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Influenza A Virus Infection Induces Muscle Wasting via IL-6 Regulation of the E3 Ubiquitin Ligase Atrogin-1

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Abstract

Muscle dysfunction is common in patients with ARDS and is associated with morbidity that can persist for years after discharge. In a mouse model of severe influenza A pneumonia, we found the pro-inflammatory cytokine IL-6 was necessary for the development of muscle dysfunction. Treatment with an FDA approved antibody antagonist to the IL-6 receptor (tocilizumab) attenuated the severity of influenza A-induced muscle dysfunction. In cultured myotubes, IL-6 promoted muscle degradation via JAK/STAT, FOXO3a and atrogin-1 upregulation. Consistent with these findings, *atrogin-1*^{+/-} and *atrogin-1*^{-/-} mice had attenuated muscle dysfunction following influenza infection. Our data suggest that inflammatory endocrine signals originating from the injured lung activate signaling pathways in the muscle that induce dysfunction. Inhibiting these pathways may limit morbidity in patients with influenza A pneumonia and ARDS.

Introduction

Influenza A pneumonia is the most common cause of morbidity from an infectious agent responsible for at least 20,000 to 50,000 deaths in the United States yearly (1, 2). The lung epithelium is the primary target of the influenza A virus (IAV) infection, and the resulting diffuse damage to the epithelium can impair gas exchange resulting in severe pneumonia and development of the Acute Respiratory Distress Syndrome (ARDS). About 30–40% of patients with ARDS die, and most survivors often suffer multiple morbidities related to their prolonged critical illness (3, 4). Muscle weakness is evident in up to 50% of ARDS survivors, where it is associated with prolonged lengths of stay in the intensive care unit and

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the development of the multiple organ dysfunction syndrome (5–7). In patients who survive ARDS, muscle weakness results in substantial morbidity and long-term reductions in quality of life (8–11).

As survival from ARDS continues to improve (12, 13), it is increasingly important to prevent muscle dysfunction and other long-term sequelae of the syndrome (14). Our understanding of the pathophysiology of muscle weakness associated with critical illness is incomplete. Some investigators have argued that skeletal muscle dysfunction in ARDS is largely attributable to the prolonged immobility associated with critical illness (14, 15). Others have shown that the ubiquitin proteasome system is activated in skeletal muscle within hours or days of the onset of critical illness, suggesting a role for active signaling processes that promote muscle dysfunction (7, 16).

Maintenance of muscle mass requires an equilibrium between the synthesis and breakdown of myofiber proteins. Muscle atrophy occurs when there is a net loss of muscle mass causing shrinkage of the myofibers. There are several different types of skeletal muscle atrophy including, sarcopenia, disuse atrophy and cachexia (9, 10, 17). The FoxO (forkhead box O) class of transcription factors have been implicated in the control of muscle degradation (18, 19). Specifically, FoxO1 and FoxO3 upregulate the muscle-specific ubiquitin ligases: MAFbx/atrogen-1 (muscle atrophy F-box protein) and MuRF1 (muscle RING finger-1), which target muscle proteins for degradation by the ubiquitin proteasome system (20, 21). The pro-inflammatory cytokine IL-6 has been reported to regulate atrogen-1 and thus promote muscle atrophy (22–24). Cellular signaling induced by IL-6 has been shown to activate the Janus kinases (JAK)/signal transducer and activator of transcription 3 (STAT3) (25). Phosphorylation of STAT3 leads to the translocation of (FoxO)3 to the nucleus leading to upregulation of atrogen-1 (19, 26).

In animal models and healthy volunteers infected with influenza A virus, IL-6 and TNF- α are rapidly detected in bronchoalveolar lavage fluid and nasal washings (27, 28). IL-6 is also increased in the serum of patients with lung injury (29, 30), and in mouse models of influenza infection (28). Increased levels of IL-6 and other inflammatory cytokines are associated with prolonged mechanical ventilation (29, 30) and increased mortality (29–31). In this study, we sought to determine whether IL-6-mediated activation of STAT3/atrogen-1 was necessary for the development of muscle dysfunction in a murine model of influenza A infection.

Materials and Methods

Reagents

All cell culture reagents were from Corning Life Sciences (Tewksbury, MA). All other chemicals were purchased from Sigma-Aldrich (St. Louis, MO).

Mice

Adult male (12–16 weeks) C57Bl/6 wild-type were obtained from The Jackson Laboratories (Bar Harbor, ME). Age-matched male *atrogen-1*^{-/-} and *atrogen-1*^{+/-} mice as well as wild-type littermates (*atrogen-1*^{+/+}) on a 129S/C57Bl/6 background have been described

previously and were obtained from Regeneron Pharmaceuticals and Stewart Lecker at Beth Israel Deaconess Medical Center and Harvard Medical School (20). Mice were provided with food and water *ad libitum*, maintained on a 14-h light/10-h dark cycle, and handled according to National Institutes of Health and Northwestern University Institutional Animal Care and Use Committee guidelines. Northwestern University Institutional Animal Care and Use Committee approved all animal experimental protocols.

Influenza A Virus and Infection

Influenza A virus (A/WSN/33 [H1N1]), a murine adapted virus, was provided by Robert Lamb, Ph.D., Sc.D., Northwestern University, Evanston, IL. Mice were anesthetized with isoflurane, their lungs were intubated with a 20-gauge angiocath (32, 33) and two 25 μ l sterile aliquots of Phosphate Buffered Saline ((PBS) Control) or influenza A virus (A/WSN/33 [H1N1]) (1500 or 150 pfu/mouse) were instilled through the catheter as we have previously described (28). We continuously observed mice infected with influenza A virus for signs of distress (slowed respiration, failure to respond to cage tapping, failure of grooming, huddling and fur ruffling). Mice that developed these symptoms were sacrificed and the death was recorded as an influenza A-induced mortality. Weight was measured at infection time and prior to harvest 7 days post-infection (dpi) and recorded as percentage of weight loss from baseline.

Administration of tocilizumab

Tocilizumab is a humanized monoclonal antibody against the interleukin-6 (IL-6) receptor; for our studies, we used a murine adapted form of tocilizumab (Genentech, South San Francisco, CA). Briefly, adult male C57Bl/6 mice were anesthetized with isoflurane and were administered tocilizumab 8mg/kg in a total volume of 150 μ l of PBS or PBS alone by a retro-orbital injection. After 24 hours, the mice were infected with IAV or PBS as described above.

Measurement of muscle dysfunction

Immediately prior to muscle harvest, forelimb skeletal muscle strength was assessed using a digital grip strength meter (Columbus Instruments, Columbus, OH) as described (34, 35). Grip strength was measured in each animal six successive times, and the average of the highest four values for each mouse was used. The same operator performed all tests. The mice were then terminally anesthetized with Euthasol (pentobarbital sodium/phenytoin sodium). The soleus and extensor digitorum longus (EDL) muscles were excised and tendon was trimmed under a microscope to assure optimal accuracy for weight measurement. The muscles were then blotted dry and weighed. Muscles were either frozen in liquid nitrogen-cooled isopentane for cryosectioning or snap-frozen in liquid nitrogen for protein extraction.

Immunohistochemistry and Fiber Size Assessment

Soleus and EDL serial transverse cryosections (8 μ m) were obtained from the Northwestern University Mouse Histology and Phenotyping Laboratory and mounted on glass slides. Sections were fixed in 4% formaldehyde, permeabilized, and blocked. Immunostaining was performed with laminin primary antibody (Sigma; Catalog: L9393; 1:50) followed by Alexa

Fluor 568-conjugated secondary antibody (Life Technologies, Carlsbad, CA; Catalog: A-11011; 1:200). Images were acquired with a Zeiss LSM 510 confocal microscope using a 40x objective (Northwestern University Center for Advanced Microscopy) and analyzed using Fiji (National Institutes of Health) (36). Fiber size and cross-sectional area (CSA) were determined as previously described (34). All analysis was performed by one blinded-operator.

Cell Culture, siRNA Transfection, Drug Treatment, Myotube Analysis and Viral Plaque Assay

C2C12 mouse myoblasts (ATCC, Manassas, VA: CRL-1772) were cultured and differentiated as we have previously have described (34). For experimental procedures, myoblasts were grown to 90–95% confluence and then media was switched to 2% horse serum DMEM (differentiation media) and renewed daily for optimal myotube formation. For siRNA transfection experiments, 90–95% confluent C2C12 myoblasts were switched to differentiation media (Day 0). On Day 1 of differentiation, the media was replaced with antibiotic-free differentiation media and transfection protocol was performed 12 hours later. On day 2 of differentiation, cells were transfected with 60 pmol of either mouse *STAT3*, *FKHRL1*, *FKHR*, or control siRNA (Scramb) pools (Santa Cruz Biotechnology, Dallas, TX) using Lipofectamine RNAiMax Transfection reagent (Thermo Fisher Scientific, Waltham, MA) according to the manufacturer's instructions. On day 4 of differentiation, cells were treated with 10 ng/ml of recombinant mouse IL-6 (BD Biosciences, San Jose, CA) for 16 hours. For the dose-response treatment of IL-6, cells were treated on day 4 of differentiation with various concentrations IL-6 (2.5, 5 and 10 ng/ml) for 16 hours. For experiments with tocilizumab, cells were treated with 10 µg/ml on Day 3 for 24 hours followed by treatment on Day 4 of differentiation with 10 ng/ml of recombinant mouse IL-6 for 16 hours. C2C12 myotubes were treated on day 4 with 2 MOI of influenza A virus (A/WSN/33 [H1N1]) for 16 hours. On day 5 of differentiation, cells were imaged and then cell lysates were obtained for Western blot analysis. Images were acquired using a Nikon Eclipse TE2000-U inverted scope with a 40X objective. Myotube diameters were quantified by one blinded-operator measuring ~375–679 tube diameters (5–10 measurements per fiber) from five random fields from a minimum of 3 independent experiments using MetaMorph Software (Molecular Devices, Sunnyvale, CA; Version 6.3) as previously described (34). Madin-Darby Canine Kidney (MDCK) cells (ATCC, CCL-34) cells were maintained as described previously and used to perform a viral plaque assay with homogenized soleus muscle or lungs from mice infected with influenza (37).

Western Blot Analysis

C2C12 myotubes were washed with ice-cold PBS and then lysed with 2X Cell Lysis Buffer (Cell Signaling Technology, Danvers, MA). Lysates were centrifuged at 20,000 x g and the supernatant was collected. Soleus and EDL muscles were lysed with ice-cold lysis buffer (Nonidet P-40 1%, glycerol 10%, NaCl 137 mM, Tris-HCl pH 7.5 20 mM) containing protease (1 Complete Mini, EDTA-free tablet, Roche) and phosphatase (sodium fluoride 30 mM, β-glycerophosphate 250 mM, sodium orthovanadate 1mM) inhibitors (38). Tissue was then homogenized with a Tissue Tearor (BioSpec Products, Inc, Bartlesville, OK) for 1 minute. Samples were centrifuged at 15,000 rpm for 10 minutes at 4° C and the supernatant

was collected. Protein concentrations were determined with Protein Assay Dye (Bio-Rad, Hercules, CA) using the Bradford method (39). Equal amount of protein were loaded on a SDS-PAGE and Western blot analysis was performed as described previously (40). Incubation with primary antibodies was performed overnight at 4°C. Immunoblots were quantified by densitometry using Image J 1.46r (National Institutes of Health) or Image Studio Software (LI-COR Inc., Lincoln, NE) (41). The following antibodies were used: Rabbit Monoclonal to Fbx32 (Abcam, Cambridge, United Kingdom; catalog: ab168372; 1:1000), Rabbit polyclonal to MuRF1 (ECM Biosciences, Versailles, KY; catalog: MP3401; 1:1000), Rabbit monoclonal to GAPDH (D16H11) (Cell Signaling; catalog: 5174s; 1:1000), Rabbit monoclonal to STAT3 (79D7) (Cell Signaling; catalog: 4904s; 1:1000), Rabbit monoclonal to FoxO3a (75D8) (Cell Signaling; catalog: 2497s; 1:1000), Rabbit polyclonal to FoxO1 (L27) (Cell Signaling; catalog: 9454s; 1:1000). Primary antibodies were detected with horseradish peroxidase-conjugated secondary antibodies including Anti-Rabbit IgG, HRP-linked (Cell Signaling; catalog: 7074s; 1:2000 dilution) and Goat-Anti-Rabbit IgG-HRP conjugate (Bio-Rad; catalog: 170-6515; 1:16,000 dilution).

Bronchoalveolar lavage (BAL) fluid and blood collection for measurement of cell count, protein and IL-6.

Animals were euthanized with Euthazol (pentobarbital sodium/phenytoin sodium) prior to BAL and blood collection. A mid-line dissection through the abdominal cavity up to the thoracic cavity was performed followed by a small incision at the edge of the diaphragm, which resulted in a fatal pneumothorax. Then, a right ventricular puncture for the collection of blood was performed. Blood was collected in a microvette microtube with EDTA (Kent Scientific) and kept on ice and then spun at 2000 rpm. Serum was collected and used for IL-6 determination by ELISA (Thermo Fisher, KMC0062) (37). Bronchoalveolar lavage (BAL) fluid was obtained through a 20-gauge angiocath ligated into the trachea through a tracheostomy. A total of 1-ml of PBS instilled into the lungs was aspirated three times to collect BAL fluid for cell counts (Cellometer K2; Nexcelom Bioscience), protein quantification (Bradford assay) and IL-6 determination by ELISA.

Statistics

Results are shown as mean \pm SEM. Data were analyzed and statistical differences were compared by a Student's *t*-test between two groups and by a one-way ANOVA with Dunnett or Tukey's post-hoc corrections for three or more groups using GraphPad Prism version 7.0 for Windows (GraphPad Software, La Jolla, CA). All *in vitro* experiments were repeated a minimum of three times. We considered specific data points outliers if they deviated by more than two times the standard deviation from the mean and they were excluded automatically from the statistical analysis. Results were considered significant when *p* value < 0.05.

Results

C57Bl/6 wild-type mice lose muscle weight and strength after influenza infection.

We infected C57Bl/6 wild-type mice with either a low dose (150 pfu/mouse) or high dose (1500 pfu/mouse) of influenza A virus (A/WSN/33 [H1N1]) (IAV) and assessed total body

weight, forelimb grip strength, soleus and EDL muscle wet weight, and fiber cross-sectional area (CSA). Compared to control (PBS-treated) mice, IAV-infected mice experienced significant weight loss after both low dose and high dose IAV infection (Figure 1A). Forelimb grip strength was significantly decreased in the IAV-infected mice (Figure 1B). In addition, mice-infected with either low or high dose influenza had significantly decreased soleus and EDL wet muscle weight (Figure 1C, 1D). Figure 1E and 1H show representative images of soleus and EDL laminin-stained cryosections that were used to determine the mean CSA. Mice infected with high dose IAV had significantly decreased mean soleus and EDL CSA (Figure 1F, 1I) with the histograms showing a left shift of frequency distribution indicating a predominance of thinner fibers in IAV-infected mice (Figure 1G, 1J).

To determine that the IAV infection was restricted to the lung, we measured by plaque assay the viral load in lungs and soleus muscle of mice infected with IAV for 7 days. As expected the viral load was almost undetectable in soleus muscle homogenates compared with lung homogenates of wild-type IAV-infected mice (soleus: 38.33 pfu/ml \pm 6.009 S.E.M.; lung: 1.25×10^7 pfu/ml \pm 8.3660×10^5 S.E.M.). A Student's *t*-test between two groups was used to show statistical differences with a $p < 0.0001$ ($n = 3-5$ mice/group).

Influenza-induced muscle dysfunction is attenuated in mice with genetic loss of atrogen-1.

The ubiquitin-proteasome system regulates muscle degradation via the E3 ubiquitin ligases, MAFbx/atrogen-1 and MuRF1 (20, 21). We set out to determine if either or both of these muscle specific-E3 ubiquitin ligases were up-regulated in our murine model of IAV infection. C57Bl/6 wild-type mice were infected with IAV (1500 pfu/mouse) and 7 days dpi the mice were sacrificed and the soleus muscles were excised. We found that atrogen-1 protein expression was significantly increased while MuRF1 levels remained unchanged (Figure 2A, 2B). To further explore the role of atrogen-1 in mediating muscle protein loss with influenza, we used mice with heterozygous or homozygous deletion of the *MAFBx/atrogen-1* gene (designated as *atrogen-1* throughout the paper) (20). Compared with their wild-type (*atrogen-1^{+/+}*) littermates, levels of atrogen-1 were significantly lower in the soleus muscle of *atrogen-1^{+/-}* and *atrogen-1^{-/-}* mice (Figure 2C). Phenotypic differences in the animals were also evident. Basally, *atrogen-1^{+/-}* had slightly reduced grip strength but was not statically different as compared to the PBS-treated *atrogen-1^{+/+}*. *Atrogen-1^{-/-}* mice baseline grip strengths were markedly decreased, but there was no difference between control (PBS-treated) and IAV-treated mice, suggesting that lower levels of atrogen-1 are protective during influenza infection (Figure 2D). While *atrogen-1^{+/+}* had significantly decreased wet muscle weight for soleus and EDL after IAV infection, wet muscle weight after influenza A infection was similar to uninfected mice in the *atrogen-1^{-/-}* mice (Figure 2E-F). The haploinsufficient *atrogen-1^{+/-}* mice showed levels of wet muscle weight loss intermediate between *atrogen-1^{+/+}* and *atrogen-1^{-/-}* (Figure 2E-F). The mean fiber CSA for the soleus and EDL muscles were decreased in the *atrogen-1^{+/+}* IAV-infected mice as compared to control (PBS-treated) mice, but the IAV-induced decrease in CSA was attenuated *atrogen-1^{+/-}* and *atrogen-1^{-/-}* mice (Figure 2G-P). Each of the *atrogen-1* genotypes infected with IAV lost body weight as compared to the PBS-treated mice, but the *atrogen-1^{+/-}* and *atrogen-1^{-/-}* weight loss was slightly less severe (Figure 3A).

Additionally, IL-6 levels were measured in serum and bronchoalveolar lavage (BAL) fluid. Total protein and cell count were also determined in the BAL fluid. IL-6 levels were markedly increased in the IAV-infected *atrogen-1^{-/-}* in both serum and BAL fluid as compared to IAV-infected *atrogen-1^{+/-}* and *atrogen-1^{-/-}* mice (3B and 3C). There was no difference amongst the different *atrogen-1* genotypes infected with IAV in total protein and cell count of the BAL fluid (3D and 3E).

The IL-6-receptor antibody tocilizumab attenuates influenza-induced muscle dysfunction.

To investigate the role of IL-6 in mediating influenza A-induced muscle atrophy *in vivo*, we treated C57Bl/6 mice with tocilizumab, which is a monoclonal antibody with specificity for the IL-6 receptor that prevents IL-6 from binding to both the soluble and membrane-bound IL-6 receptor through competitive blockade (42). C57Bl/6 mice were administered tocilizumab 8mg/kg or PBS (control) 24 hours by retro-orbital injection prior to infection with IAV (1500 pfu/mouse) or PBS and then 7 dpi weight loss and grip strength were assessed. There was no difference in total body IAV-induced weight loss between the PBS- and tocilizumab-treated mice (Figure 4A). The reduction of grip strength observed in IAV-infected mice treated with vehicle was attenuated in the IAV-infected and tocilizumab treated mice (Figure 4B). Consistent with these findings, soleus and EDL wet muscle weights were significantly reduced in influenza-infected compared with uninfected wild-type mice, while there was no significant decrease in either the soleus or EDL wet muscle weight in uninfected compared with tocilizumab treated mice (Figure 4C, 4D). The influenza A-induced induction of atrogen-1 protein was attenuated in the soleus muscle of tocilizumab-treated mice (Figure 4D). Similarly, the influenza A-induced reductions in soleus and EDL CSA were attenuated in mice that had been pre-treated with tocilizumab compared with vehicle as was the left-shift of muscle fiber size frequency distribution for both the soleus and EDL (Figure 4F-M).

IL-6 acts via STAT3, FoxO3a, and Atrogen-1 to induce skeletal muscle atrophy

To obtain insight into the mechanisms by which IL-6 induces muscle atrophy, we treated differentiated C2C12 myotubes with various doses of recombinant IL-6 (2.5, 5 and 10 ng/ml) or with 1 MOI of IAV or media alone, and blinded measurements of myotube diameters were performed 16 hours later. Treatment of C2C12 cells with IL-6 resulted in a dose-dependent reduction in myotube diameter, while treatment with IAV had no effect on fiber diameter (Figure 5A). To determine if STAT3 was involved in the IL-6-induced reduction in myotube diameter, we transfected well-differentiated C2C12 cells with scrambled siRNA or siRNA specific for *STAT3* followed by treatment with recombinant IL-6 and then measured myotube diameters. Silencing of STAT3 prevented the decrease in diameter seen with IL-6 treatment (Figure 5B). To determine whether STAT3 acts through FoxO1 or FoxO3a to reduce myotube diameter, we compared the effect of IL-6 on myotube diameter in control, FoxO1 and FoxO3a silenced cells. Silencing of FoxO3a attenuated the decline in myotube diameter induced by IL-6 (Figure 5C), but silencing of FoxO1 did not prevent the decrease in myotube diameter (Figure 5D). Consistent with our *in vivo* findings, pretreatment of myotubes with tocilizumab prevented the IL-6-induced decrease in myotube diameter as compared to the myotubes that received the mock treatment and IL-6 (Figure 5E). Moreover, treatment of myotubes with IL-6 increased the expression of atrogen-1 in

vehicle treated cells and this effect was attenuated in cells pretreated with tocilizumab (Figure 5F).

Discussion

Severe infection with the influenza A virus is an important cause of lung injury and ARDS (1, 2). A significant proportion of patients with ARDS develop skeletal muscle weakness that is associated with prolonged duration of mechanical ventilation and with increased short-term and long-term mortality (43, 44). In patients who survive ARDS, muscle dysfunction may persist for years after hospital discharge, impairing quality of life and affecting survival (1, 2, 9, 10, 17). Using a murine model of influenza A infection, we show that IL-6 released from the influenza A infected lung upregulates atrogin-1 to promote the active degradation of muscle proteins via the ubiquitin-proteasome pathway. Muscle mass and function could be preserved during influenza A infection through genetic deletion of atrogin-1 or through pharmacologic blockade of the IL-6 receptor. The fact that mice infected with IAV had negligible viral load detected in the soleus muscle homogenates and that differentiated C2C12 mouse myotubes have reduced muscle fiber size when treated with IL-6 and not when incubated with IAV, suggest an endocrine mechanism for the muscle loss observed in influenza A infected mice. The effects of IL-6 on fiber size were attenuated by knockdown of STAT3 or FoxO3a and by pharmacologic inhibition of the IL-6 receptor, but not with knockdown of FoxO1. These data suggest that muscle loss during influenza A infection is an active process initiated by endocrine signals from the injured lung.

Our data in mice are potentially relevant for humans. IL-6 and TNF- α are the first detectable cytokines after influenza infection (27, 37), and are increased in the serum of patients with lung injury (29–31). In humans with influenza A infection, IL-6 levels peak within 2 days of infection, and remain elevated for up to 7 days, with levels that correlate with symptom severity (27). In patients with ARDS, the activation of proteolytic pathways follows a similar time course, with increases in ubiquitination of muscle proteins detected within hours of onset of critical illness and remaining active for up to one week following discharge from the ICU (7, 45). Our *in vivo* measurements were performed at 1 week following infection of the influenza A virus (A/WSN/33 [H1N1]) administered intratracheally, at which point proteolytic pathways are active and muscle weakness is evident. In *atrogin-1* mice, we observed an increase in IL-6 in both serum and bronchoalveolar lavage fluid after IAV infected in all the genotypes, but even though it was profoundly increased in the *atrogin-1*^{-/-} mice the lack of atrogin-1 in the muscle is enough to protect against muscle atrophy.

While our studies demonstrate a deleterious role for IL-6 during IAV-induced lung injury, IL-6 can also be produced by muscle fibers during exercise where it acts as a myokine (46). For example, with extreme exercise, IL-6 may be sufficiently produced by working muscle to increase circulating levels up to 100-fold (47) without signs of muscle tissue damage (48) or a change in IL-6 expressing immune cells (49). During exercise, IL-6 is released from muscle in response to low glycogen stores (50) or activation of AMPK, and acts through autocrine and paracrine mechanisms to increase fat oxidation, and enhance insulin-stimulated glucose uptake through GLUT4 translocation (51). As a myokine, IL-6 also provides an endocrine signal to enhance hepatic glucose production during exercise (52).

Importantly, IL-6 is produced as the predominant myokine in response to exercise and yet, as we observed it can also induce muscle dysfunction as part of the “cytokine storm” associated with IAV infection. We speculate that differences in the kinetics of IL-6 release and its association with other pro-inflammatory cytokines may underlie the differential response of muscle to IL-6. After exercise, IL-6 levels peak rapidly then fall (53, 54) while sustained elevations are present for days after influenza infection (27). Further, IL-6 generated during exercise is not accompanied by an increase in other cytokines elevated during influenza A infection (55). Our data are consistent with those of others who have shown that prolonged exposure to recombinant IL-6 can induce muscle atrophy (22, 56), and with observations from others that high levels of IL-6 are associated with the age-related decline in muscle function due to sarcopenia (57).

We found that pharmacologic inhibition of the IL-6 receptor using the FDA approved drug tocilizumab attenuated muscle loss during influenza A infection. We did not see adverse effects of tocilizumab in the muscle as there was no difference in muscle mass of the soleus and EDL. Furthermore, we saw no change in total body weight in mice treated with tocilizumab. Nevertheless, in health, IL-6 has an important role in maintenance of muscle bulk, modulating muscle carbohydrate and lipid metabolism (47, 58–60). The differing roles for IL-6 signaling in the muscle during health and disease are poorly understood, and suggest that the effects of IL-6 in the muscle might be modulated by other factors present in the muscle microenvironment during disease. Furthermore, influenza A-induced muscle loss was only partially inhibited by tocilizumab, suggesting that other cytokines released in response to influenza A infection (e.g. IL-1 β , IL-18 and others), changes in inflammatory cell populations (e.g. Treg cells), or nutrient deprivation might independently contribute to muscle dysfunction (61–66). Influenza A infection in mice provides an excellent model system to further explore these possibilities.

We observed increased levels of atrogin-1 in muscle following influenza A infection, and heterozygous and homozygous loss of atrogin-1 resulted in a dose-dependent inhibition of influenza A induced muscle degradation. In cultured myotubes, the induction of atrogin-1 by IL-6 required the activation of STAT3 and FoxO3a, but not FoxO1. These data suggest inhibiting atrogin-1 directly or the pathways required for its induction may preserve muscle mass during influenza A infection. These results should be interpreted with some caution; however, as others have observed increased expression of mRNA for both *atrogin-1* and *MuRF1* in the skeletal muscle following influenza infection (67). Further, it is not clear whether the IL-6 and atrogin-1 mediated degradation of muscle proteins is adaptive process during influenza A infection. For example, the degradation of muscle proteins might provide metabolic substrates to fuel inflammatory cell responses to pathogen challenge. Alternatively, the systemic release of IL-6 might be a maladaptive consequence of the excessive inflammation thought to underlie the development of the multiple organ dysfunction syndrome. While our study does not answer this question, we saw no worsening in weight loss or mortality in influenza A infected atrogin-1 compared with wild type controls, nor did we observe any impact of tocilizumab on these parameters. However, tocilizumab has been associated transient neutropenia, increased body weight and elevated cholesterol in clinical trials (68, 69).

There are a number of limitations to this study. First, because the expression of atrogin-1 is largely limited to the muscle, the effects of atrogin-1 loss in other cells is less of a concern. However, it is possible that partial or complete developmental loss of atrogin-1 might activate pathways in *atrogin-1^{+/-}* and *atrogin-1^{-/-}* mice, respectively, which protect against influenza A-induced muscle loss. Further studies using inducible knockouts or pharmacologic inhibitors could address these concerns. Additionally, our model system is limited to infection with a single murine adapted influenza A virus. Studies using other strains of influenza A virus or administering tocilizumab to pigs, which are susceptible to circulating influenza A viruses could address this limitation. Finally, while IL-6 levels are often elevated in patients with ARDS secondary to other causes, the importance of IL-6 in muscle degradation in these conditions is unknown.

In summary, we provide evidence that influenza infection results in the release of IL-6 from the injured lung, which activates STAT3 and FoxO3a in the muscle to induce the expression of atrogin-1. Atrogin-1 is an E3 ligase that targets key muscle proteins for active degradation via the ubiquitin proteasome system. Thus, as depicted in Figure 6, IL-6 acts as an endocrine signal to promote active degradation of the skeletal muscle during influenza A pneumonia. Inhibition of the IL-6 receptor and genetic deletion of atrogin-1 attenuates the loss of muscle mass and function associated with influenza A infection, suggesting these pathways might be therapeutically targeted to preserve muscle mass in patients with influenza A-induced ARDS.

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Abbreviations used in this article:

IAV	influenza a virus
ARDS	acute respiratory distress syndrome
FoxO	forkhead box O
MuRF1	muscle RING finger 1
MAFbx/atrogin-1	muscle atrophy F-box protein
EDL	extensor digitorum longus
CSA	cross sectional area

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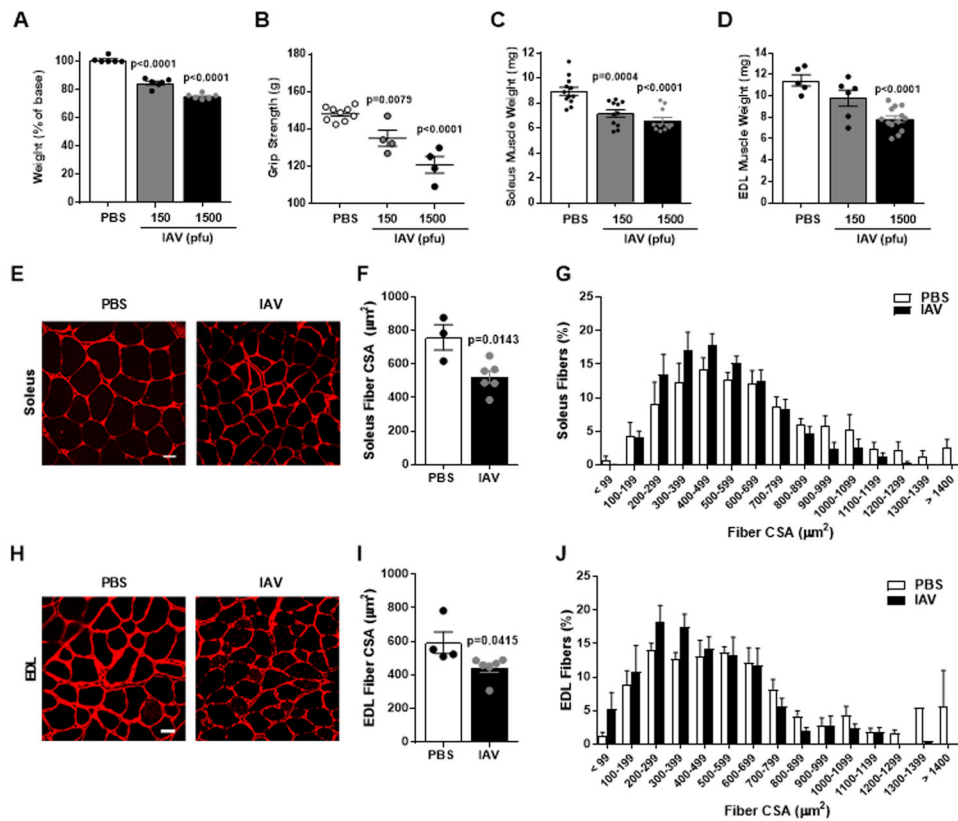


Figure 1. Wild-type mice lose muscle weight and strength after influenza infection and have reduced cross-sectional area (CSA) after influenza infection. Adult male C57Bl/6 mice were infected with 150 or 1500 pfu/mouse of influenza A Virus (IAV) or PBS (control) and 7 days post-infection (**A**) total body weight ($n=6$ mice/group) and (**B**) fore-limb grip strength were determined ($n=4-9$ mice/group). (**C**) Soleus ($n=11-12$ mice/group) and (**D**) EDL ($n=5-13$ mice/group) wet muscle weights were determined after muscles were excised, frozen and cryosectioned. (**E-J**) Soleus and EDL CSA were determined using cross-sections immunostained for laminin. (**E, H**) Representative images are shown. (40X magnification) Scale Bars = 20 μm . (**F**) Mean soleus fiber CSA ($n=3-6$ mice/group) and (**G**) a histogram depicting the fiber size distribution. (**I**) Mean EDL fiber CSA ($n=4-6$ mice/group) and (**J**) histogram depicting the fiber size distribution. Bars represent means \pm SEM. $n \geq 3$ of at least 3 independent experiments. A one-way ANOVA with Dunnett's post-hoc corrections for comparison with more than 3 groups and a Student's *t*-test between two groups were used to show statistical differences. All groups are compared to the control PBS-treated and *p*-values are shown.

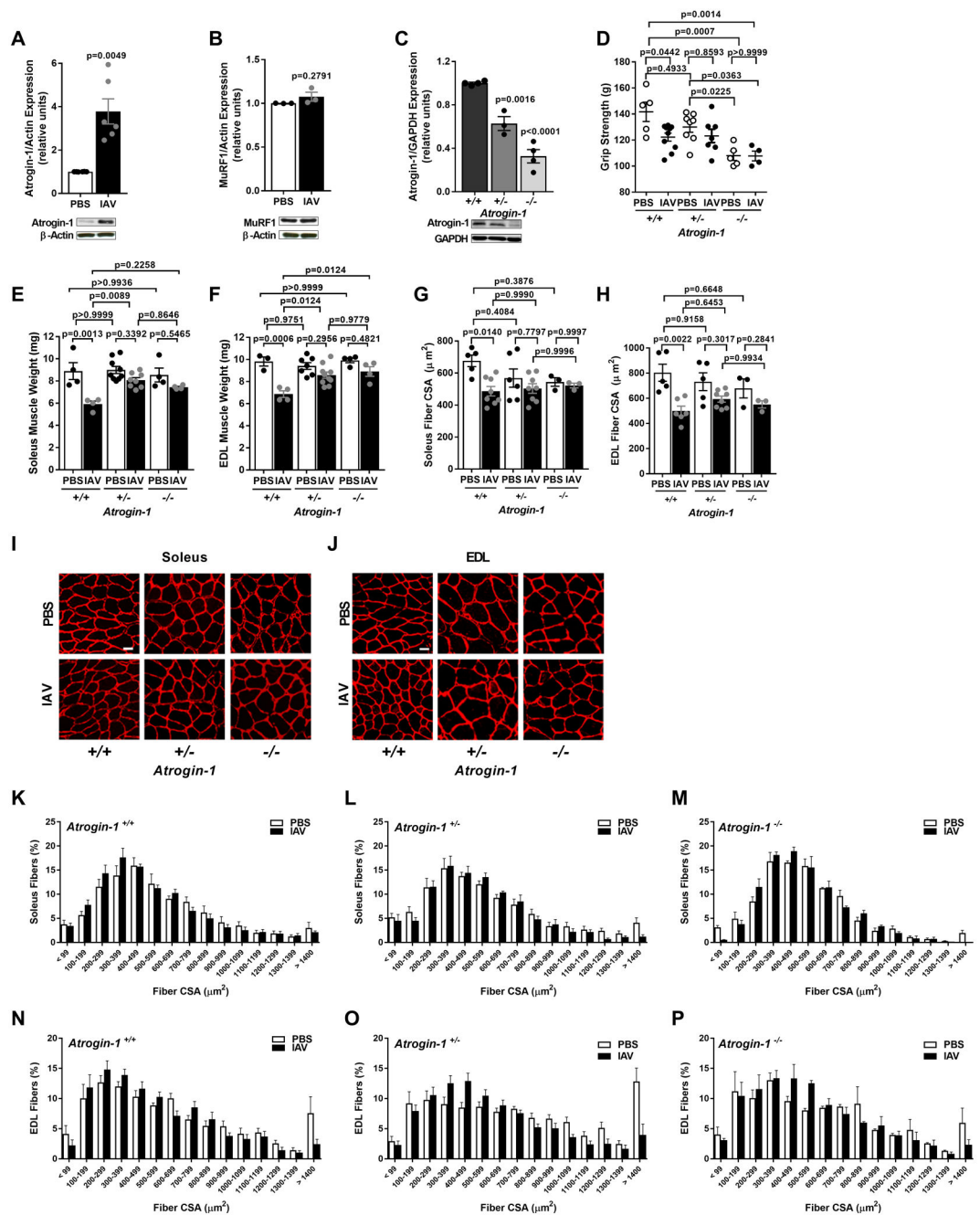


Figure 2. Influenza-induced muscle dysfunction is attenuated in mice with genetic loss of atrogenin-1.

(A-B) Adult male C57Bl/6 mice were infected with 1500 pfu/mouse of influenza A Virus (IAV) or PBS (control) and 7 days post-infection soleus muscles were excised and atrogenin-1 (n=6 mice/group) and MuRF1 (n=3 mice/group) expression was determined by Western blot. β -actin was used as loading control. A Student's *t*-test between two groups was used to show statistical differences. (C) Soleus muscle from *atrogenin-1* ($^{+/+}$, $^{+/-}$ and $^{-/-}$) mice were excised and atrogenin-1 expression was determined by Western blot. GAPDH was used as loading control. (n=3-4 mice/group). A one-way ANOVA with Dunnett's post-hoc

corrections for comparison with more than 3 groups was used to show statistical differences. **(D-P)** *Atrogin-1*^{+/+}, *Atrogin-1*^{+/-} and *Atrogin-1*^{-/-} mice infected with 1500 pfu/mouse of influenza A Virus (IAV) or PBS (control) and 7 days post-infection **(D)** fore-limb grip strength was determined. (n=4-10 animals/group). A one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. **(E)** Soleus (n=4-9 animals/group) and **(F)** EDL (n=3-9 groups/group) wet muscle weight were determined after muscles were excised, frozen and cryosectioned. A one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. **(G-P)** Soleus (n=3-8 mice/group) and EDL (n=3-8 mice/group) cross sectional area (CSA) were determined using muscle cross-sections immunostained for laminin. A one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. **(G-H)** Meansoleus and EDL fiber CSA are shown. **(I-J)** Representative images are shown. (40X magnification) Scale Bars = 20 μ m. **(K-P)** Histograms depicting the fiber size distribution for soleus muscle of **(K)** *Atrogin-1*^{+/+} **(L)** *Atrogin-1*^{+/-} and **(M)** *Atrogin-1*^{-/-} and EDL muscles of **(N)** *Atrogin-1*^{+/+} **(O)** *Atrogin-1*^{+/-} and **(P)** *Atrogin-1*^{-/-}. Bars represent means \pm SEM and p-values are shown. n = 3 from at least 3 independent experiments.

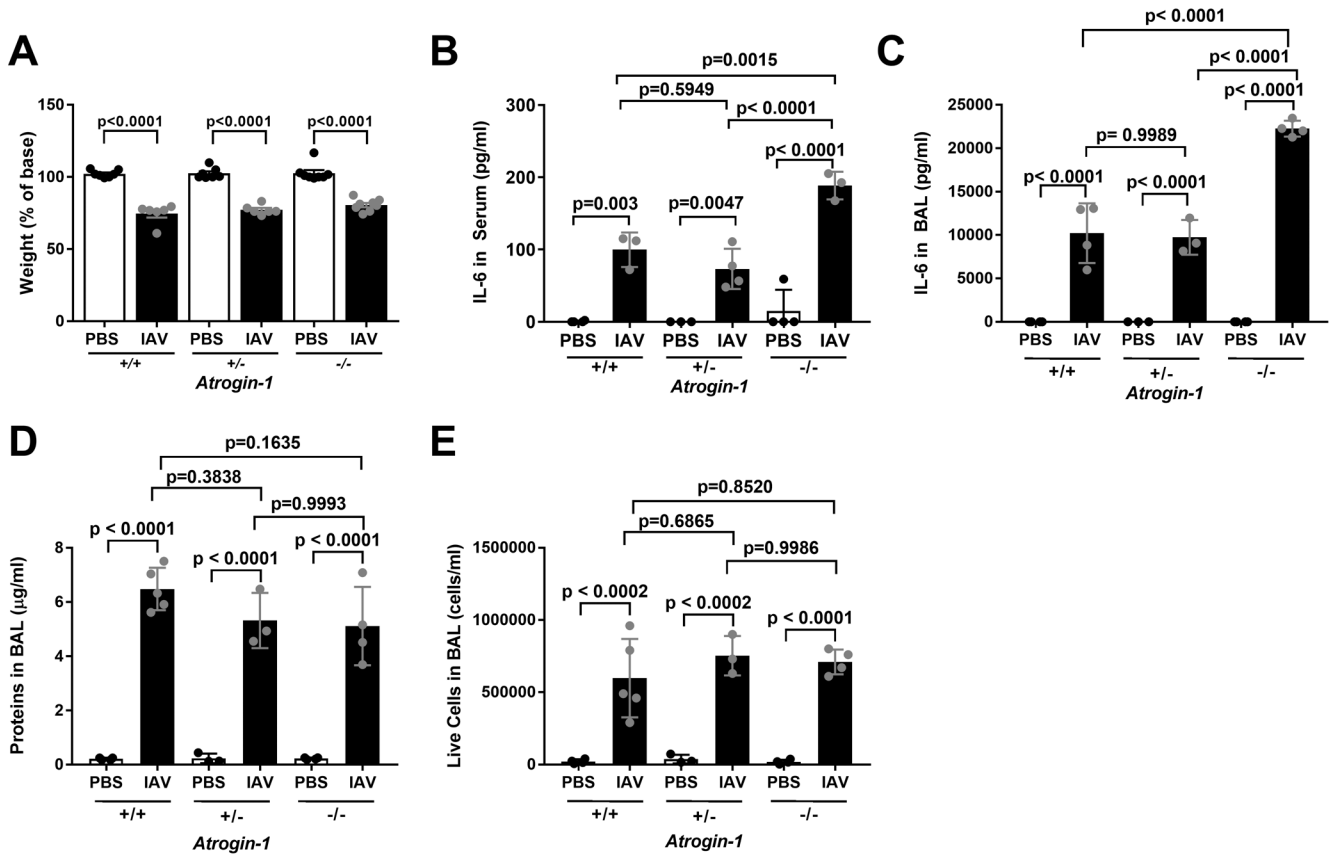


Figure 3. Influenza-induced body weight is attenuated and IL-6 is increased in serum and bronchoalveolar lavage fluid in mice with genetic loss of atrogin-1. Adult male *atrogin-1* (+/+, +/- and -/-) mice were infected with 1500 pfu/mouse of influenza A Virus (IAV) or PBS (control) and 7 days post-infection total body weight was assessed (n=6-8 mice/group). **(B)** IL-6 was measured by ELISA in the serum. **(C)** IL-6 was measured by ELISA in the bronchoalveolar lavage (BAL) fluid. **(D)** Live cells in BAL fluid were measured by trypan blue exclusion. **(E)** Proteins in BAL fluid were measured by Bradford Assay. (n=3-5 mice/group). Bars represent means \pm SEM and p-values are shown. A one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. n = 3 from at least 3 independent experiments.

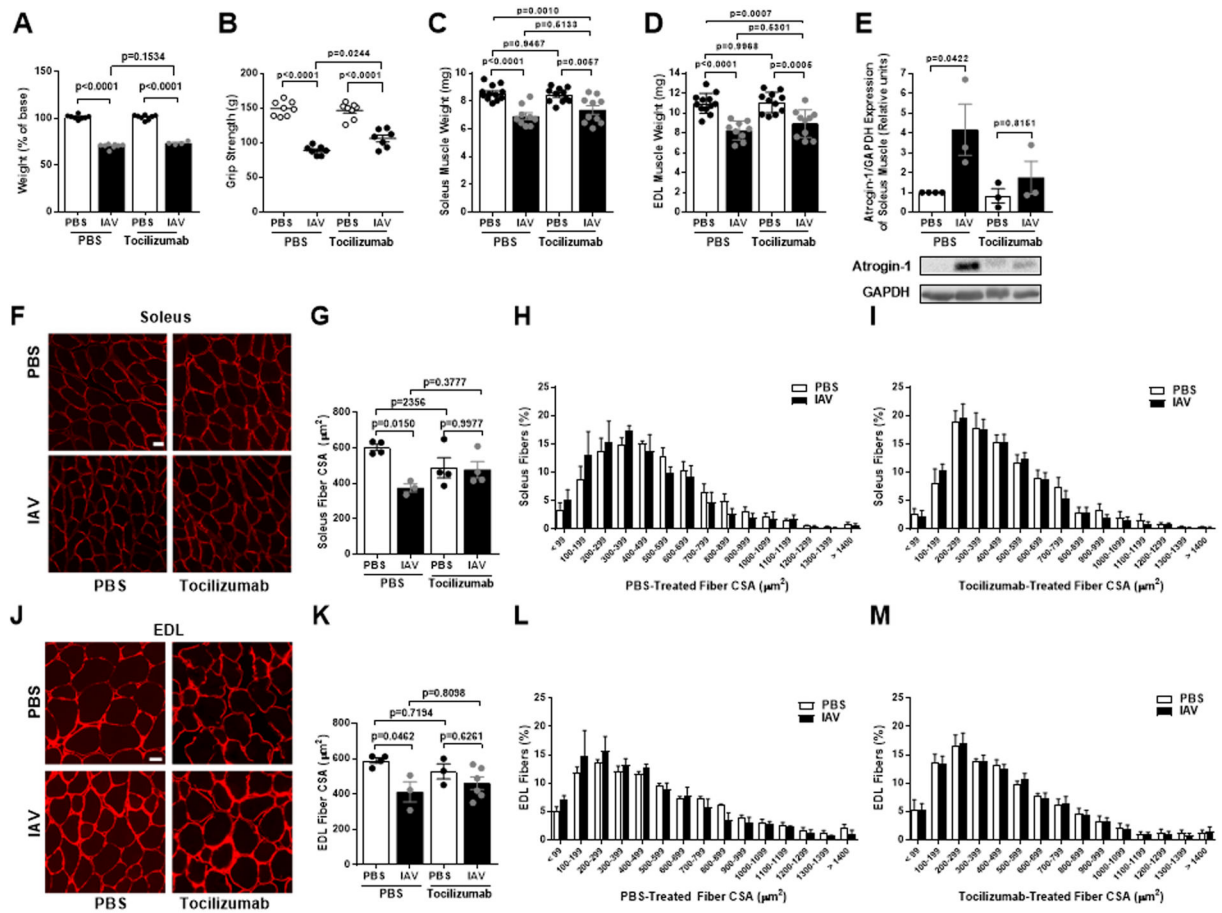


Figure 4. The IL-6-receptor antibody tocilizumab attenuates influenza-induced muscle dysfunction.

Adult male wildtype C57Bl/6 mice were pre-treated with the interleukin-6 (IL-6) receptor antibody, tocilizumab (8 mg/kg), 24 hours later infected with 1500 pfu/mouse of influenza A Virus (IAV) or PBS (control) and 7 days post-infection (A) total body weight (n=4-8 mice/group). (B) fore-limb grip strength was determined (n=7-8 mice/group). (C) Soleus (n=9-12 mice/group) and (D) EDL (n=10-12 mice/group) wet muscle weight were determined after muscles were excised, frozen and cryosectioned. For A-D, a one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. The p-values are shown. (E) Quantification and representative Western blot of atrogin-1 expression in the mouse soleus muscle of PBS and tocilizumab-treated mice (n=3 mice/group). A one-way ANOVA with Dunnett's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. All groups are compared to the control PBS-treated and p-value is shown. (F-M) Muscle cross-sections immunostained for laminin antibody were used to determine the mean soleus and EDL fiber cross-sectional area (CSA). (F, J) Representative images are shown. (40X magnification) Scale Bars = 20 μm. (G) Soleus (n=4 mice/group) and (K) EDL (n=3-6 mice/group) mean fiber cross-sectional area (CSA) are shown. A one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. (H-I, L-M) Histograms depicting the fiber size distribution are shown for (H-I) soleus and (L-M) EDL

mice treated with PBS or tocilizumab prior to influenza infection. Bars represent means \pm SEM and p-values are shown. n = 3 of at least 3 independent experiments.

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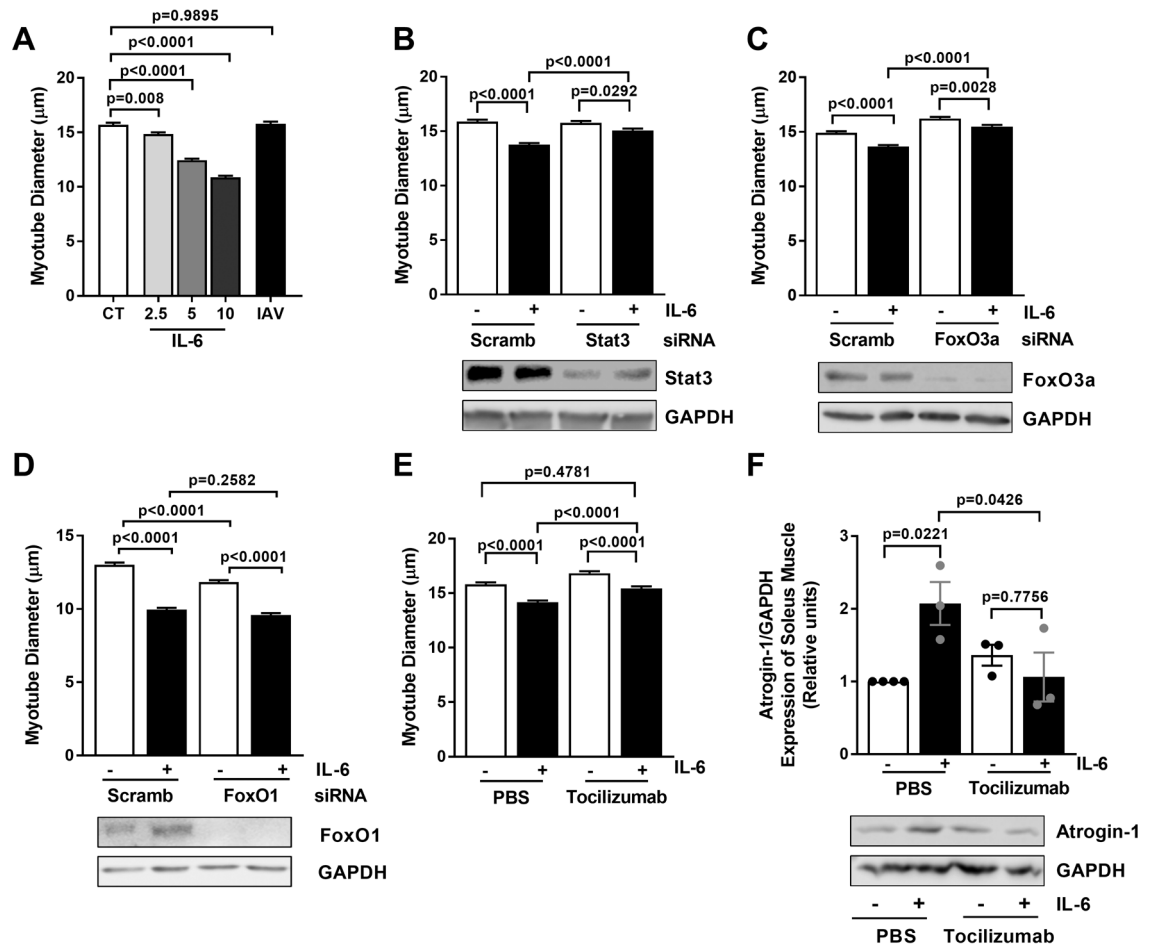


Figure 5. IL-6 acts via STAT3, FoxO3a, and Atrogin1 to induce skeletal muscle dysfunction.

(A) C2C12 myotubes were treated with recombinant IL-6 (2.5, 5, 10 ng/ml) or infected with 2 MOI IAV on Day 4 of differentiation for 16 hours. Phase contrast images were acquired on Day 5 of differentiation. Graph depicts average myotube diameters (~375 measurements/group; from at least 4 independent experiments). A one-way ANOVA with Dunnett's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. All groups are compared to the control (CT) and p-value is shown. (B-D) Myotubes were transfected with scrambled (Scramb), *STAT3* (~425 measurements/group; from at least 3 independent experiments), *FoxO3a* (~625 measurements/group; from at least 5 independent experiments) or *FoxO1* (~679 measurements/group; from at least 4 independent experiments) siRNA on Day 2 of differentiation followed by recombinant IL-6 treatment (10 ng/ml) on Day 4. Phase contrast images were obtained and cell lysates were analyzed on Day 5 of differentiation. Upper panel shows graph depicting average myotube diameter while lower panel depicts Western blot showing knock-down efficiency. GAPDH is shown as loading control. A one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. (D-E) C2C12 myotubes were treated with tocilizumab 10 μg/ml on Day 3 of differentiation for 24 hours followed by treatment with recombinant mouse IL-6 for 16 hours. (E) Graph depicts average myotube diameter (~475 measurements/group; from at least 4 independent experiments). A

one-way ANOVA with Tukey's post-hoc corrections for comparison with more than 3 groups was used to show statistical differences. **(F)** Quantification and representative Western blot of atrogin-1 expression in C2C12 myotubes. GAPDH is shown as loading control (n=3 mice/group). Bars represent means \pm SEM and p-values are shown.

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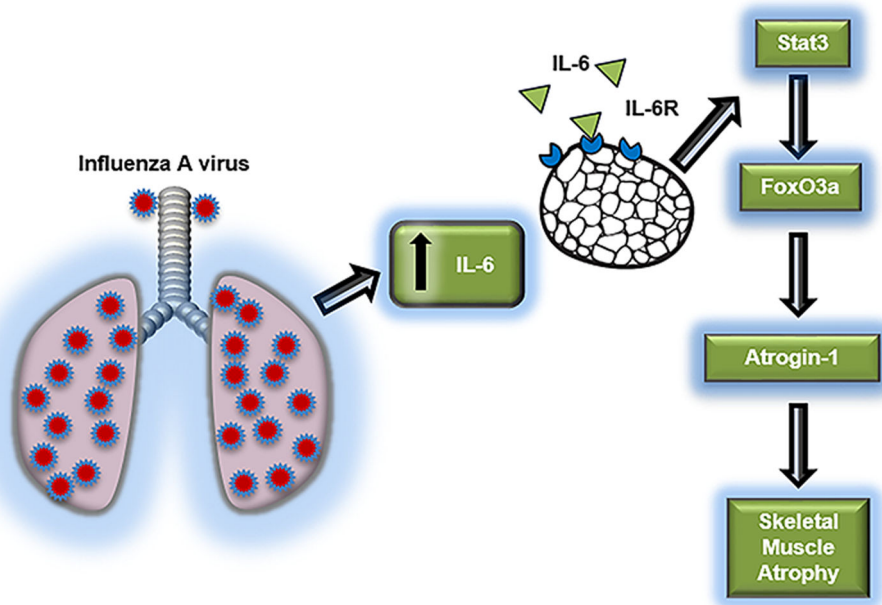


Figure 6.

In a murine model of influenza A infection, the release of IL-6 leads to the up regulation of the E3 ligase atrogin-1 leading to skeletal muscle dysfunction.