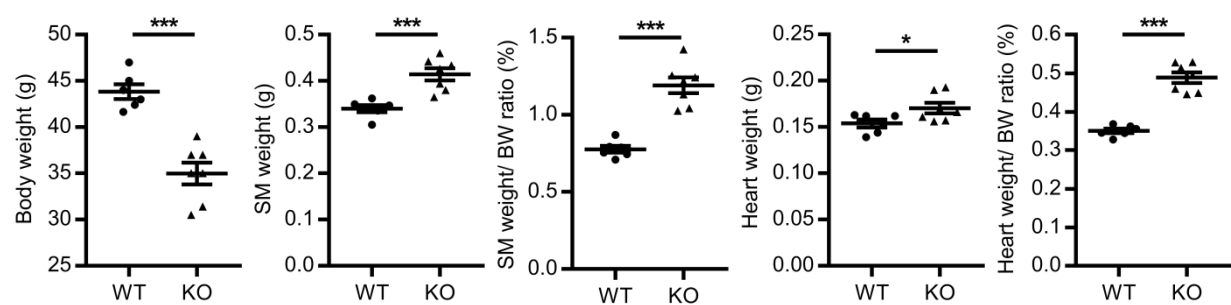
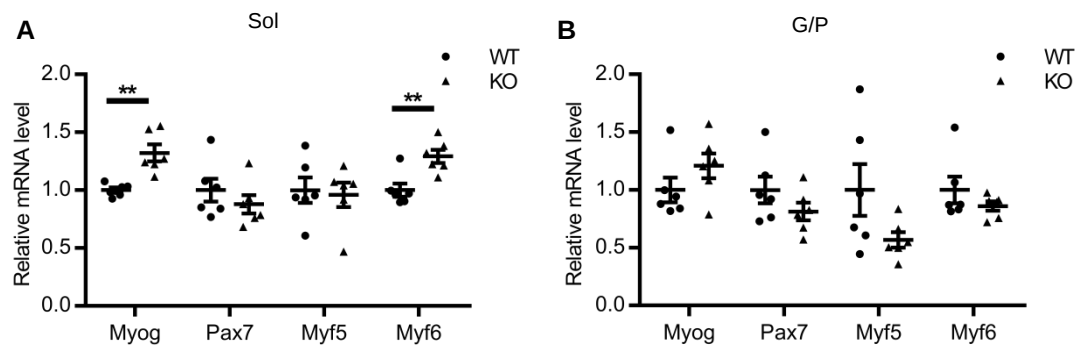


Supplemental Figure 1.



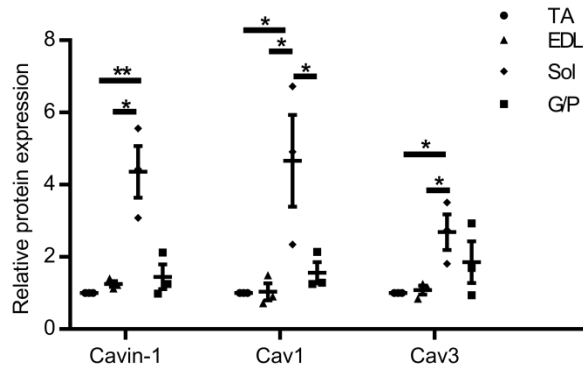
Supplemental Figure 1. Muscle weights and body weights in 7 month-old male WT and cavin-1 KO mice (n=6-7). Data are represented as mean $\pm$ SEM. 2-tailed Student's t-test was used for comparison between WT and KO mice. * $P < 0.05$ and *** $P < 0.001$.

Supplemental Figure 2.



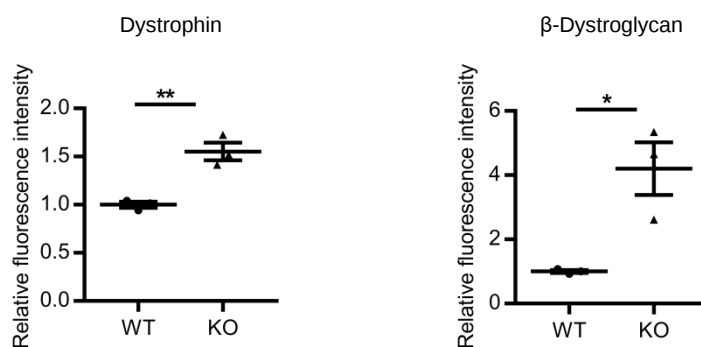
Supplemental Figure 2. mRNA expression of myogenic markers in **(A)** soleus (Sol) muscle and **(B)** gastrocnemius/plantaris (G/P) muscle of 3 month-old WT and KO mice (n=6). Data are represented as mean $\pm$ SEM. Bonferroni's multiple comparison test was used to determine the statistical significance. * $P < 0.05$ and ** $P < 0.01$.

Supplemental Figure 3.



Supplemental Figure 3. Densitometric quantification of caveolae protein levels determined by western blots in tibialis anterior (TA), extensor digitorumlongus (EDL), soleus (Sol) and gastrocnemius/plantaris (G/P) muscles from wild type mice, normalized to MyoD protein levels, indicating the highest protein expression of cavin-1, Cav1 and Cav3 in soleus muscle (n=3). Data are represented as mean $\pm$ SEM. 2-tailed Student's t-test was used for comparison between Sol and TA, EDL and G/P, respectively. * $P < 0.05$ and ** $P < 0.01$.

Supplemental Figure S4.



Supplemental Figure 4. Relative intracellular fluorescence intensity of dystrophin and β -dystroglycan on WT and cavin-1 KO muscle sections. Mean intracellular fluorescence intensity (sarcolemmal fluorescence excluded) was measure in >100 myofibers for each muscle cryosection (3 mice/group). Data are represented as mean $\pm$ SEM. 2-tailed Student's t-test was used for comparison between WT and KO. * $P < 0.05$ and ** $P < 0.01$.

Supplemental Table 1. Primer sequences used for quantitative real-time PCR

Gene	Forward Primer	Reverse Primer
Atrogin-1	AACCGGGAGGCCAGCTAAAGAACA	TGGGCCTACAGAACAGACAGTGC
MuRF-1	GAGAACCTGGAGAAGCAGCT	CCGCGGTTGGTCCAGTAG
Mstn	AGTGGATCTAAATGAGGGCAGT	GTTTCCAGGCGCAGCTTAC
Igf-1	TGGATGCTCTTCAGTTCGTG	TGGTAGATGGGGGCTGATAC
Col1a1	ACATGTTTCAGCTTTGTGGACC	TAGGCCATTGTGTATGCAGC
Col3a1	TGGTAGAAAGGACACAGAGGC	TCCAACCTCACCCCTTAGCACC
Col4a1	ACAAAAGGGTGATGCTGGAG	CTCCCTTTGTACCGTTGCAT
Col5a1	CTCCAACACCTCCAATCCAG	GTCCTCCAATCCCCTCAAAG
Col6a1	GATGAGGGTGAAGTGGGAGA	CAGCACGAAGAGGATGTCAA
Col6a2	ATGTGAGGGAGACCTGTGGA	TGTGCCTGTTTCTGACTTGG
Col6a3	CAGAACCATTGTTTCTCACT	AGGACTACACATCTTTTCAC
Fn1	TTAAGCTCACATGCCAGTGC	TCGTCATAGCACGTTGCTTC
Vim	CGGAAAGTGGAATCCTTGCA	CACATCGATCTGGACATGCTGT
Tgfb1	ATTCCTGGCGTTACCTTGG	AGCCCTGTATTCCGTCTCCT
Ndufa9	CTTTGGGCTGAGTCCATTG	TTGAGCTCCAGTGGTGTGG
Sdhb	GCACATTCACCTGTCACCAA	AGGGATTCAAGTACCCAGCAG
Cyt c	TCCATCAGGGTATCCTCTCC	GGAGGCAAGCATAAGACTGG
Cox5a	GGGTCACACGAGACAGATGA	GGAACCAGATCATAGCCAACA
Cox5b	GATGAGGAGCAGGCTACTGG	CAGCCAAACCAGATGATA
Nrf1	CTCTGCATCTCACCTCCAAAC	TCGCACCACATTCTCCAAAG
Erra	GCAGGGCAGTGGGAAGCTA	CCTCTTGAAGAAGGCTTTGCA
MtfA	TGCTAAAGATGATAGGATTCGTTACG	CGACGGATGAGATCACTTCGT
Ucp2	CCCTAATGGCTGCCTACCAA	GGGTCCAGGTCAGCATGG
Ucp3	TACCCAACCTTGGCTAGACG	GCCTGGCAATCTTTTGCTT
Mcad	CAACACTCGAAAGCGGCTCA	ACTTGCGGGCAGTTGCTTG
Lcad	GGACTCCGTTCTGCTTCCA	TGCAATCGGGTACTCCCACA
Scd1	GGCCTGTACGGGATCATACTG	GGTCATGTAGTAGAAAATCCCGAAGA
MHCI	CCTTGGCACCAATGTCCCGGCTC	GAAGCGCAATGCAGAGTCGGTG
MHCIIa	ATGAGCTCCGACGCCGAG	TCTGTTAGCATGAACTGGTAGGCG
MHCIIx	AAGGAGCAGGACACCAGCGCCCA	ATCTCTTTGGTCACTTTCCTGCT
MHCIIb	GTGATTTCTCCTGTCACCTCTC	GGAGGACCGCAAGAACGTGCTGA
Tfam	CTGCACTCTGCCATCCAAA	CTGAGCATTGCGAGGCCTTT
Nd1	GCCACCTTACAAATAAGCGCTCTC	ACGCAATTTCTGGCTCTGC
Myog	TTGCTCAGCTCCCTCAACCA	TGTGGGAGTTGCATTCACTGG
Pax7	TGCCGATATCAGGAGACTGGGTC	TTTCTCCACATCCGAGTCGC
Myf5	AAGGCTCCTGTATCCCTCAC	TGACCTTCTCAGGCGTCTAC
Myf6	GGGCCTCAAAGGCTTCGTT	CCACAGATCGTCGGAAAGCAG
36B4	GAGGAATCAGATGAGGATATGGGA	AAGCAGGCTGACTTGTTGC