Denervation atrophy is independent from Akt and mTOR activation and is not rescued by myostatin inhibition

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ABSTRACT
The purpose of our study was to compare two acquired muscle atrophies and the use of myostatin inhibition for their treatment. Myostatin naturally inhibits skeletal muscle growth by binding to ActRIIB, a receptor on the cell surface of myofibers. Because blocking myostatin in an adult wild-type mouse induces profound muscle hypertrophy, we applied a soluble ActRIIB receptor to models of disuse (limb immobilization) and denervation (sciatic nerve resection) atrophy. We found that treatment of immobilized mice with ActRIIB prevented the loss of muscle mass observed in placebo-treated mice. Our results suggest that protection from disuse atrophy is regulated by serum and glucocorticoid-induced kinase (SGK) rather than by Akt. Denervation atrophy, however, was not protected by ActRIIB treatment, yet resulted in an upregulation of the pro-growth factors Akt, SGK and components of the mTOR pathway. We then treated the denervated mice with the mTOR inhibitor rapamycin and found that, despite a reduction in mTOR activation, there is no alteration of the atrophy phenotype. Additionally, rapamycin prevented the denervation-induced upregulation of the mTORC2 substrates Akt and SGK. Thus, our studies show the denervation atrophy is not only independent from Akt, SGK and mTOR activation but also has a different underlying pathophysiological mechanism than disuse atrophy.

KEY WORDS: Skeletal muscle, Muscle atrophy pathophysiology, TGF-β signaling, Myostatin, Denervation atrophy

INTRODUCTION
Healthy skeletal muscle maintains a balance between protein synthesis and protein degradation (Glass, 2010; McKinnell and Rudnicki, 2004). Disruption of this balance, resulting from a wide variety of conditions including immobilization (or disuse), chronic obstructive pulmonary disease, starvation, denervation or renal failure, can all lead to atrophy of the skeletal muscle (Glass, 2010; McKinnell and Rudnicki, 2004; Ruegg and Glass, 2011). Many studies have indicated that there is a common molecular pathway that is disrupted in all forms of skeletal muscle atrophy regardless of their etiology (Lecker and Goldberg, 2002; Lecker et al., 2004; Sacheck et al., 2007). This paradigm is thought to revolve mainly around the insulin-like growth factor (IGF) pathway and the serine-threonine kinase protein kinase B, commonly referred to as Akt (Glass, 2010; Rommel et al., 2001; Schiaffino and Mammucari, 2011). The Akt signaling cascade can regulate muscle mass by both inhibiting protein degradation and promoting protein synthesis; overexpression of Akt can lead to both muscle hypertrophy and the prevention of atrophy (Bodine et al., 2001b; Glass, 2010; Rommel et al., 2001).

Akt is also capable of inducing hypertrophy by promoting protein synthesis through the activation of the mammalian target of rapamycin (mTOR) pathway (Bodine et al., 2001b; Glass, 2010; Laplante and Sabatini, 2012; Ma and Blenis, 2009). The mTOR pathway consists of two complexes, mTORC1 and mTORC2 (Laplante and Sabatini, 2012; Ma and Blenis, 2009). Both complexes, when activated, contain the phosphorylated form of mTOR and the shared scaffold protein mammalian lethal with Sec13 protein 8 (mLST8; also called GpL) (Inoki et al., 2005; Laplante and Sabatini, 2012; Ma and Blenis, 2009). The mTORC1 complex also includes the protein raptor and can phosphorylate eIF4E-binding protein (4E-BP1) and p70 ribosomal S6 kinase (p70S6k), both of which will lead to an increase in protein translation (Fenton and Gout, 2011; Ma and Blenis, 2009; Richter and Sonenberg, 2005). The phosphorylation and activation of p70S6k specifically in skeletal muscle, has been shown to be dependent upon a second scaffold protein, eIF3f (Csibi et al., 2009; Csibi et al., 2008). The mTORC2 complex, with the unique protein rictor, creates a positive-feedback loop for both complexes by activating Akt and the serum and glucocorticoid-induced kinase (SGK) (Inoki et al., 2005; Laplante and Sabatini, 2012; Ma and Blenis, 2009). SGK has recently been shown to maintain and preserve skeletal muscle mass by inactivating FoxO3a and activating the mTOR pathway (Andres-Mateos et al., 2013; Aoyama et al., 2005; Brunet et al., 2001).

Myostatin is a protein that is thought to disrupt the balance between protein synthesis and protein degradation of healthy skeletal muscle by inhibiting Akt (Glass, 2010; Morissette et al., 2009; Sartori et al., 2009; Trendelenburg et al., 2009). Myostatin, also known as growth and differentiation factor-8 (GDF-8), is a reviewed here.
inhibitors to treat inherited muscle disorders. Myopathies such as dystrophin-negative muscular dystrophy, limb girdle muscular dystrophy and spinal muscular atrophy, among others, have all been treated with myostatin inhibitors (Morine et al., 2010; Morrison et al., 2009; Ohsawa et al., 2006; Sumner et al., 2009; Wagner et al., 2002). However, only a few studies have attempted to use myostatin inhibitors for the treatment of acquired myopathies and they have focused mainly on systemic conditions such as cancer cachexia, diabetes, or even obesity (Guo et al., 2012; Guo et al., 2009; Koncarevic et al., 2012; Zhou et al., 2010).

In our study we wanted to test the hypothesis the soluble ActRIIB receptor, a myostatin receptor fused to an Fc domain (Lee et al., 2005) that inhibits myostatin signaling, is able to prevent single-limb, acquired muscle atrophy. We used two mouse models for this purpose: a hindlimb immobilization model (disuse atrophy) and a sciatic nerve resection model (denervation atrophy). Our results indicate that myostatin inhibition is beneficial in settings of disuse, but not denervation, atrophy. Our subsequent molecular analysis and comparison of these two atrophy models led us to the surprising conclusion that denervation atrophy is not dependent upon the activation of Akt, SGK or mTOR, suggesting that there is not a universal pathway responsible for all forms of atrophy and therefore denervation should be treated as a distinct pathogenic condition.

RESULTS

Myostatin inhibitor ActRIIB protects muscle from disuse, but not denervation, atrophy

To assess whether myostatin inhibition would protect muscle from atrophy, we used two separate mouse models. We either attached a surgical staple to immobilize one hindlimb of our mice or denervated them by surgical removal of the sciatic nerve from one hindlimb, and then treated both groups with 10 mg/kg ActRIIB for 3 weeks. Owing to the enlargement of all non-challenged muscle, ActRIIB treatment resulted in a substantial increase in total body mass in both atrophy models (Fig. 1A,C, left graphs).

The immobilized placebo-treated group had 19.5% less tibialis anterior (TA) muscle mass compared with controls (Fig. 1A, center graph). However, the ActRIIB-treated immobilized mice did not show loss of TA muscle mass compared with untreated controls (Fig. 1A, center graph). When compared with controls, the measurement of the minimum feret diameter (MFD) showed a similar pattern of loss of muscle fiber diameter in the immobilized placebo group (13.3% reduction), but not in the ActRIIB-treated mice (Fig. 1A, right graph, 1B).

In contrast to the immobilization experiment, both the placebo- and ActRIIB-treated denervated mice lost a significant amount of TA muscle mass compared with sham-operated controls (49.7% and 44.8%, respectively; Fig. 1C, center graph). Measurement of the MFD also showed that the denervated groups had 39.6% and 45.2% smaller TA muscle fibers (placebo- and ActRIIB-treated, respectively) compared with sham-operated controls (49.7% and 44.8%, respectively). Notably, the authors also show that denervation atrophy is not affected by activation of pro-growth molecules that have been shown to benefit other forms of atrophy.

Implications and future directions

These preclinical data show that myostatin inhibition can prevent disuse atrophy but not muscle atrophy caused by denervation. Thus, for myostatin inhibition to be effective, an intact nerve-muscle conduction system must be present. This is essential information for future clinical applications of myostatin inhibition. Of equal importance, these findings provide new information about the molecular basis of disuse atrophy and of denervation atrophy. Specifically, although it is widely believed that all forms of skeletal muscle atrophy follow a similar molecular pattern, this work suggests that the mechanism of denervation atrophy is different to that of other forms of muscle wasting.
Smad3. In addition, ActRIIB treatment of immobilized mice also did not alter the activation of Smad2 or Smad3 (Fig. 2A). Denervation alone induced a threefold upregulation in total Smad2 and a fivefold increase in active pSmad3, but ActRIIB treatment did not attenuate either of these (Fig. 2A). We subsequently examined the expression of the non-canonical TGF-β signaling markers extracellular-signal-regulated kinases 1 and 2 (ERK1/2). Both models of atrophy demonstrated an increase in active pERK1/2 compared with their respective controls (Fig. 2B). ActRIIB treatment prevented the activation of ERK in the immobilized model but not in the denervated model. Denervation also resulted in a significant upregulation of total ERK1/2 protein expression, something not observed in the immobilized model (Fig. 2B).

Myostatin also influences the expression of other proteins involved in muscle growth and regeneration, including myogenin and p21. Myostatin promotes the expression of p21 and inhibits the expression of myogenin, both of which will negatively regulate the differentiation of muscle precursor cells and therefore muscle growth (Langley et al., 2002; Ohsawa et al., 2006). In both models of atrophy, we found that p21 expression is unchanged, both with and without ActRIIB treatment. However, loss of myogenin was observed in the immobilization model but not in the immobilized treated with ActRIIB. Denervation alone induced a significant upregulation in myogenin, but was not changed further by ActRIIB treatment (supplementary material Fig. S1).

We found that muscle disuse results in an upregulation of the non-canonical TGF-β signaling marker pERK1/2, which was prevented by ActRIIB treatment. In addition, another marker of myostatin signaling, myogenin, is sensitive to immobilization but not in ActRIIB-treated mice. ActRIIB treatment did not inhibit the upregulation of canonical or non-canonical TGF-β signaling markers or any other marker of myostatin signaling in denervated muscle.

**Akt and SGK are dysregulated in disuse and denervation atrophy**

Next we examined the expression levels of Akt in both atrophy models. We found that immobilization with or without ActRIIB treatment did not alter the amount of active pAKT (Fig. 3). An upregulation in total Akt, however, was observed in immobilized mice treated with ActRIIB compared with controls. Conversely, in both the placebo- and ActRIIB-treated denervated models showed a threefold upregulation.
of SGK expression when compared with sham-operated controls (Fig. 3).

We next examined the expression and activation levels of FoxO3a, a downstream target of SGK and Akt. Both SGK and Akt have been shown to have equal affinity to phosphorylate FoxO3a at T32, but Akt preferentially phosphorylates S253, the site that will inactivate the protein (Brunet et al., 2001). In both atrophy models, with or without ActRIIB treatment, we found no significant difference in phosphorylation at either the T32 or S253 site or in total expression levels of FoxO3a (Fig. 4A). Expression levels of the FoxO3a target atrogin-1 further substantiated these results. We found no difference in expression in either immobilization model, but both placebo- and ActRIIB-treated denervated muscle had reduced atrogin-1 levels (Fig. 4A).

The Akt-mTOR-FoxO3a pathway is also known to play an important part in the regulation of autophagy in skeletal muscle (Schiaffino and Mammucari, 2011; Zhao et al., 2008). Examination of autophagy markers in immobilized mice showed no change between the control, placebo-treated and ActRIIB-treated groups (Fig. 4B). Denervation, however, induced a significant increase in several markers of autophagy, including Lamp2, LC3b, ATG7 and p62 (Fig. 4B). Treatment with ActRIIB did not further change the expression of autophagy markers in denervated muscle.

Our results showed that SGK, not Akt, is lost as a result of immobilization, but not when the muscle is protected by treatment with ActRIIB. Levels of both SGK and Akt were significantly upregulated in denervated muscle with or without ActRIIB treatment. Moreover, a significant upregulation in autophagy markers was observed in denervation, but not disuse, atrophy.

The mTOR signaling pathway is upregulated in denervation atrophy

The mTOR signaling cascade plays an important role in skeletal muscle maintenance by promoting protein synthesis (Glass, 2010; Ma and Blenis, 2009). We therefore examined the activation of both the mTORC1 and mTORC2 complexes in our models of atrophy and ActRIIB treatment.

We found no change in the mTOR complex components p-mTOR, total mTOR, raptor and rictor in immobilized mice with or without ActRIIB treatment (Fig. 5A). However, a scaffold protein common between both mTOR complexes, GβL, was downregulated with immobilization, but was maintained at control levels in ActRIIB-treated immobilized muscle (supplementary material Fig. S2). A substrate of mTORC1, p70S6k, and a second scaffold protein, eIF3f, which facilitates the interaction between mTORC1 and p70S6k, also showed reduced expression in immobilized muscle (Fig. 5B and supplementary material Fig. S2). The loss of p70S6k and eIF3f expression was not seen in immobilized mice treated with ActRIIB (Fig. 5B). A second mTORC1 substrate, 4E-BP1, did not show any change in phosphorylation or total abundance in placebo- or ActRIIB-treated immobilized mice compared with controls (supplementary material Fig. S2).

The denervated mice demonstrated a significant upregulation in several components of both the mTORC1 and mTORC2 complexes (Fig. 5A). The three- to fivefold upregulation of p-mTOR, total mTOR, raptor and rictor was unchanged by ActRIIB treatment of denervated muscle (Fig. 5A). The scaffold proteins GβL and eIF3F were also upregulated in both denervation models compared with sham-operated controls (supplementary material Fig. S2). Downstream of the mTORC1 complex, however, we found that denervation results in a significant, almost threefold, drop in phosphorylation at the T389 activation site of p70S6k, despite no change in total p70S6k expression (Fig. 5B). ActRIIB treatment of denervated mice did not prevent the loss of p70S6k activation. Both placebo and ActRIIB treatment of denervated mice led to an upregulation in total and phosphorylated 4E-BP1 (supplementary material Fig. S2).

Active p-mTOR can also regulate autophagy by the phosphorylation and inactivation of ULK1 (the mammalian homolog of Caenorhabditis elegans ATG1), a protein known to act as an inhibitor of autophagy. The denervated mice demonstrated a significant upregulation of ATG1, a protein known to act as an inhibitor of autophagy.
negatively regulate p70S6k (Egan et al., 2011; Lee et al., 2007). We found no change in total ULK1 expression or phosphorylation at the mTOR-specific inhibitory S757 site in immobilized mice, with or without ActRIIB treatment, compared with controls (supplementary material Fig. S3). Similarly, sham-operated, denervated and ActRIIB-treated denervated muscle did not show any difference in total ULK1 expression levels. However, both placebo- and ActRIIB-treated denervated mice produced a significant drop in phosphorylation at the S757 inhibitory site compared with sham-operated controls (supplementary material Fig. S3).

We found that, compared with controls, muscle disuse did not change the levels of the main mTOR-complex components. However, immobilization did lead to reduced expression of p70S6k and several mTOR-associated scaffold proteins. Immobilized mice were protected from the loss of these proteins by treatment with ActRIIB. In contrast, denervation atrophy resulted in a significant upregulation in nearly all components of the mTOR pathway except for p70S6k – changes that were not prevented by ActRIIB treatment. Although total p70S6k expression did not change, denervation led to the loss of active p-p70S6k, which might be due to an increase in active ULK1.

**Fig. 3. Loss of SGK, but not Akt, is observed in disuse atrophy, but upregulation of both occurs in denervation atrophy.** Western blot analysis of TA muscle protein lysates. No change in active pAkt is seen between the control (‘C’), immobilized (‘I’) and immobilized with ActRIIB treatment (‘I+A’) groups. Total Akt is increased in immobilized mice treated with ActRIIB compared with controls. SGK expression is decreased with immobilization, but not when mice are treated with ActRIIB. Both denervation alone (‘D’) and with ActRIIB treatment (‘D+A’) lead to a substantial upregulation in active pAkt, total Akt and SGK when compared with sham-operated controls (‘S’). Quantitative analysis of blots is displayed in the graph with arbitrary units of mean ± s.e.m. *P<0.01 and †P<5.0×10^{-5} with respect to controls. Lines indicate where intervening lanes have been removed from a single image to show the most representative band for that treatment group.

Rapamycin treatment does not alter the denervation atrophy phenotype

The upregulation of mTOR signaling observed in denervated muscle could be a compensatory mechanism employed to prevent further atrophy or could be contributing to the pathogenic phenotype (Ramos et al., 2012). In order to clarify this difference in our model, we next treated the denervated mice with the mTOR inhibitor rapamycin (2 mg/kg body weight) for 3 weeks.

We found that, compared with sham-operated controls, denervation resulted in a 50.2% loss of TA muscle mass that was not prevented by treatment with rapamycin (Fig. 6A, top graph). MFD measurements also showed that muscle fiber size was reduced by 39.8% and 41.1% in denervated muscle with or without rapamycin treatment, respectively, compared with sham-operated controls (Fig. 6A, bottom graph, 6B).

We then looked at the mTOR signaling pathway in TA muscle protein lysates. As previously demonstrated, muscle denervation led to an increase in p-mTOR, total mTOR, raptor, rictor, GβL and eIF3f compared with sham-operated controls (Fig. 6C and supplementary material Fig. S4). However, compared with the placebo group, rapamycin treatment of denervated mice resulted in a nearly twofold reduction in active p-mTOR. A trend towards reduced raptor expression was also seen in rapamycin compared with placebo-treated denervated mice, but it did not reach significance (Fig. 6B). We observed that, of the scaffold proteins, rapamycin treatment of denervated mice reduced the expression of GβL, but not eIF3f (supplementary material Fig. S4). No change was seen between the denervated and denervated with rapamycin treatment in the mTORC1 substrates p70S6k and 4E-BP1 (Fig. 6D and supplementary material Fig. S4). We also noted that the expression of rictor was not upregulated in denervated mice given rapamycin treatment, compared with sham-operated controls (Fig. 6B). A complete knock down of p-mTOR was achieved by treating denervated mice with a higher dose of rapamycin (10 mg/kg body weight); however, because the phenotype remained the same as with the low-dose treatment, we continued our analysis using the more physiological dose of 2 mg/kg body weight (supplementary material Fig. S5).

Because both Akt and SGK are substrates of the mTORC2 complex, we also examined the expression levels of these proteins in our rapamycin-treated denervation model. Denervation alone resulted in an upregulation of pAkt, total Akt, and SGK (Fig. 7A). However, we found that rapamycin treatment prevented the increase in active pAkt and SGK expression, but not the increase in total Akt expression, caused by denervation alone (Fig. 7A). We once again observed no difference in expression or activation of FoxO3a in denervated muscle, with or without rapamycin treatment. In addition, rapamycin treatment did not prevent the loss of atrogin-1 expression observed with denervation alone (Fig. 7B).

Our data demonstrate that rapamycin treatment does not rescue denervation atrophy despite inhibiting the activation of mTOR in the atrophic muscle. In addition, rapamycin treatment prevented the upregulation of pAkt and SGK – but not the downregulation of atrogin-1 – that is normally seen in denervation atrophy.

**DISCUSSION**

The ability to prevent or treat acquired forms of skeletal muscle atrophy has the potential for wide-reaching benefits to millions of patients. Immobilization alone is a natural complication from many primary conditions, including limb casting, reduced movement when ill or bed rest, all of which can lead to atrophy of the skeletal muscles. The loss of muscle mass can prolong recovery from the primary condition of the patient and increase rehabilitation time. Myostatin inhibitors are excellent candidates for the treatment of acquired muscle atrophies owing to their dramatic and immediate
effect on muscle. Indeed, numerous pharmaceutical companies are, in fact, currently working on developing myostatin inhibitors for the treatment of a variety of muscle disorders. A careful analysis of how these inhibitors influence muscle mass under various and unique pathological conditions is an essential step towards bringing them to the clinic.

Our studies show that myostatin inhibition has the potential for clinical application in the prevention of disuse atrophy. This protection was demonstrated by the preservation of both muscle mass and fiber diameter in immobilized mice (Fig. 1A). However, when this type of treatment does advance to the clinic, it will have to be taken into consideration that currently available myostatin inhibitors cause an increase in muscle mass in the body. Of equal importance are our findings that myostatin inhibition is not effective against atrophy when the neuromuscular connection has been lost (Fig. 1C). This result was somewhat surprising given that the mouse model for amyotrophic lateral sclerosis (ALS), a disease that results in loss of muscle innervation, showed improvement with myostatin inhibition (Morrison et al., 2009). In light of our results, however, this phenomenon is probably due to the heterogeneity of the innervated and denervated muscle fibers that are a consequence of this disease. The innervated muscles would benefit and become larger as a result of myostatin inhibition and be able to compensate for the unaffected, non-innervated fibers. In addition, despite previous research suggesting that denervation leads to an increase in myostatin transcript and protein expression, our data indicate that this might not be the main reason for the loss of muscle mass (Baumann et al., 2003; Liu et al., 2007; Shao et al., 2007). Owing to the extensive post-translational processing and modification that occurs to myostatin before it is a mature protein, it is possible that the previous studies have measured non-functional protein or untranslated mRNA (Lee, 2004).

Our subsequent analyses of the TGF-β signaling pathway in immobilized and denervated groups showed that myostatin inhibition did not reduce the canonical TGF-β signaling markers, pSmad2 and pSmad3, in either model (Fig. 2A). This is in contrast to previously published work demonstrating that myostatin itself will increase Smad activation in skeletal muscle and one report showing that ActRIIB will lower pSmad2 levels in a non-wild-type mouse (Langley et al., 2002; Rebbapragada et al., 2003; Sartori et al., 2009; Zhou et al., 2010). Our data might instead suggest that it is the downregulation of non-canonical TGF-β signaling markers that is of greater importance in understanding the effect that ActRIIB has on the muscle. However, it is possible that, owing to the long duration of our experiment, we have missed the ActRIIB-induced downregulation of Smad activation in our models.
We also explored the Akt-FoxO3a-mTOR pathway because myostatin itself can inhibit its activation and it is another commonly investigated atrophy regulation pathway (Glass, 2010; Trendelenburg et al., 2009). Neither of our atrophy mouse models, however, demonstrated the expected increase in activation and phosphorylation of Akt when treated with ActRIIB (Fig. 3) (Morissette et al., 2009; Trendelenburg et al., 2009). This complements a previous study showing that loss of Akt expression does not attenuate the muscle hypertrophy response to ActRIIB treatment (Goncalves et al., 2010). In addition, in our immobilization model, we found that SGK, not Akt, exhibited the expected pattern of loss of expression with atrophy but not in mice protected by ActRIIB treatment (Fig. 3). Our results suggest that Akt and SGK signaling to further clarify our results. We first examined the expression and activation of FoxO3a, the main transcription factor needed for the activation of atrogenes, a set of E3 ubiquitin ligases known to be highly involved in muscle atrophy (Lecker and Goldberg, 2002; Zhao et al., 2008). Despite loss or overexpression of either Akt or SGK, we found no difference in the phosphorylation of FoxO3a in either atrophy model (Fig. 4A). A FoxO3a atrogen target, atrogin-1, is also unchanged in the immobilized mice and is actually decreased in denervated muscle. We conclude that, at 3 weeks, neither FoxO3a nor atrogin-1 are mediating either form of atrophy or providing protection from disuse atrophy with ActRIIB treatment. It is possible that the involvement of FoxO3a and atrogin-1 are more immediate and occur at an earlier time point than we analyzed. A temporal involvement of atrogin-1 is indeed likely because an atrogin-1 knockout mouse is partially protected from denervation atrophy (Bodine et al., 2001a).

We also examined markers of another type of protein degradation regulated by the Akt-mTOR-FoxO3a pathway, autophagy. Because immobilization does not change any of the autophagy markers studied, we propose that autophagy is not a regulator of long-term disuse atrophy. Conversely, denervation led to a significant increase in all pro-autophagy markers examined, as has been previous described in the literature (Fig. 4B) (O’Leary and Hood, 2009; Zhao et al., 2007). It is reasonable to speculate that this pronounced
increase in autophagy has such a detrimental effect on the muscle that it is interfering with the ability of ActRIIB to prevent denervation atrophy.

Because Akt and SGK can regulate the pro-growth mTOR pathway, we also examined whether this pathway was dysregulated in our models of atrophy. We found that, although denervation does cause a substantial increase in most components of the mTOR complexes, levels of these proteins were all unchanged in immobilized muscle compared with controls (Fig. 5A). We did observe, however, that both atrophies bring about a loss of activity of the mTORC1 substrate p70S6k. Although that is suggestive that both immobilization and denervation lead to decreased protein synthesis, this is brought about through different mechanisms in the two atrophies. Immobilization leads to a loss p70S6k expression, possibly mediated by the loss of the scaffold protein eIF3f (Csibi et al., 2009; Csibi et al., 2008), but denervation resulted in a loss of p70S6k activation – possibly due to an increase in ULK1 activity (Egan et al., 2011; Lee et al., 2007).

A recent report demonstrated that treating a laminin-deficient mouse model with an mTOR inhibitor reduced autophagy and significantly improved the aberrant muscle phenotype of the mice (Ramos et al., 2012). Similarly, we also treated our denervated mouse model with the mTOR inhibitor rapamycin to test whether the upregulation of mTOR that we observed is a compensatory mechanism or is contributing to the atrophy phenotype. We found that, compared with placebo-treated mice, rapamycin had no effect on the denervation atrophy phenotype despite downregulating mTORC signaling in the atrophic muscle (Fig. 6A,B).

Of great interest were our findings that rictor, more so than raptor, is sensitive to rapamycin treatment in our denervation model (Fig. 6C). This complements several reports suggesting that examination of both the mTORC1 and mTORC2 complexes is necessary to fully understand the \textit{in vivo} effects of rapamycin (Lamming et al., 2012; Sarbassov et al., 2006; Ye et al., 2012). We found this to be the case in our models, because rapamycin completely prevented the denervation-induced upregulation of rictor. In addition, the mTORC2 substrates Akt and SGK also exhibited reduced activation with rapamycin treatment compared with placebo-treated denervated muscle. In fact, when compared with sham-operated controls, rapamycin-treated denervated mice demonstrated no significant upregulation in active pAkt and SGK (Fig. 7A).

Previous research has indicated that several different pathways can contribute to atrophy pathogenesis. For example, upregulation of PGC-1α has been shown to be important for preventing sarcopenic and ALS-induced muscle atrophy, and activation of the IL-6–STAT3 pathway has been linked to the muscle wasting seen in cancer cachexia (Bonetto et al., 2012; Bonetto et al., 2011; Da Cruz et al., 2012).
et al., 2012; Wenz et al., 2009). However, it should be noted that the vast majority of papers indicate that it is the IGF-Akt-FoxO3a pathway that is the most important for the progression of muscle atrophy (Lecker and Goldberg, 2002; Lecker et al., 2004; Sacheck et al., 2007). Although it is well-established that Akt can play a substantial role in muscle hypertrophy and atrophy under certain circumstances, it is becoming clear that denervation is an exception to this rule. Even though some evidence has indicated that increasing Akt activation might be beneficial to the muscle during early stages of denervation atrophy (Bodine et al., 2001b), our results nonetheless outline an important point for future treatment approaches – Akt, mTOR and SGK do not represent promising therapeutic targets for individuals suffering from prolonged denervation. A detailed timecourse assessment of the molecular events that occur as denervation atrophy progresses could clarify how certain proteins and pathways are involved in the pathogenesis. Indeed, we are currently in the process of conducting a cohesive study looking at these various signaling pathways at multiple time points after denervation.

In conclusion, our studies bring two very significant contributions to the field of skeletal muscle atrophy. First, disuse atrophy, but not denervation atrophy, can be prevented by the application of a myostatin inhibitor. Myostatin inhibitors are currently being tested in clinical trials and will soon be available for use by physicians. The knowledge that myostatin inhibition is not a ubiquitous treatment option for all forms of atrophy will aid in providing targeted therapies to individual patients. Secondly, denervation atrophy is independent from Akt, mTOR and SGK activation, and therefore does not follow the usual molecular pathway paradigm for muscle wasting. Our study highlights the fundamental differences in the pathophysiology of disuse and denervation atrophy, and emphasizes the importance of developing unique strategies for their treatment.

MATERIALS AND METHODS

Animal models
Mice were housed in facilities in accordance with the Animal Care and Use Committee of Johns Hopkins University School of Medicine safety protocols. All animals used for experiments were 2-month-old male, C57BL/6 mice (Jackson Laboratories, Bar Harbor, ME, USA). Mice were immobilized by use of a surgical staple (Autosuture Royal 35W stapler, Mansfield, MA, USA) attached to the hind foot held against the tibia as previously described (Caron et al., 2009). Mice were denervated by surgical removal of ~6 mm of sciatic nerve from one hindlimb. Control mice were sham operated in the same way, but without the removal of the nerve. Wounds were closed with surgical staples. The mice were monitored for infection, then anesthetized with isoflurane and euthanized after 3 weeks. Skeletal muscles were dissected and either flash-frozen in liquid nitrogen or imbedded in optimal cutting temperature (OCT) compound (Tissue-Tek, Torrance, CA, USA) for cryostat sectioning. Experiments were repeated at least twice with a minimum of five mice per treatment group. Data shown are averages across all experimental sets. All TA mass measurements are normalized to tibia length.
Drug delivery
Mice were given intraperitoneal injections of soluble ActRIIB at a dosage of 10 mg/kg body weight on days 1, 4, 8 and 15 in a volume of 500 μl PBS. Control and placebo mice were injected with 500 μl PBS on the same days. Soluble ActRIIB was obtained from Dr Se-Jin Lee and was produced as previously described (Lee et al., 2005). Rapamycin (LC Laboratories, Woburn, MA, USA) was dissolved in DMSO and mice were given daily intraperitoneal injections at a dosage of 2 mg/kg or 10 mg/kg body weight diluted to 200 μl in saline solution (saline with 0.2% carboxymethylcellulose and 0.25% Tween80). Control and placebo mice received an equal volume of DMSO diluted to 200 μl in saline solution.

Sectioning and staining
Tissue imbedded in OCT was sectioned with a cryostat (Microm HM 550) into 10-mm sections. Slides were stained by blocking in 5% BSA/PBS then incubated overnight with primary antibody against laminin-1 (Chemicon, Temecula, CA, USA) in 1% BSA/PBS at 4°C. The slides were washed three times for 5 minutes with 1% BSA/PBS, incubated 90 minutes at room temperature with appropriate secondary antibody (Alexa Fluor 488, Invitrogen, Grand Island, NY, USA), washed again three times for 5 minutes with 1% BSA/PBS, and finally mounted with hard set mounting media (Vector Laboratories, Burlingame, CA, USA). Immunofluorescent pictures were taken with an Eclipse i80 microscope (Nikon, Melville, NY, USA) at 10× magnification for myofiber measurement analysis. Myofiber size was determined by measuring the minimum feret diameter (MFD) of 700-1000 fibers for three to five mice per treatment group using Nikon NS elements 2.0 software. Representative pictures of myofiber size were taken at 20× magnification.

Western blots
Flash-frozen tissue was homogenized using a polytron in Total Protein Extraction with protease inhibitors (Roche, San Francisco, CA, USA). Protein concentration was obtained using the Pierce BCA protein assay (Thermo Scientific, Rockford, IL, USA). 20 μg of total protein lysates were separated on 4-12% gradient Bis-Tris mini gels or 18% Tris-Glycine mini gels (Invitrogen, Grand Island, NY, USA) then wet transferred to nitrocellulose membrane. Nitrocellulose membranes were blocked for 30 minutes at room temperature in either 5% BSA/PBST or 5% milk/PBST then incubated with primary antibodies overnight at 4°C. Membranes were then washed twice for 20 minutes in PBST, incubated for 1 hour at room temperature with appropriate secondary antibody, and then washed twice for 20 minutes using PBST. Membranes were then developed with SuperSignal West Femto or Dura secondary antibody, and then washed twice for 20 minutes using PBST. Antibodies were obtained from Cell Signaling, Boston, MA, USA (Smad2 #5339, PERK1/2 #2022720, ERK1/2, p-FOXO3a #3221, mTOR S2448, rictor, raptor, Gβλ, p-4E-BP1 #370, 4E-BP1, pAKT #4513, Akt, ATG7, LC3b, SGK, pULK1 Th575), Novus Biologicals, Littleton, CO, USA (eIF3, Lamp2), Epitomics, Burlingame, CA, USA (pSmad3 S423/S425, Smad3), Invitrogen, Grand Island, NY, USA (pSmad5 S507/506, Abcam, Cambridge, MA, USA (p70S6K, p-p70S6K #2587, Myogenin, FoxO3a, p62, ULK1), Millipore, Billerica, MA, USA (pFOXO3a #3221, p21), ECM Biosciences, Versailles, KY, USA (atrogin-1), and Santa Cruz Biotechnology, Dallas, TX, USA (GAPDH). Secondary antibodies were obtained from GE Healthcare, Pittsburgh, PA, USA (anti-rabbit, anti-mouse, anti-rat) and Sigma-Aldrich, St Louis, MO, USA (anti-goat).

Statistical analysis
All figures represent a comparison of control, placebo immobilized/denervated, and treated immobilized/denervated mice. All figure graphs are expressed as means ± s.e.m. and plotted using Excel. Statistical analyses were done using R statistical software to generate a one-way ANOVA followed by Tukey’s HSD pairwise comparison for P-values. P-value ≤ 0.5×10^-2 was considered significant.

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Competing interests
The authors declare no competing financial interests.

Author contributions
E.M.M. performed all experiments, analysis, and writing of the manuscript. E.A.-M., R. Mejias, J.L.S. assisted in experimental design, data analysis, and writing of the manuscript. E.M.M., R. Mi and J.S.-P. performed the denervation surgeries. S.Y. aided in fiber size measurements. S.-J.L., A.H. and E.M.M. contributed to experiment design and writing of the manuscript. R.D.C. designed and developed the original experimental plan.

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