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Novel Therapeutic Strategy for the Prevention of Bone Fractures

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### 14. ABSTRACT

Falls and debilitating bone fractures are a major problem for veterans, and more than 40,000 veterans suffered hip fractures from 2000-2002. Falls are the main etiological factor in more than 90% of fractures, and so treatments that can improve muscle strength while at the same time increasing bone mass will significantly reduce fracture-related morbidity and mortality. Myostatin is a factor that induces muscle wasting and suppresses bone formation. Our data collected thus far demonstrate i) myostatin suppresses proliferation of aged, but not young, myoblasts, ii) myostatin is elevated with age in muscles composed primarily of slow-twitch fibers (e.g. soleus), and iii) myostatin increases muscle mass and muscle fiber size in aged mice. These findings suggest that myostatin inhibitors may have potential for suppressing muscle wasting and improving muscle repair in older individuals, but their effect on bone may be less significant.

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**Introduction**

Loss of muscle mass with age is implicated in age-related bone loss, and muscle frailty contributes to an increased incidence of falls and fractures. Yet, the molecular mechanisms underlying age-related muscle wasting, and the ability of muscle to promote bone formation and fracture healing, are unknown. We have focused our research on the role of myostatin (GDF-8) in muscle-bone interactions in order to develop more effective treatment and prevention strategies for muscle injury, frailty, and bone fracture. We have previously shown that myostatin deficiency increases bone strength and biomineralization throughout the skeleton, and that a new myostatin inhibitor (propeptide) increases both muscle mass and bone formation (Hamrick et al., 2007, 2010; Elkasrawy and Hamrick, 2010). Our research therefore suggests that myostatin is a key factor regulating both myogenesis and osteogenesis. Although some studies have found no association between age and myostatin transcript levels in skeletal muscle (Marcell et al., 2001), others reveal a marked elevation in skeletal muscle myostatin expression with aging in humans (Leger et al., 2008). Additional research indicates that circulating levels of myostatin increase with age in men and women, and are highest in people aged 60-90 (Yarasheski et al., 2002). The latter finding suggests that myostatin is implicated in the sarcopenia of aging, hence myostatin inhibitors are likely to be useful pharmacological agents for treating age-related muscle atrophy as well as bone loss.

The goal of our CDMRP-sponsored research is to better characterize myostatin's role in age-related bone loss, so that targeted therapies to prevent bone fractures by enhancing muscle and bone strength can be developed. We hypothesize that the expression of myostatin and its receptor are elevated with aging in bone and muscle, which antagonizes the osteogenic and myogenic capacity of stem cells in these tissues, but that myostatin inhibitors will reverse this age-related decline in musculoskeletal function. Year 1 of the project was to determine how the expression of myostatin and its antagonist follistatin change with age in musculoskeletal tissues, whereas the goal of year 2 was to determine the effects of myostatin on anabolic pathways in primary bone-derived stromal cells and skeletal muscle myoblasts in vitro. Year 3 studies tested the hypothesis that a myostatin inhibitor could enhance muscle and bone mass in aged animals in vivo. Year 4 (extension) examined the effects of aging on myostatin signaling in human bone marrow stem cells, and the effects of a myostatin inhibitor on behavioral assessments of musculoskeletal function in aged mice.

Findings to date demonstrate that myostatin is elevated in skeletal muscle of aged mice, but only in slow- but not fast-twitch muscles. Myostatin is also elevated in bone marrow from aged mice, whereas in patient samples activin A increases with age more so than myostatin. Myostatin suppressed proliferation of aged, but not young, myoblasts, and also increased the expression of differentiation markers in myoblasts from aged mice. Myostatin did not dramatically alter osteogenic differentiation of either young or aged bone marrow stromal cells; however, myostatin does have a significant inhibitory effect on proliferation of these stem cells. Our new in vivo data using a myostatin inhibitor (propeptide) show that blocking myostatin function in vivo increases muscle mass, grip strength, and fiber size in aged mice, but does not alter bone mass, strength, or parameters of bone formation & resorption. These data suggest that targeting myostatin may have significant therapeutic potential for improving muscle function in older adults, perhaps leading to the prevention of falls and fractures.

**Body**

**Aim 1 (months 1-12). Determine how the expression of myostatin, its receptor, and the myostatin antagonist follistatin change with age in musculoskeletal tissues.**

**Task 1.** Human bone marrow aspirates will be collected from relatively young (18-30) and older (50-70) patients in the MCG orthopaedic clinic. Samples from younger patients are collected as waste by-products during ACL reconstructions, whereas those from older patients are discarded during total knee and hip replacement surgery.
We investigated age-related changes in the activin A-myostatin-follistatin system using bone marrow samples from young (<50 years, n=7) and older (>70 years, n=10) knee arthroplasty patients. Supernatant samples were analyzed using ELISA. Results indicate that follistatin and myostatin levels are not significantly altered with age in human bone marrow supernatants, whereas activin A levels increased increased by more than 120% in human bone marrow (Fig. 1). The marked increase in activin A levels with age in the patient samples was associated with a similar increase in the activin A: follistatin ratio.

![Figure 1](chart1.png)

**Figure 1.** Protein levels (right) of activin A, myostatin, and follistatin in bone marrow supernatants from young (<50 years) and older (>70 years) knee arthroplasty patients determined using ELISA assays.

We also examined gene expression in bone marrow-derived stem cells to explore age-associated changes in expression of the myostatin receptor (ActRIIB), to determine how this may impact sensitivity of bone cells to the factors shown in Figure 1. Results indicate that, consistent with the data shown below in Figure 4, expression of the myostatin (and activin) receptor increases with age in BMSCs (Fig. 2).

![Figure 2](chart2.png)

**Figure 2.** Relative gene expression in BMSCs harvested from young (<50 years) and older (>70 years) knee arthroplasty patients (n=3 per group) determined using real-time PCR.

Task 1 status: Complete.

**Task 2.** Bone aspirates and muscle samples are collected from mice 12, 18 and 24 months of age, 15 mice per age group (total = 45 mice).

Protein levels of activin A, myostatin, and follistatin in skeletal muscle and bone marrow of young and aged mice were determined using ELISA. We excluded the 18 mo age group because we found in PCR assays that this group was consistently an outlier. ANOVAs performed on activin A and follistatin normalized for total protein, and the ratio of normalized activin A: follistatin, revealed no significant changes with age in either the soleus or extensor digitorum longus muscles. Levels of normalized myostatin showed a slight but non-significant decrease in the EDL with age (Fig. 3A), whereas SOL showed a significant increase in normalized myostatin with age (Fig. 3A). Likewise, the ratio of normalized myostatin: follistatin
showed a significant decrease with age in EDL (Fig. 3B), but a significant increase with age in SOL (Fig. 3B). Two-factor ANOVA with age (12 or 24 mo) and muscle (EDL or SOL) as the two factors showed significant age*muscle interaction effects for both normalized myostatin (p<.05) and the myostatin: follistatin ratio (p<.01), with myostatin levels decreasing with age in the EDL but increasing with age in SOL.

**Figure 3.** Protein levels measured using ELISA of myostatin (A; normalized by BCA) and relative to follistatin (B) in the extensor digitorum longus muscle (EDL) and soleus muscle (SOL) of young (12 mo) and aged (24 mo) mice. Protein levels measured using ELISA of myostatin (C; normalized by BCA) and relative to follistatin (D) in bone marrow supernatant of young (12 mo) and aged (24 mo) mice.

Comparisons of normalized protein levels obtained from bone marrow supernatant revealed no significant differences between older and young mice for activin A, follistatin, or the activin A: follistatin ratio. Normalized myostatin is significantly increased in mouse bone marrow with increasing age (Fig. 3C), as is the ratio of normalized myostatin: follistatin (Fig. 3D). Gene expression data showed slight increases in expression of the myostatin receptor (ActRIIB) with age in both soleus muscles and BMSCs of mice (Fig. 4).
Figure 4. Expression of the myostatin receptor (ActRIIB, or Acvr2b) in the soleus muscle (left graph), extensor digitorum longus muscle (EDL, middle graph), and in bone marrow stromal cells (BMSCs, right graph) from young (6-12 mo) and aged (24 mo) mice. Expression shows a slight but non-significant increase with age in muscle and bone tissues.

Task 2 status. Complete.

Milestone 1: Age-related changes in the expression of myostatin, its receptor, and the myostatin antagonist follistatin in muscle and bone tissues will be defined. Milestone 1 has been reached, with the conclusion that myostatin levels in bone marrow and slow twitch muscle fibers increase slightly with age, as does expression of the myostatin receptor in BMSCs of both humans and mice. Activin A levels also increase with age in human bone marrow, but as shown below this does not have a detrimental effect on bone marrow stem cells and may actually enhance their proliferation.

Aim 2 (months 12-24). Determine the effects of myostatin on anabolic pathways in primary bone-derived stromal cells and skeletal muscle myoblasts in vitro.

Task 1 (months 12-24). Human bone marrow stromal cells from young and aged patients have been collected during year 1 for specific aim 1. In years 2 and 3 these cells will be treated with myostatin, and myostatin inhibitors, in dose-response studies.

Results demonstrate that myostatin treatment of BMSCs actually increases proliferation, which is similar to the effect observed with activin treatment of mouse myoblasts discussed below. The lower dose of propeptide also increased proliferation, although this effect was not seen at the higher dose. Age of donors did not affect the proliferative response (Fig.5).
Figure 5. Proliferation assays of primary human BMSCs isolated from young and aged patients treated with myostatin (left graph) or myostatin inhibitor (right graph). BM70=18 yrs, BM74=40 yrs, BM92=72yrs, BM81=74 yrs.

We also examined the effects of myostatin and myostatin inhibitor on BMSCs in vitro, and found that these treatments had no impact on expression of osteogenic factors whether young or old cells are used (Fig. 6).

Figure 6. Expression of the osteogenic markers osteocalcin (top) and Cbfa1/runx2 (bottom) after treatment with either myostatin or a myostatin inhibitor in young and old BMSCs. Altered myostatin signaling does not have a significant impact on these factors.

Task 1 status: Complete.
Task 2. Primary myoblasts and BMSCs from young and aged mice will have been collected during year 1 for specific aim 1. In year two these cells will be treated with myostatin, and myostatin inhibitors, in dose-response studies.

*Primary myoblast experiments for aim 2, task 2: For culture and treatment of primary myoblasts, tibialis anterior muscles were dissected and placed in sterile PBS. The muscle was minced with a sterile scalpel under aseptic conditions. Minced muscle was digested in 0.2% collagenase type II (Gibco) for 1 hour with frequent shaking followed by digestion in 1x trypsin for 30 minutes. The slurry was pelleted and trypsin supernatant removed. The slurry was re-suspended in proliferation medium. Upon completion of enzymatic digest, slurry was poured over a 70µm cell strainer (Fisher) to remove any remaining connective tissue. The cells were then added to collagen type I (BD Bioscience) coated T-25 flasks. Primary myoblasts were allowed to attach for 72 hours. Cells were then maintained in proliferation medium (PM): DMEM (Hyclone) supplemented with 10% fetal bovine serum, 10% horse serum, 1% penicillin / streptomycin, and 0.5% chick embryo extract (Sera Labs U.K.). Medium was changed every 48 hours until T-25 flask was confluent. Once confluent, cells were trypsinized and counted using NucleoCounter (New Brunswick Scientific). Cells were then plated in a 96 well plate at 5,000 cells/cm². Cells were allowed to attach in proliferation medium for 48 hours. Proliferation medium was removed, cells washed with PBS, and DMEM supplemented with 1% I.T.S was added followed by either control (PBS) or high or low dose activin A (50 ng/ml, 100 ng/ml), follistatin (100 ng/ml, 1000 ng/ml) or myostatin (100 ng/ml, 1000 ng/ml) (R&D Systems, Minneapolis). Doses follow those utilized by He et al. (2005) for activin A and Zhu et al. (2007) for myostatin and follistatin. After 24 hours of treatment, MTT reagent was added according to manufacturer’s protocol and O.D. was determined 2 hours later.

For differentiation assays, cells were isolated and cultured for one week until confluent as described above. Cells were then trypsinized and plated in 12 well plates at 5,000 cells/ml and allowed to attach overnight in proliferation medium. PM was removed, cells washed with PBS. DMEM supplemented with 1% I.T.S followed by the addition of Mstn, Fstn, Activin A or control. Cells were maintained in treatment for 48 hour then harvested in TRIZOL® reagent (Invitrogen) for RNA isolation and subsequent cDNA synthesis (Bio-Rad). 50-100 ng of cDNA was amplified in duplicates in each 40-cycle reaction using an iCycler™ (Bio-Rad) with annealing temperature set at 60ºC, ABsolute™ QPCR SYBR® Green Fluorescein Mix (ABgene, Thermo Fisher Scientific), and custom-designed qRT-PCR primers (Table 1). A melt curve was used to assess the purity of amplification products. mRNA levels were normalized to β-Actin/18S and gene expression was calculated as fold change using the comparative CT method. If not otherwise indicated, treated groups were compared to PBS control groups.

ANOVA showed significant treatment effects for activin A, myostatin, and follistatin treatment of primary myoblasts (Fig. 7A). Activin A produced a significant dose-response increase in proliferation in myoblasts from younger mice but not older mice; however, the treatment*age interaction was not significant. Follistatin also increased proliferation in dose-response manner in young myoblasts, but the effect was attenuated in myoblasts from older mice (Fig. 7B). There was a significant (p=.001) treatment*age interaction effect for follistatin treatment where the treatment effect was much greater in younger myoblasts compared to older cells. Myostatin treatment also increased proliferation in young myoblasts compared to untreated cells, whereas in older myoblasts myostatin decreased proliferation at the low dose (Fig. 7C). There was also a significant (p<.001) treatment*age interaction effect for myostatin, where myostatin treatment increased proliferation in young myoblasts but decreased proliferation in the older cells.
Figure 7. Proliferation assays of primary myoblasts isolated from young (12 mo, left column) and aged (24 mo, right column) mice treated with activin A (A), follistatin (B), or myostatin (C).

RT-PCR data revealed no marked changes in the expression of differentiation markers MyoD, myogenin, or myosin heavy chain (MHC) with Activin A treatment. MyoD and MHC expression were also unaffected by follistatin treatment of young and aged myoblasts; however, follistatin stimulated a significant increase in myogenin expression in aged but not young myoblasts (Fig. 8A). ANOVAs demonstrated a significant age effect for myogenin expression (p<.05), and there was also a significant treatment * age interaction for myogenin expression. Myostatin treatment had no effect on MyoD or myogenin expression in either young or aged myoblasts, but myostatin did produce a significant increase in MHC expression in aged but not young myoblasts (Fig. 8B).
Primary BMSC experiments for Aim 2, task 2: In Aim 1 we found that activin A was also elevated with age in bone marrow, and so we extended our treatments in aim 2 to include activin A, myostatin, and follistatin. Bone marrow aspirates were flushed from femora and magnetic nanoparticles conjugated to anti-mouse CD11b, CD45R/B220, and Pan DC monoclonal antibodies were used to remove hematopoietic-lineage cells. A round of positive-selection was then performed using anti-Sca-1 microbeads. Enriched BMSCs were cultured in proliferation medium (DMEM supplemented with 10% heat-inactivated FBS) in T-75 flasks until ~80% confluent. Cells were then lifted with trypsin/EDTA, plated in 96-well plates at a density of 5,000 cells/well in proliferation medium, and allowed to attach for 24 h. Proliferation medium was removed, cells washed with PBS, and DMEM supplemented with 2% heat-inactivated FBS (for BMSCs) was added followed by control (PBS), activin A, follistatin, or myostatin (all from R&D Systems, Minneapolis) at the same doses noted above for primary myoblasts. After 24 h of treatment, MTS reagent was added according to the manufacturer's protocol (Promega, Madison, WI) and absorbance at 492 nm was read 2 h later. Osteogenic differentiation and Alizarin Red S (ARS) staining was performed as described previously (Zhang et al., 2008).

ANOVA showed a significant (P<.001) age effect in each treatment group, with younger BMSCs overall having higher values for proliferation than older BMSCs, irrespective of the treatment (Fig. 9). Proliferation assays showed no effects of activin treatment on either young or old BMSCs (Fig. 9A). Two-factor ANOVA revealed a significant treatment*age interaction for follistatin, with follistatin having no impact on proliferation in older BMSCs but reducing proliferation in young BMSCs (Fig. 9B). Myostatin significantly decreased BMSC proliferation in both age groups (Fig. 9C); however, this effect was greater in the aged cells, and this treatment*age interaction was also significant (P<.05) for myostatin.

Figure 8. Gene expression of the differentiation marker myogenin (top) and myosin heavy chain (MHC) in young (12M, left column) and aged (24M, right column) primary myoblasts treated with follistatin (A) and myostatin (B; GDF-8).
Figure 9. Results of proliferation assays following treatment of primary bone marrow stromal cells (BMSCs) with activin A (A; activin), follistatin (B; Fstn), and myostatin (C; Mstn). Means with different superscripts differ significantly from one another (P<.05).

Differentiation assays using alizarin red staining to detect mineralization revealed that activin treatment significantly increased mineralization of young and older BMSCs (Fig. 10). The effect of age on mineralization was significant (P<.001) for activin, with the younger cells consistently showing greater mineralization in response to treatment. Treatment effects were less pronounced in aged BMSCs, with the low dose of activin increasing alizarin red staining but the other doses showing no significant effect (Fig. 10C, D). The treatment*age interaction was significant (P<.01) for activin using two-factor ANOVA. In the myostatin experiments, the higher dose of myostatin significantly decreased mineralization in younger cells but not in older cells, and the treatment*age interaction was also significant for myostatin (P<.01). Finally, follistatin treatment did not significantly affect mineralization either young or older BMSCs (Fig. 10).

Figure 10. Alizarin red staining of bone marrow stromal cells cultured in osteogenic conditions. Images of wells (top row) and quantification of staining (bottom row) in BMSCs from young mice (A, B; 12 Mo BMSCs) and older mice (C, D; 24 Mo BMSCs) treated with Activin A (Act), Myostatin (Myo) or Follistatin (Foll) at 50 ng/ml (50), 100 ng/ml (100) or 1000 ng/ml (1000). ***P<.001, *P<.05 relative to same-aged PBS controls.

Task 2 status: Completed. There are two myostatin inhibitors that can be used, one that is myostatin-specific (propeptide) and one that binds both myostatin and activin A (decoy receptor). Prior to determining which is optimal we needed, in aim 2, to determine the effects of...
myostatin and activin on muscle and bone cells. Our data indicate that activin A can have positive effects on myoblast proliferation, and activin also enhances mineralization. Thus, a decoy receptor is not an optimal inhibitor because of its effect of blocking activin activity, hence a myostatin-specific inhibitor (propeptide) the optimal therapeutic to be validated in aim 3.

Milestone 2: The optimal treatment and dose of myostatin inhibitor for enhancing osteogenesis and myogenesis in aged BMSCs and myoblasts will have been determined in vitro. Our data in aim 2 indicate that the myostatin propeptide is the optimal inhibitor because of the contrasting effects of myostatin vs activin A in vitro.

Aim 3 (months 24-36). Determine the effects of myostatin inhibitors on muscle and bone aging in vivo.

Task 1 (months 24-30). Mice 22 months of age will be treated with a myostatin propeptide. Effects on behavioral performance and measures of bone and muscle anabolism will be determined.

Animals & treatments: C57BL6 mice were purchased from the aged rodent colony at the National Institute on Aging, National Institutes of Health (USA) at 22 months of age and delivered to Georgia Regents University, Augusta GA. Animals were allowed to acclimate for one week and were maintained at the Laboratory Animal Service Facility of Georgia Regents University. An earlier dose-response study was used to evaluate the efficacy of a myostatin propeptide in vivo (Hamrick et al., 2010). Adult mice (5-6 mo.) were treated with the propeptide at 0, 10, 20, or 50 mg/kg at day 0, 5, and 10 and then sacrificed one week after the last treatment. Those data showed that propeptide treatment increased fore- and hindlimb muscle mass by 10% at the 10 mg/kg dose and increased muscle mass by more than 15% at the 20 mg/kg dose, but the 50 mg/kg dose did not increase muscle mass beyond the increase observed in the 20 mg/kg group (Hamrick et al., 2010). The 20 mg/kg dose was therefore used in this study. Mice were divided into two treatment groups: a vehicle group (VEH; n=14) and a myostatin propeptide group (PRO; n=15). Mice received i.p. injections every five days for 25 days with a dosage of 20 mg/kg body weight at a volume of 0.2 ml. We used a four- rather than eight-week treatment period because the animals were old enough we were concerned they might die of natural causes prior to completion of the study. Myostatin propeptide [4.48mg/ml] was obtained from Pfizer Inc (Cambridge, MA, USA). Mice were given calcein i.p. injections to label actively mineralizing bone surfaces four days and 24 hours prior to sacrifice.

Myostatin propeptide increases muscle mass and fiber size in aged mice: Body weight of the vehicle- and propeptide-treated animals was similar at the end of the study (Fig 11A). Each treatment group did, however, lose some weight over the treatment period but this was less dramatic for the treated animals, such that their decrease in body weight from day 0 to day 25 was significantly less than that of the vehicle-treated mice (Fig. 11B).

Figure 11. Body mass (A) and change in body weight (B) for animals treated with either saline (VEH) or myostatin propeptide (PRO) weekly for over four weeks.
Muscle mass of the tibialis anterior was significantly increased in the treated mice, both absolutely (Fig. 12A) and relative to body weight (Fig. 12B). Fiber size of the predominantly fast-twitch extensor digitorum longus (EDL) muscle was also significantly increased by more than 15% in the treated mice (Fig. 12C), whereas the increase in muscle fiber size in the predominantly slow-twitch soleus (SOL) muscle was also increased significantly (P<.05) but by a lesser magnitude (~5%).

**Figure 12.** Muscle parameters for mice treated with saline (VEH) or myostatin propeptide (PRO) weekly for a period of four weeks. (A) Tibialis anterior mass, (B) tibialis anterior mass relative to body weight, (C) extensor digitorum longus fiber diameter (EDL).

Propeptide treatment produced a slight but non-significant increase in the expression of myostatin itself, as well as expression of myosin heavy chain and IGF-1 (Fig. 13). Surprisingly, expression of the ubiquitin ligases Murf1 and Mafbx was significantly increased with propeptide treatment (Fig. 13), and the PCR data were further validated by Western blot (Fig. 13).

**Figure 13.** Real-time PCR data (top row) for mice treated with saline (VEH) or myostatin propeptide (PRO) weekly for a period of four weeks showing increased expression of Murf1 and Mafbx in PRO-treated mice (left), and increased expression of BMP-2 in mice treated with propeptide (right). *P<.05. Western blot data from muscle (bottom) showing increased Murf1 and Mafbx in treated muscle.

Myostatin inhibitor significantly increases grip strength in aged mice. We assessed musculoskeletal performance in vehicle (control) and propeptide (pro) treated mice using behavioral outcome measures in Small Animal Behavior Core (SABC) at Georgia Regents University. Rotarod tests, measures of free cage activity, and grip strength tests were performed over a period of approximately 5 days both before and after the four-week treatment.
period. Results demonstrate that rotarod performance was similar for control and treated groups both before and after treatment, however grip strength was significantly increased in the treated mice (Figure 14).

**Figure 14.** Behavioral performance data for mice treated with saline (Con) or myostatin propeptide (Pro) both per- and post-treatment showing similar measures for rotarod (A) and open-field activity (B), but increased grip strength (C) in the propeptide treated mice. *P<.05.

**Myostatin inhibitor does not alter bone formation or bone strength in aged mice:** MicroCT data from the tibia show that bone mineral density is actually slightly higher (3%) in the tibias of vehicle-treated mice (Table 1), but other parameters such as bone volume relative to total volume, trabecular number, and trabecular thickness are similar between the two groups (Table 1). Likewise, three-point bending tests of tibias show that ultimate force, stiffness, and toughness (energy to fracture) are also similar between the vehicle- and propeptide-treated mice (Table 1).

**Table 1.** microCT and biomechanical testing of the proximal tibia for mice treated with saline (VEH) or myostatin propeptide (PRO; 20 mg/kg). BMD=bone mineral density, BV/TV=bone volume relative to total volume, Tb.Th=trabecular thickness, Tb.N=trabecular number, Fu=ultimate force, U=energy-to-fracture, S=stiffness.

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Bone histomorphometry data reveal that osteoblast and osteoclast numbers do not differ between the experimental groups (Table 2). Fluorochrome labeling showed double-labels in only three mice from each group, and so single-labeled surfaces were compared. Actively mineralizing surfaces were also similar between the two groups of mice (Table 2).

Table 2. Bone histomorphometry data for the distal femur of mice treated with saline (VEH) or myostatin propeptide (PRO; 20 mg/kg). N.Ob/BS=osteoblast number per bone surface, MS/BS=mineralizing surface (single-label) relative to bone surface, N.Oc/BS=osteoclast number per bone surface.

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</table>

Gene expression data show no significant differences in the expression of osteogenic genes Osx or Runx2 with propeptide treatment, however the expression of BMP-2 is increased in animals receiving the propeptide (Fig. 13).

Our data show that PRO treatment significantly increases muscle fiber size and muscle mass, both absolutely and relative to body weight. In contrast bone volume, bone strength, and histomorphometric parameters of bone formation and bone resorption were unchanged with PRO treatment. Our findings are consistent with previous studies utilizing a myostatin antibody in aged mice showing that targeting myostatin increases muscle fiber size and mass; however, our data differ from work recently published by Chiu et al. (2013, J. Gerontol. A Biol. Sci. Med. Sci) utilizing a decoy myostatin receptor (ActRIIB-Fc) showing that ActRIIB-Fc appears particularly effective at increasing bone density and bone formation. The anabolic effects of ActRIIB-Fc on aged bone are likely due to the ability of this molecule to antagonize other ligands besides myostatin, such as activin or bone morphogenetic proteins. There are several additional points to consider though when comparing the effects of the two inhibitors. If increases in muscle mass with inhibitor treatment might prevent or delay, not necessarily reverse, bone loss then a myostatin propeptide or antibody could be a safe and effective prophylactic approach for age-related bone loss. Perhaps more importantly, as noted below, the decoy receptor does not appear to be safe.

Task 1 status: The overall goals of task 1 were completed, and we were able to complete the grip strength and rotarod studies with the cost-extension into FY14; however, although the grip strength measures showed a significant change with treatment, the other behavioral measures were quite similar. We have developed an IACUC-approved video system for analyzing gait mechanics which we believe will have greater resolution than the rotarod and ambulatory tests, and we proposed to apply this technology in the extension period for FY15. In addition, our serum assays showed significantly elevated IL-6, an inflammatory cytokine, in propeptide-treated mice. This cytokine is regulated by several important microRNAs (miRNAs) of the let7 family in skeletal muscle, and we are seeking an extension into FY15 to analyse these miRNAs.

Task 2 (months 30-36). Mice 22 months of age will be treated with a soluble decoy myostatin receptor for 8 weeks. Effects on behavioral performance and measures of bone and muscle anabolism will be determined.

Task 2 status. Since the CDMRP application was submitted, Acceleron decided to cease their Phase II trial of the soluble decoy myostatin receptor (ACE-03) because of safety issues (http://quest.mda.org/news/ace-031-clinical-trials-duchenne-md-stopped-now). Acceleron then announced that they will no longer pursue ACE-031 (http://www.acceleronpharma.com/2013/05/acceleron-and-shire-conclude-collaboration-on-ace-031/) this drug because of the safety issues. We therefore decided not to pursue task 2, since
the molecule now appears to be unsafe in humans, and because our data in aim 2 indicate that activin A can have positive effects on myoblast proliferation, and activin also enhances mineralization. Thus, a decoy receptor is not an optimal inhibitor because of its effect of blocking activin activity.

**Milestone 3:** The optimal myostatin inhibitor for enhancing osteogenesis and myogenesis in aged rodents will be determined in vivo. The data presented in aim 3, when taken in consideration of the Acceleron trials noted above, suggest that a myostatin propeptide is likely to be a safe and effective molecule for enhancing muscle mass with aging.

**Key Research Accomplishments in Year 4:**

- We published *(Experimental Gerontology)* the first study on the effects of myostatin propeptide in aged rodents.
- The propeptide treatment data were presented in the form a plenary podium presentation at the national meeting of the American Society for Bone & Mineral Research in Baltimore, MD.
- Our findings were also presented in the form an invited keynote address as the Fesler-Lampert Chair in Aging Studies and the Center on Aging, University of Minnesota, Minneapolis, Minnesota.

**Reportable Outcomes:**

**Manuscripts:**


Abstracts from Professional Presentations:


Invited Seminars:

2013 The Fesler-Lampert Chair in Aging Studies and the Center on Aging, University of Minnesota, Minneapolis, Minnesota.

2013 College of Health Sciences, University of Delaware, Newark, Delaware.

2013 Symposium on Muscle & Mobility, Abbott Nutrition, Columbus, Ohio.

2013 Progress in Bone Biology: Ageing & the Skeleton, Bone Academy, Vienna, Austria.


2012 The Hilton Head Workshop on Regenerative Medicine, Hilton Head, SC.
Conclusions:
Falls and debilitating bone fractures are a major problem for veterans, and more than 40,000 veterans suffered hip fractures from 2000-2002. Men have a higher fracture-related mortality than women, and one out of every three male veterans that sustains a hip fracture dies within one year. Falls are the main etiological factor in more than 90% of fractures, and so treatments that can improve muscle strength while at the same time increasing bone mass will significantly reduce fracture-related morbidity and mortality. Myostatin is a factor that induces muscle wasting and suppresses bone formation. Our data collected thus far demonstrate i) myostatin suppresses proliferation in aged, but not young, myoblasts, ii) myostatin is elevated with age in muscles composed primarily of slow-twitch fibers (e.g. soleus), and iii) myostatin increases muscle mass and muscle fiber size in aged mice. These findings suggest that myostatin inhibitors may have potential for suppressing muscle wasting and improving muscle repair in older individuals, but their effect on bone may be less significant.

References:


**Appendices:** Articles recently published.
Effects of the activin A–myostatin–follistatin system on aging bone and muscle progenitor cells

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A B S T R A C T

The activin A–myostatin–follistatin system is thought to play an important role in the regulation of muscle and bone mass throughout growth, development, and aging; however, the effects of these ligands on progenitor cell proliferation and differentiation in muscle and bone are not well understood. In addition, age-associated changes in the relative expression of these factors in musculoskeletal tissues have not been described. We therefore examined changes in protein levels of activin A, follistatin, and myostatin (GDF-8) in both muscle and bone with age in C57Bl6 mice using ELISA. We then investigated the effects of activin A, myostatin and follistatin on the proliferation and differentiation of primary myoblasts and mouse bone marrow stromal cells (BMSCs) in vitro. Myostatin levels and the myostatin:follistatin ratio increased with age in the primarily slow-twitch mouse soleus muscle, whereas the pattern was reversed with age in the fast-twitch extensor digititorum longus muscle. Myostatin levels and the myostatin:follistatin ratio increased significantly (+75%) in mouse bone marrow with age, as did activin A levels (+17%). Follistatin increased the proliferation of primary myoblasts from both young and aged mice, whereas myostatin increased proliferation of younger myoblasts but decreased proliferation of older myoblasts. Myostatin reduced proliferation of both young and aged BMSCs in a dose-dependent fashion, and activin A increased mineralization in both young and aged BMSCs. Together these data suggest that aging in mice is accompanied by changes in the expression of activin A and myostatin, as well as changes in the response of bone and muscle progenitor cells to these factors. Myostatin appears to play a particularly important role in the impaired proliferative capacity of muscle and bone progenitor cells from aged mice.

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1. Introduction

Aging is associated with a number of changes in the musculoskeletal system, including progressive deterioration of articular cartilage in the form of osteoarthritis, loss of muscle mass in the form of sarcopenia, and loss of bone density and strength in the form of osteoporosis. Muscle weakness and frailty contribute directly to postural instability, which in turn increases the risk for falls, and falls are the main etiological factor in more than 90% of bone fractures. The more than 1.5 million osteoporotic fractures a year in the US place a significant burden on the healthcare system, and also contribute to significant morbidity and poor quality of life. Treatments that can improve muscle strength and at the same time increase bone mass will therefore significantly reduce fracture-related morbidity and mortality.

The activin A–myostatin–follistatin system is believed to play an important role in musculoskeletal growth, development and aging. Myostatin (GDF-8) and activin A bind type II activin receptors and signal through a transforming growth factor-beta signaling pathway involving SMAD phosphorylation. Activin is thought to bind with greater affinity to the type II A activin receptor (ActRIIA) and myostatin to the type II B receptor (ActRIIB), but both are involved in the regulation of muscle mass (Gilson et al., 2009; Lee et al., 2010). Follistatin antagonizes both myostatin and activin A activity, and mice overexpressing follistatin in skeletal muscle show a more dramatic phenotype than mice lacking myostatin alone (Lee et al., 2005). These data suggest that alterations in either myostatin or activin A with aging or disuse can have significant effects on muscle mass, and these may be further influenced by relative levels of follistatin. Although myostatin is not highly expressed by bone cells, loss of myostatin function is associated with increased bone density in mice (Elkasaw and Hamrick, 2010; Morisette et al., 2009). The increased bone density of mice lacking myostatin is likely multifactorial, and may result not only from the indirect effects of increased muscle mass (Hamrick, 2011, 2012) but also from increased circulating levels of IGF-1 (Williams et al., 2011). The type II A and type II B activin receptors are both expressed by chondrocytes and osteoblasts, and activin A has been observed to inhibit mineralization by osteoblasts in vitro whereas follistatin can increase mineralization (Eijken et al., 2007). Furthermore, inhibiting activin A in...
vivo using a decoy soluble activin A receptor (ActRIIA) increases bone formation in mice (Pearsall et al., 2008).

The activin A–myostatin–follistatin system therefore appears to play a number of important roles in muscle as well as in bone metabolism. Although some studies have found no association between age and myostatin transcript levels in skeletal muscle (Marcell et al., 2001), others reveal a marked elevation in skeletal muscle myostatin expression with aging in humans (Leger et al., 2008). Additional research suggests that circulating levels of myostatin increase with age in men and women, and are highest in people aged 60–90 (Yarasheski et al., 2002). The latter finding may implicate myostatin in the sarcopenia of aging, hence myostatin inhibitors could be useful pharmacological agents for treating age-related muscle atrophy as well as bone loss. Indeed, a myostatin inhibitor has been shown to improve muscle regeneration in aged mice (Siriett et al., 2007), and a recent study reveals that muscle-derived stem cells from older male patients show a 165% higher level of myostatin expression compared to stem cells from younger patients (McKay et al., 2012). Myostatin levels can also be reduced in skeletal muscle with resistance exercise in older men (Dalbo et al., 2011), and aerobic exercise can decrease myostatin expression in older women (Konopka et al., 2010). While there has been absolutely little information exists, however, with regard to tissue-specific changes in activin A and follistatin levels with aging or how the response of muscle and bone cells to these factors is altered with age.

We have previously shown that aged C57BL6 mice share a number of key features in common with the aging human musculoskeletal system including an age-related decline muscle mass, both less is known about activin A and follistatin. Circulating activin A levels have been observed to increase with age, whereas circulating follistatin levels did not show age-associated changes (Baccarelli et al., 2001; Hurwitz and Santoro, 2004). Very little information exists, however, with regard to activin A and follistatin levels with aging or how the response of muscle and bone cells to these factors is altered with age.

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### 2. Materials and methods

#### 2.1. Animals

C57BL6 mice were purchased from aged rodent colony at National Institute on Aging, National Institutes of Health (USA) at 12 and 24 months of age and delivered to Georgia Health Sciences University, Augusta, GA. Animals were allowed to acclimate for approximately two weeks and were maintained at the Laboratory Animal Service Facility of the Georgia Health Sciences University. Animals were sacrificed by CO₂ overdose and thoracotomy following procedures approved by the Institutional Animal Care and Use Committee of Georgia Health Sciences University.

#### 2.2. ELISA assays

Whole extensor digitorum longus or soleus muscle was dissected from C57BL6 mice at either 12 or 24 months of age and snap-frozen in liquid nitrogen. Each muscle was placed in 1 ml phosphate buffered saline (PBS) and subjected to homogenization using Fisherbrand Tissuehomizer® rotary homogenizer until large pieces of muscle were no longer visible. Samples were subjected to two freeze–thaw cycles to disrupt the plasma membrane then centrifuged briefly. Samples were separated into 250 μl aliquots and stored at −80 °C until assayed. Follistatin and Activin A ELISA kits were purchased from R&D Systems and assays performed according to manufacturer’s protocol without sample dilution. Myostatin ELISA kits were purchased from Alpco diagnostic and also performed according to manufacturer’s protocol, but samples for myostatin assay were diluted 5-fold. Total protein was measured for each sample using Pierce BCA Protein Assay Kit (Thermo Scientific) according to manufacturer’s protocol. Briefly: after thawing samples for ELISA, 25 μl of sample was incubated undiluted with 200 μl of Pierce® working reagent for 20 min at 37 °C. Absorbance was recorded at 590 nm wavelength and total protein concentrations were calculated based on standard curve generated with bovine serum albumin.

Left and right femora were dissected from C57BL6 mice at either 12 or 24 months of age and placed in PBS. Samples were immediately minced using fine-point surgical scissors. Samples were vortexed vigorously and triturated with a pipette to liberate marrow from bone. Samples were centrifuged briefly, supernatant was collected, and bone fragments were discarded. Supernatant was centrifuged at

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**Table 1**

<table>
<thead>
<tr>
<th>Name</th>
<th>Sequence</th>
<th>Amplicon size</th>
</tr>
</thead>
<tbody>
<tr>
<td>Myogenin</td>
<td>Fwd: GGAAGTCCTGTTCGGTGAC</td>
<td>150</td>
</tr>
<tr>
<td></td>
<td>Rev: CGGCGCGAGTTCTCCAC</td>
<td></td>
</tr>
<tr>
<td>MysD</td>
<td>Fwd: GCTCTGACAAAGTCACTAGG</td>
<td>184</td>
</tr>
<tr>
<td></td>
<td>Rev: GGTCCAGGGTCCACAG</td>
<td></td>
</tr>
<tr>
<td>Myosin, heavy polypeptide 3 (Mhy3)</td>
<td>Fwd: ACGTACAGGGTGCTGACAGCGG</td>
<td>90</td>
</tr>
<tr>
<td>BMP-2</td>
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<td>83</td>
</tr>
<tr>
<td></td>
<td>Rev: TCAATGCTGCGCGTCACTAGG</td>
<td></td>
</tr>
<tr>
<td>Osteocalcin</td>
<td>Fwd: ATTAGACACCCCTGTGACATAC</td>
<td>120</td>
</tr>
<tr>
<td></td>
<td>Rev: CGAATGCCTGTGACATAC</td>
<td></td>
</tr>
</tbody>
</table>

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![Fig. 1](image_url)
1500 g for 5 min to pellet cells. Supernatant was collected, aliquoted, and stored at −80 °C until assayed. Specific ELISA assays and total protein were performed and measured as described above for skeletal muscle homogenates.

2.3. Primary myoblast isolation, culture, and proliferation and differentiation assays

Mice 12 and 24 months of age were euthanized and tibialis anterior muscle dissected and place in sterile PBS. The muscle was minced with a sterile scalpel under aseptic conditions. Minced muscle was digested in 0.2% collagenase type II (Gibco) for 1 h with frequent shaking followed by digestion in 1× trypsin for 30 min. The slurry was pelleted and trypsin supernatant removed. The slurry was re-suspended in proliferation medium. Upon completion of enzymatic digest, slurry was poured over a 70 μm cell strainer (Fisher) to remove any remaining connective tissue. The cells were then added to collagen type I (BD Bioscience) coated T-25 flasks.

Primary myoblasts were allowed to attach for 72 h. Cells were then maintained in proliferation medium (PM): DMEM (Hyclone) supplemented with 10% fetal bovine serum, 10% horse serum, 1% penicillin/streptomycin, and 0.5% chick embryo extract (Sera Labs U.K.). Medium was changed every 48 h until T-25 flask was confluent. Once confluent, cells were trypsinized and counted using NucleoCounter (New Brunswick Scientific). Cells were then plated in a 96 well plate at 5000 cells/cm². Cells were allowed to attach in proliferation medium for 48 h. Proliferation medium was removed, cells washed with PBS, and DMEM supplemented with 1% insulin-transferrin–sodium selenite (ITS) was added followed by either control (PBS) or high or low dose activin A (50 ng/ml and 100 ng/ml), follistatin (100 ng/ml and 1000 ng/ml) or myostatin (100 ng/ml and 1000 ng/ml) (R&D Systems, Minneapolis). Doses follow those utilized by He et al. (2005) for activin A and Zhu et al. (2007) for myostatin and follistatin. After 24 h of treatment, MTS reagent was added according to the manufacturer’s protocol (Promega, Madison) and absorbance at 492 nm was read 2 h later.

For differentiation assays, cells were isolated and cultured for one week until confluent as described above. Cells were then trypsinized and plated in 12 well plates at 5000 cells/ml and allowed to attach overnight in proliferation medium. PM was removed, cells washed with PBS, and DMEM-supplemented with 1% ITS was added followed by the addition of Mstn, Fstn, Activin A or control. Cells were

![Graphs showing results of proliferation assays following treatment of primary myoblasts with activin A (A; activin), follistatin (B; Fstn), and myostatin (C; Mstn). Means with different superscripts differ significantly from one another (P<.05). Error bars represent one standard deviation and sample size includes eight replicates per group.](image-url)
maintained in treatment for 48 h then harvested in TRIZOL® reagent (Invitrogen) for RNA isolation and subsequent cDNA synthesis (Bio-Rad). 50–100 ng of cDNA was amplified in duplicates in each 40-cycle reaction using an iCycler™ (Bio-Rad) with annealing temperature set at 60 °C, ABSolute™ QPCR SYBR® Green Fluorescein Mix (ABgene, Thermo Fisher Scientific), and custom-designed qRT-PCR primers (Table 1). A melt curve was used to assess the purity of amplification products. mRNA levels were normalized to β-Actin/18S and gene expression was calculated as fold change using the comparative CT method. If not otherwise indicated, treated groups were compared to PBS control groups.

2.4. Primary bone marrow stromal cell (BMSC) isolation, culture, and proliferation and differentiation assays

Bone marrow aspirates were flushed from mouse femora and tibias and BMSCs isolated using magnetic bead sorting as previously described (Zhang et al., 2008). Briefly, magnetic nanoparticles conjugated to anti-mouse CD11b, CD11c, CD45R/B220, and Pan DC monoclonal antibodies were used to remove hematopoietic-lineage cells and those that were negative for these four antigens remained in the solution and were collected and subjected to a round of positive-selection using anti-Sca-1 microbeads.

Enriched BMSCs were cultured in proliferation medium (DMEM supplemented with 10% heat-inactivated FBS) in T-75 flasks until ~80% confluent. Cells were then lifted with trypsin/EDTA, plated in 96-well plates at a density of 5000 cells/well in proliferation medium, and allowed to attach for 24 h. Proliferation medium was removed, cells washed with PBS, and DMEM supplemented with 2% heat-inactivated FBS (for BMSCs) was added followed by control (PBS), activin A, follistatin, or myostatin (all from R&D Systems, Minneapolis) at the same doses noted above for primary myoblasts. After 24 h of treatment, MTS reagent was added according to the manufacturer’s protocol (Promega, Madison, WI) and absorbance at 492 nm was read 2 h later.

Osteogenic differentiation and Alizarin Red S (ARS) staining was performed as described previously (Zhang et al., 2008). In brief, enriched BMSCs were plated in 24-well plates at a density of 5000 cells/cm² in proliferation medium and allowed to attach for 24 h. Cells were then treated with osteogenic induction medium (DMEM) supplemented with 5% FBS, 0.25 mM ascorbic acid, 0.1 μM dexamethasone, and 10 mM β-glycerophosphate (all from Sigma-Aldrich) for 14 d. For ARS staining, differentiated BMSCs were washed with PBS and fixed in 3% paraformaldehyde (Sigma-Aldrich) for 30 min. Cells were stained with 40 mM ARS pH 4.1 (Sigma-Aldrich) for 15 min followed by washing with excess dH2O. Stained plates were scanned for visualization. For quantitative destaining, 10% acetic acid was added for 30 min. Samples were transferred to a 1.5 ml microcentrifuge tube, overlaid with mineral oil (Sigma-Aldrich), heated to 85 °C for 10 min, and transferred to ice for 5 min. Following centrifugation at 14,000 rpm for 15 min, supernatants were removed and neutralized with 10% ammonium hydroxide. Aliquots were transferred to a 96-well plate and absorbance was read at 405 nm (Gregory et al., 2004). Expression of osteogenic

Fig. 3. Real-time PCR data for primary myoblast expression of myogenin (A) and myosin heavy chain (B) in response to treatment with follistatin (Fstn, top) and myostatin (Mstn, bottom). Means with different superscripts differ significantly from one another (P<.05). Error bars represent one standard deviation and sample size includes four-six replicates per group.
genes was assessed using primers for bone morphogenetic protein-2 (BMP-2) and osteocalcin (OCN) shown in Table 1.

2.5. Statistical analysis

Two-factor ANOVAs were performed with treatment and age as the two factors. Fisher’s least significant difference (LSD) test was used for post-hoc, pairwise comparisons. All statistical analyses were performed using SYSTAT® software.

3. Results

3.1. Protein levels of activin A, myostatin, and follistatin in skeletal muscle

ANOVaras performed on activin A and follistatin normalized for total protein, and the ratio of normalized activin A: follistatin, revealed no significant changes with age in either the soleus or extensor digitorum longus muscles. Levels of normalized myostatin showed a slight but non-significant decrease in the EDL with age (Fig. 1A), whereas SOL showed a significant increase in normalized myostatin with age (Fig. 1A). Likewise, the ratio of normalized myostatin: follistatin showed a significant (~40%) decrease with age in EDL (Fig. 1B), but a significant (~four-fold) increase with age in SOL (Fig. 1B). Two-factor ANOVA with age (12 or 24 mo) and muscle (EDL or SOL) as the two factors showed significant age × muscle interaction effects for both normalized myostatin (p < .05) and the myostatin:follistatin ratio (p < .01), with myostatin levels decreasing with age in the EDL but increasing with age in SOL.

3.2. Effects of activin A, myostatin, and follistatin on the proliferation and differentiation of primary myoblasts

Overall there was a significant (P < .01) age effect for both myostatin and follistatin treatment, in which young cells were more proliferative in response to treatment than older cells. ANOVAs showed significant treatment effects for activin A, myostatin, and follistatin (Fig. 2A). Activin A produced a significant dose–response increase in proliferation in myoblasts from younger mice but not older mice; however, the treatment × age interaction was not significant. Follistatin also increased proliferation in dose–response manner in young myoblasts, but the effect was attenuated in myoblasts from older mice (Fig. 2B). There was a significant (p = .001) treatment × age interaction effect for follistatin treatment where the treatment effect, approximately two-fold at the highest dose, was much greater in younger myoblasts compared to older cells. Myostatin treatment also increased proliferation in young myoblasts compared to untreated cells, whereas in older myoblasts myostatin decreased proliferation at the low dose (Fig. 2C). There was also a significant (p < .001) treatment × age interaction effect for myostatin, where myostatin treatment increased proliferation in young myoblasts but decreased proliferation in the older cells.

RT-PCR data revealed no marked changes in the expression of differentiation markers MyoD, myogenin, or myosin heavy chain (MHC) with Activin A treatment. MyoD and MHC expression were also unaffected by follistatin treatment of young and aged myoblasts; however, follistatin stimulated a significant increase in myogenin expression in aged but not young myoblasts (Fig. 3A). ANOVAs demonstrated a significant age effect for myogenin expression (p < .05) with older cells generally showing higher levels of myogenin expression, and there was also a significant treatment × age interaction for myogenin expression. Myostatin treatment had no effect on MyoD or myogenin expression in either young or aged myoblasts, but myostatin did produce a significant increase in MHC expression in aged but not young myoblasts (Fig. 3B).

3.3. Protein levels of activin A, myostatin, and follistatin in bone marrow supernatants

Comparisons of normalized protein levels obtained from bone marrow supernatant revealed no significant differences between older and young mice for activin A, follistatin, or the activin A: follistatin ratio. Normalized myostatin is significantly (~80%) increased in mouse bone marrow with increasing age (Fig. 4A), as is the ratio of normalized myostatin: follistatin (+60%; Fig. 4B).

3.4. Effects of activin A, myostatin, and follistatin on the proliferation and differentiation of primary bone marrow stromal cells (BMSCs)

ANOVaras showed a significant (P < .001) age effect in each treatment group, with younger BMSCs overall having higher values for proliferation than older BMSCs, irrespective of the treatment (Fig. 5). Proliferation assays showed no effects of activin treatment on either young or old BMSCs (Fig. 5A). Two-factor ANOVA revealed a significant treatment × age interaction for follistatin, with follistatin having no impact on proliferation in older BMSCs but reducing proliferation in young BMSCs (Fig. 5B). Myostatin significantly decreased BMSC proliferation in both young cells (~15%) and aged cells (~40%); however, this effect was greater in the aged cells, and this treatment × age interaction was also significant (P < .05) for myostatin (Fig. 5C).

Differentiation assays using alizarin red staining to detect mineralization revealed that activin treatment significantly increased mineralization of young and older BMSCs (Fig. 6). The effect of age on mineralization was significant (P < .001) for activin, with the
younger cells consistently showing greater mineralization in response to treatment. Treatment effects were less pronounced in aged BMSCs, with the low dose of activin increasing alizarin red staining but the other doses showing no significant effect (Fig. 6C,D). The treatment × age interaction was significant (P < .01) for activin using two-factor ANOVA. In the myostatin experiments, the higher dose of myostatin significantly decreased mineralization in younger cells but not in older cells, and the treatment × age interaction was also significant for myostatin (P < .01). Finally, follistatin treatment did not significantly affect mineralization either young or older BMSCs (Fig. 6).

4. Discussion

The ELISA assays using muscle and bone samples from aged mice provided results that were unexpected for two reasons. First, it is well-known that fast-twitch (type II) muscle fibers atrophy with age (Holloszy et al., 1991) and a number of studies have implicated myostatin in muscle atrophy (McKay et al., 2012). In addition, Hennebry et al. (2009) found that mice lacking myostatin showed an increase in type I fibers and decrease in type II fibers, and blocking myostatin function in vivo using a myostatin antibody has been shown to increases type II fiber size in aging mice (Murphy et al., 2010). The fact that we observed a decline in myostatin levels with age in the predominantly fast-twitch extensor digitorum longus muscle of mice, but a relative increase in myostatin levels in the slow-twitch (type I) soleus, was therefore not anticipated. Nevertheless, it has been shown that mice lacking myostatin lose the same percentage of muscle mass with age as normal mice (Morissette et al., 2009), and muscles composed of different fiber types are all larger in aged myostatin-deficient mice compared to those of same-aged wild-type mice (Jackson et al., 2012). Given that both myostatin levels and the myostatin:follistatin ratio increased with age in soleus muscles of mice compared to extensor digitorum longus, myostatin inhibitors may have a greater impact on preserving muscle mass with age in those muscles composed predominantly of slow- rather than fast-twitch fibers.

The second in vivo finding that was also unexpected was the significant rise in myostatin protein with age in bone marrow supernatants. This was not expected because myostatin is primarily secreted by muscle and not by osteoblasts or marrow stromal cells. Yarasheski et al. (2002) did, however, find that serum myostatin increased with age in older women, and Szulc et al. (2012) found that serum myostatin increased in men until age 57 and then declined. The rise in bone marrow myostatin that we observed with age in mice likely reflects an increase...
in circulating myostatin levels with age. The elevated concentrations of myostatin in the bone marrow microenvironment of aged mice is significant given the findings discussed below regarding the effects of myostatin on proliferation of bone marrow stromal cells. Moreover, while myostatin was found to be most highly expressed in skeletal muscle progenitors early in development (McPherron et al., 1997), data presented here and elsewhere suggest that myostatin levels may rise again later in life. Thus, the interplay among activin A, myostatin, and follistatin is dynamic and changes across the aging spectrum.

The cell proliferation experiments yielded findings that were consistent with those of previous studies with regard to the effects of age and treatment. First, in both primary myoblasts and BMSCs, age effects were significant regardless of treatment. That is, aged myoblasts and BMSCs were less proliferative overall in response to treatment whether the cells were exposed to activin A, myostatin, or follistatin. This does not appear to be related to expression of ActRIIB, since we have observed using PCR (data not shown) that ActRIIB expression does not differ significantly either between muscle and bone cells of mice at similar ages, or between musculoskeletal tissues of young versus aged mice. Second, in the case of primary myoblasts, all of the ligands generally increased proliferation of young myoblasts at lower doses. Activin A was previously found to suppress embryonic muscle development in chicks (He et al., 2005), and to inhibit proliferation of stem cells derived from adult muscle (Nomura et al., 2008). Myostatin has also been shown previously to be a potent inhibitor of myoblast proliferation (McFarlane et al., 2011), and while myostatin increased proliferation of young myoblasts it decreased proliferation of aged myoblasts. The effects of these ligands on proliferation were quite different in BMSCs, where follistatin and myostatin both suppressed proliferation of young BMSCs, and myostatin significantly suppressed the proliferation of aged BMSCs. These data are consistent with our previous findings showing that BMSCs of mice lacking myostatin were more proliferative in vitro than those from normal mice (Elkasrawy et al., 2011). Probably the most important observation from these in vitro experiments is that, in the case of both aged myoblasts and aged BMSCs, myostatin consistently suppressed cell proliferation. Aged muscle precursors have an impaired capacity for proliferation (Conboy et al., 2003; Machida and Booth, 2004), as do aged BMSCs (Kretlow et al., 2008). Blocking myostatin function in aged animals may therefore have significant potential for improving the repair and regeneration of both muscle and bone by improving the proliferative capacity of both myo- and osteo-progenitor cells.

Findings from our differentiation assays are generally consistent with the idea that myostatin inhibits proliferation and promotes terminal differentiation of aged myoblasts, as myostatin treatment increased myosin heavy chain expression in aged myoblasts. The osteoblast differentiation and mineralization assays revealed that while activin A increased mineralization, myostatin exposure suppressed mineralization in vitro (Fig. 6). These experiments where BMSCs were treated with myostatin are also consistent with our previous studies using BMSCs from myostatin-deficient mice, which showed that Mstn−/− BMSCs had an increased capacity for mineralization in vitro and impaired potential for adipocyte differentiation (Hamrick et al., 2007; Kellum et al., 2009). The activin A differentiation experiments with BMSCs differ somewhat from those of Eijken et al. (2007) who found that activin A treatment strongly inhibited mineralization in osteoblast cultures, whereas follistatin increased mineralization. Our data, on the other hand, showed that osteogenic culture of BMSCs in the presence

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**Fig. 6.** Alizarin red staining of bone marrow stromal cells cultured in osteogenic conditions (A–D) and PCR data for osteogenic genes (E–F) from BMSCs treated with activin A. Images of wells (A, C) and quantification of staining (B, D) in BMSCs from young mice (A, B; 12 mo BMSCs) and older mice (C, D; 24 mo BMSCs) treated with Activin A (Act), Myostatin (Myo) or Follistatin (Folli) at 50 ng/ml (50), 100 ng/ml (100) or 1000 ng/ml (1000). ***P<.001 and *P<.05 relative to same-aged PBS controls. Error bars represent one standard deviation and four replicates are included per treatment group.
of activin A increased mineralization whereas follistatin had no effect on mineralization in either young or old BMSCs. It is likely that our findings differ from those of Eijken et al. (2007) because our experiments examined the effects of activin A and follistatin on the differentiation of osteoprogenitors, whereas Eijken et al. treated differentiated osteoblasts. It is therefore important to acknowledge here that, while our study examined the proliferation and differentiation of progenitor cell populations, ligands of the activin A–myostatin–follistatin system may have different effects on target cells at different stages of development, differentiation, and maturation.

5. Conclusions

Our experiments using aged mice as a model system for investigating the role of activin A, myostatin, and follistatin in musculoskeletal aging suggest that these ligands are likely to be involved in the altered capacity for tissue formation and repair that is observed in aged rodents and humans. Specifically, primary myoblasts and bone marrow stromal cells of aged animals were found to be less proliferative than those of younger animals in response to most of the treatments utilized. Myostatin levels in bone marrow and in soleus muscles increased with age in mice, and myostatin treatment suppressed proliferation of both aged myoblasts and aged BMSCs. Myostatin also inhibited mineralization of BMSCs under osteogenic culture conditions. Together these data suggest that targeting myostatin in aged animals may improve the proliferative capacity of muscle and bone progenitor cells, perhaps enhancing the potential for muscle and bone repair and regeneration in the therapeutic setting.

Acknowledgments

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M. Bowser et al. / Experimental Gerontology 48 (2013) 290–297
A Myostatin Inhibitor (Propeptide-Fc) Increases Muscle Mass and Muscle Fiber Size in Aged Mice but Does not Increase Bone Density or Bone Strength

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Abstract

Loss of muscle and bone mass with age are significant contributors to falls and fractures among the elderly. Myostatin deficiency is associated with increased muscle mass in mice, dogs, cows, sheep and humans, and mice lacking myostatin have been observed to show increased bone density in the limb, spine, and jaw. Transgenic overexpression of myostatin propeptide, which binds to and inhibits the active myostatin ligand, also increases muscle mass and bone density in mice. We therefore sought to test the hypothesis that in vivo inhibition of myostatin using an injectable myostatin propeptide (GDF8 propeptide-Fc) would increase both muscle mass and bone density in aged (24 mo) mice. Mice were injected weekly (20 mg/kg body weight) with recombinant myostatin propeptide-Fc (PRO) or vehicle (VEH; saline) for four weeks. There was no difference in body weight between the two groups at the end of the treatment period, but PRO treatment significantly increased mass of the tibialis anterior muscle (+7%) and increased muscle fiber diameter of the extensor digitorum longus (+16%) and soleus (+6%) muscles compared to VEH treatment. Bone volume relative to total volume (BV/TV) of the femur calculated by microCT did not differ significantly between PRO- and VEH-treated mice, and ultimate force (Fu), stiffness (S), toughness (U) measured from three-point bending tests also did not differ significantly between groups. Histomorphometric assays also revealed no differences in bone formation or resorption in response to PRO treatment. These data suggest that while developmental perturbation of myostatin signaling through either gene knockout or transgenic inhibition may alter both muscle and bone mass in mice, pharmacological inhibition of myostatin in aged mice has a more pronounced effect on skeletal muscle than on bone.

Keywords
sarcopenia; osteoporosis; fractures; anabolic therapy

1. Introduction

Globally, the size of the aging population is increasing rapidly, and as a corollary the prevalence of age-related musculoskeletal disorders such as osteoarthritis, sarcopenia, and
osteoporosis is also increasing (Sanchez-Riera et al., 2010). A primary contributor to the morbidity and mortality associated with aging is an increased frequency of falls, and falls are often accompanied by bone fractures. Indeed, falls are the primary factor in more than 90% of bone fractures (Jarvinen et al., 2008). In many cases bone fractures limit subsequent capacity for normal daily activities, ambulation, and independence, ultimately leading to assisted living situations which can be financially burdensome. The disability following a fall and fracture also contributes directly to an increase in comoribities such as respiratory infections, which in turn contribute to greater overall age-related mortality (Bertram et al., 2011).

The growth, development, and aging of muscle and bone are closely linked. Pediatric gains in bone mass are normally preceded by gains in muscle mass, and loss of muscle mass with age typically precedes peak rates of loss in bone density (Hamrick et al., 2010a). The close coupling of muscle and bone across the lifespan therefore suggests that changes in one tissue may be mechanistically linked with changes in another. Indeed, there are multiple mechanisms linking the two tissues such as mechanical loading, muscle-derived trophic factors (myokines), as well as systemic factors such as sex steroids and growth factors that have anabolic effects on both muscle and bone. The functional and perhaps molecular integration between the two tissues therefore suggests that therapeutic strategies targeting one particular tissue may have positive effects on the other, or that certain pharmacologic approaches (e.g., androgen-receptor modulators, vitamin D receptor agonists) could positively impact both tissues at once (Hamrick 2010, 2011, 2012).

Given the very close linkages between muscle and bone referenced above, we sought to test the hypothesis that pharmacologic inhibition of myostatin (GDF-8) could increase both muscle and bone mass in an aged animal model. Our rationale for pursuing this hypothesis is based on several key observations. The first is that mice lacking myostatin show increased muscle mass as well as increased bone density in the spine, limb, and jaw (Elkasrawy et al., 2010). The second is that recent studies have demonstrated that a myostatin antibody (LeBrasseur et al., 2009; Murphy et al., 2010) and a decoy myostatin receptor (Chiu et al., 2013) can increase muscle mass in aged mice. The decoy receptor (ActR-IIB) was also found to increase bone formation and bone density (Chiu et al., 2013). These findings suggest that therapeutic modulation of myostatin in vivo may be an effective strategy for preserving muscle and bone mass with age, and so we employed a mouse model to evaluate this hypothesis. Specifically, we have previously shown that C57BL6 mice lose significant muscle mass, bone density, and bone strength with age, such that mice 24 months of age show a marked decrease in these measures compared to mice at 12 months of age (Hamrick et al., 2006a). This study utilizes myostatin propeptide treatment in aged (24 months) C57BL6 mice as an in vivo model for assessing the potential effects of myostatin inhibition on age associated muscle and bone loss. The propeptide fragment is utilized here because it has previously been shown to enhance muscle and bone repair in vivo, and binds the active myostatin ligand with high affinity (Bogdanovich et al., 2005; Hamrick et al, 2010b).

2. Materials & Methods

2.1 Production and validation of myostatin propeptide

The myostatin propeptide-Fc fusion protein was produced in CHO cells as described previously (Jiang et al., 2004). To measure the activity of myostatin and the efficacy of the myostatin propeptide a luciferase reporter gene assay was developed where the vector pGL3(CAGA)12 – neo was stably transfected into A204 (human rhabdomyosarcoma) cells. Addition of myostatin to the A204 CAGA cells, and the binding of myostatin to its receptors, initiates the Smad signal transduction pathway and activates the luciferase reporter gene. The level of activation is proportional to the luciferase activity and the linear
portion of the activity curve is in the ng/ml range (Fig. 1), which is what is expected for a protein in the TGFβ superfamily. The addition of the myostatin propeptide prevents the binding of myostatin to its receptors, and the IC50 for the propeptide is approximately 2.0 nM (Fig. 1).

2.2 Animals and treatment for aging study

C57BL6 mice were purchased from the aged rodent colony at the National Institute on Aging, National Institutes of Health (USA) at 22 months of age and delivered to Georgia Regents University, Augusta GA. Animals were allowed to acclimate for one week and were maintained at the Laboratory Animal Service Facility of Georgia Regents University. An earlier dose-response study was used to evaluate the efficacy of a myostatin propeptide in vivo (Hamrick et al., 2010b). Adult mice (5–6 mo.) were treated with the propeptide at 0, 10, 20, or 50 mg/kg at day 0, 5, and 10 and then sacrificed one week after the last treatment. Those data showed that propeptide treatment increased fore- and hindlimb muscle mass by 10% at the 10 mg/kg dose and increased muscle mass by more than 15% at the 20 mg/kg dose, but the 50 mg/kg dose did not increase muscle mass beyond the increase observed in the 20 mg/kg group (Hamrick et al., 2010b). The 20 mg/kg dose was therefore used in this study. Mice were divided into two treatment groups: a vehicle group (VEH; n=14) and a myostatin propeptide group (PRO; n=15). Mice received i.p. injections every five days for 25 days with a dosage of 20 mg/kg body weight at a volume of 0.2 ml. Myostatin propeptide [4.48mg/ml] was obtained from Pfizer Inc (Cambridge, MA, USA). Mice were given calcein i.p. injections to label actively mineralizing bone surfaces four days and 24 hours prior to sacrifice.

2.3 Tissue collection

Animals were euthanized by CO2 overdose and thoracotomy following procedures approved by the Institutional Animal Care and Use Committee of Georgia Regents University. The extensor digitorum longus (EDL) and soleus (SOL) muscles from one limb were dissected out, cut in half and embedded in OCT for cryostat sectioning and muscle fiber size measurements. The tibialis anterior from one limb was dissected out, weighed, snap frozen and homogenized for RT-PCR of the following myogenic markers: myostatin, Mafbx, Murf 1, MHC, and IGF-1. The right tibias were disarticulated and fixed in 70% ethanol for μCT and plastic sectioning while the left tibias were stored damp at minus 20°C for biomechanical testing followed by RT-PCR for bone formation and osteoblast differentiation markers Runx2, Osx and BMP-2. The femora were fixed in 4% paraformaldehyde, decalcified and embedded in paraffin for sectioning and stained for osteoclasts (TRAP kit from Sigma 386A-1KT) and osteoblasts (Celestine blue/van Geison’s).

2.4 Bone Histomorphometry

Osteoblast and osteoclast counts were performed as previously described (Wenger et al., 2010) on 4–5 μm sections after the specimens were decalcified in 4% EDTA for 1 week, dehydrated, cleared in xylene, then embedded in paraffin. Osteoblasts were counted on sections stained using von Giessen, and osteoclasts counted on sections stained for tartrate-resistant acid phosphatase (TRAP) activity. Standardized peripheral locations from the metaphysis were measured in a fixed region of interest. Mineralizing surfaces were measured from calcine-labeled, undecalced bone sections. Tibias fixed in 70% ethanol were dehydrated and embedded in methyl methacrylate and sectioned in the horizontal (transverse) plane. Sections were viewed using a Zeiss Axiosplan2 fluorescent microscope and captured using a SPOT® digital camera to image labeled bone surfaces. Forming surface was calculated as the percentage of non-eroded, single-labeled surface/total surface × 100 (MS/BS).
2.5 Biomechanical testing and micro-computed tomography (uCT)

Left tibias were stored damp at −20°C before being allowed to thaw at room temperature in PBS for 1 hr. Specimens were tested in three-point bending using a Vitrodyne V1000 Material Testing system as described previously (Hamrick et al., 2006b, 2008). Tibias were mounted antero-posteriorly on stainless steel fixtures 5mm apart, approximately 2.5 mm either side of center. Testing was linear displacement control with a displacement speed of 10 μm/sec using a Transducer Techniques 5 kg load cell. Structural, or extrinsic, properties including ultimate force (Fu; height of curve) and stiffness (S; slope of curve) were calculated from load-displacement curves. MicroCT images of the right tibia were scanned using Bruker SkyScan1174 compact micro-CT (Belgium), software version 1.5 (build 9) using NRRecon version 1.6.4.8 for reconstructed images.

2.6 PCR and Western blotting

Total RNA was extracted using Trizol and cDNA was synthesized using Quantitect reverse transcription kit (catalog no. 205310; Qiagen). Expression was analyzed quantitatively by means of the Quantitect SYBR Green PCR kit (catalog no. 204143, Qiagen), and QuantiTect Primer Assays. We used specific primers provided by QuantiTect Primer Assays for Myostatin, Murf1, MaFbx, MHC, IGF-1, Runx2, Osx, BMP-2 and 18S, GAPDH and β-actin (internal controls; Table 1). Half of each extensor digitorum longus muscle was snap-frozen in liquid nitrogen for Western blotting. Each muscle was placed in phosphate buffered saline (PBS) and subjected to homogenization using Fisherbrand Tissuemiser® rotary homogenizer until large pieces of muscle were no longer visible. Samples were subjected to two freeze–thaw cycles to disrupt the plasma membrane then centrifuged briefly. Protein concentrations were measured using a commercial BCA reagent (Pierce, Rockford, IL) to ensure equal loading. 30 μg of proteins from whole tissue lysates were mixed 1:1 with 2× sample buffer and then applied to 4–20 % polyacrylamide gels. Samples were electrophoretically separated and transferred to nitrocellulose membrane (Invitrogen, Carlsbad, CA). The membranes were incubated with specific primary antibodies MURF1 (Abcam cat. 77577) or MAFbx (Santa Cruz Biotech cat. 166806) and then incubated with anti-mouse or anti-rabbit peroxidase-conjugated secondary antibodies (Santa Cruz, CA). After the incubation, the membranes were washed three times for 15 min each with 1× TTBS solution and then incubated with 1 ml of chemiluminescence reagent (Invitrogen). The protein bands were visualized using X-ray films (Fisher Scientific, Rochester, NY).

2.7 Statistical analysis

Single-factor ANOVA with treatment group as the factor was used to for pairwise comparisons of morphometric and histomorphometric parameters. For analysis of gene expression data, the control genes of 18S and Actin were averaged to obtain an average control gene for muscle tissue while GAPDH was used as the control gene for bone. Difference in control gene Ct expression between GDF-8 and vehicle was assessed using a two-sample t-test. Delta Ct values for each treatment group were calculated as ΔCt = CtTarget gene − CtControl gene. The difference in ΔCt expression between GDF-8 and vehicle was assessed using a two-sample t-test. The magnitude of the difference between the groups was estimated using deltadelta Ct values for each target gene and these were calculated as ΔΔCt = ΔCtGDF-8 − ΔCtvehicle and fold change was calculated as 2 to the power −ΔΔCt. SAS® version 9.3 (SAS Institute, Inc., Cary, NC) was used for all analyses and alpha=0.05 was used to determine statistical significance.
3. Results

3.1 Myostatin propeptide increases muscle mass and fiber size in aged mice

Body weight of the vehicle- and propeptide-treated animals was similar at the end of the study (Fig. 2A). Each treatment group did, however, lose some weight over the treatment period but this was less dramatic for the treated animals, such that their decrease in body weight from day 0 to day 25 was significantly less than that of the vehicle-treated mice (Fig. 2B). Muscle mass of the tibialis anterior was significantly increased in the treated mice, both absolutely (Fig. 3A) and relative to body weight (Fig. 3B). Fiber size of the predominantly fast-twitch extensor digitorum longus (EDL) muscle was also significantly increased by more than 15% in the treated mice (Fig. 3C,D), whereas the increase in muscle fiber size in the predominantly slow-twitch soleus (SOL) muscle was also increased significantly (Fig. 3E) but by a lesser magnitude (~5%). Propeptide treatment produced a slight but non-significant increase in the expression of myostatin itself, as well as expression of myosin heavy chain and IGF-1 (Fig. 4A). Surprisingly, expression of the ubiquitin ligases Murf1 and Mafbx was significantly increased with propeptide treatment (Fig. 4A), and the PCR data were further validated by Western blot (Fig. 4B).

3.2 Myostatin inhibitor does not alter bone formation or bone strength in aged mice

MicroCT data from the tibia show that bone mineral density is actually slightly higher (3%) in the tibias of vehicle-treated mice (Table 2), but other parameters such as bone volume relative to total volume, trabecular number, and trabecular thickness are similar between the two groups (Table 2). Likewise, three-point bending tests of tibias show that ultimate force, stiffness, and toughness (energy to fracture) are also similar between the vehicle- and propeptide-treated mice (Table 2). Bone histomorphometry data reveal that osteoblast and osteoclast numbers do not differ between the experimental groups (Table 3). Fluorochrome labeling showed double-labels in only three mice from each group, and so single-labeled surfaces were compared. Actively mineralizing surfaces were also similar between the two groups of mice (Table 3). Gene expression data show no significant differences in the expression of osteogenic genes Osx or Runx2 with propeptide treatment, however the expression of BMP-2 is increased in animals receiving the propeptide (Fig. 4C).

4. Discussion

Pharmacological inhibition of myostatin has, to date, been pursued using a variety of in vivo approaches. These include utilization of myostatin-specific antibodies (Bogdanovich et al., 2002; Wagner et al., 2008; LeBrasseur et al., 2009; Murphy et al., 2010), a decoy myostatin receptor (ActRIIB-Fc; Lee et al., 2005; Bialek et al. 2008; Borgstein et al., 2009; Chiu et al., 2013), and myostatin propeptide (Bogdanovich et al., 2005; Hamrick et al., 2010b). Published data now exist in which each of these therapies has been evaluated in aged rodents, so that some comparison of the different approaches can be undertaken. Our data using a myostatin propeptide are consistent with data from studies using myostatin antibodies, where these myostatin inhibitors were found to have significant, positive effects on muscle mass, fiber size, and muscle force production. Specifically, LeBrasseur et al. (2009) used a slightly higher dose (25 mg/kg) than we used in our study (20 mg/kg), but also used weekly injections of a myostatin inhibitor (PF-354 antibody) over a period of 4 weeks in mice 24 months of age. They too found a moderate (<10%) in muscle mass and a significant increase in muscle mass relative to body weight (+12–17%), as we did for the tibialis anterior muscle relative to body weight (+~15%). Murphy et al. (2010), like LeBrasseur et al. (2009), used weekly doses of the PF-354 antibody but used a lower dose (10 mg/kg) for a longer treatment period—14 weeks of treatment starting in mice aged 18 months. These authors found increases in overall muscle mass (<10%) in the gastrocnemius.
and quadriceps of the aged mice following 14 weeks of treatment, and a significant increase in (+12%) in muscle fiber cross-sectional area of the tibialis anterior muscle, that were similar in magnitude to the changes we observed with propeptide treatment. Together, these studies using myostatin antibodies and our study using the myostatin propeptide show similar increases in muscle fiber size and muscle mass using either a low dose (10 mg/kg) over a longer (14 week) treatment period, or a higher dose (20–25 mg/kg) over a shorter treatment period (4 weeks).

Data from in vivo studies using either the myostatin antibody or the propeptide differ in two important ways from those utilizing the decoy myostatin receptor (ActRIIB-Fc). First, a 10 mg/kg dose of ActRIIB-Fc administered twice weekly for four weeks increased tibialis anterior mass by 30% and quadriceps mass by 25% (Chiu et al., 2013). These increases in muscle mass are much greater than those observed with either the myostatin antibody or propeptide, which as noted above generated increases in total muscle mass of <10%. It is possible that these differences could be due to the more frequent administration of the ActRIIB-Fc, but the ActRIIB-Fc dose is much lower than that used in either our study or the study by LeBrasseur et al. (2009), suggesting that the ActRIIB-Fc is a more potent molecule for increasing muscle mass in aged mice. The reason for the greater potency of the ActRIIB-Fc for increasing muscle mass is likely because this molecule can bind several ligands in addition to myostatin, including activin, BMP-3, BMP-7, BMP-9, BMP-10, and GDF-11 (Souza et al., 2008). Some of these molecules, such as activin, are also likely to play a role in regulating muscle mass, which is further indicated by the fact that ActRIIB-Fc treatment can increase muscle mass in mice that lack myostatin altogether (Lee et al., 2005). The second way in which our data differ from those using ActRIIB-Fc is related to the effects on bone. ActRIIB-Fc treatment was previously documented to increase bone formation and bone mass in young, growing mice (Bialek et al., 2008; Yan et al., 2008), and the data from Chiu et al. (2013) are consistent with this earlier report in showing that ActRIIB-Fc increases bone density and serum markers of bone formation in aged mice after just 30 days of treatment. In contrast, our data revealed no bone effects with myostatin propeptide treatment. These data may indicate that, as proposed by Chiu et al. (2013), the anabolic effects of ActRIIB-Fc on bone are due to antagonizing effects on ligands other than myostatin, such as various BMPs or activin.

Previous work in our lab showed that myostatin can inhibit the proliferation of aged bone marrow stromal cells (Bowser et al., 2013), that bone marrow stromal cells from mice lacking myostatin show increased proliferation (Elkawrawy et al., 2011), and that myostatin can inhibit chondrogenesis in vivo and in vitro (Elkawrawy et al., 2012). These data may at least in part explain the increased fracture callus size following osteotomy in mice lacking myostatin (Kellum et al., 2009), and increased fracture callus bone volume in mice treated with myostatin propeptide following osteotomy (Hamrick et al., 2010). That is, myostatin seems to play a key role in musculoskeletal injury repair, one in which myostatin secretion from muscle is elevated following muscle damage, and then mediates the repair response in adjacent bone by modulating progenitor cell proliferation (Elkawrawy et al., 2012). On the other hand, myostatin appears to have a more limited role in mature, intact bone. This is indicated by the fact that myostatin itself is not expressed at a significant level by osteoblasts (Digirolamo et al., 2011), and that myostatin inhibition via propeptide treatment in adult mice does not significantly alter osteoblast number, mineralizing surfaces, or bone strength (Tables 1 and 2). Thus, therapeutic targeting of myostatin specifically via antibody or propeptide treatment may have clinical application in the context of improving muscle mass alone, or improving the healing of muscle and bone following injury, but is not likely to have a significant impact on bone formation in the intact, aged animal. In contrast, the decoy myostatin receptor (ActRIIB-Fc) appears capable of increasing muscle mass, bone formation, and bone strength in aged rodents, suggesting that this molecule may have
potential clinical use for age-associated loss of both muscle and bone in the form of sarcopenia and osteoporosis.

Muscle and bone are closely associated spatially and in terms of structure and function during growth, development, and aging. Muscle in particular has been considered a driving force for bone modeling and remodeling, in that muscle is the primary source of mechanical stimuli for bone and bone tissue is thought to adapt its gross structure in response to muscle-derived stimuli. Thus, targeting muscle therapeutically is thought to be one approach for improving bone health, simply by enhancing the mechanical relationship between muscle and bone. On the other hand, a large portion of osteoporotic fractures do not occur in individuals with low bone density as measured by two-dimensional densitometry, and so fall prevention alone may be another strategy for reducing falls and fall-associated morbidity and mortality in the elderly (Jarvinen et al., 2008). Behavioral interventions such as resistance exercise or nutritional interventions such as vitamin D supplementation (Girgis et al., 2013) may improve muscle strength and/or neuromuscular control and proprioception, perhaps reducing fall risk. The extent to which myostatin inhibition may augment such strategies remains relatively unexplored. Mice are relatively small in body weight and their bones are capable of withstanding loads many times their own body mass—for example it takes more than 2 kg of force to fracture the tibia of a 32 g mouse (Fig. 1, Table 2). Thus, increases in muscle mass in these small mammals may not significantly alter the mechanical environment of their bones. Additional studies in patient populations are needed to determine the extent to which therapeutic targeting of muscle alone via a myostatin antibody or propeptide, perhaps in conjunction with an exercise regimen, could reduce the incidence of bone fractures versus a molecule such as ActRIIB-Fc, that may potentially increase the mass and strength of both muscle and bone.

5. Conclusions

We tested the hypothesis that in vivo inhibition of myostatin using an injectable myostatin propeptide (GDF8 propeptide-Fc) would increase both muscle mass and bone density in aged (24 mo) mice. Our goal was to evaluate this potential therapeutic for its capacity to increase both muscle and bone mass in the setting of age-associated sarcopenia and osteoporosis. Mice were injected weekly (20 mg/kg body weight) with recombinant myostatin propeptide-Fc (PRO) or vehicle (VEH; saline) for four weeks. The data show that PRO treatment significantly increases muscle fiber size and muscle mass, both absolutely and relative to body weight. In contrast bone volume, bone strength, and histomorphometric parameters of bone formation and bone resorption were unchanged with PRO treatment. Our findings are consistent with previous studies utilizing a myostatin antibody in aged mice showing that targeting myostatin increases muscle fiber size and mass; however, our data differ from work utilizing a decoy myostatin receptor (ActRIIB-Fc) to inhibit myostatin function in that ActRIIB-Fc appears particularly effective at increasing bone density and bone formation whereas the propeptide does not. The anabolic effects of ActRIIB-Fc on aged bone are likely due to the ability of this molecule to antagonize other ligands besides myostatin, such as activin or bone morphogenetic proteins. Clinical trials evaluating the potential of these molecules to prevent falls and fractures are needed to determine the optimal approaches for reducing musculoskeletal diseases and complications in the elderly.

Acknowledgments

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Figure 1.
Myostatin-induced luciferase activity declines significantly with increasing concentration of myostatin propeptide. The pGL3(CAGA)12 – neo reporter vector contains 12 CAGA boxes previously reported to be TGF-β-responsive elements (Dennler et al. (1998) EMBO J. 17:3091–3100), a neo resistance gene, and the basic luciferase reporter plasmid pGL3 (Promega Corporation).
Figure 2.
Body mass (A) and change in body weight (B) for animals treated with either saline (VEH) or myostatin propeptide (PRO) weekly for over four weeks.
**A**

**Tibias Anterior Mass**

\[ P < 0.05 \]

**B**

**Normalized Tibialis Anterior Mass**

\[ P < 0.01 \]

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Figure 3.
Muscle parameters for mice treated with saline (VEH) or myostatin propeptide (PRO) weekly for a period of four weeks. (A) Tibialis anterior mass, (B) tibialis anterior mass relative to body weight, (C) extensor digitorum longus fiber diameter (EDL), (D) alpha-laminin stained cryostat sections of the EDL, (E) soleus muscle fiber diameter.
Figure 4.
Real-time PCR data for mice treated with saline (VEH) or myostatin propeptide (PRO) weekly for a period of four weeks showing increased expression of Murf1 and Mafbx in PRO-treated mice (A), Western blot showing similar increases in Murf1 and Mafbx with PRO-treatment (B), and increased expression of BMP-2 in mice treated with propeptide (C). *P<.05.
### Table 1

Nucleotide sequences of mouse primers used for RT-PCR

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microCT and biomechanical testing of the proximal tibia for mice treated with saline (VEH) or myostatin propeptide (PRO; 20 mg/kg).

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<td>0.11±0.01</td>
<td>.47</td>
</tr>
<tr>
<td>Tb. N</td>
<td>0.59±0.14</td>
<td>0.54±0.16</td>
<td>.23</td>
</tr>
<tr>
<td>Fu (kg)</td>
<td>2.21±.40</td>
<td>2.18±.34</td>
<td>.39</td>
</tr>
<tr>
<td>U (kg/um²)</td>
<td>740.6±417.5</td>
<td>670.3±309</td>
<td>.31</td>
</tr>
<tr>
<td>S (g/um)</td>
<td>4.6±2.0</td>
<td>4.7±2.0</td>
<td>.44</td>
</tr>
</tbody>
</table>

BMD=bone mineral density, BV/TV=bone volume relative to total volume, Tb.Th=trabecular thickness, Tb.N=trabecular number, Fu=ultimate force, U=energy-to-fracture, S=stiffness.
Table 3
Bone histomorphometry data for the distal femur of mice treated with saline (VEH) or myostatin propeptide (PRO; 20 mg/kg).

<table>
<thead>
<tr>
<th>Parameter</th>
<th>VEH (n=15)</th>
<th>PRO (n=14)</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td>N.Ob/BS</td>
<td>27.26±17.49</td>
<td>25.09±9.31</td>
<td>.14</td>
</tr>
<tr>
<td>MS/BS</td>
<td>0.41±0.17</td>
<td>0.43±0.13</td>
<td>.34</td>
</tr>
<tr>
<td>N.Oc/BS</td>
<td>6.33±2.61</td>
<td>6.18±3.82</td>
<td>.38</td>
</tr>
</tbody>
</table>

N.Ob/BS=osteoblast number per bone surface, MS/BS=mineralizing surface (single-label) relative to bone surface, N.Oc/BS=osteoclast number per bone surface.
Myostatin (GDF-8) Signaling in Progenitor Cells and Applications to Bone Repair

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Word count:
5892 words
1 tables
5 figures
68 references
<table>
<thead>
<tr>
<th>Abbreviation</th>
<th>Description</th>
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<tr>
<td>ActRIIB</td>
<td>type IIB activin receptor</td>
</tr>
<tr>
<td>Acvr2b</td>
<td>type IIB activin receptor</td>
</tr>
<tr>
<td>Alk4/5</td>
<td>activin-like kinase receptor 4 and 5</td>
</tr>
<tr>
<td>BMP-9, -10, 11</td>
<td>bone morphogenetic protein-9, -10, -11</td>
</tr>
<tr>
<td>BMSC</td>
<td>bone marrow stromal cell</td>
</tr>
<tr>
<td>Dkk1</td>
<td>gene encoding Dickkopf-related protein 1</td>
</tr>
<tr>
<td>ERK</td>
<td>extracellular signal-regulated kinase</td>
</tr>
<tr>
<td>FLRG</td>
<td>follistatin-related gene</td>
</tr>
<tr>
<td>Fzd3</td>
<td>gene encoding the frizzled-3 protein</td>
</tr>
<tr>
<td>GDF-8</td>
<td>growth and differentiation factor-8, or myostatin</td>
</tr>
<tr>
<td>GDF-11</td>
<td>growth and differentiation factor-11</td>
</tr>
<tr>
<td>Gsk3a</td>
<td>gene encoding glycogen synthase kinase 3 alpha</td>
</tr>
<tr>
<td>MAPK</td>
<td>mitogen-activated protein kinase</td>
</tr>
<tr>
<td>microCT</td>
<td>micro-computed tomography</td>
</tr>
<tr>
<td>TGF-β</td>
<td>transforming growth factor-beta</td>
</tr>
<tr>
<td>Wnt4</td>
<td>Wingless-type MMTV integration site family, member 4</td>
</tr>
<tr>
<td>Wnt5a</td>
<td>Wingless-type MMTV integration site family, member 5a</td>
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Abstract

Myostatin is a member of the TGF-β superfamily of growth and differentiation factors and is widely recognized as a potent regulator of skeletal muscle mass and regeneration. It has, however, recently been shown that myostatin is expressed in tissues aside from muscle including fat and tendon. In addition, studies monitoring gene expression in the fracture callus following injury have shown that myostatin is highly expressed in the early phases of fracture healing. Furthermore, other research has demonstrated that mice lacking myostatin show increased bone density and a marked increase in fracture callus size following fibula osteotomy. Yet, the basic mechanisms by which myostatin may impact bone healing after injury have not been fully elucidated. Our in vitro studies using 3D aggregate culture to induce chondrogenic differentiation of bone marrow derived stromal cells (BMSCs) reveal that myostatin suppresses the proliferation and chondrogenic differentiation of BMSCs by altering the expression of Sox-9 and various Wnt-related factors. These in vitro studies suggest that myostatin may directly impair bone regeneration, a hypothesis that is also supported by our recent finding that exogenous myostatin reduces fracture callus bone volume. Together, the data suggest that myostatin may represent a novel therapeutic target for management of orthopedic trauma where both bone and muscle are damaged and, furthermore, that myostatin inhibitors may enhance fracture healing and improve recovery following musculoskeletal injury.
Introduction

Myostatin (GDF-8) was first identified as a new member of the transforming-beta growth factor (TGF-beta) superfamily of growth and differentiation factors that was highly expressed in skeletal muscle progenitors (McPherron et al. 1997). The functional characterization of myostatin using a myostatin-knockout mouse led to the surprising observation that loss of normal myostatin signaling produced a dramatic increase in muscle mass (McPherron et al. 1997). Since the original discovery of myostatin, its role in muscle development and muscle regeneration has been explored using a variety of in vivo and in vitro approaches (Zimmers et al. 2002, Reis-Porszaz et al. 2003, Li et al., 2008, McFarlane et al., 2011). These studies have generally shown that myostatin plays an important role in the proliferation and differentiation of myoblasts, and that inhibiting myostatin expression improves muscle regeneration and increases muscle fiber size and number (Lee 2004, Burks and Cohn 2011).

It is now clear, however, that myostatin is expressed in many tissues aside from muscle (Allen et al. 2008, Mendias et al. 2008), and that it has important functions in cells ranging from fibroblasts to adipocytes. This raises the possibility that myostatin is involved in the differentiation of mesenchymal stem cells (MSCs), since mice lacking myostatin show significant alterations in tissues of mesenchymal origin. Specifically, myostatin deficient mice show decreased body fat (McPherron and Lee 2002, Hamrick et al. 2006), increased bone mineral density (Elkasrawy and Hamrick 2010), and tendons and ligaments that are relatively weak and exhibit cellular hypoplasia (Mendias et al. 2008, Fulzele et al. 2010). In addition, other studies show that the myostatin receptor (Acvr2b, or ActRIIB) is expressed in bone marrow-derived stromal cells (Hamrick et al. 2007), and myostatin can regulate the differentiation of C3H 10T(1/2) mouse mesenchymal multi-potent cells (Rebbapragada et al. 2003, Artaza et al. 2005). Finally, GDF-11, a member of the TGF-β superfamily that is structurally and functionally very similar to myostatin (McPherron et al. 2009), has been shown to decrease chondrogenic and myogenic differentiation of MSCs in micromass cultures (Gamer et al., 2001).
Previous work therefore suggests that myostatin is an important factor in regulating not only the fate of mesenchymal progenitors but also the proliferation, differentiation, and function of mesenchymal derivatives such as adipocytes, chondrocytes, myoblasts, and fibroblasts. These observations have obvious implications for bone repair, since myostatin is also expressed in bone during the early phases of fracture healing (Cho et al. 2002). This chapter reviews the basic biology of myostatin signaling, its role in mediating the proliferation and differentiation of mesenchymal progenitors, and the effects of myostatin and myostatin inhibition on bone repair.

**Myostatin Signaling Through the Type IIB Activin Receptor (Acvr2b, or ActRIIB)**

Myostatin circulates in a latent form bound to a propeptide, which is cleaved by a BMP1/Tolloid matrix metalloproteinase, releasing the active form capable of binding the type IIB activin receptor. Follistatin and follistatin–related gene (FLRG) are other proteins that can bind and inhibit the activity of myostatin by maintaining its latency (Lee 2004, Tsuchida 2004). Myostatin has previously been shown to bind the type IIB activin receptor (Acvr2b, or ActRIIB) with high affinity (Lee 2004), with activin itself having a higher affinity for the type IIa activin receptor (Donaldson et al. 1992). Other ligands are, however, also known to bind Acvr2b along with myostatin. In mice these include BMP-9, -10, and -11 along with Inhibinβ and TGF-β1. In human serum the same ligands were found bound to Acvr2b, with the exception of TGF-β1 (Souza et al. 2008).

Acvr2b is a serine/threonine kinase receptor, which recruits and phosphorylates activin-like kinase receptor 5 (ALK5) or ALK4 (Rebbapragada et al 2003). This consequently activates and phosphorylates Smad2/3, which dissociate from the ligand/receptor complex to bind with the co-Smad, Smad4, allowing translocation of the Smad complex to the nucleus where it targets several DNA binding proteins to regulate transcriptional response (Lee 2004) (Fig. 1). Smad4 also activates Smurf-mediated ubiquitination of the ActRIIB/ALK4-5 receptor complex, and promotes its proteosomal degradation (Bradley et al. 2008). Although myostatin, like other
TGF-β family members, is most well known to activate Smad signaling, it can also activate the p38 MAPK and ERK1/2 pathways (Allendorph et al. 2006, Steelman et al. 2000, Ekaza and Cabello 2007). The p53 MAPK and ERK1/2 pathways work on promoting the survival of mature muscle synthetia by the action of p53, which blocks apoptosis. ERK1/2 also activates p21, which arrests myoblast proliferation (Bradley et al., 2008). Autocrine myostatin signaling is regulated in part through a negative feedback loop, where myostatin activation of smad2/3 stimulates expression of the inhibitory smad Smad7, which in turn inhibits myostatin signal transduction (Zhu et al. 2004).

A number of studies now provide evidence that myostatin may also signal through the Wnt pathway. For example, Wnt4 expression is significantly elevated in skeletal muscle from myostatin-deficient mice (Steelman et al. 2006), and myostatin enhances nuclear translocation of beta-catenin and formation of the Smad3-beta-catenin-TCF4 complex in human mesenchymal stem cells (Guo et al., 2008). These findings are significant from the perspective of bone healing and regeneration, since components of the Wnt pathway such as Wnt4, Wnt5a, and Wnt5b, are known to be upregulated during the process of fracture repair (Hadjigyrou et al. 2002; Zhong et al. 2006). In addition, beta-catenin-mediated Wnt pathway signaling is also involved in endochondral bone formation (Zhong et al. 2006; Chen et al. 2007a, 2007b). The Wnt pathway ultimately regulates the levels of beta-catenin in the cytoplasm and promotes its nuclear translocation, where it acts in conjunction with other transcription factors to promote different gene transcription (Akiyama 2000).

**A Role for Myostatin in Regulating Stem Cell Differentiation**

A growing body of evidence reveals that myostatin plays a role in regulating myogenesis, adipogenesis and tissue fibrosis by altering the differentiation of progenitor cells of mesenchymal origin. Given the double-muscled phenotype of mice lacking myostatin, it is not surprising that myostatin suppresses cell proliferation during myogenic differentiation of C2C12 progenitors (Taylor et al. 2001, Rios et al. 2001). Myostatin also induces the differentiation of
myofibroblasts in C3H 10T1/2 mesenchymal stem cells (Artaza et al. 2008). In vivo, inhibition of myostatin using an antibody decreases apoptosis and caspase-3 expression in skeletal muscle (Murphy et al. 2010). Myostatin likewise increases proliferation and extracellular matrix synthesis of primary muscle fibroblasts (Li et al. 2008), as well as fibroblasts from human knee ligaments (Fulzele et al. 2010). These data indicate that myostatin is a pro-fibrotic factor that contributes to the accumulation of collagen-rich fibrous tissue after muscle injury (Zhu et al. 2007).

While myostatin appears to inhibit myogenesis and stimulate fibrosis, it also inhibits adipogenic differentiation of both human bone marrow-derived stromal cells (Guo et al, 2008) and adipogenic differentiation of mouse pre-adipocytes (Kim et al. 2001) and mesenchymal stem cells (Rebbapragada et al. 2003). In human stromal cells, myostatin induced nuclear translocation of beta-catenin, altered the expression of several Wnt/beta-catenin pathway genes, and suppressed expression of the adipogenic transcription factors PPAR gamma and C/EBP alpha (Guo et al 2008). In mouse MSCs and pre-adipocytes, adipogenesis was induced with BMP-7, and myostatin treatment suppressed this BMP7 induced effect (Rebbapragada et al. 2003). The inhibitory effect of myostatin on adipogenesis in these experiments could be reversed by blocking ALK5. In contrast, other studies have shown that myostatin can directly stimulate adipogenesis in C3H10T1/2 cells (Artaza et al. 2005, Feldman et al 2006). These apparently contradictory findings are difficult to reconcile, particularly when congenital absence of myostatin is associated with a dramatic decrease in body fat (Lin et al. 2002, McPherron and Lee 2002). Further investigation into myostatin’s role in adipocyte differentiation is clearly needed.

We have previously shown that absence of myostatin increases the osteogenic differentiation of primary bone marrow stromal cells from mice (Hamrick et al. 2007). This finding is perhaps consistent with some of the in vitro experiments above demonstrating that myostatin may favor the adipogenic differentiation of mesenchymal progenitors; hence, absence
of myostatin or myostatin inhibition would be expected to direct MSCs toward the osteogenic or myogenic lineages. This hypothesis is supported by the finding that myostatin inhibitors increase bone formation and bone mass in mice (Bialek et al. 2008, Ferguson et al. 2009), and also improve bone repair following osteotomy (Hamrick et al. 2010). We have also found that myostatin treatment inhibits the chondrogenic differentiation of bone marrow-derived stromal cells by suppressing the expression of Sox-9, and by altering the expression of a number of Wnt-related genes. Specifically, exogenous myostatin reduced the expression of type II collagen and the proliferation of growth plate chondrocytes as well as bone marrow stromal cells in vitro (Elkasrawy et al. 2011a). The Wnt signaling factor Dkk1 is significantly upregulated with myostatin treatment during TGF-β1-induced chondrogenesis, and GSK3A (responsible for beta-catenin instability) is also upregulated, whereas the Wnt ligand binding protein Fzd3 is significantly downregulated with myostatin treatment (Fig. 2). Dkk1 is a molecule that inhibits the Wnt signaling pathway by binding to and antagonizing LRP5/6, forming a ternary complex with LRP6 that induces its rapid endocytosis and removal from the plasma membrane (Mao et al. 2002). GSK3A is another factor that is involved in Wnt signaling, which is a component of the GSK3/Axin/APC proteosomal complex responsible for ubiquitination of β-catenin to prevent its accumulation in the cytoplasm and translocation into the nucleus. Fzd3, a transmembrane protein that can bind Wnt ligands including Wnt5a, and Wnt5a is known to be a molecule that plays a key role in chondrogenesis (Church et al. 2002, Yang et al. 2003), is downregulated in the presence of myostatin during TGF-β1-induced chondrogenesis. In addition, as noted above, GDF-11, a member of the TGF-β superfamily that is structurally very similar to myostatin (McPherron et al. 2009), has been shown to decrease chondrogenic and myogenic differentiation of MSCs in micromass cultures (Gamer et al. 2001).

TGF-β1 is a growth factor that is well known to enhance several key processes in the chondrogenic pathway, including cellular condensation, adhesion, and extracellular matrix production during chondrogenesis (James et al. 2009). TGF-β1 and myostatin can both bind the
same co-receptor, ALK5 (Rebbapragada et al. 2003, Derynck and Feng 1997). We have shown that Sox-9 was down regulated with myostatin treatment, a treatment effect that was also seen with ALK5 inhibition during TGF-β1-induced chondrogenesis (Elkasrawy et al. 2011a). We also showed that GSK3α, responsible for beta-catenin instability, is upregulated with myostatin treatment during TGF-β1-induced chondrogenesis (Fig. 2). These data suggest that myostatin could be competitively inhibiting ALK5-mediated TGF-β1 signaling (Figure 3). TGF-β signaling up-regulates Sox-9 and Wnt5a, and increases nuclear accumulation and stability of beta-catenin during chondrogenesis (Zhou et al., 2004; Kawakami et al., 2006; Lorda-Diez, 2009). Fzd3 is a transmembrane proteins that bind Wnt ligands (Chen and Struhl, 1999). These experimental findings indicate that myostatin might be capable of competitively inhibiting ALK5-mediated TGF-β1 signaling, or alternatively that myostatin may stimulate the expression of certain factors that could actively inhibit TGF-β1, such as Smad7 (e.g. Zhu et al. 2004; Iwai et al. 2008).

**Implications for Bone Repair**

The first indication that myostatin might play a key role in bone repair and regeneration came from our work in myostatin-knockout mice, which revealed that fracture callus bone volume was significantly increased in mice lacking myostatin following fibular osteotomy (Fig. 4, Kellum et al. 2009). The idea that myostatin might in some way suppress or inhibit bone healing was further supported by another study in which we showed that a myostatin inhibitor (propeptide) increased fracture callus bone volume in mice two weeks following osteotomy (Hamrick et al. 2010a). Fracture healing in rodents involves three key phases that have been well-described by numerous authors: an initial inflammatory phase, a chondrogenic phase, and an osteogenic phase. The in vivo findings in mice lacking myostatin and in mice treated with a myostatin inhibitor are generally supportive of the in vitro data referenced above, suggesting that myostatin acts to suppress chondrogenesis directly (Table 1). Our immunolocalization studies showed that myostatin is highly expressed in the region of the wound blastema 12 and 24 hours following osteotomy, and that exogenous myostatin reduces fracture callus bone volume.
(Elkasrawy et al. 2011b). This is one mechanism by which myostatin may influence bone repair (Fig. 5, Table 1), but we believe there are also several others which we describe below. In addition to suppressing chondrogenesis during fracture healing, myostatin may also mediate the differentiation of muscle-derived progenitors that contribute to callus formation. It has been recognized for decades that the muscle bed itself plays a crucial role in fracture healing (Pritchard and Ruzicka 1950), and that cells derived from muscle may in fact migrate into the site of injury and promote bone healing (Schindeler et al. 2009, Liu et al., 2010). Cells harvested from muscle after exposure to an adjacent fracture are highly osteogenic and form bone nodules in vitro (Glass et al. 2011). Thus, muscle not only facilitates bone repair by providing a vascular bed and perhaps trophic factors (see below), but it is a possible source of progenitor cells that can differentiate into osteoblasts, further improving bone healing (Glass et al. 2011). Given that myostatin is a potent factor that can mediate the differentiation of muscle-derived cells, it is likely that elevated levels of myostatin following injury may drive muscle-derived stromal cells toward a myofibroblast lineage rather than toward an osteogenic fate. In this way local or systemic myostatin could inhibit bone healing by limiting the supply of osteoprogenitors available from the pool of muscle-derived stromal cells (Fig. 5).

A third manner in which myostatin may alter bone healing is by regulating muscle regeneration and hence the secretion of muscle-derived paracrine factors (“myokines”; Hamrick 2011). Studies in which skeletal muscle is transplanted into cardiac muscle support the notion that myofibers secrete a number of paracrine factors that are not only osteogenic (Hamrick 2011) but also promote the survival of neighboring muscle fibers (“paracrine theory of transplantation”; Perez-Ilzarbe et al. 2008). Factors that are actively secreted by myotubes in vitro include insulin-like growth factor 1 (IGF-1), osteonectin (SPARC), and basic fibroblast growth factor (FGF-2) (Hamrick, 2011). All of these factors have been observed to increase bone formation when injected peripherally, and deficiency of these factors is associated with low bone mass (Hamrick, 2011). Elevated levels of myostatin impair muscle regeneration and
induce muscle atrophy, which would in turn be expected to reduce the secretion of factors normally produced by healthy myofibers such as IGF-1 and FGF-2 (Hamrick et al, 2010b). Recently it was also shown that mice lacking myostatin have elevated serum levels of IGF-1, and that this was associated with elevated expression of IGF-1 in the liver (Williams et al. 2011). Thus, elevated secretion of myostatin, which is known to occur with disuse, infection, burns, or AIDS- and cancer-related cachexia, may indirectly impair bone healing and bone formation by decreasing circulating levels of liver-derived IGF-1 (Williams et al. 2010).

Summary
Since the discovery of myostatin in 1997 our understanding of its biology and function has grown considerably so that we now recognize a broad range of potential roles for myostatin in musculoskeletal growth, development, and regeneration. These include effects of myostatin on tendon and ligament development and healing (Mendias et al. 2008, Eliasson et al. 2009, Fulzele et al. 2010), adipogenesis and fat deposition (McPherron and Lee 2002, Hamrick et al. 2006), and bone density and bone healing (Morissette et al. 2009; Kellum et al. 2009), in addition to the well-documented effects of myostatin on muscle mass and regeneration (Lee 2004). These and other in vivo and in vitro studies now point to a role for myostatin in the proliferation and differentiation of mesenchymal progenitors. These effects are mediated by TGF-beta signaling through the type IIB activin receptor, involving phosphorylation of Smad2/3, but our own work in chondrocytes and the work of others in MSCs and myoblasts suggest that myostatin also signals through the Wnt pathway. Various myostatin inhibitors have, not surprisingly, been observed to increase muscle mass, decrease fat mass, and increase bone formation. These findings suggest that myostatin represents a potential therapeutic target for improving tissue repair and regeneration in musculoskeletal diseases and following musculoskeletal injury.
Acknowledgments

Many of the findings discussed in this chapter are derived from our research on myostatin which has been supported by the National Institutes of Health (AR049717), the Office of Naval Research (N000140810197), and the Department of the Army (USAMRMC PR093619). We are grateful to Drs. Paul Yaworsky, Alexandra McPherron, and Li-Fang Liang for many helpful discussions regarding the biology of myostatin. Ethan Kellum, Phonepasong Arounleut, Matthew Bowser, David Immel, Craig Byron, Penny Roon, Donna Kumiski, and Cathy Pennington provided valuable technical assistance and support for many of the studies reviewed in this paper.
Summary Points

- Myostatin, also known as growth and differentiation factor-8 (GDF-8), was discovered in 1997.

- Myostatin plays a number of roles in musculoskeletal growth, development, and regeneration.

- These include effects of myostatin on tendon and ligament development and healing, adipogenesis and fat deposition, and bone density and bone repair.

- Blocking myostatin signaling is well-known to increase muscle mass and improve muscle regeneration.
Dictionary

**Wnt pathway:** The wnt signaling pathway is most well known for its role in regulating cell fate and cell-cell interactions during embryonic development in both vertebrates and invertebrates. The term “wnt” refers to the name of the gene implicated in the wingless mutation in fruit flies (Drosophila).

**Endochondral ossification:** Long bones such as the humerus and femur of the mammalian skeleton first form as condensations of cartilage, which then undergo a process of ossification to form a bony skeleton. This process is referred to as endochondral ossification, and a similar process occurs during bone healing, where a “soft” callus of cartilage is replaced by a “hard” bony callus.

**Bone marrow stromal cells:** Bone marrow includes a variety of cell types, including cells that can give rise to tissue types of mesodermal origin (e.g., muscle, fat, bone). These cells in bone marrow are referred to as bone marrow stromal cells.

**Myokines:** It is now known that muscle can secrete a number of different growth factors and cytokines that can affect other organs, and these muscle-derived factors are referred to as myokines.
References


Guo, W. et al. 2008. The effects of myostatin on adipogenic differentiation of human bone marrow-derived mesenchymal stem cells are mediated through cross-communication between Smad3 and Wnt/beta-catenin signaling pathways. J Biol. Chem. 283: 9136-9145.


Table 1. Key Facts Related to Myostatin and Bone Healing

1. The receptor for myostatin is expressed in bone marrow stromal cells, and chondrocytes as well as muscle cells express myostatin.

2. Following traumatic musculoskeletal injury, myostatin is most highly expressed by injured muscle cells in the first 24-48 hours following the injury.

3. Congenital absence of myostatin increases the volume of cartilage and bone in the fracture callus, whereas delivery of exogenous myostatin impairs muscle and bone healing.

4. In vitro experiments show that myostatin can inhibit the chondrogenic differentiation of bone marrow stromal cells.

5. Blocking myostatin signaling immediately following musculoskeletal injury is therefore a potential therapeutic strategy for improving the healing of both muscle and bone.

Legend to Table 1. This table lists key facts that are related to the role of myostatin in bone repair, including direct roles for myostatin on osteo- and chondroprogenitors, and indirect roles mediated by muscle-derived factors.
Figure Caption Titles and Legends

Figure 1 Title. The myostatin signaling pathway.

Figure 1 Legend. Myostatin (M) binds the type IIB activin receptor (ActRIIB) which recruits the type I co-receptor ALK5, phosphorylating Smad2/3 which translocate to the nucleus where they regulate target gene expression. A number of factors thought to be involved in the proliferation and differentiation of mesenchymal stem cells are regulated through this TGF-beta signaling pathway.

Figure 2 Title. Effects of myostatin treatment on gene expression.

Figure 2 Legend: Exposure of bone marrow-derived mesenchymal stem cells to exogenous myostatin alters the expression of several factors involved in the Wnt signaling pathway.

Figure 3 Title. Hypothetical model showing mechanisms by which myostatin may mediate TGF-beta induced chondrogenesis.

Figure 3 Legend. TGF-beta 1 and myostatin compete for recruitment of ALK5. TGF-beta stimulates Wnt5a activation of the chondrogenic cascade, whereas myostatin may attenuate this effect.

Figure 4 Title. Effects of myostatin deficiency on fracture healing.

Figure 4 Legend. MicroCT images of the fibula fracture callus in normal mice (left two panels) and mice lacking myostatin (right two panels).

Figure 5 Title. General model summarizing effects of myostatin inhibition (left) and myostatin expression (right) on musculoskeletal repair and regeneration.

Figure 5 Legend. Blocking myostatin signaling following injury of muscle and bone is likely to enhance bone healing in two ways (left side of figure): directly, because suppression of
myostatin activity will enhance chondrogenesis and indirectly, because enhancing muscle
regeneration is likely to restore the normal secretion of paracrine, osteogenic trophic factors
(e.g., IGF-1) from muscle. Myostatin expression following musculoskeletal injury inhibits bone
healing in two ways (right side of figure), first by suppressing chondrogenesis directly and
second by increasing fibrosis in injured muscle thereby reducing secretion of osteogenic
paracrine factors from muscle tissue.
Figure 1
Figure 2
Figure 3
Figure 4
Figure 5

Myostatin inhibitor

- Direct regulatory effects
- Accelerated Fracture repair
- Improved muscle regeneration

Musculoskeletal injury

Myostatin expression

- Direct regulatory effects
- Fibrotic tissue formation
- Delayed or non-union

Indirect effect
REVIEW

The skeletal muscle secretome: an emerging player in muscle–bone crosstalk

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In vitro and in vivo studies provide evidence that a variety of growth factors and cytokines are actively secreted by muscle tissue. Muscle can therefore function as an endocrine and paracrine organ. These peptides characterize the muscle secretome, and many muscle-derived factors such as insulin-like growth factor-1, basic fibroblast growth factor, interleukin-15, myostatin and secreted protein acidic and rich in cysteine (osteonectin) are also known to have significant effects on bone metabolism. The factors secreted by muscle may vary according to muscle activity, in that muscle contraction, muscle atrophy or traumatic muscle injury can alter the type and relative abundance of particular factors released from muscle cells. The molecular and cellular pathways by which muscle-derived factors affect different types of bone cells (for example, osteoblasts, osteoclasts and osteocytes) are, however, poorly understood. Nevertheless, these findings further underscore the complex nature of muscle–bone interactions, and highlight the importance of integrating muscle biology and physiology into our understanding of bone growth, development and aging.

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latter hypothesis for the following reasons. First, it is clear that intact muscle flaps are a rich source of secreted growth factors. Second, although muscle does provide important vascular support for bone, skin has a higher vascular density than muscle, yet fasciocutaneous skin flaps do not have the same anabolic effects on healing bone as muscle flaps. Third, it is clear that conditioned medium from cultured muscle cells has positive effects (for example, increasing extracellular matrix synthesis) on cultured chondrocytes, revealing that muscle is a source of secreted growth factors both in vivo and in vitro. Finally, the muscle secretome has become increasingly well characterized. We now know that muscle secretes a wide variety of growth factors, cytokines and molecules involved in extracellular matrix remodeling. Moreover, different laboratories have independently identified the same factors secreted from muscle, further validating the existence of a well-defined muscle secretome. Some of these factors have been referred to as ‘myokines’ by different authors; however, several muscle-derived factors (for example, IGF-1, FGF-2, IL-6) are secreted in abundance by other tissues, so the term myokine should not be taken to imply that they are muscle-specific.

Our growing knowledge of the muscle secretome has important implications for bone biology, as it presents new opportunities for targeting muscle in order to better develop the therapeutic program for aging and healing bone. This review seeks to highlight the muscle-derived factors that may impact bone metabolism, and also propose future directions for research aimed at advancing our current understanding of muscle–bone crosstalk.

Growth Factors Actively Secreted by Skeletal Muscle

Insulin-like growth factor-1 (IGF-1). Wound exudates from intact muscle flaps contain high levels of IGF-1 and immunohistochemistry has been used to localize IGF-1 along the muscle–periosteal interface of mouse forelimbs. These in vivo findings are consistent with in vitro studies in which proteomic approaches have been used to detect IGF-1 in conditioned medium from cultured myotubes. IGF-1 expression, measured as an increase in IGF-1 mRNA, is increased in skeletal muscle with muscle contraction, and elevated levels of IGF-1 protein are detected in serum following resistance exercise. It is well established that IGF-1 has important osteogenic effects on the skeleton and IGF-1 is also involved in myofiber hypertrophy, suggesting that muscle-derived IGF-1 may couple both muscle and bone anabolism.

Basic fibroblast growth factor (FGF-2). FGF-2 lacks the conventional signal sequence for export out of cells via the classic exocytotic pathway, and it has been shown that mechanically induced plasma membrane disruption is one mechanism by which FGF-2 is released from myocytes both in vivo and in vitro. Eccentric, lengthening contractions are particularly effective for releasing FGF-2 stored in the cytosol of muscle cells, and levels of FGF-2 detected in conditioned medium from cultured myotubes are increased with mechanical stretching. It is important to note here that the muscle ‘injury’ that occurs during eccentric muscle contraction involves plasma membrane disruptions that are followed by membrane repair. These types of mechanical disruptions are argued here to involve a different cascade of molecular signaling events than traumatic muscle injuries that are associated with muscle regeneration. The latter type of mechanical and structural disruption is much more severe, and involves cleanup of necrotic tissue by macrophages, expression of inflammatory factors and activation of satellite cells. FGF-2 has positive effects on bone formation in estrogen-deficient rodents and is a well-known osteogenic factor. Mechanically induced release of FGF-2 following eccentric contraction and plasma membrane disruption is another potential pathway by which physical activity and bone formation may be coupled physiologically.

Myostatin (GDF-8). Although factors such as IGF-1 and FGF-2 are secreted by a number of tissues in addition to muscle, myostatin is most abundant in muscle tissue and appears to be secreted primarily by muscle. Thus, myostatin can be considered a bona fide myokine. Conditions associated with elevated levels of myostatin expression include disuse atrophy, cancer and AIDS-related cachexia and increased circulating levels of glucocorticoids. Myostatin treatment induces muscle wasting and myostatin deficiency increases muscle mass. We have recently shown that myostatin is highly expressed by injured myofibers following traumatic extremity injury, and that local application of exogenous myostatin increases skeletal muscle fibrosis and inhibits bone repair. These data are consistent with studies referenced above, demonstrating that intact muscle flaps enhance bone repair, whereas coverage of bone injuries with damaged muscle does not have the same positive effects on bone healing. These in vivo data are also consistent with our in vitro studies showing that myostatin treatment suppresses the proliferation and chondrogenic differentiation of bone marrow-derived stromal (stem) cells. In addition, blocking the myostatin using a recombinant propeptide improves muscle regeneration and fracture healing following orthopaedic trauma. Although severe muscle trauma and muscle damage increase myostatin expression, which in turn impairs bone healing, eccentric muscle contraction and exercise both decrease myostatin expression in skeletal muscle. These studies suggest that although IGF-1 and FGF-2 are muscle-derived factors that can have significant, positive effects on bone formation, myostatin is a factor released from muscle during traumatic
Table 1 Growth factors, cytokines and other peptides secreted by muscle, the factors that influence their secretion and their potential effects on bone metabolism

<table>
<thead>
<tr>
<th>Muscle-derived peptides</th>
<th>Factors that stimulate peptide secretion</th>
<th>Role(s) in bone metabolism</th>
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<tr>
<td>Growth factors</td>
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<tr>
<td>IGF-1</td>
<td>Resistance exercise</td>
<td>Stimulates bone formation</td>
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<tr>
<td>FGF-2</td>
<td>Eccentric muscle contraction</td>
<td>Stimulates bone formation</td>
</tr>
<tr>
<td>GDF-8</td>
<td>Muscle damage, cachexia, atrophy</td>
<td>Suppresses chondrogenesis and fracture healing</td>
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<tr>
<td>Extracellular matrix molecules</td>
<td></td>
<td>Promotes bone mineralization, fracture callus remodeling, bone formation Cleaving of procollagen and possibly heterotopic ossification</td>
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<tr>
<td>SPARC</td>
<td>Resistance exercise, muscle regeneration</td>
<td></td>
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<tr>
<td>MMP-2</td>
<td>Resistance exercise and re-loading</td>
<td></td>
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<tr>
<td>BMP-1</td>
<td>Blast trauma to muscle</td>
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<td>Inflammatory cytokines</td>
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<tr>
<td>IL-6</td>
<td>Physical activity and muscle contraction</td>
<td>Bone resorption and turnover</td>
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<tr>
<td>IL-7</td>
<td>Physical activity and muscle contraction</td>
<td>Bone resorption</td>
</tr>
<tr>
<td>IL-15</td>
<td>Resistance exercise</td>
<td>Increase bone mass, decrease adiposity</td>
</tr>
</tbody>
</table>

Abbreviation: IL, interleukin.

and catabolic conditions that may inhibit and suppress bone repair (Figure 1).

Factors Involved in Extracellular Matrix Remodeling Secreted by Muscle

Secreted protein acidic and rich in cysteine (SPARC, or osteonectin). SPARC is a glycoprotein that is abundant in the extracellular matrix of various tissues including bone and skeletal muscle and is involved in tissue repair, remodeling of the extracellular matrix, and promoting collagen mineralization by osteoblasts. One of the more surprising aspects of research on the muscle secretome is the consistency with which SPARC is detected as a factor secreted by isolated muscle cells. A number of different research groups using both human and rodent-derived muscle cells have identified SPARC in conditioned medium from cultured myotubes. SPARC secretion is increased following resistance exercise and myotube hypertrophy, but SPARC is also highly expressed following injury and during muscle regeneration. Additional research is needed to determine the role of muscle-derived SPARC in bone formation, but the studies reviewed above suggest that exercise-induced SPARC secretion could potentially have a role in enhancing bone formation and mineralization (Table 1).

Matrix metalloproteinase-2 (MMP-2). Mice lacking MMP-2 are known to show bone loss and reduced bone density, and absence of MMP-2 effects later stages of fracture callus remodeling. MMP-2, similar to SPARC, is found in both muscle as well as bone, and studies using cultured myotubes reveal that MMP-2 is actively secreted by myotubes in vitro. Insulin treatment of rat myotubes increases secretion of MMP-2, and serum levels of MMP-2 are elevated in diabetic patients suffering from hyperinsulinemia. MMP-2 expression increases with exercise and with re-loading following hindlimb suspension, and decreases with injury, but is elevated in skeletal muscle following disuse. These data show that MMP-2, like SPARC, has an important role in muscle tissue remodeling during various repair events. The role of muscle-derived MMP-2 in bone metabolism is also unclear, but muscle-derived MMP-2 could potentially couple both muscle and bone turnover (Table 1).

Bone morphogenetic protein-1 (BMP-1). BMP-1 is not a true bone morphogenetic protein but rather is a protease that cleaves the propeptide fragments ofprocollagens I, II and III. BMP-1 is secreted from cultured primary human myotubes in vitro, and its secretion is decreased in rat myotubes exposed to very high contractions of insulin. Recently, high levels of BMP-1 protein and mRNA were detected in muscle biopsies from patients who had experienced blast trauma in the combat setting. This is significant from the perspective of muscle–bone crosstalk because blast trauma is associated with a high incidence of heterotopic ossification, a condition where bone forms within muscle tissue. Additional work is needed to better understand the role(s) of muscle-specific BMP-1 secretion in normal and pathological bone formation, but BMP-1 could represent a potential therapeutic target for the prevention of heterotopic ossification (Table 1).

Inflammatory Factors that are Secreted from Muscle During Exercise

Interleukin-6, -7 and -15 (IL-6, IL-7, IL-15). The term ‘myokine’ was originally coined in reference to IL-6, a factor that Pedersen and colleagues determined was released from muscle during exercise, and had important effects on other tissues including the liver and adipose depots. Type I (slow-twitch) fibers express high levels of IL-6, and IL-6 levels are increased in serum with exercise. IL-6 can stimulate expression of the anti-inflammatory factor IL-10, and mice lacking IL-6 develop obesity and insulin resistance. IL-6 is often considered a pro-resorptive cytokine for bone, but mice lacking IL-6 do not show an osteopenic phenotype and IL-6 may facilitate bone formation during conditions of high bone turnover. IL-7 is also widely considered to be an osteoclastogenic cytokine, and IL-7 is actively secreted by muscle cells. Interestingly, many of the studies cited above (for example, Chan et al., Norheim et al., Henningsen et al., Yoon et al.) that utilized in vitro cultures of myotubes to characterize the muscle secretome failed to identify IL-6 and -7 in conditioned medium. These observations raise the possibility that there are other myokines left to be identified that may be discovered through novel alternative in vivo and in vitro approaches. Finally, IL-15 is highly expressed in muscle tissue and is upregulated following resistance exercise. Transgenic mice overexpressing IL-15 in skeletal muscle that show elevated circulating IL-15 levels also show decreased fat mass and increased bone mass. Importantly, these mice did not differ in lean mass or body weight from normal controls, suggesting that the increased bone mass was not due to any alterations in mechanical factors.

Summary and Future Research

A paracrine role for skeletal muscle is not necessarily a new concept, as experiments where skeletal muscle was trans-
planted into cardiac tissue have revealed that skeletal muscle implants are a source of trophic factors supporting the survival of surrounding myocardial cells. The paracrine and endocrine effects of muscle on bone are, however, only now beginning to become more well defined. We are at a very early stage in our understanding of how the muscle secretome impacts bone and other organs, and hence a number of outstanding questions remain. These include questions as to how muscle-derived factors impact particular cell types. For example, how do factors secreted from myocytes during muscle contraction and muscle hypertrophy influence bone resorption by osteoclasts, bone formation by osteoblasts or the adipogenic differentiation of bone marrow-derived stem cells? Similarly, how do various modes of muscle contraction, such as concentric vs eccentric contraction, alter myokine expression or secretion? Finally, how do unloading, microgravity or prolonged bedrest impact the muscle secretome?

These questions all point to the larger issue of how muscle activity and metabolism impact the overall systemic environment to which cells of bone and other tissues are exposed. For example, it is clear that exposing muscle of aged animals to circulating factors from younger animals improves the regenerative capacity of muscle. Alternatively, how do changes in muscle, be they catabolic or anabolic, alter the systemic milieu to affect other organs such as bone? Preserving muscle mass decreases mortality and improves survival in tumor-bearing mice, and loss of lean mass is an important predictor of health outcomes following burns and with chronic wounds. The positive effects of muscle mass on health outcomes are related at least to the fact that muscle is the primary source of free amino acids in the body; however, it is also likely that certain myokines may also have a role. For example, treatment of a tumor cell-line with mouse serum following exercise, or with conditioned medium from myotubes following electrical stimulation, reduces tumor cell proliferation and increases apoptosis. These effects of muscle-derived serum and media on tumor cells were attributed to the fact that myotubes secrete the anti-tumor protein oncostatin M. This finding has significant implications for cancer and bone (for example, osteosarcoma), and future studies might be directed at elucidating the interface between myokines, metastasis of cancer to bone and tumor growth in bone tissue.

Another outstanding question concerns the interactions of muscle-derived factors with organs other than bone. Skeletal muscle hypertrophy and hyperplasia increase serum IGF-1 levels in mice lacking myostatin. The increase in circulating IGF-1 levels is associated with elevated liver-derived IGF-1 in these mice with no change in muscle-specific IGF-1 expression. Thus, heritable variation in muscle mass has the potential to dramatically alter growth factor production by other organs, which is likely to have broad systemic effects on tissues such as bone and cartilage. Exercise has recently been shown to increase secretion of follistatin by hepatocytes, and follistatin is a potent inhibitor of both activin and myostatin, factors that suppress muscle hypertrophy and impair muscle regeneration. Follistatin also increases osteoblast mineralization in vitro, raising the possibility that interactions among myokines and hepatokines may influence bone metabolism both directly, through their effects on bone cells and indirectly, by modulating growth factor and cytokine production in other organs. A similar relationship may exist between muscle and fat, where myokines can induce lipolysis, that would presumably affect circulating levels of adipokines such as leptin, which can in turn significantly alter bone formation and resorption.

In conclusion, in vivo and in vitro studies now demonstrate that muscle can function as an endocrine and paracrine organ. The factors secreted by muscle may vary according to muscle activity, such as concentric and eccentric contraction, disuse or damage in the form of traumatic injury. Factors actively secreted by muscle range from growth factors to inflammatory cytokines, and these peptides have the potential to alter bone metabolism in a variety of ways. Additional research is needed to better define the molecular and cellular pathways through which muscle-derived factors affect different types of bone cells, and anabolic and catabolic processes, in bone tissue. Nevertheless, the studies reviewed here further underscore the complex nature of muscle–bone interactions, and highlight the importance of integrating muscle biology and physiology into our understanding of bone growth, development and aging.

Conflict of Interest

The author declares no conflict of interest.

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