

## ***ACTN3* genotype influences muscle performance through the regulation of calcineurin signaling**

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Research Article

Muscle biology

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# ACTN3 genotype influences muscle performance through the regulation of calcineurin signaling

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**$\alpha$ -Actinin-3 deficiency occurs in approximately 16% of the global population due to homozygosity for a common nonsense polymorphism in the *ACTN3* gene. Loss of  $\alpha$ -actinin-3 is associated with reduced power and enhanced endurance capacity in elite athletes and nonathletes due to “slowing” of the metabolic and physiological properties of fast fibers. Here, we have shown that  $\alpha$ -actinin-3 deficiency results in increased calcineurin activity in mouse and human skeletal muscle and enhanced adaptive response to endurance training.  $\alpha$ -Actinin-2, which is differentially expressed in  $\alpha$ -actinin-3-deficient muscle, has higher binding affinity for calsarcin-2, a key inhibitor of calcineurin activation. We have further demonstrated that  $\alpha$ -actinin-2 competes with calcineurin for binding to calsarcin-2, resulting in enhanced calcineurin signaling and reprogramming of the metabolic phenotype of fast muscle fibers. Our data provide a mechanistic explanation for the effects of the *ACTN3* genotype on skeletal muscle performance in elite athletes and on adaptation to changing physical demands in the general population. In addition, we have demonstrated that the sarcomeric  $\alpha$ -actinins play a role in the regulation of calcineurin signaling.**

## Introduction

$\alpha$ -Actinin-3 is one of the major components of the skeletal muscle Z-disk in fast-twitch muscle fibers (1) and interacts with multiple structural, metabolic, and signaling proteins (2, 3). Homozygosity for a common nonsense polymorphism in the *ACTN3* gene (R577X) results in complete  $\alpha$ -actinin-3 deficiency in an estimated 16% of the global population (4) and is associated with variations in human muscle performance. The *ACTN3* 577XX-null genotype is markedly underrepresented in elite sprint and power athletes (5–9) and is associated with reduced muscle strength and sprint performance in nonathlete cohorts (10–13), suggesting that  $\alpha$ -actinin-3 deficiency has a detrimental effect on the optimal function of fast muscle fibers. In contrast, the *ACTN3* 577XX genotype is overrepresented in elite endurance athlete cohorts (5, 14, 15), suggesting a beneficial effect on endurance capacity. Recent studies in athletes and nonathletes further suggest that the *ACTN3* genotype influences the adaptive response of skeletal muscle to exercise training (10, 16).

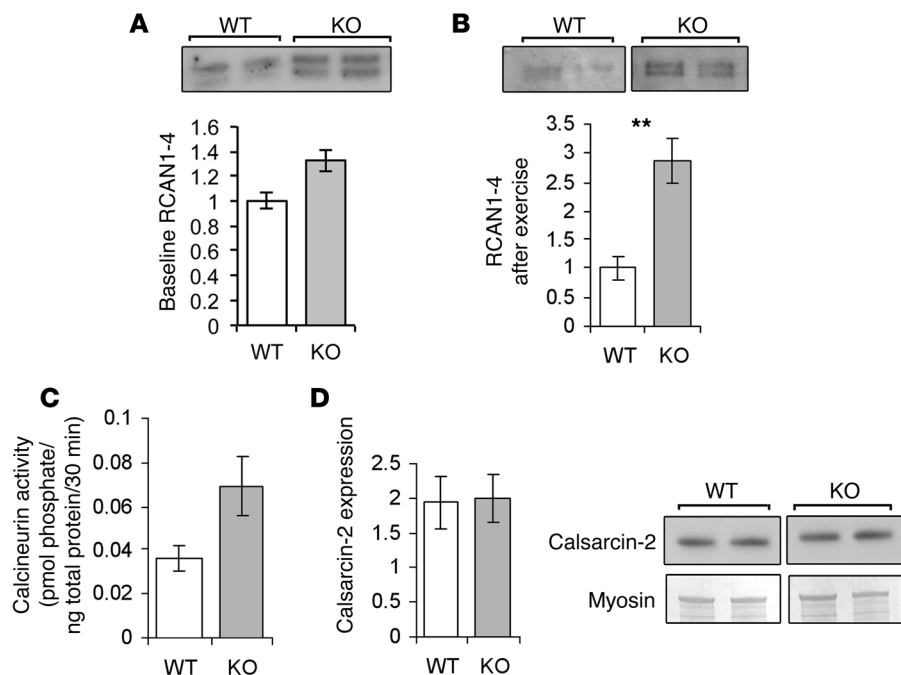
The *Actn3* KO mouse model mimics the phenotypic effects of  $\alpha$ -actinin-3 deficiency in humans (17). The closely related sarcomeric isoform  $\alpha$ -actinin-2 compensates for the absence of  $\alpha$ -actinin-3 and is expressed in all fiber types in *Actn3* KO mice, similarly to *ACTN3* 577XX humans. Compared with WT mice, *Actn3* KO mice have substantially lower grip strength, increased recovery from fatigue, and enhanced endurance exercise performance associated with increased levels of glycogen and a shift in fast muscle fiber properties toward a slow-twitch, oxidative phenotype, without

changes to fiber typing as defined by myosin heavy chains (17–19). The metabolic phenotype in the *Actn3* KO mouse is similar to that seen in skeletal muscle following exercise training, suggesting that *Actn3* KO muscle is “pretrained” for endurance performance.

The molecular basis for the effects of  $\alpha$ -actinin-3 deficiency on fast muscle fiber properties is not yet known. One potential downstream target is the calcium- and calmodulin-dependent protein phosphatase calcineurin, which indirectly associates with sarcomeric  $\alpha$ -actinins at the Z-disk via their mutual binding to calsarcins (20–23). It is well established that calcineurin signaling plays a critical role in the remodeling of skeletal muscle; it specifically induces activation of the slow myogenic program and a shift in muscle fiber attributes toward a slow-twitch, oxidative phenotype (24, 25) and confers fatigue resistance in endurance running (26). The expression of activated calcineurin in fast glycolytic skeletal muscles has been shown to induce a shift in glucose metabolism toward decreased glucose oxidation and increased insulin-stimulated glucose incorporation into glycogen (27). It has recently been shown that calsarcin-2, which, like  $\alpha$ -actinin-3, is expressed exclusively in fast-twitch muscle, inhibits calcineurin signaling in vivo. Mice deficient in calsarcin-2 display enhanced endurance capacity and their muscles exhibit a shift toward slower fiber properties, similar to the phenotype observed in the *Actn3* KO mouse model (28). There were no changes in the expression of  $\alpha$ -actinin-2 and -3 in the muscles of calsarcin-2 KO mice, raising the possibility that the  $\alpha$ -actinins act upstream of calsarcin-2 as key regulators of muscle fiber phenotype (28). We hypothesized that the phenotypic changes observed in  $\alpha$ -actinin-3-deficient fast fibers are due to increased calcineurin activity, which in turn is regulated by the differential binding of calsarcin-2 to sarcomeric  $\alpha$ -actinins.

**Conflict of interest:** The authors have declared that no conflict of interest exists.

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**Figure 1**

Calcineurin signaling is increased in  $\alpha$ -actinin-3-deficient muscles. **(A)** Muscles of KO mice on a C57BL/6J background showed a 1.3-fold increase in RCAN1-4 protein expression compared with WT muscles, suggesting increased calcineurin activity ( $P = 0.057$ ;  $n = 3-4$  for both groups). **(B)** In exercised mice on a 129 background, RCAN1-4 was 2.9-fold upregulated in the KO group ( $P = 0.004$ ;  $n = 6$  for both groups). **(C)** Direct assays of calcineurin activity confirmed a 1.9-fold increase in calcineurin signaling in exercised KO muscles ( $P = 0.093$ ;  $n = 6$  for both groups). **(D)** Calsarcin-2 expression was similar between WT and KO muscles (mean  $\pm$  SEM;  $**P < 0.01$ , Mann-Whitney  $U$  test). In **B** and **D**, the lanes were run on the same gel but were noncontiguous.

In this study, we demonstrate that calcineurin activity is increased in the absence of  $\alpha$ -actinin-3 and provide a molecular mechanism to explain the slower metabolic, physiological, and functional phenotypes associated with  $\alpha$ -actinin-3 deficiency. Increased calcineurin activity is also associated with increased muscle plasticity, as demonstrated by an enhanced adaptive response to endurance training in *Actn3* KO mice and an increased switch in muscle fiber type from fast-twitch glycolytic fibers toward fast-twitch oxidative fibers. We further demonstrate that calsarcin-2, a key inhibitor of calcineurin activation, preferentially binds to  $\alpha$ -actinin-2 over  $\alpha$ -actinin-3 and that  $\alpha$ -actinin-2 outcompetes calcineurin for binding to calsarcin-2. We propose that the upregulation of  $\alpha$ -actinin-2, in the absence of  $\alpha$ -actinin-3, releases calcineurin from calsarcin-2 inhibition and alters the metabolic phenotype of fast glycolytic (2B) muscle fibers. Taken together, our data suggest a novel role for the sarcomeric  $\alpha$ -actinins in the phenotypic regulation of skeletal muscle fibers at baseline and in response to physical demands via modulation of calcineurin-dependent signaling.

**Results**

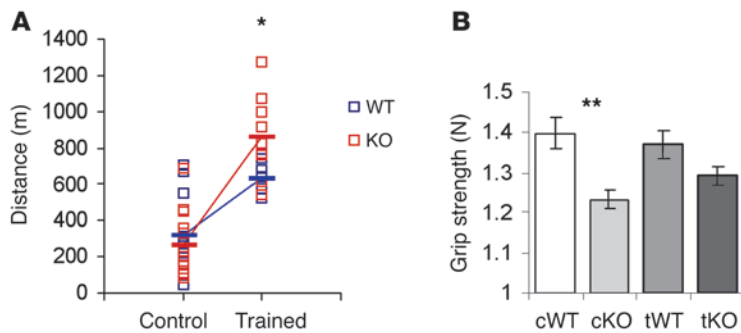
*Calcineurin activity is increased in  $\alpha$ -actinin-3-deficient muscles.* To determine whether the phenotypic changes in  $\alpha$ -actinin-3-deficient muscle are associated with increased calcineurin activity, we first examined the expression of RCAN1-4, a downstream regulator of calcineurin, which is specifically increased upon activation of calcineurin and, in turn, stimulates calcineurin signaling when expressed at physiological levels (29). Above physiological levels, RCAN1-4 also acts in a negative feedback loop to inhibit further calcineurin activity (30, 31). The expression of *RCAN1* is tightly controlled by a cluster of 15 NFAT binding sites and is thus a sensitive endogenous calcineurin reporter gene (30). At baseline, *Actn3* KO muscles showed a 1.3-fold increase in the levels of RCAN1-4 compared with WT muscles, however, this failed to reach formal significance ( $P = 0.057$ ) (Figure 1A). As calcineurin is known to play a critical role in promoting the transition of fast-twitch muscle

fibers toward a slow-twitch phenotype in response to low-amplitude, sustained calcium activity (32), we further examined the levels of RCAN1-4 in muscles from WT and KO mice 3 days after a single bout of endurance running; this approach has previously been used to identify gene expression changes in muscle in response to endurance exercise (33). Our results show that the muscles of *Actn3* KO mice have significantly elevated expression of RCAN1-4 after exercise compared with the muscles of WT mice (2.9-fold;  $P = 0.004$ ) (Figure 1B), which is consistent with potentiated calcineurin activity in the absence of actinin-3.

To confirm that calcineurin activity is increased in unexercised and exercised *Actn3* KO muscles, we directly assayed the total phosphatase activity and the calcium-independent phosphatase activity in lysates of muscles. Calcineurin activity is calcium dependent and is determined by subtracting the level of calcium-independent phosphatase activity from the total phosphatase activity. Consistent with RCAN1-4 results, exercised *Actn3* KO muscles demonstrated a 1.9-fold increase in calcineurin activity compared with strain-matched WT muscles (Figure 1C). There were no differences in the activity of calcium-independent phosphatases between WT and *Actn3* KO samples (data not shown), indicating a specific influence of sarcomeric  $\alpha$ -actinin isoform expression upon calcium-dependent phosphatase (calcineurin) activity in skeletal muscle.

Since calsarcin-2 has previously been demonstrated to inhibit calcineurin activity (28), we investigated whether differences in calcineurin activity between WT and *Actn3* KO muscles were associated with altered expression of the calsarcin-2 protein. We found that calsarcin-2 expression was not altered in *Actn3* KO muscles compared with WT muscles (Figure 1D), suggesting that sarcomeric  $\alpha$ -actinins regulate calcineurin activity via a mechanism that does not depend on differential expression of calsarcin-2.

*$\alpha$ -Actinin-3 deficiency potentiates the adaptive response to endurance training.* Increased calcineurin activity is associated with activation of the slow myogenic program and an enhanced response of skel-

**Figure 2**

Endurance performance and grip strength of WT and KO mice after endurance training. (A) The endurance performance of untrained control WT (cWT) and control KO mice (cKO) were not significantly different at 15 to 16 weeks of age, however, the endurance performance of KO mice was significantly greater than WT mice after training. Trained WT mice (tWT) ran twice the distance of the cWT mice, while trained KO mice (tKO) ran 3.3 times the distance of untrained KO mice. (B) No significant increase in grip strength was detected in either WT or KO mice with endurance training. WT mice demonstrated greater grip strength than KO mice in both the untrained and trained groups (mean  $\pm$  SEM; \* $P < 0.05$  for WT versus tKO; \*\* $P < 0.01$ , Mann-Whitney  $U$  test;  $n = 6-8$  for all groups).

etal muscle to increased physical demands. Since  $\alpha$ -actinin-3 deficiency is associated with enhanced endurance performance in elite athletes (5, 14), we hypothesized that increased calcineurin activity in  $\alpha$ -actinin-3-deficient muscles would contribute to a greater adaptive response to endurance training. To test this hypothesis, we compared the performance of WT and *Actn3* KO mice following 4 weeks of treadmill training with age-matched (16 weeks of age), untrained control cohorts (Figure 2A). We have previously shown that KO mice run farther than WT mice at 8 weeks of age (17), but this difference in baseline endurance performance is lost by 16 weeks of age (34). However, after 4 weeks of exercise training, 16-week-old *Actn3* KO mice were able to run significantly farther than the trained WT mice ( $P = 0.026$ ). Specifically, trained WT mice ran twice the distance of WT controls, while trained *Actn3* KO mice ran 3.3 times farther than *Actn3* KO controls. Significant interaction between genotype and training confirms an increased response to endurance training in *Actn3* KO mice ( $F = 5.32(1, 38)$ ,  $P = 0.027$ ), consistent with a significantly increased response to training in *Actn3* KO mice. In contrast, grip strength was not significantly altered with training in either WT or *Actn3* KO mice (Figure 2B). Similar to our previous results, untrained WT mice had a greater average grip strength compared with untrained *Actn3* KO mice ( $P = 0.003$ ) (18). A similar trend was observed for the trained group ( $P = 0.088$ ).

*Actn3* KO muscles exhibit increased oxidative enzyme activity and altered fiber type adaptation in response to endurance training. To examine the response to endurance training in greater detail, we assessed the activity of metabolic enzymes in the quadriceps muscles of sedentary and trained WT and *Actn3* KO mice. The quadriceps muscle was examined because it is one of the most activated muscles during uphill running (35). Consistent with previously published data, untrained control *Actn3* KO mice showed higher activities of enzymes involved in the glycolytic pathway (hexokinase [HK]) (Figure 3A), the mitochondrial aerobic pathway (citrate synthase [CS], succinate dehydrogenase [SDH]) (Figure 3, B and C), and the fatty acid oxidation pathway ( $\beta$ -hydroxyacyl CoA dehydrogenase

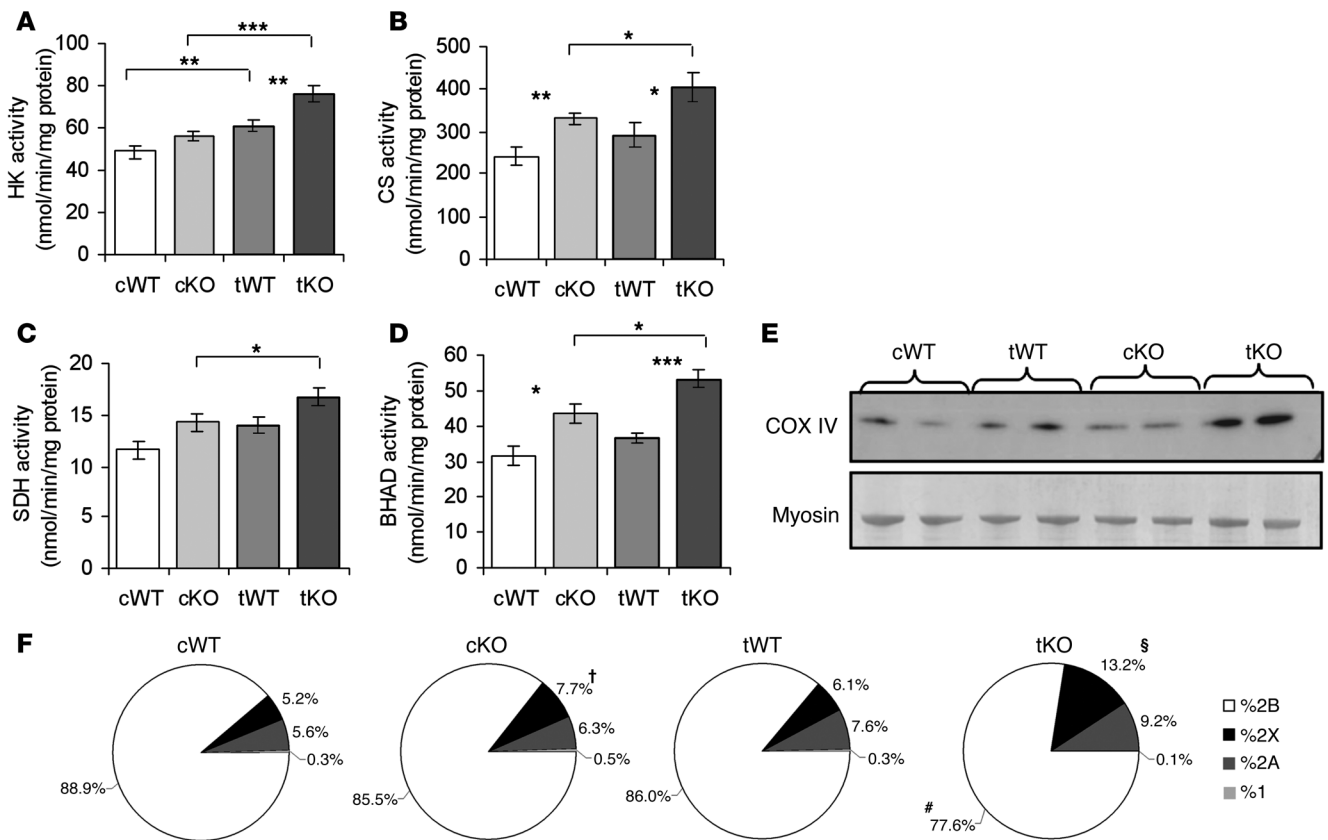
[BHAD]) (Figure 3D) (18). Following endurance training, the activity of CS, SDH, BHAD, and HK increased in both WT and *Actn3* KO muscles. Western blot analysis of the mitochondrial protein cytochrome c oxidase subunit IV (COX IV) similarly showed an increase in expression in response to endurance training (Figure 3E). The activities of oxidative enzymes in WT muscle after training were similar to the muscles of sedentary *Actn3* KO mice, supporting our proposal that *Actn3* KO muscle is “pretrained” for endurance performance.

Since increased calcineurin activity promotes adaptation toward the slow myogenic program in response to physical demands, we investigated whether *Actn3* KO muscles demonstrate greater changes in fiber size and fiber type proportions in response to endurance training in comparison with WT muscles. Consistent with previously published data (18), fiber type proportions (as defined by myosin heavy chain isoforms) did not differ between the muscles of untrained WT and *Actn3* KO mice (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI67691DS1). However, the effect of endurance training on fiber type proportions differed between the WT and *Actn3* KO muscles. Endurance training had a minimal effect on the proportions of 2B and 2X fibers in WT quadriceps; changes did not reach formal statistical significance (2B:  $-6.5\%$ ,  $P = 0.082$ ; 2X:  $2.0\%$ ,  $P = 0.537$ ). In the *Actn3* KO quadriceps, there was a  $9.1\%$  decrease in 2B fibers ( $P = 0.004$ ) and a  $5.2\%$  increase in 2X fibers ( $P = 0.04$ ). Similar increases in 2A fibers (in which  $\alpha$ -actinin-3 is not usually expressed) were observed in both WT and KO muscles ( $4.8\%$  WT,  $P = 0.008$ ;  $3.8\%$  KO,  $P = 0.015$ ). These results suggest that *Actn3* KO muscles have an increased propensity for fiber-type conversion toward a slower phenotype. Training also induced greater hypertrophy of 2X and 2A fibers in *Actn3* KO muscles (2X:  $24.4\%$ ,  $P = 0.051$ ; 2A:  $26.5\%$ ,  $P = 0.051$ ) compared with WT (2X:  $13.5\%$ ,  $P = 0.093$ ; 2A:  $5.5\%$ ,  $P = 0.589$ ), such that 2X fibers were found to be significantly larger in KO muscles compared with WT muscles after training (Supplemental Figure 2).

To examine the combined effect of changes in fiber size and number in each fiber type, we analyzed the total cross-sectional area (CSA) occupied by fibers of each type as a proportion of the overall muscle cross section (Figure 3F). Interestingly, control *Actn3* KO muscles showed a trend toward lower 2B CSA proportions ( $-3.3\%$ ,  $P = 0.093$ ) and significantly higher 2X CSA proportion ( $2.5\%$ ,  $P = 0.041$ ) compared with WT muscles. Endurance training did not significantly alter the CSA proportions of any fiber type in WT muscles. In contrast, *Actn3* KO muscles showed a significantly reduced 2B CSA of  $8.0\%$  ( $P = 0.041$ ) and an increased 2X CSA proportion of  $5.5\%$  ( $P = 0.009$ ) in response to training, resulting in further differences between WT and *Actn3* KO in 2B (WT:  $85.1\%$ , *Actn3* KO:  $77.6\%$ ;  $P = 0.004$ ) and 2X (WT:  $6.3\%$ , *Actn3* KO:  $13.2\%$ ;  $P = 0.004$ ) CSA proportions. This was confirmed by significant genotype and training interaction for 2X CSA using two-way ANOVA ( $F = 5.71(1, 19)$ ,  $P = 0.027$ ). Together, these results suggest that *Actn3* KO muscles demonstrate greater plasticity in relation to fiber-type shifts in response to endurance training.

*Muscles from  $\alpha$ -actinin-3-deficient humans exhibit increased calcineurin activity.* To confirm that our observations in *Actn3* KO mouse muscle are relevant to and reproducible in humans, we examined muscle biopsies from humans with *ACTN3* 577RR





**Figure 3**

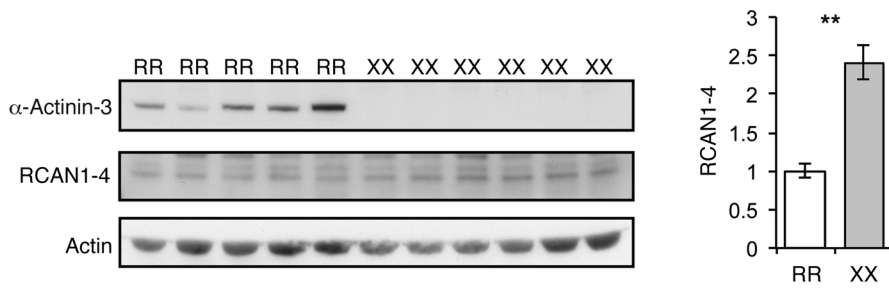
Enhanced response to exercise training in KO mice. Training increased activities of (A) HK, (B) CS, (C) SDH, and (D) BHAD in both WT and KO muscles. (E) Exercise training increased the level of COX IV in both WT and KO muscles, with KO muscles demonstrating higher levels of COX IV than WT in both untrained and trained states. (F) The CSA occupied by each fiber type is presented as a proportion of the overall muscle cross section. cKO muscles demonstrated 3.3% lower 2B CSA proportions ( $P = 0.093$ ) and 2.5% higher 2X ( $P = 0.041$ ) CSA proportions compared with cWT muscles. tWT muscles showed no significant alterations in CSA proportions for any fiber type compared with cWT muscles, while tKO muscles demonstrated significant decreases in 2B and increases in 2X CSA compared with cKO muscles (mean  $\pm$  SEM; \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ; † $P < 0.05$  for cWT versus cKO; # $P < 0.05$ ; § $P < 0.01$  for cKO versus tKO, Mann-Whitney  $U$  test;  $n = 6-7$  muscles; greater than 5,000 fibers per muscle for all groups).

genotypes (WT,  $n = 5$ , all female, age range 27–67 years, mean age 47 years) and 577XX ( $\alpha$ -actinin-3 deficient,  $n = 6$ , all female, age range 33–77 years, mean age 49 years) for differences in calcineurin activity. Consistent with our findings in the *Actn3* KO mouse, muscles from *ACTN3* 577XX humans showed increased RCAN1-4 expression compared with the muscles of *ACTN3* 577RR humans ( $P = 0.004$ ) (Figure 4).

*Calsarcin-2 preferentially binds to  $\alpha$ -actinin-2.* The expression of calsarcin-2 (which inhibits calcineurin activity; ref. 28) did not differ between the muscles of WT and *Actn3* KO mice and hence cannot account for the altered calcineurin activity associated with  $\alpha$ -actinin-3 deficiency. Since the sarcomeric  $\alpha$ -actinins and calcineurin share the same N-terminal binding site on calsarcin (23), we hypothesized that the observed differences in calcineurin activity between WT and *Actn3* KO muscles could be associated with a differential binding affinity between calsarcin-2 and the  $\alpha$ -actinin isoforms. Since previous studies of the interaction of calsarcin-2 with the sarcomeric  $\alpha$ -actinins have not differentiated between  $\alpha$ -actinin-2 and -3 (22), we sought to define the binding affinity of the calsarcins to  $\alpha$ -actinin-2 and -3 individually using a yeast two-hybrid assay.

Our results showed that the N-terminal region of calsarcin-2 (amino acids 1–110 and 1–90) binds to both  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3, but there is preferential binding for  $\alpha$ -actinin-2 (Figure 5A). We also examined the binding affinities of the sarcomeric  $\alpha$ -actinins for the corresponding regions of calsarcin-1 and calsarcin-3, but did not observe growth above background levels (empty vector; pGAD10), suggesting that interaction with the sarcomeric  $\alpha$ -actinins in this region is unique to calsarcin-2. However, full-length calsarcin-1 and calsarcin-3, as well as calsarcin-2, interact with the rod domain of full-length  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3. There was no preferential binding to  $\alpha$ -actinin-2 or  $\alpha$ -actinin-3 for the calsarcins at this site (Supplemental Figure 3). These results suggest that while the rod domain of sarcomeric  $\alpha$ -actinins interacts readily with the C terminus of the calsarcins, the N-terminal of calsarcin-2 shows a preference for binding with  $\alpha$ -actinin-2.

We further quantified the interactions between the N-terminal of calsarcin-2 (1 to 110) to full-length  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3 in the yeast two-hybrid system by liquid  $\beta$ -galactosidase assays (Figure 5B). The  $\beta$ -galactosidase reaction for both  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3 was greater than that of the empty vector alone, indicating that calsarcin-2 binds to both sarcomeric  $\alpha$ -actinins.



**Figure 4** Calcineurin signaling is increased in  $\alpha$ -actinin-3-deficient human muscles. Muscles of *ACTN3* 577XX humans showed a significant increase in the expression of RCAN1-4 when compared with *ACTN3* 577RR humans ( $n = 5$  and 6 females, respectively;  $P = 0.004$ ; mean  $\pm$  SEM;  $**P < 0.01$ , Mann-Whitney  $U$  test). Histogram shows the relative density of bands normalized to actin and with the RR mean set to 1 (AU). 577RR, homozygous for the *ACTN3* R allele.

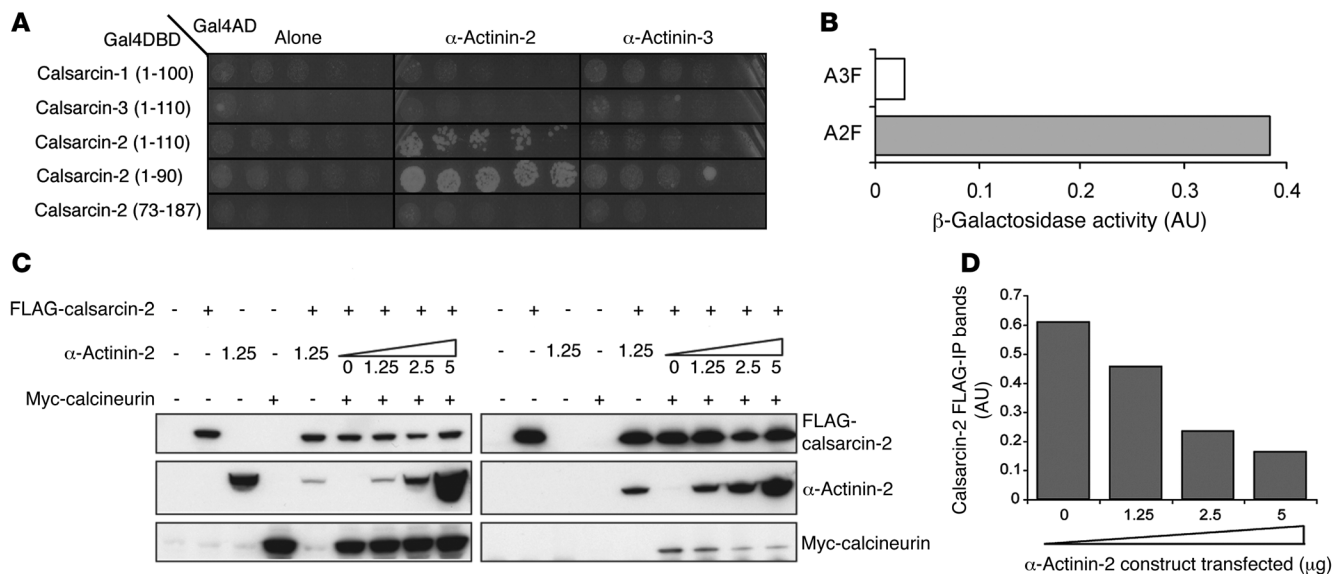
However, the interaction with full-length  $\alpha$ -actinin-2 was 13.6-fold higher compared with  $\alpha$ -actinin-3, consistent with a stronger affinity for binding of calsarcin-2 to full-length  $\alpha$ -actinin-2 in preference to  $\alpha$ -actinin-3.

$\alpha$ -Actinin-2 competes with calcineurin for binding to calsarcin-2. Since  $\alpha$ -actinin-2 is the only isoform expressed in *Actn3* KO mouse muscle and human *ACTN3* 577XX muscle and is upregulated more than 2-fold compared with WT mouse muscle (36), we hypothesized that the preferential binding of calsarcin-2 to  $\alpha$ -actinin-2 could influence the tethering and inhibition of calcineurin by calsarcin-2. To test this hypothesis, we cotransfected FLAG-tagged cal-

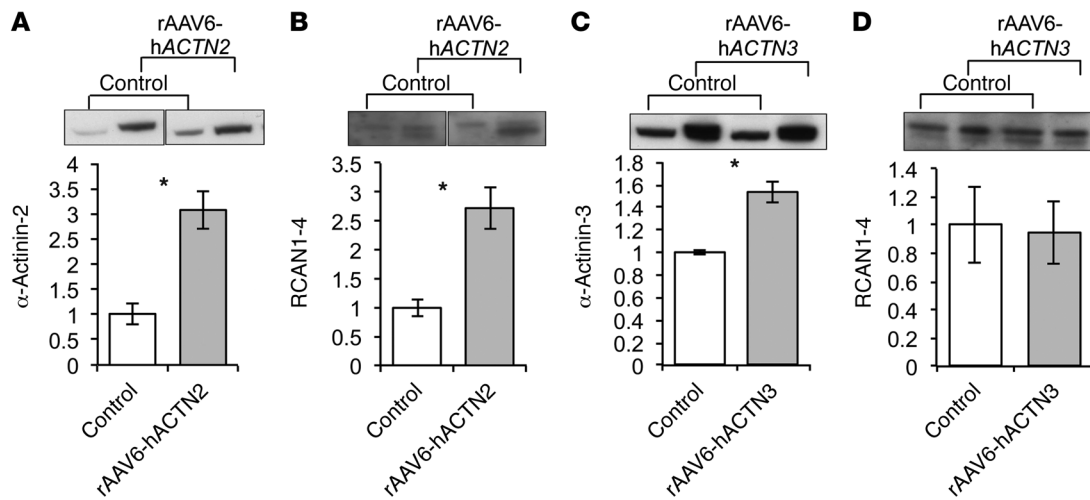
sarcin-2 and Myc-tagged calcineurin into COS-1 cells in the presence of increasing amounts of  $\alpha$ -actinin-2 (Figure 5, C and D, and Supplemental Figure 4); cell extracts were subsequently immunoprecipitated with anti-FLAG affinity gel to extract proteins that had bound to calsarcin-2. As predicted, our results show that the amount of calcineurin that coimmunoprecipitates with calsarcin-2 is inversely proportional to the level of  $\alpha$ -actinin-2 present, suggesting that increasing concentrations of  $\alpha$ -actinin-2 inhibit binding of calcineurin to calsarcin-2. This, in turn, would result in increased calcineurin activity.

*RCAN1-4 is increased in WT mouse muscle overexpressing  $\alpha$ -actinin-2.* To confirm that

differential expression of  $\alpha$ -actinin-2 directly results in enhanced calcineurin signaling in vivo, we overexpressed  $\alpha$ -actinin-2 by injecting the tibialis anterior muscles of WT mice with recombinant AAV6 vectors packaging a CMV-*bACTN2* or a CMV-*bACTN3* expression cassette. Western blot analyses showed that  $\alpha$ -actinin-2 was increased by 3.1-fold in muscles treated with CMV-*bACTN2* compared with control WT muscles injected with empty vector (Figure 6A). Consistent with our in vitro data, RCAN1-4 was also increased (2.7-fold) compared with control WT muscles, indicating that increased calcineurin signaling is associated with increased  $\alpha$ -actinin-2 expression (Figure 6B). In contrast, RCAN1-4



**Figure 5**  $\alpha$ -Actinin-2 expression influences the inhibition of calcineurin activity via preferential binding to calsarcin-2. (A) The N-terminal calsarcin-2 constructs, but not the corresponding N-terminal region on calsarcin-1 and -3, interact with  $\alpha$ -actinin-2 in yeast two-hybrid assays. In addition, there was higher-affinity binding of calsarcin-2 to  $\alpha$ -actinin-2 than to  $\alpha$ -actinin-3; this was a 13.6-fold difference when quantitated by liquid  $\beta$ -galactosidase assays (B). (C) COS-1 cells were transfected with or without vectors encoding FLAG-calsarcin-2 and Myc-calcineurin and varying amounts (0–5  $\mu$ g) of untagged  $\alpha$ -actinin-2. Whole-cell extracts (5% input) showed the expression of each protein in COS-1 cells. Immunoprecipitations using anti-FLAG M2 affinity gel demonstrated decreasing coimmunoprecipitation of calcineurin with calsarcin-2, with increasing concentrations of  $\alpha$ -actinin-2, indicative of  $\alpha$ -actinin-2 competitive inhibition of the calsarcin-2 interaction with calcineurin. (D) Densitometric quantitation of the decrease in calcineurin coimmunoprecipitation by calsarcin-2 with increasing amounts of  $\alpha$ -actinin-2 from the final four lanes for the FLAG immunoprecipitation blots (presented as a ratio of calcineurin/calsarcin-2).



**Figure 6** Overexpression of  $\alpha$ -actinin-2 in mouse muscle increases RCAN1-4 expression. (A and B) Overexpression of  $\alpha$ -actinin-2 (3.1-fold) through i.m. injection of  $1 \times 10^{11}$  vector genomes (vg) of rAAV6-CMV-*mActn2* resulted in a 2.7-fold increase in RCAN1-4 expression ( $P = 0.028$ ). (C and D) In contrast, a 1.5-fold overexpression of  $\alpha$ -actinin-3 through injection of  $1 \times 10^{11}$  vg of rAAV6-CMV-*mActn3* did not significantly alter RCAN1-4 expression in injected muscles (mean  $\pm$  SEM; \* $P < 0.05$ , Mann-Whitney *U* test;  $n = 4$  for all samples). In A and B, the lanes were run on the same gel but were noncontiguous.

expression was not significantly altered in muscles injected with CMV-*hACTN3*, despite demonstrating a 1.5-fold upregulation of  $\alpha$ -actinin-3 (Figure 6, C and D).

### Discussion

Our finding that  $\alpha$ -actinin-2 competes with calcineurin for binding with calsarcin-2 indicates that the  $\alpha$ -actinins act cooperatively with calsarcin-2 in skeletal muscle to regulate calcineurin signaling. Since  $\alpha$ -actinin-2 is the only sarcomeric  $\alpha$ -actinin isoform expressed in the *Actn3*-null muscles of KO mice and *ACTN3* 577XX humans, we propose that increased calcineurin signaling in  $\alpha$ -actinin-3-deficient muscles is the result of the liberation of calcineurin from the inhibitory influence of calsarcin-2 (summarized in Figure 7). This could account for the observed increase in calcineurin activity in *Actn3* KO muscles and, in turn, for the shift toward increased oxidative metabolism in fast muscle fibers and enhanced endurance capacity at baseline, as well as an enhanced adaptive response and greater shift in fiber type following endurance training. Furthermore, our studies suggest a novel and important role for the  $\alpha$ -actinins in the phenotypic regulation of muscle fibers and provide tantalizing insights into the specialized functions of  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3. The sarcomeric  $\alpha$ -actinins diverged 250–300 million years ago following gene duplication, and *ACTN3* has since developed restricted expression in fast muscle fibers (37). While the sarcomeric  $\alpha$ -actinins have been traditionally classified as “structural” muscle proteins, it is increasingly evident that they also have significant roles in the regulation of metabolic and signaling pathways. The relative amounts of  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3 in slow and fast muscle fibers likely influence the downstream characteristics of individual fibers in terms of their metabolic profile, contractile properties, and fiber-type specification.

### Methods

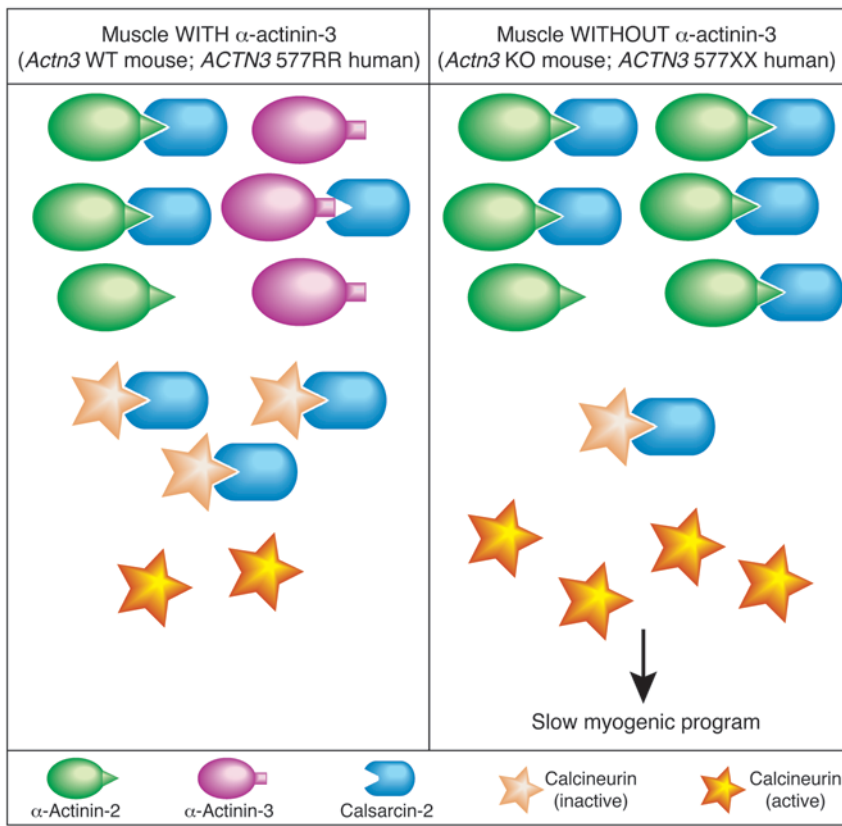
**Animals.** Mice used were of a 129X1/SvJ or C57BL/6J genetic background. The metabolic and performance phenotypes that were previously reported in 129X1/SvJ (17, 18) have been replicated and confirmed in the C57BL/6J

and C57BL/10ScSn backgrounds that were bred for ten generations (Supplemental Figure 5). All animals were fed food and water ad libitum and were maintained on a 12-hour light/12-hour dark cycle from 7:00 am to 7:00 pm.

**Immunoblotting.** Muscle sections were cut from frozen muscles, and BCA protein assays were performed on homogenates to allow equal sample loading for protein content. Samples were loaded onto precast minigels (Invitrogen) and separated by SDS-PAGE, then transferred to polyvinylidene fluoride membranes (Millipore). These were then blocked with 5% skim milk, 1x PBS, 0.1% Tween 20, probed with the indicated antibodies, and developed with ECL reagents (Amersham Biosciences). The primary antibodies used include: RCAN/DSCR1 (1:500; D6694; Sigma-Aldrich); calsarcin-2 (1:10,000; C22920; BD Biosciences); COX IV (1:2,000; 20E8; Molecular Probes); porin (1:5,000; 20B12; Molecular Probes);  $\alpha$ -actinin-2 (1:500,000; 4A3; gift from A. Beggs, Children’s Hospital Boston, Boston Massachusetts, USA); and  $\alpha$ -actinin-3 (1:1,500; 5A2; gift from A. Beggs, Children’s Hospital Boston). Following probing, PVDF membranes were stained with Coomassie brilliant blue (Sigma-Aldrich) to generate final myosin loading controls. Densitometry of immunoblot bands was performed using ImageJ image processing software (NIH).

**Calcineurin assay.** A calcineurin assay kit (207007; Calbiochem) was used according to the manufacturer’s instructions for a 30-minute reaction time on muscle homogenates. Results were normalized to total protein, as determined by BCA assay.

**Endurance performance testing and training.** Endurance performance of 6 WT and 9 KO male mice were tested by treadmill running using a previously described protocol (17) on a motorized treadmill (Columbus Instruments). Mice were tested at 10 weeks and 16 weeks of age, 3 days after the conclusion of the training period. All animals were trained and tested by an operator blinded to the mouse genotype. Mice were tested and trained between 9:00 am and 12:00 pm within the first 6 hours of their dark cycle. Endurance training by motorized treadmill running began when mice were 12 weeks of age, 5 days per week for 4 weeks, with training intensity initiated at an 8% grade at 15 meters per minute for the first 2 days and increased to a 10% grade thereafter. The running speed was increased to 20 meters per minute by day 8 of training, and the duration of training



**Figure 7**

Schematic of sarcomeric  $\alpha$ -actinin regulation of calcineurin signaling.  $\alpha$ -Actinin-2 is differentially expressed in muscles without  $\alpha$ -actinin-3. Higher-affinity binding of calsarcin-2 for  $\alpha$ -actinin-2 over  $\alpha$ -actinin-3 and competitive binding of  $\alpha$ -actinin-2 for calsarcin-2 lead to the increased release of calcineurin from calsarcin-2 inhibition in the absence of  $\alpha$ -actinin-3. We propose that increased free (active) calcineurin activates downstream signaling and the slow myogenic program in  $\alpha$ -actinin-3–deficient muscle.

was increased from 45 to 75 minutes by day 11, such that mice would run 1.5 kilometers per day during the final 2 weeks of training. Mice were closely monitored for the duration of the training and gently nudged when needed to encourage continued participation in the training to completion. To ensure the health of the mice, the body weight of each mouse was monitored at the beginning of each week, and each mouse was inspected for injuries prior to and after each round of treadmill running.

**Grip strength tests.** The forearm grip strength of mice was tested using a grip strength meter (Columbus Instruments) according to a previously described method (18).

**Enzyme assays.** The activity of lactate dehydrogenase (LDH) (Enzyme Commission [EC] number 1.1.1.27), BHAD (EC 1.1.1.35), CS (EC 4.1.3.7), SDH (EC 1.3.5.1), and HK (EC 2.7.1.1) in WT and KO muscles was determined using previously described methods (18). For all assays, the linearity with time and dependence on the amount of extract added were confirmed. All assays for a particular enzyme were performed in triplicate on the same day. Activities for each sample in the absence of substrate were measured for each sample and were subtracted from these values.

**Fiber measurements and fiber typing.** Fiber measurements and fiber typing were performed on immunostained muscle sections using MetaMorph software (Molecular Devices), as previously described (38). For immunohistochemistry analysis, transverse 8- $\mu$ m sections were cut from the midsection of frozen quadriceps muscles. All sections were blocked with AffiniPure Fab Fragment Goat Anti-mouse IgG (1:25; Jackson ImmunoResearch) for 1 hour to prevent cross-reaction with endogenous mouse antibodies. Sections were then incubated for 1 hour at room temperature with primary antibodies against myosin heavy chain 2B (used neat, BF-F3; DSHB, University of Iowa, Iowa City, Iowa, USA), 2A (used neat, SC71; DSHB, University of Iowa), and type 1 (1:300; MAB1628; Chemicon), which were prelabeled using the Zenon labeling kit (Molecular Probes)

according to the manufacturer's instructions. All samples were also costained with dystrophin (dys6-10; 1:2,000; L. Kunkel, Boston Children's Hospital, Harvard Medical School, Boston Massachusetts, USA) to define fiber borders. Images were captured using an Olympus BX50 microscope and ProgRes software (SciTech).

**cDNA cloning and sequencing.** All PCR inserts for yeast two-hybrid analysis were amplified from the human skeletal muscle cDNA and subcloned into pGBT9 and pGAD10 vectors (Clontech). Primer design for calsarcin-1 (NM\_016599), calsarcin-2 (NM\_021245), calsarcin-3 (NM\_133371),  $\alpha$ -actinin-2 (NM\_001103), and  $\alpha$ -actinin-3 (NM\_001104) were based on their respective GenBank template sequences indicated here in parentheses. Sequences of the completed constructs were confirmed by DNA sequencing.

**Yeast two-hybrid analysis.** Yeast two-hybrid analysis was performed as indicated in the Yeast Protocols Handbook (Clontech). The pGAD10  $\alpha$ -actinin-2 (aa 1-894) and  $\alpha$ -actinin-3 (aa 1-901) constructs were cotransformed with various pGBT9 constructs containing calsarcin-1 (aa 1-100), calsarcin-2 (aa 1-110), or calsarcin-3 (aa 1-110) inserts. Transformants were plated and incubated for 3 days at 30°C on minimum yeast medium lacking leucine and tryptophan. Transformants were then spotted as a dilution series on minimum yeast medium lacking leucine, tryptophan, and histidine and were incubated at 30°C. The plates were observed over 5 days to assess the growth strength.

**Liquid  $\beta$ -galactosidase assay.** The liquid  $\beta$ -galactosidase assay was carried out according to the method outlined in the Yeast Protocols Handbook (Clontech). Transformants were grown in 5 ml of selective liquid media overnight at 30°C. Two milliliters of culture was transferred to 8 ml of yeast extract/peptone/dextrose (YPD) medium and incubated for an additional 5 hours. Culture (1.5 ml of each) was spun, and the pellet was washed and then resuspended in 100  $\mu$ l of buffer 1 (23.8 mg/ml HEPES, 9 mg/ml NaCl, 0.65 mg/ml L-aspartate, 10 mg/ml BSA, 0.05% Tween 20, v/v,





pH 7.25). Cells were lysed in five rounds of freeze-thawing in liquid nitrogen followed by the addition of 700  $\mu$ l of buffer 1 containing chlorophenol red  $\beta$ -D-galactopyranoside (1.36 mg/ml), and the mixture was incubated at 30°C until the samples turned red. The reaction was stopped by the addition of 500  $\mu$ l of 3 mM ZnCl<sub>2</sub>, and absorbance was read at 570 nm. For each sample, AUs were calculated to correct for culture OD and incubation time.

**Competitive coimmunoprecipitation.** COS-1 cells were maintained in DMEM supplemented with 10% FCS. Confluent cells in a 10-cm tissue culture dish were transiently transfected with the following amount of expression plasmids: 0.5  $\mu$ g of pMT3-FLAG-calsarcin-2, 0.25  $\mu$ g of pMT3-Myc-calcineurin subunit A, and an increasing dosage (0, 1.25, 2.5, and 5  $\mu$ g) of pMT3- $\alpha$ -actinin-2, using the FuGENE 6 transfection reagent (Roche Molecular Biochemicals). Empty pMT3 vector was added to appropriate transfections to ensure that the total micrograms of plasmid transfected was the same for all wells. This resulted in the expression of calsarcin-2 recombinant protein fused with an N-terminal FLAG epitope and calcineurin subunit A fused with an N-terminal Myc epitope. Forty-eight hours after transfection, cells were harvested in 250  $\mu$ l of NP-40 lysis buffer containing 50 mM Tris (pH 8.0), 150 mM NaCl, 1% NP-40, 0.2 mM PMSF, and protease inhibitor (Sigma-Aldrich). Cells were incubated in NP-40 lysis buffer at 4°C for 30 minutes, and cell debris was removed by centrifugation. Ten microliters of this whole-cell extract was reserved as 5% protein input. The remaining 200  $\mu$ l of whole-cell extracts was precleared by incubating with Protein G Sepharose beads (GE Healthcare). The unbound portion of the whole-cell extract was immunoprecipitated for 1 hour at 4°C using Anti-FLAG M2 Affinity Gel (Sigma-Aldrich). Beads were washed with NP-40 lysis buffer prior to protein elution with SDS-DTT. The eluent was separated on SDS-PAGE and immunoblotted using anti-FLAG (Sigma-Aldrich), anti-Myc (Santa Cruz Biotechnology Inc.), or 4B3 rabbit antibody (gift from A. Beggs from Children's Hospital Boston) for calsarcin-2, calcineurin subunit A, and  $\alpha$ -actinin-2, respectively.

**Generation of rAAV6-CMV-bACTN2 and rAAV6-CMV-bACTN3 expression cassettes.** Full-length human ACTN2 and ACTN3 were cloned into recombinant AAV6 expression cassettes containing AAV2 inverted terminal repeats using standard cloning techniques. The coding sequences of  $\alpha$ -actinin-2 and  $\alpha$ -actinin-3 were cloned into the SacII site of the pAAV-CMV-SV40 plasmid. Methodologies for control vector creation, HEK293 cell transfection for rAAV6 vectors, vector purification, and titer quantification can be found in Gregorevic et al. (39).

**Intramuscular delivery of rAAV6 expression cassettes.** To transduce skeletal muscles in vivo, recombinant adenoassociated viral vectors (rAAV6 vectors) were injected directly into the tibialis anterior of WT mice using Hamilton syringes. Mice were anesthetized with isoflurane, and 1  $\times$  10<sup>11</sup> vector genomes of either rAAV-aa2 or rAAV-aa3 were injected in 30  $\mu$ l of HBSS directly into the anterior compartment of the hind limb. The contralateral

limb (used as a control) was injected with the empty vector lacking a functional gene. Four weeks after injection, mice were sacrificed via cervical dislocation, and the tibialis anterior muscles were excised and processed.

**Statistics.** Statistics were performed using SPSS software (version 15; SPSS Inc.). Because of small sample sizes, normality could not be assumed, so nonparametric statistics were applied. Kruskal-Wallis tests were applied for multiple group comparisons, followed by post-hoc Mann-Whitney *U* tests for pairwise group comparisons. Two-way ANOVA was also applied to determine whether genotype influences the training response. All results are presented as the mean  $\pm$  SEM. The  $\alpha$  level was set at *P* < 0.05.

**Study approval.** This study was approved by the Animal Care and Ethics Committee of the Children's Medical Research Institute and The Children's Hospital at Westmead and by the Sydney Children's Hospitals Network Human Research Ethics Committee. Informed consent was obtained from all individuals included in this research.

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