

# RNA-targeting Cas Systems Correct Molecular and Physiological Features of Myotonic Dystrophy Type I

Ranjan Batra<sup>1,2</sup>, David Nelles<sup>1,2</sup>, Daniela Martino Roth<sup>1</sup>, Maurice Swanson<sup>3</sup>,  
Gene Yeo<sup>2</sup>, Kathie Bishop<sup>1</sup>

(1) Locana, Inc. 3545 John Hopkins Ct. San Diego, CA 92121

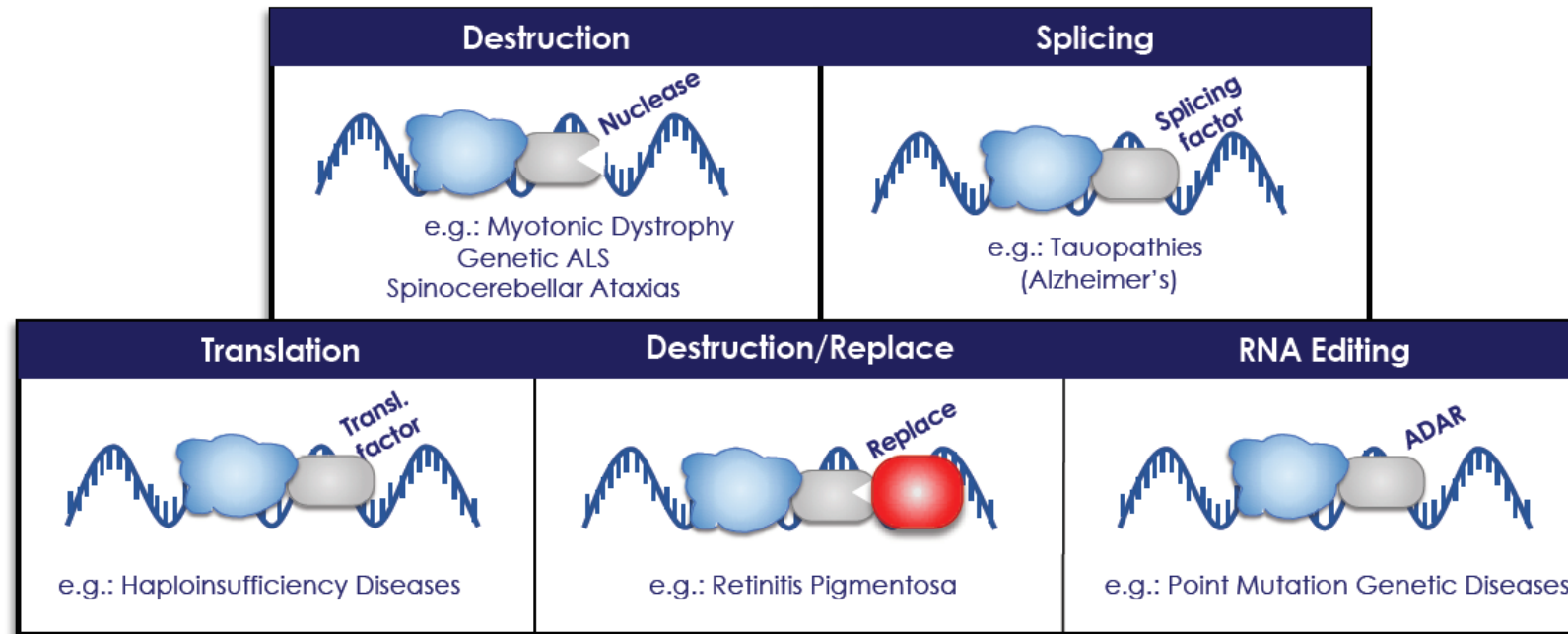
(2) Cellular and Molecular Medicine, University of California, San Diego Neurosciences Graduate Program, San Diego, CA

(3) Center for Neurogenetics and Genetics Institute, University of Florida, College of Medicine, Gainesville, FL 32610, USA



# Introduction

## Locana's RNA Targeting Platform



- Locana has families of RNA binding proteins and a menu of effector domains, which combined present an array of options to address human diseases.
- Using different effector modules, the system can be built to knockdown toxic RNA, upregulate mRNA translation, alter splicing patterns, edit RNA, and knockdown mutant RNA while simultaneously replacing with a normal copy.
- In the current study, RNA Targeting Cas (RCas) systems were evaluated for their knockdown potential for disease related RNAs

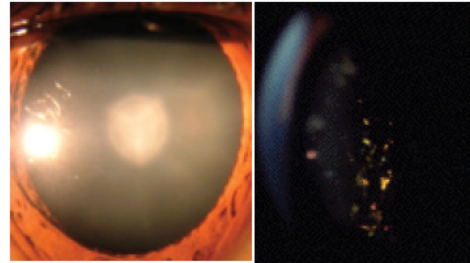
# Myotonic Dystrophy is a Multisystemic Disease



Myotonia  
Muscular Atrophy



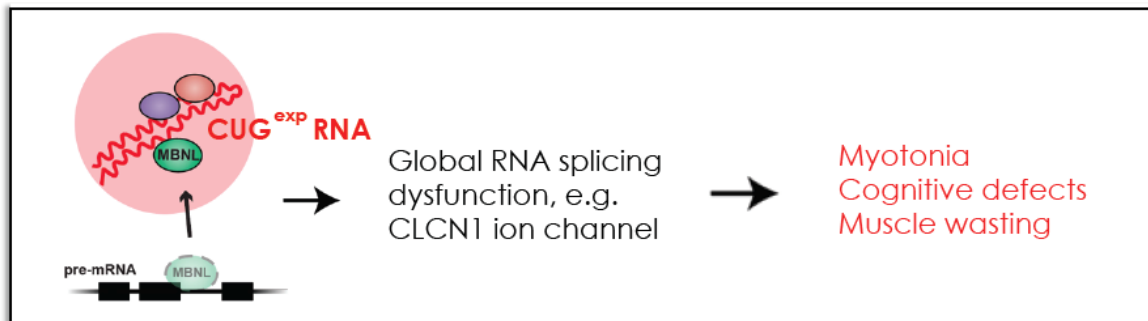
Facial muscle  
weakness  
Developmental  
delays in brain  
function



Cataracts



Congenital Myotonic  
Dystrophy



## Myotonic Dystrophy Etiology

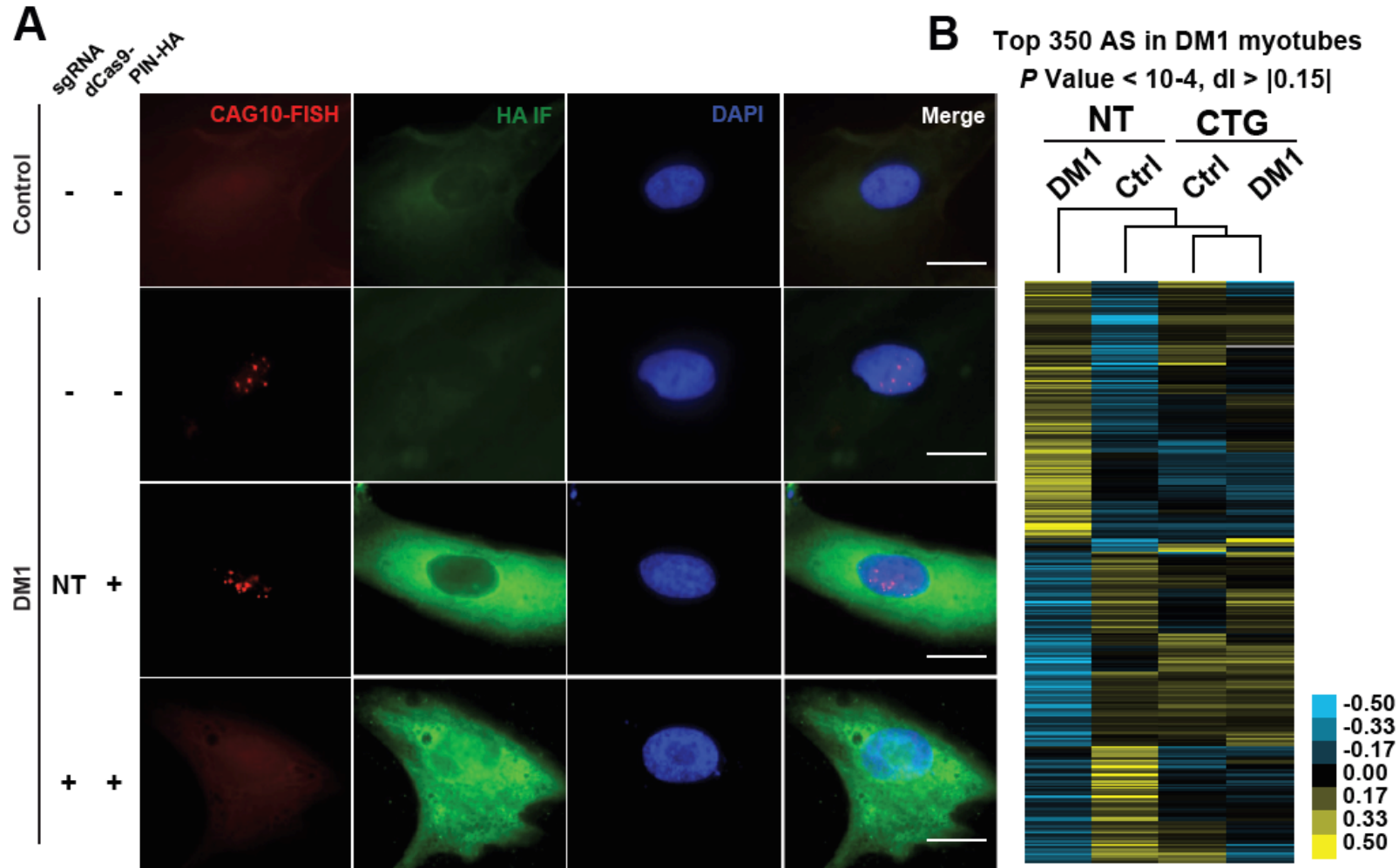
- Myotonic Dystrophy (DM1) is caused by a CTG repeat expansion in the 3' untranslated region of the DMPK gene.

- RNA transcripts containing the CUG repeat expansions sequester muscleblind-like (MBNL) proteins which are the regulators of the alternative splicing switch from fetal to adult isoforms.

- Loss of MBNL proteins in different tissues result in alternative splicing dysregulation which are the major cause of symptoms of DM1.

- Targeting and Eliminating expanded CUG repeats is a therapeutic strategy for DM1.

# RCas9 Corrects DM1 Pathology in Patient Muscle Cells



## Study Design and Results

- Primary myoblasts were isolated from bicep biopsies from DM1 patients and controls.
- RCas9 system targeting CUG repeats was introduced using Lentivirus.

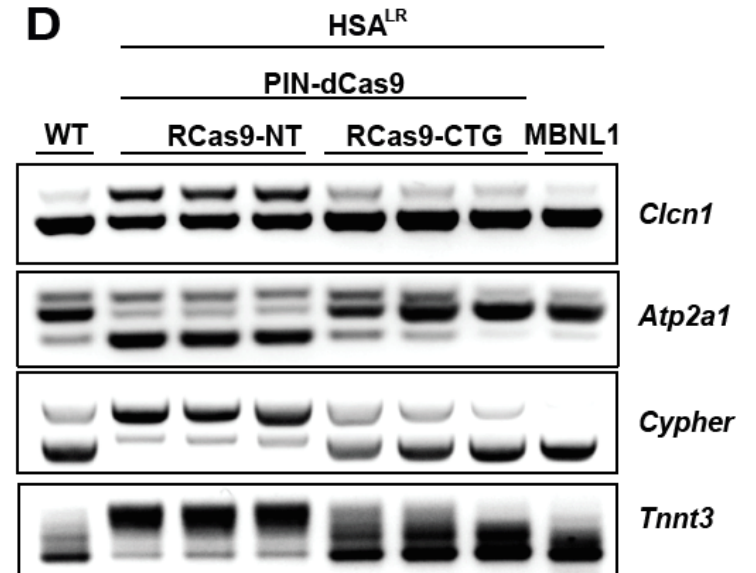
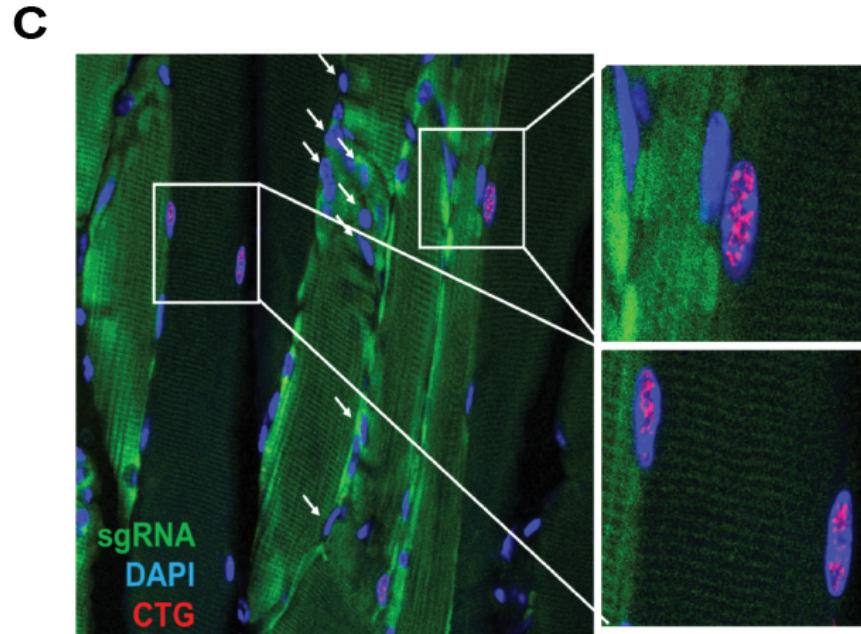
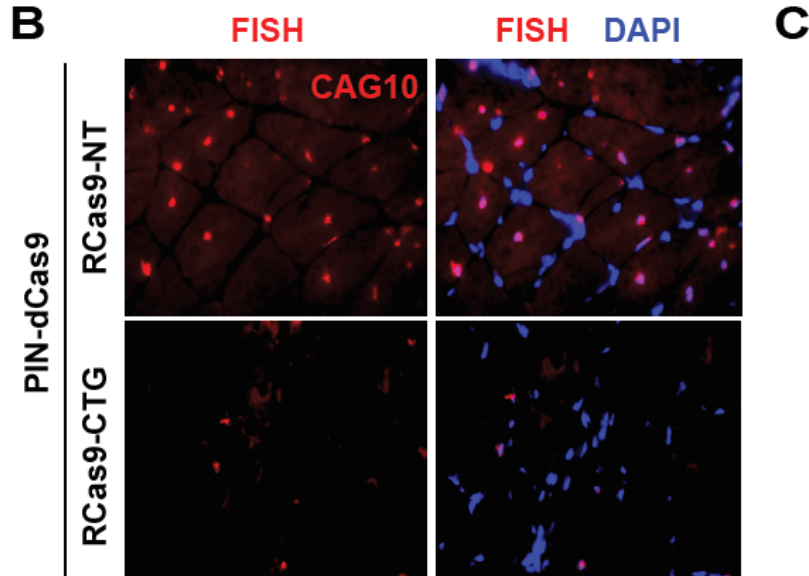
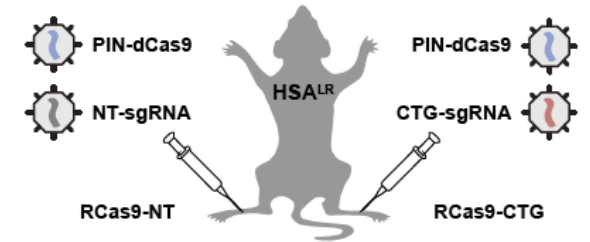
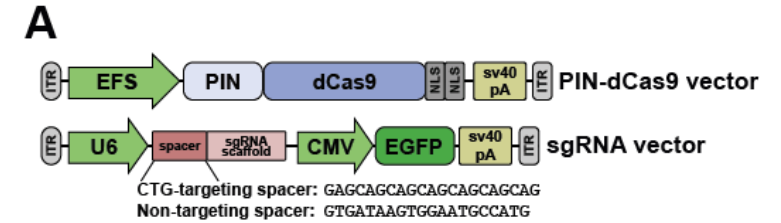
(A) RNA FISH was performed to image RNA-foci typical of DM1 and IF was used to visualize Cas9 tagged with an HA tag. RCas9-CTG eliminated CUG repeats.

(B) Myoblasts differentiated into myotubes were subjected to RNA-seq and DM1-related alternative splicing (AS) changes were hierarchically clustered.



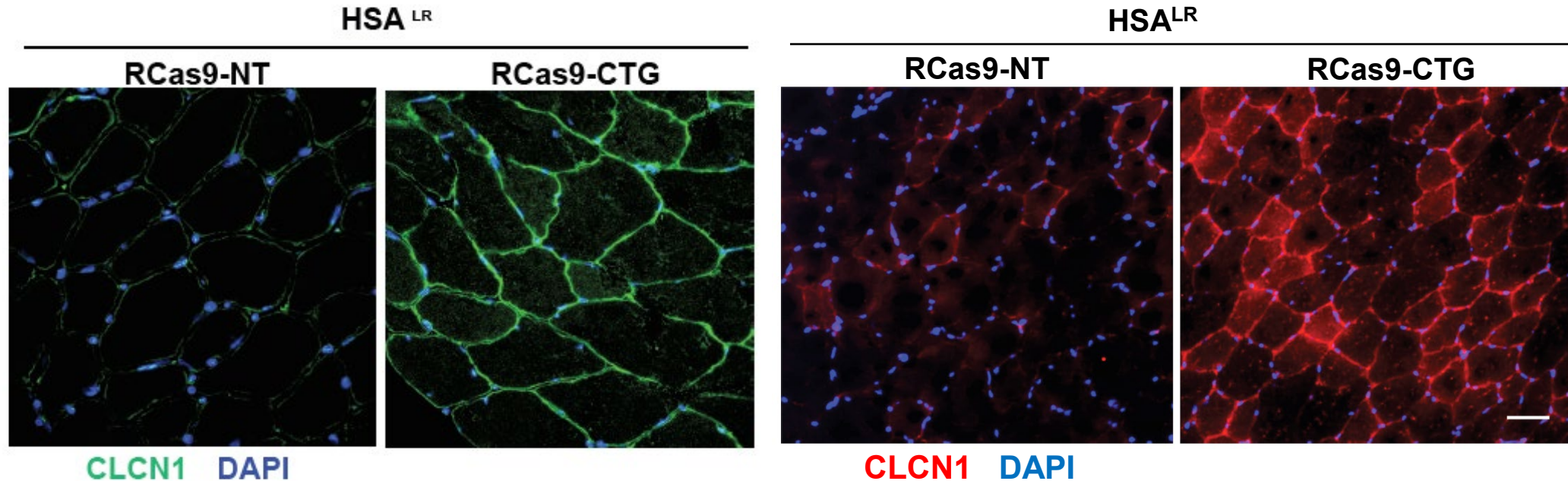
# Intramuscular AAV9-RCas9 Eliminates CUG Repeat Foci and Reverses Hallmark Splicing Defects in Adult DM1 Skeletal Muscle

**Study Design:** HSA<sub>LR</sub> mouse model of DM1 expresses ~250 CTG repeats under the control of human skeletal actin (ACTA1) promoter. Non-Targeting RCas9 (RCas9-NT) and CUG-Targeting RCas9 (RCas9-CTG) were injected in contralateral tibialis anterior (TA) muscles of HSA<sub>LR</sub> mice. The RCas9 system was delivered with two AAV9 vectors (2.5\*10<sup>10</sup> vg each; schematic A) containing Cas9 linked with the SMG6 RNA endonuclease PIN and CUG targeting single guide RNA (sgRNA) and GFP, respectively. Muscles were examined 4 weeks post injection.



# Intramuscular RCas9 Corrects CLCN1 Expression in DM1 Mice

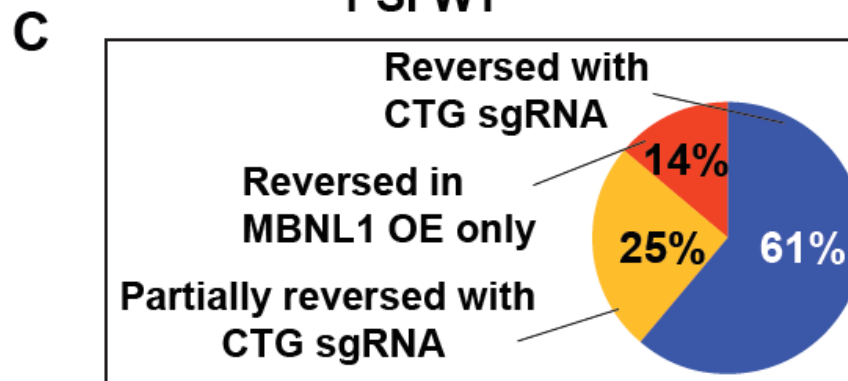
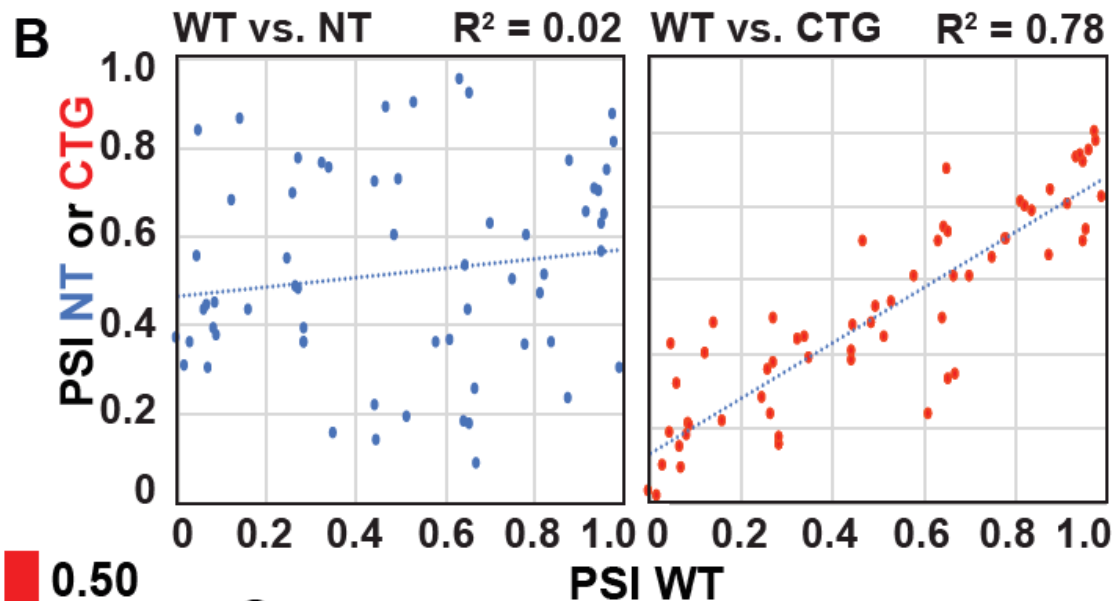
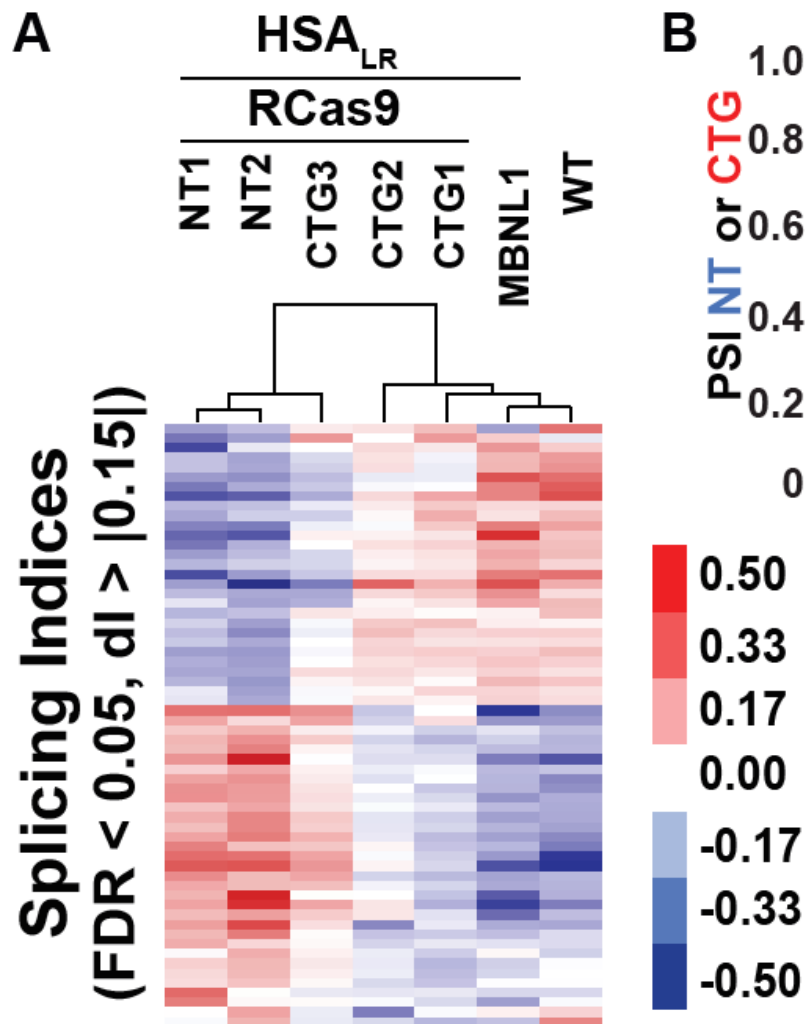
E



**Results:** (A) Schematic of Intramuscular Injection of RCas9-NT and RCas9-CTG into contralateral TA muscles. (B) RNA-FISH of contralateral TA muscles show elimination of toxic CUG RNA-foci with RCas9-CTG treatment. (C) Longitudinal section of TA shows that transduced fibers expressing RCas9-CTG (green) show clearance of CUG RNA foci whereas non-transduced fibers (dark) retain CUG RNA foci. (D) Reversal of DM1-related missplicing in RCas9-CTG treated muscle. Re-constitution of Chloride channel CLCN1 staining on RCas9-CTG treatment.

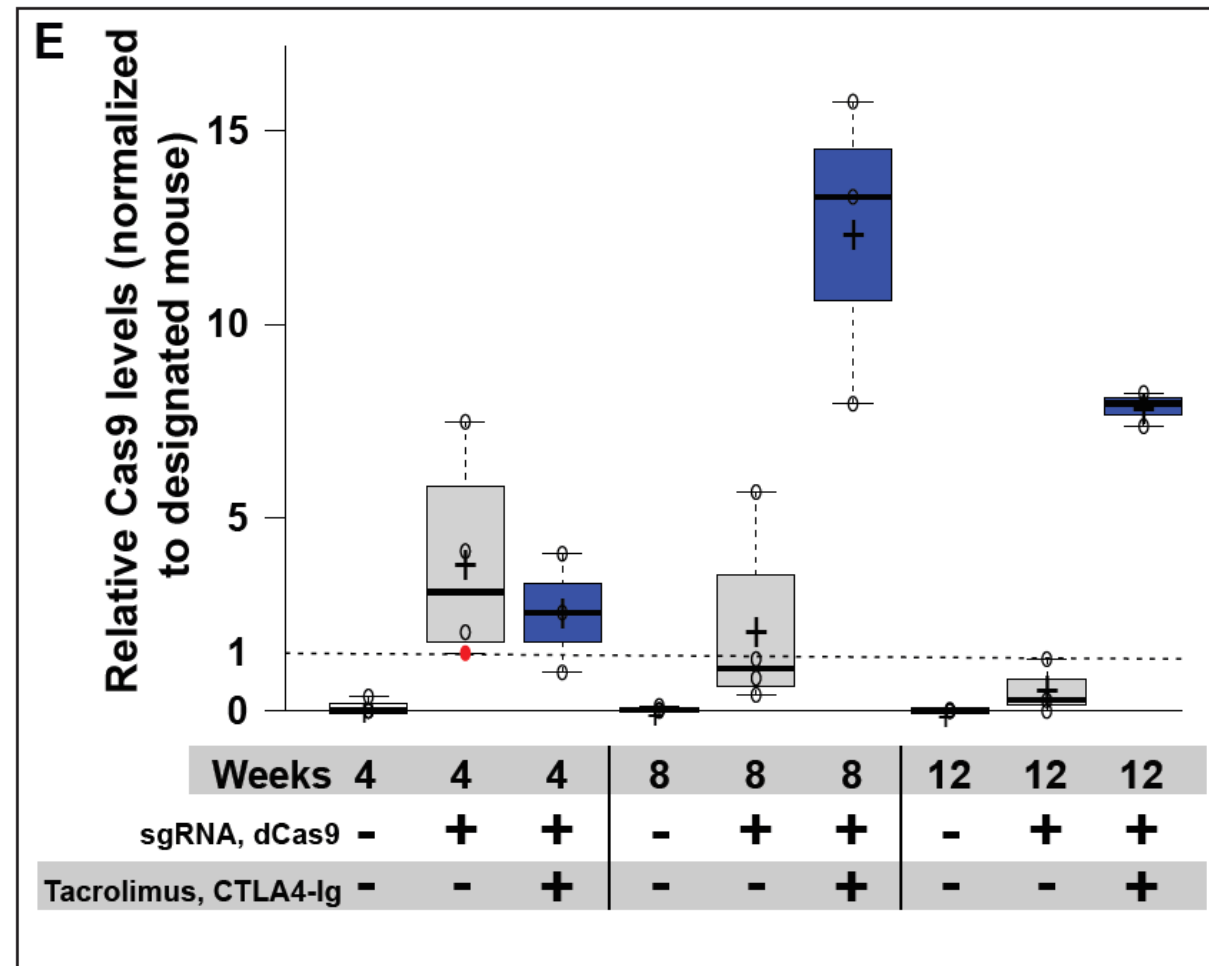
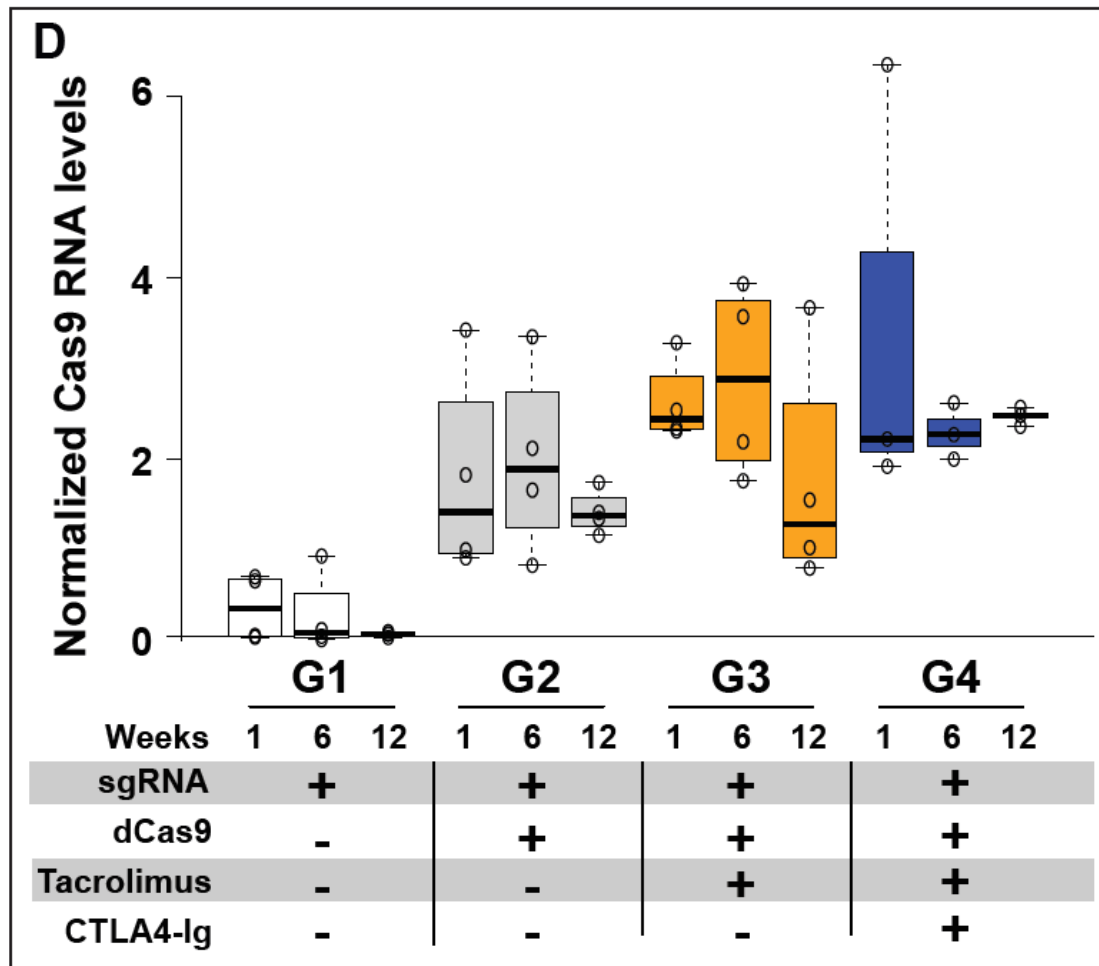


# RCas9 Reverses Transcriptome-wide DM1-Associated Missplicing



**Results:** (A) Hierarchical Clustering of exon splicing indices (percent spliced in or PSI) of HSA<sub>LR</sub> TA treated with either RCas9-NT (NT), RCas9-CTG (CTG) or MBNL1 (positive control). RCas9-CTG treated muscle clusters with MBNL1 and WT groups. (B) Scatter plots of PSI values of wildtype (WT) vs. NT (HSA<sub>LR</sub>) or WT vs. CTG (HSA<sub>LR</sub>) shows marked reduction of splicing defects after RCas9-CTG (CTG) treatment. (C) ~86% of top 350 DM1-related AS changes show partial reversal (25%) or complete reversal (61%) of splicing defects.

# Transient Immunosuppression Tolerizes RCas9-CTG Expression

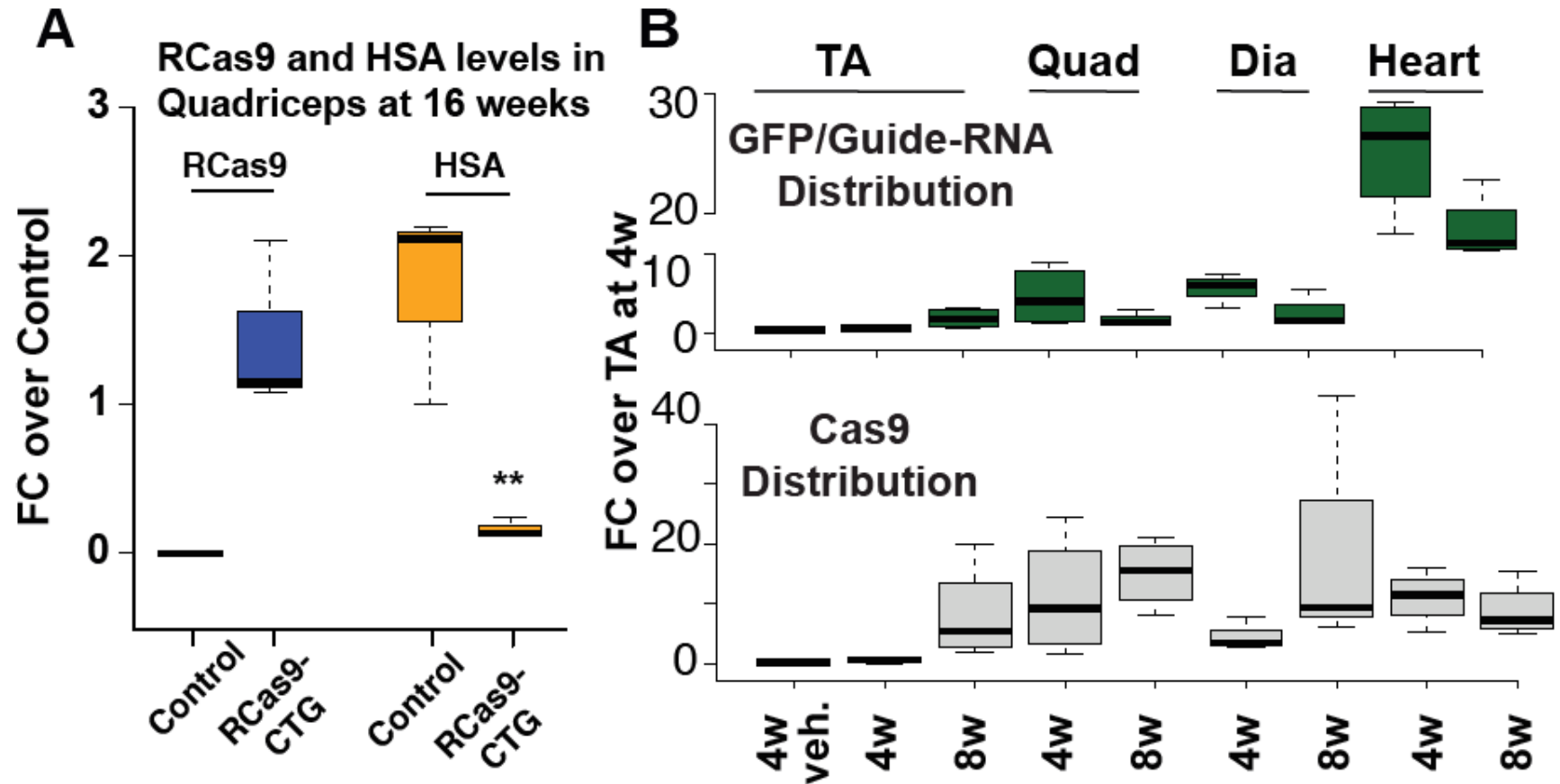


**Results:** (D) RCas9 levels were sustained in wildtype mice after TA injection for >12 weeks with or without immunosuppression with Tacrolimus and CTLA4-Ig (T-cell pathway). (E) RCas9 levels were sustained for >12 weeks post IM injection with immunosuppression with Tacrolimus and CTLA4-Ig but not without immunosuppression.



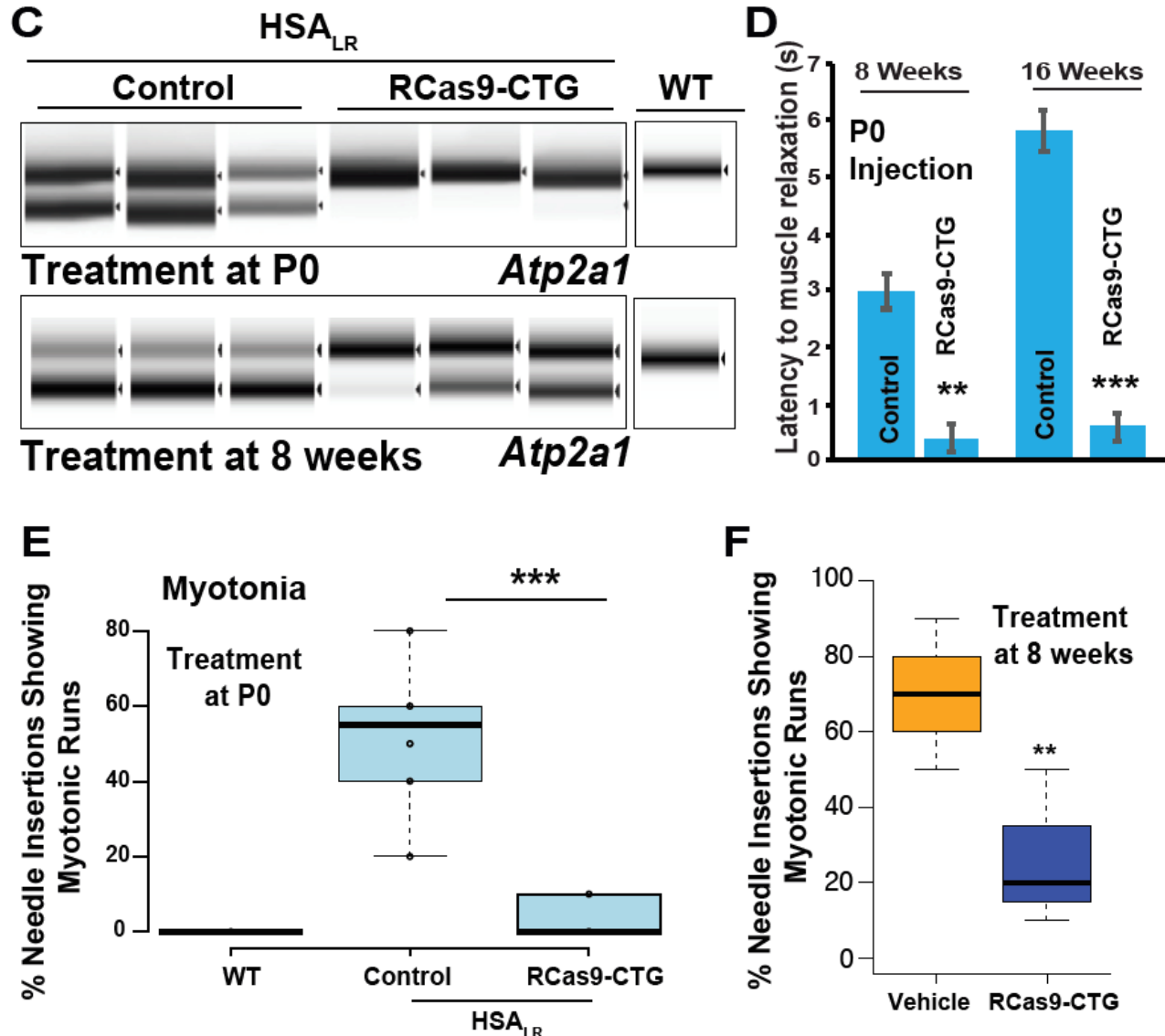
# Systemic RCas9 Corrects DM1-related Abnormalities

**Study Design:** Either P0 neonatal or adult HSALR mice were injected with either vehicle (control) or RCas9-CTG systemically in the temporal vein (1011 vg) or lateral tail vein (5\*1011 vg/kg) respectively. Mice were subjected to functional tests for mytonia and electromyography tests and tissues were processed at various time points (4-16 weeks) as indicated and analysis of RCas9-CTG expression and splicing biomarkers was carried out.



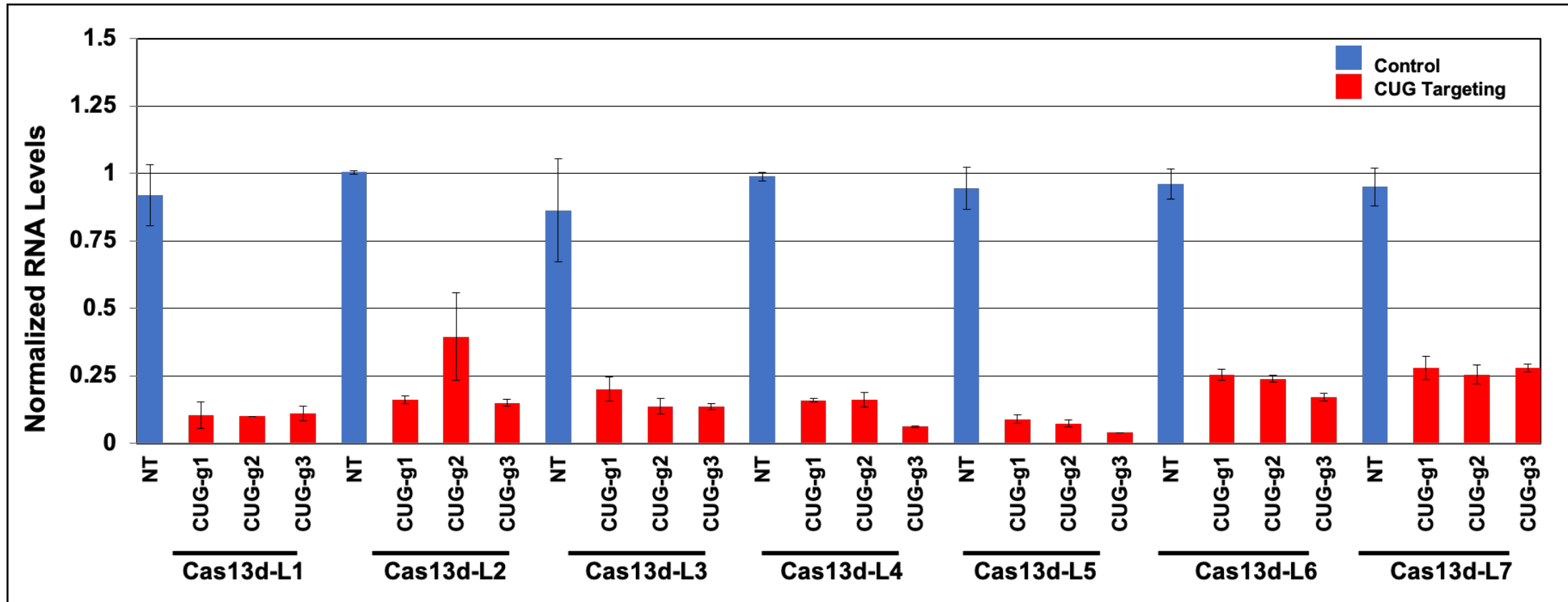
**Results:** (A) Boxplot shows mRNA levels of RCas9 and HSA (CUG repeat levels) in the quadriceps of treated HSALR mice 16 weeks after treatment of P0 HSALR mice. N=3; Student's T-test  $p < 0.01$ . (B) Box-plots show distribution mRNA levels of guide-RNA-GFP and Cas9 levels in various indicated tissues at 4 and 8 weeks after lateral tail vein injection (N=3).

# Systemic RCas9 Corrects DM1-related Abnormalities



**Results:** (C) Splicing of alternative exon 22 in *Atp2a1* assessed by RT-PCR in quadriceps. (D) Latency to hind-limb relaxation at 8 and 16w post-treatment of P0 HSA<sub>LR</sub> mice. (E and F) Electromyography (EMG, Gastrocnemius) of wildtype, vehicle, and AAV9-RCas9-CTG treated P0 HSA<sub>LR</sub> mice at 8 weeks post injection (E; N=5, Z-score = 2.8,  $p < 0.01$ ) or vehicle and RCas9-CTG treated adult mice 4 weeks post injection (F; T-test  $p$  value  $< 0.05$ ).

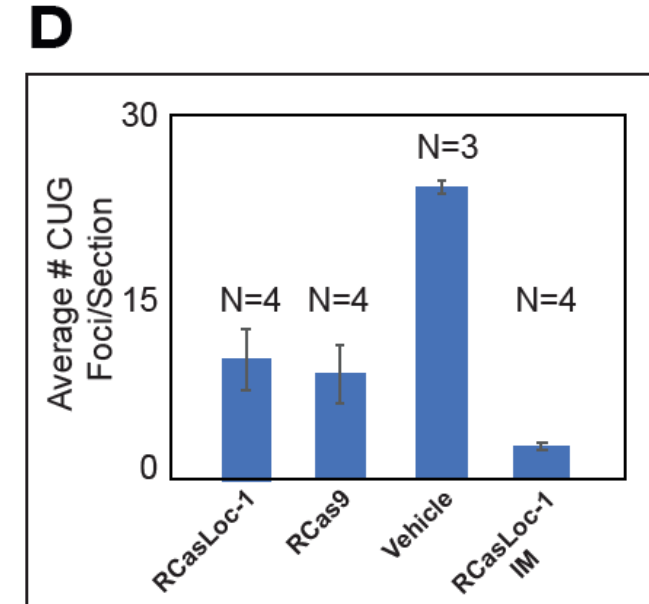
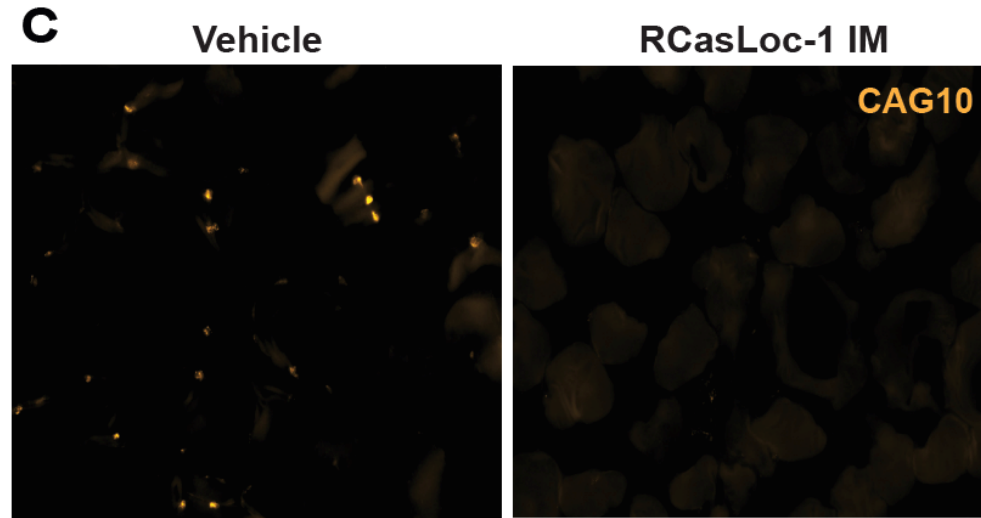
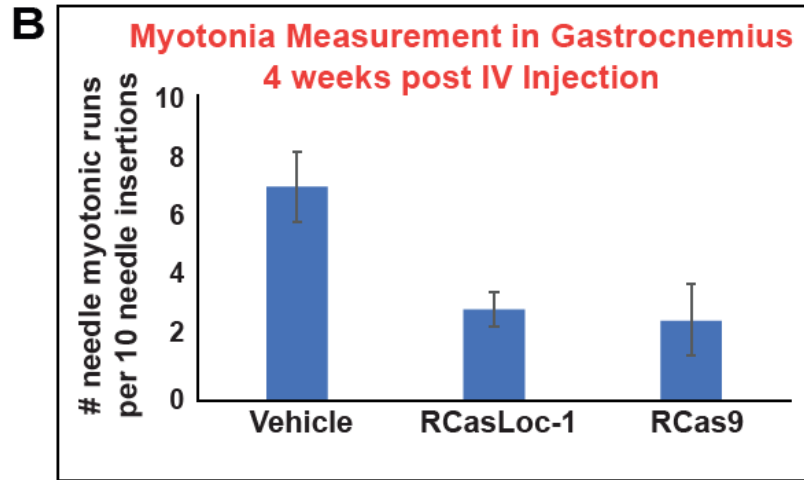
# Locana's Proprietary RNA Targeting Cas Proteins are Effective in Eliminating Expanded CUG RNA



**Study Design:** Various RNA Targeting Cas proteins were tested with cognate guide RNAs for three different frames of CUG repeats in vitro (cells).

**Results:** (A) Various RCas systems were tested for their CUG-targeting activity in vitro with transient transfection and RT-qPCR.

# Locana's Proprietary RNA Targeting Cas Proteins are Effective in Eliminating Expanded CUG RNA



**Study Design:** Locana's proprietary unitary (packed in a single AAV) Cas13d-L2 was tested against the RCas9 system with intramuscular (tibialis anterior,  $5 \times 10^{10}$  vg) and systemic delivery (lateral tail vein,  $10^{12}$  vg). Myotonia was measured with EMG and RNA foci were quantified 4 weeks post injection.

**Results:** (B) HSALR mice were injected in the lateral tail vein with AAV9 carrying RCas9 or RCasLoc-1. EMG was performed and number of myotonic runs were recorded for 10 needle insertions. Both RCas9 and RCasLoc-1 (Cas13dL2) decreased myotonia. (C) RNA-FISH was performed to stain for CUG RNA foci in HSALR TA and we observed a decrease in RNA foci with RCas9 and RCasLoc-1.



# Conclusions

## Conclusions

- RNA Targeting Cas9 targets and cleaves CUG RNA foci *in vitro* and *in vivo*.
- IM of RCas9 reverses DM1-related splicing abnormalities and restores CLCN1 staining.
- RCas9 levels were sustained for >12 weeks post IM injection in wildtype and HSA<sub>LR</sub> mice when co-administered with Tacrolimus and CTLA4-Ig.
- Systemic injection of RCas9 leads to reduction in toxic CUG RNA foci, splicing defects, and myotonia.
- RNA Targeting Cas systems packaged in a **unitary AAV** vector show *in vitro* and *in vivo* CUG elimina-