

ORIGINAL ARTICLE

Sarcolemmal targeting of nNOS μ improves contractile function of *mdx* muscle

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Abstract

Nitric oxide (NO) is a key regulator of skeletal muscle function and metabolism, including vasoregulation, mitochondrial function, glucose uptake, fatigue and excitation–contraction coupling. The main generator of NO in skeletal muscle is the muscle-specific form of neuronal nitric oxide synthase (nNOS μ) produced by the NOS1 gene. Skeletal muscle nNOS μ is predominantly localized at the sarcolemma by interaction with the dystrophin protein complex (DPC). In Duchenne muscular dystrophy (DMD), loss of dystrophin leads to the mislocalization of nNOS μ from the sarcolemma to the cytosol. This perturbation has been shown to impair contractile function and cause muscle fatigue in dystrophic (*mdx*) mice. Here, we investigated the effect of restoring sarcolemmal nNOS μ on muscle contractile function in *mdx* mice. To achieve this, we designed a modified form of nNOS μ (NOS-M) that is targeted to the sarcolemma by palmitoylation, even in the absence of the DPC. When expressed specifically in *mdx* skeletal muscle, NOS-M significantly attenuates force loss owing to damaging eccentric contractions and repetitive isometric contractions (fatigue), while also improving force recovery after fatigue. Expression of unmodified nNOS μ at similar levels does not lead to sarcolemmal association and fails to improve muscle function. Aside from the benefits of sarcolemmal-localized NO production, NOS-M also increased the surface membrane levels of utrophin and other DPC proteins, including β -dystroglycan, α -syntrophin and α -dystrobrevin in *mdx* muscle. These results suggest that the expression of NOS-M in skeletal muscle may be therapeutically beneficial in DMD and other muscle diseases characterized by the loss of nNOS μ from the sarcolemma.

Introduction

Nitric oxide is directly involved in many aspects of skeletal muscle function, including vasomodulation (1), mitochondrial biogenesis (2), glucose uptake (3,4), suppression of oxidative stress (5), activation of cyclic GMP synthesis (6) and modulation of excitation–contraction coupling via nitrosylation of the ryanodine receptor (7). As nitric oxide (NO) regulates several important functions in healthy muscle, it is not surprising that perturbations of NO production and cellular localization are associated with muscle diseases, including the muscular dystrophies.

In skeletal muscle fibers, NO is predominantly produced by two isoforms of neuronal nitric oxide synthase (nNOS) in

response to binding of Ca⁺⁺/calmodulin (CaM). These isoforms, nNOS μ and nNOS β , are both products of the NOS1 gene. nNOS μ is very similar to the brain form, nNOS α , but contains the muscle-specific μ insert of 34 residues, located between the oxygenase and reductase catalytic domains, and immediately adjacent to the CaM-binding site (8). nNOS μ is primarily associated with the sarcolemma through interaction with dystrophin and α -syntrophin (9,10). nNOS β differs from nNOS α in that it lacks the amino terminal PDZ scaffolding domain (11). In skeletal muscle, nNOS β is expressed at much lower levels than nNOS μ but is highly localized to the Golgi apparatus in close juxtaposition with soluble guanylate cyclase and protein kinase G, suggesting

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that activation of nNOS β is important for an NO-cGMP signaling pathway at the Golgi (12). Mice lacking both nNOS μ and nNOS β have smaller muscles with reduced specific force and high susceptibility to fatigue (12).

Restricted spatial NO signaling is likely determined by localization of NOS. Although a highly diffusible molecule, NO action is limited to areas near its site of synthesis by the high concentration of scavengers (e.g. myoglobin and glutathione). Although some nNOS μ is present in the cytosol, most of the enzyme is restricted to the sarcolemma in normal muscle. During muscle atrophy, nNOS μ becomes mislocalized moving from the sarcolemma to the cytosol where the NO produced can activate genes that induce muscle atrophy pathways (13). In Duchenne muscular dystrophy (DMD), a severe, progressive and fatal muscle disease caused by mutations in dystrophin, nNOS μ is also lost from the sarcolemma (14). This mislocalization results in a lack of sarcolemmal NO production that prevents the override of sympathetic vasoconstriction resulting in functional ischemia of the muscle (15). Experiments in mice show that loss of sarcolemmal nNOS μ not only leads to functional ischemia but also makes the muscle more susceptible to fatigue (16) and damage from eccentric contractions (17).

The *mdx* mouse serves as a dystrophin-deficient mouse model of DMD. It shares many of the key characteristics of DMD including utrophin upregulation, functional ischemia, increased fatigue and damage from eccentric contractions even though the continuous degeneration/regeneration cycles are not observed in adult *mdx* limb muscles. Studies published over the past decade or more have suggested that transgenic overexpression of nNOS α in *mdx* muscle improves the dystrophic pathology, even when not targeted to the sarcolemma (18,19,20). However, a major limitation of these studies was the lack of any measurements of muscle function, which is ultimately the most relevant determinant of mobility and longevity in DMD patients. Given the important regulatory role of nNOS μ on normal skeletal muscle contractility (21), we aimed to test whether sarcolemmal nNOS μ targeting was required for improved function of dystrophic muscles.

Here, we report that a novel form of nNOS μ , targeted to the sarcolemma by addition of a palmitoylation sequence, is remarkably effective at improving functional aspects of skeletal muscle including resistance to eccentric contraction-induced injury and fatigue. In contrast, expression of similar amounts of non-targeted nNOS μ , which remains in the cytosol, shows no improvement. Interestingly, we also demonstrate that targeted nNOS μ effectively upregulates sarcolemmal expression of utrophin and other members of the dystrophin protein complex (DPC). Together, our results demonstrate that targeted nNOS μ is a promising therapeutic approach for DMD and other neuromuscular diseases that exhibit reduced or absent sarcolemmal nNOS μ .

Results

Generation of transgenic mice

To compare the effects of general nNOS μ with sarcolemma-targeted nNOS μ expression, we generated two lines of transgenic mice. Both used the human skeletal actin (HSA) promoter to drive transcription (Fig. 1A). This promoter is only active in skeletal muscle and is turned on early in embryonic muscle development and maintained throughout mature skeletal muscle (22). A hemagglutinin tag (HA) was placed at the C-terminal end of nNOS μ in both lines. To one transgene we added a C-terminal palmitoylation signal sequence from the K-Ras oncogene (RAS) to target nNOS μ to the sarcolemma. Hereafter, this construct will be referred to as NOS-M, whereas NOS denotes the non-targeted,

nNOS μ -HA construct. We obtained three founder mice expressing NOS and two founders expressing NOS-M. Each founder line was bred onto the *mdx* background (C57Bl/10ScSn) for at least three generations before use in experiments with non-transgenic littermates as controls.

Both NOS and NOS-M were expressed well in skeletal muscle (Fig. 1B, C). Western blots of tissue homogenates showed high expression of the nNOS transgene products in both tibialis anterior (TA) and diaphragm muscles. The relative amounts of NOS and NOS-M were similar. Immunofluorescence studies of these two muscles show that transgenically expressed NOS is only faintly labeled at the sarcolemma whereas NOS-M shows robust sarcolemmal labeling (Fig. 1B, C).

Analysis of muscle function

TA *in situ*

Dystrophic muscles are particularly vulnerable to damage from eccentric contractions (23). We therefore tested the performance of skeletal muscle from WT, *mdx*, and both transgenic lines (bred onto the *mdx* background) during eccentric contractions. These experiments were performed *in situ* with the knee immobilized and the distal tendon of the TA muscle attached to a force transducer (24). A major advantage of this preparation is that *in vivo* vascular function is maintained, thus preserving proper oxygenation. Maintenance of *in vivo* oxygen levels is necessary when studying functions involving NO (21). Isometric force was measured during sequential contractions initiated by direct stimulation of the sciatic nerve (Fig. 2). The dramatic reduction of force in the *mdx* mice is evident as early as the third contraction. After five contractions, the force generated by the *mdx* and *mdx*-NOS muscle fell to ~40% of the initial force whereas for the *mdx*-NOS-M muscle, force only dropped to ~60% of its initial value (note WT drops to ~90% of initial force). From eccentric contractions 4–10, the force produced by *mdx*-NOS-M was significantly greater than *mdx* and *mdx*-NOS (see Fig. 2). Thus, sarcolemmal expression of nNOS provides considerable protection against force loss in dystrophic muscle, particularly during the early eccentric contractions, where most of the force loss occurs in regular *mdx* mice.

We then determined the effect of sarcolemmal nNOS μ restoration on TA skeletal muscle-specific force (force normalized to cross-sectional area). The specific force of TA muscle was significantly increased in NOS-M mice at low (30 and 50 Hz) stimulation frequencies but not at higher frequencies compared with *mdx*-NOS and normal *mdx* mice (Fig. 3A). In addition to eccentric contractions, muscle fatigue is another major cause of muscle weakness in dystrophic muscles (25). Muscle force was substantially higher (20%) for *mdx*-NOS-M muscles compared with those of *mdx*-NOS and regular *mdx* mice over 4 min of fatiguing contractions (Fig. 3B). In addition, recovery from fatigue was significantly greater for *mdx*-NOS-M muscle than the other two groups of *mdx* mice, whereas *mdx*-NOS mice had a small but significantly greater recovery than regular *mdx* mice (see Fig. 3B).

Diaphragm *ex vivo*

We also measured diaphragm function using small muscle strips *ex vivo* (26). Here, specific force at 120 Hz stimulation was not improved in *mdx* muscles by either the NOS-M or NOS transgene (Fig. 4A). Diaphragm fatigue over 1 min of repetitive stimulation was also not different between the groups (Fig. 4B). However, force recovery after fatigue was significantly greater for the *mdx*-NOS-M muscles compared with both *mdx*-NOS and regular *mdx*, whereas *mdx*-NOS was significantly lower than regular *mdx* mice (see Fig. 4B).

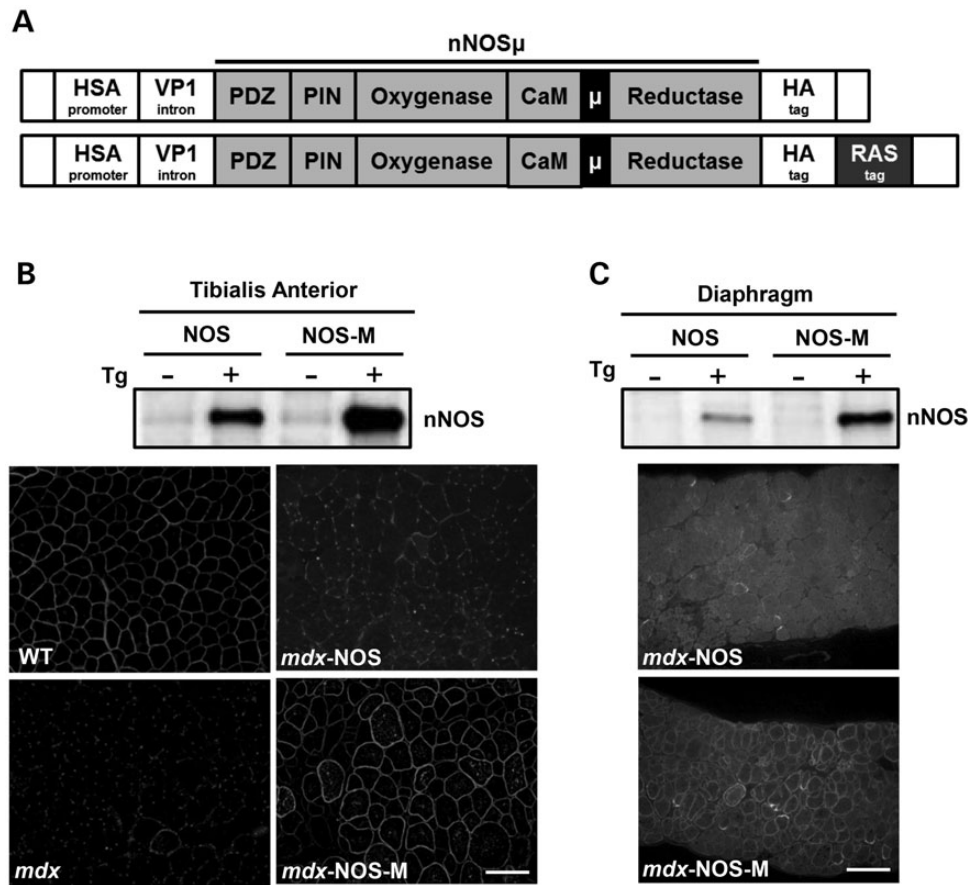


Figure 1. Design of transgenes and expression of nNOS μ in muscles. (A) Schematic representation of the two nNOS μ transgenes with the regular NOS (top) and the modified, NOS-M version (lower) showing the addition of the palmitoylation (RAS) tag (see text for details). (B) TA and (C) diaphragm representative western blots and immunostaining showing nNOS μ from NOS and NOS-M transgenic mice. Note the strong sarcolemmal staining in NOS-M but not NOS muscles. Scale bar = 100 μ m.

From these physiological studies, we conclude that nNOS μ targeted to the sarcolemma markedly improves the contractile function of *mdx* muscle during eccentric contractions and muscle fatigue. In fact, the magnitude of these improvements is similar to those attained by treatments that upregulate utrophin and current dystrophin gene therapy approaches (27,28,29). In contrast, untargeted nNOS μ shows no improvement in protection against force loss from eccentric contractions or fatigue in TA, and while these muscles show a small improvement in TA fatigue recovery compared with regular *mdx* muscle, fatigue recovery is worse than *mdx* mice in diaphragm muscles.

NOS-M targets utrophin and dystrophin-associated proteins to the sarcolemma

Transgenic expression and upregulation of utrophin improves the dystrophic pathology in the *mdx* mouse (27,30). As increased NO bioavailability upregulates utrophin levels in *mdx* muscle (31,32), we tested the hypothesis that NOS-M improves *mdx* contractile function by increasing sarcolemmal utrophin levels. We found that NOS-M markedly increases expression (Fig. 5A) and sarcolemmal localization (Fig. 5B) of utrophin in *mdx* TA muscles. In addition, several other key proteins of the DPC including β -dystroglycan, α -dystrobrevin-1 and -2 and α -syntrophin are also upregulated and localized at the sarcolemma. Therefore, our results

indicate that proper localization of nNOS μ on the sarcolemma is required for the observed increase in utrophin and proteins of the DPC. Whether these restorations are mediated by NOS-M scaffolding, by increased NO bioavailability, or both, requires further study. The pathway mediating the upregulation of utrophin and DPC proteins increase does not appear to involve increased transcription/mRNA stabilization. We found no significant difference in mRNA levels of utrophin or α -dystrobrevin when comparing muscle RNA isolated from *mdx* and *mdx-NOS-M* mice (Supplementary Material, Fig. S1).

Discussion

We have developed a novel approach to restore nNOS μ at the sarcolemmal membrane in skeletal muscle. Introduction of a K-RAS palmitoylation signal sequence at the carboxy terminus of nNOS μ causes the modified enzyme (NOS-M) to be localized on the surface membrane, as expected for proteins that are palmitoylated. The rationale for this approach is based on the hypothesis that NO bioavailability is determined in large part by the localization of its site of synthesis. In contrast to unmodified nNOS μ (NOS), NOS-M improves two important aspects of contractile function in *mdx* muscle: resistance to eccentric contraction-induced force loss and improved resistance to muscle fatigue. Eccentric damage and fatigue are important causes of muscle weakness in skeletal muscle of *mdx* mice and DMD boys.

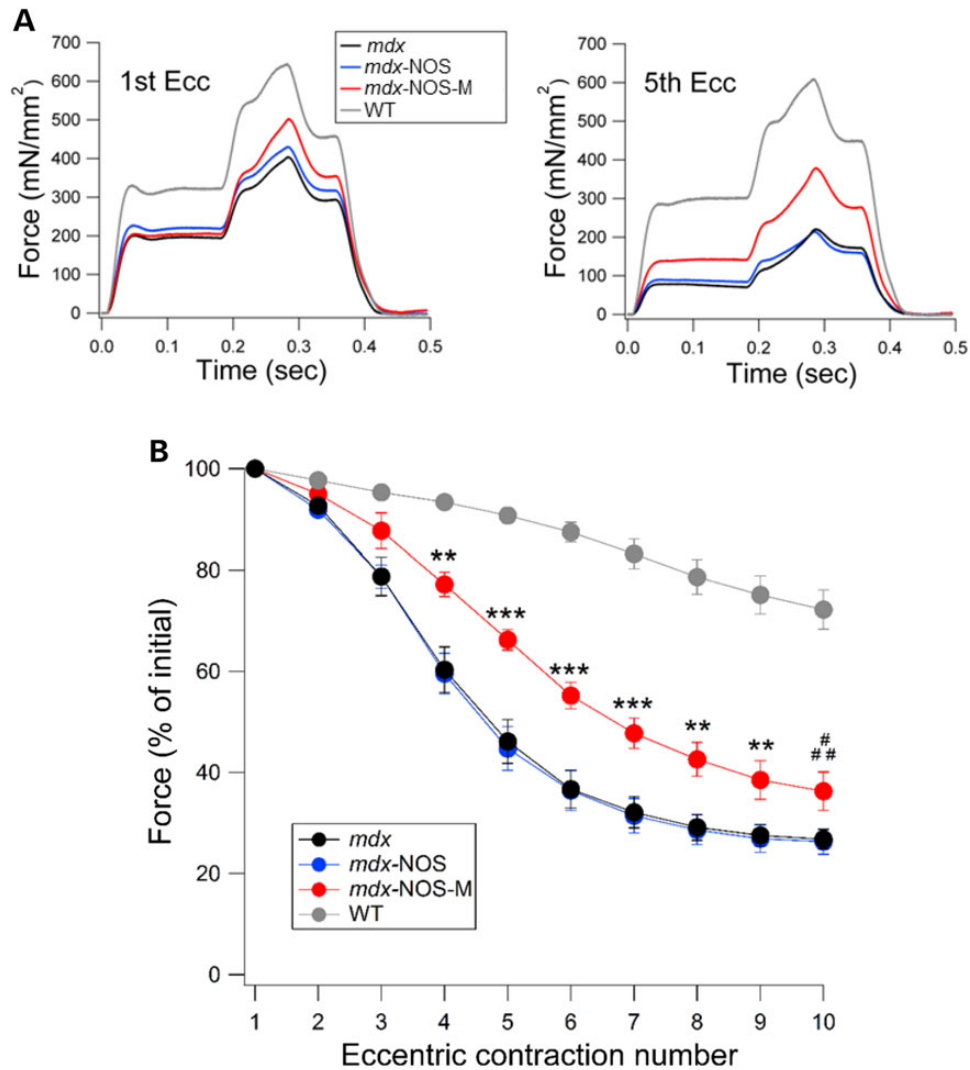


Figure 2. NOS-M reduces force loss from eccentric contractions in *mdx* TA muscle. (A) Representative traces showing force production during the first and fifth eccentric contractions for each group of mice. (B) Pooled data showing the isometric force production over 10 eccentric contractions, expressed as a percent of first contraction. ***P* < 0.01 *mdx-NOS-M* versus *mdx* and *mdx-NOS*, ****P* < 0.001 *mdx-NOS-M* versus *mdx* and *mdx-NOS*, #*P* < 0.05 *mdx-NOS-M* versus *mdx*, ##*P* < 0.01 *mdx-NOS-M* versus *mdx-NOS*.

Our results conflict with results published by Tidball and colleagues, in which they claim that improvement of pathological features of *mdx* muscle by transgenic expression of nNOS does not require sarcolemmal localization (18,19,20). Several differences in approach and analyses may explain the conflicting results. First, the Tidball group used nNOS α , the brain form of nNOS, rather than the muscle-specific form, nNOS μ . The function of the μ insert is not known but its location between the oxygenase and reductase catalytic regions and adjacent to the CaM binding site suggests an important role in regulating NO production. Second, the transgenic expression levels of nNOS α in their experiments were exceptionally high, especially in the 'supernatant fraction'. High levels of untargeted, cytosolic NOS may have deleterious effects caused, for example, by inappropriate non-physiological nitrosylation of proteins. Also, translocation of nNOS μ from the sarcolemma to the cytosol is a key factor in muscle atrophy during mechanical unloading (13). Finally, no increase in utrophin localization on the sarcolemma was observed in *mdx* muscle from transgenic muscle expressing nNOS α (20). In fact, total utrophin levels were reduced, compared with *mdx*

muscle. Although some pathological features were improved by nNOS α expression in *mdx* muscle (reduced inflammation and reduction of infiltrating macrophages) early in the disease process, no measurements of contractile function were conducted.

Upregulation of utrophin, a homolog of dystrophin, has received major attention as a possible therapeutic target, based on initial studies showing that muscle-specific expression of utrophin is highly effective in reducing the pathological deficits caused by the loss of dystrophin. One advantage of utrophin expression, compared with genetic treatments with micro-dystrophin constructs, is the greatly reduced likelihood of eliciting an immune response. A potential disadvantage is that the utrophin complex fails to restore nNOS μ to the sarcolemma (33), perhaps because of differences in the repeat 16–17 region that is essential for nNOS μ association with the DPC (22). In addition to increasing sarcolemmal utrophin levels, NOS-M has the added benefit of restoring nNOS μ to its proper location. NOS-M does not appear to affect transcription or transcript stabilization of utrophin but may increase utrophin

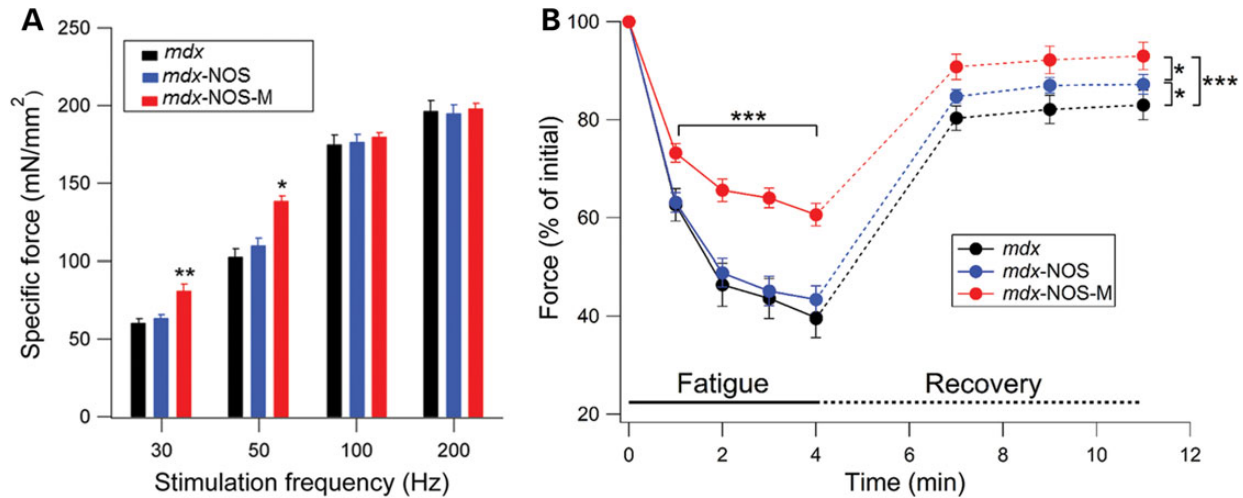


Figure 3. NOS-M enhances low-frequency force, protects against muscle fatigue and improves force recovery in *mdx* TA muscle. (A) Specific force values at various stimulation frequencies for each group. * $P < 0.05$ *mdx-NOS-M* versus *mdx* and *mdx-NOS*, ** $P < 0.01$ *mdx-NOS-M* versus *mdx* and *mdx-NOS*. (B) Mean force levels measured each minute during 4 min of fatigue (solid lines) and during fatigue recovery (dotted lines). For fatigue, *** $P < 0.001$ *mdx-NOS-M* versus *mdx* and *mdx-NOS*. For fatigue recovery, * $P < 0.05$, *** $P < 0.001$.

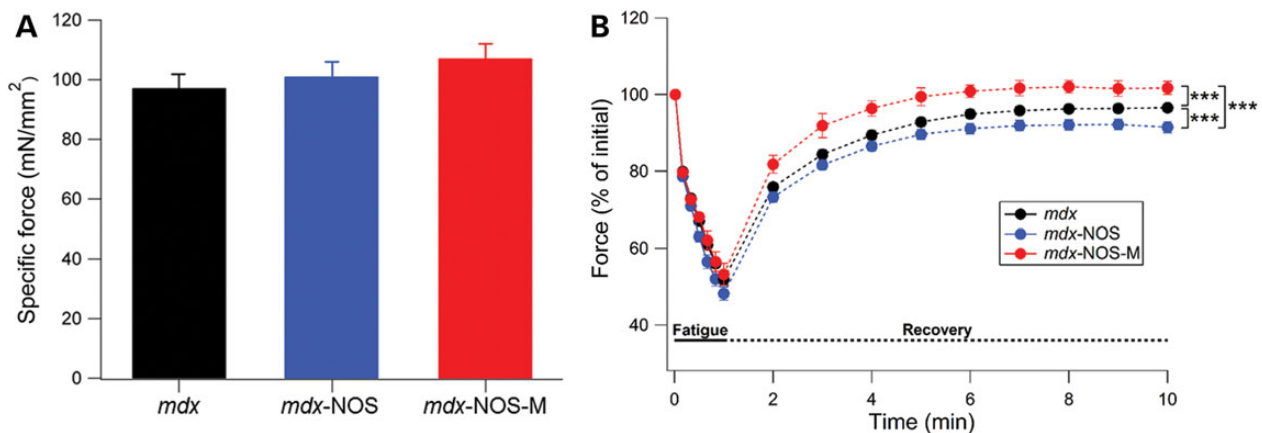


Figure 4. NOS-M improves fatigue recovery in *mdx* diaphragm muscle. (A) Specific force for diaphragm muscles during 120-Hz stimulation. There were no significant differences between any groups. (B) Mean force levels during 1 min of fatigue (solid lines) and for 10 min of fatigue recovery (dotted lines). For fatigue recovery, *** $P < 0.001$.

levels via S-nitrosylation-mediated inhibition of calpain (34), or by serving as a scaffold that stabilizes utrophin and associated proteins in a sarcolemmal complex. Regardless of the mechanism by which utrophin levels are increased, the observation that NOS-M increases levels of utrophin (and associated proteins) and partially restores normal function in dystrophic muscle makes it a potential therapeutic candidate. As NOS-M is much smaller than dystrophin/utrophin, it may serve as a more amenable protein for use in DMD-directed gene therapy.

Although our studies were focused on the *mdx* mouse model of DMD, NOS-M may also have therapeutic value in other muscle diseases in which nNOS μ expression and localization are perturbed. These include forms of limb-girdle muscular dystrophy (primarily the sarcoglycanopathies) (35), amyotrophic lateral sclerosis (36), myasthenia gravis (37), disuse atrophy (13,38), sarcopenia (39) and muscle mitochondrial diseases (40). Development of methods for viral delivery of NOS-M to skeletal muscle will be needed to test the therapeutic effectiveness in these devastating human muscle diseases.

Materials and Methods

nNOS μ cloning and transgenic mice generation

nNOS μ cDNA was derived from total skeletal muscle RNA using reverse transcription-PCR and the presence of the μ insert confirmed by sequencing. We used PCR to add a single heme-agglutinin (HA) tag epitope to the C-terminal of nNOS μ or the HA tag preceded by the palmitoylation signal sequence from the K-ras oncogene (amino acid sequence: KDGKKKKKSKTKCVIM) (41). These two nNOS μ constructs were cloned into the NotI/PacI sites of pBSX-HSAvpA vector (42,43) (a gift from Drs. Jeff Chamberlain and Stephen Hauschka, University of Washington). The pBSX-HSAvpA vector contains the HSA promoter that provides skeletal muscle-specific expression of the nNOS μ transgenes. Linearized constructs were injected into oocytes of a cross of the C57BL/6 and C3H mouse strains (Transgenic Resources Program, University of Washington). Lines of mice expressing the transgenes were bred onto WT and *mdx*, both on the C57BL/10ScSn background, for a minimum of three generations. All

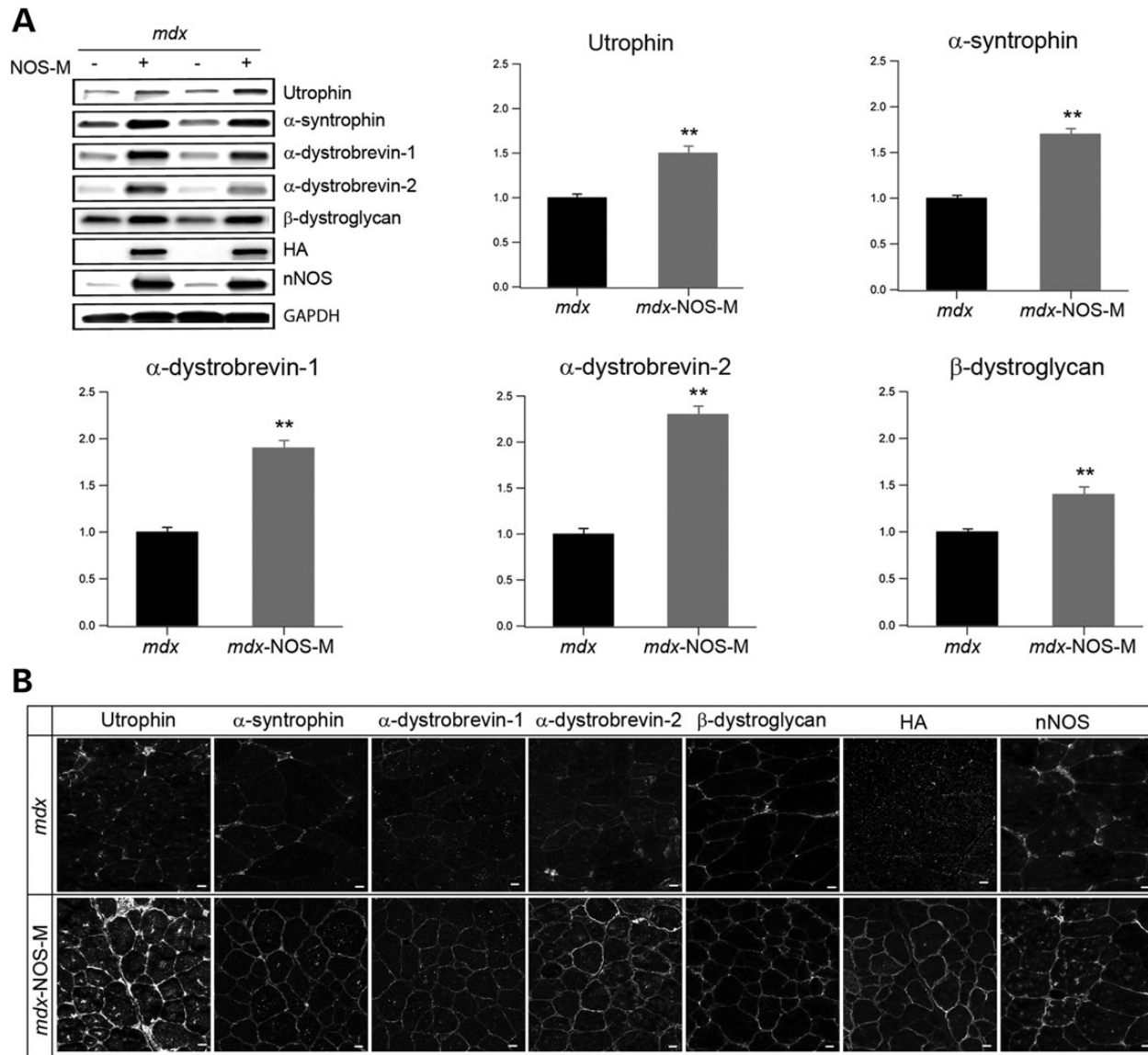


Figure 5. NOS-M increases the expression and sarcolemmal localization of utrophin and DPC proteins. (A) Representative western blots (top left) and pooled data showing the levels of utrophin and DPC proteins from TA muscles of *mdx* and *mdx-NOS-M* mice. Data were pooled from 5–10 mice/group. The western blots also show the levels of nNOS μ and the HA tag associated with the transgene. ** $P < 0.01$. (B) Representative immunostaining images showing the increased sarcolemmal localization of utrophin and DPC proteins in *mdx-NOS-M* muscle. Scale bar = 10 μ m.

experimental procedures performed on mice were approved by the Institutional Animal Care and Use Committee at the University of Washington.

In situ analysis of skeletal muscle contractile function

Anesthetized mice were positioned on a 37°C heated platform, the distal tendon of the TA muscle was surgically isolated, the knee joint was restrained with a surgical needle and the distal tendon attached to the lever arm of a servomotor (Aurora Scientific, ON, Canada). The TA muscle was activated by stimulation of the sciatic nerve using two needle electrodes. The muscle was adjusted to an optimum length (L_0) to produce the maximum tetanic force. While held at L_0 , the TA was stimulated every 2 min at increasing frequencies (10 to 200 Hz) to generate force frequency curves, to obtain the maximal tetanic force (P_0). Then, both L_0 and TA mass were recorded and used to normalize to the physiological cross-

sectional area ($[L_0 \times \text{density}]/\text{mass}$) and calculate specific tetanic force. For each mouse, the left TA was used for testing resistance to exercise-induced fatigue, whereas the right was used to test susceptibility to contraction-induced injury. To test resistance to exercise-induced fatigue, TA muscles were subjected to a series of repeated isometric stimulations (200 Hz) at 2-s intervals for 4 min. Recovery from fatigue was assessed by recording force output at 3, 5 and 7 min after the completion of the fatigue period.

Resistance to eccentric contraction (ECC)-induced injury

TA muscles were subjected to a series of 10 lengthening (eccentric) contractions. Muscles were maximally stimulated (200 Hz) for 175 ms at L_0 to achieve maximal isometric tension, and then stretched while contracting by 20% of L_0 at a velocity of two fiber lengths per second. The total stimulation time for each contraction was 350 ms. Each eccentric contraction was

separated by 1 min. The maximum isometric force measured before the stretch was recorded for each contraction and expressed as a percent of the first contraction.

Ex vivo analysis of diaphragm muscle function

Diaphragm muscle strips, 2- to 4-mm wide were used for *ex vivo* function (26). Briefly, strips were dissected in physiological buffer, bubbled with 95% O₂ and 5% CO₂ (pH 7.4). The central tendon of the strip was attached to the lever of a servomotor (Aurora Scientific), and the rib bone was attached to a stainless steel hook. Muscle stimulation was provided by two platinum electrodes, attached to the inside walls of the chamber, which are connected to a stimulator (Aurora Scientific). A length-force curve was measured using 120 Hz stimulation to establish the optimum length (L_0), and specific force was later determined by dividing force by cross-sectional area. The muscle strip was then subjected to a fatigue protocol. Here, the muscle was stimulated at 120 Hz, every second for 60 s. Recovery of force following fatigue was measured at 1-min intervals up to 10 min.

Antibodies

Our antibodies to utrophin, α -syntrophin and α -dystrobrevin have been described previously (44). We used the following commercial antibodies. Rabbit polyclonal antibodies: glyceraldehyde-3-phosphate dehydrogenase (Millipore), nNOS (Invitrogen) and HA (Sigma). Mouse monoclonal antibodies: pan dystrobrevin (BD Biosciences) and β -dystroglycan (Novacastra). Goat polyclonal anti HA (Abcam).

Western blotting

Muscle extracts in 1% Triton X-100 were prepared as described previously (44). Protein concentrations were determined using the bicinchoninic acid reagent (Pierce), and equal amounts of muscle protein were used for SDS-PAGE. Proteins were transferred to a PVDF membrane (Millipore) and the primary antibodies detected using HRP-conjugated secondary antibodies (Jackson ImmunoResearch) and ECL substrate (Pierce) with a Protein Simple Fluorochem M imager. Band intensity was determined using Image J software and plotted with the average *mdx* value set to 1. *N*-values (where *N* refers to an individual mouse) were as follows: for utrophin-*mdx* *N* = 9, *mdx*-NOS-M *N* = 10, for syntrophin dystrobrevin and dystroglycan *mdx* *N* = 5, *mdx*-NOS-M *N* = 9.

Immunohistochemistry

Muscles from 8- to 9-week-old mice were dissected, embedded in Tissue Tek OCT, frozen in liquid nitrogen-cooled isopentane. Cryostat sections (10 μ m thick) were incubated with primary antibody at 4°C overnight then with Alexa Fluor-conjugated secondary antibodies (Invitrogen) for 1 h at room temperature. Images were obtained using a Zeiss 510 Meta confocal microscope at the W.M. Keck Center for Advanced Studies in Neural Signaling at the University of Washington.

Data analysis

In all experiments, transgenic mice were compared with non-transgenic littermates. As different lines were obtained, and results from different lines for the same transgene performed very similarly, we used pooled data. Thus, C57Bl10 and *mdx* non-transgenic data correspond to negative littermates of all

transgenic lines. Data from cytosolic NOS transgenic mice correspond to the average of three different lines. All values are expressed as the mean \pm SEM, and statistical significance was set at $P < 0.05$. For two experimental groups, a Student's *t*-test was used for statistical analysis. For more than two groups, two-way ANOVA was used. Where an ANOVA was significant, an LSD *post hoc* test was used to evaluate differences between independent variables (Data Desk, Ithaca, New York).

Supplementary Material

Supplementary Material is available at HMG online.

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Conflict of Interest statement: None declared.

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