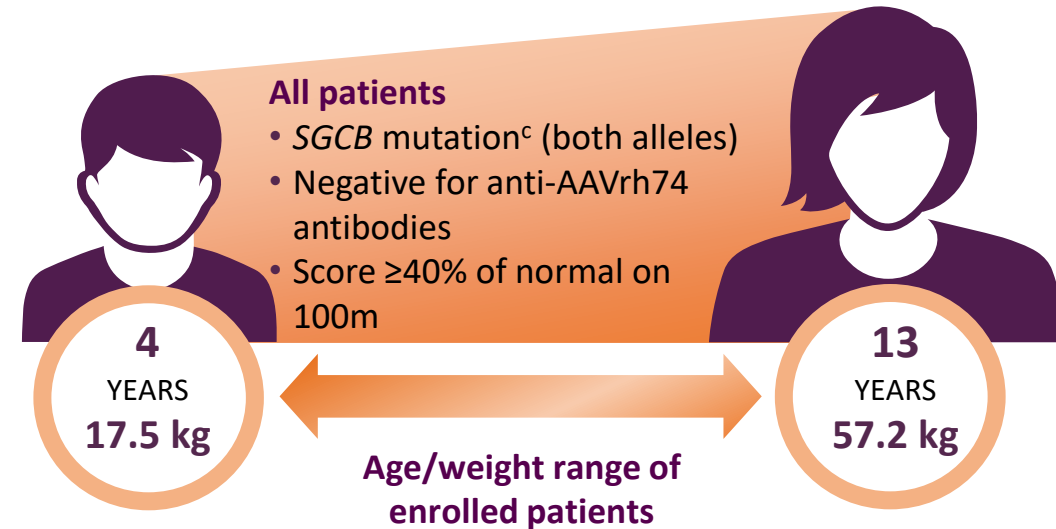
Screening

60 days initial monitoring

3 years safety monitoring

EOS visit

● Muscle biopsy



- **Primary endpoint:** Safety
- **Secondary endpoint:** SGCB expression at Week 8
- **Other endpoints**
  - Change in creatine kinase (CK) from baseline
  - Functional endpoints (NSAD and timed tests: 100m, 10m, 4-stair climb, and time to rise)

10m=10-m timed test; 100m=100-m timed test; EOS=end of study; NSAD=North Star Assessment for Limb-girdle Type Muscular Dystrophies.

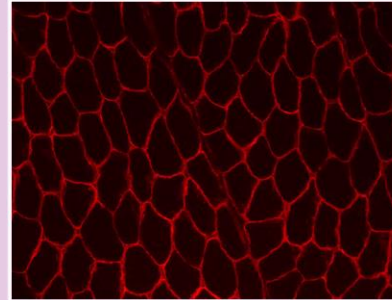
<sup>a</sup> $1.85 \times 10^{13}$  vg/kg measured using linear reference plasmid DNA quantitative polymerase chain reaction (qPCR); supercoiled reference DNA equivalent is  $5 \times 10^{13}$  vg/kg;

<sup>b</sup> $7.41 \times 10^{13}$  vg/kg measured using linear reference plasmid DNA qPCR; supercoiled reference DNA equivalent is  $2 \times 10^{14}$  vg/kg;

<sup>c</sup>Patients 1–5 had missense mutations in exons 3–6, and Patient 6 had a nonsense mutation.

# Robust expression and sarcolemmal localization of SGCB at Day 60 post infusion

Normal control



Patient:

E01

E02

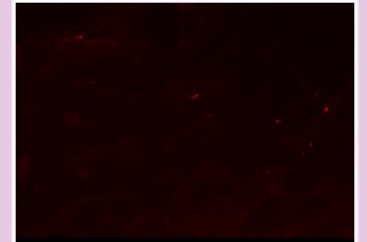
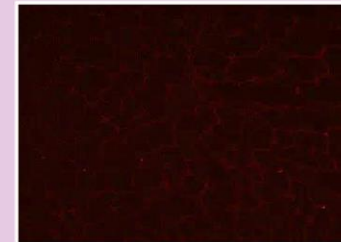
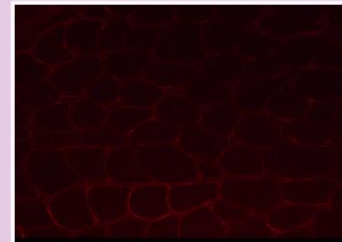
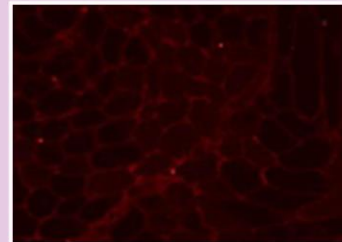
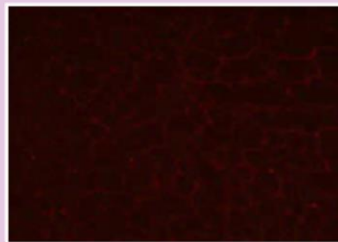
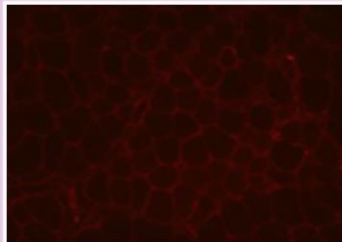
E03

E04

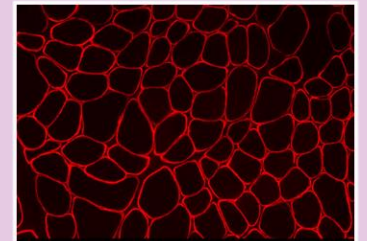
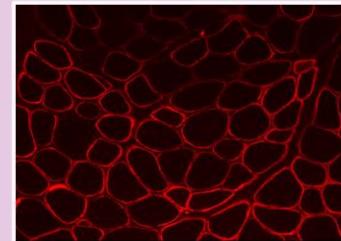
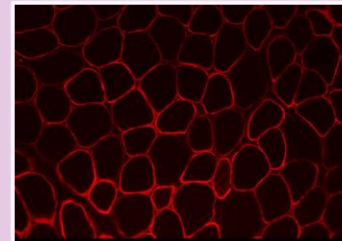
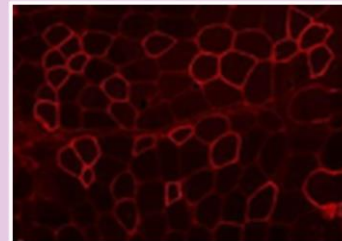
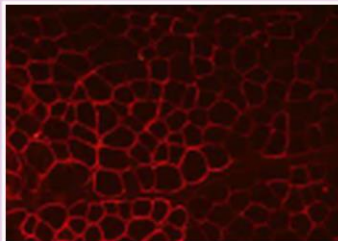
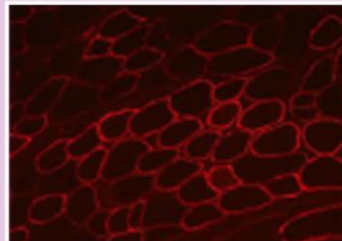
E05

E07

Pretreatment



Post treatment



# Expression of SGCB at 60 days post infusion was sustained for 2 years in Cohort 1

## SGCB protein expression at skeletal muscle sarcolemma

**Cohort 1**  
( $1.85 \times 10^{13}$  vg/kg)<sup>a</sup>

Time point	Western blot % NC (SD)	% SGCB+ fibers % NC (SD)	SGCB intensity % NC (SD)	Vector copies per nucleus (SD)	
				qPCR	ddPCR
Day 60	36 (2.7)	51 (10.6)	47 (9.5)	0.59 (0.4)	—
Year 2	54 (16.1)	48 (22.0)	35 (22.9)	0.13 (0.1)	0.46 (0.4)

**Cohort 2**  
( $7.41 \times 10^{13}$  vg/kg)<sup>b</sup>

Time point	Western blot % NC (SD)	% SGCB+ fibers % NC (SD)	SGCB intensity % NC (SD)	Vector copies per nucleus (SD)	
				qPCR	ddPCR
Day 60	62 (8.7)	72 (6.2)	73 (21.8)	4.24 (2.8)	2.26 (0.9)

**A dose-response in full-length SGCB protein expression was observed at Day 60 and sustained at 2 years in Cohort 1**

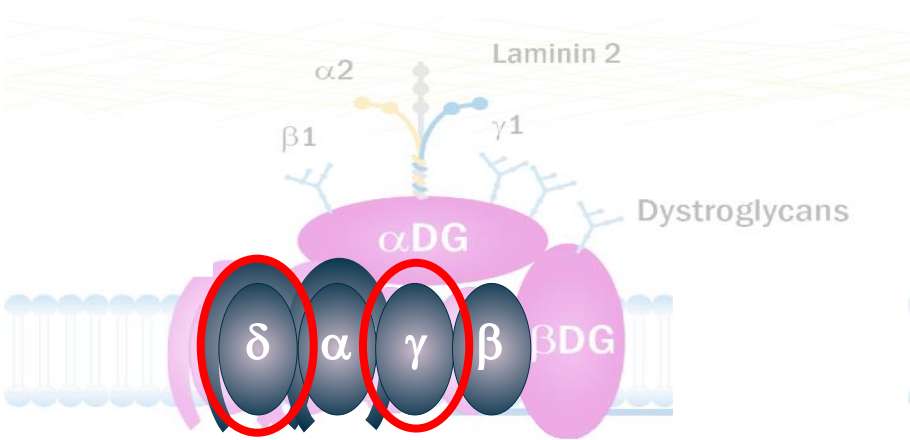
ddPCR=droplet digital PCR; NC=normal control; PCR=polymerase chain reaction; qPCR=quantitative PCR; SGCB= $\beta$ -sarcoglycan. Values are mean (SD).

<sup>a</sup> $1.85 \times 10^{13}$  vg/kg measured using linear reference plasmid DNA qPCR; supercoiled reference DNA equivalent is  $5 \times 10^{13}$  vg/kg;

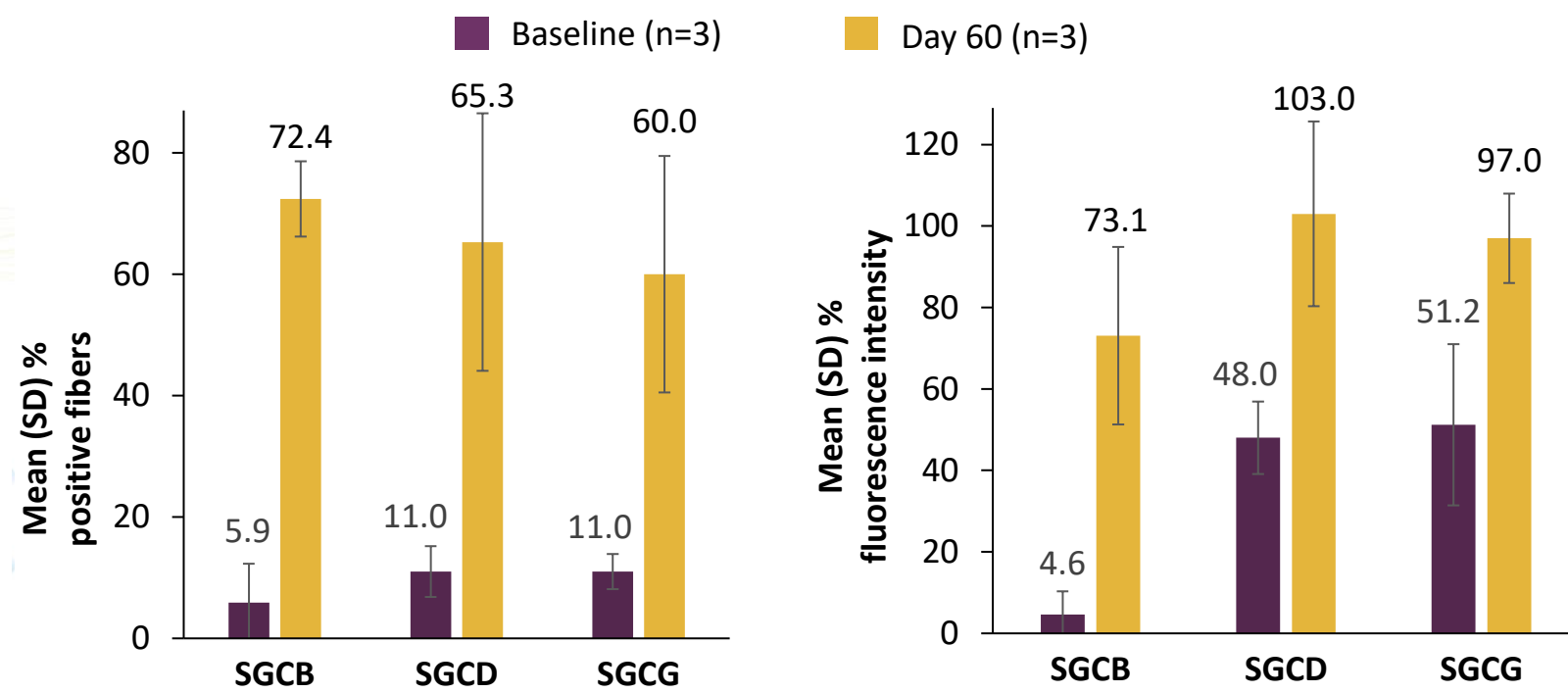
<sup>b</sup> $7.41 \times 10^{13}$  vg/kg measured using linear reference plasmid DNA qPCR; supercoiled reference DNA equivalent is  $2 \times 10^{14}$  vg/kg.

# Expression of SGCB results in increased expression of SGCD and SGCG, demonstrating reconstitution of the sarcoglycan complex within the DAPC

## Sarcoglycan complex

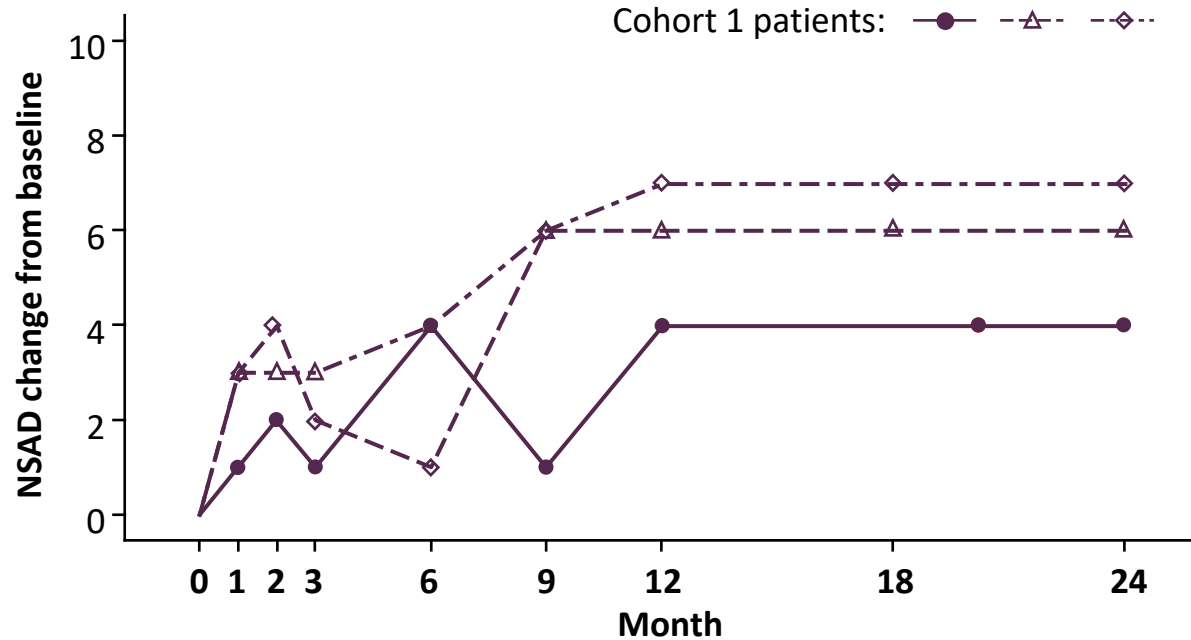


## Expression of the dystrophin-associated protein complex (DAPC) (Cohort 2, Day 60)

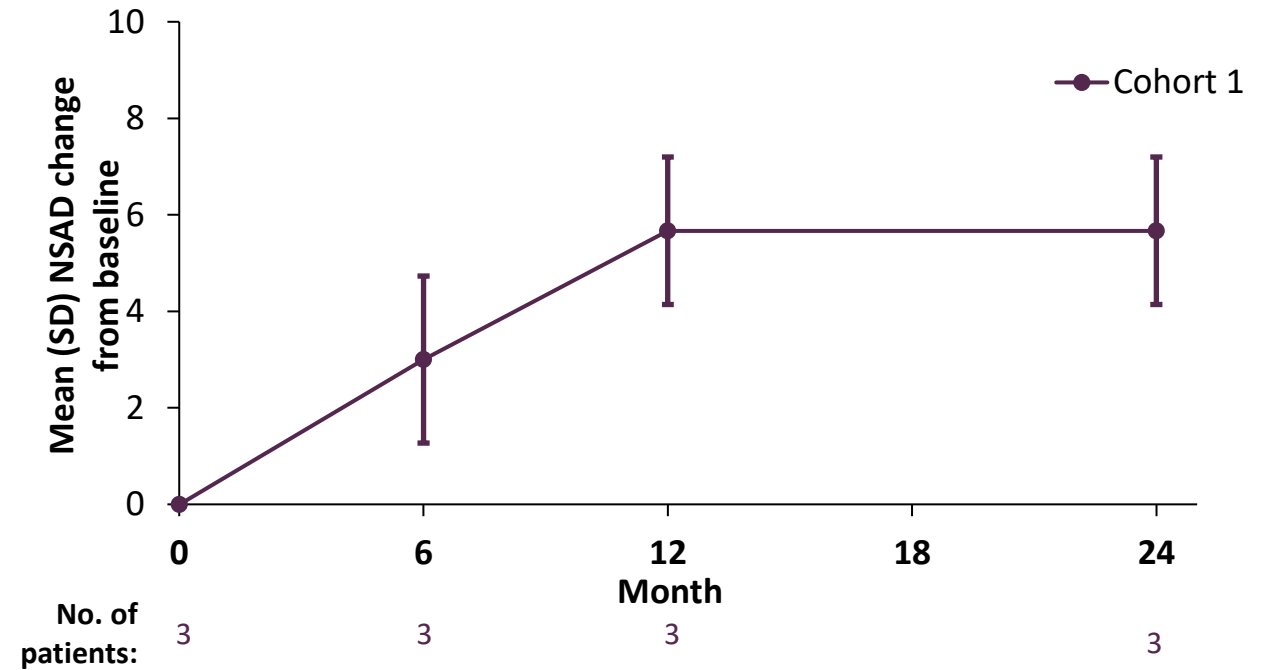


# Cohort 1: SRP-9003 treatment resulted in improvement in NSAD total score sustained for 2 years

Individual patient data



Mean data

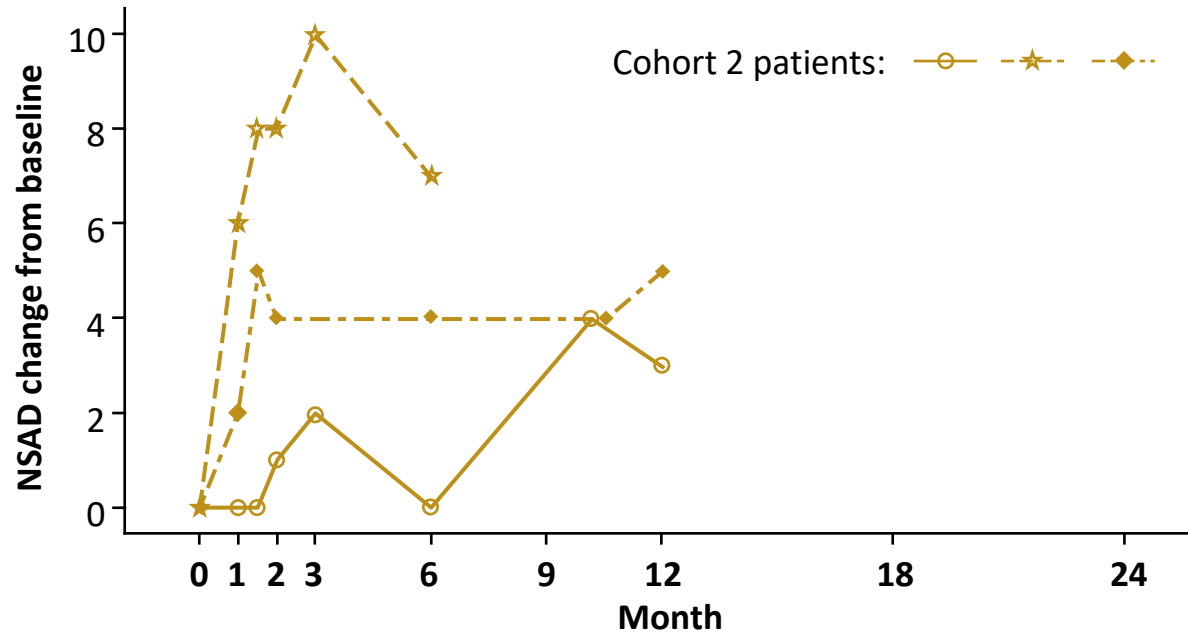


Patients treated with SRP-9003 in Cohort 1 demonstrated improvements in NSAD that were sustained for 2 years

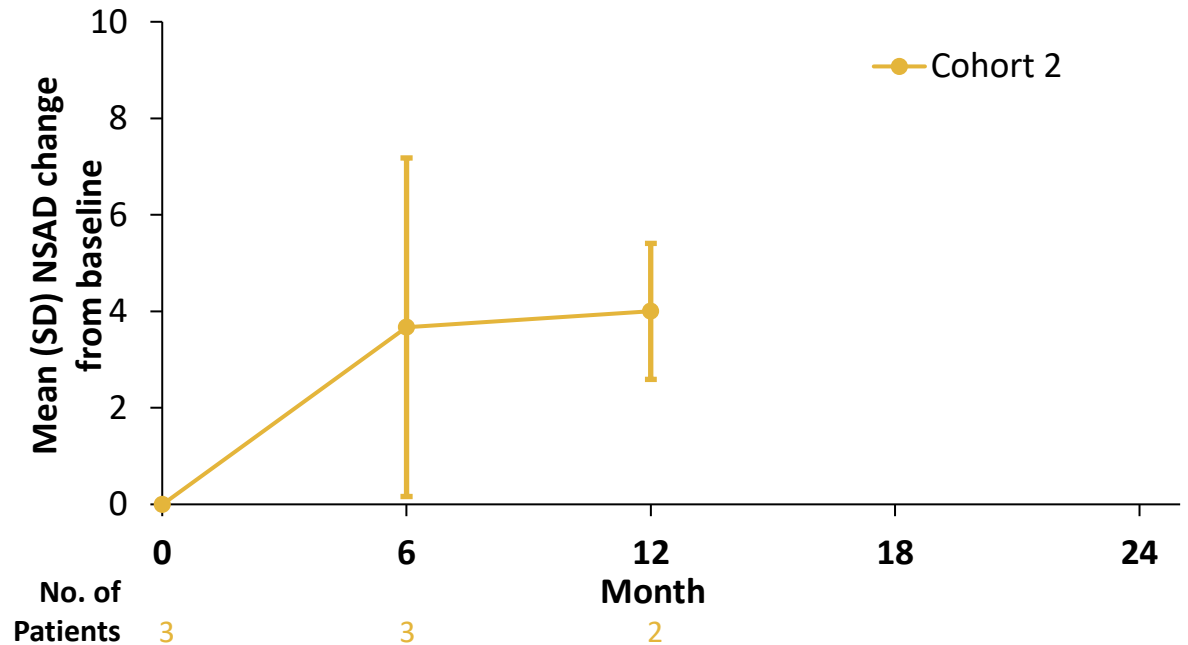


# Cohort 2: SRP-9003 treatment improved NSAD total score

Individual patient data



Mean data



Patients treated with SRP-9003 in Cohort 2 demonstrated improvements in NSAD up to 1 year

# SRP-9003 treatment resulted in sustained improvements in timed function tests

Mean (SD) change from baseline (s)	Cohort 1 ( $1.85 \times 10^{13}$ vg/kg) <sup>a</sup>			Cohort 2 ( $7.41 \times 10^{13}$ vg/kg) <sup>b</sup>	
	6 months	12 months	24 months	6 months	12 months
Time to rise	-0.2 (0.8)	-0.8 (0.4)	-0.6 (0.2)	-1.3 (0.9)	-1.1 (1.1)
4-stair climb	-0.5 (0.4)	-0.5 (0.3)	-0.3 (0.4)	-0.4 (0.3)	-0.4 (0.0)
100m	-3.8 (2.9)	-5.3 (3.2)	-2.8 (6.4)	-6.3 (6.7)	-7.9 (5.4)
10m	-0.6 (0.3)	-0.6 (0.2)	-0.2 (0.5)	-0.6 (0.6)	-0.6 (0.2)

Negative numbers correspond to faster test times.

**Patients treated with SRP-9003 demonstrated improvements over baseline in timed function tests, which were sustained for 2 years in Cohort 1 and 1 year in Cohort 2**

10m=10-m timed test; 100m=100-m timed test.

<sup>a</sup> $1.85 \times 10^{13}$  vg/kg measured using linear reference plasmid DNA quantitative polymerase chain reaction (qPCR), supercoiled reference DNA equivalent is  $5 \times 10^{13}$  vg/kg;

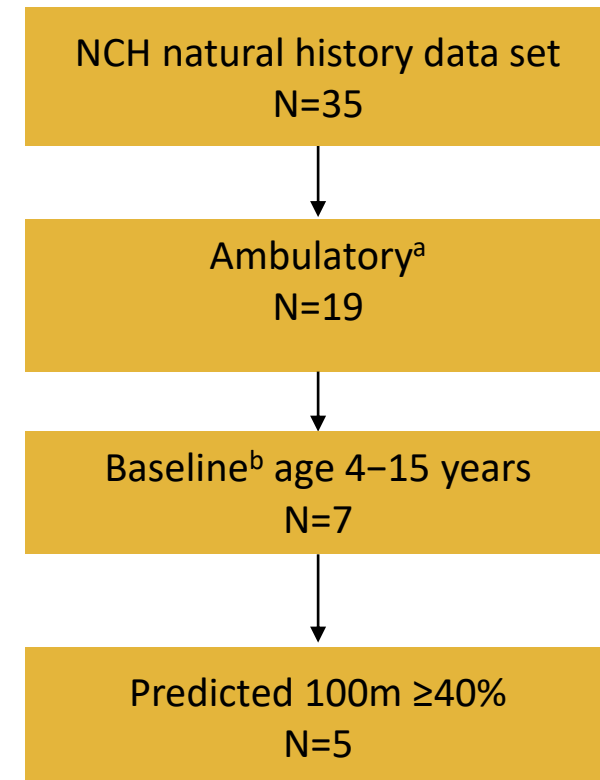
<sup>b</sup> $7.41 \times 10^{13}$  vg/kg measured using linear reference plasmid DNA qPCR, supercoiled reference DNA equivalent is  $2 \times 10^{14}$  vg/kg.

# Selection of natural history control cohort for comparison with SRP-9003-101

**Selected comparison cohort from Nationwide Children's Hospital (NCH) natural history data set, based on the same key inclusion criteria as in Study SRP-9003-101**

- Ambulatory status (yes)
- Baseline age (4–15 years)
- Predicted 100m ( $\geq 40\%$ )

## NCH LGMD2E/R4 natural history cohort



# Baseline comparison of SRP-9003–treated patients vs natural history cohort

## Baseline characteristics comparison

	SRP-9003-101 (N=6)	NCH (N=5)
Age (years)	10.0 (3.5)	9.8 (3.2)
Male, n (%)	3 (50)	3 (60)
NSAD score	41.2 (3.7)	49.0 (3.9)
100m (s)	51.4 (10.5)	38.9 (3.9)
10m (s)	5.1 (0.9)	4.4 (0.3) <sup>a</sup>

Values are mean (SD) unless noted otherwise.

Age and sex are well balanced between 9003-101 patients and the NCH control cohort

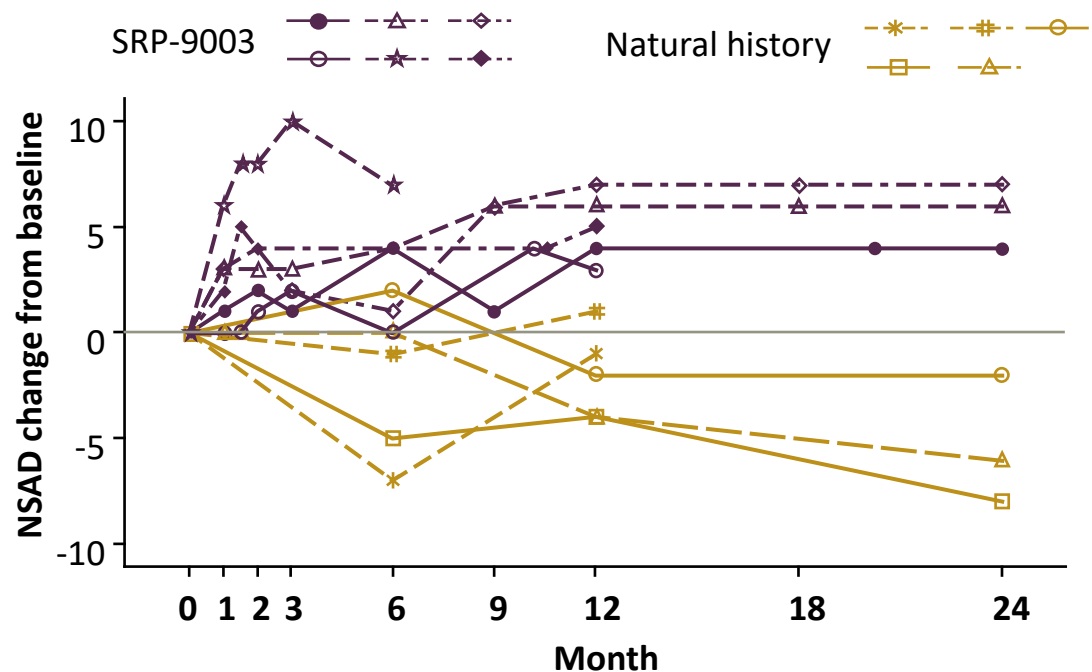
Baseline functional endpoints scores are *higher* in the NCH control cohort

Explored alternative matching criteria; however, still unable to achieve balanced baseline functions

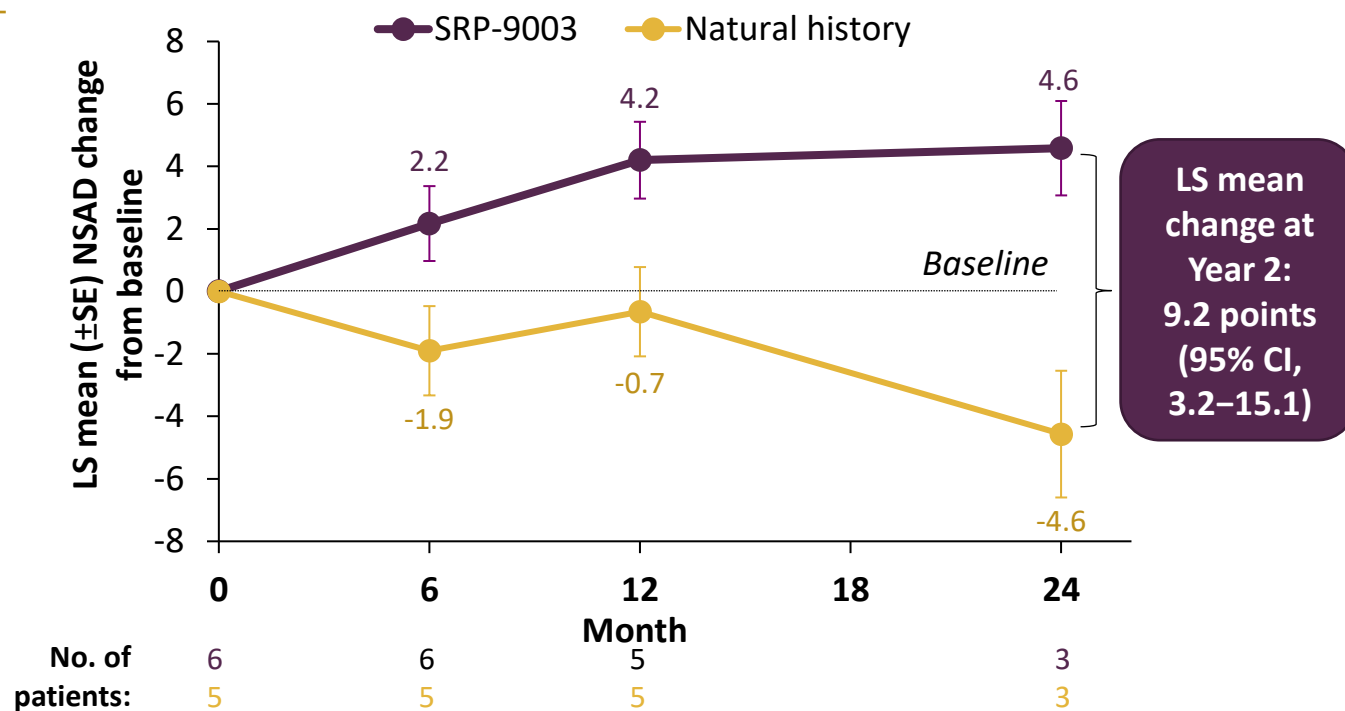
Therefore, used mixed-model repeated measures (MMRM) analysis to adjust for baseline function

# SRP-9003–treated patients display an improvement in total NSAD score vs natural history

NSAD change from baseline: Individual patient data



NSAD change from baseline: LS means by cohort



**Patients treated with SRP-9003 demonstrated clinically meaningful improvements in functional outcomes in an exploratory comparison vs an LGMD2E/R4 natural history cohort, as measured by NSAD**

LGMD2E/R4=limb-girdle muscular dystrophy type 2E/R4; LS=least-squares; MMRM=mixed-model repeated measures; NSAD=North Star Assessment for Limb-girdle Type Muscular Dystrophies. MMRM analysis included fixed effects for treatment arm, visit, and treatment arm by visit interaction, and baseline NSAD, baseline 100m, and baseline 10m as continuous covariates; the first-order autoregressive structure was used for variance-covariance matrix of within-patient errors.

# Safety results reinforce favorable safety profile, with no new safety signals

## Cohort 1 as of January 14, 2021 (n=3)

- 2 patients had elevated liver enzymes, 1 of which was designated a serious adverse event (SAE) and associated with transient increase in bilirubin
  - Occurred during or after steroid tapering; resolved within days following supplemental steroid treatment
- 1 patient experienced mild vomiting which resolved within 1 day without treatment

## Cohort 2 as of January 14, 2021 (n=3)

- Majority of AEs were mild to moderate (e.g., vomiting, pain in extremity) and resolved
- 1 treatment-related SAE observed
  - Dehydration resulting from vomiting 3 days after infusion; resolved within 2 days with treatment
- 1 patient had mildly elevated gamma glutamyl transferase (GGT)
  - Returned to within normal limits while on tapering dose of steroids; GGT levels did not increase after steroid treatment
- No stopping/discontinuation rules were triggered by AEs
- One of the participants in this trial died unexpectedly due to a recreational accident unrelated to the study

## Both cohorts

- No other laboratory abnormalities were suggestive of safety concerns
  - No decreases in platelet counts observed outside the normal range
  - No clinical sequelae associated with complement activation

**Results show no new safety signals, and treatment-related AEs occurred early and were transient and manageable**

# Conclusions

- This interim analysis reinforces the favorable safety profile of systemically administered SRP-9003
- SRP-9003 showed efficient transduction and drove robust, dose-dependent SGCB protein expression in all patients at Day 60, resulting in reconstitution of the sarcoglycan complex; SGCB expression was sustained up to 2 years
- Creatine kinase decreased by 77% at Year 2 in Cohort 1 and 74% at Year 1 in Cohort 2 (data not presented)
- Patients treated with SRP-9003 demonstrated improvements over baseline in NSAD and timed function tests that were sustained up to 2 years in Cohort 1 and 1 year in Cohort 2
- Exploratory post hoc analysis showed SRP-9003-treated patients had clinically meaningful improvements in functional outcomes, as measured by NSAD, compared with a natural history cohort
- The observed durable treatment effect provides proof of concept and supports further clinical assessment of SRP-9003 gene transfer therapy in patients with LGMD2E/R4

## Key Takeaway:

**Persistence of SRP-9003 in transduced muscle continues to drive meaningful levels of SGCB expression over time, leading to sustained functional improvements**

# Acknowledgments

- **Sarepta Therapeutics, Inc., and the authors thank the patients and their families for their participation in the study**
- **Sarepta Therapeutics, Inc., thanks Linda Lowes, Jerry R. Mendell, and their teams at Nationwide Children's Hospital for their contributions to the study and providing data on the LGMD2E/R4 natural history cohort**
  - Lindsay Alfano
  - Megan Iammarino
  - Brenna Powers
  - Natalie Reash
  - Kiana Shannon