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Supplemental Information

Gene Therapy via *Trans*-Splicing for *LMNA*-Related

Congenital Muscular Dystrophy

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Total Lmna

Supplemental information

Figure S1: Protein evaluation of trans-splicing in vitro.

(A) Representative western blot analysis for FLAG and lamin A/C expression relative to Ponceau red from C2C12 cells transfected or not (NT) with FLAG-prelaminA, various PTM-BD and PTM-BDinv. No trans-splicing event is detected by western blot. (B) Immunofluorescence analysis of C2C12 transfected with PTM-BD4 and PTM-BD9. Cells were double-stained with anti-FLAG (green) and anti-Lamin A/C (red) antibodies. Nuclei were stained with DAPI (blue). Arrows point on FLAG+ nuclei. (Scale bar: 10 µm).

Figure S2: Evaluation of the transduction rate in mouse primary myotubes.

Immunofluorescent confocal micrographs of WT mouse primary myotubes transduced with AAV2/9-CMV-GFP and stained for GFP (green) and Desmin (red). Nuclei were stained with DAPI (blue). (Scale bar: 100 µm).

<u>Figure S3:</u> Detection of repaired *Lmna* mRNA induced by 5' *trans*-splicing in mouse $Lmna^{\Delta K32/\Delta K32}$ primary myotubes.

Immunofluorescence analysis of *Lmna*^{$\Delta K32/\Delta K32} myotubes transduced with AAV2/9-PTM-BD4$ and PTM-BD9. Cells were double-stained with anti-FLAG (green) and anti-lamin A/C (red)antibodies (**A**) or with anti-lamin A/C (green) and anti-emerin (red) antibodies (**B**). Nucleiwere stained with DAPI (blue). Arrows show relocalization of lamin A/C to nuclear envelope.(Scale bar: 10 µm).</sup>

Figure S4: In vitro validation of Lmna 5' trans-splicing in primary mouse mutant myotubes.

Western blot analysis for lamin A/C expression relative to Ponceau red from primary mouse WT myotubes and mutant (Δ K32) myotubes transduced either with AAV2/9-PTM-0, -BD3 or - BD4. Transduction with AAV2/9-PTM-BD3 and -BD4 induces a small increase in lamin A/C protein level.

Figure S5: Improve in emerin and lamin B1 localization after *Lmna* 5' trans-splicing in vitro.

Immunofluorescence against emerin (**A**) or lamin B1 (**B**) in non-transduced (NT) or PTM-BD3transduced primary mutant (Δ K32) myotubes. Nuclei are counterstained with DAPI. Stars highlighted nuclei with corrected emerin or lamin B1 localization. Pictures on the right are a magnification of areas delimited with white doted squares. (Scale bar: 50 µm).

Figure S6: Evaluation of trans-splicing efficiency in vivo.

(A) Hematoxylin eosin staining of striated muscles sections from WT mice 50 days postsystemic injection with AAV2/9 expressing PTM-BD3. (Scale bar: 50 μm). *TA*: *Tibialis anterior*; *Gast: Gastrocnemius*. (B) Immunofluorescence analysis of *Tibialis anterior* sections from WT mice 50 days post-systemic injection with AAV2/9-PTM-0, -PTM-BD4 and PTM-BD4inv. Cells were double-stained with anti-FLAG (green) and anti-Lamin A/C (red) antibodies. Nuclei were stained with DAPI (blue). (Scale bar: 10 μm). (C) PCR analysis was performed with primers specific for *trans*-spliced *Lmna* (F-FLAG and R-*Lmna* exon 7), PTM (F-FLAG and R-*Lmna* exon 1) or total *Lmna* (F-*Lmna* exon 1 and R-*Lmna* exon 1) on RNA extracts from 15 days-old mouse heart, *Gastrocnemius* and liver after AAV2/9-PTM-BD4.

<u>Supplemental Table 1</u>: Sequence of the different binding domains targeting intron 5 of *Lmna* pre-mRNA.

BD1	CCCTGGAAAGCCAGACAGGCATCAGATTCAGAACCATAGCATGTGGTCCTGGAGTCCCTCCC
BD2	CATGTCCAGTCCAAGGGAAGCTCTATCTCCAAGTAGGGAATGCGGCTGGGTTTGTAACTCAGTGAGAA AGCACCTATCTACCAAGCAAGAGGCCTCTGGTTCAGTCTCCGGCAGGCA
BD3	TCAGTGAGAAAGCACCTATCTACCAAGCAAGAGGCCTCTGGTTCAGTCTCCGGCAGGCA
BD4	AGTTAAGGGTGACCTTGAATCTCTGATACTTCAACCTGAGATAAAAGGCATGCACCACCCAGTTCCCTGTGG TCCTGGGGATAGAAGCCAGGCCATCGTGTATAATAAGCAAGC
BD5	CTGCCTCCCGAGTGCTGGGATTAAAGGCCTGTGCTATTACAACCTGGCTTCTTTTCTTTC
BD6	TGTTTGTTTGTGGTATTTTGGTTTGGTTTGGTTTTTGGTTTTTCAAGACAGGGTTTCTCTGTATAGCCCTGG CTGTCCTGGAACTCACTTTGTAGACCAGGCTGGCCTCGAACTCAGAAATCCTCCTGCCTCTGCCTCCCGAGT GCTGGG
BD7	CACTGCAGTGTCATGACAGACCTGGGAGTCCCGGAGGACAAGAATGTTCATCATTTTCCACTTTTTTTT
BD8	TCTGTCTGAAACCTCTCATTCCGAGGCTAGACCCCCATCTTCTTTTGTCACAAATAGTGTAGCTATTGTGAC AAATAATATCGACACAAGTAACATTTCACACTGCAGTGTCATGACAGACCTGGGAGTCCCGGAGGACAAGAA TGTTCA
BD9	AAGGGCTGATGTCGATGAAGAGGGAGGGGGGCACAGGGGCAGGGGAGGGGACACTCAGGATCTGTAGGCTCTCT CTAGAGCCACCATGGCCTTATCCAGTGACCAGGAAGCCCTTCCTAATGTCTGTC

<u>Supplemental Table 2:</u> PCR primers used to amplify trans-spliced *Lmna* mRNA, total *Lmna* mRNA and PTM mRNA.

Primer	Sequence
F-FLAG	ATGGACTACAAGGACGACGA
F- <i>Lmna</i> ex1	GCCAGCTCTACCCCACTGT
R- <i>Lmna</i> ex1	CAGACTCAGTGATGCGAAGG
F- <i>Lmna</i> ex2	GGGGACTTGTTGGCTGCGCA
R- <i>Lmna</i> ex6-1	TCCAGGGCCAGCTTGATGTCCAG
R- <i>Lmna</i> ex9-2	CCATCTCTCGCTCTTTCTCA
R- <i>Lmna</i> ex7	CCGCACGAACTTTCCCTCTT