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Increased autophagy accelerates colchicine-induced muscle toxicity

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Abbreviations: HMG-CoA, 3-hydroxy-3-methylglutaryl-coenzyme A; HMGCR, 3-hydroxy-3-methylglutaryl-CoA reductase; IBMPFD, inclusion body myopathy associated with Paget disease of the bone and fronto-temporal dementia; i.p., intraperitoneal; PBS, phosphate-buffered saline; TA, tibialis anterior

Colchicine treatment is associated with an autophagic vacuolar myopathy in human patients. The presumed mechanism of colchicine-induced myotoxicity is the destabilization of the microtubule system that leads to impaired autophagosome-lysosome fusion and the accumulation of autophagic vacuoles. Using the MTOR inhibitor rapamycin we augmented colchicine’s myotoxic effect by increasing the autophagic flux; this resulted in an acute myopathy with muscle necrosis. In contrast to myonecrosis induced by cardiotoxin, myonecrosis induced by a combination of rapamycin and colchicine was associated with accumulation of autophagic substrates such as LC3-II and SQSTM1; as a result, autophagic vacuoles accumulated in the center of myofibers, where LC3-positive autophagosomes failed to colocalize with the lysosomal protein marker LAMP2. A similar pattern of central LC3 accumulation and myonecrosis is seen in human patients with colchicine myopathy, many of whom have been treated with statins (HMGCR/HMG-CoA reductase inhibitors) in addition to colchicine. In mice, cotreatment with colchicine and simvastatin also led to muscle necrosis and LC3 accumulation, suggesting that, like rapamycin, simvastatin activates autophagy. Consistent with this, treatment of mice with four different statin medications enhanced autophagic flux in skeletal muscle in vivo. Polypharmacy is a known risk factor for toxic myopathies; our data suggest that some medication combinations may simultaneously activate upstream autophagy signaling pathways while inhibiting the degradation of these newly synthesized autophagosomes, resulting in myotoxicity.

Introduction

Most cases of toxic myopathy are the consequence of drugs commonly used for therapeutic interventions.1 Muscle tissue is particularly vulnerable to drug-associated toxicity because of its relative abundance, high blood flow, and metabolic rate. One example of a toxic myopathy is colchicine-induced myopathy.2 Colchicine is commonly used for the treatment of gouty arthritis, often at high initial doses. Colchicine myopathy is characterized by myalgias, muscle weakness, and occasional rhabdomyolysis.2 The mechanism of colchicine-associated muscle toxicity is poorly understood. Colchicine destabilizes microtubules, which are required for both endosomal and autophagosomal maturation,3 the resultant accumulation of endosomal and autophagic debris has been previously proposed as the pathogenic mechanism for colchicine myopathy.3

Macroautophagy (autophagy) is the bulk degradation of cytoplasmic proteins and organelles.3 This process includes multiple steps, many of which require intact microtubules: (1) phagophore initiation and expansion, (2) autophagosome formation, and (3) autophagosome fusion with acidic lysosomes.5 Skeletal muscle is sensitive to stimuli that activate autophagic degradation, such as nutrient deprivation and exercise.6,7 In addition, several drugs such as rapamycin activate autophagy in mouse skeletal muscle.8 If the accumulation of autophagic debris due to impaired microtubule trafficking is indeed the pathogenic mechanism of colchicine-induced myopathy, it would be reasonable to expect that inducing autophagy in the setting of colchicine treatment should exacerbate the disease pathology/progression.

Statins (HMGCR [3-hydroxy-3-methylglutaryl-CoA reductase] inhibitors) are among the most commonly prescribed medications for the reduction of cholesterol,9 but their use is limited by muscle toxicity. Statin use has been associated with muscle pain, exercise intolerance, muscle weakness, and rhabdomyolysis.10 Interestingly, the risk of statin-associated muscle toxicity increases with age and medication usage.10 In fact, many acute cases of muscle toxicity have been reported in patients concomitantly taking statins and colchicine.11-20 Statins reportedly induce...
autophagy in vitro, lending credence to our hypothesis that active autophagy is the potential driving force behind colchicine-induced muscle toxicity. To explore the impact of enhanced autophagy in colchicine-induced myopathy, we evaluated muscle pathology in mice treated with rapamycin or simvastatin in the setting of colchicine.

**Results**

**Rapamycin accelerates colchicine-induced muscle toxicity**

Previous studies have evaluated the toxicity of colchicine in a rodent model and found that four weeks of intraperitoneal (i.p.) colchicine at a dose of 0.4 mg/kg/d caused a vacuolar myopathy. Similarly, we found that i.p. colchicine at the same dose for 40 d caused myopathy with rare vacuoles, and the accumulation of autophagic markers LC3 and SQSTM1 in mouse skeletal muscle (Fig. 1; see Fig. S1 for isotype control IgG staining). In some cases, LC3 and SQSTM1 colocalized to the central region of a myofiber (Fig. 1E). To investigate whether autophagy induction accelerated colchicine-induced muscle toxicity, we treated 24 mice for 10 d with vehicle, i.p. colchicine (0.4 mg/kg/d), i.p. rapamycin (10 mg/kg/d), or i.p. colchicine and rapamycin combination; following treatment, we analyzed the tibialis anterior (TA) and quadriceps muscles by routine histochemistry. TA and quadriceps muscles from mice treated for 10 d with colchicine or rapamycin alone appeared similar to control mouse muscle when visualized via H&E or congo red, with no evidence of myopathic features (not shown). Therefore, in the case of colchicine, treatment periods longer than 10 d are necessary to induce the vacuolar changes seen in colchicine myopathy. In contrast, mice treated with 0.4 mg/kg/d i.p. colchicine and 10 mg/kg/d i.p. rapamycin for 10 d showed abnormal fibers with rare vacuolar structures, as evidenced by H&E, congo red, acid phosphatase, and succinate dehydrogenase staining (Fig. 2A–D). To confirm that the observed abnormal fibers were necrotic and that we had indeed accelerated the muscle toxicity induced by colchicine, we immunostained muscle sections with an antibody to complement c5b9, a marker of necrotic muscle fibers (Fig. 2E–H). The percentage of fibers that stained positive for c5b9 increased significantly in mice treated with both drugs compared with vehicle-, rapamycin-, or colchicine-treated groups (Fig. 2I). These data suggest that enhancing autophagy with rapamycin augments the pathogenesis of colchicine, leading to a necrotic autophagic myopathy.

**Colchicine plus rapamycin-treated mouse muscle accumulates autophagic structures**

Given that we enhanced autophagosome biogenesis with rapamycin and at the same time blocked autophagosome degradation with colchicine, we reasoned that the pathological changes seen in the skeletal muscle of colchicine plus rapamycin-treated mice may be due to the accumulation of nondegraded...
Figure 2. Rapamycin treatment augments pathology in a mouse colchicine myopathy model. Histochemical analysis of TA muscle from mice treated with colchicine (0.4 mg/kg/d) and rapamycin (10 mg/kg/d) for 10 d. (A and B) Congo red, (C) acid phosphatase, and (D) succinate dehydrogenase staining demonstrate a myopathy with scattered necrotic fibers (arrows). (E and H) TA muscle from (E) vehicle-, (F) rapamycin-, (G) colchicine- or (H) colchicine plus rapamycin-treated mice immunostained with an antibody to complement C5b9; arrows denote necrotic fibers. (I) Quantification of the percentage of C5b9-positive fibers per 20x field in muscle from mice treated with vehicle, rapamycin, colchicine or colchicine plus rapamycin for 10 d. *P < 0.05, the Student unpaired t-test between treatment and control. n = 5/condition. Scale bar: 50 µm.
autophagosomes. To test this hypothesis, we immunoblotted TA muscle lysates from vehicle-, rapamycin-, colchicine- or colchicine plus rapamycin-treated mice for the autophagic substrates SQSTM1 and LC3 (Fig. 3A). Upon phagophore formation, LC3-I can become lipidated and attach to the growing phagophore membrane as LC3-II.\textsuperscript{4} SQSTM1 and LC3-II levels were increased in colchicine-treated mice and to a greater extent in colchicine plus rapamycin-treated mouse muscle (Fig. 3A). Ubiquitinated proteins can be degraded via autophagy and are thus autophagic substrates.\textsuperscript{4} Consistent with this, the level of ubiquitinated proteins was elevated in colchicine plus rapamycin-treated mouse muscle (Fig. 3A). In addition, two other autophagic

![Figure 3. Autophagic substrates accumulate in the muscle of colchicine plus rapamycin-treated mice. (A) Immunoblot using antibodies to LC3, SQSTM1, ubiquitinated proteins, BNIP3, BECN1, and actin of TA muscle from 3–4 mo old mice treated i.p. with vehicle, 10 mg/kg/d rapamycin (rapa), 0.4 mg/kg/d colchicine (colch), or both drugs (colch+rapa) for 10 d. Blot is a representative of 3 independent experiments. (B) Quantitative PCR analysis of RNA from TA muscle obtained from 3–4 mo old mice treated i.p. with vehicle (veh) or colchicine+rapamycin (colc+rap) for 10 d. Data are represented as fold change as compared with vehicle treatment. \( P \) values were obtained from the Student unpaired \( t \)-test between treatment and control. \( n = 3/condition. \) (C) Immunoblot using antibodies described above in addition to RIPK1 kinase and GAPDH of TA muscle injected with vehicle or 100 \( \mu \)L of 12 \( \mu \)M cardiototoxin and harvested 2 d later. (D) TA muscle from vehicle-, rapamycin-, colchicine-, or colchicine plus rapamycin-treated mice immunostained with an antibody to LC3 or SQSTM1. DAPI-stained nuclei are blue; the sarcolemma of some fibers is outlined. Scale bar: 50 \( \mu \)m.]
proteins, BNIP3 and BECN1/Beclin 1, were increased in TA muscle lysates from mice treated with both colchicine and rapamycin compared with mice treated with vehicle, rapamycin alone, or colchicine alone (Fig. 3A). The accumulation of autophagic substrates such as LC3-II and SQSTM1 may not solely be caused by a decrease in their degradation but also as a result of enhanced transcription of the corresponding genes. To test this, we performed quantitative RT-PCR expression analysis using total RNA isolated from TA muscle obtained from vehicle-treated and colchicine plus rapamycin-treated mice (Fig. 3B). Consistent with enhanced autophagosome biogenesis, the expression of MAP1LC3A, MAP1LC3B, and SQSTM1 were increased by ~3–4 fold (Fig. 3B). In addition, the expression of the lysosomal/endosomal protein LAMP1 was increased by ~3 fold (Fig. 3B). In contrast, the expression of BNIP3 was not different between groups (Fig. 3B).

We do not think that the accumulation of autophagic substrates was a consequence of muscle necrosis, since muscle treated for two days with cardiotoxin showed a modest change in LC3-II levels and a decrease in SQSTM1 levels (Fig. 3C). Immunohistochemical staining of LC3, SQSTM1, and BNIP3 in cardiotoxin-treated muscle demonstrated that these autophagic markers were indeed altered in myofibers and not infiltrating cells (Fig. S2). Interestingly, BECN1 and BNIP3, as well as the necrotic signaling kinase RIPK1, were elevated after two days of cardiotoxin-induced muscle injury (Fig. 3C).

To define the localization of accumulated autophagic proteins in treated mouse muscle, we performed immunohistochemistry with antibodies to LC3, SQSTM1, and BNIP3; in addition, we used an antibody to the lysosomal marker, LAMP2. All autophagic markers were increased in colchicine plus rapamycin-treated mouse muscle when compared with muscle from the control, rapamycin-, and colchicine-treated mice (Fig. 3D and not shown). Scattered fibers showed an increase in SQSTM1-positive puncta in colchicine alone-treated muscle, but other autophagic markers were not increased above that seen with vehicle or rapamycin treatment (Fig. 3D). Interestingly, these autophagic proteins were localized to the central region of myofibers (Fig. 4) rather than throughout the sarcoplasm, as seen in cardiotoxin-treated muscle.
To further characterize this, we performed several immunofluorescence colocalization studies. As expected, LC3 and LAMP2 did not colocalize in colchicine plus rapamycin-treated mouse muscle, consistent with a failure in autophagosome-lysosome fusion (Fig. 5A). In contrast, SQSTM1 colocalized with LC3, suggesting that the loading of autophagosomes was not impaired (Fig. 5B). Finally, SQSTM1 and ubiquitinated proteins strongly colocalized, indicating that association of SQSTM1 with ubiquitinated substrates was not affected (Fig. 5C). Electron microscopy of similarly treated muscle demonstrated the accumulation of vacuolated structures, many of which had double membranes and contained cellular contents, consistent with autophagosomes. These structures were found within the central region of fibers between myofibrils (Fig. 5D and E).

Simvastatin-induced autophagy accelerates colchicine-induced muscle toxicity

We previously reported the pathological features of seven patients with colchicine-associated myopathy. Polypharmacy is a well-established risk factor for colchicine muscle toxicity, and commonly prescribed drugs, such as the cholesterol-lowering agents statins, are likely to play an important role. Indeed, two of seven patients in our original patient cohort (#22 and #25) and two of two patients diagnosed with colchicine myopathy since completion of that study (four of nine or 44% in total) were on both colchicine and a statin at the time of muscle biopsy. Immunohistochemical staining of these patients’ muscle biopsies showed a pattern of LC3 and SQSTM1 deposition in myofibers similar to that seen in colchicine plus rapamycin-treated mice (a representative biopsy is shown in Fig. 6; compare with a muscle sample from control patient with no muscle disease, Fig. S3). A detailed quantification of the percentage of LC3- and SQSTM1-immunopositive myofibers in these human specimens has been previously reported. We therefore reasoned that statin medications, like rapamycin, may accelerate and/or potentiate colchicine toxicity.

To test this, mice were divided into four treatment groups of six mice each: vehicle alone, 0.4 mg/kg/d colchicine, 10 mg/kg/d simvastatin, or both colchicine and simvastatin daily for 10 d. Following treatment, mice were sacrificed and TA and quadriceps muscles were analyzed via western blot, routine histopathology.
and immunohistochemistry. Similar to the muscle pathology seen in mice treated with colchicine plus rapamycin (Figs. 2 and 3) and human muscle pathology in patients treated with colchicine plus statin (Fig. 6), muscle from mice treated with colchicine plus simvastatin displayed myopathic features, with scattered necrotic fibers and LC3 accumulation within the center of myofibers (Fig. 7B and D). This pattern was not seen in the other treatment groups, including the group treated with simvastatin alone when compared with vehicle-treated mice. As we have previously demonstrated, there was accumulation of LC3-II in mice that received colchicine for 2 d, indicating a block in basal autophagy (Fig. 7A and C). To confirm necrosis and quantify the augmentation of colchicine toxicity by simvastatin, we immunostained tissue with an antibody to c5b9 and quantified the percentage of positive fibers (Fig. 7E and F). Similar to colchicine plus rapamycin, colchicine plus simvastatin treatment resulted in an increase in necrotic fibers when compared with vehicle-, colchicine- or simvastatin-treated mice (Fig. 7G). Immunoblotting for autophagic substrates in colchicine plus simvastatin-treated mouse muscle also demonstrated an increase in LC3-II, SQSTM1, and ubiquitinated proteins when compared with any of the control groups (Fig. 7H); these data raise the possibility that, like rapamycin, statins activate autophagy.

**Statins induce autophagy in mouse skeletal muscle**

Previous studies have demonstrated that statins can induce autophagy in cell culture models of skeletal muscle, but it has not been established whether this occurs in vivo in mature skeletal muscle. To evaluate this, we treated mice with four different statin drugs (simvastatin, atorvastatin, lovastatin, and pravastatin) for 7 d prior to isolating skeletal muscle and performing an immunoblot using an anti-LC3 antibody. To confirm that these drugs truly enhanced autophagic flux, we treated similar mice with 0.4 mg/kg colchicine on days 6 and 7 and then harvested TA muscles for immunoblot. LC3-II levels were unchanged in muscle from mice treated with statins alone when compared with muscle from vehicle-treated mice. As we have previously demonstrated, there was accumulation of LC3-II in mice that received colchicine for 2 d, indicating a block in basal autophagy (Fig. 8A and B). Consistent with an increase in autophagic flux, skeletal muscle lysates from mice receiving a statin for 7 d and colchicine on days 6 and 7 displayed increased accumulation of LC3-II when compared with the colchicine group, suggesting that statins enhance autophagic flux in vivo.

The myotoxic effect of colchicine is well known, but it has been difficult to pinpoint its pathogenic mechanism. The bulk of the relevant mechanisms proposed so far to explain colchicine-induced muscle toxicity hinge on the disruption of the microtubule network. In this study, we report data which support an expansion of the paradigm regarding the mechanism of colchicine-induced muscle toxicity from mere disruption of the microtubule network to a functional and/or synergistic relationship between colchicine and autophagy induction. We have shown that mice treated for 10 d with both colchicine and autophagic inducers such as rapamycin or simvastatin display (1) increased accumulation of autophagic substrates and ubiquitinated proteins; and (2) worsening of the colchicine-induced muscle pathology when compared with mice injected with colchicine alone. Thus, the emerging hypothesis is that colchicine-induced muscle toxicity is a result of autophagic vacuole build-up due to enhanced autophagosome formation and impaired autophagic degradation; this possibility is supported by the fact that autophagic markers LC3 and SQSTM1 robustly accumulate in the muscle of patients with colchicine-induced autophagic vacuolar myopathy.

Colchicine and chloroquine have both been associated with toxic vacuolar myopathies. In the case of chloroquine, lysosomal de-acidification is the presumed mechanism by which autophagic vacuoles accumulate and fail to be degraded. Previous
rodent models of chloroquine-induced myopathy have treated mice for several weeks with chloroquine to induce pathology. Interestingly, denervation of muscle hastens the development of an autophagic myopathy resulting in pathology over the course of days rather than weeks. Denervation is a potent inducer of muscle atrophy and autophagy, an increase in autophagy in the setting of impaired autophagic degradation may explain the synergism between chloroquine treatment and denervation in experimental models of chloroquine myopathy. Many vacuolar myopathies are due to mutations in proteins associated with lysosomal function and accumulate autophagic proteins within the myofiber similarly to the pathology seen in colchicine-associated myopathy. For example, Danon disease, Pompe disease, and X-linked myopathy with excessive autophagy all accumulate autophagic debris. Whether the activation of autophagy is beneficial or detrimental for these diseases is not known. We recently demonstrated that enhancing autophagy with rapamycin worsened the vacuolar pathology, protein aggregate accumulation and weakness in a mouse model of inclusion body myopathy associated with Paget disease of the bone and fronto-temporal dementia (IBMPFD). IBMPFD is caused by mutations in valosin containing protein, which is required for maturation of autophagosomes to autolysosomes. Interestingly, in the case of IBMPFD, X-linked myopathy with excessive autophagy and sporadic inclusion body myositis, MTORC1 activity is diminished, suggesting enhanced autophagosome formation.
current study lends further support to a mechanism of disease in which activation of autophagy drives the vacuolar pathology in some myopathies.

Statins initiate autophagy in cultured cells.\(^{21,33}\) The mechanism by which statins induce autophagy likely relates to their mechanism of action within the HMGCR pathway. Specifically, depletion of geranylgeranyl diphosphate by statins and bisphosphonates can induce autophagy.\(^{21,34}\) Moreover, farnesyl transferase inhibitors and geranylgeranyl transferase inhibitors both induce autophagy in vitro.\(^{35,36}\) Our studies confirm these cell culture models and demonstrate that statins can effectively enhance autophagic flux in vivo in mouse skeletal muscle. In this context, it is worth noting that UCSF muscle pathology practice has seen a sharp increase in the number of colchicine and hydroxychloroquine myopathy cases in the last 5–6 y, coinciding with a large increase in statin use (based on a CDC survey entitled “Health, United States, 2010,” the percentage of statin-using adults \(\geq\) 45 y of age increased from 2% in the period of 1988–1994 to 25% in the period 2005–2008, and is likely even higher today).

Risk factors for statin muscle toxicity are higher dosages, concomitant use of certain drugs that can inhibit statin metabolism, and more recently polymorphisms in the gene \(\text{SLCO1B1/OATP1B1}\) that encodes solute carrier organic anion transporter family, member 1B1, and mediates the uptake of various drugs including statins.\(^{37,38}\) It is conceivable that the accelerated muscle pathology seen in our studies is due to increased statin drug concentration in the setting of colchicine administration. However, based on the findings of our study, we propose that mutations or polymorphisms in genes within the autophagy pathway may also be risk factors for statin myotoxicity.

**Materials and Methods**

**Antibodies and reagents for mouse studies**

Antibodies for BECN1 (Cell Signaling Technologies, 3738), GAPDH (Cell Signaling Technologies, 2118), LAMP2 (Santa Cruz Biotechnology, sc-18822), BNIP3 (Abcam, ab10433), Actin (Sigma, A2066), LC3B (Sigma, L7543) for western blotting, LC3 (Nanotools, 0231-100/LC3-5F10) for immunostaining, RIPK1/RIP1 (BD Transduction Laboratories, 610458), c5b9 (Dako, M0777), and antibodies to SQSTM1 (Proteintech, 18420-1-AP) for western blotting, and to SQSTM1 (Sigma, P0067) for immunostaining were obtained from the indicated sources. For the autophagic assessment of necrotic fibers in Figure S2, the VECTOR M.O.M. immunodetection Kit Flourescein (Vector Laboratories, FMK-2001) was used to quench endogenous background IgG. Rapamycin (LC Laboratories R-5000) and cardiotoxin (Sigma, C9759) were also purchased. Simvastatin (1965), atorvastatin (3776), lovastatin (1530), and pravastatin (2318) were obtained from TOCRIS.

**Animals and drug treatments**

All experiments were conducted using 3–4 mo old C57BL/6 mice from Jackson Laboratory. Animal studies were approved by the Animal Studies Committee of Washington University School of Medicine. Upon arrival, mice were housed in a temperature-controlled environment with 12 h light/dark cycles, where they...
received food and water ad libitum. All drugs were injected intraperitoneally. Rapamycin was dissolved in 100% ethanol at 62.5 mg/ml, stored at -20 °C, and diluted to 2.5 mg/ml in vehicle (5% PEG 400, 5% Tween 80, and 4% ethanol) just before injection. Colchicine was dissolved in water, stored at -20 °C as a stock solution at a concentration of 4 mg/ml, and diluted to 0.1 mg/ml before injection. Simvastatin, atorvastatin, lovastatin, and pravastatin were administered at a dose of 10 mg/kg body weight. Simvastatin and lovastatin were dissolved in 100% ethanol and stored at -20 °C as stocks; on the day of experiment, they were diluted with PBS immediately prior to injection. Atorvastatin was dissolved in DMSO and stored as stock solution, which was diluted with PBS immediately prior to injection. Pravastatin was dissolved in water. For all drug treatments, control mice received equal volumes of respective vehicles. Mice were subjected to i.p. injection of both rapamycin (10 mg/kg body weight) and colchicine (0.4 mg/kg body weight) daily for 10 d. Age-match littermates were injected with vehicle (5% Tween 80, 5% polyethylene glycol, and 4% ethanol), rapamycin alone, or colchicine alone as controls. Similar experiments were performed with simvastatin (10 mg/kg body weight) in place of rapamycin. Cardiotoxin (from Naja mossambica mossambica) was dissolved in water and stored at -20 °C as stock solution, which was diluted in PBS to a working concentration of 12 μM prior to experiment. For each mouse, tibialis anterior muscles were injected with 100 μL of 12 μM cardiotoxin and muscles were harvested 2 d later.

Western blotting

Muscle tissues were homogenized using RIPA lysis buffer (50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 1% NP-40, 0.25% Na-deoxycholate, 1 mM EDTA) supplemented with protease inhibitor cocktail (Sigma-Aldrich, s8820-20TAB) and then centrifuged at 14,000 g for 10 min. Protein concentrations were determined using BCA protein assay kit (Thermo Fisher Scientific, 23225). Aliquots of lysates were solubilized in Laemmli sample buffer and equal amounts of proteins were separated by 12% SDS-PAGE. Proteins were transferred to nitrocellulose membrane and the membrane was blocked with 5% nonfat dry milk in phosphate-buffered saline (PBS) for 1 h. The membrane was then incubated with primary antibodies in 5% nonfat dry milk overnight at 4 °C; after incubation with the appropriate secondary antibody conjugated with horseradish peroxidase, enhanced chemiluminescence (Immunopure, 31430) was used for protein detection. If necessary, blots were stripped using Re-blot strong solution (Millipore, 2504) and reprobed with the appropriate antibodies. Autoradiographs of immunoblots were scanned using an Epson 636 Expression scanner; immunoblots were obtained using the G:BoxChemi XT4, Genesys Version 1.1.2.0 (Syngene).

Gene expression analyses

Total RNA was isolated from TA muscle with SV Total RNA isolation kit (Promega, Z3100) according to the manufacturer’s instructions. The concentration and quality of the total RNA isolated was determined using a Nanodrop spectrophotometer. cDNA was synthesized using the RT1 First Strand kit (Qiagen, 330401). Gene expression levels were analyzed by real-time PCR on an Applied Biosystems model 7500 (software v2.0.5) using the RT1 SYBER Green ROX qPCR mastermix (Qiagen, 330522) and the mouse autophagy RT1 Profiler PCR array (Qiagen, 330231 PAMM-084ZA), which contains primers for the expression of 84 autophagy genes arranged in a staggered format. Data were normalized to the average of 4 housekeeping genes, Gapdh, Actb/β-actin, B2m/β2-microglobulin, and Gusb/β-glucuronidase and expressed in terms of $2^{-ΔΔCt}$.

Histochemistry, immunohistochemistry, and electron microscopy

Isolated muscle was mounted using tragacanth gum (Sigma, G1128) and quick frozen in liquid nitrogen-cooled 2-methylbutane. Samples were stored at -80 °C until sectioning into 10-μm sections. H&E, congo red, acid phosphatase, and succinate dehydrogenase staining was performed as previously described. Briefly, muscle sections were affixed to slides incubated for 10 min in ice-cold acetone, mounted with Prolong Gold + DAPI (Invitrogen, P36931), and examined using a fluorescence microscope (Nikon 801 upright+ and Roper Scientific EZ monochrome CCD camera with deconvolution software analysis [NIS Elements, Nikon]). Nonfluorescent images were taken with a 5 megapixel color CCD (Nikon). For electron microscopy, TA muscle was harvested, rinsed briefly in PBS and then placed immediately in Karnovsky fixative at 4 °C for 24 h. Fixed muscle was embedded in plastic and sectioned for EM imaging. Images were taken using a Jeol 1400 and a Gatan Orius 832 Digital Camera. Image processing and analysis were done with (NIS Elements 4.0) software and Adobe Photoshop.

Quantification of necrotic fibers

The percentage of c5b9-positive fibers was quantified per 20× field. Twenty randomly selected microscopy fields from 10 TA muscles from 5 mice/condition were selected and the total number of myofibers and the number of immunoreactive fibers were counted and expressed as a percentage.

Human subjects

Study design was reviewed and approved by the University of California San Francisco Committee on Human Research; the informed consent requirement was waived given the noninvasive nature of the study and a minimal potential for harm to study participants. Immunoperoxidase staining for LC3 (mouse monoclonal antibody, clone 5F10, Nanotools, 0231-100) and SQSTM1 (guinea pig polyclonal antibody, Progen Biotechnik, GP62) was performed on formalin-fixed, paraffin-embedded tissue samples using the Ventana Benchmark XT automated slide preparation system as described previously. Images were taken with a DP72 digital camera on a BX41 bright-field light microscope using cellSens Entry 1.6 software (all by Olympus) and were edited with Adobe Photoshop CS5 Version 12.1.32.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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