

# Role of satellite cells versus myofibers in muscle hypertrophy induced by inhibition of the myostatin/activin signaling pathway

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**Myostatin and activin A are structurally related secreted proteins that act to limit skeletal muscle growth. The cellular targets for myostatin and activin A in muscle and the role of satellite cells in mediating muscle hypertrophy induced by inhibition of this signaling pathway have not been fully elucidated. Here we show that myostatin/activin A inhibition can cause muscle hypertrophy in mice lacking either syndecan4 or Pax7, both of which are important for satellite cell function and development. Moreover, we show that muscle hypertrophy after pharmacological blockade of this pathway occurs without significant satellite cell proliferation and fusion to myofibers and without an increase in the number of myonuclei per myofiber. Finally, we show that genetic ablation of *Acrv2b*, which encodes a high-affinity receptor for myostatin and activin A specifically in myofibers is sufficient to induce muscle hypertrophy. All of these findings are consistent with satellite cells playing little or no role in myostatin/activin A signaling in vivo and render support that inhibition of this signaling pathway can be an effective therapeutic approach for increasing muscle growth even in disease settings characterized by satellite cell dysfunction.**

activin receptors | GDF-8 | follistatin

**M**yostatin (MSTN) is a transforming growth factor- $\beta$  family member that acts as a negative regulator of skeletal muscle mass. *Mstn*<sup>-/-</sup> mice exhibit an approximate doubling of skeletal muscle mass throughout the body as a result of a combination of increased numbers of muscle fibers and increased muscle fiber sizes (1). The function of myostatin is highly conserved among mammals: naturally occurring mutations in the *MSTN* gene resulting in increased muscling have been identified in cattle (2–5), sheep (6), dogs (7), and humans (8).

The identification of myostatin and its biological function immediately suggested the possibility that inhibitors of this pathway may have clinical applications for treating patients with muscle loss. Indeed, there has been considerable focus on elucidating the molecular mechanisms underlying myostatin activity, with the goal of identifying strategies for pharmacological intervention. A number of key regulatory components of this signaling system have been identified, including inhibitory extracellular binding proteins, such as follistatin (9), FSTL-3 (10), GASP-1/GASP-2 (11), and the myostatin propeptide (9, 12), as well as myostatin receptors, which include both the type II receptors, ACVR2 and ACVR2B (9, 13), and the type I receptors, most likely ALK4 and ALK5 (14). The elucidation of the myostatin regulatory system has led to the development of a wide range of myostatin inhibitors that are active in vivo, and postnatal elimination of myostatin activity in mice either by systemic administration of these inhibitors (13, 15–18) or by induced muscle-specific deletion of the *Mstn* gene (19) has been shown to cause significant growth of muscle fibers, demonstrating the critical importance of this pathway in limiting muscle growth

in adult animals. Finally, the function of myostatin in muscle seems to be redundant with that of at least one other TGF- $\beta$  family member (13, 20), including activin A (21).

Although the fundamental role of myostatin and activin A in negatively regulating muscle growth has been firmly established, there is considerable debate as to the identities of their cellular targets. A critical question is whether these ligands act in vivo by signaling to satellite cells, which are the stem cells resident in adult muscle, or directly to myofibers. Defining the role of satellite cells in mediating myostatin/activin A signaling and the effects of myostatin/activin A inhibition is important not only for understanding the basic biology of skeletal muscle growth but also for pursuing clinical applications based on targeting this pathway. A major question has been whether therapies based on myostatin/activin A inhibition will have beneficial effects in clinical settings in which satellite cells are largely dormant or exhausted, such as in muscular dystrophy or age-related sarcopenia. In this regard, conflicting results have been reported in a number of studies that have examined the effect of myostatin/activin A loss or inhibition on satellite cells in vivo (22–27). Here, using various combinations of genetic and pharmacological approaches in mice, we examine the contribution of satellite cells to muscle hypertrophy induced by myostatin/activin A inhibition.

## Results

If inhibition of myostatin/activin A activity results in muscle hypertrophy by causing activation and fusion of satellite cells to myofibers, one prediction is that the effect of blocking this pathway would be attenuated in mice in which satellite cells are defective. To test this prediction, we examined two mutant strains that have previously been reported to have defects in muscle regeneration.

We first examined the effect of blocking myostatin/activin A signaling in mice lacking syndecan4 (*Sdc4*). *Sdc4* is expressed by satellite cells (28), and *Sdc4*<sup>-/-</sup> mice are relatively healthy with

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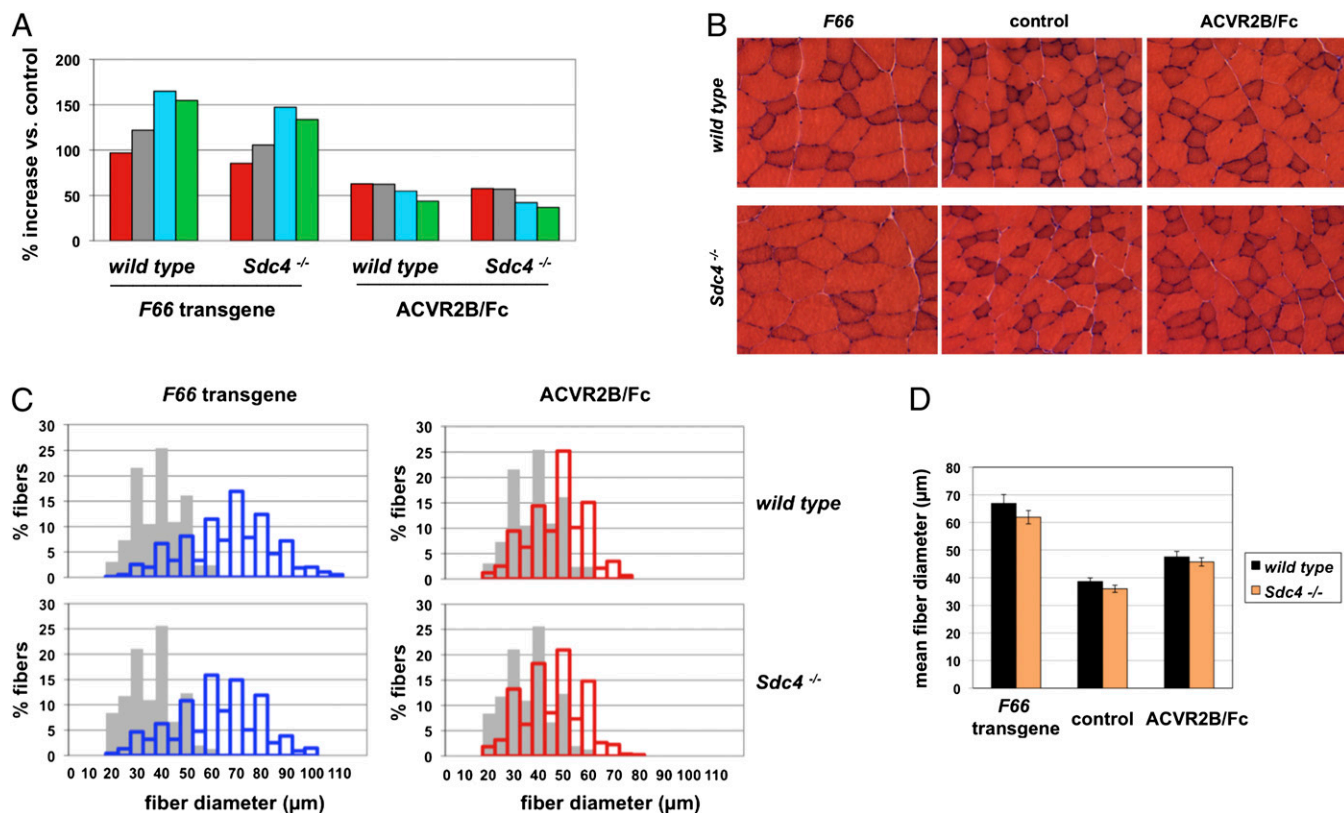
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**Fig. 1.** Effect of blocking the myostatin/activin A pathway in *Sdc4*<sup>-/-</sup> mice. (A) Percentage increase in weights of the pectoralis (red), triceps (gray), quadriceps (blue), and gastrocnemius/plantaris (green) muscles of WT and *Sdc4*<sup>-/-</sup> mice resulting either from the presence of the *F66* transgene or from administration of ACVR2B/Fc (10 mg/kg for 4 wk). Actual weights, SEs, and *P* values are shown in Table 1. (B) Hematoxylin and eosin-stained sections prepared from gastrocnemius muscles showing the hypertrophy induced by the *F66* transgene and by ACVR2B/Fc. (C) Distribution of fiber sizes in gastrocnemius muscles of mice carrying the *F66* transgene (blue bars) or injected with ACVR2B/Fc (red bars) compared with control mice (gray bars) either lacking the transgene or injected with vehicle. (D) Mean fiber diameters in the gastrocnemius muscles.

normal body weights (29) but exhibit a severe defect in muscle regeneration after chemical injury owing to a failure of satellite cell activation (30). We used both genetic and pharmacological approaches to determine the effect of blocking myostatin/activin A signaling in *Sdc4*<sup>-/-</sup> mice. For the genetic approach, we examined the effect of overexpressing the myostatin/activin A inhibitor follistatin by crossing *Sdc4* mutants to *F66* transgenic mice, which express follistatin under the control of a myosin light chain promoter and enhancer (20). We elected to use the *F66*

transgene rather than the *Mstn* deletion mutation in these studies for two reasons. First, we showed previously that whereas increases in both fiber numbers and fiber sizes contribute significantly to increased muscling in *Mstn*<sup>-/-</sup> mice (1), the increases in muscle mass seen in *F66* transgenic mice result almost entirely from muscle fiber hypertrophy (20). Second, because follistatin has a relatively broad range of specificity in that it is capable of blocking multiple ligands, including both myostatin and activin A, we reasoned that the *F66* transgenic approach would allow us

**Table 1.** Effect of the follistatin transgene (*F66*) and the soluble ACVR2B receptor (ACVR2B/Fc) on muscle weights (mg) in *Sdc4* mutant mice

Variable	Pectoralis	Triceps	Quadriceps	Gastrocnemius
<i>Sdc4</i> <sup>+/+</sup> (n = 11)	72.6 ± 1.6	94.2 ± 2.0	190.6 ± 3.7	134.9 ± 2.3
<i>Sdc4</i> <sup>+/+</sup> + <i>F66</i> (n = 13)	143.0 ± 4.3*	209.1 ± 9.5*	504.9 ± 17.6*	343.7 ± 9.6*
Effect of <i>F66</i> (%)	+96.9	+122.0	+164.9	+154.7
<i>Sdc4</i> <sup>-/-</sup> (n = 12)	71.2 ± 1.9	92.3 ± 2.1	182.4 ± 3.7	128.2 ± 2.2
<i>Sdc4</i> <sup>-/-</sup> + <i>F66</i> (n = 14)	131.9 ± 4.8*	189.9 ± 8.4*	450.9 ± 18.5* <sup>†</sup>	299.5 ± 9.3* <sup>‡</sup>
Effect of <i>F66</i> (%)	+85.4	+105.7	+147.2	+133.7
<i>Sdc4</i> <sup>+/+</sup> + PBS (n = 3)	65.3 ± 2.2	86.0 ± 3.1	171.7 ± 8.0	128.3 ± 4.5
<i>Sdc4</i> <sup>+/+</sup> + ACVR2B/Fc (n = 3)	106.3 ± 7.6 <sup>§</sup>	139.7 ± 10.4 <sup>§</sup>	265.7 ± 17.0 <sup>§</sup>	184.3 ± 14.9 <sup>§</sup>
Effect of ACVR2B/Fc (%)	+62.8	+62.4	+54.8	+43.6
<i>Sdc4</i> <sup>-/-</sup> + PBS (n = 3)	69.3 ± 2.1	91.0 ± 3.2	188.7 ± 5.9	136.3 ± 2.9
<i>Sdc4</i> <sup>-/-</sup> + ACVR2B/Fc (n = 3)	109.3 ± 2.9 <sup>¶</sup>	143.0 ± 1.4 <sup>¶</sup>	267.7 ± 3.9 <sup>¶</sup>	186.3 ± 7.1 <sup>¶</sup>
Effect of ACVR2B/Fc (%)	+57.7	+57.1	+41.9	+36.7

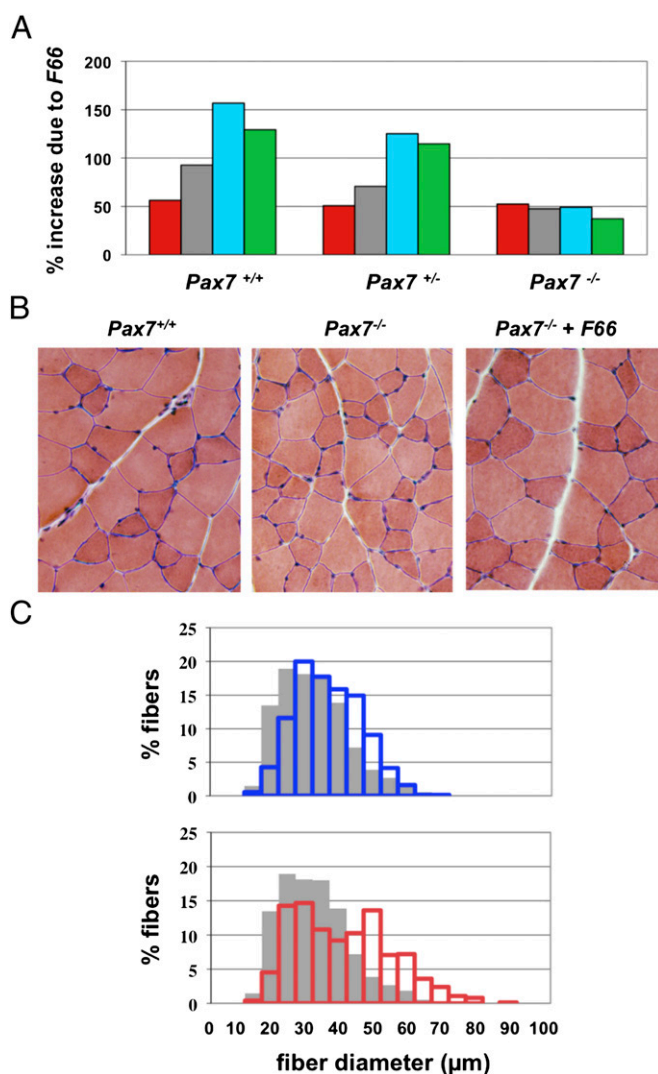
\**P* < 0.001 vs. *Sdc4*<sup>+/+</sup>; <sup>†</sup>*P* < 0.05 vs. *Sdc4*<sup>+/+</sup> + *F66*; <sup>‡</sup>*P* < 0.01 vs. *Sdc4*<sup>+/+</sup> + *F66*; <sup>§</sup>*P* < 0.05 vs. *Sdc4*<sup>+/+</sup> + PBS; <sup>¶</sup>*P* < 0.001 vs. *Sdc4*<sup>-/-</sup> + PBS; <sup>||</sup>*P* < 0.01 vs. *Sdc4*<sup>-/-</sup> + PBS.

**Table 2. Effect of the follistatin transgene (*F66*) on muscle weights (mg) in *Pax7* mutant mice**

Variable	Pectoralis	Triceps	Quadriceps	Gastrocnemius
<i>Pax7</i> <sup>+/+</sup> (n = 12)	81.8 ± 3.4	107.7 ± 4.5	212.3 ± 8.9	144.0 ± 5.7
<i>Pax7</i> <sup>+/+</sup> + <i>F66</i> (n = 13)	127.8 ± 5.5*	207.6 ± 14.2*	545.3 ± 35.0*	330.2 ± 21.1*
Effect of <i>F66</i> (%)	+56.2	+92.8	+156.8	+129.3
<i>Pax7</i> <sup>+/-</sup> (n = 15)	79.5 ± 1.7	108.0 ± 1.9	208.6 ± 3.5	140.7 ± 2.7
<i>Pax7</i> <sup>+/-</sup> + <i>F66</i> (n = 22)	119.9 ± 3.3 <sup>†</sup>	184.5 ± 7.4 <sup>†</sup>	469.5 ± 18.5 <sup>†</sup>	302.0 ± 10.5 <sup>†</sup>
Effect of <i>F66</i> (%)	+50.7	+70.8	+125.1	+114.7
<i>Pax7</i> <sup>-/-</sup> (n = 9)	40.1 ± 2.1*	48.3 ± 1.7*	115.0 ± 6.6*	81.7 ± 4.9*
<i>Pax7</i> <sup>-/-</sup> + <i>F66</i> (n = 13)	61.1 ± 2.5 <sup>‡</sup>	71.3 ± 2.6 <sup>‡</sup>	171.4 ± 5.9 <sup>‡</sup>	112.1 ± 4.0 <sup>‡</sup>
Effect of <i>F66</i> (%)	+52.2	+47.5	+49.0	+37.2

\**P* < 0.001 vs. *Pax7*<sup>+/+</sup>; <sup>†</sup>*P* < 0.001 vs. *Pax7*<sup>+/-</sup>; <sup>‡</sup>*P* < 0.001 vs. *Pax7*<sup>-/-</sup>.

to observe effects of inhibiting the general signaling pathway rather than inhibiting just myostatin.



**Fig. 2.** Effect of the *F66* transgene in *Pax7*<sup>-/-</sup> mice. (A) Percentage increase in weights of the pectoralis (red), triceps (gray), quadriceps (blue), and gastrocnemius/plantaris (green) muscles of WT, *Pax7*<sup>+/-</sup>, and *Pax7*<sup>-/-</sup> mice resulting from the presence of the *F66* transgene. Actual weights, SEs, and *P* values are shown in Table 2. (B) Hematoxylin and eosin-stained sections prepared from gastrocnemius muscles showing the hypertrophy induced by the *F66* transgene in *Pax7*<sup>-/-</sup> mice. (C) Distribution of fiber sizes in gastrocnemius muscles of WT mice (blue bars) and *Pax7*<sup>-/-</sup> mice without (gray bars) or with (red bars) the *F66* transgene.

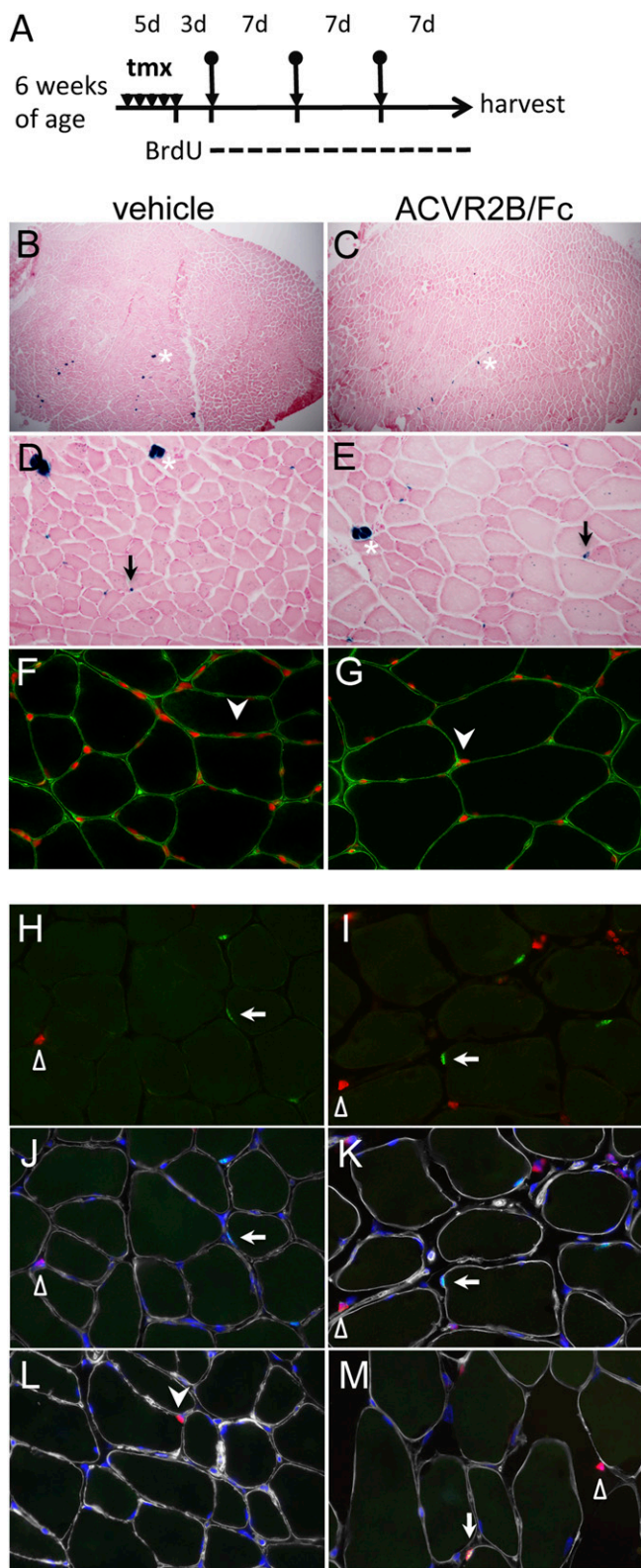
Consistent with our previous report, the *F66* transgene in a WT background caused increases in muscle mass ranging from 97% to 165%, depending on the specific muscle group (Fig. 1A and Table 1), which resulted from dramatic increases in muscle fiber sizes (Fig. 1B and C). Morphometric analysis of the gastrocnemius muscle showed that the mean fiber diameter in *F66* transgenic mice was increased by 73% compared with that in WT mice (Fig. 1D). We observed similar effects of the *F66* transgene in an *Sdc4*<sup>-/-</sup> background (Fig. 1A and Table 1). The magnitude of the increases in muscle weights caused by the *F66* transgene in the *Sdc4*<sup>-/-</sup> background was comparable to that seen in a WT background. Moreover, the extent of muscle fiber hypertrophy induced by the *F66* transgene in *Sdc4*<sup>-/-</sup> mice (Fig. 1B) was similar to that seen in a WT background in terms of the distribution of fiber sizes (Fig. 1C) and in terms of mean fiber diameter (Fig. 1D).

For the pharmacological approach, we examined the effect of blocking myostatin/activin A in adult *Sdc4*<sup>-/-</sup> mice using a soluble form of the activin type IIB (ACVR2B) receptor in which the extracellular ligand binding domain was fused to an Fc domain (ACVR2B/Fc; ref. 13). We showed previously that ACVR2B/Fc is a potent myostatin/activin A inhibitor capable of inducing significant muscle growth when administered systemically to adult mice. As shown in Fig. 1 and Table 1, the ACVR2B/Fc inhibitor induced significant muscle hypertrophy not only in WT mice but also in *Sdc4*<sup>-/-</sup> mice. As in the case of the *F66* transgene, the magnitude of the increases in muscle weights seen in *Sdc4*<sup>-/-</sup> mice (37–58%, depending on the specific muscle group analyzed) and the shift in the distribution in fiber sizes resulting from ACVR2B/Fc administration were comparable to those observed in WT mice (Fig. 1 and Table 1). Hence, muscle hypertrophy induced by myostatin/activin A inhibition either by follistatin or by ACVR2B/Fc is *Sdc4* independent.

It is possible that muscle hypertrophy induced by myostatin/activin A inhibition is dependent on satellite cell activity that is distinct from *Sdc4*-dependent activity required for muscle regeneration after chemical injury. For this reason, we examined the effect of myostatin/activin A inhibition in a mouse strain in which satellite cells are severely depleted, namely, *Pax7*<sup>-/-</sup> mice. Unlike *Sdc4*<sup>-/-</sup> mice, *Pax7*<sup>-/-</sup> mice are considerably smaller than WT mice and mostly die within the first 2 wk after birth (31). Skeletal muscles of *Pax7*<sup>-/-</sup> mice are significantly smaller overall and contain fibers with reduced diameters (32).

For our studies, we used a *Pax7* mutant line that we generated independently by gene targeting. Specifically, we replaced most of the *Pax7* coding sequence with an rtTA tetracycline activator cassette (Fig. S1). Consistent with previous reports, mice homozygous for our *Pax7* mutant allele were severely wasted, and most died before weaning. Some mutant animals on a hybrid background, however, were viable to adulthood, and these mice were small and had muscle weights that were reduced by 43–55% compared with WT mice (Table 2). As in the *Sdc4* studies, we examined the effect of overexpressing follistatin in *Pax7*<sup>-/-</sup> mice





**Fig. 3.** No substantial myofiber incorporation from satellite cells after ACVR2B/Fc-induced muscle hypertrophy. (A) Schema of experimental time line. Bullet-arrows indicate the time of vehicle or ACVR2B/Fc administration; d, day; tmx, tamoxifen. Vehicle and ACVR2B/Fc-treated samples are indicated at top. (B–E) X-gal stained (blue) TA muscle sections from lineage-labeled *Pax7<sup>CE/+</sup>, R26R<sup>LacZ</sup>* mice treated with vehicle (B and D) and ACVR2B/Fc (C and E), counterstained with Nuclear Fast Red; B and C, low magnification; D and E, high magnification; asterisks, spindle myofibers; arrows,  $\beta$ -gal<sup>+</sup>

by crossing in the *F66* transgene. As shown in Fig. 2A and Table 2, the *F66* transgene had a significant effect even in the absence of Pax7, with muscle weights in *F66, Pax7<sup>-/-</sup>* mice being 37–52% higher than those of *Pax7<sup>-/-</sup>* mice. Analysis of muscle sections revealed significant hypertrophy induced by the *F66* transgene (Fig. 2B), and morphometric analysis revealed a clear shift in the distribution of fiber sizes toward larger fibers (Fig. 2C). The *F66* transgene affected fibers throughout the size spectrum. At the lower end of the spectrum, less than 5% of fibers in *F66, Pax7<sup>-/-</sup>* mice had diameters  $\leq 20 \mu\text{m}$  compared with approximately 15% of fibers in *Pax7<sup>-/-</sup>* mice. At the higher end of the spectrum, nearly 36% of fibers in *F66, Pax7<sup>-/-</sup>* mice had diameters  $\geq 50 \mu\text{m}$ , compared with less than 9% of fibers in *Pax7<sup>-/-</sup>* mice. Overall, the relative effect of the *F66* transgene was lower in *Pax7<sup>-/-</sup>* mice than in WT mice, which was not unexpected given the severe muscle phenotype resulting from the absence of Pax7. Nevertheless, inhibition of myostatin/activin A by follistatin clearly caused significant increases in muscle growth in the mutant mice.

These studies demonstrated that inhibition of the myostatin/activin A pathway can lead to significant muscle hypertrophy even in mice severely compromised for satellite cell function (i.e., *Sdc4* mutants) or number (i.e., *Pax7* mutants). To complement these satellite cell deficiency studies, we next investigated the role of satellite cells in mediating the effect of myostatin/activin A inhibition by directly monitoring the behavior and fate of satellite cells after pharmacological blockade of this pathway. For these studies, we used an inducible cell marking-lineage tracing system for adult satellite cells (33). We have previously shown that combining a *Pax7-CreER<sup>T2</sup>* knockin allele (*Pax7<sup>CE</sup>*) with a Cre reporter LacZ allele (*R26R<sup>LacZ</sup>*; ref. 33) allows tamoxifen-inducible permanent marking of PAX7-positive satellite cells, which can be easily identified by X-gal staining. Moreover, lineage tracing of these marked cells revealed that they are capable of fusing to myofibers either during the perinatal period or after injury in adult mice, which can be visualized by X-gal staining in myofibers.

We examined the effect of myostatin/activin A inhibition by treating these mice with ACVR2B/Fc (scheme outlined in Fig. 3A). Three days after tamoxifen-induced cell marking, we initiated three weekly i.p. injections of ACVR2B/Fc and killed the mice 1 wk after the last injection. As expected, systemic administration of the soluble receptor to tamoxifen-treated *Pax7<sup>CE</sup>, R26R<sup>LacZ</sup>* mice caused significant muscle growth of all muscle groups examined (Table S1). We focused our analysis on the tibialis anterior (TA), where this marking-tracing system has been extensively characterized. TA muscles showed a 35% increase in wet weight upon treatment with the soluble receptor. Despite this dramatic muscle hypertrophy, however, we did not observe any significant fusion of satellite cells into the myofibers as assessed by X-gal staining (Fig. 3B–E). Consistent with the very low number of X-gal-stained myofibers, the number of X-gal-stained satellite cells relative to total myofibers was similar in ACVR2B/Fc and PBS injected mice (Table 3).

Thus, both the satellite cell functional deficiency mouse models and the forward satellite cell marking strategy were consistent with myostatin/activin A inhibition-induced muscle hypertrophy requiring minimal input from satellite cells. To further confirm

satellite cells. (F and G) Immunostaining of dystrophin (green) to delineate myofiber boundaries for identifying myonuclei (arrowheads) stained by DAPI (red). (H–M) Immunostaining of PAX7 (green), BrdU (red), and Laminin (white), counterstained with DAPI (blue); H and I, overlaid images of PAX7 and BrdU; J and K, same images further overlaid with Laminin and DAPI; L and M, rare examples of a PAX7<sup>+</sup>BrdU<sup>+</sup> satellite cell and a PAX7<sup>+</sup>BrdU<sup>+</sup> myonucleus, respectively. For H–M, arrows, filled arrowheads, and open triangles indicate satellite cells, myonuclei, and interstitial cells, respectively. Color scheme for antigens is the same as in H–K.

**Table 3. Myonuclei and satellite cells in ACVR2B/Fc-injected mice**

Variable	PBS	ACVR2B/Fc
X-gal stained sections*		
Total myofibers	3,893 ± 411	2,694 ± 480
LacZ+ myofibers	6.0 ± 2.9	5.0 ± 4.0
LacZ+ mononuclear cells	156.3 ± 6.3	96.5 ± 6.8
LacZ+ mononuclear cells/myofiber	0.041 ± 0.003	0.038 ± 0.007
DAPI/dystrophin stained sections <sup>†</sup>		
Total myofibers	1,094 ± 132	823 ± 183
Total myonuclei	580 ± 75	425 ± 76
Myonuclei/myofiber	0.523 ± 0.22	0.511 ± 0.21
% Central nucleated fibers	0.28 ± 0.08	0.26 ± 0.10

\*n = 4 mice per group; quantification of eight representative fields per muscle.

<sup>†</sup>n = 3 mice per group; quantification of 15 representative fields per muscle.

these findings, we used two additional methods to assess the extent of satellite cell fusion to myofibers in mice treated with ACVR2B/Fc. First, we quantified numbers of myonuclei and central nucleated fibers by staining sections with both DAPI and dystrophin. As shown in Fig. 3 F and G and Table 3, we found no significant differences in either the number of myonuclei (i.e., DAPI+ nuclei within the dystrophin boundary) per myofiber or the percentage of central nucleated fibers in mice injected with ACVR2B/Fc compared with mice injected with vehicle. Second, we modified the experimental scheme outlined in Fig. 3A by feeding mice with BrdU supplemented in drinking water after the first ACVR2B/Fc injection and throughout the duration of the study. As shown in Fig. 3 H and I, nuclei in which BrdU had been incorporated into the newly replicated DNA were readily detected. To identify newly divided satellite cells, we coimmunostained for PAX7, and to identify BrdU-positive myonuclei (Pax7-negative), we coimmunostained for laminin to outline the myofibers. As shown in Fig. 3 H–M and Table 4, we detected very few BrdU-positive satellite cells and very few BrdU-positive myonuclei in muscles of mice injected with ACVR2B/Fc. Hence, the results of all of these experiments indicate that the soluble receptor caused significant muscle hypertrophy in these mice, seemingly with very little satellite cell recruitment.

Given the lack of evidence for satellite cell contribution to muscle hypertrophy induced by myostatin/activin A inhibition, we tested whether blocking myostatin/activin A signaling in myofibers is sufficient to induce muscle hypertrophy. Specifically, we examined the effect of targeting myostatin/activin A receptors in myofibers. In previous studies we had shown that myostatin is capable of binding the two activin type II receptors, ACVR2 and ACVR2B, but has a higher affinity for ACVR2B (9). ACVR2B has also been shown to be a high-affinity receptor for activin A (34). To ablate ACVR2B function selectively in myofibers, we generated mice carrying a conditional deletion allele for *Acrv2b*, in which we

**Table 4. Numbers of Pax7- and BrdU-positive cells in ACVR2B/Fc-injected mice**

Cells counted	Mononuclear cells			Myofibers	
	Pax7+	BrdU+	Pax7+/BrdU+	Total analyzed	BrdU+
PBS sample 1	102	79	0	1,263	0
PBS sample 2	104	61	0	1,358	1
PBS sample 3	105	62	1	1,417	1
ACVR2B/Fc sample 1	104	438	1	1,353	2
ACVR2B/Fc sample 2	101	294	2	847	1
ACVR2B/Fc sample 3	101	288	1	863	1

flanked exons 2–4 with LoxP sites (Fig. 4A). Our rationale was that removal of exons 2–4 by cre-mediated recombination would delete the entire ligand-binding domain, the transmembrane domain, and part of the cytoplasmic domain and would put the remaining downstream coding sequence out of frame. Hence, deletion of exons 2–4 would almost certainly result in a null allele. We then targeted *Acrv2b* in myofibers by breeding mice carrying the floxed allele to transgenic mice expressing cre recombinase from a myosin light chain promoter and enhancer (*MLC-cre*; ref. 35).

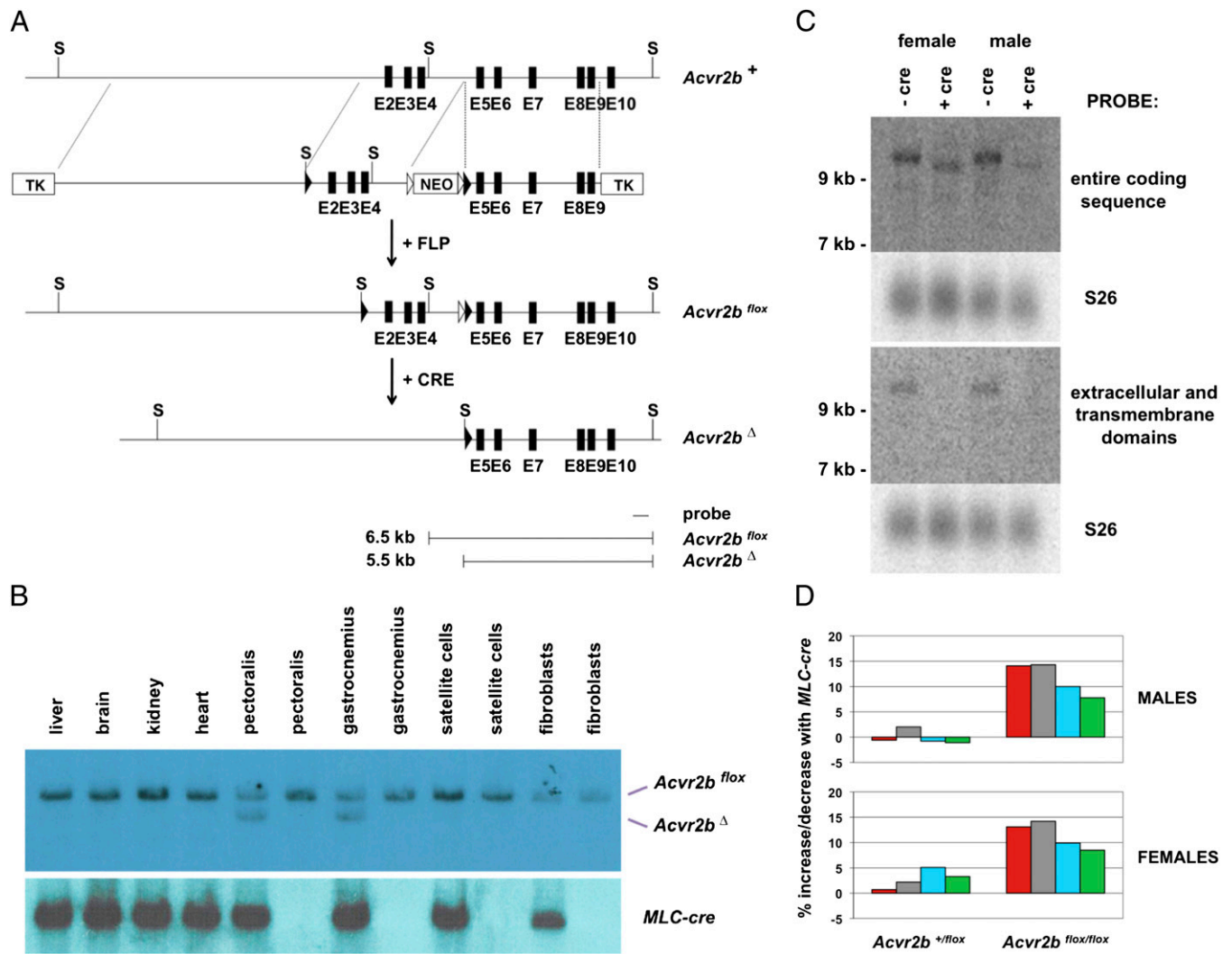
To determine the specificity of the *MLC-cre* transgene in directing recombination of the floxed *Acrv2b* allele, we carried out Southern analysis of DNA isolated from various tissues. Recombination between the LoxP sites was detected only in skeletal muscles of mice carrying the *MLC-cre* transgene (Fig. 4B); in these muscles, approximately half of the genomic DNA had undergone recombination, which is the maximal amount that would be predicted for myofiber-specific targeting, given the known fraction of myonuclei relative to total nuclei in muscle (36). As further confirmation that recombination in muscle was restricted to the myofibers, we freshly isolated satellite cells from these animals (*Materials and Methods*). Using this procedure, we obtained preparations of cells that were 88–89% Pax7-positive. We also harvested fibroblasts during the satellite cell isolation procedure for an additional negative control. As shown in Fig. 4B, no recombination was detected in either satellite cells or fibroblasts from *MLC-cre* mice, supporting that the *MLC-cre* transgene directs efficient recombination specifically in myofibers. We further examined the efficiency of recombination by Northern blot analysis of muscle RNA isolated from *Acrv2b<sup>flx/flx</sup>* mice either with or without the *MLC-cre* transgene. As shown in Fig. 4C, a probe corresponding to the entire coding sequence of ACVR2B detected a single RNA species with an apparent size of greater than 9 kb in *Acrv2b<sup>flx/flx</sup>* mice lacking cre. In *Acrv2b<sup>flx/flx</sup>* mice carrying the *MLC-cre* transgene, the probe detected a smaller RNA with an electrophoretic mobility consistent with cre-mediated deletion of exons 2–4. Moreover, unlike the full-length RNA, the smaller RNA species was not detected using a probe corresponding to the extracellular and transmembrane domains of the ACVR2B. These results demonstrate that the *MLC-cre* transgene was effective in inducing recombination of the *Acrv2b<sup>flx</sup>* allele, and the absence of the full-length transcript in muscles of *Acrv2b<sup>flx/flx</sup>*, *MLC-cre* mice further suggests that myofibers are the primary site of *Acrv2b* expression in muscle.

We then assessed the effect of eliminating ACVR2B in myofibers by comparing muscle weights in *Acrv2b<sup>+/flx</sup>* and *Acrv2b<sup>flx/flx</sup>* mice with and without the *MLC-cre* transgene. As shown in Fig. 4D and Table 5, mice that were homozygous for *Acrv2b<sup>flx</sup>* and that also carried the *MLC-cre* transgene exhibited statistically significant differences in muscle size, with muscle weights being increased by 8–14% in males and 9–14% in females, depending on the specific muscle group. Given that we have previously shown that ACVR2B is functionally redundant with the other type II activin receptor, namely ACVR2, in terms of regulation of muscle growth (13), we presume that targeting both receptors in myofibers would lead to an enhanced response. Nevertheless, our data show that eliminating even just ACVR2B function in myofibers is sufficient to induce muscle hypertrophy, consistent with myofibers being a target for myostatin/activin A signaling.

## Discussion

Although considerable progress has been made in elucidating the regulatory and signaling mechanisms of myostatin at the molecular level, there is considerable controversy as to the cell types in muscle that are direct targets of myostatin signaling in vivo. A variety of studies have demonstrated that myostatin can regulate both proliferation and differentiation of myoblasts and satellite cells in culture (22, 37–43), and these findings, taken together with the phenotype of *Mstn<sup>-/-</sup>* mice, have led to the





**Fig. 4.** Effect of targeting *Acvr2b* in myofibers. (**A**) Gene targeting strategy. Locations of exons 2–10 (E2–E10) are shown as black boxes, FRT (Flippase recognition target) sites are denoted by open triangles, and LoxP sites are denoted by filled triangles. The neo cassette was removed by crossing mice carrying the targeted allele with transgenic mice expressing FLP (Flippase) recombinase in the germline, resulting in an *Acvr2b*<sup>flox</sup> allele. Recombination between the LoxP sites results in a deletion allele (*Acvr2b*<sup>Δ</sup>) lacking exons 2–4. Southern blot analysis of genomic DNA digested with *ScaI* (S) and hybridized using the probe indicated is predicted to give 6.5-kb and 5.5-kb fragments for the *Acvr2b*<sup>flox</sup> and *Acvr2b*<sup>Δ</sup> alleles, respectively. (**B**) Southern blot analysis of genomic DNA isolated from various tissues or cell preparations (as indicated) using either the probe shown in A or a probe for the *MLC-cre* transgene. (**C**) Northern blot analysis of total RNA (25 μg) isolated from the gastrocnemius muscle using probes corresponding either to the entire coding sequence or to the extracellular and transmembrane domains of ACVR2B. Blots were probed with S26 (ribosomal protein) as a loading control. (**D**) Percentage increase or decrease in weights of the pectoralis (red), triceps (gray), quadriceps (blue), and gastrocnemius/plantaris (green) muscles of *Acvr2b*<sup>+flox</sup> or *Acvr2b*<sup>flox/flox</sup> mice resulting from the presence of the *MLC-cre* transgene. Actual weights, SEs, and *P* values are shown in Table 5.

model that a major role for myostatin is to maintain satellite cells in quiescence. Consistent with this model, muscle hypertrophy induced by electroporation of a follistatin expression construct was found to be blunted in mice that had been subjected to local  $\gamma$  irradiation, implying a role for satellite cells in this process (24). Indeed, one study reported that muscles of mice lacking myostatin have increased numbers of satellite cells, as well as an increased proportion of satellite cells in an activated state (22), and a second recent study reported that pharmacological inhibition of the myostatin pathway in mice could induce satellite cell proliferation (26). Several other studies, however, reported that the increased muscle mass seen either in *Mstn*<sup>−/−</sup> mice (23), in muscles electroporated with a dominant negative *Acvr2b* expression construct (25), or in mice treated with a myostatin inhibitor (27) occurs in the absence of satellite cell activation and fusion. Moreover, both *in vitro* and *in vivo* studies have sug-

gested that myostatin is capable of acting directly on the myofibers themselves. In particular, purified myostatin has been shown to be capable of inhibiting protein synthesis (39) and reducing myotube diameter (44) when added to differentiated myotubes in culture, and overstimulation of this pathway in mice by implantation of myostatin-expressing cells (45) or by electroporation of constructs expressing either myostatin (46) or a constitutively active type I receptor (25) has been shown to induce muscle fiber atrophy.

Here, we have used complementary approaches to investigate further the role of satellite cells in mediating the effects of blocking this pathway *in vivo*. In previous studies, we had shown that muscle growth is regulated by not only myostatin but by multiple members of the TGF- $\beta$  family, including activin A (13, 20, 21). For the studies presented here, we attempted to maximize the effect of blocking this pathway in muscle by using inhibitors and approaches capable of targeting both myostatin and activin A. Specifically, we

**Table 5. Effect of the targeting *Acvr2b* in myofibers on muscle weights (mg)**

Variable	Pectoralis	Triceps	Quadriceps	Gastrocnemius
<b>Males</b>				
<i>Acvr2b</i> <sup>+flox</sup> (n = 23)	71.3 ± 2.1	91.0 ± 2.3	189.1 ± 5.6	129.2 ± 4.1
<i>Acvr2b</i> <sup>+flox</sup> + <i>MLC-cre</i> (n = 28)	70.9 ± 1.7	92.8 ± 2.1	187.6 ± 4.2	127.8 ± 2.7
Effect of <i>MLC-cre</i> (%)	-0.6	+2.0	-0.8	-1.1
<i>Acvr2b</i> <sup>flox/flox</sup> (n = 27)	70.2 ± 1.7	92.4 ± 1.9	191.9 ± 4.4	130.4 ± 2.6
<i>Acvr2b</i> <sup>flox/flox</sup> + <i>MLC-cre</i> (n = 20)	80.1 ± 1.9 <sup>*,†</sup>	105.6 ± 2.3 <sup>†,‡</sup>	211.1 ± 4.8 <sup>*,§</sup>	140.6 ± 3.2 <sup>¶,  </sup>
Effect of <i>MLC-cre</i> (%)	+14.1	+14.3	+10.0	+7.8
<b>Females</b>				
<i>Acvr2b</i> <sup>+flox</sup> (n = 15)	44.2 ± 1.5	66.7 ± 1.7	128.5 ± 2.9	87.2 ± 2.3
<i>Acvr2b</i> <sup>+flox</sup> + <i>MLC-cre</i> (n = 25)	44.5 ± 0.9	68.2 ± 1.1	135.0 ± 2.6	90.1 ± 1.7
Effect of <i>MLC-cre</i> (%)	+0.7	+2.2	+5.1	+3.3
<i>Acvr2b</i> <sup>flox/flox</sup> (n = 21)	42.7 ± 1.2	63.5 ± 1.0	129.6 ± 2.3	86.7 ± 1.4
<i>Acvr2b</i> <sup>flox/flox</sup> + <i>MLC-cre</i> (n = 31)	48.3 ± 0.9 <sup>†,¶</sup>	72.5 ± 0.9 <sup>*,†</sup>	142.4 ± 2.2 <sup>†,‡</sup>	94.1 ± 1.7 <sup>§,¶</sup>
Effect of <i>MLC-cre</i> (%)	+13.1	+14.2	+9.9	+8.5

\**P* < 0.01 vs. *Acvr2b*<sup>+flox</sup>; †*P* < 0.001 vs. *Acvr2b*<sup>flox/flox</sup>; ‡*P* < 0.001 vs. *Acvr2b*<sup>+flox</sup>; §*P* < 0.01 vs. *Acvr2b*<sup>flox/flox</sup>; ¶*P* < 0.05 vs. *Acvr2b*<sup>+flox</sup>; ||*P* < 0.05 vs. *Acvr2b*<sup>flox/flox</sup>.

used a pharmacological approach in which we administered a soluble form of the ACVR2B receptor to adult mice and genetic approaches in which we either overexpressed follistatin as a transgene or ablated the *Acvr2b* gene specifically in muscle. First, we show that muscle hypertrophy induced by myostatin/activin A inhibition can occur in mice that are null for either *Sdc4* or *Pax7*, both of which are deficient in satellite cell activity. Second, we show that pharmacological blockade of the myostatin/activin A pathway using ACVR2B/Fc can cause significant muscle growth with little or no fusion of satellite cells to the growing myofibers and with little or no change in the number of myonuclei per fiber. Finally, we show that blockade of myostatin/activin A signaling specifically in myofibers by genetically targeting the high-affinity receptor, ACVR2B, is sufficient to induce muscle hypertrophy, consistent with myofibers being a direct target for myostatin/activin A signaling. The results of all of these studies suggest that satellite activation and fusion to myofibers do not play a significant role in muscle growth induced solely by inhibition of the myostatin/activin A pathway in adult mice. Additional studies will be required to determine whether the role of myostatin/activin A signaling may be more complex in physiologic settings in which satellite cells are activated by other stimuli, such as after exercise or muscle injury; in such settings, for example, it is possible that inhibition of myostatin/activin A signaling to myofibers may influence satellite cell function either by release of secondary signals that act directly on activated satellite cells or by making the myofibers more permissive for satellite cell fusion.

These findings have direct implications not only for understanding the cellular mechanisms underlying muscle growth but also for targeting the myostatin/activin A pathway for clinical applications. Although inhibitors of this pathway are being pursued as potential therapeutic agents for a variety of different diseases leading to muscle loss, a critical question is whether this strategy will be effective in disease states in which the satellite cell population may already be compromised. For example, for diseases like muscular dystrophy, a widely held view is that although satellite cell activation can compensate for the degenerative process in early stages of the disease, this pool of cells is eventually depleted, thereby leading to an accelerated rate of disease progression at later stages of the disease. Similarly, satellite cell exhaustion has been proposed to be a major contributor to the muscle loss that occurs during aging. Our data indicate that satellite cells are not a primary target for myostatin/activin A signaling and suggest that myostatin and activin A regulate muscle homeostasis predominantly by signaling directly to myofibers. If this is the manner in which these ligands func-

tion, many different diseases affecting muscle should potentially be responsive to inhibition of this pathway irrespective of satellite cell loss or dysfunction in these disease settings.

## Materials and Methods

All animal experiments were carried out in accordance with protocols that were approved by the Institutional Animal Care and Use Committees at the Johns Hopkins University School of Medicine, the Carnegie Institution for Science, or Baylor College of Medicine. *Pax7* and *Acvr2b* targeting constructs were generated from phage clones isolated from a 129 SvJ genomic library. Embryonic stem cell targeting was carried out using either R1 (kindly provided by A. Nagy, R. Nagy, and W. Abramow-Newerly, Samuel Lunenfeld Research Institute, Toronto, Canada) or E14Tg2A (BayGenomics) cells. Blastocyst injections of targeting clones were carried out either by the Johns Hopkins Transgenic Core Facility or by the Baylor Genetically Engineered Mouse Core. *F66* transgenic mice and *Sdc4* mutant mice were maintained on a C57BL/6 background, and *Pax7* mutant and *Acvr2b* flox mice were maintained on a hybrid C57BL/6 and 129/SvJ background. The ACVR2B/Fc fusion protein was expressed in Chinese hamster ovary cells, purified from the conditioned medium using a protein A Sepharose column, and administered by weekly i.p. injections. Control mice were injected with vehicle (PBS). Tamoxifen preparation, storage, and injection regimen were as previously described (33). BrdU was supplemented in the drinking water as previously described (47) immediately after the first ACVR2B/Fc and vehicle injections and for the entire duration.

For measurement of muscle weights, individual muscles from both sides of 10-wk-old mice were dissected, and the average weight was used for each muscle. Morphometric analysis of the gastrocnemius muscle was carried out as previously described (20). For plotting the distribution of fiber sizes, all data for a given genotype were pooled. X-gal histochemistry and immunostaining were carried out as previously described (33). Images were taken under a Nikon E800 scope with a Canon EO5 camera. Immunostaining was carried out with mouse anti-PAX7 (1:10; DSHB), rabbit anti-Laminin (1:2,000; Sigma), mouse anti-dystrophin (1:1,000; Abcam), and sheep anti-BrdU (1:1,000; AbCam) primary antibodies, followed by host-specific secondary antibodies conjugated with Alexa 488, 568, and 647 (all at 1:1,000; Molecular Probes). DAPI (0.4 µg/mL; Sigma) was used to stain nuclei before mounting in Fluoromount G (Southern Biotech). Images were taken under an Axioskope with a monochrome AxioCam. Image processing and quantification were performed by using Photoshop or Metamorph.

Satellite cell isolation was performed essentially as described by Sherwood et al. (48) using collagenase and dispase digestion of hindlimb muscles. Dissociated cells were filtered through 40-µm restrainers and plated onto Matrigel (BD Biosciences) coated dishes for 48 h in culture. They were then trypsinized and consecutively preplated twice on noncoated dish for 30 min each. Attached cells were designated fibroblasts, and the unattached cell fraction was designated satellite cells, both of which were immediately harvested for DNA isolation. A small fraction of satellite cells was plated onto four-well chamber slides coated with Matrigel for 6 h, fixed, and subjected to Pax7 immunostaining as described above to evaluate the enrichment of satellite cells.

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