

Numb-deficient satellite cells have regeneration and proliferation defects

Rajani M. George^{a,1}, Stefano Biressi^{b,1}, Brian J. Beres^a, Erik Rogers^a, Amanda K. Mulia^a, Ronald E. Allen^c, Alan Rawls^a, Thomas A. Rando^{b,d,2}, and Jeanne Wilson-Rawls^{a,2}

^aSchool of Life Sciences, Arizona State University, Tempe, AZ 85287-4501; ^bPaul F. Glenn Laboratories for the Biology of Aging and Department of Neurology and Neurological Sciences, Stanford University School of Medicine, Stanford, CA 94305; ^cSchool of Animal and Comparative Biomedical Sciences, University of Arizona, Tucson, AZ 85721-0038; and ^dNeurology Service and Rehabilitation Research and Development Center of Excellence, Veterans Affairs Palo Alto Health Care System, Palo Alto, CA 94304

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The adaptor protein Numb has been implicated in the switch between cell proliferation and differentiation made by satellite cells during muscle repair. Using two genetic approaches to ablate *Numb*, we determined that, in its absence, muscle regeneration in response to injury was impaired. Single myofiber cultures demonstrated a lack of satellite cell proliferation in the absence of Numb, and the proliferation defect was confirmed in satellite cell cultures. Quantitative RT-PCR from Numb-deficient satellite cells demonstrated highly up-regulated expression of p21 and Myostatin, both inhibitors of myoblast proliferation. Transfection with Myostatin-specific siRNA rescued the proliferation defect of Numb-deficient satellite cells. Furthermore, overexpression of Numb in satellite cells inhibited Myostatin expression. These data indicate a unique function for Numb during the initial activation and proliferation of satellite cells in response to muscle injury.

myogenesis | stem cell | skeletal muscle | conditional mutation

Satellite cells represent a muscle-specific stem cell population that allows for muscle growth postnatally and is necessary for muscle repair (1). In response to muscle-fiber damage, quiescent satellite cells that lie along the myofibers under the plasma-lemma are activated and proliferate. Proliferating satellite cells have a binary fate decision to make—they can differentiate into myoblasts and intercalate into myofibers by fusion to repair the damaged muscle or they can renew the satellite cell population and return to a quiescent state (2–4). Quiescent satellite cells express paired box 7 (*Pax7*), but low or undetectable levels of the myogenic regulatory factors *Myf5* and *MyoD* (5, 6). Activated satellite cells robustly express *Pax7* and *MyoD/Myf5*, but a subset will subsequently down-regulate the myogenic regulatory factors in the process of satellite cell self-renewal (7). Recent studies have demonstrated that, in vivo, *Pax7*-positive cells are necessary for muscle repair (8, 9).

Notch signaling is an important regulator of satellite cell function; it is implicated in satellite cell activation, proliferation (2, 10, 11), and maintenance of quiescence (12, 13). Expression of constitutively active *Notch1* results in maintenance of *Pax7* expression and down-regulation of *Myod/Myf5* whereas inhibition of Notch signaling leads to myogenic differentiation (10, 14). In fact, conditional ablation of *Rbpj* embryonically results in hypotrophic muscle (15), and, if ablated in the adult, satellite cells undergo spontaneous activation and precocious differentiation with a failure of self-renewal (12, 13). In adult muscle, the Notch ligand, Delta-like1 (*Dll1*), is expressed on satellite cells, myofibers, and newly differentiating myoblasts and is necessary for repair (10, 11, 16). In aged muscle, impairment of regeneration is due, in part, to a failure of *Dll1* expression (17).

Numb encodes four proteins with molecular masses of 65, 66, 71, and 72 kDa by alternative splicing of two exons (18, 19). The Numb proteins are cytoplasmic adaptors that direct ubiquitination and degradation of Notch1 by recruiting the E3 ubiquitin ligase Itch to the receptor (18–22). Numb is a cell-fate determinant that mediates asymmetric cell division, leading to selective Notch

inhibition in one daughter cell and its subsequent differentiation whereas the other daughter has active Notch signaling and remains proliferative (10). Embryonically, Numb is expressed in the myotome whereas Notch1 is limited to the dermomyotome (23, 24). This pattern suggests that the expression of Numb in one daughter cell allows entry into the myogenic lineage. Indeed, overexpression of Numb embryonically increases the number of myogenic progenitors in the somite (25, 26).

Numb expression increases during the activation and proliferative expansion of satellite cells, becoming asymmetrically segregated in transit-amplifying cells and leading to asymmetric cell divisions (10, 27). These observations led to a model in which Numb inhibits Notch signaling in one daughter satellite cell, allowing it to undergo myogenic differentiation. The molecular switch that controls the decision of satellite cell progeny to continue proliferating or to differentiate is not well understood. This process seems to be controlled by a decrease of Notch signaling due to increased expression of Numb and an increase in Wnt signaling (10–14, 17, 28). In these studies, we examined the role of Numb in satellite cell function by genetic deletion of Numb from myogenic progenitors and satellite cells. Our observations reveal that Numb is necessary for satellite cell-mediated repair. Furthermore, Numb-deficient satellite cells have an unexpected proliferation defect due to an up-regulation of Myostatin. These data indicate a unique role for Numb in regulating the activation and proliferation of satellite cells.

Significance

This study discloses a role for Numb in the activation and proliferation of adult muscle satellite cells and a unique function in the regulation of the muscle mass determinant Myostatin. Using two different genetic approaches to ablate *Numb*, one that ablated *Numb* in the myogenic lineage developmentally leading to reduced muscle mass. We determined that, in Numb-deficient muscle, regeneration was impaired, there was reduced stem cell proliferation, and there was an up-regulation of Myostatin. Overexpression of Numb suppressed Myostatin expression, and Myostatin-specific siRNA rescued the proliferation defect. These studies increase our knowledge of the signaling pathways involved in stem cell function and raise the possibility of regulating the Numb/Myostatin balance as a therapeutic approach to enhance muscle regeneration.

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¹R.M.G. and S.B. contributed equally to this work.

²To whom correspondence may be addressed. E-mail: Jeanne.Wilson-Rawls@asu.edu or rando@stanford.edu.

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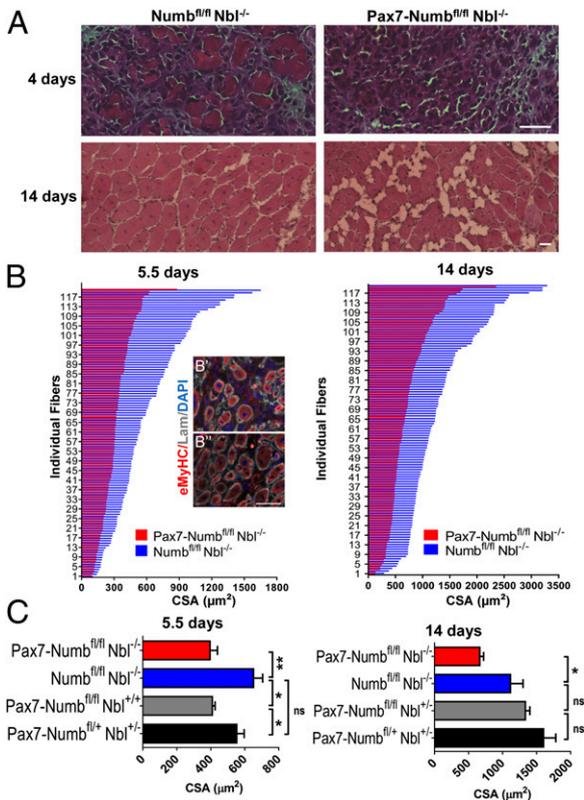


Fig. 1. Reduced muscle mass in mice lacking Numb in the Pax7-lineage after injury. (A) TA muscles from Pax7-Numb^{fl/fl} Nbl^{-/-} and Numb^{fl/fl} Nbl^{-/-} adult mice isolated 4 or 14 d after BaCl₂ injury and H&E stained. (Scale bars: 50 μm.) (B) CSA of regenerating fibers at 5.5 and 14 d post BaCl₂ injury in muscles from Pax7-Numb^{fl/fl} Nbl^{-/-} and Numb^{fl/fl} Nbl^{-/-} littermates. Representative sections of Pax7-Numb^{fl/fl} Nbl^{-/-} and Numb^{fl/fl} Nbl^{-/-} TA muscles at 5.5 dpi stained with antibodies against eMyHC and Laminin (Lam) respectively in B' and B''. Nuclei were DAPI stained. (Scale bars: 50 μm.) (C) The mean of the average CSA calculated (n > 3) at 5.5 and 14 dpi. Pax7-Numb^{fl/fl} Nbl^{-/-} showed a statistically significant reduction in regenerating fiber size at both time points. When Numb, but not Nbl, was ablated, a significant reduction in fiber size was observed at 5.5 dpi but not at 14 dpi. Data are the mean ± SEM. *P < 0.05; **P < 0.01; ns, not significant.

Results

To determine whether Numb plays a critical role in muscle regeneration, it was genetically excised in the myogenic lineage using mice with Cre recombinase inserted in the Pax7 locus (Pax7^{ICNm}) (29) interbred with mice with a floxed Numb allele that deletes exons 4 and 5 (30) and a Numbl-like (Nbl) null mutation (31, 32) (Fig. S1) (we will refer to the Pax7^{ICNm/+} Numb^{fl/fl} Numbl-like^{-/-} mice as Pax7-Numb^{fl/fl} Nbl^{-/-}). When bred into the R26R^{YFP} line (33), the recombination frequency in single-fiber culture was ~100% of Pax7-positive (Pax7^{+ve}) cells, based on YFP expression (Fig. S2 A and B). The high level of recombination was confirmed by FACS analysis detecting YFP^{+ve} cells in the Vcam^{+ve}/Cd45^{-ve}/Cd31^{-ve}/Sca1^{-ve} population isolated from muscle pre- and post-BaCl₂ injury (Fig. S2 C and D). The Pax7-Numb^{fl/fl} Nbl^{-/-} mice weighed significantly less than Numb^{fl/fl} Nbl^{-/-} or Pax7-Numb^{fl/+} Nbl^{+/-} mice at 1 mo of age (Fig. S3). Also, the tibialis anterior (TA) muscles weighed significantly less, and their myofibers had a smaller average cross-sectional area (CSA) than controls (Fig. S3 C–F), indicating a potential role for Numb in the growth of muscle.

TA muscles of Pax7-Numb^{fl/fl} Nbl^{-/-} and Numb^{fl/fl} Nbl^{-/-} mice were injured with BaCl₂ and harvested at different time points. At 5.5 d postinjury (dpi), there were significantly fewer embryonic myosin heavy chain (eMyHC) positive fibers detected in the

Pax7-Numb^{fl/fl} Nbl^{-/-} muscle than in Numb^{fl/fl} Nbl^{-/-} controls (Fig. 1B). At 14 dpi, Numb/Nbl-deficient muscle demonstrated fatty deposition, increased endomysial connective tissue, and infiltration of fibrotic cells, but controls healed with no visible fibrotic or degenerative changes (Fig. 1A). Consistently, the CSA of regenerating fibers was significantly smaller in Pax7-Numb^{fl/fl} Nbl^{-/-} TA muscles than in Numb^{fl/fl} Nbl^{-/-} muscles (Fig. 1C). These differences persisted 2 mo postinjury (Fig. S4). These data indicate that the loss of Numb in the Pax7-lineage, including satellite cells, results in a defective repair response in adult muscle.

The observed deficits in myofiber size, muscle, and body weight may be due to the fact that Numb regulates the size of the pool of myogenic progenitor cells during development (25, 26). It was possible that a developmental deficit affected postnatal repair; therefore, mice with a ubiquitously expressed, tamoxifen (TMX)-inducible Cre recombinase transgene (CAGG ER-Cre, referred to as ER) (34, 35) were bred to mice with floxed alleles of Numb, deleting exon 1, and Numbl-like, deleting exons 3–5, Numb^{tm1zili/tm1zili} Nbl^{tm1zili/tm1zili}, referred to as Nbl^{fl/fl} Nbl^{fl/fl} (Fig. S1) (36, 37). In these mice, Numb is expressed normally during growth, thus avoiding potential developmental effects. Adult mice were treated with TMX, and the quadriceps femoris (QF) muscles were injured with cardiotoxin (CTX) and were examined histologically at 10 dpi. In regenerating muscle from ER-Nbl^{fl/fl} Nbl^{fl/fl} and ER-Nbl^{fl/fl} Nbl^{+/+} mice, there were degenerative changes, including collagen deposition, myofiber degradation, and infiltration of fibrotic cells, that were not evident in ER-Nbl^{+/+} Nbl^{fl/fl}, double heterozygous, or no Cre controls (Fig. 2A–E). The CSA of regenerating myofibers was significantly smaller in ER-Nbl^{fl/fl} Nbl^{fl/fl} muscle than controls, consistent with that observed with Pax7-Numb^{fl/fl} Nbl^{-/-} mice (Fig. 2F). Our data indicate that, regardless of the method of injury, genetic lesion, Cre driver, or muscle examined, Numb-deficient muscle had a defective repair response. Although we cannot completely rule out a role for Numb in other cell types in regenerating muscle, the phenocopy of the defects noted using the ubiquitous and Pax7-specific Cre alleles indicates that Numb is intrinsically necessary for satellite cell function in regenerative myogenesis.

Numb, by virtue of its ability to regulate Notch signaling, has been implicated as part of the molecular switch that controls the decision of satellite cells to continue proliferating or to differentiate (10, 24–26, 28, 38–41). The current model posits that a loss of Numb should result in increased proliferation of the progenitor cells. A lack of differentiating satellite cells would

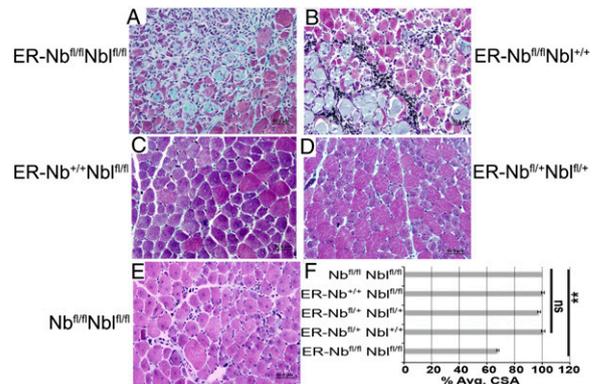


Fig. 2. Impaired regeneration in muscle lacking Numb post-TMX induction of ER-Cre recombinase. (A–E) Representative sections show aberrant repair in Numb-deficient muscle including increased collagen deposition, fibrotic cells, and necrotic myofibers. TMX-treated mice had the left QF injected with CTX, and muscles were harvested 10 dpi and trichrome stained. Genotypes are as indicated (n = 5 per genotype). (Scale bars: 50 μm.) (F) The average CSA as a percentage of control muscle from a minimum of 200 fibers per muscle (n > 5 mice per genotype) at 10 dpi. ER-Nbl^{fl/fl} Nbl^{fl/fl} muscle had a significant decrease in fiber size compared with controls (**P < 0.0001).

result in aberrant repair. To examine proliferation, single muscle fibers were isolated from *Pax7-Numb^{fl/fl} Nbl^{-/-}* and *Numb^{fl/fl} Nbl^{-/-}* mice that were bred with the *R26R^{YFP}* line (Fig. 3A). There was no significant difference in the average number of satellite cells per fiber in Numb-deficient muscles compared with controls at time 0. After 72 h in culture, *Pax7-Numb^{fl/fl} Nbl^{-/-}* fibers unexpectedly had significantly fewer satellite cells than controls (Fig. 3A and B and Fig. S5). When muscles were examined at 4 dpi, there were significantly fewer Pax7⁺ cells in sections of *Pax7-Numb^{fl/fl} Nbl^{-/-}* than in *Numb^{fl/fl} Nbl^{-/-}* mice (Fig. 3C and D). However, the proportion of Pax7⁺ cells that were also Ki67⁺ was the same (Fig. 3D).

The cause of the intriguing loss of satellite cells was tested directly in vitro; mononucleated satellite cells were isolated from QF muscles of *ER-Nb^{fl/fl} Nbl^{fl/fl}* mice and were treated with 4-hydroxytamoxifen (4OH-T) or ethanol, in vitro, then assayed by genomic qPCR to determine the level of *Numb* excision. Only cells that demonstrated recombination levels >90% and were >90% Pax7⁺ (Fig. S6) were used for analysis. In culture, control cells proliferated regardless of treatment, but Numb-deficient cells demonstrated a dramatic decrease in cell number (Fig. 4A and Fig. S7). Daily inspection revealed no obvious cell death, and there was no increase in apoptosis in 4OH-T-treated *ER-Nb^{fl/fl} Nbl^{fl/fl}* cells, as determined by Caspase 3/7 activity (Fig. 4B). In the absence of niche support, Numb-deficient cells did not proliferate. These data reveal an unexpected unique function for Numb in the regulation of satellite cell proliferation.

To determine the molecular basis of the proliferation defect, we compared gene expression in *ER-Nb^{fl/fl} Nbl^{fl/fl}* satellite cells

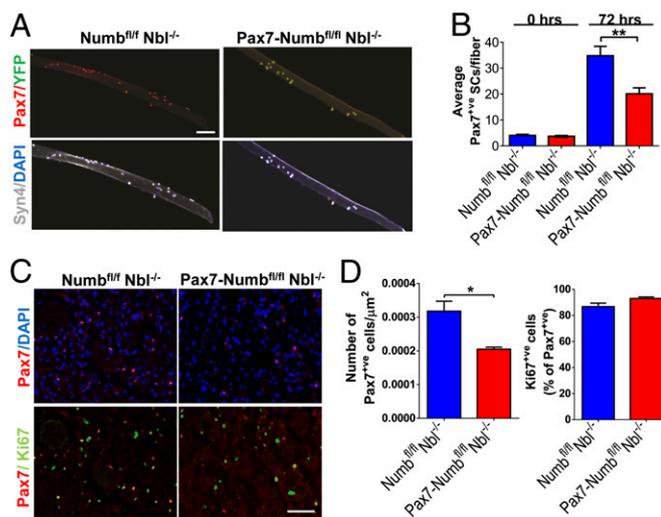


Fig. 3. Decreased numbers of satellite cells in single-fiber cultures and injured muscles from mice lacking Numb in the Pax7-lineage. (A) Single fibers and associated satellite cells from EDL muscles of *Numb^{fl/fl} Nbl^{-/-}* and *Pax7-Numb^{fl/fl} Nbl^{-/-}* mice bred with *R26R^{YFP}* mice were cultured for 66 h and stained with antibodies recognizing Syn4, Pax7, and GFP. Nuclei were DAPI stained. (Scale bar: 100 μ m.) Note that ~98% of the Syn4⁺ cells also express Pax7. A majority of Pax7⁺ satellite cells from *Pax7-Numb^{fl/fl} Nbl^{-/-}* are YFP⁺, but YFP was not detected in *Numb^{fl/fl} Nbl^{-/-}* mice. (B) Pax7⁺ cells were quantified immediately (0 h) or after 72 h of culture (50 fibers per time point). Significantly fewer satellite cell progeny, $P < 0.001$, were found on the fibers from the *Pax7-Numb^{fl/fl} Nbl^{-/-}* muscle after 72 h. (C) Pax7 and Ki67 IF of TA muscles from *Numb^{fl/fl} Nbl^{-/-}* and *Pax7-Numb^{fl/fl} Nbl^{-/-}* mice 4 d after BaCl₂ injury. Nuclei were stained with DAPI. (Scale bar: 50 μ m.) (D) The average number of Pax7⁺ cells per μ m² and the percentage of Pax7⁺ cells that are Ki67⁺ were quantified 4 d after BaCl₂ injury. At least four fields were evaluated for each genotype ($n = 3$ experiments). There are fewer Pax7⁺ cells in *Pax7-Numb^{fl/fl} Nbl^{-/-}* muscle, but a similar proportion are proliferating (Ki67⁺), in both *Pax7-Numb^{fl/fl} Nbl^{-/-}* and control muscle at this time point.

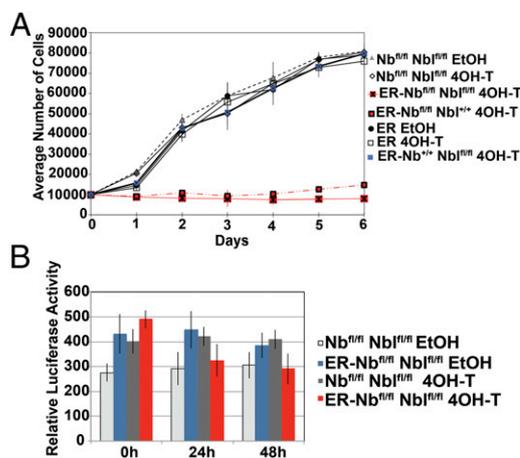


Fig. 4. Numb-deficient satellite cells have a proliferation deficit. (A) The loss of *Numb* resulted in a static cell number. Satellite cells were plated in growth medium after treatment with 4OH-T or ethanol; genotypes are as indicated. Triplicate wells were trypsinized daily, stained with 0.4% Trypan Blue, and counted. Data are mean number of cells \pm SD ($n = 3$). (B) Caspase 3/7 activity showed that Numb-deficient satellite cells had no increase in apoptosis in culture. *ER-Nb^{fl/fl} Nbl^{fl/fl}* and *Nb^{fl/fl} Nbl^{fl/fl}* satellite cells after 4OH-T or ethanol treatment; data are mean Luciferase levels \pm SD ($n = 4$).

treated with vehicle or 4OH-T. We were specifically interested in the expression of Notch signaling pathway genes given the well-known role of Numb in Notch regulation. Our analysis showed that there was no significant difference in the expression of Notch1, or its ligand Dll1. Also, there was only a minor increase in the expression of the Notch target genes *Hes1* and *Hes6* (Table S1). There was no significant difference in the expression levels of *Myod* and *Pax7*. Intriguingly, the cell-cycle regulator *p21* and *Myostatin* (*Mstn*), both of which inhibit myoblast proliferation (4, 41–50), were significantly up-regulated in 4OH-T-treated cells (Table S1).

Myostatin is necessary for the balance between proliferation and differentiation of muscle progenitors embryonically and postnatally (41–50). *Myostatin* signaling activates expression of *p21*, inhibiting myoblast proliferation (41–43, 48–50). Based on the high level of *Mstn* mRNA in 4OH-T-treated *ER-Nb^{fl/fl} Nbl^{fl/fl}* satellite cells, we next determined whether it was also elevated in vivo. QF muscles of TMX or vehicle-treated mice were injected with CTX, and satellite cells were harvested 5 dpi and analyzed by Western blot. The Numb-deficient satellite cells had high levels of *Myostatin*, which was undetectable in controls (Fig. 5A).

The ability of Numb to affect *Mstn* expression in cells was further examined by overexpression of Numb in primary myoblasts. Cells were transfected with plasmids that express Numb-GFP fusion proteins (AcGFP-Numb) (Fig. S8), and then FACS sorted 2 d posttransfection for GFP (Fig. 5B and C). The expression of *Mstn* and *Hes1* was determined by qRT-PCR comparing GFP⁺ and GFP⁻ fractions. Two different isoforms of Numb (65 and 66 kDa) were evaluated; in both cases, GFP⁺ cells demonstrated significantly less *Mstn* mRNA than control cells (Fig. 5D). These data indicate that Numb can regulate *Mstn* levels. Interestingly, *Hes1* expression was not significantly decreased by increased Numb (Fig. 5D), suggesting that Notch signaling is not the primary target of Numb in proliferating satellite cells. We next determined whether knockdown of *Mstn* would rescue the proliferation defect. Muscle-derived *ER-Nb^{fl/fl} Nbl^{fl/fl}* cells treated in vitro with either 4OH-T or vehicle and transfected with three different *Mstn*-specific siRNAs demonstrated > 85% knockdown, but the scramble siRNA had no effect on *Mstn* levels (Fig. 6A). Treatment of satellite cells with *Mstn*-specific siRNA resulted in increased proliferation of Numb-deficient and control cells (Fig. 6B). These data indicate a previously unrecognized function for Numb in the regulation of *Mstn*

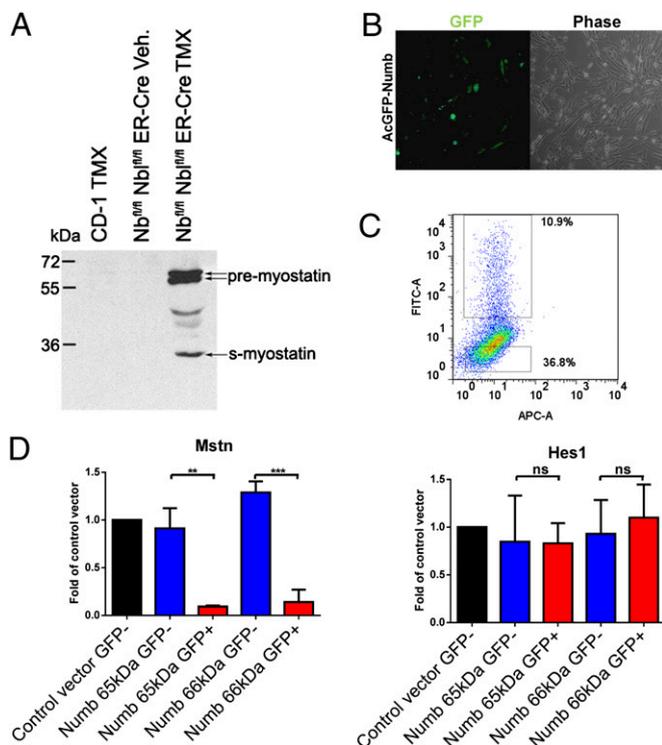


Fig. 5. Numb affects Myostatin expression in myoblasts. (A) Western blot of satellite cells harvested 5 d post-CTX injury shows that only Numb-deficient satellite cells expressed detectable levels of Myostatin. Total protein lysates were analyzed using an anti-Myostatin antibody; genotypes are as indicated ($n = 3$). (B) Micrograph of primary myoblasts 1 d after transfection with AcGFP-Numb (66-kDa isoform). (C) FACS plot 2 d after transfection with AcGFP-Numb (65-kDa isoform). The APC channel was used to evaluate autofluorescence. (D) qRT-PCR of Mstn and Hes1 expression in the GFP^{ve} and GFP^{ve} fractions of myoblasts transfected with AcGFP-Numb. Expression is reported as average fold induction over control vector-transfected cells. A similar trend was observed with two different Numb isoforms ($n = 3$). ** $P < 0.01$; *** $P < 0.001$; ns, not significant.

expression and, as a result, in the control of myogenic progenitor proliferation.

Discussion

Skeletal muscle repair is dependent on the proliferative expansion of satellite cells and the balance between the proliferation of myogenic progenitors and their differentiation into muscle fibers. Numb participates in protein ubiquitination (18–22) and

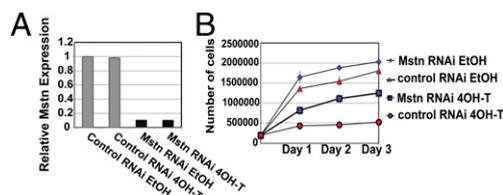


Fig. 6. Mstn knock-down rescued the proliferation defect in Numb-deficient satellite cells. (A) *ER-Nb^{fl/fl} Nb^{fl/fl}* satellite cells that were 4OH-T or ethanol treated were transfected with Mstn-specific or scramble siRNAs. Mstn expression was determined by qRT-PCR and Mstn-specific siRNAs knockdown expression by 85%. Data are relative gene expression \pm SD ($n = 3$). (B) Proliferation assays demonstrate that Mstn-specific siRNA rescues the proliferation defect. *ER-Nb^{fl/fl} Nb^{fl/fl}* satellite cells were 4OH-T or ethanol treated and transfected with Mstn or scramble siRNA. Daily, triplicate wells were counted; data are average cell number \pm SD, ($n = 3$).

is a regulator of binary cell fate (51). The Numb expression pattern in activated satellite cells predicted a role for this gene in regulating skeletal muscle repair. We directly tested this role by conditionally deleting *Numb* to examine the impact that a lack of Numb has on satellite cell function and muscle regeneration. Numb-deficient muscles demonstrated defective myofiber repair (Figs. 1 and 2), significantly fewer satellite cells were found on single fiber cultures (Fig. 3 and Fig. S5), and isolated satellite cells that had Numb deleted in vitro had a proliferation defect (Fig. 4 and Fig. S7). The loss of only *Nbl* did not result in aberrant repair or proliferation. The current model of Numb function in adult muscle is that it regulates asymmetric division of satellite cell progeny and the decision point between differentiation or continued proliferation (10, 16, 40, 51, 52). Our studies show that Numb also has a role much earlier than this cell-fate decision: controlling activation-associated proliferative expansion of satellite cell progeny.

Numb can negatively regulate Notch signaling (20–22, 32, 36, 53–56). However, in keeping with the observation that Notch signaling was not significantly affected in the somite of Numb-overexpressing transgenic mice (25), we found very limited changes in the expression of Notch genes in the myogenic cells isolated from Numb KO mice, leading us to screen for alternative mediators. Our data suggest that Mstn is regulated by Numb. After injury, Numb-deficient muscle demonstrated high levels of Mstn compared with control muscle. Overexpression of two Numb isoforms, p65 and p66, decreased Mstn levels. Mstn-specific siRNA rescued the proliferation deficit in Numb-deficient satellite cells. These data demonstrate that the increased expression of Mstn is responsible for the lack of proliferative expansion of Numb-deficient satellite cells. Mstn is an important regulator of postnatal myogenesis; it is expressed in quiescent satellite cells and inhibits their activation (41, 43). It also inhibits myoblast proliferation, inducing differentiation (42, 44, 48–50). *Mstn* loss-of-function mutations in mice, cattle, and sheep result in dramatically increased muscle mass (57–62). *Mstn*^{-/-} mice demonstrate enhanced muscle repair, even in senescent animals (63, 64).

Recent work has demonstrated that Mstn regulates myoblast differentiation through the Notch pathway (43), and it was proposed that a balance between Notch and Smad3, the downstream effector of Mstn, regulated regenerative competence of satellite cells (43). Furthermore, in aging satellite cells, this balance is changed, leading to impaired regeneration (38). This interaction is likely altered in the absence of Numb, but, as there was no significant change in the expression levels of the canonical Notch genes, it is difficult to conclude that Notch is the target that results in the proliferation defect. It is possible that Mstn governs satellite cell proliferation through still undefined mechanisms not relying on changes in the Smad3/Notch balance. The recent observation that the atrophy seen in *Smad3*^{-/-} muscles could be due to altered Mstn levels supports this idea (65). Mstn may signal independently of Smad3 via either Smad2 or other pathways such as the phosphatidylinositol 3-kinase Wnt and c-Jun N-terminal kinase pathways (66–68).

The molecular mechanisms used by Numb to regulate Mstn expression remain to be defined. In mammals Numb regulates several signaling pathways with different modalities, mainly affecting the ubiquitin and endocytosis network (54). For example, Numb regulates differentiation of cerebellar granule cell progenitors (GCPs) by targeting Gli1, an effector of Hedgehog signaling, to the proteasome, allowing GCPs to differentiate (69). Numb interacts with Mdm2 to hamper ubiquitination of p53 (70). Numb interacts with the endocytic machinery to regulate EGF, transferrin (71), and integrin (72) trafficking. Numb binds to activated TrkB, a receptor for BDNF, and causes endocytic recycling of this receptor (73). Numb also acts as a scaffold for aPKC in the cytoplasm, promoting both BDNF-dependent activation of aPKC and neural precursor cell migration (73). Numb interacts with E-cadherin and regulates its localization. Similarly, knock-down of Numb in MDCK cells destabilizes E-cadherin-based cell adhesion and potentiates sensitivity to hepatocyte growth factor

(HGF) (74, 75). Intriguingly, we recently reported that high concentrations of HGF inhibit satellite cell proliferation by inducing Mstn (76). This observation opens the possibility that HGF-dependent signaling could link Numb to Mstn expression.

Androgens and anabolic steroids reduce muscle loss caused by immobilization and spinal-cord injury. Interestingly, recent reports show that the steroid Nandrolone, which counteracts denervation-dependent atrophy, induces increased Numb expression in denervated muscles (77). Mstn up-regulation has been observed in gastrocnemius muscles after denervation and is therefore implicated in the associated atrophy (78, 79). This finding, along with the observation reported here that Numb overexpression reduces Mstn levels, opens the possibility that the beneficial effects of Nandrolone are mediated by a change in the Numb/Mstn balance. Increased levels of Mstn (80, 81) and decreased expression of Numb (82) have been observed during aging, suggesting that this balance could play a role in age-dependent sarcopenia. These findings suggest that a link between Numb and Mstn could control biological events beyond the regulation of the regenerative capacity of satellite cells after injury.

Materials and Methods

Mice. *Numblike* (*Nbl*^{-/-}) (31, 32) and conditional *Numb* mutant (*Numb*^{fl/fl}) (30) mice were provided by W. Zhong (Yale University, New Haven, CT). *Pax7*-Cre mice (*Pax7*^{Cre/m}) (29) were provided by C. Keller (Oregon Health and Science University, Portland, OR). R26R^{YFP} (33), *Numb*^{tm1Zll/tm1Zll}, *Numb*^{tm1Zll/tm1Zll} (36), and the TMX-inducible Cre recombinase line *CAGG ER-Cre* (34, 35) were obtained from The Jackson Laboratory. Mice were housed and experiments were done, under protocols approved by the Institutional Animal Care and Use Committees at the Veterinary Medical Unit at the VA Palo Alto Health Care Systems and Arizona State University and in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals.

Treatment with BaCl₂, TMX, and CTX. Adult (≥60 d old) mice were injected intraperitoneally with 0.4 mg of TMX (Sigma-Aldrich) dissolved in corn oil twice a day for 5 d. After 28 d, QF muscles were injected with 10 μM CTX (Calbiochem) with 0.02% India ink suspension to mark the injection site. For BaCl₂ injury (Sigma-Aldrich), adults were anesthetized by isoflurane inhalation. TA muscles were injected with 50 μL of BaCl₂ solution (1.2% in sterile 0.9% NaCl). Muscles were harvested from age- and sex-matched control and mutant animals.

Morphometry, Histology, and Immunofluorescence. Muscles were flash frozen in isopentane, and midbelly cryostat sections (8 μM) were stained with hematoxylin/eosin (H&E) or with antibodies against Laminin (Sigma-Aldrich), eMyHC (DHSB), GFP (Aves), MyoD (Becton Dickson), Pax7 (DHSB), and Ki67 (Becton Dickson), as described (83). Nuclei were DAPI stained. Technical details are in *SI Materials and Methods*. The area of 120 fibers in four stochastically chosen microscopic fields for each TA/QF was measured using ImagePro Plus software (Media Cybernetics). At least three muscles were analyzed per genotype and time point. Gomori's trichrome stain was done on paraffin-embedded muscle (5 μm) sections at 10 dpi. Details are in *SI Materials and Methods*.

Single-Fiber Isolation and Culture. Single fibers were isolated from EDL muscles as described (28) and fixed immediately or cultured for 72 h in Ham's F-10 with 20% (vol/vol) FBS and 5 ng/mL FGF (Atlanta Biological). Fibers were fixed in 2% paraformaldehyde for immunofluorescence (IF) with antibodies against Syn4 (gift of B. Olwin, University of Colorado, Boulder CO), Pax7, and GFP. Pax7/YFP⁺ cells associated with single fibers in two experiments were quantified for each condition.

Satellite Cell Preparation. Satellite cells were isolated as described (84). Details are in *SI Materials and Methods*. To activate Cre recombinase, cells were treated with 1 μM 4OH-T (Sigma-Aldrich) dissolved in ethanol for 48 h.

FACS Analysis. Satellite cells were isolated from the hindlimb muscles, and FACS was done as described (12), using a FACSAria III (BD Biosciences). Antibodies used to identify the satellite cells recognized CD31, CD45, Sca1, and CD106. Details are in *SI Materials and Methods*. Physical parameters (SSC and FSC) and DAPI dilactate (250 μg/μl; Invitrogen) excluded dead cells.

Proliferation and Caspase 3/7 Assays. Satellite cells were plated at 1 × 10⁵ cells per mL on Matrigel-coated 96-well plates in triplicate for both assays. Daily, cells were trypsinized, stained with 0.4% Trypan Blue (CellGro), and counted using a hemocytometer (*n* = 3). A Caspase-Glo 3/7 assay (Promega) was used to quantify apoptosis per the manufacturer (*n* = 3).

Western Blots. Muscles of TMX- or vehicle-treated mice were injected with CTX, satellite cells were isolated 5 dpi, and total protein lysates were made. Proteins were detected with mouse anti-Mstn (1:1,000; Abcam) and visualized with anti-mouse-AP antibody using ECL substrate (GE Healthcare) (*n* = 3). Technical details are in *SI Materials and Methods*.

Overexpression Studies. Primary myoblasts were isolated from Sv129 mice (Charles River) as described (83). Myoblasts and HEK 293T [American Type Culture Collection (ATCC)] cells were plated on laminin/ collagen-coated wells and transfected with Lipofectamine 2000 (Invitrogen) per the manufacturer. Two days posttransfection, cells were fixed in 4% (wt/vol) paraformaldehyde for IF with antibodies against Numb (1:100, Abcam) and GFP (1:500) or processed for FACS. Mouse Numb cDNAs (65- and 66-kDa isoforms) from C2C12 cells (ATCC) were cloned into pAcGFP-C1 vector (Clontech). The AcGFP-Numb fusion proteins were verified by sequencing.

qRT-PCR. Satellite cell RNA was isolated using TRIzol (Invitrogen). RNA was treated with DNase I before cDNA synthesis using SuperScriptIII (Invitrogen). cDNA was quantified using transcript-specific, intron-spanning primers and Sybergreen I (Eurogentec) on an ABI7900HT thermocycler (Applied Biosystems). Samples were normalized to Gapdh using ΔΔC_t analysis. Data are mean relative expression ± SD.

siRNA. Satellite cells were seeded at 1 × 10⁵ cells per mL and transfected 24 h later with 50 nM Mstn specific or scramble siRNA in 1.5 mL of Lipofectamine RNAiMax (Invitrogen). Total RNA was isolated 3 d posttransfection, and qRT-PCR was done as described.

Statistical Analysis. For morphometry, two-tailed *t* tests were used to calculate the significance of observed differences. For qRT-PCR, body/muscle weight, and number of satellite cells on single fibers, data were analyzed using two-tailed *t* tests with significance of *P* < 0.05 at the 95% confidence level. For caspase and proliferation assays, significance was determined using one-way ANOVA and the post hoc Tukey test.

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