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Supplementary Information

A HaloTag-TEV genetic cassette for mechanical phenotyping of native proteins

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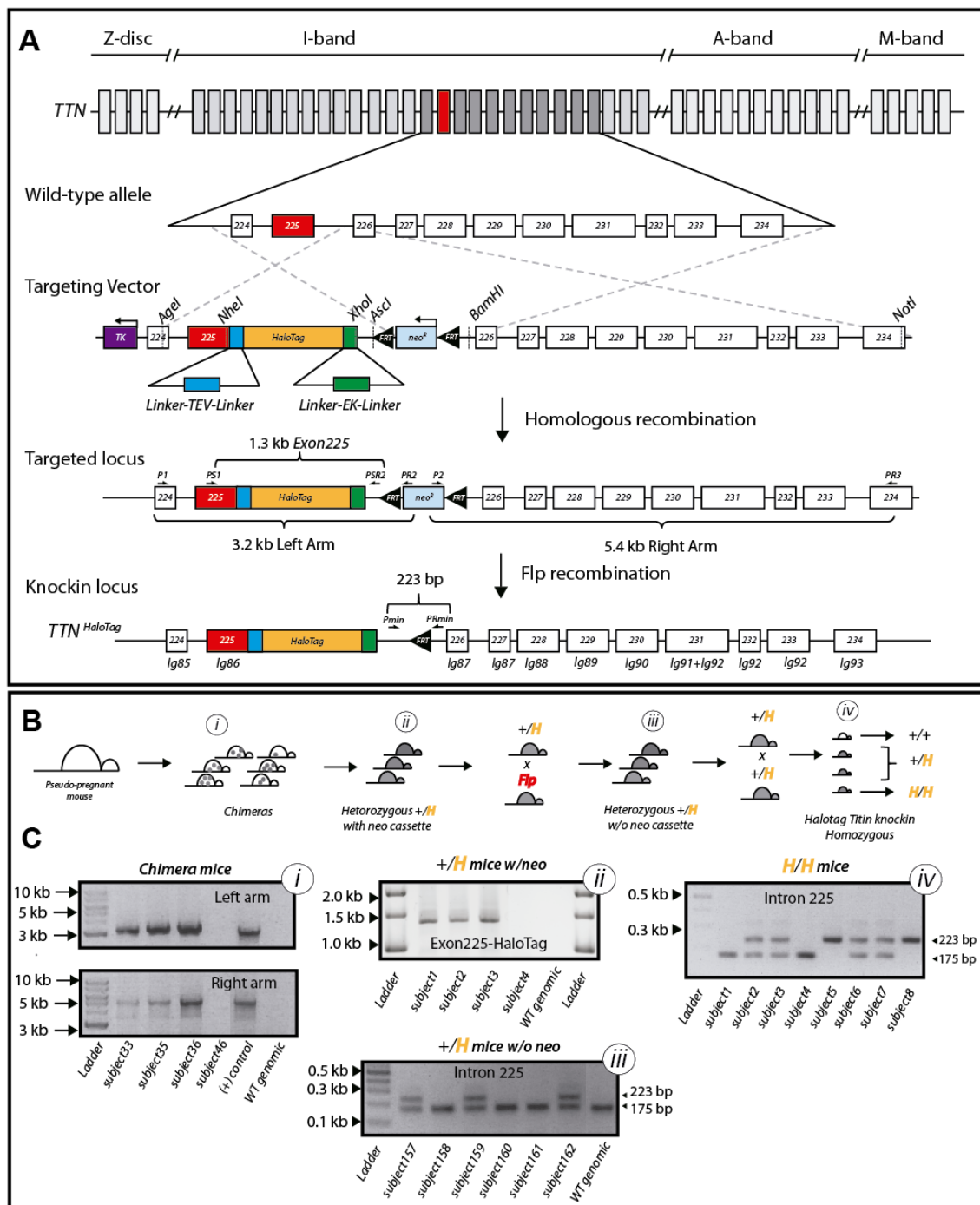
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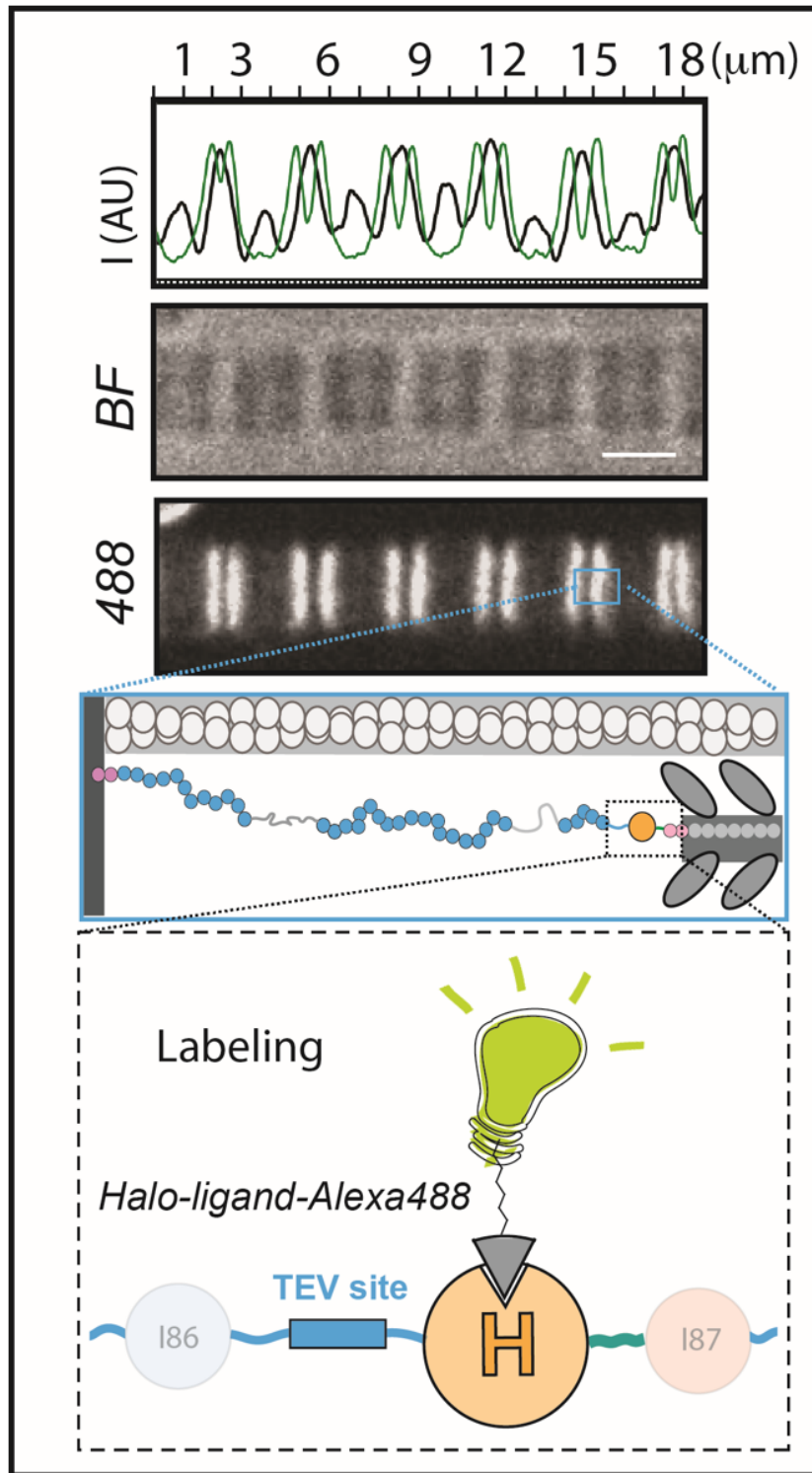
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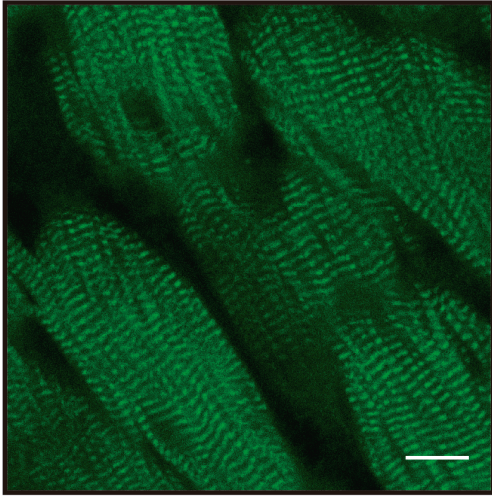


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 2
 3 **Supplementary Figure S1. Generation of knock-in mice with the HaloTag-TEV cassette**
 4 **inserted into the titin gene.** (A) The targeting vector contains the *TTN* sequence between exons
 5 224 and 234. The HaloTag gene is inserted downstream of exon 225, flanked by linkers
 6 including TEV and EK sites. A Neo resistance gene flanked by FRT elements was inserted in
 7 intron 225. Restriction sites and primers used during the different genetic engineering steps are
 8 shown, together with the size in base pairs (bp) of relevant fragments. (B) Strategy followed to
 9 get homozygous knock-in mice (H/H) from recombinant ES cells. The different experimental
 10 stages are labeled i-iv, see below. (C) We used PCR amplification of genomic DNA to confirm
 11 (i) the presence of the HaloTag in the *TTN* gene in chimera mice (primers *P1* and *PR2*, *left arm*;
 12 primers *P2* and *PR3*, *right arm*), (ii and iii) the presence and subsequent removal of the neo
 13 resistance by crossing with Flp mice (primers *PS1* and *PSR2*, Exon225-HaloTag; *Pmin* and
 14 *PRmin*, Intron225), and (iv) the generation of the homozygous H/H titin mouse (*Pmin* and
 15 *PRmin*). We used the vector construct and wild-type genomic DNA as controls.

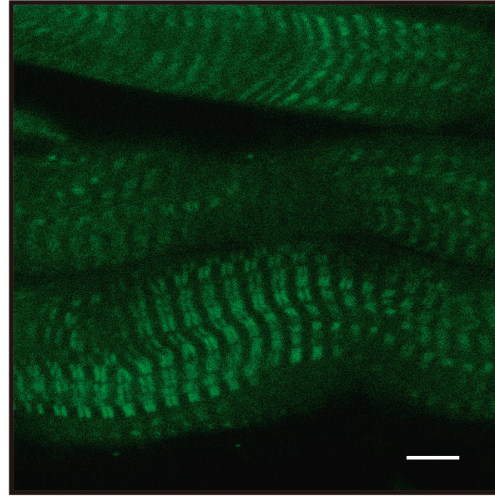


Supplementary Figure S2. Spinning disk microscopy. Under bright field illumination, the A- and I-band sections are easily distinguished along the myofibril as dark and clear regions, respectively (middle panel, BF). The fluorescent signal coming from Alexa488-labeled HaloTag-TEV titin appears as doublets at the I/A band interface (bottom panel, 488). The top panel shows intensity profiles (Alexa488 fluorescence in green and brightfield intensity in black). *Insets:* Cartoons showing the location of the HaloTag-labeled titin. Scale bar, 2.5 μm .

Confocal

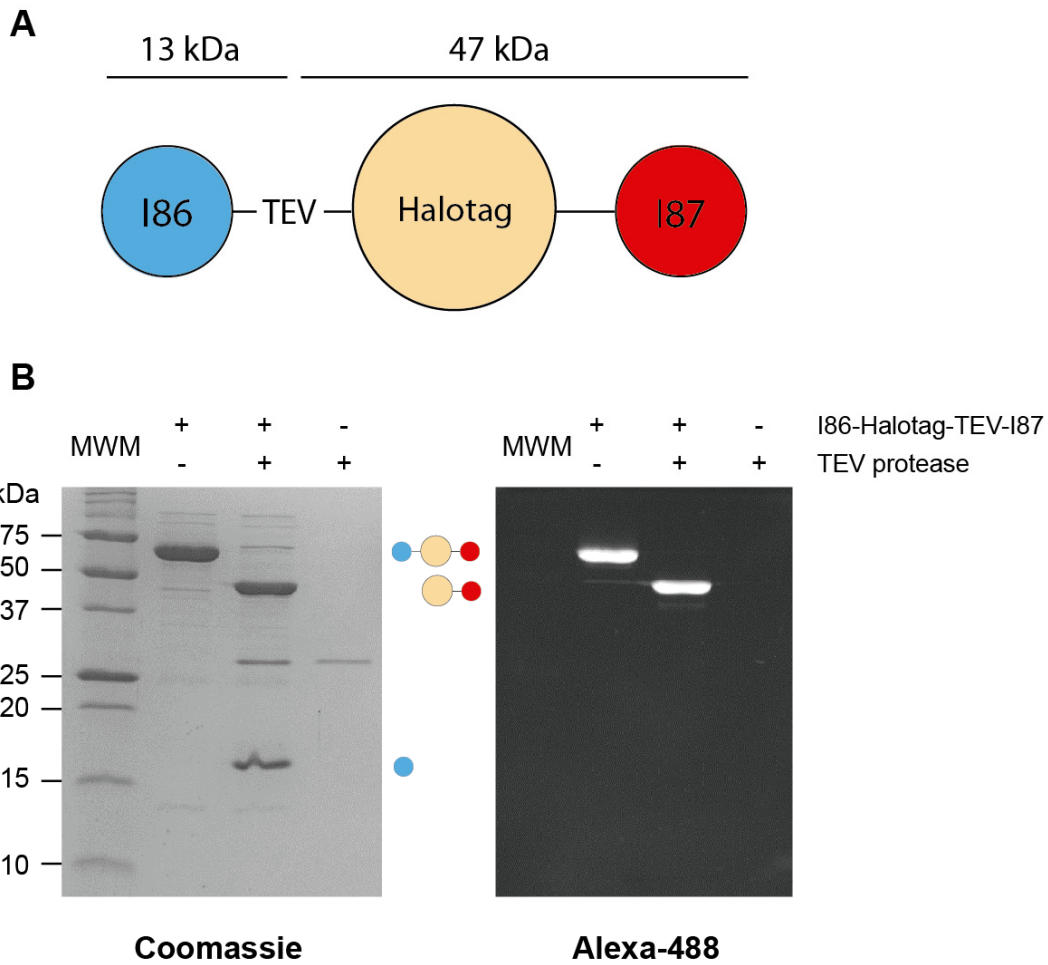


STED



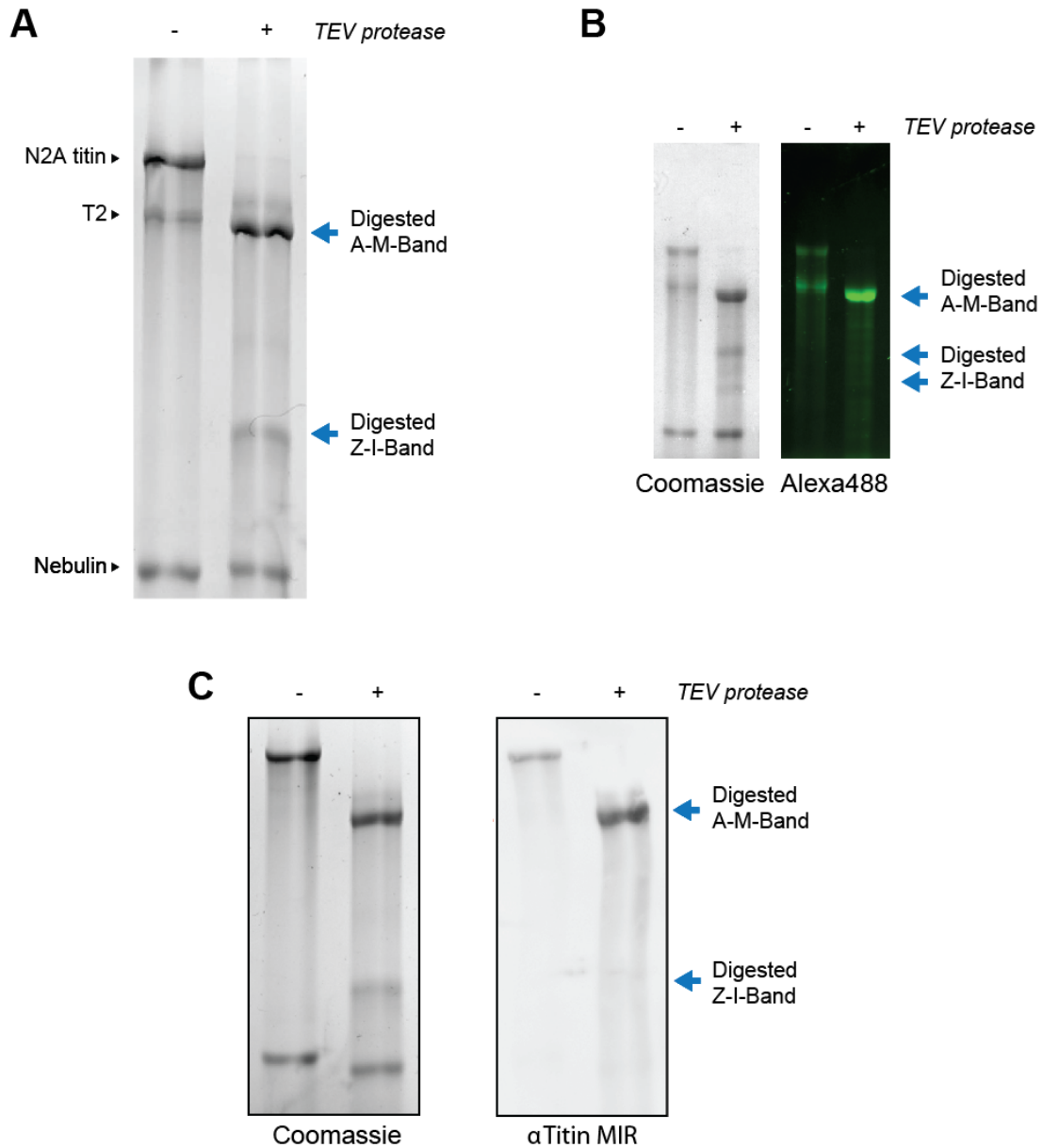
Supplementary Figure S3. The HaloTag-TEV cassette is correctly inserted in cardiac titin. The heart of a homozygous HaloTag-TEV titin mouse was incubated with HaloTag Oregon Green ligand, fixed and clarified. Although these samples show higher autofluorescence than skeletal preparations (**Figure 1B**), there is strong labeling in bands as expected from the location of the HaloTag insertion in titin (**Figure 1A**). Staining in doublets can also be observed, although to a lesser extent than in skeletal muscle, probably reflecting shorter I-bands in cardiac sarcomeres. Scale bars: 10 μm (Confocal) and 5 μm (STED).

1

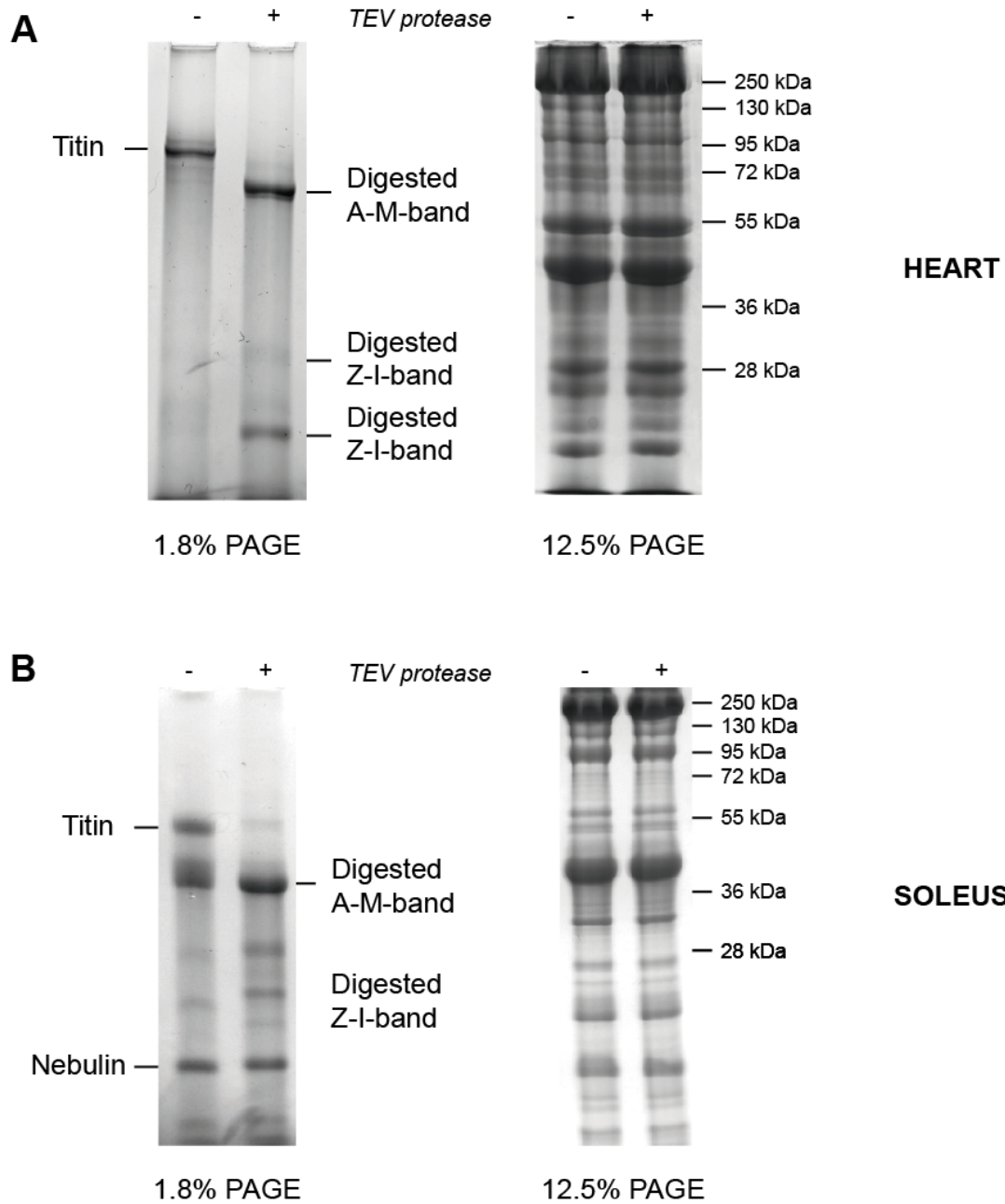


Supplementary Figure S4. Digestion of recombinant HaloTag-TEV titin fragment. (A) Scheme of I86-HaloTag-TEV-I87 (60 kDa) showing the size of the fragments that result from TEV digestion. (B) I86-HaloTag-TEV-I87 was treated or not with TEV protease (28 kDa) at 34°C for 1 hour, and results were analyzed by 17% SDS-PAGE. Digestion resulted in the appearance of two new bands at the expected mobility (*left*, Coomassie staining). Specificity of TEV-cleavage is demonstrated by labeling with HaloTag Alexa488 ligand, which only reacts with HaloTag-containing bands (*right*, Alexa488 fluorescence).

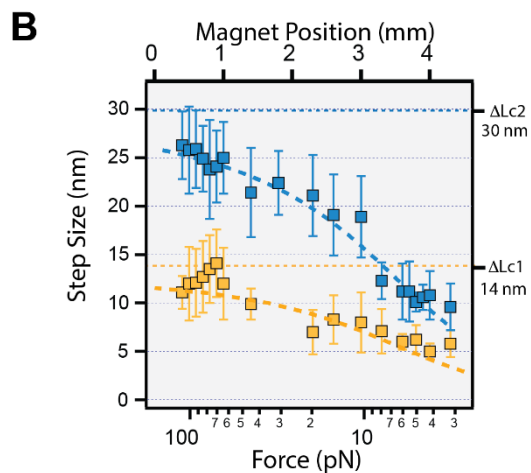
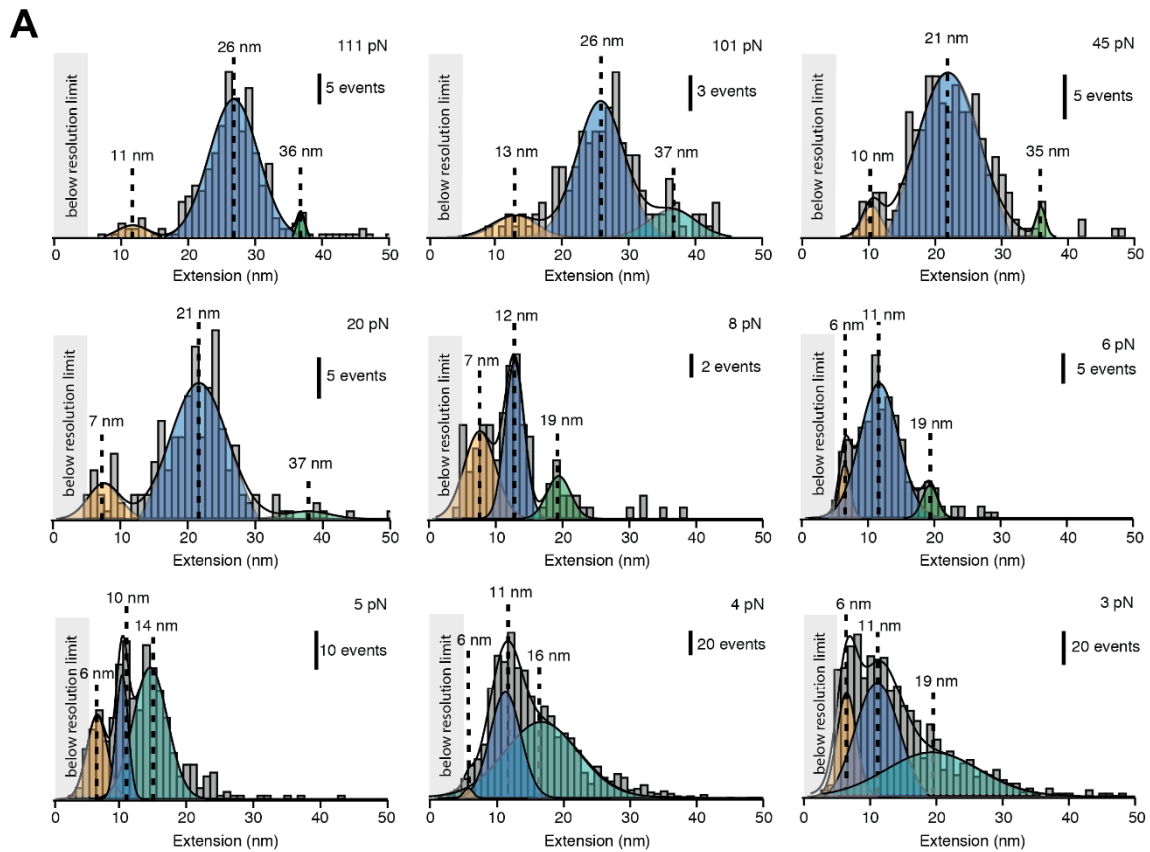
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1
2 **Supplementary Figure S5. TEV digestion of skeletal muscles from homozygous HaloTag-**
3 **TEV titin mice, as analyzed using 1.8% acrylamide SDS-PAGE gels. (A)** Results of TEV-
4 digestion of isolated soleus myofibrils (Coomassie staining). **(B)** Analysis of the digestion of
5 psoas myofibrils with TEV. Both Coomassie staining and HaloTag Alexa488 ligand are used to
6 visualize proteins. The HaloTag-specific Alexa488 ligand only labels the digested A-M-band
7 fragment **(C)** Equivalent results are obtained in TEV digestions of soleus myofibrils, as analyzed
8 by western blot using the MIR antibody, which recognizes the A-band segment of titin.
9



1
2 **Supplementary Figure S6. Effect of TEV treatment on the protein composition of striated**
3 **muscle from homozygous HaloTag-TEV titin mice. (A) Left:** 1.8% SDS-PAGE shows TEV-
4 induced specific digestion of cardiac titin. *Right:* the pattern of protein bands in 12.5% SDS-
5 PAGE gels remains unaffected by TEV treatment. **(B)** Equivalent results are obtained with
6 soleus samples. In this particular experiment, specific assignment of Z-I-band fragments is
7 hindered by some non-specific degradation of titin already present in the -TEV sample.



1

2 **Supplementary Figure S7. (Un)folding step sizes.** (A) Single titin molecules extracted from
 3 gastrocnemius muscles were pulled at different forces between 3 and 111 pN, and the size of the
 4 unfolding and refolding events was measured. We found three main populations of step sizes
 5 (solid lines are Gaussian fits to the data). (B) Force-dependency of the step sizes for the two
 6 populations corresponding to single-domain unfolding events, and fits to the worm like chain
 7 model of polymer elasticity (dashed lines). We obtain $\Delta L_C = 30 \pm 1$ nm (blue) and $\Delta L_C = 14 \pm 1$
 8 nm (yellow).

9

1 **Supplementary Text S1. Sequence of the targeting vector to introduce Halotag-TEV in**
2 **titin.** Exons and Introns are shown in grey and white background, respectively.

3
4 Exon224—Exon225—**TEV**—HaloTag—**EK**—NeoFRT—Exon226—Exon227—Exon228—
5 Exon229—Exon230—Exon231—Exon232—Exon233—Exon234

6
7 **Exon224**

8 GTGGTGACACGTT CAGAAGGAAGAGTTCACACGCTCACCC TGAGGGATGTGAAGCTAGAA
9 GATGCTGGCGAAGTCCAAC TAACTGCAAAGGATTTCAAAC TCAGGCCAATCTCTTTGTG
10 AAAG

11 gtaattagaaaacttattcctaaatacacacaatgaagacaatcacaacctctttat
12 tgtgggagagtaccacagattattaaaatcaagtttccaaaagtctaatttttacat
13 ggcataacaaaatgcagggttttttgtttttgtttttgttttaactataagcagt
14 gtagtacatgcctttaaaccagcactcaggaggcagaggcaggtggatctctgtgag
15 caaggtcagcctggtctacagagtgagttccaggacatgcagggccacacatagaa
16 tctcagaaaagaaagacaaagggatggattaaaactaacatgggttttatactta
17 tcttctgtctctgttttattattgtatttttactagttcctaagtacaactgctaaa
18 aatgattaccagtcctatgttatgaatcatttttagccattatctttatgcttcat
19 taattttctgtgaagctgcaatcagcagaggattgccacaagtttgaaccagcctg
20 tacatagtgggtttccaggatagctaggaaaaggggaaaaaacaacaacaaacat
21 agaacagtggtttattttcctctgtatataataatcaatgacaacacatcattgtg
22 aaaacatctaagtgaaaaatagctttacaacaagcaactgagacagaagcagccc
23 agatgccagagggaaatcagagaaagttctggaactcttggaaggaaacagcta
24 gtatgattctttcag

25 **Exon225**

26 AACCCCGGTTGAGTTCAC TAAAGCCTCTTGAGGACCAGACGGT CGAAGAGGAGGCCACTG
27 CAGTACTGGAGTGTGAAGTATCCAGAGAAAATGCCAAAGT GAAATGGTTCAA
28 AATGGGA CAGAAATCCTCAAAGCAAGAAGTATGAAATCGTTGCTGATGGCAGGGT
29 CAGGAAGCTCA TTATTCATGGTTGTACCC CAGAGGATATCAAACGTACACTTGT
30 GATGCTAAAGATTTA AGACCTCCTGTAACCTGAATGTTGTTCTG

31 **HaloTag**

32
33 GCTAGCGACAACACCACACCTGAG **GAGGACCTGTACTTCCAGAGC** GACAACACCACACCCGAGGCC
34 GAAATCGGAACAGGCTTCCCTTTTCGACCCCCATTATGTGGAAGTGCTGGGCGAGAGGATG
35 CACTACGTGGATGTTGGACCCAGGGATGGCACCCCTGTGCTGTTCC TGCATGGCAACCC
36 ACCAGCAGCTACGTGTGGAGGAACATACCCCCATGTTGCTCCTACACATAGATGCATC
37 GCTCCAGATCTGATTGGAATGGGAAAGAGCGATAAAACCTGATCTGGGATATTTTTTCGAT
38 GACCATGTGAGATTTATGGATGCTTTTCATTGAAGCTCTGGGACTGGAAGAAGTGGTGC
39 TGATTTCATGATTGGGGAAGCGCTCTGGGATTTTCATTGGGCTAAAAGAAAATCCTGAA
40 AAGTGAAGGAATTGCTTTTATGGAATTCATCAGGCCTATCCCTACCTGGGACGAATGGCCA
41 GAATTCGCCAGGGAGACCTTCCAGGCCCTCCGGACAACAGACGTGGGCAGAAAGCTGATC
42 ATCGATCAGAACGTGTTTCATCGAGGGAACCC TGCCATGGGAGTGGTCAGGCCCTGACC
43 GAGGTGGAGATGGACCACTATAGGGAGCCCTTCC TGAACCCGTTGACCGGGAGCCCTG
44 TGGAGGTTCCCTAACGAGCTGCCATCGCCGGAGAGCCCGCAACATCGTGGCCCTGGTG
45 GAGGAGTACATGGATTGGCTGCACCAGAGCCCTGTGCCAAGCTGCTGTTCTGGGCGACA
46 CCCGGCGTGATCCCCCTGCCAAGCCGCTAGACTGGCTAAGAGCCTGCCAACCTGC
47 AAAGCTGTGGACATCGGCCCTGGACTGAATCTGCTGCAGGAAGACAACCCCGACCTGATC
48 GGCTCTGAGATCGCCAGGTGGCTGAGCACCC TGGAGATCAGCGGC GACAACACCACACC
49 GAG **GACGACGACGACAAG** GACAACACCACCCCGAGACTCGAG

50 gtaagtattcctccacaggacttggcatttgaagtcattgtagccaaaacaaccaacac
51 atgtaatgcatgcccctcttggcaactcacagatggatgctgacctaactgatacttc
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59 aactgctacctagctaatttgccttgccttttcttcttgcctcagaaaacattttttgc
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1 cgctctactgtatcatcctgtagagtagccccttgggatttcttccccttgctgctctac
2 ttctcttggagaaggctcagccgtggtggcttatcttctagccttccagctagcattgg

NeoFRT

3
4 GAAGTTCTATACTTTCTAGAGAATAGGAACTTCGGAATAGGAACTTCAGTGGCTATGGCAGGGCTTGCC
5 GCCCCGACGTTGGCTGCGAGCCCTGGGCCTTCACCCGAACCTGGGGGGTGGGGTGGGGAAAAGGAAGAAA
6 CGCGGGCGTATTGGCCCAATGGGGTCTCGGTGGGGTATCGACAGAGTGCCAGCCCTGGGACCGAACCC
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9 AAGGCGATAGAAGGCGATGCGCTGCGAATCGGGAGCGGCATACCGTAAAGCACGAGGAAGCGGTGAGCC
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11 CCAGCCGGCCACAGTCGATGAATCCAGAAAAGCGGCCATTTTCCACCATGATATTCGGCAAGCAGGCATC
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16 TCGCCCAATAGCAGCAGTCCCTTCCCGCTTCAGTGACAACGTCGAGCACAGCTGCGCAAGGAACGCCCG
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23 AACCACACTGCTCGACATTGGGTGGAAACATTCAGGCCCTGGGTGGAGAGGCTTTTTGCTTCTCTTGcA
24 AAACCACACTGCTCGCAATTCGAAGTTCCTATACTTTCTAGAGAATAGGAACTTCGGAATAGGAACTTC
25

26 ttctcttggagaaggctcagccgtggtggcttatcttctagccttccagctagcattgg
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40 catttgaattccaacattattctcttacag

Exon226

41
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43 [CTCGGTTTCGAGTGCAGAAATTTCCAAAGAAAATGAGAAG](#)

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45 tactattgacaaaatgcaaggaaagaagaactagtaagcctttctgacttccatttt
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57 aaggacccttgggatactagttcatgtcatggatactgtgtaaggaacaacatgaaaac
58 tgattttaagaaatctatttgtcttaaaatagat

Exon227

59
60 [GTTTCAGTGGTTTAAAGATGGTGTGCTGAAATTA AAAAGGGCAAAAAGTATGACATCATTTCT](#)
61 [AAGGAGCAGTACGAATTCCTGTCTATCAACAAAATGTCTACTGAATGATGAAGCAGAATAT](#)

1 TCCTGTGAAGTGAGGACAGCAAGAACTTCCGGCATGCTGACAGTCC TAG
2 gtgaatgtgaaggcttttcttttactaagcatactagtcaggaacccaacttccagttt
3 actgactgcatctcggttcttttctgcag

4 **Exon228**

5 AAGAAGAAGCTGTCTTCACAAAAAATCTTGCCAACCTTGAAGTTAGTGAAGGAGACACTA
6 TCAAACTGGTGTGTGAAGTCTCCAAGCCTGGGGCAGAAGTGATTTGGTACAAAGGGGATG
7 AGGAGATCATCGAAACAGGGAGATTTGAAATACCTTACTGATGGAAGGAAGAGAATCTTGA
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9 GAACGGACAGCAAAGTCAAAGTACACG
10 gtatgaaatctcagtggaagggcattttcttttcatccattcttgtcgtattaagata
11 caaatgcccgatggttttctactgtcttccaactttttctttccag

12 **Exon229**

13 AACTTGCTGCTGAGTTCATCTCGAAGCCTCAAAACCTTGAATTTCTTGAAGGAGAAAAGG
14 CTGAGTTTGTCTGCACTATCTCAAAGGAAAGCTTCGAAGTTCAGTGAAGAGGGGATGATC
15 AGACACTTGAATCTGGAGATAAATATGACATCATTGCTGATGGCAAAAAGAGAGTCTTAG
16 TTGTAAAGGATGCCACATTACAAGACATGGGCACCTACGTAGTCATGGTTGGGGCTGCCA
17 GAGCCGACGCTCACCTGACAGTCATTG
18 gtaagtttgtccttgcctccctgcaggcttaaatctgtagttttgatccttcatcatact
19 acaaagtcttttgattgtttacag

20 **Exon230**

21 AAAAACTCAGGATCATAGTTCCCTCTTAAGGACACCAAGGTGAAGGAACAACAAGAGGTTG
22 TCTTCAACTGCGAAGTCAATACTGAAGGTGCCAAAGCCAAATGGTTCAGAAATGAAGAAG
23 CCATATTTGATAGTTCAAAAATACATCATTCTCCAAAAGACCTGGTCTACACCCTCAGAA
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26 gtatgtgacatacatgacactagagaattttcccgcagcaatttatattatgcccgacaa
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28 **Exon231**

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30 TCACATTCTGGTGCAAGGTGAATCGTCTCAATGTGACACTGAAGTGGACCAAAAATGGAG
31 AAGAAGTGGCTTTTGACAACCGTATATCATACCGAATTGATAAGTACAAACACTCTCTAA
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33 AATCCGTGGCAGAGCTGCTCATCATAGAAGCCCCAACAGAATTCGTGGAGCACCTGGAAG
34 ACCAGACGGTCACAGAGTTTATGACGCTGTCTTCTCCTGCCAGCTCTCCAGAGAGAAAG
35 CGAATGTA AAAATGTTACAGAAATGGAAGAGAAATCAAGGAAGGCAAAA
36 gtacgcaaaacgtgtctgcctccctctgttgttctgtgtgtactttgacatcagacatt
37 tgcctaatactcatgctacaagctaactaatcattcaatctcttctttag

38 **Exon232**

39 ATACAAGTTTGAAGAGGATGGGAGCATCCACAGGCTCATCATAAAAGACTGCAGGCTGGA
40 GGATGAGTGTGAATACGCTTGTGGTGTAGAGGACCGCAAGTCCCAGCTAGACTTTTTGT
41 AGAAG
42 gttagttattggcttcaaggataattctagctgaagtgacaatctttttacaatgctaata
43 aaaaatacaaacacatatctgtttttatttttacattttctctccag

44 **Exon233**

45 AAATTCAGTTGAGATTATCAGGCCCTCTCAAGACATTTCTTGAAGCCCCCTGGTGCAGACG
46 TTATCTTCTTGGCTGAGCTCAACAAAGATAAAAGTGGAGGTCCAATGGCTTAGAAATAACA
47 TGATCGTCGTCCAGGGTGACAAGCACCCAGATGATGAGTGAAGGAAAGATACACAGGCTAC
48 AGATTTGTGATATTAAGCCACGTGACCAGGGCGAATACAGATTCATTGCCAAAGATAAAG
49 AAGCCAGGGCTAAACTTGAATTAGCAG
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53 tgtcaaaccactgttattcctggcatgggcagctgctcatgacttcttagaagagttcct
54 cactcaactgataggcaactcctgtggctcttttaaatatgaaacctgcataggtttccc
55 cccaaagccccattgtacatgagagaaagctgtctgacttatcctgaactaagtatgcag
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58 catcttcaccaaaccactgtttccaaacatgaatagttaggagtaaaacaataacaaca
59 caacaagctattataactctccaacagaatgcactgctatgaaactactgtggttatgat
60 aatactgtggtttatcaaatgtatagaaaaaaaacctatgtcatagtatgactatagta
61 tataactcaaacatgtttgtgtgttctctttgtcactag

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Exon234

```
CTGCACCTAAAATCAAGACAGCTGATCAAGATCTCGTCGTTGATGCTGGCCAGCCTCTGA  
CAATGGTGGTACCCTATGATGCCTACCCCAAAGCAGAAGCTGAATGGTTTAAAGAGAACG  
AACCTCTATCTACAAAAACCGTTGACACTACGGCTGAGCAGACTTCTTTCAGAATCTCAG  
AAGCCAAGAAGGACGACAAGGGGAGGTATAAAATCGTGCTTCAGAACAAGCATGGGAAAG  
CAGAGGGCTTCATCAATTTACAAGTTATTG
```

1 **Supplementary Text S2. Sequence of the I86-HaloTag-TEV-I87 recombinant construct.**
2 The QS peptide bond cleaved by TEV is indicated. Please note that DNA sequence do not
3 match the murine DNA because it was codon-optimized for optimal protein expression in
4 *E.coli*.

6 I86-TEV site-Halotag-EK site-I87

7 **cDNA:**

8 ATGAGAGGATCGCATCACCATCACCATCACGGATCCCTCCGGTTGAATTTACCAAACCGCTGG
9 AAGATCAGACCGTTGAAGAAGAAGCAACCGCAGTTCTGGAATGTGAAGTTAGCCGTGAAAATGC
10 CAAAGTGAAATGGTTTTAAAAACGGCACCGAAATCCTGAAAAGCAAGAAATATGAAATTGTGGCC
11 GATGGTCGTGTGCGCAAACCTGATTATTCATGGTTGTACACCGGAAGATATCAAGACCTATACCT
12 GTGATGCCAAAGATTTCAAACAGCTGCAATCTGAATGTTGTTCTGGCAAGCGATAATACCAC
13 TCCGGAA **GAGGATCTGTATTTTCAGAGT** GATAATACAACCCCTGAA **GCAGAAATCGGTACTGGC**
14 **TTTCCATTCGACCCCATTTATGTGGAAGTCTGGGCGAGCGCATGCACTACGTGATGTTGGTC**
15 **CGCGCGATGGCACCCCTGTGCTGTTCTGCACGGTAACCCGACCTCCTCCTACGTGTGGCGCAA**
16 **CATCATCCCGCATGTTGCACCGACCCATCGCTGCATTGCTCCAGACCTGATCGGTATGGGCAA**
17 **TCCGACAAACCAGACCTGGGTTATTTCTTCGACGACCACGTCGGCTTCATGGATGCCTTCATCG**
18 **AAGCCCTGGGTCTGGAAGAGGTGTCCTGGTCAATCACGACTGGGGCTCCGCTCTGGGTTTTCCA**
19 **CTGGGCCAAGCGCAATCCAGAGCGCGTCAAAGGTATTGCATTTATGGAGTTCATCCGCCCTATC**
20 **CCGACCTGGGACGAATGGCCAGAATTTGCCCGGAGACCTTCCAGGCCTCCGCACCACCGACG**
21 **TCGGCCGCAAGCTGATCATCGATCAGAACGTTTTTATCGAGGGTACGCTGCCGATGGGTGTCGT**
22 **CCGCCCCTGACTGAAGTCGAGATGGACCATTACCGCGAGCCGTTCTGAATCCTGTTGACCGC**
23 **GAGCCACTGTGGCGCTTCCCAAACGAGCTGCCAATCGCCGGTGAGCCAGCGAACATCGTCGCGC**
24 **TGGTCGAAGAATACATGGACTGGCTGCACCACTCCCTGTCCCGAAGCTGCTGTTCTGGGGCAC**
25 **CCCAGGCGTTCTGATCCCACCGCCGAAGCCGCTCGCTGGCCAAAAGCCTGCCTAACTGCAAG**
26 **GCTGTGGACATCGGCCCGGGTCTGAATCTGCTGCAAGAAGACAACCCGGACCTGATCGGCAGCG**
27 **AGATCGCGCGCTGGCTGTGACGCTCGAGATTTCCGGCGATAACACGACACCTGAAGATGATGA**
28 **TGATAAAGACAATACGACACCGGAAACACGTGCACCGCATGTGGAATTTCTGCGTCCGCTGACC**
29 **GATCTGCAGGTAAAGAAAAAGAAACCGCACGTTTTGAATGCGAGATCAGCAAAGAAAATGAAA**
30 **AGGTGCAGTGGTTTTAAGATGGTGCCGAAATCAAAAAGGCAAAAATACGACATCATCTCCAA**
31 **AGGTGCCGTTTCGTATTCTGGTTATTAACAAATGTCTGCTGAACGATGAAGCCGAATATAGCTGT**
32 **GAAGTTCGTACCGCACGTACCAGCGGTATGCTGACCAGATCTTAA**

33 **Protein:**

34 MRGSHHHHHHGS PPVEFTKPLEDQTVEEEEATAVLECEVSRENAKVKWFKNGTEILKSKKYEIVA
35 DGRVRKLI IHGCTPEDIKTYTCDAKDFKTSNCLNVV LASDNTTPE **EDLYFQ'** SDNTTPEAEIGT
36 GPFDPHYVEVLGERMHYVDVGP RDGTPVLF L HGNPTSSYVWRNI IPHVAPTHR CIAPDLIGMG
37 KSDKPDLGYFFDDHVR FMDAFIEALGLEEVV LVIHDWGSALGFHWAKRNP ERVKGIAFMEFIRP
38 IPTWDEWPEFARET FQAFRTTDVGRKLI IDQNVFIEGTLPMGVVRPLTEVEMDHYREPFLNPVD
39 REPLWRFPNELPIAGEPANIVALVEEYMDWLHQSPVPKLLFWGTPGVLI PPAAEARLAKSLPNC
40 KAVDIGPGLNLLQEDNPD LIGSEIARWLSTLEISGDNTTPE **DDDDK** DNTTPE **TRAPHVEFLRPL**
41 **TDLQVKEKETARFECEISKENEKVQWFKDGAIEIKKGKKYDIISKGAVRILVINKLLNDEAEYS**
42 **CEVRTARTSGMLTRS**

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- 1 **Supplementary Video S1. 3D reconstruction of HaloTag-TEV clarified muscle fibers.**
- 2 Following specific labeling of gastrocnemius muscle with Oregon Green Halo ligand, fixation
- 3 and clarification, a 250- μm -deep Z-stack was obtained using multiphoton microscopy (127
- 4 images). Individual images were intensity corrected and the software Imaris was used to
- 5 produce the 3D animation.