

**Fig. 4.** Intensity of scattered light (488 nm) as a function of scattering angle for dispersions of the first four clusters. Solid lines are calculations of the theoretical scattering from randomly oriented idealized packings of 844-nm spheres (31). The data and curves are offset for clarity. Curve 1, single spheres; 2, sphere doublets; 3, triangles; 4, tetrahedra. The agreement between theory and experiment shows that the clusters are well separated and stable.

ade (11). The varied and unusual polyhedra of this sequence illustrate how certain symmetries, including fivefold rotational symmetry, can arise solely from compression and packing constraints and need not require interparticle attractions. It may well be possible to define other types of constraints in microsphere systems and in so doing to discover new sphere-packing motifs.

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#### Supporting Online Material

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Materials and Methods  
References

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## Cell Therapy of $\alpha$ -Sarcoglycan Null Dystrophic Mice Through Intra-Arterial Delivery of Mesoangioblasts

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Preclinical or clinical trials for muscular dystrophies have met with modest success, mainly because of inefficient delivery of viral vectors or donor cells to dystrophic muscles. We report here that intra-arterial delivery of wild-type mesoangioblasts, a class of vessel-associated stem cells, corrects morphologically and functionally the dystrophic phenotype of virtually all downstream muscles in adult immunocompetent  $\alpha$ -sarcoglycan ( $\alpha$ -SG) null mice, a model organism for limb-girdle muscular dystrophy. When mesoangioblasts isolated from juvenile dystrophic mice and transduced with a lentiviral vector expressing  $\alpha$ -SG were injected into the femoral artery of dystrophic mice, they reconstituted skeletal muscle in a manner similar to that seen in wild-type cells. The success of this protocol was mainly due to widespread distribution of donor stem cells through the capillary network, a distinct advantage of this strategy over previous approaches.

There are currently three main experimental approaches to therapy of muscular dystrophies (1). Gene therapy focuses on the development of new vectors capable of delivering

efficiently the missing gene to the postmitotic nuclei of the muscle fibers in vivo. The pharmacological approach aims to restore the protein complex that is altered in many forms of muscular dystrophy through different strategies ranging from skipping mutated exons to increasing the synthesis of cognate proteins such as utrophin. The cell therapy approach aims to functionally rescue the tissue by delivery of cells (2); these may be satellite cells (which regenerate new fibers after damage to the muscle) or pluripotent stem cells (which have been shown to differentiate into skeletal muscle both in vitro and in vivo). However, the limited self-renewal and migratory capacity of dystrophic satellite cells and our modest knowledge of stem cell biology have so far hampered the success of the cell therapy approach.

We recently identified a class of vessel-associated fetal stem cells that can differ-

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entiate into most mesoderm (but not other germ layer) cell types when exposed to certain cytokines or to differentiating cells of a mesodermal tissue. We termed these cells "mesoangioblasts" (3). Mesoangioblasts can be grown extensively in culture (more than 50 passages), yet they are growth inhibited at confluence, do not grow in soft agar, and are not tumorigenic in nude mice assays (4). Different clonal lines of mesoangioblasts show similar, although nonidentical, growth and differentiation potential: When subjected to microarray analysis, they show profiles of gene expression that not only are similar among themselves, but also resemble those recently reported for hematopoietic, neural, and embryonic stem cells (5, 6). In keeping with their differentiation potential, mesoangioblasts predominantly express genes enriched in mesoderm, some of which are receptors and signaling molecules for mesoderm, inducing molecules such as bone morphogenetic protein (BMP), Wnt, and Notch (7).

When injected into the blood circulation, mesoangioblasts accumulate in the first capillary filter they encounter and are able to migrate outside the vessel, but only in the presence of inflammation, as in the case of dystrophic muscle. Indeed, mesoangioblasts express many receptors for inflammatory cytokines and are able to migrate in vitro and in vivo in response to HMGB-1 (8), a nuclear protein that is released by necrotic cells and acts as a potent inflammatory cytokine (9). We thus reasoned that if these cells were injected into an artery, they would accumulate into the capillary filter and from there into the interstitial tissue of downstream muscles.

To test this possibility, we injected  $5 \times 10^5$  [ $^{14}\text{C}$ ]thymidine-labeled (10) wild-type mesoangioblasts into the right femoral artery (11) of 1-month-old  $\alpha$ -SG null mice ( $n = 2$ ) (12). After 24 hours,  $30 \pm 7\%$  of the injected cells were detected in the muscles downstream of the injected artery, whereas  $<3\%$  of donor cells were detected in the same muscles when injection had occurred through the tail vein ( $n = 2$ ) or intramuscularly ( $n = 2$ ). Tissue distribution of donor cells was analyzed 24 hours after injection through the femoral artery ( $n = 4$ ); 1,1'-dioctadecyl-3,3,3',3'-tetramethylindocarbocyanine perchlorate (DiI)-labeled mesoangioblasts were detected outside the vessel wall and within the extracellular matrix of all downstream skeletal muscles, especially in areas where degeneration and regeneration were occurring (fig. S1A). Analysis of serial sections throughout the muscle length (fig. S1C) revealed that DiI-labeled cells accumulated mainly in the quadriceps ( $3.1 \pm 0.4\%$  of

total nuclei,  $n = 2$ ), gastrocnemius ( $2.3 \pm 0.3\%$ ,  $n = 2$ ), and soleus ( $1.2 \pm 0.2\%$ ,  $n = 2$ ) and to a lesser extent in other muscles such as the tibialis anterior ( $0.8 \pm 0.2\%$ ,  $n = 2$ ) and extensor digitorum longus (EDL) ( $0.3 \pm 0.1\%$ ,  $n = 2$ ). In contrast, satellite cells from transgenic mice ubiquitously expressing green fluorescent protein (GFP) (13), delivered through the femoral artery ( $n = 4$ ), could not cross the endothelial layer and remained adherent to the vessel wall (fig. S1B). When male mesoangioblasts, previously labeled with a lentiviral vector expressing nuclear LacZ (14), were injected into the femoral artery of female  $\alpha$ -SG null mice ( $n = 3$ ), polymerase chain reaction analysis of two reporter genes (*Sry* and *LacZ*) confirmed the presence of the host cells in all the hindlimb skeletal muscles analyzed 2 weeks after injection, but not in the diaphragm (fig. S1D). At this time, foci of mesoangioblasts were mainly located in the microcapillary network, as revealed by fluorescence in situ hybridization (FISH) analysis of the Y chromosome (15) of male wild-type mesoangioblasts (fig. S1E, arrows) injected into  $\alpha$ -SG null female mice. Several regenerating, centrally nucleated fibers showed the presence of Y chromosome-positive nuclei underneath the fiber basal lamina (double arrows in inset, fig. S1E); both FISH and immunostaining with antibodies to  $\beta$ -galactosidase identified positive nuclei in the center of regenerating fibers (arrowheads, fig. S1, E and F). Several DiI-labeled mesoangioblasts were detected at the periphery of fibers, where they expressed M-cadherin (fig. S1G) and c-Met (16), typical markers of satellite cells (17). Similarly labeled mesoangioblasts were also detected in several vessels near areas of regeneration, where they expressed  $\alpha$ -smooth muscle actin (arrowhead, fig. S1H) or platelet endothelial cell adhesion molecule (arrowheads, fig. S1, I and J); other labeled mesoangioblasts that did not express these markers were also present in the sections (arrows, fig. S1, I and J). These results show that mesoangioblasts can diffuse from the arterial tree into skeletal muscle, where they are incorporated into regenerating fibers or express markers of smooth muscle, satellite cells, or endothelium in vivo.

**Restoration of  $\alpha$ -sarcoglycan and dystrophin-glycoprotein complex by wild-type mesoangioblasts.** We next investigated whether injection of wild-type mesoangioblasts in the muscles of  $\alpha$ -SG null mice may restore expression of  $\alpha$ -SG and of the whole dystrophin complex (18). Two months after a single injection of  $5 \times 10^5$  wild-type male mesoangioblasts ( $n = 4$ ) into female  $\alpha$ -SG null mice, many areas of the quadriceps, tibialis anterior, soleus, gastrocnemius, and EDL ex-

pressed  $\alpha$ -SG, whereas the protein was completely absent from the muscles of the untreated null mouse (an example for the quadriceps is shown in Fig. 1A). FISH analysis revealed male nuclei adjacent to  $\alpha$ -SG-positive membrane (Fig. 1B) and longitudinal sections, which allowed us to estimate that  $\alpha$ -SG expression extended for at least  $50 \mu\text{m}$  away from the last Y chromosome-positive nucleus detected, consistent with a certain mobility of the protein within the plasma membrane and outside of the transcriptional domain of a specific nuclear cluster. Transverse sections of control,  $\alpha$ -SG null, and treated  $\alpha$ -SG null muscle at higher magnification confirmed the reliability of the method used (Fig. 1B).

All the muscle fibers that expressed  $\alpha$ -SG after mesoangioblast injection also expressed  $\beta$ -,  $\gamma$ -, and  $\delta$ -sarcoglycans and dystrophin (an example is shown for the soleus in Fig. 1C). Western blot analysis performed on similarly treated mice ( $n = 3$ ) confirmed the presence of the  $\alpha$ -SG protein in skeletal muscle of  $\alpha$ -SG null mice treated with wild-type mesoangioblasts. Protein expression in membrane fractions of gastrocnemius or quadriceps from treated mice was comparable to that in wild-type muscles (Fig. 1D). A similar pattern was maintained up to at least 3 months after a single mesoangioblast injection. Comparable levels of integrin  $\beta 1$  subunit and myosin heavy chain (MyHC) were detected in all the different samples analyzed (Fig. 1D). Taken together, these results show that intra-arterial mesoangioblast delivery was effective in restoring expression of  $\alpha$ -SG protein and of the other members of the dystrophin-glycoprotein complex in treated  $\alpha$ -SG null mice. No immune reaction occurred against reconstituted fibers, even though low-titer serum antibodies to  $\alpha$ -SG were detected in treated mice (fig. S2A). On the basis of our previous data, we predicted that very few mesoangioblasts might have reached other muscles (such as those of the noninjected leg or the body wall), and consequently few if any  $\alpha$ -SG-positive fibers should be present in these muscles. Immunofluorescence analysis of the corresponding muscles of the contralateral leg and of the diaphragm in the same injected mice revealed rare  $\alpha$ -SG-positive fibers (16). Several sparse Y chromosome-positive undifferentiated donor cells persisted in the liver and lung 2 months after intrafemoral injection (16).

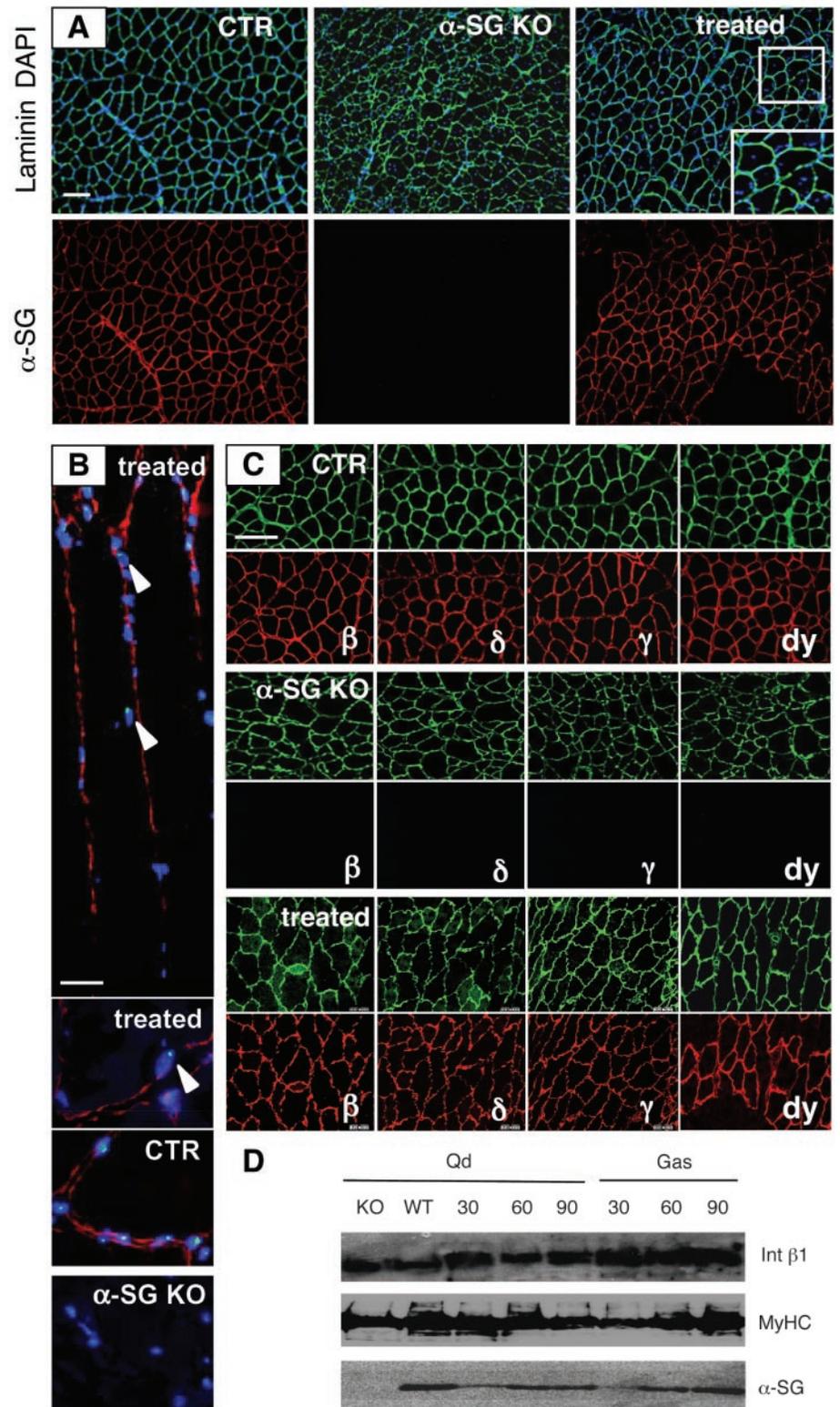
**Long-term effect of donor wild-type mesoangioblasts.** We then injected  $5 \times 10^5$  male wild-type mesoangioblasts three times (at 40-day intervals) into the femoral artery of 2-month-old  $\alpha$ -SG null female mice ( $n = 6$ ). Animals were analyzed 4 months after the first injection (at 6 months of age). Histological analysis of skeletal muscle tissue

of mice treated by three injections showed an increased number of apparently normal fibers and reduction of the necrotic areas and of cellular infiltrates (Fig. 2A). Consistent with the histology, immunofluorescence analysis revealed the widespread presence of  $\alpha$ -SG

throughout the whole soleus muscle (more than 50% of the fibers), in contrast with the total absence of signal in untreated  $\alpha$ -SG null mice (fig. S3A). We found markedly decreased uptake of Evans Blue dye in skeletal muscle of  $\alpha$ -SG null mice treated with me-

soangioblasts relative to untreated mice (Fig. 2B), demonstrating a preserved integrity of the sarcolemma (18). Finally, staining with Azan Mallory revealed a marked reduction of the fibrosis (stained in blue by the dye) in the muscles of treated mice (Fig. 2C). Even after

**Fig. 1.** Expression of  $\alpha$ -SG and other dystrophin-associated proteins in  $\alpha$ -SG null mice after intra-arterial delivery of wild-type mesoangioblasts. **(A)** Low magnification of quadriceps from control (CTR) mice (left),  $\alpha$ -SG null ( $\alpha$ -SG KO) mice (center), and treated  $\alpha$ -SG null mice (injected with wild-type mesoangioblasts 2 months before killing) (right). Large areas of the treated muscle expressed  $\alpha$ -SG after staining with a specific antibody (red). Sections were also stained with antibodies to laminin (green) and with 4',6'-diamidino-2-phenylindole (DAPI) (blue). Inset: Higher magnification of treated muscle with centrally located nuclei. **(B)** Longitudinal (upper) and transverse (lower) sections of the tibialis anterior of treated mice stained with antibodies to sarcoglycan (red) and hybridized with a probe recognizing the Y chromosome (green). Sections were also stained with DAPI. The merged images reveal donor Y chromosome-positive nuclei (arrowheads) adjacent to  $\alpha$ -SG-positive membranes. Sections from wild-type male (CTR) and  $\alpha$ -SG null female ( $\alpha$ -SG KO) mice are also shown as controls. **(C)** High magnification of soleus from the same treated mice reveals extensive reconstitution of the dystrophin complex after double staining with antibodies to laminin (green, upper panels) or to sarcoglycans and dystrophin (dy) (red, lower panels). **(D)** Western blot analysis of proteins isolated from a postnuclear membrane fraction of two large muscles (quadriceps and gastrocnemius) from  $\alpha$ -SG null mice 30, 60, and 90 days after injection of wild-type mesoangioblasts. The filters were reacted with antibodies to  $\alpha$ -SG (12); MyHC (3) and  $\beta$ 1 integrin are shown as internal controls. Scale bar, 100  $\mu$ m (A and C), 20  $\mu$ m (B).



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three consecutive injections, very few  $\alpha$ -SG-positive fibers were present in the contralateral soleus (fig. S3C). Consistent with the immunofluorescence analysis, Western blot revealed significant accumulation of  $\alpha$ -SG protein at high level in the treated right quadriceps but undetectable levels of the protein in the contralateral untreated left quadriceps (fig. S3B).

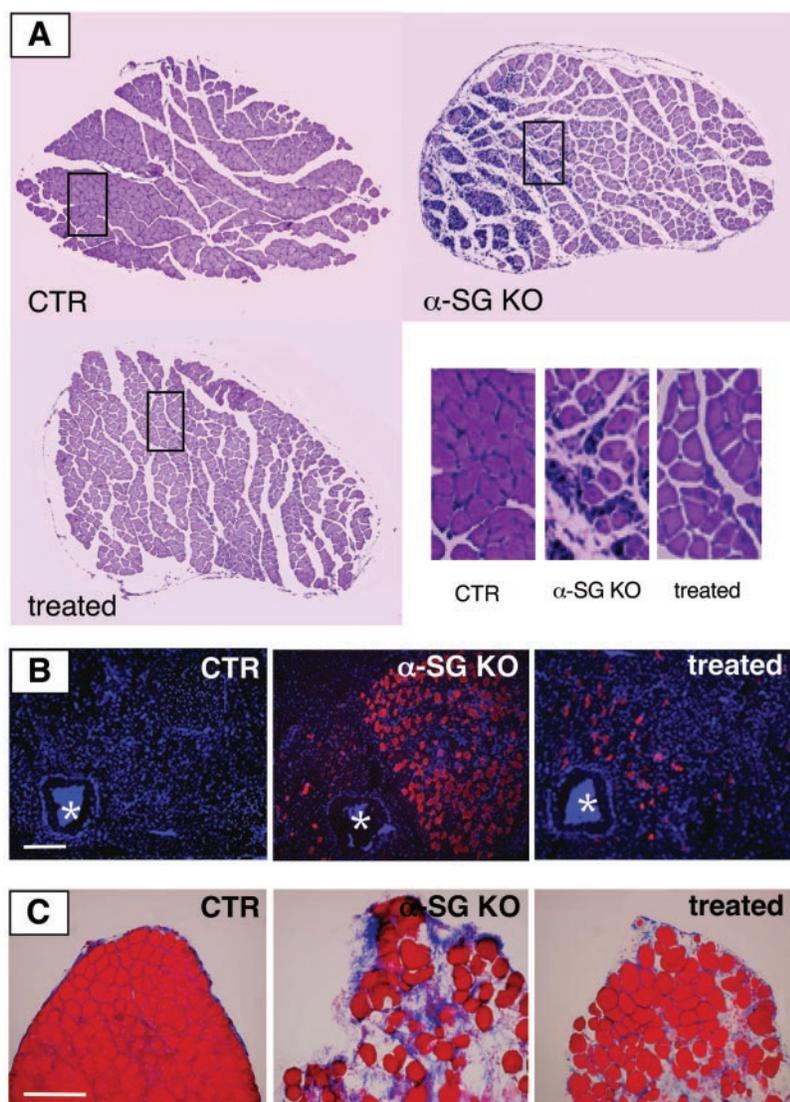
**Complete functional recovery of treated muscle.** The physiology of hindlimb muscles was studied in 6-month-old  $\alpha$ -SG null mice treated with three injections of wild-type mesoangioblasts, and the results obtained indicate that the amelioration of skeletal muscle morphology sustains a recovery of function. Previous studies have suggested

that functional analysis of whole skeletal muscles *in vitro* is not a sensitive approach to showing impairment of skeletal muscle function in  $\alpha$ -SG null mice. In fact, notwithstanding a clear impairment in animal motility, tetanic force of whole muscles of  $\alpha$ -SG null mice was not significantly decreased (12).

To better assess skeletal muscle function, we dissected a large population ( $n = 277$ ) of individual muscle fibers from gastrocnemius muscles of control ( $n = 3$ ),  $\alpha$ -SG null ( $n = 3$ ), and mesoangioblast-treated  $\alpha$ -SG null ( $n = 3$ ) mice, and measured their cross-sectional area (CSA), specific force (Po/CSA), and maximum shortening velocity ( $V_o$ ) (Fig. 3, A and B). All fibers used were identified on the basis of

MyHC isoform composition (19, 20). Because the large majority of fibers contained MyHC-2B (~80%), only type 2B fibers were used for comparison. The analysis of the distribution of CSAs in the three populations of fibers revealed an increase in fiber size in  $\alpha$ -SG null mice relative to control mice, as well as a partial recovery of normal size in treated  $\alpha$ -SG null mice (Fig. 3B). The median of CSAs was higher for  $\alpha$ -SG null mice ( $5066 \mu\text{m}^2$ ) than for controls ( $3472 \mu\text{m}^2$ ) and was intermediate for treated  $\alpha$ -SG null mice ( $3922 \mu\text{m}^2$ ); a similar trend was also clear for the 75th percentile ( $8074 \mu\text{m}^2$  for  $\alpha$ -SG null mice,  $4983 \mu\text{m}^2$  for control mice, and  $5654 \mu\text{m}^2$  for treated  $\alpha$ -SG null mice). Finally, both the range of CSAs between the 25th and 75th percentile and the smallest and largest CSA values strongly suggest that the range of variability of CSAs was larger for  $\alpha$ -SG null mice than for controls and was intermediate for treated  $\alpha$ -SG null mice. Po/CSA was significantly lower ( $P < 0.001$ ) in single muscle fibers from  $\alpha$ -SG null mice ( $42.01 \pm 21.63 \text{ kN/m}^2$ ,  $n = 106$ ) than in single muscle fibers from controls ( $61.76 \pm 24.80 \text{ kN/m}^2$ ,  $n = 82$ ). Po/CSA significantly ( $P < 0.001$ ) recovered to normal values in treated  $\alpha$ -SG null mice ( $65.48 \pm 27.02 \text{ kN/m}^2$ ,  $n = 89$ ). (For more details of the mean Po/CSA values of the fibers from each animal, see table S2.) No difference was found in  $V_o$  among the three fiber groups. Because all fibers were type 2B fibers, the latter result is fully consistent with the well-known dependence of  $V_o$  on MyHC isoform composition (21). The lack of difference for  $V_o$  suggests that neither muscular dystrophy nor mesoangioblast treatment affected the kinetics of actomyosin interaction (22). Morphometric analysis of centrally nucleated (i.e., regenerating) fibers indicated that treated mice had a size distribution different from that of untreated dystrophic mice (Fig. 3C). Large, degenerating fibers were absent and a single-mode distribution comparable with normal fibers was observed, which suggests that when regenerating fibers incorporate mesoangioblasts, they are less susceptible to further degeneration.

We also counted the total number of fibers in soleus muscles from control ( $n = 4$ ),  $\alpha$ -SG null ( $n = 4$ ), and treated  $\alpha$ -SG null mice ( $n = 4$ ). Cross cryosections of soleus muscles from the right leg (the treated leg in the treated  $\alpha$ -SG null mice) were stained with hematoxylin and eosin (H&E). Care was taken to cut sections that extended to the edge of muscles. Two sections from the portion of the muscle with the largest CSA were chosen for analysis. A significant decrease ( $-43.42 \pm 4.98\%$ ) of total fiber number ( $P < 0.0004$ ) was observed in  $\alpha$ -SG null mice ( $n = 460.05 \pm 40.57$ ) relative to control mice ( $n =$



**Fig. 2.** Morphology of long-term treated  $\alpha$ -SG null dystrophic muscles after three consecutive injections of wild-type mesoangioblasts. (A) H&E staining of the soleus of a 6-month-old control wild-type (CTR) and similarly aged  $\alpha$ -SG null mice ( $\alpha$ -SG KO) and treated  $\alpha$ -SG null mice. Higher magnification of the sections is also shown. The untreated dystrophic muscle showed a large area of necrosis and disrupted morphology that was significantly reduced after treatment. (B) Evans Blue dye injection into the tail vein. Red identifies damaged fibers that have taken up the dye; nuclei are revealed in blue by DAPI staining; tibial bone is marked with a white star. (C) Azan-Mallory staining reveals accumulation of extracellular scar tissue (blue). Scale bars, 100  $\mu\text{m}$ .

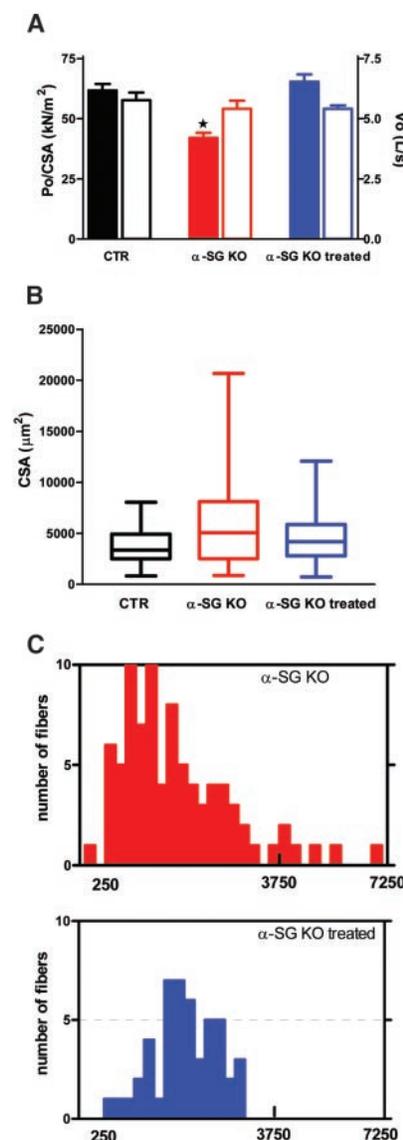
813.75 ± 92.84). A significant ( $P < 0.004$ ) recovery (+55.97 ± 23.59%) of total fiber number was observed in treated  $\alpha$ -SG null mice ( $n = 718.25 \pm 108.64$ ) relative to  $\alpha$ -SG null mice, suggesting a more effective regeneration of muscle fibers. At 6 months of age, dystrophic mice show reduced motility in the cage; in contrast, treated mice maintained a certain degree of spontaneous motility despite a monolateral treatment. To quantify the extent of residual motility, we tested control, dystrophic, and mesoangioblast-treated dystrophic mice ( $n = 4$  for each group) by forced run on the rotarod (23) at a fixed speed (1.6 m/min for 4 min; 2 min run + 1 min rest + 2 min run). On average, the control mice fell off the rotarod only 3.5 (±1.29) times during the 4-min test and never stopped running before the end of the test. The  $\alpha$ -SG null mice could hardly run on the rotarod [i.e., fell off 12.25 (±2.22) times per minute] and stopped running after 1 min. The treated  $\alpha$ -SG null mice fell off 9.75 (±1.71) times per minute and could run for 3 min before stopping. These data suggest that functional recovery of treated muscles led to a partial increase in motility, likely because only one leg had been treated.

**Morphological and functional recovery by autologous, genetically corrected mesoangioblasts.** To test whether autologous, genetically corrected stem cells may represent a possible model for the therapy of muscular dystrophy, we isolated mesoangioblasts from vessels of juvenile dystrophic mice (15 days). Mesoangioblasts are obtained at much lower frequency from adult than from embryonic vessels and grow at a lower rate (division time, 24 versus 12 hours), but they are equivalent to their embryonic counterparts in terms of gene expression and differentiation potency (4). We produced a third-generation lentiviral vector (14) expressing the mouse  $\alpha$ -SG cDNA and GFP cDNA as a reporter gene (hPGK-GFP- $\alpha$ -SG vector). After 48 hours of virus assembly in 293T packaging cells, we collected the supernatant containing the viral particles and used it to infect mesoangioblasts at a multiplicity of infection (MOI) of 200 for 24 hours. Under these conditions, more than 90% of the cell population was efficiently transduced. Furthermore, 293T cells as well as transduced  $\alpha$ -SG null mesoangioblasts efficiently produced the recombinant protein (Fig. 4A) at a level comparable to normal skeletal muscle (24), a key requirement for preventing  $\alpha$ -SG-dependent cytotoxicity. When  $\alpha$ -SG null mesoangioblasts were transduced with this vector, more than 90% of the population expressed GFP without changing their growth rate or acquiring features of transformed cells (16). Transduced  $\alpha$ -SG null mesoangioblasts efficiently differentiated into myotubes coex-

pressing GFP and MyHC when cocultured with uninfected C2C12 myoblasts (Fig. 4B). Four months after three intra-arterial injections of  $5 \times 10^5$  transduced  $\alpha$ -SG dystrophic mesoangioblasts, many fibers expressed a strong cytosolic diffuse GFP signal and  $\alpha$ -SG on the membrane (Fig. 4C). In contrast, very few GFP- $\alpha$ -SG-positive fibers were detected in the contralateral muscles of the noninjected leg (Fig. 4D). Western blot analysis (Fig. 4E) revealed accumulation of the  $\alpha$ -SG protein in the membranes of the  $\alpha$ -SG null gastrocnemius muscle treated with genetically corrected mesoangioblasts. After long-term treatment with genetically corrected mesoangioblasts,  $\alpha$ -SG null mice had restored specific force in individual fibers of their gastrocnemius muscle (Fig. 4F) and also showed an ameliorated motility on the rotarod test (16), similar to the effect observed with dystrophic mice treated with wild-type mesoangioblasts. No immune reaction was de-

tected against the GFP protein, although low-titer antibodies were present in the serum (fig. S2B). Indeed, immunostaining with antibodies to MAC-1, which identifies infiltrating macrophages, revealed reduced numbers in the muscles of treated  $\alpha$ -SG null mice (fig. S2C).

**Future prospects.** These data indicate that mesoangioblasts represent a promising approach to cell therapy of primary myopathies. In comparison with skeletal myoblasts, mesoangioblasts show the ability to cross the endothelium and migrate extensively in the tissue interstitium, where they are recruited by regenerating muscle fibers, thus reconstituting the dystrophin-glycoprotein complex. Although this is also the fate of blood-borne progenitors from the bone marrow (25–29), the frequency of this event is too low to result in noticeable amelioration of the dystrophic phenotype (26, 29). In contrast, mesoangioblasts can be expanded in vitro and directly delivered in large numbers through the arte-



**Fig. 3.** Functional properties and cross-sectional areas of individual muscle fibers of long-term treated  $\alpha$ -SG null dystrophic muscles after three consecutive injections of wild-type mesoangioblasts. (A) Specific tension (Po/CSA) and maximum shortening velocity (Vo), and (B) distribution of cross-sectional areas (CSAs) of a population of 277 single muscle fibers isolated from gastrocnemius muscles of CTR mice ( $n = 3$ ),  $\alpha$ -SG null mice ( $n = 3$ ), and treated  $\alpha$ -SG null mice ( $n = 3$ ). Methods for determining CSA, force (Po), and Vo of isolated muscle fibers are described in detail in the online supplement (19, 20). As in all mice studied, the large majority of fibers (~80%) from gastrocnemius muscles contained MyHC-2B; only type 2B fibers were used for comparison. In (A), mean values of Po/CSA (solid bars) and Vo (open bars) are shown for type 2B single skinned muscle fibers from the three groups of mice: CTR (black bars),  $\alpha$ -SG KO (red bars), and treated  $\alpha$ -SG KO (blue bars). Po/CSA values were significantly lower in  $\alpha$ -SG KO mice than in CTR and treated  $\alpha$ -SG KO mice, as indicated by the asterisk. [Statistical significance of the differences between mean values of Po/CSA and of Vo was assessed by one-way analysis of variance (ANOVA) followed by the Student-Newman-Keuls test to determine which groups were significantly different ( $P < 0.05$ ) from the others.] Details of the Po/CSA values and of animal-to-animal variation are reported in the online supplement. In (B), the box-and-whiskers plot shows the median of CSAs or 50th percentile (indicated by the line in the middle of the box), the 75th percentile (i.e., the top of the box), the range of CSAs (i.e., the width of the box), and the smallest and largest CSAs (i.e., the top and bottom of the whiskers). (C) Distribution of CSA values of centrally nucleated (i.e., regenerating) muscle fibers from  $\alpha$ -SG KO (upper histogram, red bars) and treated  $\alpha$ -SG KO mice (lower histogram, blue bars) determined on H&E-stained cross cryosections of bundles of gastrocnemius muscles from three mice per group.

## RESEARCH ARTICLES

rial circulation, with no need for a complex procedure such as bone marrow transplantation. Intra-arterial injections are simple and safe procedures in patients where they may be repeated frequently, in contrast to the mouse, where a surgical procedure is required.

In comparison with mesoangioblasts, viral vectors do not cross the endothelium and require intramuscular delivery; they transduce muscle fibers very efficiently, but only in the injected area because of their limited diffusion (30, 31). Finally, a number of studies indicate amelioration of the dystrophic phenotype by expression of biologically active molecules such as insulin-like growth factor-I or neutralizing antibodies for myostatin (32, 33). These strategies, which do not lead to gene replacement, may be part of a future combination therapy by preserving muscle integrity and thus improving the efficacy of cell therapies.

Before clinical application of these results can be envisioned, several points remain to be resolved. Human mesoangioblasts have been isolated from fetal vessels

but not yet from the patient's own vessels (4). Our work with juvenile mice indicates that this will ultimately be possible. Recently described adult mesoderm progenitors (34) may also represent a useful source, although it has not been shown that they can cross the vessel wall. As far as gene replacement is concerned, lentiviral vectors still pose safety concerns for future clinical application, even though a clinical trial using these vectors has been approved (35); however, the use of muscle-specific promoters active only in postmitotic cells and the addition of suicide genes should improve their safety. Moreover, lentiviral vectors are limited in the size of insert they accommodate, so that for Duchenne muscular dystrophy, microdystrophin (36) or a small RNA causing exon skipping (37) would have to be used (because full-length dystrophin cDNA is too large). Instead of gene replacement, heterologous mesoangioblasts from fetal vessels could be considered under an appropriate regime of immune suppression, if studies in mice should

show that only low doses or transient treatment are required.

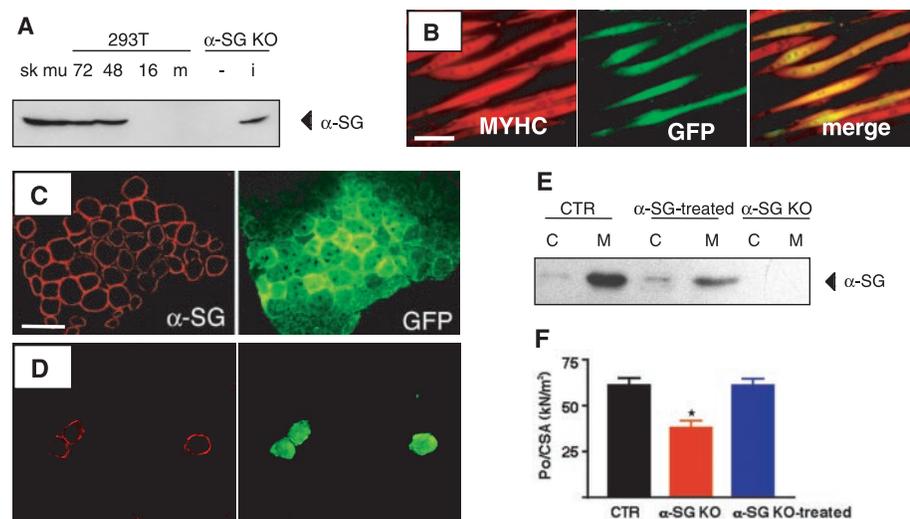
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### Supporting Online Material

www.sciencemag.org/cgi/content/full/1082254/DC1  
Materials and Methods  
Figs. S1 to S3  
Tables S1 to S3  
References

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**Fig. 4.** In vitro gene transfer into dystrophic mesoangioblasts and tissue replacement with transduced cells. (A) Western blot analysis of packaging 293T (at 72, 48, and 16 hours after transfection), mock transfected (m), and  $\alpha$ -SG null mesoangioblasts either untreated (-) or transduced (i) with a third-generation lentiviral vector expressing  $\alpha$ -SG cDNA under the transcriptional control of the PGK promoter and followed by an IRES (internal ribosomal entry site)-GFP, reacted with a monoclonal antibody to  $\alpha$ -SG (sk mu,  $\alpha$ -SG from skeletal muscle). (B) Immunofluorescence analysis of a coculture of transduced mesoangioblasts with control C2C12 myoblasts, showing a large number of MyHC/GFP-positive myotubes that incorporated GFP-positive mesoangioblasts. Scale bar, 10  $\mu$ m. (C) Immunofluorescence with antibodies to  $\alpha$ -SG (red) of the gastrocnemius muscle injected three times with  $5 \times 10^5$  dystrophic, transduced mesoangioblasts through the femoral artery revealed many  $\alpha$ -SG/GFP-positive fibers (green). Scale bar, 100  $\mu$ m. (D) Immunofluorescence with antibodies to  $\alpha$ -SG (red) of the contralateral, noninjected gastrocnemius muscle revealed very few  $\alpha$ -SG/GFP-positive fibers (green). (E) Western blot analysis of proteins isolated from postnuclear membrane (M) and cytosolic (C) fractions of the quadriceps muscles from  $\alpha$ -SG null mice injected three times with  $5 \times 10^5$  dystrophic, transduced mesoangioblasts through the femoral artery. The same fractions from control (CTR) and dystrophic ( $\alpha$ -SG KO) untreated mice are also shown. (F) Specific tension (Po/CSA) of single muscle fibers ( $n = 95$ ) of gastrocnemius muscles from CTR mice (black bar),  $\alpha$ -SG null mice (red bar), and  $\alpha$ -SG null mice treated with transduced mesoangioblasts (blue bar). Po/CSA values were significantly lower in  $\alpha$ -SG KO mice than in CTR and treated  $\alpha$ -SG KO mice, as indicated by the asterisk. Details of the Po/CSA values and of animal-to-animal variation are reported in the online supplement.

## MATERIALS and METHODS

### Cells and Culture conditions

The mouse myogenic cell line C2C12 (ATCC) was grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with heat-inactivated 10% fetal calf serum (FCS), 100 IU/ml penicillin and 100 µg/ml streptomycin. Myogenic differentiation was induced by replacing 10% FCS with 2% horse serum in the culture medium. Murine satellite cells were isolated and cultured as previously described (S1). Briefly, muscle was finely minced with scissors, digested with 2 mg/ml dispase, 0.1 mg/ml collagenase in phosphate buffered saline (PBS) for 45' at 37°C, washed in complete medium, and pipetted to obtain a single-cell suspension. Cell suspension was pre-plated twice to deplete early adhering non myogenic cells and then cultured on collagen coated dishes in DMEM containing 10% horse serum and 3% chick embryo extract. Mesoangioblasts were isolated from explants of mouse embryonic and juvenile vessels as previously described and grown in RPMI supplemented with heat-inactivated 10% fetal calf serum (FCS), 100 IU/ml penicillin and 100 mg/ml streptomycin (3). Briefly, explants were cultured for a week, then dissociated to a single cell suspension and cloned a feeder layer of mitomycin C-treated STO fibroblasts. On average, 30-50 clones developed from an explant of embryonic aorta, and only 5-10 from post-natal juvenile vessels. Virtually, all growing clones showed the same cobblestone morphology and the same pattern of marker expression (i.e. CD34, Flk-1, Sca-1, Thy1 and c-Kit). Several embryonic clones (named D15, D16 and D63) were selected for the experiments reported here, but only data relative to D16 (male) are shown. Post-natal mesoangioblasts were also isolated from 15 day old  $\alpha$ -SG null mice. On average more clones (up to 30 per explant) were obtained from vessels of dystrophic muscles, but many of these underwent senescence and only few continued to grow at a rate indistinguishable from their wild type counterparts (average division time: 24 h). One of these clones was used in the experiments reported here. Mesoangioblasts were transduced in vitro with third generation lentiviral vectors (I3) encoding either reporter genes (nuclear LacZ or GFP) or the  $\alpha$ -SG/GFP fusion protein (see below). In some experiments, mesoangioblasts were labeled with Dil (Molecular Probe, USA) as described (S2) and then injected through the femoral artery of mice. Cells number was assessed by direct cell count, and cell viability by Trypan blue dye (Sigma) exclusion.

### Immunoblotting

Cells ( $1.5 \times 10^6$  cells) were washed three times with PBS and lysed for 5 min with lysis buffer (50 mM Tris/HCl, pH 7.4, 1 mM EDTA, 1 mM EGTA 1% Triton X-100, 1 mM, and protease inhibitor cocktail (Sigma)). Tissue samples from Gastrocnemius, and Quadriceps of control, mesoangioblast-treated and untreated  $\alpha$ -SG null mice were homogenized with the same lysis buffer and centrifuged at 1000g for 20 min at 20°C to discard nuclei and cellular debris. Protein concentration was determined by bicinchoninic acid (BCA) protein assay (Pierce) using bovine serum albumin as standard. The membrane fraction was obtained by ultracentrifugation (70000g for 40 min at 4°C). Either total homogenate or membrane (pellet) and the cytosolic (supernatant) fractions were separated by 10% dodecyl sulphate polyacrylamide gel electrophoresis (SDS PAGE). For Western blot analysis, proteins were transferred to Immobilon (Amersham) membranes, saturated with 1% Bovine Serum Albumin (BSA), 0.1% Tween-20 (Sigma) in PBS (blocking solution) and reacted overnight at 4°C with  $\alpha$ -SG monoclonal antibody (Novocastra) at 1:100 dilution, or with MF20 monoclonal antibody (S3) at 1:20 dilution, or with anti  $\beta$ -1 integrin (a gift from I. De Curtis; S4) at 1:100 dilution. The filters were washed three times (15 minutes each at RT) with blocking solution

and then reacted with anti-mouse secondary antibody conjugated with horse radish peroxidase (HRP) IgG (Amersham) at 1:1000 dilution for 1 hour at RT, washed three times and finally visualized with the ECL immunoblotting detection system (Pierce) (S5). In the experiments aimed to detect an immune reaction to  $\alpha$ -SG or to GFP, membrane fractions were isolated as described above from rabbit muscle, separated by 10% SDS-PAGE and transferred to nitrocellulose as described above. Filters were reacted with various dilutions of the sera from mesoangioblast-treated animal or with anti- $\alpha$ -SG (see above) or with anti-GFP rabbit antibody (Molecular Probes) at 1:500 dilution. The procedure was as described above except that a secondary anti rabbit IgG, HRP conjugated (Sigma) was used.

### Immunofluorescence

Cells were grown on glass coverslips until sub-confluent, washed with PBS and fixed with 4% paraformaldehyde at room temperature for 10 minutes. Muscle samples from control,  $\alpha$ -SG null or mesoangioblast-treated  $\alpha$ -SG null mice were frozen in liquid nitrogen cooled isopentane and serial 10 mm thick sections were cut with a Leyca cryostat. Cells were permeabilized with 0.1% Triton X-100, 0.2% BSA in PBS for 10 minutes at RT, while tissue sections were incubated without detergent. Cells and tissue sections were washed three times with 0.2% BSA in PBS, and incubated overnight at 4°C with the following primary antibodies: rabbit polyclonal anti  $\alpha$ -SG at 1:20 dilution (S6) anti  $\beta$ -SG, anti  $\delta$ -SG, anti  $\gamma$ -SG rabbit polyclonal antibodies at 1:20 dilution (I1); anti-dystrophin monoclonal antibody Dys2 (Novocastra) at 1:50 dilution; anti laminin monoclonal or polyclonal antibodies (Sigma) at 1:100 dilution; MF20 antibody at 1:5 dilution. Anti M-Cadherin monoclonal antibody was a gift from A. Starzinski-Powitz (S7), anti PECAM monoclonal antibody was a gift from E. Dejana (S8) and anti Smooth alpha actin monoclonal antibody was from Sigma. After incubation, samples were washed three times with BSA/PBS and then incubated with the appropriate FITC or TRITC conjugated anti mouse or anti-rabbit IgG for 1 hour at RT. After three final washes, the cover slips were mounted on glass slides using mowiol in PBS and analyzed under a fluorescent microscope (Nikon). Other tissue sections or cells were stained with X-Gal as described (S2) or with Hematoxylin-Eosin or with Azan Mallory according to standard procedures.

### Intra-artery delivery of mesoangioblast in $\alpha$ -SG KO mice

Approximately  $5 \times 10^5$  mesoangioblasts or satellite cells were injected into  $\alpha$ -SG null C57BL/6 dystrophic mice. Mice were anesthetized with an intraperitoneal injection of physiologic saline (10 ml/kg) containing ketamine (5 mg/ml) and xylazine (1 mg/ml) and a limited incision on the medial side of the leg was performed. Cells were injected via a 0.20-mm-diameter needle inserted into the femoral artery. The needle was connected to a peristaltic pump by a heparinated Tygon tube (Ika Labortechnik). This Tygon tube was connected to a sterile Eppendorf tube containing  $1.6 \times 10^6$  cells in 200 µl. Cells were delivered by laminar flow (5 µl) over a period of 10 s. The blood flow was not stopped before or during this procedure. There was no visible damage to the vessel wall during or after operation. The body wall muscle was closed with sutures and the skin with surgical staples. Animals were sacrificed at different times after the injection. In some experiments three consecutive injections at 40 days interval were performed.

### In situ fluorescence hybridization

After immunostaining, fluorescent in situ hybridization (FISH) was performed according to the manufacturer instructions of Vysis (Cep Y: Vysis Inc). Briefly, the processed sections were post-fixed with paraformaldehyde 4% for 5 min at RT and denatured in 70% formamide, 2xSSC at 72°C for 3 min. After dehydration in ethanol, sections were incubated overnight at 42°C with denatured

FITC-CEP y chromosome probe. After incubation, sections were washed three times (for 10 min each at 46°C) in 50% formamide in 2xSSC, once with 2xSSC and once for 5 min with 0.1% NP-40 in 2xSSC. Sections were counterstained with DAPI and analyzed with a Nikon fluorescence microscope. The kit is based on a probe recognizing Y chromosome specific tandem repeat DNA sequences and therefore fluorescence is diffused over the nucleus except than in mitotic cells.

### **Methods used to study force (Po), cross sectional area (CSA) and maximum shortening velocity (Vo)**

#### Single fiber dissection, solutions and experimental set-up

The methods used for single fiber dissection, the solutions and the experimental set-up have been previously used and described in detail (S9, 18). Briefly, muscle bundles to be used for single fiber analysis were pinned at the bottom of a Petri dish covered with Sylgard and filled with a 1:1 mixture of skinning solution and glycerol. The Petri dish was stored at -20°C for up to three weeks. On the day of the experiment, a bundle was transferred to a dish containing skinning solution maintained at 10°C and, under a stereomicroscope (Wild M3 10-60 x magnification), single fibers were manually dissected. At the end of the dissection the skinning solution was replaced with a new one containing 1% Triton X-100 to ensure membrane solubilization. After 1 hour the dish was refilled with a solution without Triton. Segments ~1mm of length were cut from the fibers and light aluminum clips were applied at both ends of the segments to attach them to the beams of the force transducer (AE 801 SensoNor, Horten, Norway) and of the isotonic lever (model 101 vibrator, Ling Dynamic System, Royston, UK) in the experimental setup.

Skimming (5 mM EGTA, pCa 9.0), relaxing (5 mM EGTA, pCa 9.0), pre-activating (EGTA 0.5 mM, pCa 9.0) and activating (EGTA 5 mM pCa 4.5) solutions were prepared as previously described (S9). The experimental set-up enabled quick transfer of the muscle fibers from the first, larger chamber (~0.4 ml), containing skinning solution, to three, smaller chambers (70  $\mu$ l), containing relaxing, preactivating, and activating solution. The electromagnetic puller, could either keep the length of the fiber segment constant to elicit isometric contractions, or impose to the specimen quick releases of preset amplitude completed in 2 ms. A stereomicroscope was fitted over the apparatus to view the fiber at 20-60X magnification during the mounting procedure and during the experiment. The set up was placed on the stage of an inverted microscope (Axiovert 10, Zeiss, Germany). As the floors of the muscle chambers were made by cover slips, specimens could be viewed at 320X magnification through the eyepieces of the microscope. A video camera (MICAM HRS, System Sud, Les Ulis, France), fitted to the camera tube of the microscope and connected through an A/D converter (Cyclope, System Sud, Les Ulis, France) to a computer (Olivetti M24), allowed to view on a TV screen at approximately 1000X magnification, and to store digitized images of the specimen during experiment. The signals from the force and displacement transducer were visualized on the screen of a storage oscilloscope (mod 5113, Tektronix, Beaverton, Oregon, USA) and on a chart recorder (WR3701, Graphtec, Japan). The signals after A/D conversion (interface CED 1401 plus, Cambridge, UK) were fed into a personal computer and stored into the hard disk. Data storage, recall and analysis were done using Spike 2 (CED, Cambridge, UK) analysis software implemented on the computer.

#### Single fiber analysis

As skinned fibers lack plasma membrane, they need to be activated by exposure to solution containing the activating ion, Ca<sup>++</sup>. As intact fibers cannot be dissected from skeletal muscles of small mammals, skinned fibers have been widely used to study contractile properties of muscle fibers from small mammals and humans and have proved to be very reliable specimen for this kind of analysis (19,

S10). In mechanical experiments in the present work, temperature was set at 12 °C. Sarcomere length (S.L.) was determined by counting striations in segments of known length at 320X magnification and set at 2.5  $\mu$ m by varying fiber length at rest. Cross sectional area (CSA) of the specimen was determined assuming a circular shape from the mean of the three diameters measured at 320X magnification, without correction for swelling. For force (Po) and maximum shortening velocity (Vo) determinations fibers were first transferred to pre-activating solution for at least 2 minutes and then maximally activated (pCa 4.45) for about 40-60 sec. To determine Vo, slack-test maneuvers were employed (S11). Details of Vo determinations have been reported previously (17). Vo was expressed in fiber length per second (Us), Po in mg and Po/CSA in kN/m<sup>2</sup>. At the end of the mechanical experiment fibers were put in 20 ml of standard buffer (S12) and stored at -20 °C for subsequent analysis of MyHC isoform content.

#### Fiber typing and myosin isoform identification

The MyHC isoform composition of the single muscle fiber segments used in mechanical experiments was determined by polyacrylamide-gel electrophoresis on 6% polyacrylamide slab gels after denaturation in SDS (SDS-PAGE) with a procedure previously described (S12,18). In the MYHC region, four bands corresponding to the four adult MyHC isoforms (MyHC-1, MyHC-2A, MyHC-2X, and MyHC-2B) could be separated. In relation to the presence of one or two bands in the MyHC region, fibers were classified in four pure fiber types and three hybrid fiber types: 1, 2A, 2X, 2B (or pure fibers), 1-2A, 2AX and 2XB (or hybrid fibers). The same electrophoretical protocol followed by densitometric analysis of MyHC bands was used to determine the MyHC isoform composition of whole muscle samples as previously described (S13). Gastrocnemius muscle of the mice used in the present work were found to contain mostly MyHC-2B and 80% of the single muscle fibers studied contained MyHC-2B.

#### **Statistics**

Values were expressed as means  $\pm$  standard deviation (SD). Statistical significance of the differences between means was assessed by one way analysis of variance (ANOVA) followed by the Student-Newman-Keuls test to determine which groups were statistically significantly different from the others. When only two groups had to be compared, unpaired Student's t test was used. A probability of less than 5% (P<0.05) was considered to be statistically significant.

Tables

**Table I.**  
Engraftment of donor cells into  $\alpha$ -SG null dystrophic mice.

Tissue of engrafted mice	Mesoangioblasts	
	Wt	$\alpha$ -SGC ko, genetically corrected
<b>Tibialis Anterior (all)</b>	<b>-:2/16; ++:8/16; +++:6/16</b>	<b>-:1/5; ++:2/5; +++:2/5</b>
TA (4 wks)	-:1/4; ++:2/4; +++:1/4	-:1/2; ++:1/2
TA (8 wks)	-:1/6; ++:4/6; +++:1/6	ND
TA (24 wks*)	++:2/6; +++:4/6	++: 1/3; +++:2/3
<b>Soleus (all)</b>	<b>-:2/16; ++:5/16; +++:9/16</b>	<b>-:1/5; ++:2/5; +++:2/5</b>
Soleus (4 wks)	-:1/4; ++:2/4; +++:1/4	-:1/2; ++:1/2
Soleus (8 wks)	-:1/6; ++:4/6; +++:1/6	ND
Soleus (24 wks*)	++:2/6; +++:4/6	++: 1/3; +++:2/3
<b>Gastrocnemius (all)</b>	<b>-:2/16; ++:2/16; +++:12/16</b>	<b>-:1/5; +++:4/5</b>
Gas. (4wks)	-:1/4; ++:1/4; +++:2/4	-:1/2; +++:1/2
Gas. (8 wks)	-:1/6; ++:1/6; +++:4/6	ND
Gas. (24 wks*)	+++6/6	+++ 3/3
<b>Quadriceps (all)</b>	<b>-:2/16; +++:14/16</b>	<b>-:1/5; +++:4/5</b>
Quad. (4 wks)	-:1/4; +++:3/4	-:1/2; +++:1/2
Quad. (8 wks)	-:1/6; +++:5/6	ND
Quad. (24 wks*)	+++6/6	+++ 3/3
<b>Diaphragm (all)</b>	<b>-:12/14; +:2/14</b>	<b>-:5/5;</b>
Diaphr. (4 wks)	-:3/4; +:1/4;	-: 2/2
Diaphr. (8 wks)	-:6/6;	ND
Diaphr. (24 wks*)	-:3/4; +:1/4	-:3/3
<b>Plantaris (8 wks)</b>	<b>-:1/2; ++:1/2</b>	<b>ND</b>
<b>Contra-lateral TA (8 wks)</b>	<b>-:4/6; +:2/6</b>	<b>-:2/5; +:3/5</b>
<b>Kidney (4 wks)</b>	<b>++: 2/4 (PCR)</b>	<b>ND</b>
<b>Lung (4 wks)</b>	<b>++: 2/4 (PCR)</b>	<b>ND</b>
<b>Liver (4 wks)</b>	<b>++: 2/4 (PCR)</b>	<b>ND</b>
<b>Bone marrow (4 wks)</b>	<b>-:4/4 (PCR)</b>	<b>ND</b>

Unless otherwise specified, donor cell derived fibers were detected by Immunofluorescence (with antibodies recognizing  $\alpha$ -SG). -: no positive fibers detected; +: less than 5%  $\alpha$ -SG positive fibers on at least 5 non serial sections; ++: from 5 to 20% positive fibers; +++: more than 20% positive fibers. ND: not done. Few animals did not show any donor cell engraftment, possibly because of technical problems of the intra-femoral artery injection. Animals sacrificed at 4 or 8 weeks had received only one injection; \*animals sacrificed at 24 weeks had received three consecutive injections (see methods for details).

**Table II.** Details on specific force (Po/CSA) of single muscle fibers from control mice (CTR),  $\alpha$ -SG null ( $\alpha$ -SG KO) mice and mesoangioblast treated  $\alpha$ -SG null ( $\alpha$ -SG KO-treated) mice reported in Fig. 3.

The following table reports the mean values of Po/CSA ( $\pm$ SD) for each of the three animals per group used for Fig. 3 and the overall mean of Po/CSA of Fig. 3.

Animal N°	Po/CSA (kN/m <sup>2</sup> )		
	CTR	$\alpha$ -SG KO	$\alpha$ -SG KO treated with WT mesoangioblasts
1	60.69 $\pm$ 22.17 n=25	46.72 $\pm$ 24.46*	73.08 $\pm$ 32.14 n=22
2	60.83 $\pm$ 23.84 n=25	38.41 $\pm$ 20.43*	61.01 $\pm$ 22.19 n=38
3	63.58 $\pm$ 31.11 n=32	37.50 $\pm$ 15.73*	65.58 $\pm$ 28.24 n=29
<b>Total</b>	<b>61.76<math>\pm</math>24.80 n=82</b>	<b>42.01<math>\pm</math>21.63 n=106</b>	<b>65.48<math>\pm</math>27.02 n=89</b>

\* significantly weaker than any control or  $\alpha$ -SG KO treated mice. Note that single muscle fibers from any of the three  $\alpha$ -SG KO mice were significantly weaker than single muscle fibers from any of the three  $\alpha$ -SG KO treated mice, whereas no statistically significant difference was found within each group (control,  $\alpha$ -SG KO and  $\alpha$ -SG KO treated), and between  $\alpha$ -SG KO treated and control groups.

**Table III.** Details on specific force (Po/CSA) of single muscle fibres from control mice (CTR),  $\alpha$ -SG null ( $\alpha$ -SG KO) mice and

mesoangioblast treated  $\alpha$ -SG null ( $\alpha$ -SG KO treated) mice reported in Fig. 4E.

Animal N°	Po/CSA (kN/m <sup>2</sup> )		
	Control	$\alpha$ -SG KO	$\alpha$ -SG KO treated by autologous genetically corrected mesoangioblasts
1	62.87 $\pm$ 25.1 n=25	43.48 $\pm$ 20.22 n=21	60.28 $\pm$ 23.32 n=26
2		35.32 $\pm$ 22.00 n=11	62.60 $\pm$ 22.19 n=12
<b>Total</b>	<b>62.87<math>\pm</math>25.1 n=25</b>	<b>40.70<math>\pm</math>20.86 n=32</b>	<b>61.00<math>\pm</math>22.20 n=38</b>

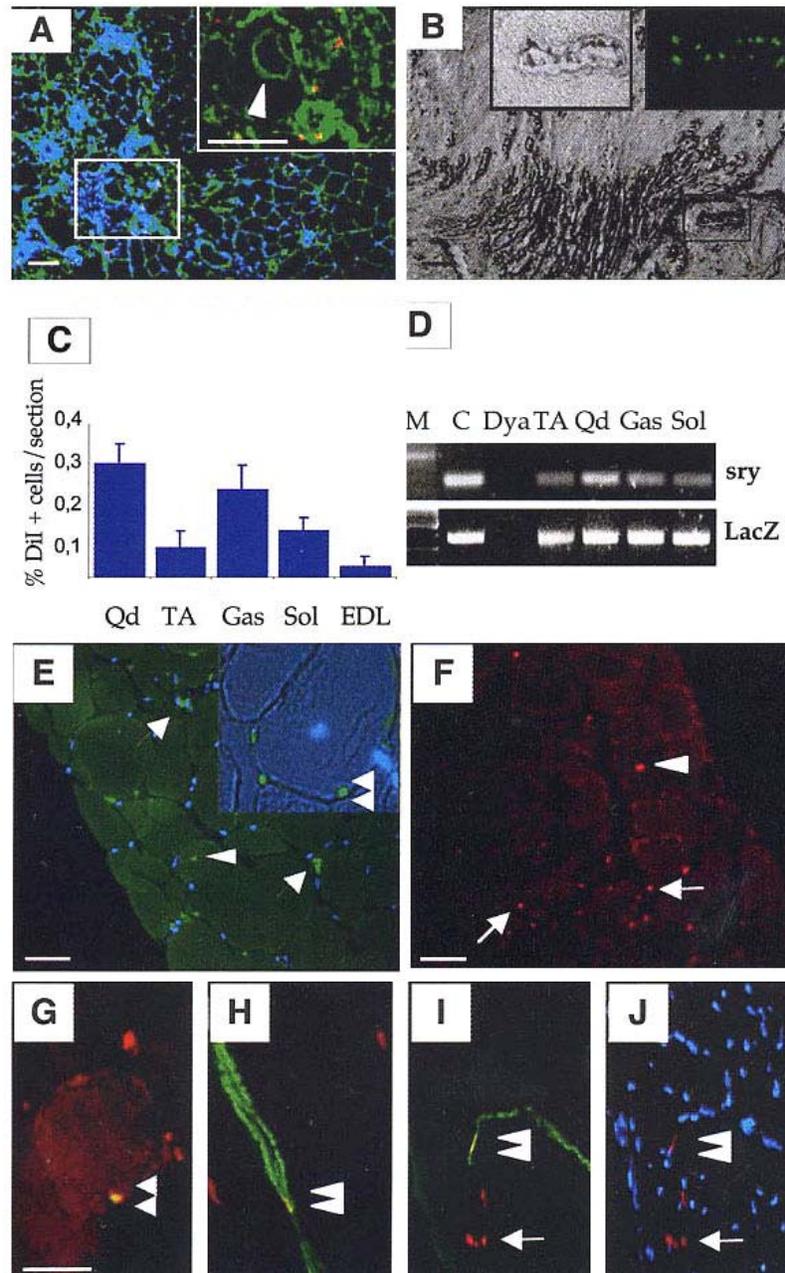
The table below reports the data of the one control, the two dystrophic and the two dystrophic, treated mice used for Fig. 4E. The number of animals analyzed is lower than in Fig. 3 because control and  $\alpha$ -SG KO animals used for Fig 4 belong to the same group as those of Fig. 3A and have very similar Po/CSA values and because both mice treated by autologous genetically corrected mesoangioblasts showed a recovery of Po/CSA identical to that shown in the animals treated by WT meso-angioblast reported in Fig 3.

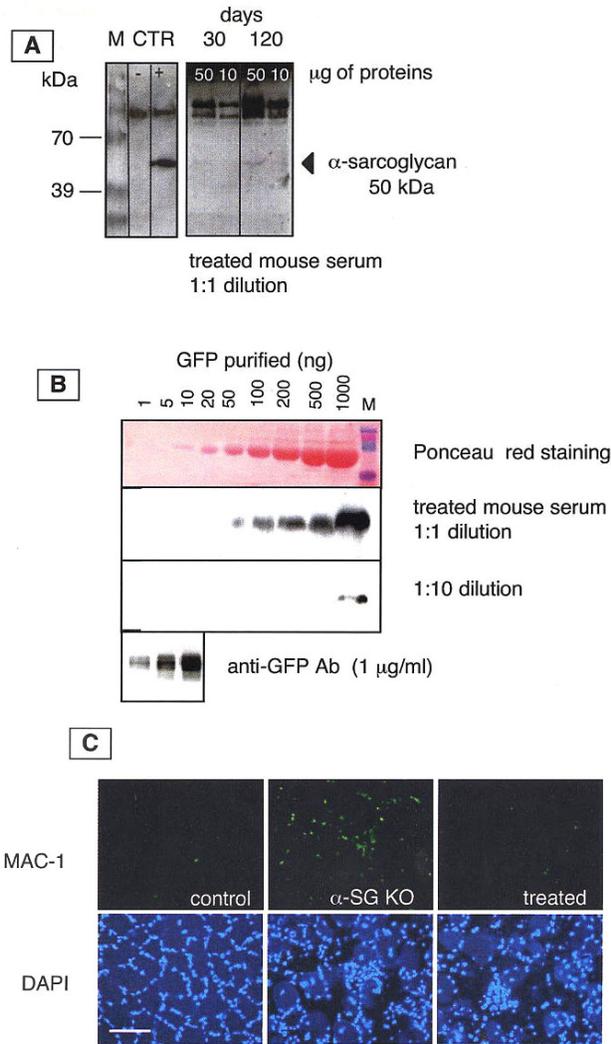
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## Figure legends

**Fig. S1.** Tissue distribution and differentiation of mesoangioblasts after intra-arterial delivery. Wt, male mesoangioblasts were transduced with a lentiviral vector encoding for nuclear LacZ (*13*) or labeled with Dil and then injected into the femoral artery of  $\alpha$ -SG null dystrophic female mice. **(A)** Low magnification of an area of degeneration/regeneration of the TA, stained with anti-laminin antibodies (green) and Dapi (blue). A higher magnification (inset) revealed many Dil labeled mesoangioblasts (red) outside of the small vessels and within the cellular infiltrate. **(B)** Similarly injected satellite cells isolated from GFP mice (*12*) did not cross the vessel wall and remained adherent to the endothelium for a few hours after the injection; high magnification of injected cells, both in phase contrast and in fluorescence, is shown in the insets. **(C)** Distribution Dil positive cells, 5 days after intra-arterial injection, counted in five non adjacent cross-sections of the TA, the Quadriceps, the Gastrocnemius the Soleus and the EDL muscles of the injected leg and expressed as percentage of total cells (identified after Dapi staining of the same sections). **(D)** PCR with oligos specific for the *sry* and *LacZ* genes reveals the presence of donor cells in the TA, the Quadriceps, the Gastrocnemius and the Soleus of the injected limb but not in the diaphragm. DNA from a male MLCSF-nLacZ mouse was used as a positive control. **(E)** FISH analysis of Y chromosome positive cells revealed a widespread distribution near the capillary network (arrows) and, at higher magnification (inset), both outside and underneath (double arrow) the basal lamina of regenerated fibers, identified by a centrally located nucleus (blue after Dapi staining). A Y chromosome positive nucleus is shown by an arrowhead in the center of a regenerating fiber. **(F)** Staining with an antibody against ( $\beta$ -galactosidase (red) also revealed mesoangioblasts near the capillary network (arrows) but also incorporated into the center of a regenerated fiber (arrowhead). **(G)** Double staining antibodies against M-Cadherin (a marker of satellite cells) and p-galactosidase. Double arrow shows a cell localized at the periphery of a fiber (identified by background fluorescence) that appears yellow in the merged image, revealing that some of ( $\beta$ -gal labeled mesoangioblasts (red), express markers typical of satellite cells. **(H,I,J)** Double staining with antibodies against ( $\beta$ -galactosidase (red) and smooth alpha actin (green in H) or PECAM (green in I) revealed that a small percentage of injected mesoangioblasts expressed smooth muscle or endothelial markers (double arrowheads showing cells that appear yellow in the merged images) while the majority did not (arrows). Nuclei were stained blue with Dapi in J. A, B, bar = 200  $\mu$ m; E-J, bar = 50  $\mu$ m.





**Fig. S2.** Immune response to GFP and  $\alpha$ -SG proteins after mesoangioblast injection. **(A)** Western Blot analysis showing the presence of antibodies against  $\alpha$ -SG protein in the serum of the mice treated with wild type mesoangioblasts. Only undiluted serum of treated animals weakly reacted with  $\alpha$ -SG protein from a membrane fraction (50  $\mu$ g) of rabbit skeletal muscle after a long (120 days) and very weakly after a short (30 days) treatment. No signal was detected with serum from untreated mice (- lane CTR panel) while a strong band was observed using a monoclonal antibody against  $\alpha$ -SG as a positive control (+ lane CTR panel). **(B)** Western Blot analysis showing the presence of antibodies against GFP protein in the serum of the mice treated with Lenti-PGK $\alpha$ -SG-GFP transduced-mesoangioblasts. The serum of treated animals revealed 50 or more ng of purified GFP protein when undiluted (second lane) and only 1 mg when diluted 1:10 (third lane), indicating a very weak response; in comparison commercial anti-GFP antibody at 1:2000 dilution readily detected 1 ng of the same protein (fourth lane). Red Ponceau staining of the filter is shown in the first lane. No signal was detected with serum from untreated mice (data not shown). **(C)** Immunofluorescence images of the Gastrocnemius of a 6 month-old wild type (control) mouse, of an  $\alpha$ -SG null animal of the same age ( $\alpha$ -SG KO) and of an  $\alpha$ -SG null animal treated as described above (treated). Sections were reacted with a monoclonal antibody against MAC-1, a specific membrane protein of macrophages (green); sections were also stained with Dapi (blue). The presence of macrophages in the skeletal muscle tissue of treated dystrophic mice (treated) was reduced at levels comparable with control; bar = 100  $\mu$ m.

