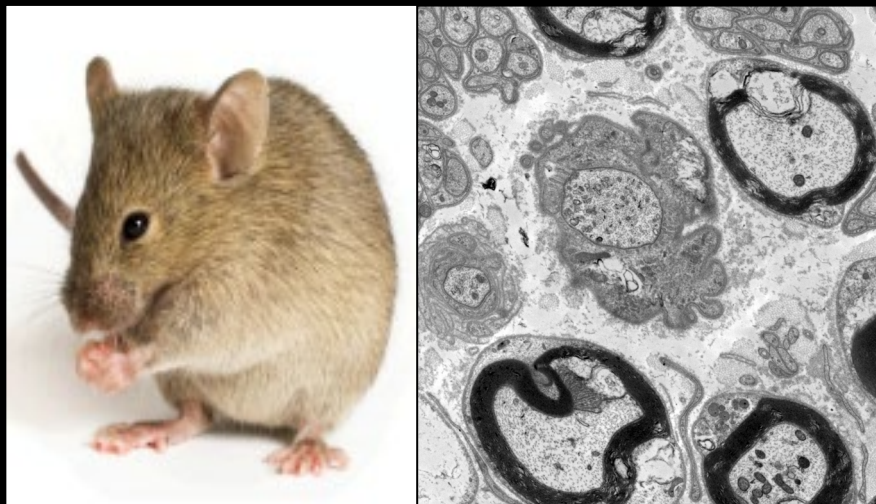
A grayscale electron micrograph of neural tissue, showing various cellular structures, myelin sheaths, and axons. The image is used as a background for the text.

Precision mouse models of childhood ALS caused by excessive sphingolipid synthesis

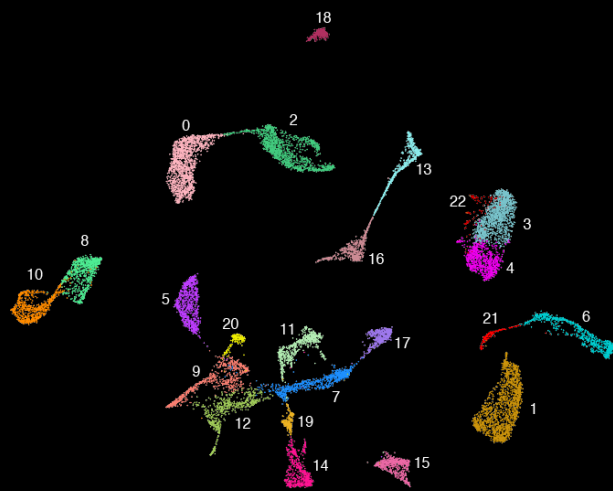
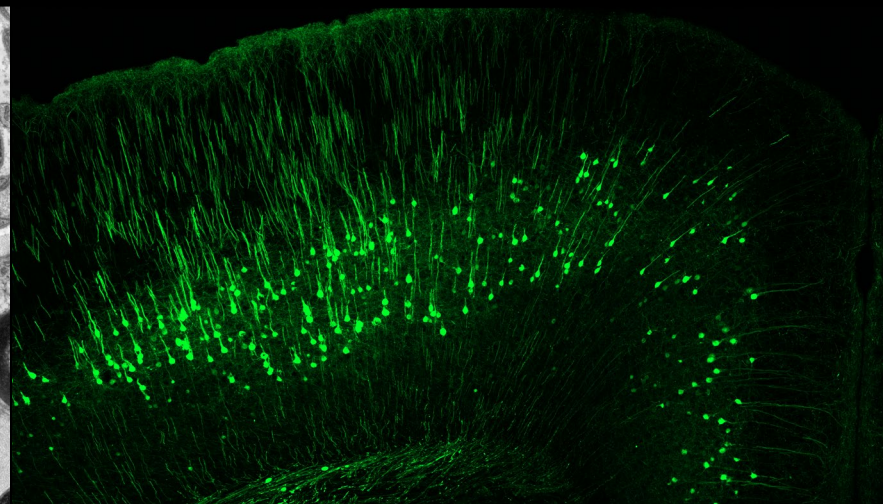
Claire Le Pichon, PhD
National Institutes of Health
NICHD Council Meeting
Sept 12, 2022

claire.lepichon@nih.gov

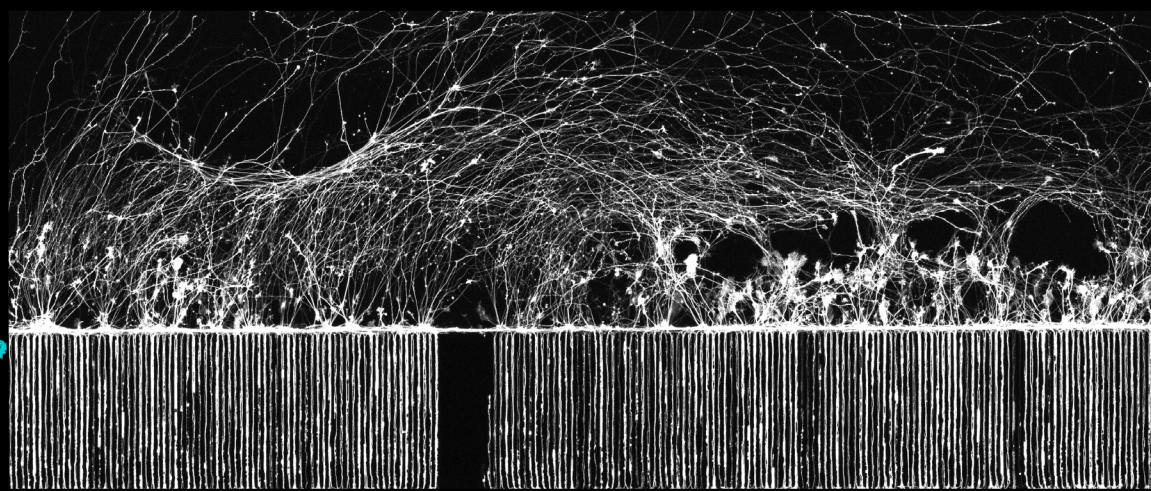
Precision mouse models of disease



Traumatic injury

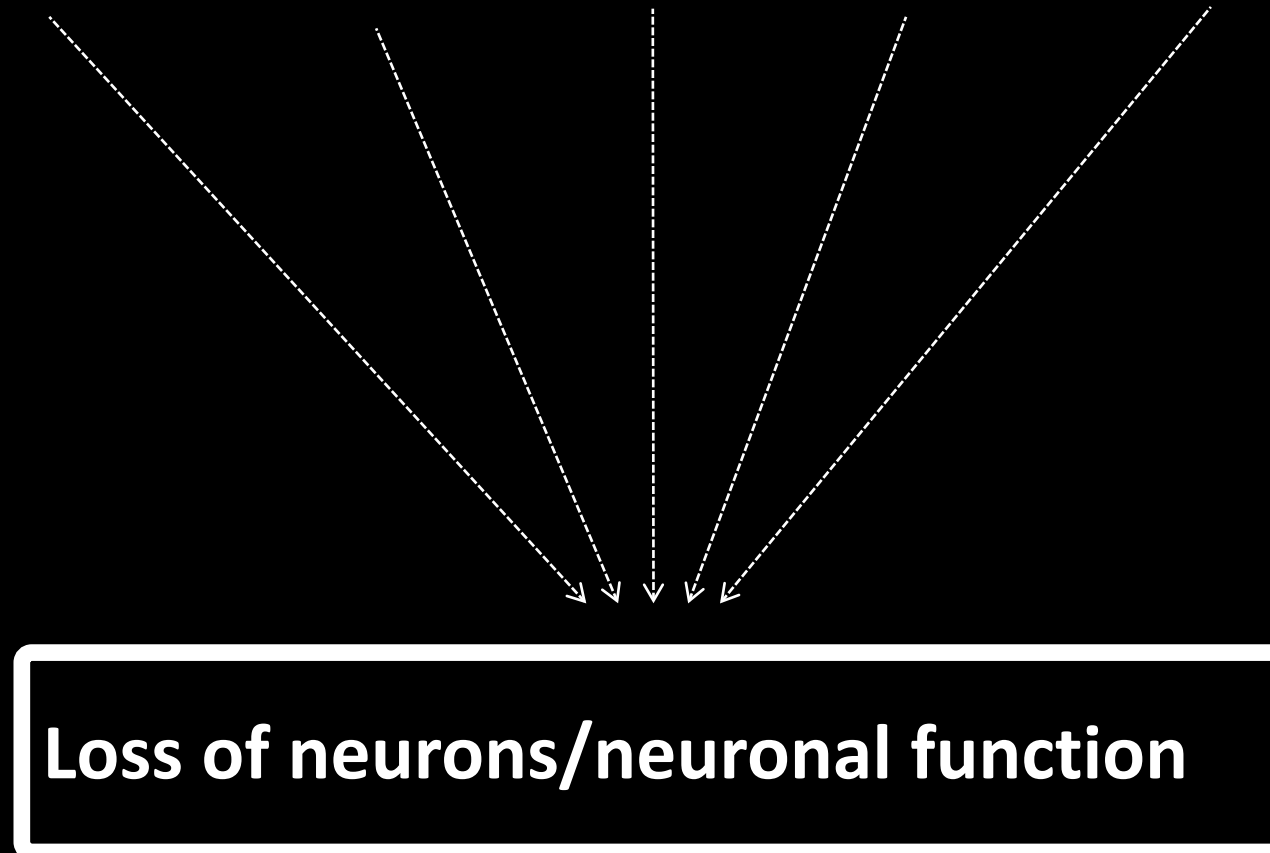
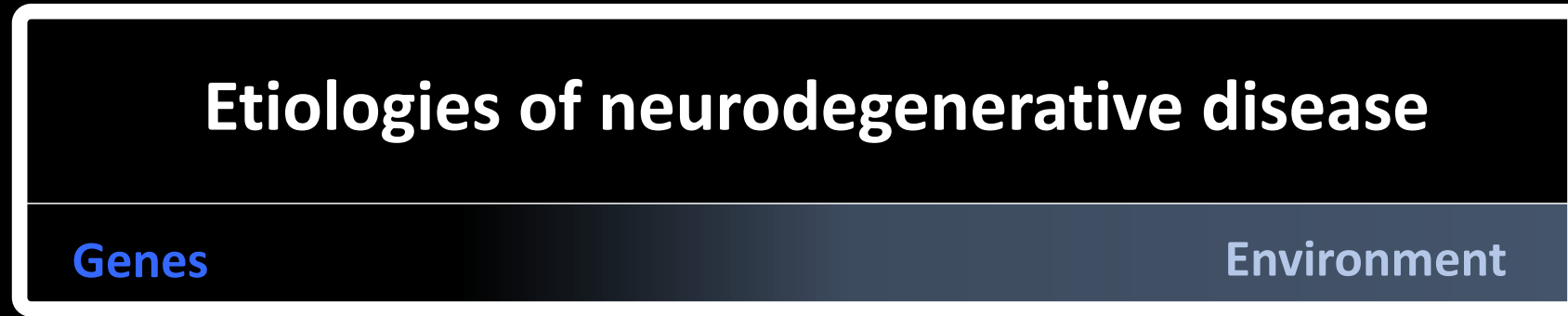


Cell type specificity



Axon injury responses in human neurons

Causes of neurodegeneration are still poorly understood



Amyotrophic lateral sclerosis

Placeholder for background on ALS

Mutations in SPTLC1 cause juvenile ALS

(serine palmitoyltransferase 1)

nature
medicine

ARTICLES

<https://doi.org/10.1038/s41591-021-01346-1>



Childhood amyotrophic lateral sclerosis caused by excess sphingolipid synthesis

Payam Mohassel¹, Sandra Donkervoort^{1,26}, Museer A. Lone^{2,26}, Matthew Nalls^{1,26}, Kenneth Gable^{3,26}, Sita D. Gupta^{3,26}, A. Reghan Foley¹, Ying Hu¹, Jonas Alex Morales Saute⁴, Ana Lucila Moreira⁵, Fernando Kok⁶, Alessandro Intronà⁷, Giancarlo Logroscino^{7,8}, Christopher Grunseich⁹, Alec R. Nickolls¹, Naemeh Pourshafie⁹, Sarah B. Neuhaus¹, Dimah Saade¹, Andrea Gangfuß¹⁰, Heike Kölbl¹⁰, Zoe Piccus¹¹, Claire E. Le Pichon¹¹, Chiara Fiorillo¹², Cindy V. Ly¹³, Ana Töpf¹⁴, Lauren Brady¹⁵, Sabine Specht¹⁴, Aliza Zidell¹⁶, Helio Pedro¹⁷, Eric Mittelman¹⁸, Florian P. Thomas¹⁸, Katherine R. Chao¹⁹, Chamindra G. Konersman²⁰, Megan T. Cho²¹, Tracy Brandt²¹, Volker Straub¹⁴, Anne M. Connolly²², Ulrike Schara¹⁰, Andreas Roos¹⁰, Mark Tarnopolsky¹⁵, Ahmet Höke²³, Robert H. Brown²⁴, Chia-Hsueh Lee²⁵, Thorsten Hornemann², Teresa M. Dunn^{3,23} and Carsten G. Bönnemann¹✉

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Research

JAMA Neurology | **Original Investigation**

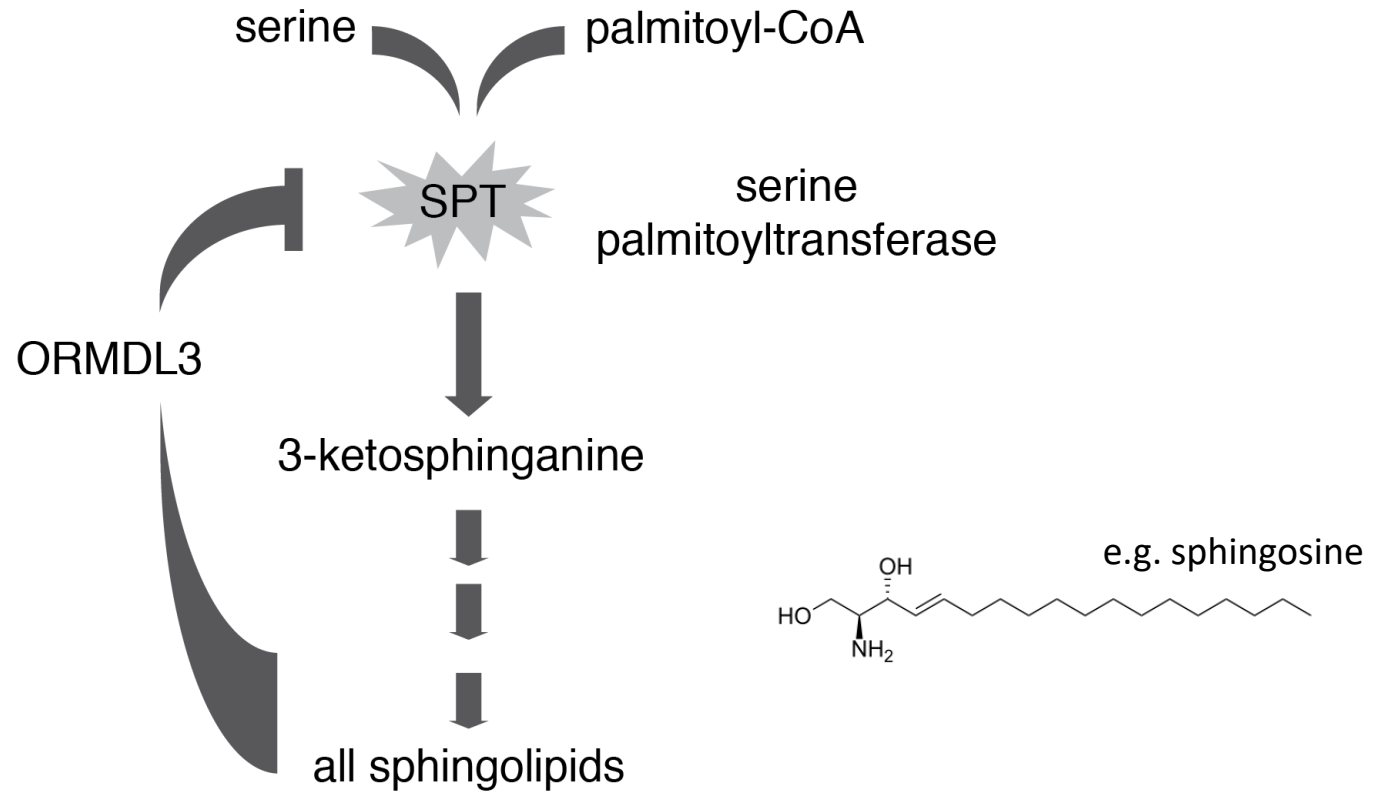
Association of Variants in the *SPTLC1* Gene With Juvenile Amyotrophic Lateral Sclerosis

Janel O. Johnson, PhD; Ruth Chia, PhD; Danny E. Miller, MD, PhD; Rachel Li, MD; Ravindran Kumaran, PhD; Yevgeniya Abramzon, BSc; Nada Alahmady, PhD; Alan E. Renton, PhD; Simon D. Topp, PhD; J. Raphael Gibbs, PhD; Mark R. Cookson, PhD; Marya S. Sabir, BSc; Clifton L. Dalgard, PhD; Claire Troakes, PhD; Ashley R. Jones, PhD; Aleksey Shatunov, PhD; Alfredo Iacoangeli, PhD; Ahmad Al Khleifat, PhD; Nicola Ticozzi, MD, PhD; Vincenzo Silani, MD; Cinzia Gellera, PhD; Ian P. Blair, PhD; Carol Dobson-Stone, PhD; John B. Kwok, PhD; Emily S. Bonkowski, ScM; Robin Palvadeau, MSc; Pentti J. Tienari, MD; Karen E. Morrison, MD; Pamela J. Shaw, MD; Ammar Al-Chalabi, PhD; Robert H. Brown Jr, MD, PhD; Andrea Calvo, PhD; Gabriele Mora, PhD; Hind Al-Saif, MD; Marc Gotkine, MBBS; Fawn Leigh, MD; Irene J. Chang, MD; Seth J. Perlman, MD; Ian Glass, MB ChB, MD; Anna I. Scott, PhD; Christopher E. Shaw, MD; A. Nazli Basak, PhD; John E. Landers, PhD; Adriano Chiò, PhD; Thomas O. Crawford, PhD; Bradley N. Smith, PhD; Bryan J. Traynor, MD, PhD; and the FALS Sequencing Consortium; American Genome Center; International ALS Genomics Consortium; and ITALSGEN Consortium

JAMA Neurol. 2021;78(10):1236-1248. doi:10.1001/jamaneurol.2021.2598

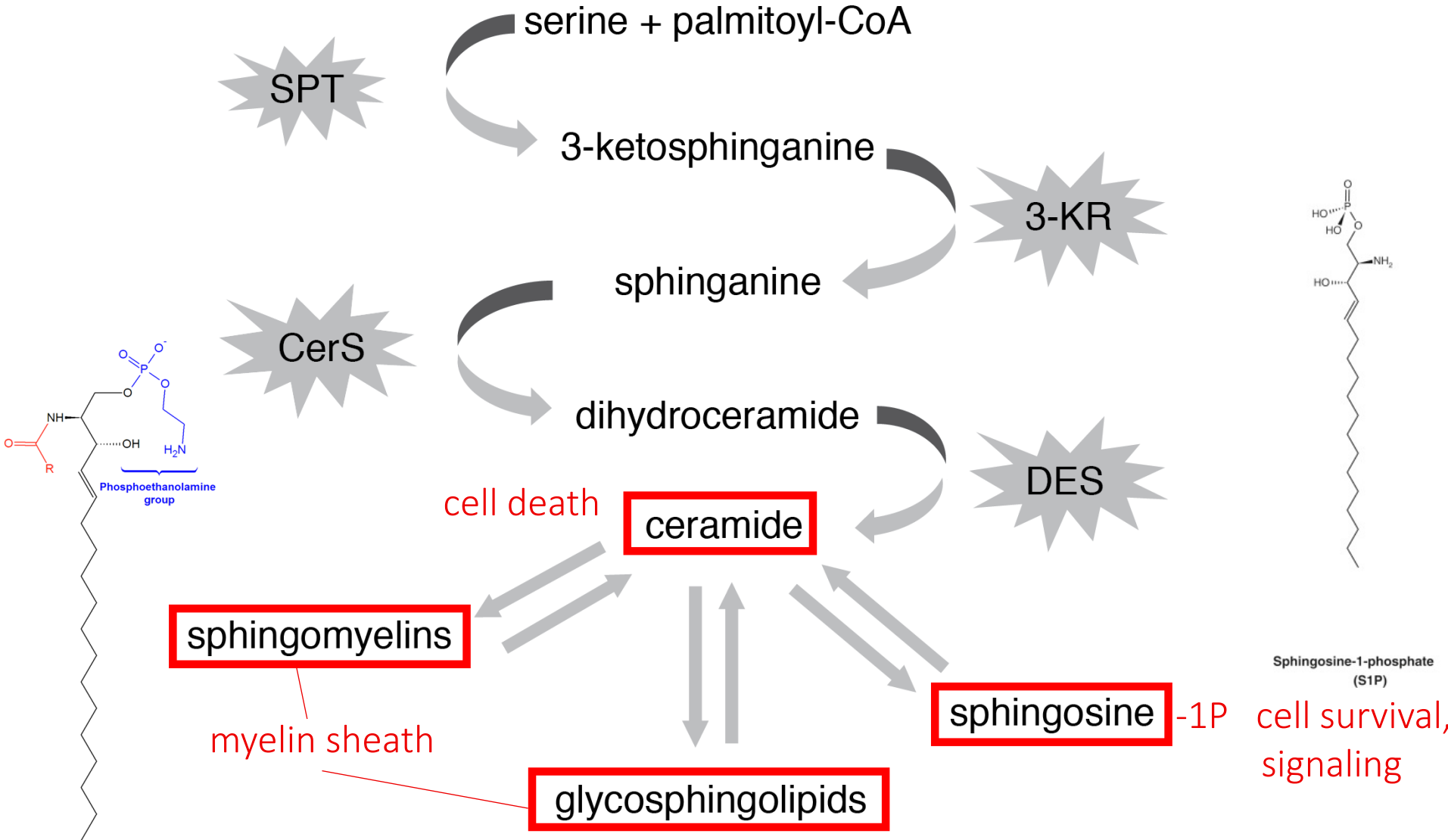
Published online August 30, 2021.

All sphingolipid biosynthesis is initiated by SPT



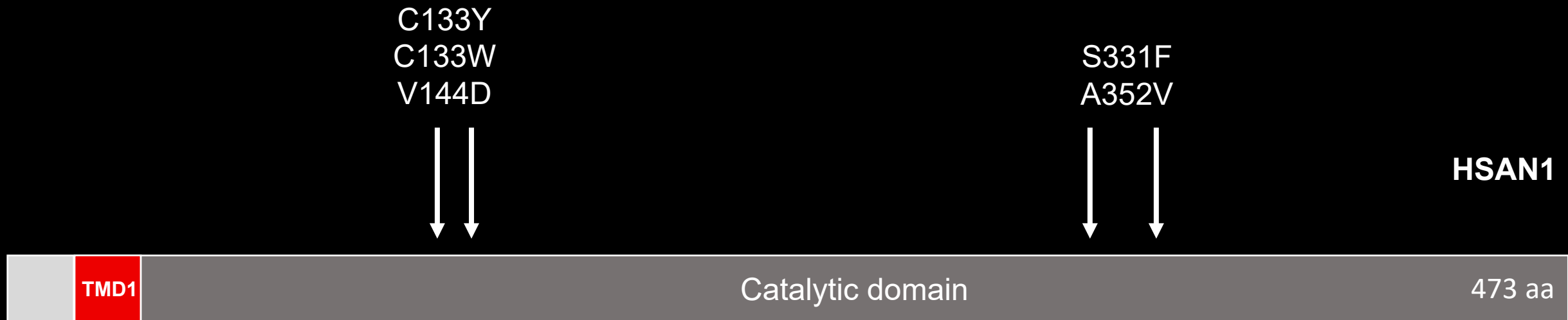
Essential lipids of membranes in mammalian cells
Wide spectrum of functions

Sphingolipid biosynthesis pathway

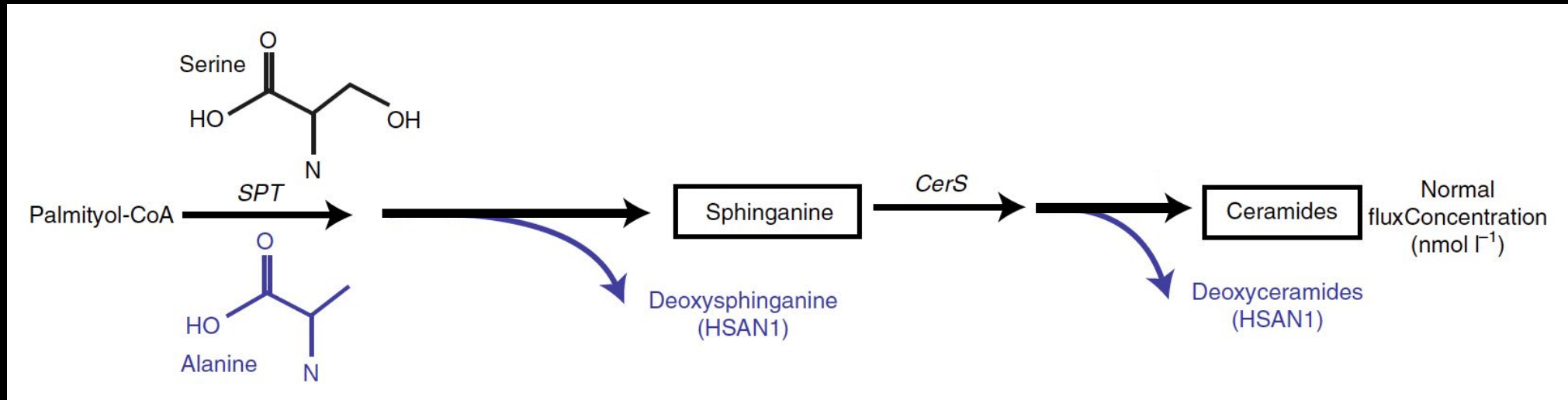


Location of disease-causing mutations in SPTLC1

- Previously known SPTLC1 mutations cause HSAN1
(Hereditary Sensory and Autonomic Neuropathy type 1)



Metabolomic alterations resulting from HSAN1 mutations in SPTLC1



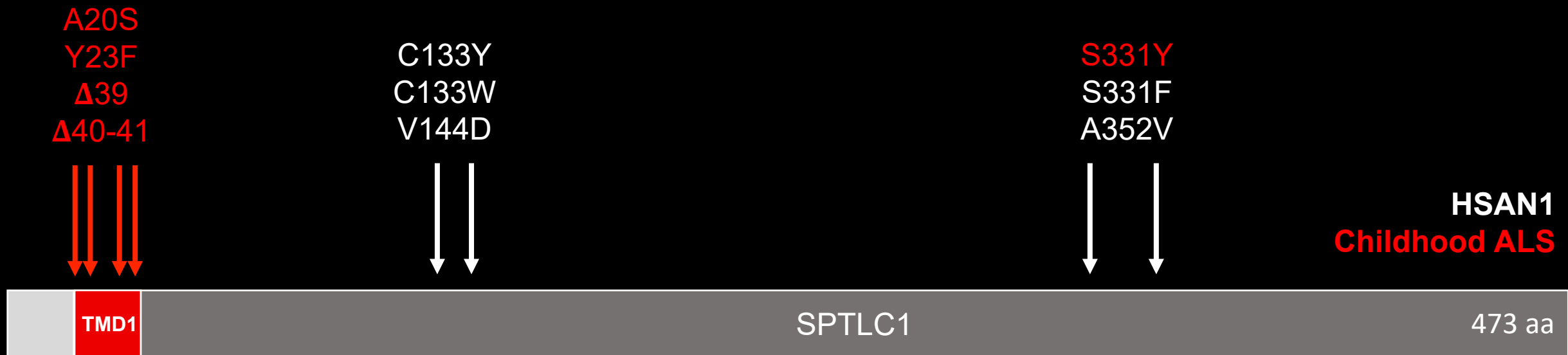
Location of disease-causing mutations in SPTLC1

- Previously known SPTLC1 mutations cause HSAN1

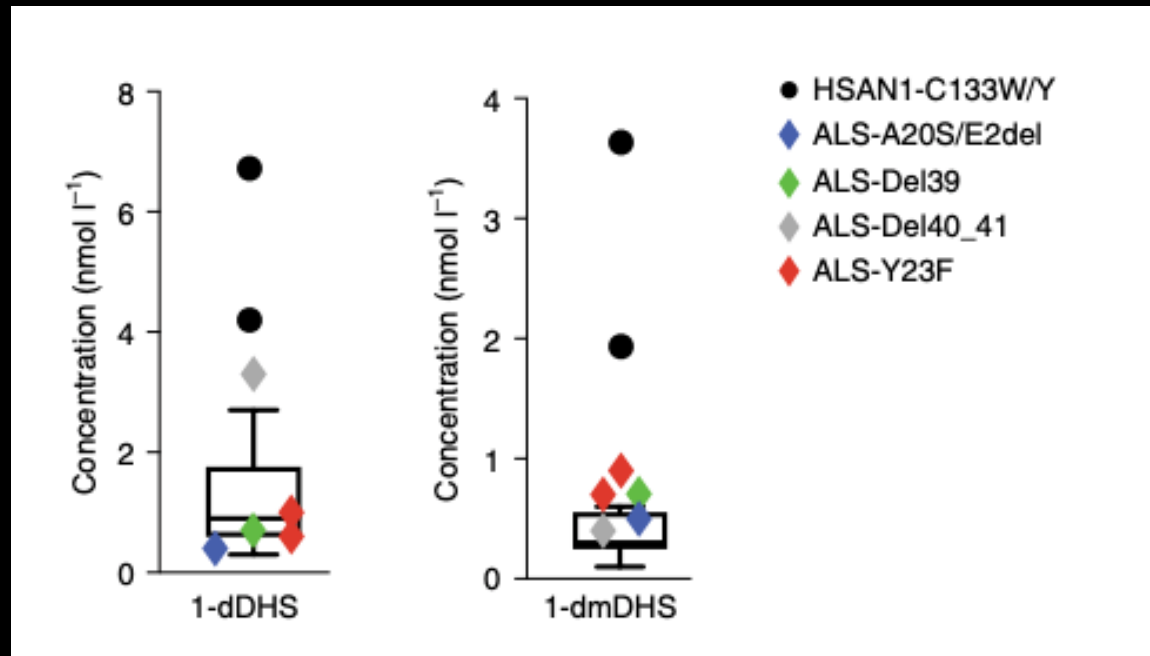


Location of disease-causing mutations in SPTLC1

- Previously known SPTLC1 mutations cause HSAN1
- The childhood ALS-linked mutations cluster in a different region: transmembrane domain 1 (TMD1)

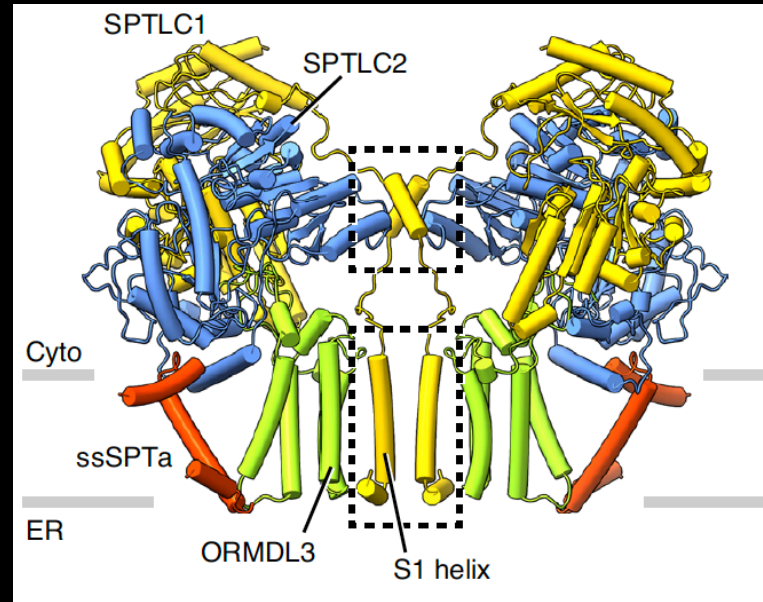


Metabolomic alterations resulting from ALS vs HSN1 mutations in SPTLC1

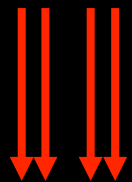


Sphingolipidomic analysis of patient serum

Location of disease-causing mutations in SPTLC1



A20S
Y23F
Δ39
Δ40-41



C133Y
C133W
V144D



S331Y
S331F
A352V



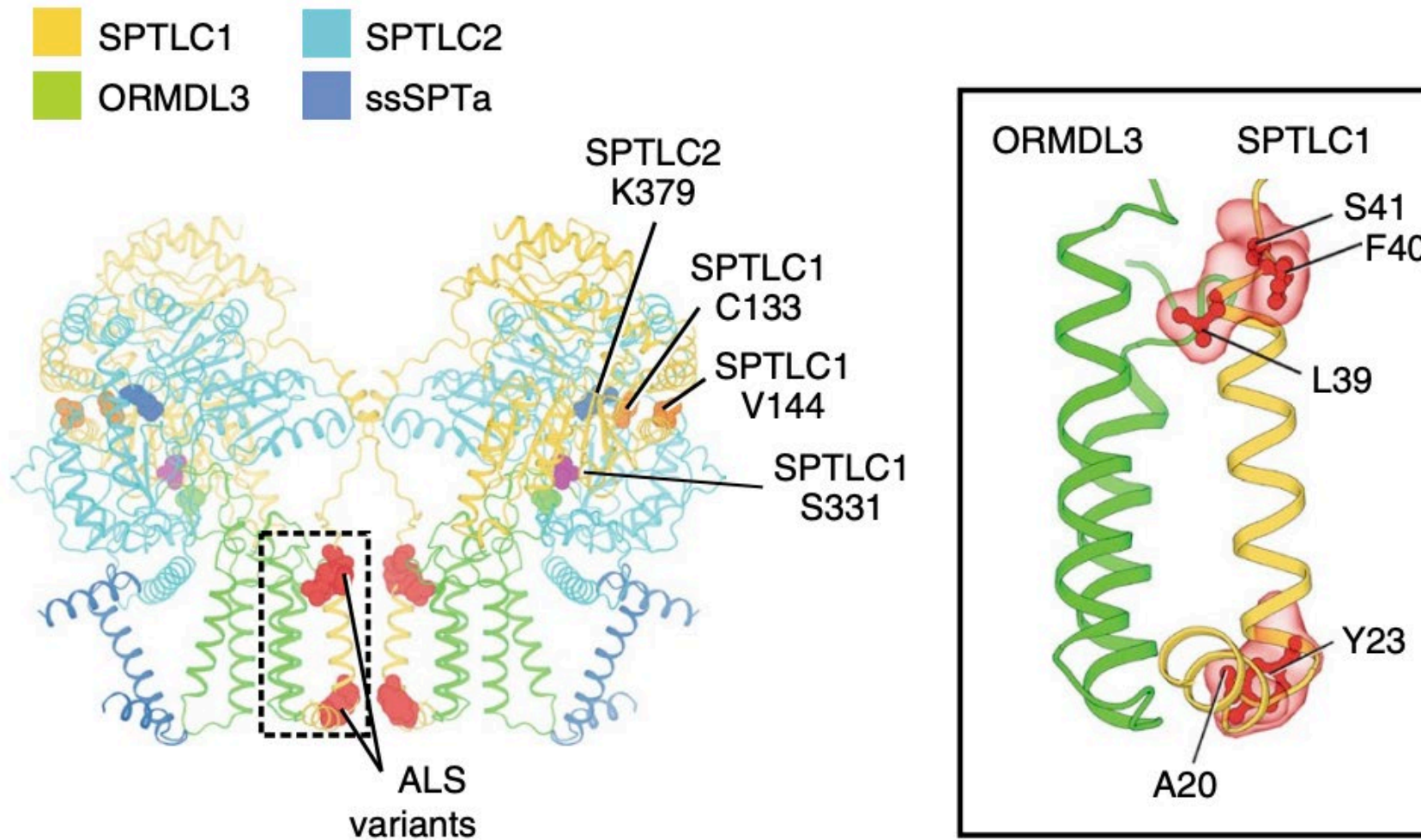
HSAN1
Childhood ALS

TMD1

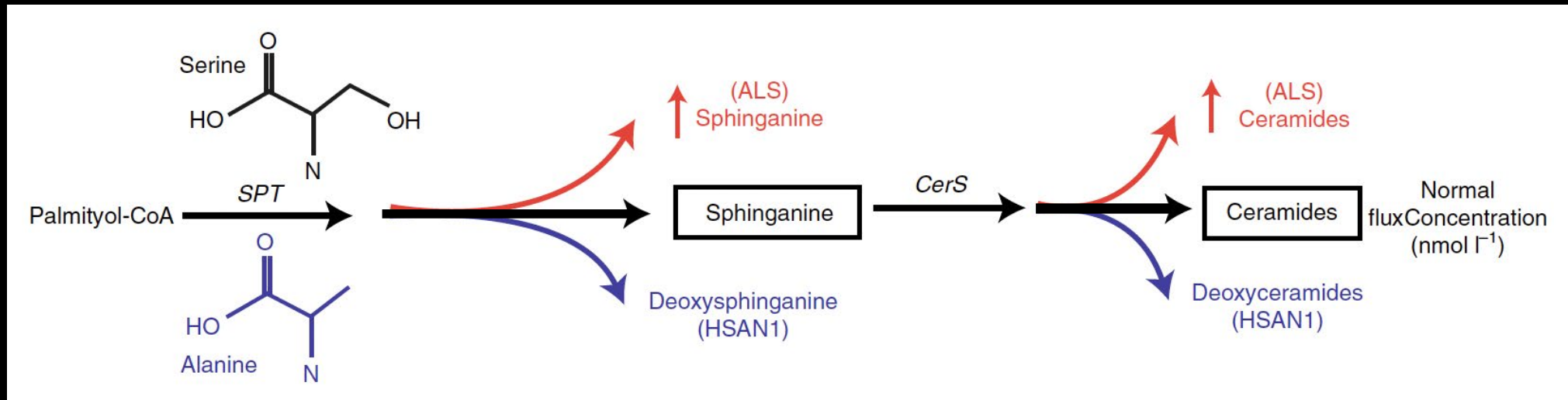
Catalytic domain

473 aa

Location of ALS mutations within SPTLC1 structure



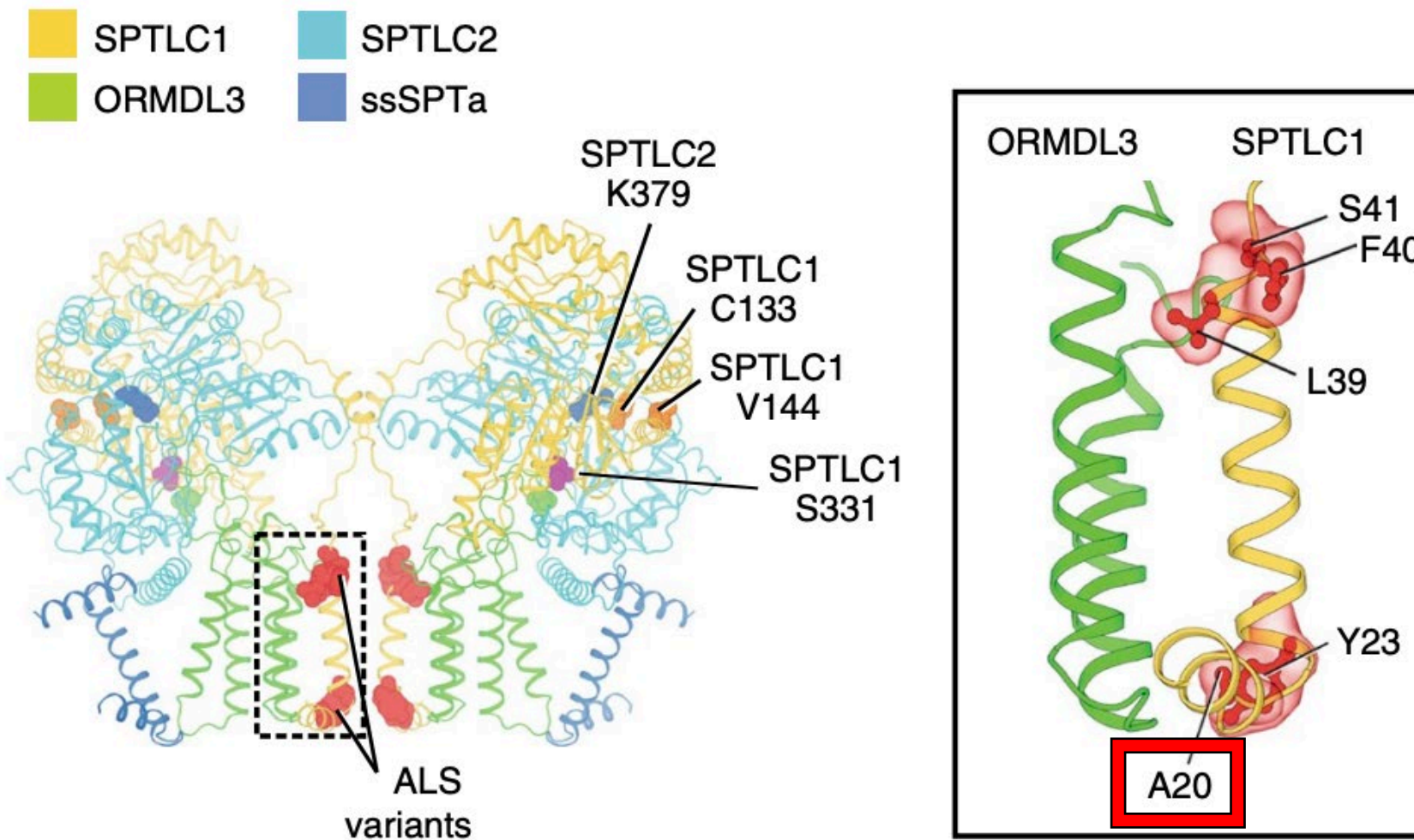
Metabolomic alterations resulting from ALS vs HSN1 mutations in SPTLC1



Hypotheses - mice engineered with these ALS mutations will exhibit:

1. elevated sphingolipid levels
2. ALS-like neurodegeneration

SPT-ORMDL3 interaction occurs at TMD1

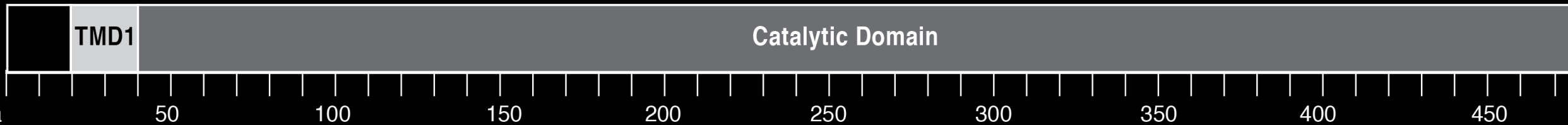
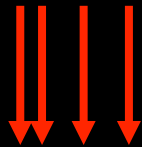


High conservation of TMD1 (encoded by exon 2)

Gene editing approach in mouse to knock in the mutation

AA number	20	23	38	39	40	41		
<i>Homo sapiens</i>	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F	S	K T Y K
<i>Pan troglodytes</i>	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F	S	K T Y K
<i>Macaca mulatta</i>	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F	S	K T Y K
<i>Felis catus</i>	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F	S	K T Y K
<i>Mus musculus</i>	L Y E A	P A Y	H L I L E G I L I L W I I R	L	V	F	S	K T Y K
<i>Gallus gallus</i>	F Y E A	P A Y	H L I L E G I L I L W I I R	L	I	F	S	K T Y K
<i>Takifugu rubripes</i>	F Y E A	P A Y	H L I L E G I L I L W I F R	L	L	F	S	K T Y K
<i>Danio rerio</i>	F Y E A	P A Y	H L I L E G F L I L W I I R	L	L	F	S	K T Y K
<i>Xenopus tropicalis</i>	F Y E A	P A Y	H L I L E G I L I L W I I R	L	I	F	S	K T Y K

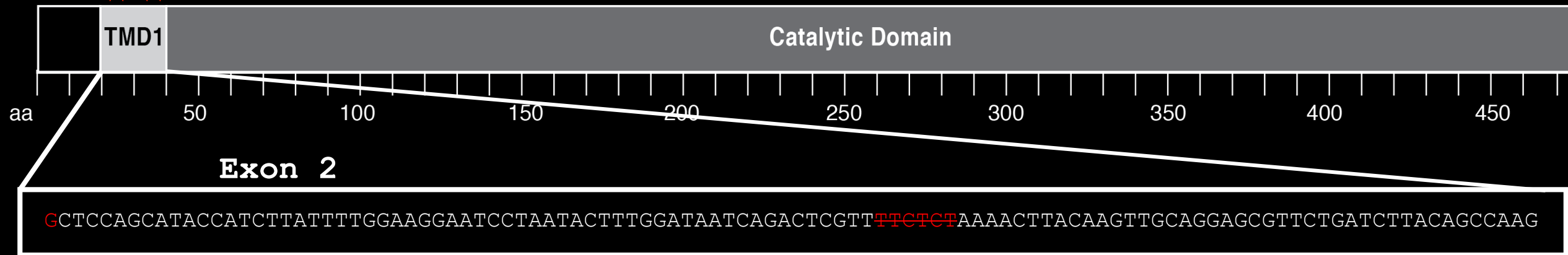
A20S
Y23F
Δ39
Δ40-41



High conservation of TMD1 (encoded by exon 2) Gene editing approach

AA number	20	23	38	39	40	41	
Homo sapiens	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F S	K T Y K
Pan troglodytes	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F S	K T Y K
Macaca mulatta	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F S	K T Y K
Felis catus	L Y E A	P A Y	H L I L E G I L I L W I I R	L	L	F S	K T Y K
Mus musculus	L Y E A	P A Y	H L I L E G I L I L W I I R	L	V	F S	K T Y K
Gallus gallus	F Y E A	P A Y	H L I L E G I L I L W I I R	L	I	F S	K T Y K
Takifugu rubripes	F Y E A	P A Y	H L I L E G I L I L W I F R	L	L	F S	K T Y K
Danio rerio	F Y E A	P A Y	H L I L E G F L I L W I I R	L	L	F S	K T Y K
Xenopus tropicalis	F Y E A	P A Y	H L I L E G I L I L W I I R	L	I	F S	K T Y K

A20S

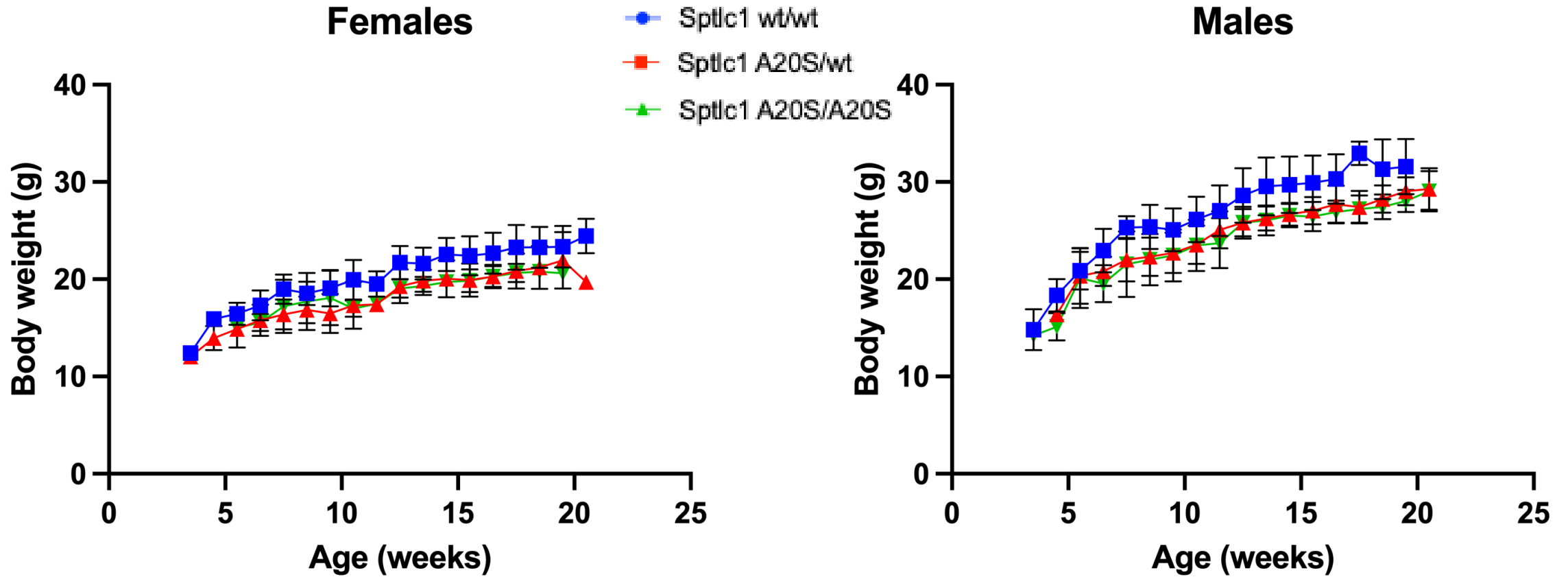


G->T
A20S

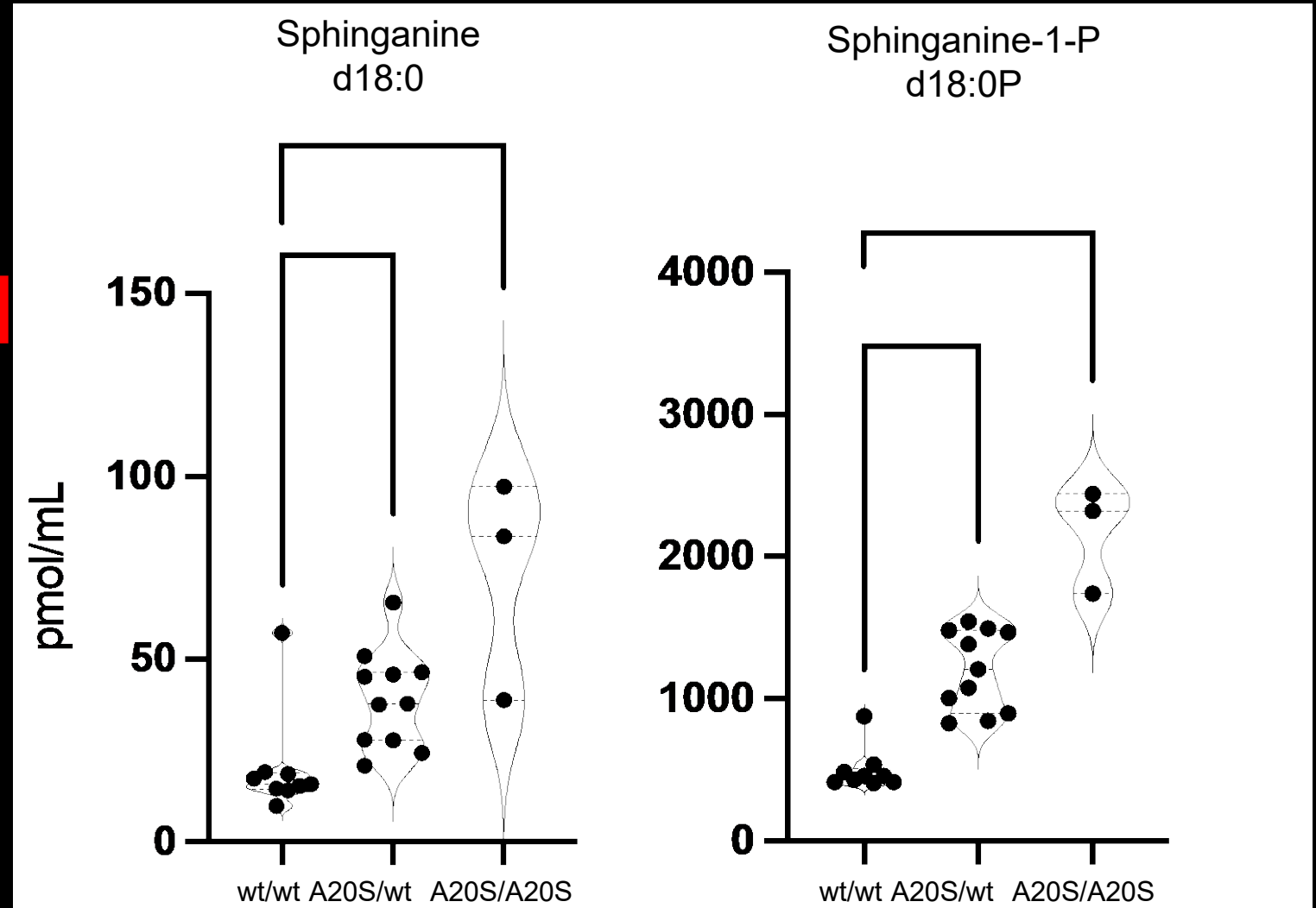
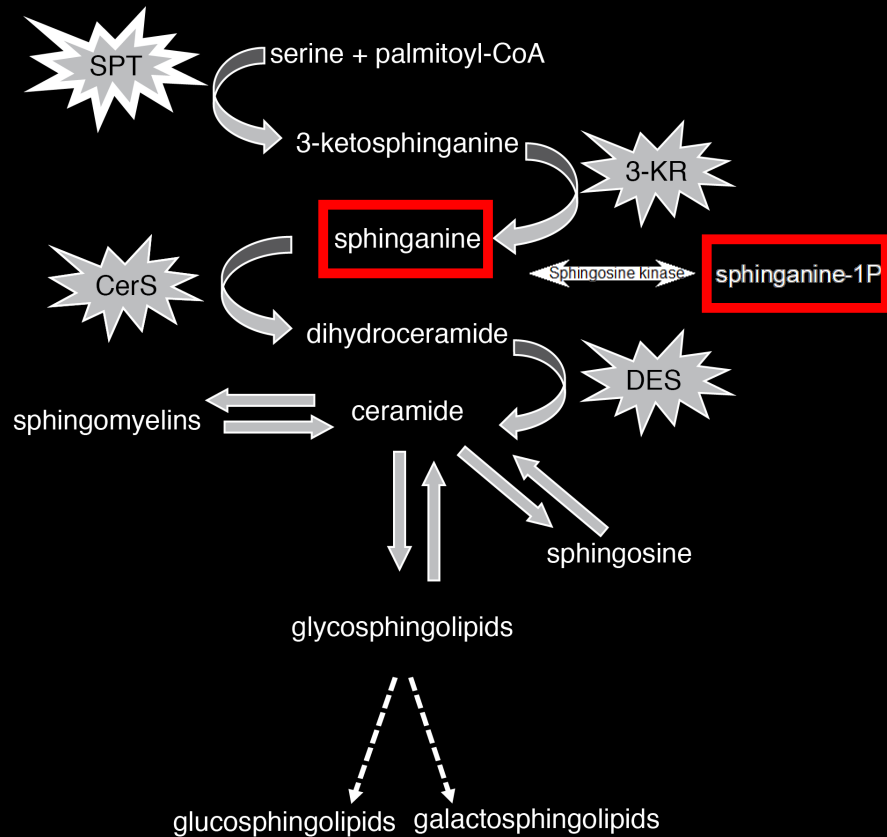
Characterization of Sptlc1^{A20S} knock-in mouse line

A20S	del 40-41
Normal lifespan	Homozygotes runted and die early (5-6 weeks old)
Mild motor symptoms	Tremors, muscle atrophy
Progressive motor neuron, nerve and muscle pathology (but no obvious motor neuron cell body loss)	Misdevelopment and neurodegeneration (more similar to a juvenile syndrome)
Homozygotes exhibit more severe pathology	Histological characterization ongoing

A20S carriers have slightly lower body weight

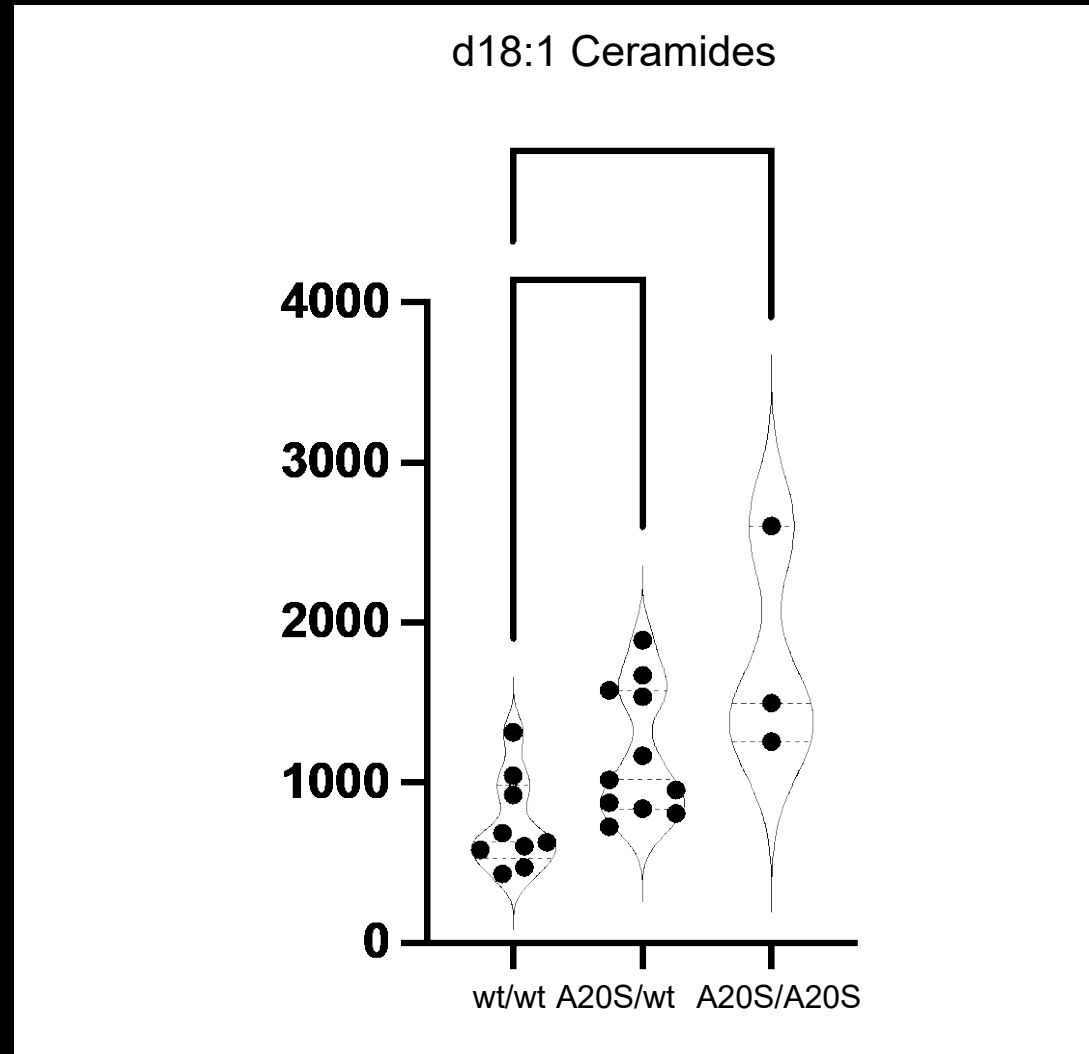
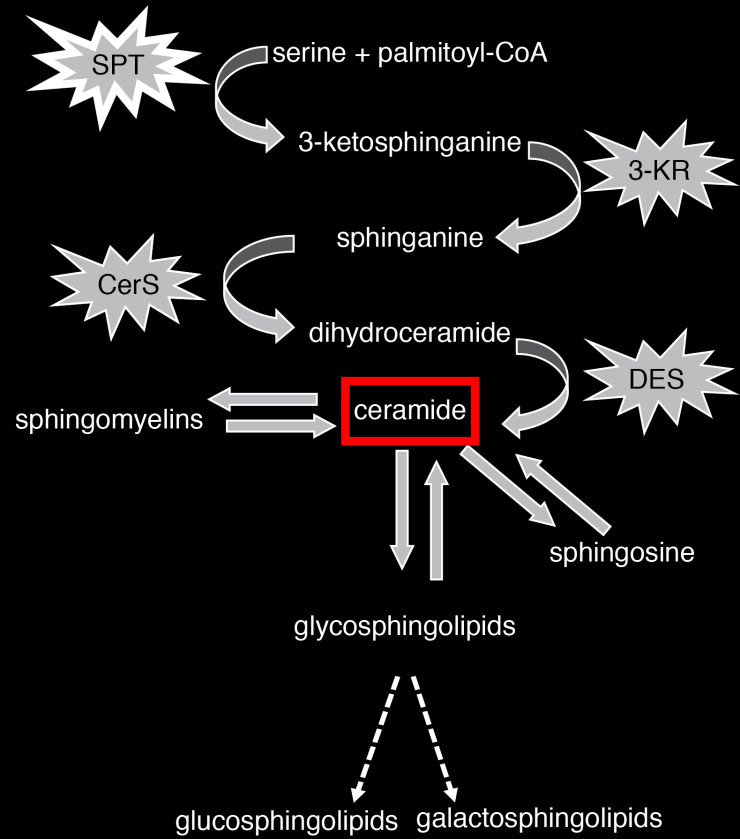


Sptlc1^{A20S} animals produce excess sphingolipids



10 to 20-week-old serum

Sptlc1^{A20S} animals produce excess sphingolipids

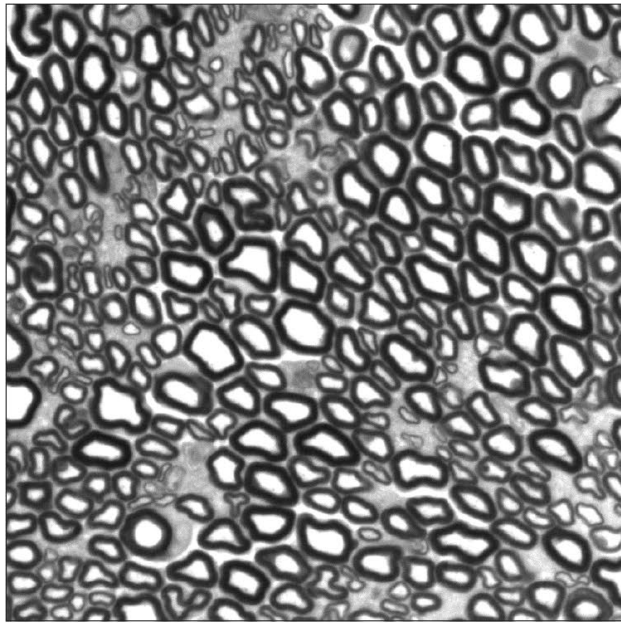


10 to 20-week-old serum

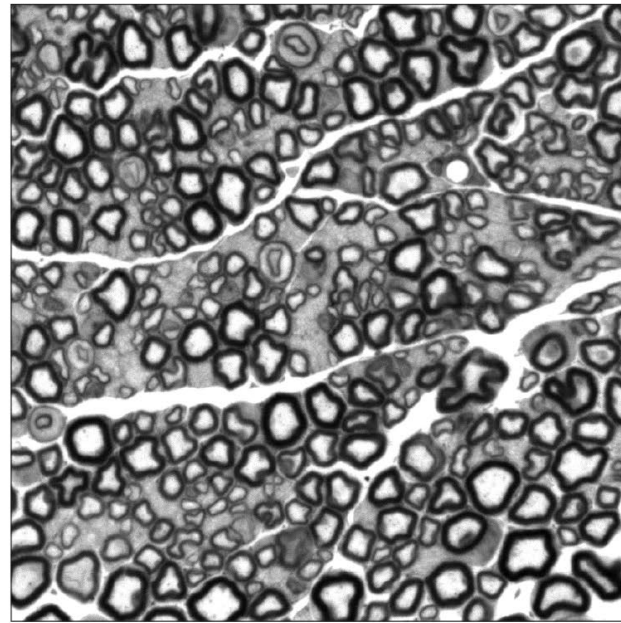
Sciatic nerve pathology in Sptlc1 A20S mutants

Sptlc1

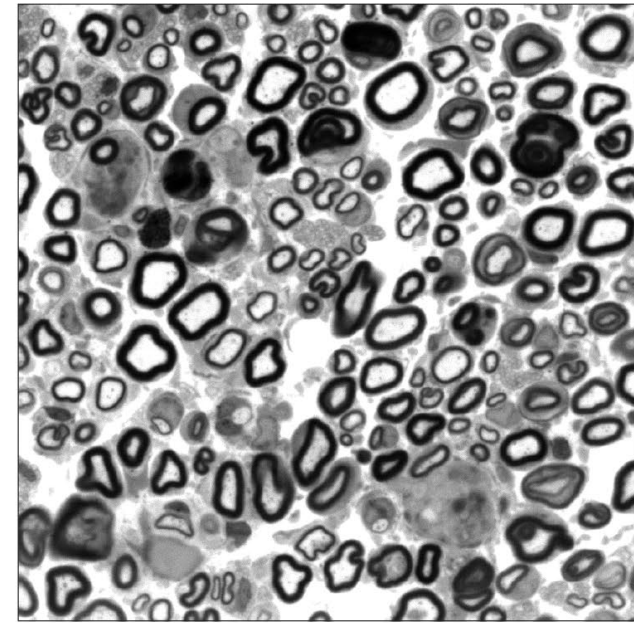
wt/wt



A20S/wt



A20S/A20S



9-week old, myelin stain

25 μ m

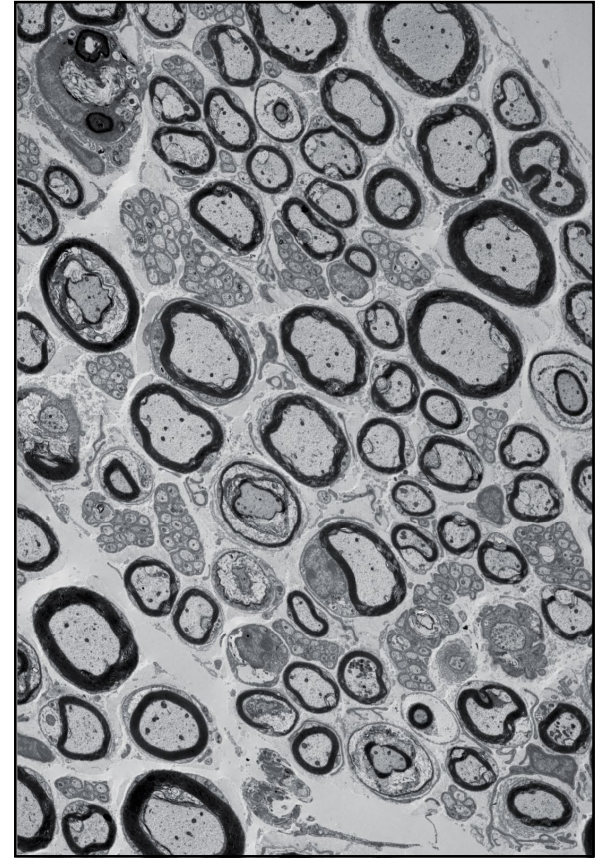
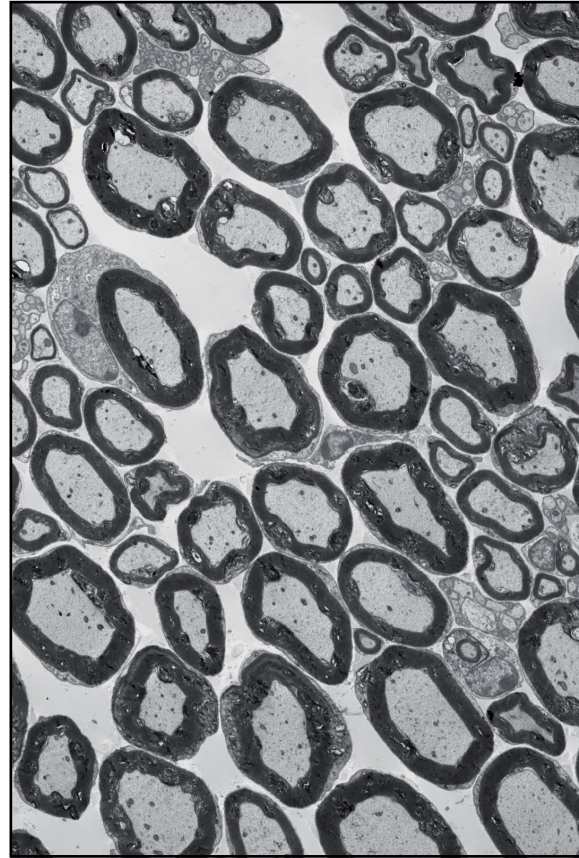
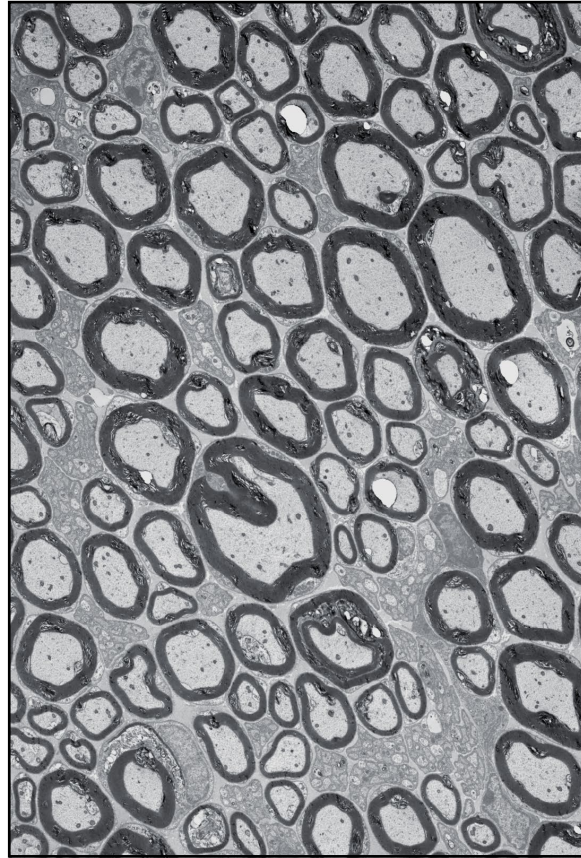
Sciatic nerve pathology in Sptlc1 A20S mutants

Sptlc1

wt/wt

A20S/wt

A20S/A20S



19-week old
Electron microscopy (EM)

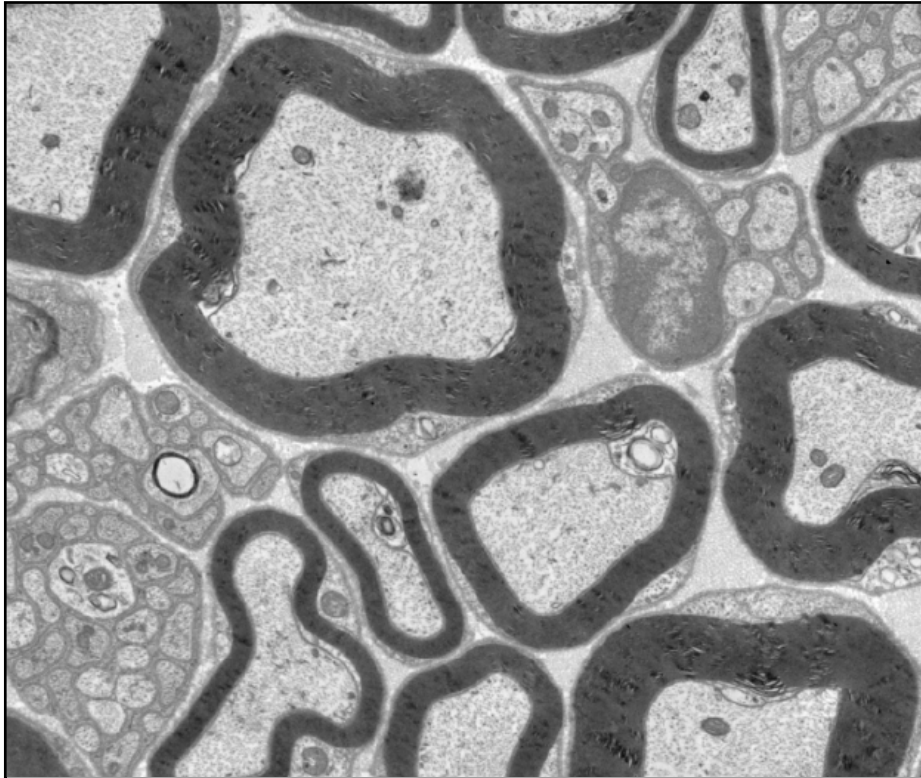
6 μ m

Degenerative pathology in $Sptlc1^{A20S}$ mutant nerve

Sptlc1

wt/wt

Healthy Schwann cells and axons



A20S/A20S

Schwann cell pathology /axonal sprouting



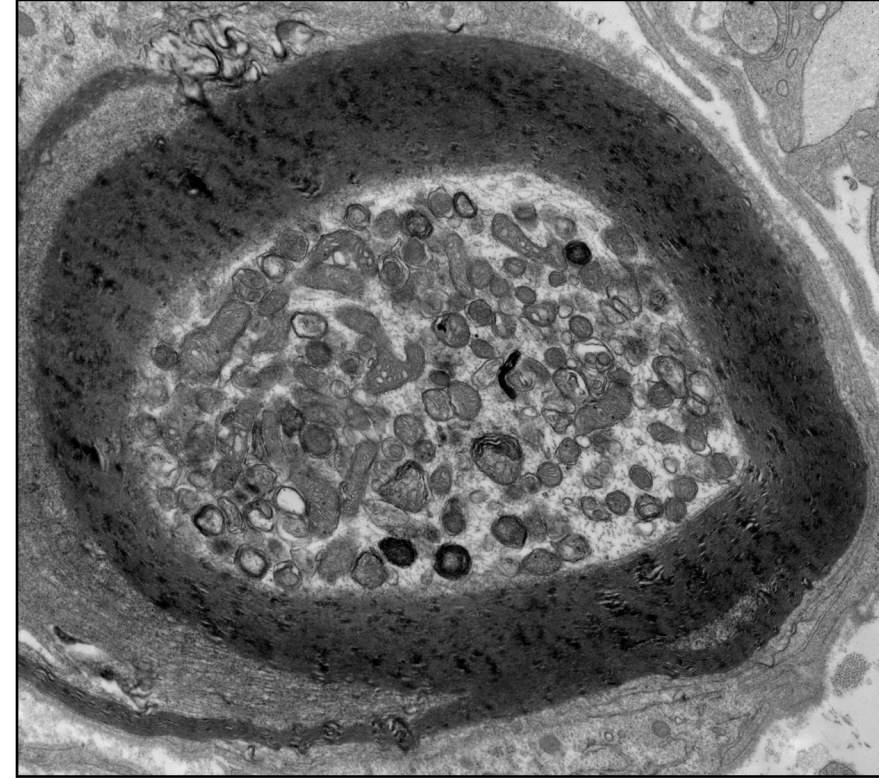
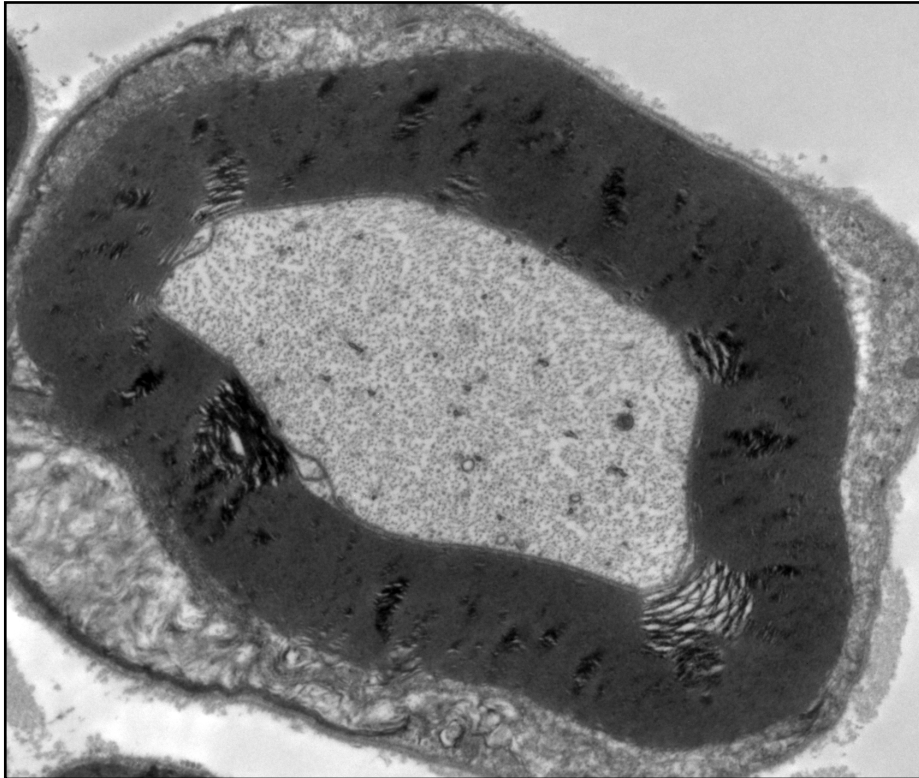
19-week old EM, littermates

Degenerative pathology in *Sptlc1*^{A20S} mutant nerve

Sptlc1

wt/wt
Healthy axon

A20S/A20S
Axonal organelle accumulations



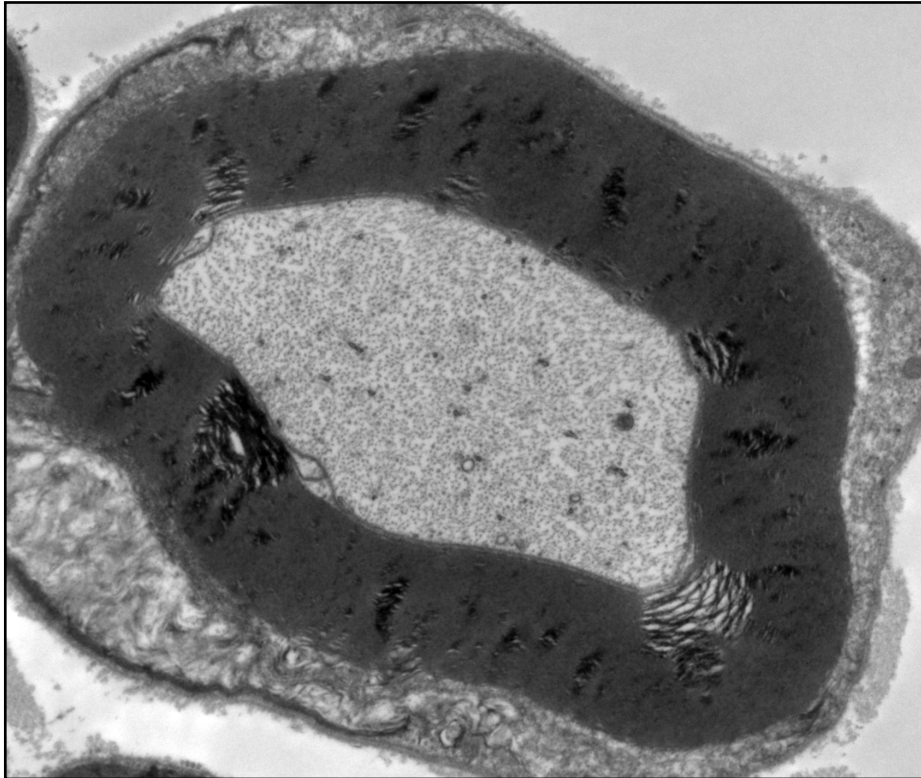
0.5 μ m

19-week old EM, littermates

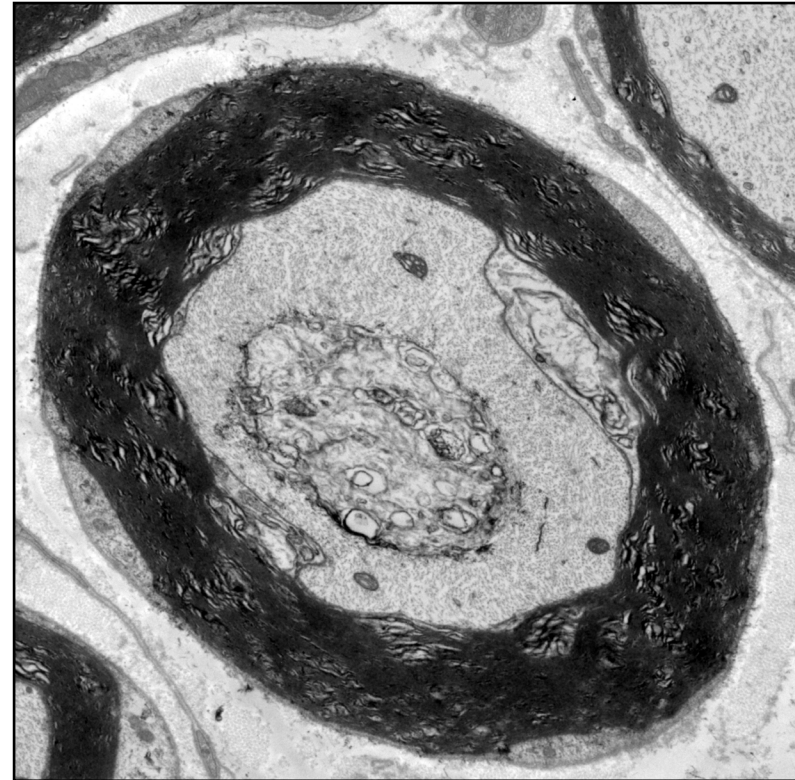
Degenerative pathology in *Sptlc1*^{A20S} mutant nerve

Sptlc1

wt/wt
Healthy axon



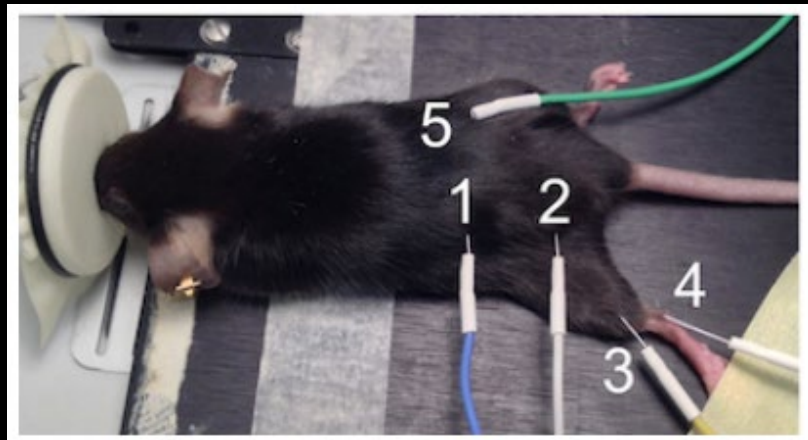
A20S/A20S
Degenerating axon



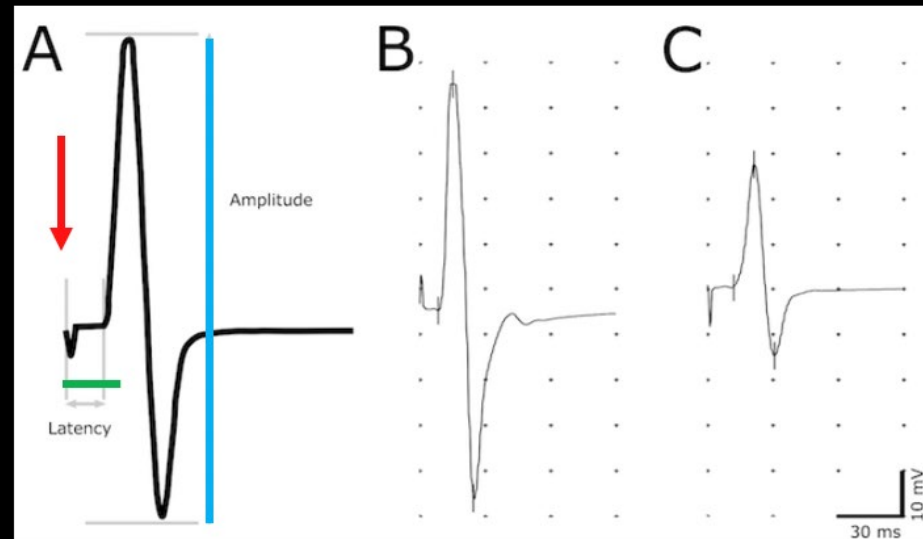
1 μ m

19-week old EM, littermates

Compound Muscle Action Potential (CMAP) recordings reflect functional connectivity between nerve and muscle



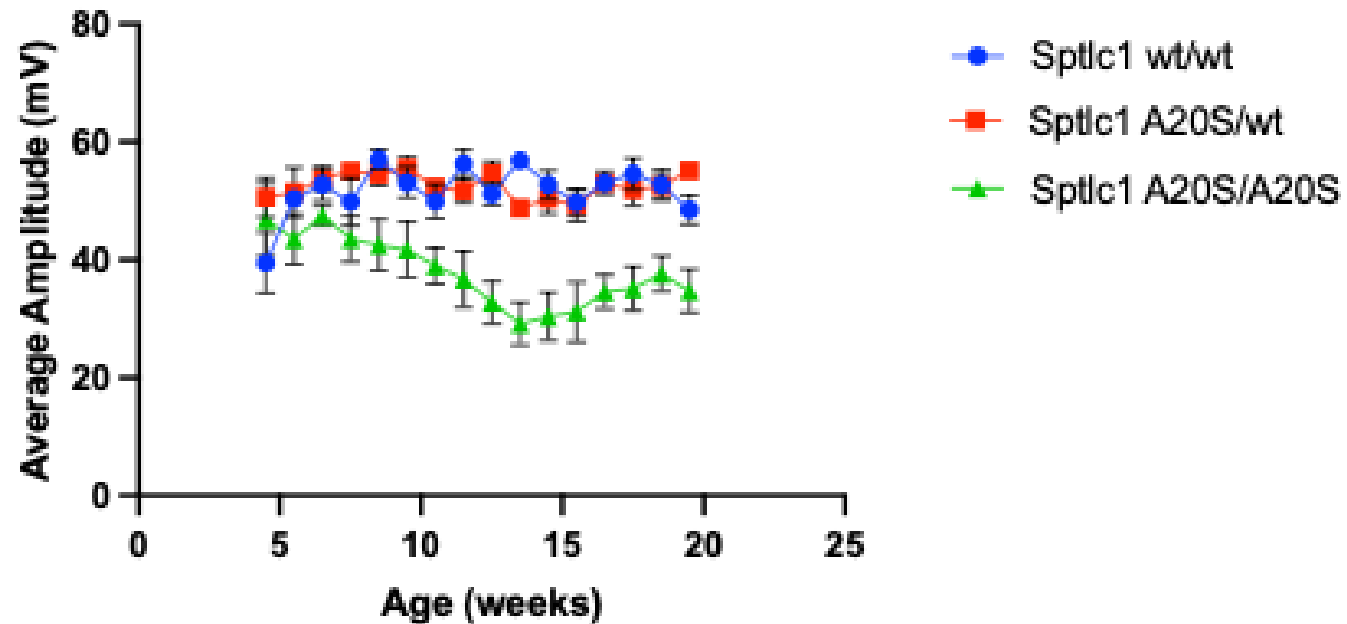
- 1+2: stimulating electrodes
- 3: recording electrode
- 4: reference
- 5: ground



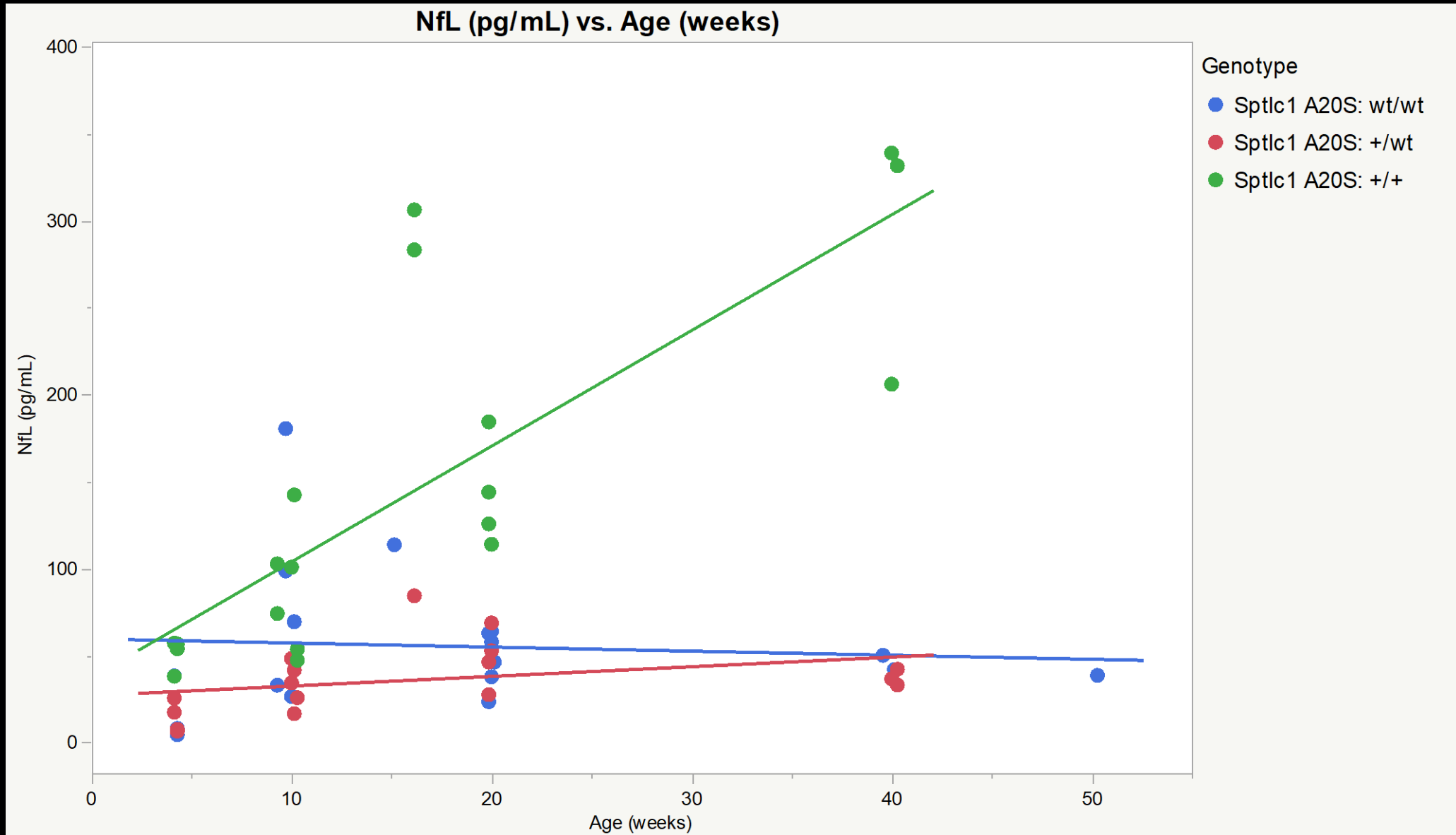
Stimulation ($t=0$)
Latency to peak
Max amplitude

A+B: waveform in WT animals
C: diminished amplitude in SOD1 animal

Nerve to muscle communication is disrupted in homozygotes



A20S homozygotes have increased NfL in serum



The Sptlc1 A20S mouse model can be used for preclinical testing

Placeholder for prevention study
Placeholder for treatment study

Closing thoughts

- Mutations in SPTLC1 associated with ALS produce neurodevelopmental and/or neurodegenerative phenotypes in gene-edited mice
 - Ongoing characterization (motor neuron counts, electrophysiology, muscle histology)
 - Planning prevention study and treatment study using inhibitors of SPTLC1 in *Sptlc1*^{A20S/A20S} mice
- Mutations in SPTLC1 have also been detected in sporadic ALS patients (Johnson et al., 2021)
- Possibility of a pathogenic role for disrupted sphingolipid metabolism in sporadic ALS
- Potential of sphingolipids as disease biomarkers for ALS

Le Pichon lab

Current members

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Hanna Silberberg

Mor Alkaslasi

Jorge Gómez-Deza

Matthew Nebiyou

Josette Wlaschin

Sangeetha Hareendran

Eliza Lloyd

Emily Clark



Alumni

Aditya Santoki

Austin Gable

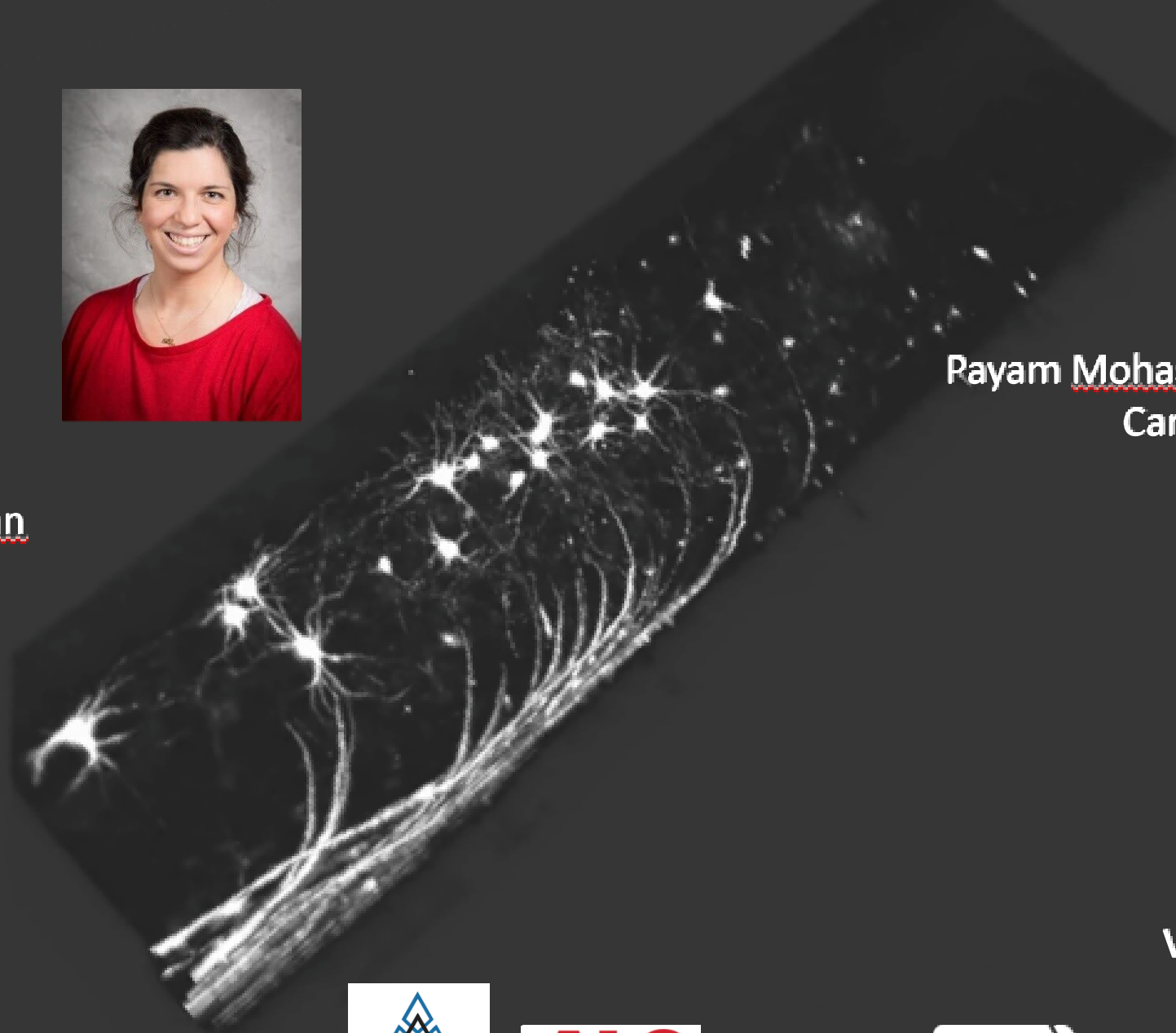
Stacey Slavutsky

Li Chen

Leana Ramos

Caroline Donahue

Jacob Gluski



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Kenneth Gable

Payam Mohassel, NIH / Johns Hopkins
Carsten Bönnemann, NINDS

Bryan Traynor, NIA

Nick Ryba, NIDCR

NIMH Transgenic Core
NHLBI Transgenic Core

NICHD Microscopy Core
Vincent Schram, Chip Dye



Eunice Kennedy Shriver National Institute
of Child Health and Human Development