

Figure S1. GFP reporter is expressed after heat-shock induction. Fluorescence dissecting microscope images of model flies (*hs-Gal4 UAS-i(CTG)480>UAS-GFP*) show strong fluorescence when transgene expression was induced (A) in comparison to uninduced counterparts (B).

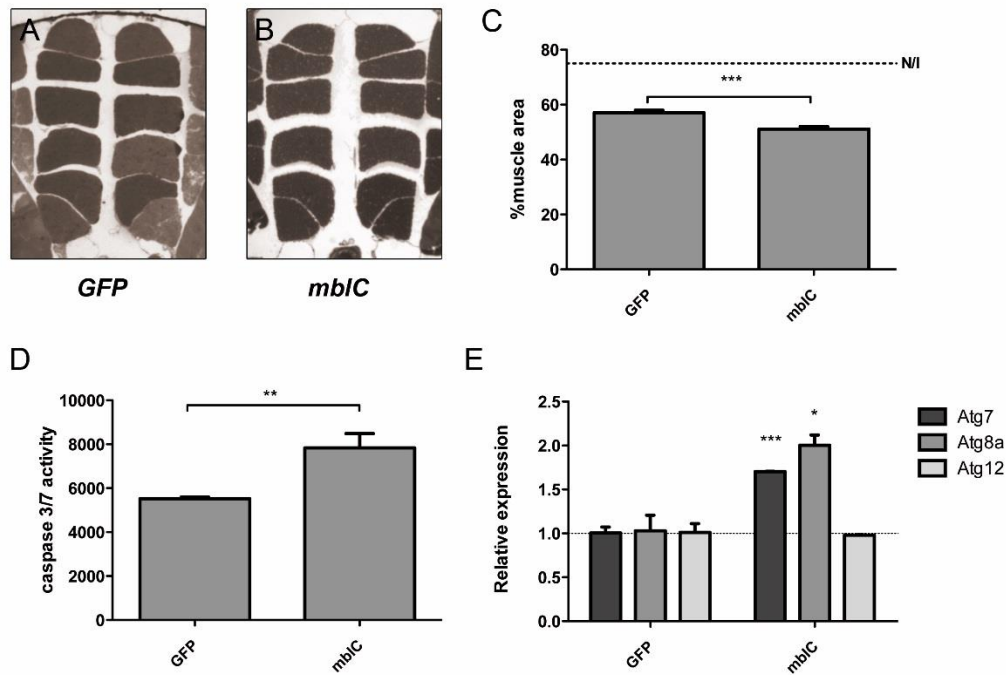


Figure S2. *mbIC* overexpression induces muscle atrophy. (A-B) Representative sections of resin-embedded thoraces of flies overexpressing the reporter GFP (A) or *mbIC* (B) under the control of the *hs-Gal4* driver. (C,D) Quantification of the mean percentage of muscle area per genotype indicates that *mbIC* overexpression induces both loss of muscle area, as well as the activation of caspase 3/7 activity. (E) Analysis of expression levels of *Atg7*, *Atg8a* and *Atg12* showed an increased expression of *Atg7* and *Atg8a* when *mbIC* was overexpressed. (* $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$). Error bars are standard deviations.

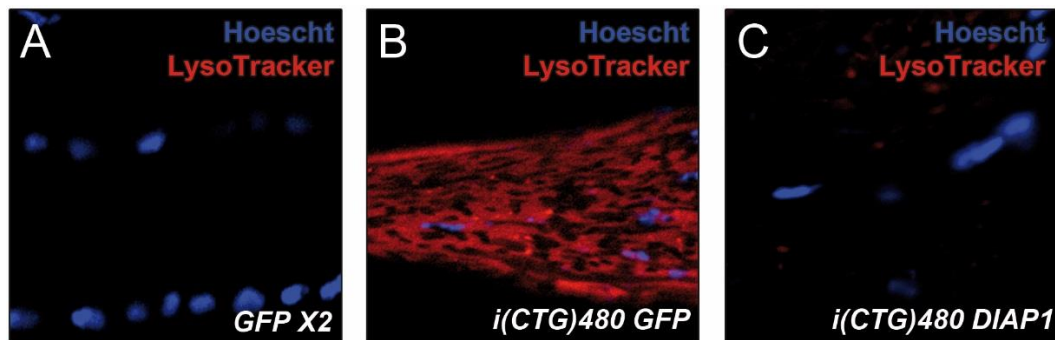


Figure S3 *DIAP1* overexpression in model flies reduces autophagy. Confocal fluorescence images of *Drosophila* IFM. Staining with LysoTracker (red) denotes that coexpression of *DIAP1* and *i(CTG)480* (*hs-Gal4 UAS-i(CTG)480>UAS-DIAP1*; C) results in autolysosome staining decrease when compared to DM1 model flies (*hs-Gal4 UAS-i(CTG)480>UAS-GFP*; B) to almost control levels (*hs-Gal4>UAS-GFP UAS-GFP*; A). Nuclei were counterstained with Hoechst (blue).

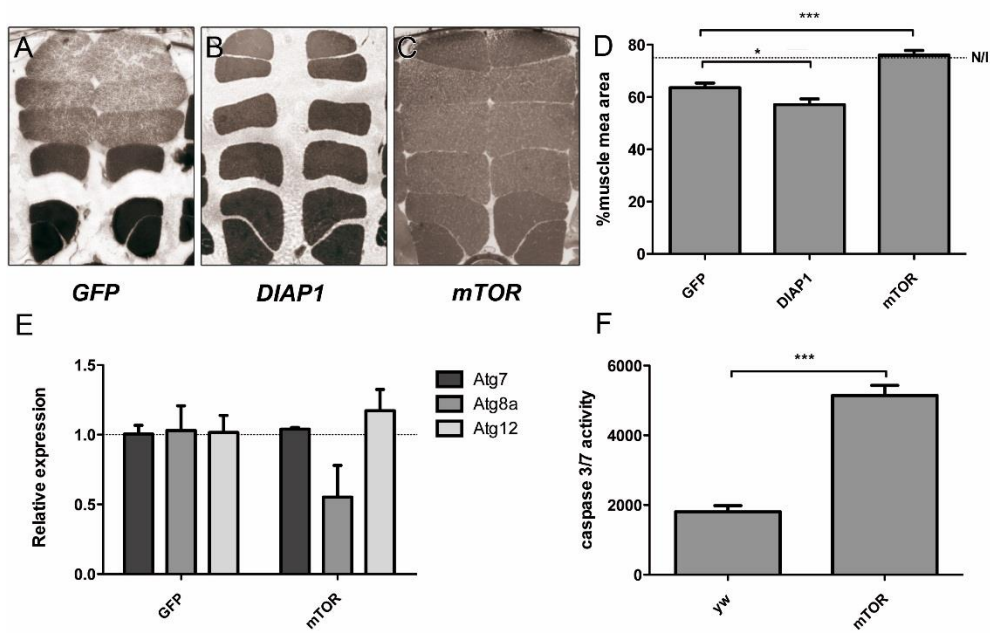


Figure S4. *DIAP1* or *mTOR* overexpression produces opposite effects on IFM muscle area in heat shocked wild type flies. (A-C) Sections of resin-embedded thoraces of flies of the indicated genotypes under the control of the *hs-Gal4* driver. The overexpression of *DIAP1* negatively affected muscle area whereas *mTOR* increased muscle area to values comparable to uninduced fly muscles (N/I) of control flies. (D) The quantification of mean muscle area confirmed that the differences were statistically significant. (E) The q-PCR analysis revealed that *mTOR* overexpression (*hs-Gal4*>*UAS-mTOR*) had no effect on autophagy-related gene expression, however, (F) Caspase 3/7 activity was dramatically increased. Statistically significant differences are denoted by asterisks (* P <0.05, *** P <0.001). Graphs show means \pm s.e.m.

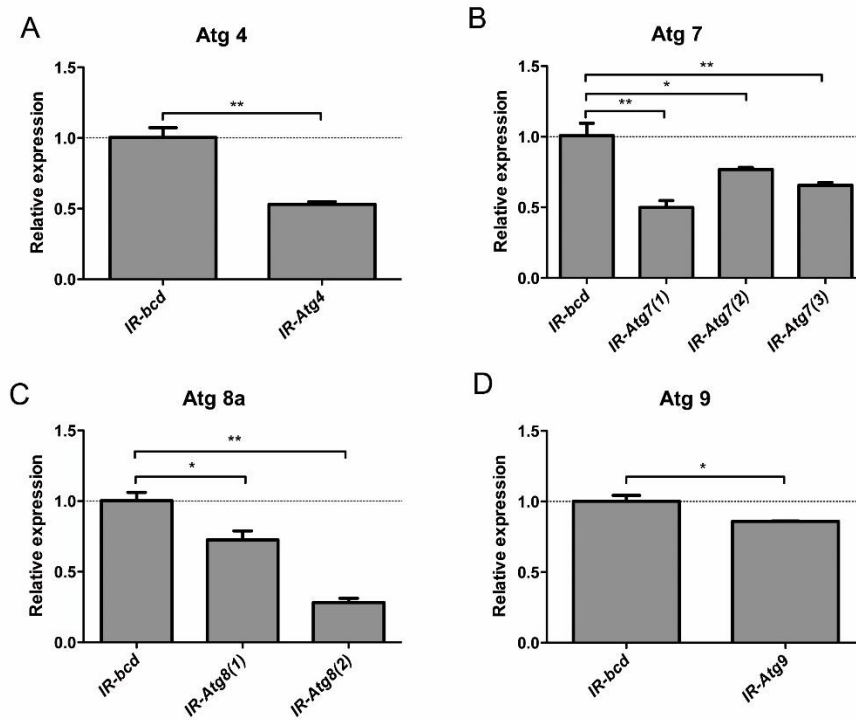


Figure S5. Quantification of relative expression levels of Atg genes upon expression of the indicated RNAi lines. Statistically significant differences are denoted by asterisks (* $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$). Error bars are standard deviations.

Table S1. Skeletal muscle biopsies from DM1 patients and unaffected controls.

Patients				Controls	
Gender	Age	CTG expansion (kb)	MIRS ⁽¹⁾	Gender	Age
Male	51	3	3	Male	58
Male	29	2	2	Male	48
Male	28	0.3	2	Male	31
Male	27	1.1	1	Male	31
Male	27	0.75	2	Male	24
Male	20	1	3	Male	25

(1) Muscular Impairment Rating Scale according to Mathieu, J. (1)

Table S2. Sequence and melting temperature (T_m) of PCR primers used for standard and quantitative PCR.

Primer pair	Forward (5'→3')	T _m (°C)	Reverse (5'→3')	T _m (°C)
<i>UAS</i>	GGAAAGTCCTTGGGGTCTTC	54	GGAAGTGAATGGGAGCA	52
<i>GAL4</i>	CACCGACGCTAATGATGTTG	52	TTTGTTTTCTGCCTCCACTG	54
<i>Atg4</i>	GCGCTCTTCGAGATCAGTCA	65	CCTGCCGCTCTCTTCAACTA	65
<i>Atg7</i>	CATAGCCTGTTTCAGCGGCCGT	71	CCGCTTGAATTCGGAGATTCCCCTG	70
<i>Atg8a</i>	ATCCAGACCGTGTGCCCGTCAT	74	ACCGACGGTCAGGTCGGAAGG	74
<i>Atg9</i>	CCACATCGAGGACCTTGACTC	60	CCACTACAATCACTGTGAATCCG	65
<i>Atg12</i>	TCGATGCCAGCGAGCAAATTTTCCT	70	GCCCCACGCCTGATTCTTGCA	72
<i>Rp49</i>	ATGACCATCCGCCAGCATAAC	55	ATGTGGCGGGTTCGCTTGTTT	55
<i>AKT1S1</i>	AGCCACAGAGACAGAGACC	58	CGTCCTCATCCATCACAAG	58
<i>AKT2</i>	CTCACACAGTCACCGAGAGC	58	TGGGTCTGGAAGGCATACTT	58
<i>ATG9A</i>	TTTGCTCAGATGGATGTTTCG	58	TCCTCAGCTTGCTGGTACTT	59
<i>BCL2</i>	CACCTGTGGTCCACCTGAC	58	CTGGACATCTCGGCGAAG	59
<i>BIRC7</i>	CCGGTCAAAGGAAGAGACTT	58	TGCGTCTCCGGTTCTTC	58
<i>GAPDH</i>	AGCCACATCGCTCAGACAC	58	CGCCAATACGACCAAT	58
<i>LAMP2</i>	AATGGCACAGTGAGCACAAA	59	GAGATGGCACAGTGGTGTGT	58
<i>mTOR</i>	TGCTGGAAGCCTTTGTCTATG	59	CGCTTGTTCCTTTGGTATT	59
<i>NKX3-2</i>	GGTGGGGTTTTCCCTGAG	59	GAAATTCTGAGGATTCAGGCTATG	59
<i>VPS52</i>	CGGCTCCGGGTCAAGG	62	CTTCTGGAGGATAAACTCTCGGAT	60
<i>VPS37</i>	CAGAAGCAAAGCTGGAGA	58	TCCACCTGCAGAAGGTCTAAC	59

1 Mathieu, J., Boivin, H., Meunier, D., Gaudreault, M. and Begin, P. (2001) Assessment of a disease-specific muscular impairment rating scale in myotonic dystrophy. *Neurology*, 56, 336-340.