

Supplementary information

Functional disruption of the dystrophin gene in Rhesus Monkey Using CRISPR/Cas9

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Supplementary Figure legends

Figure Legends

Figure S1. Off-target analysis via T7EN1 assay. DNA gel data showing no cleavage by T7EN1 for the potential targeted DNAs. WT is a negative control, PC is a positive control.

Figure S2. Histological analysis of monkey muscles. (A) Anti-dystrophin immunostaining (left) and H&E staining (right) of 8-year-old monkey muscle sections. Scale bars: 50 (left) and 20 (right) μm . **(B)** H&E staining of skeletal muscles of full-term stillborn WT and Mut-2 monkeys. Scale bars: 20 μm .

Supplementary Table 1. Potential off-target sequences and their genes.

Supplementary Table-1

Potential off-target	Genbank Sequence ID	Sequences (potential targeted region bold, mismatch: red)	Primer Sequences (5' to 3')	Amplification (bp)
1	NW_001098157.1 chr12	tatggTg ATAAACTGACTCTTGG	TCATTTCCAGGCACACTTGA TTGACTCAGTGAGGGGACCT	538bp
2	NW_001122901.1 chr8	aaagcag ATAAACTGACTCTTGG	ACCAGAAGGCAGGTGAGAGA ATACCGAGAGCAAGGTGGTG	577bp
3	NW_001108738.1 chr1	Ga AAACAT ac AACTGACTCTTGG	TGCCAGGAAACAGGGTAAAG TGTTTGCCTGCAGTCATTTCC	511bp
4	NW_001096616.1 chr11	aaaATCATa AAATGACTCTTGG	GCCCAGAGTGTAATCTGCTCA TCAAATTAAGTTAACAAAAAGGAAGG	502bp
5	JV543636.1 chr12	GGAt ATCATAAA agGACa CTTGG	A TCTGTTGTTGGGTTCTACAAATG CTGCAGCCTGACTGACAGAG	538bp
6	JV638520 chr1	GGAAATCATct ACTG c CTCTTGG	TCCCTTCCTGCCTTAAATCC TCCCAACAGGCAAGAAAATC	552bp
7	NM_001261353.1 chrX	tGAAATCAaAAg CTG at TCTTGG	GCCAAAGAGAAATCAAAACAGAA TCCTAGGAATTTTTCCATTTCA	509bp
8	JU481420.1 chrX	GGAAATgAg AAACTGACT Ca TGG	ATGGCCAATGTTGTGTGAGA TAGCCAGCTCCTCACCAGTT	525bp
9	NW_001114282.1 chr3	tGAAATCAaAAg CTG at TCTTGG	TCCTTTTTTCTCATTCTGCAA CAGATTACACTCTGGGCCATT	552bp
10	NW_001124211.1 chr9	GGAAf TCAT tg cCTGACTCTTGG	GGGCTACCCAATCTGACAAA GGGCTACAATCTGGCTTCAG	534bp
11	NW_001122901.1 chr8	GGt AAT g ATAAA a TGACT f TTGG	CCTGACCTGACAAAGCCATT GCAGTGTCCAGGGTTACCAC	565bp
12	NW_001101665.1 chr15	GcAAc TC ac AAACTG ac aCTTGG	CCACCTCCCACATAAACCAC TGAGGCCTTCATGCCTAGAT	596bp
13	NW_001118155.1 chr5	Ga AAA f TCATAAACT Ga CT c TGG	AAGCCACGCTTTTTCAACTC TGATCAGAGGAATGGATGGAG	590bp



