

RGK protein-mediated impairment of slow depolarization-dependent Ca^{2+} entry into developing myotubes

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Abbreviations: EC, excitation-contraction; ECCE, excitation-coupled Ca^{2+} entry; NFAT, nuclear factor of activated T cells; RGK, Rad-Rem-Rem2-Gem/Kir; RyR1, type 1 ryanodine-sensitive intracellular Ca^{2+} release channel; SR, sarcoplasmic reticulum

Three physiological functions have been described for the skeletal muscle 1,4-dihydropyridine receptor ($\text{Ca}_v1.1$): (1) voltage-sensor for excitation-contraction (EC) coupling, (2) L-type Ca^{2+} channel, and (3) voltage-sensor for slow depolarization-dependent Ca^{2+} entry. Members of the RGK (Rad, Rem, Rem2, Gem/Kir) family of monomeric GTP-binding proteins are potent inhibitors of the former two functions of $\text{Ca}_v1.1$. However, it is not known whether the latter function that has been attributed to $\text{Ca}_v1.1$ is subject to modulation by RGK proteins. Thus, the purpose of this study was to determine whether Rad, Gem and/or Rem inhibit the slowly developing, persistent Ca^{2+} entry that is dependent on the voltage-sensing capability of $\text{Ca}_v1.1$. As a means to investigate this question, Venus fluorescent protein-fused RGK proteins (V-Rad, V-Rem and V-Gem) were overexpressed in “normal” mouse myotubes. We observed that such overexpression of V-Rad, V-Rem or V-Gem in myotubes caused marked changes in morphology of the cells. As shown previously for YFP-Rem, both L-type current and EC coupling were also impaired greatly in myotubes expressing either V-Rad or V-Gem. The reductions in L-type current and EC coupling were paralleled by reductions in depolarization-induced Ca^{2+} entry. Our observations provide the first evidence of modulation of this enigmatic Ca^{2+} entry pathway peculiar to skeletal muscle.

Introduction

Three distinct functions for the skeletal muscle 1,4-dihydropyridine receptor ($\text{Ca}_v1.1$) have been described: (1) it serves as the voltage-sensor for the excitation-contraction (EC) coupling,^{1,3} (2) it conducts L-type Ca^{2+} current,³ and (3) it supports slow, depolarization-dependent Ca^{2+} entry that has been proposed to permeate independently of the conventional channel pore.^{4,5} Of these functions, the fundamental abilities of $\text{Ca}_v1.1$ to engage EC coupling by gating the type 1 ryanodine receptor (RyR1) and to conduct L-type current are well-documented.⁶ However, the specific route of Ca^{2+} permeation for depolarization-dependent entry remains a matter of debate.^{7,8}

Depolarization-dependent Ca^{2+} entry, also referred to as Excitation-Coupled Ca^{2+} Entry (ECCE), was first identified in *dysgenic* ($\text{Ca}_v1.1$ null) myotubes expressing a pore mutant with a minimal ability to conduct Ca^{2+} .⁴ Since this type of Ca^{2+} entry was absent in naïve *dysgenic* and *dyspedic* (RyR1 null) myotubes,^{4,5,9} the existence of a distinct Ca^{2+} channel whose gating was coupled to depolarization-dependent rearrangements in the EC coupling apparatus was proposed. Later on, the abundance of genetic

(i.e., absence in *dysgenic*, *dyspedic* and $\text{Ca}_v\beta_1$ null myotubes) and pharmacological (i.e., sensitivity to Gd^{3+} , La^{3+} , 2-aminoethyl diphenylborate, SKF 96356, nifedipine and dantrolene) characteristics shared by depolarization-dependent Ca^{2+} entry and the conventional L-type means of Ca^{2+} entry led to the idea that these pathways were, for the most part, one and the same.^{10,11} Paradoxically, these genetic and pharmacological commonalities between the Ca^{2+} entry pathways have confounded a definitive test of the latter view and, for this reason, the molecular basis of depolarization-dependent Ca^{2+} entry remains unsettled.^{7,8}

Rem, a member of the RGK (Rad, Rem, Rem2, Gem/Kir) family of small GTP-binding proteins, profoundly inhibits skeletal muscle EC coupling by interacting with the L-type Ca^{2+} channel complex. Early work showed that exogenous expression of Rem in C_2C_{12} cells almost ablated voltage-dependent SR Ca^{2+} release.¹² Subsequent work performed with cultured myotubes demonstrated that such inhibition occurs without affecting SR Ca^{2+} store content or function of RyR1, and is accompanied a substantial reduction in L-type current amplitude.¹³ Even though the inhibitory effects of Rem on the EC coupling voltage-sensor and L-type channel functions of $\text{Ca}_v1.1$ are well documented,

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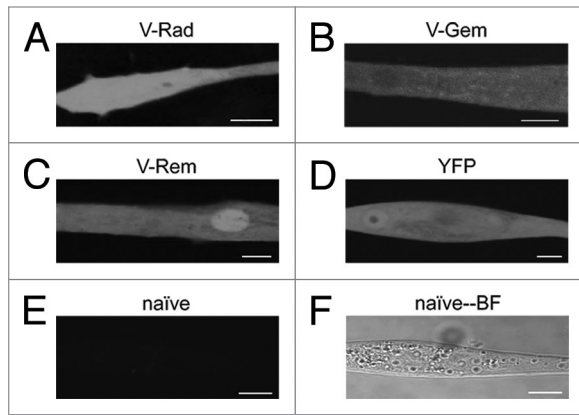


Figure 1. Exogenous expression of Venus-tagged RGK protein constructs in cultured myotubes. Confocal fluorescence images of normal mouse myotubes overexpressing V-Rad, V-Gem, V-Rem and YFP are shown as labeled in panels (A–D). Fluorescence and brightfield images of a naïve normal myotube are shown in panels (E and F), respectively. The image in (E) was acquired with identical laser settings as the image in (A). Bars-10 μ m.

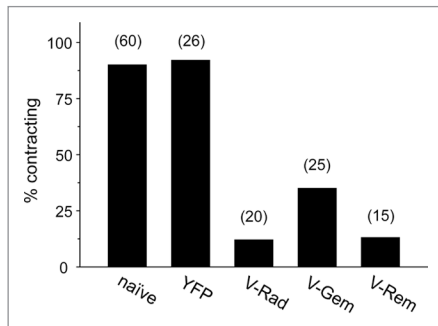


Figure 2. RGK proteins blunt EC coupling in cultured myotubes. Overexpression of a V-RGK protein construct reduces the fraction of myotubes contracting in response to a 100 V, 10 ms electrical stimulus. For each group, the total number of myotubes tested is indicated above each bar.

the ability of RGK proteins to modulate slow depolarization-dependent Ca^{2+} entry remains unknown. In this study, we have found that Rad, Gem and Rem all profoundly reduce depolarization-dependent Ca^{2+} entry in developing myotubes.

Results

RGK proteins alter myotube morphology and inhibit EC coupling

As a means to investigate the effects of the RGK proteins thought to be endogenously expressed in skeletal muscle (i.e., Rad, Gem and Rem)^{14–18} on the known functions of $Ca_v1.1$, we overexpressed one of these muscle RGK protein isoforms in cultured “normal” myotubes and confirmed expression via a fused Venus fluorescent protein tag. When viewed with confocal microscopy, V-Rad, V-Gem and V-Rem each displayed a diffuse fluorescence distribution similar to that of unfused YFP (Fig. 1A–D); very little, if any, background fluorescence

was present in naïve myotubes (Fig. 1E and F). Interestingly, the V-RGK proteins all appeared to have deleterious effects on myotube morphology. In general, V-RGK protein expressing myotubes had central protuberances and were flat and narrow compared with adjacent non-transfected myotubes.

We, with others, previously demonstrated that YFP-Rem reduces the frequency of myotube contractions without altering SR Ca^{2+} stores or RyR1 function, indicating a specific effect on $Ca_v1.1$.¹³ To test whether Rad and/or Gem are also capable of inhibiting EC coupling in myotubes, we elicited contractions with electrical field stimulation (100 V, 10 ms). Naïve normal myotubes and normal myotubes expressing non-fused YFP both readily contracted upon stimulation (54 of 60 and 24 of 26 myotubes tested, respectively; Fig. 2). In contrast, exogenous expression of V-Rad, V-Gem or V-Rem each caused a marked reduction in the fraction of contracting myotubes (3 of 25, 7 of 20, and 2 of 15 myotubes tested, respectively).

Rad, Rem and Gem inhibit L-type currents in myotubes

Although YFP-Rem is known to reduce L-type current density in myotubes,¹³ the abilities of Rad and Gem to blunt skeletal muscle L-type current have not been investigated. For this reason, we recorded L-type currents from myotubes expressing either V-Rad or V-Gem; we also deemed it necessary to assess the effects of V-Rem on the L-type current in order to compare with its effects on depolarization-dependent Ca^{2+} entry (see below). In control experiments, naïve normal myotubes produced large, slowly activating L-type currents that were not different than those observed in normal myotubes expressing YFP (-12.4 ± 0.9 pA/pF; $n = 16$ and -11.7 ± 0.7 pA/pF; $n = 16$, respectively, at +30 mV; $P > 0.05$, unpaired t test; Fig. 3A). By comparison, L-type currents were reduced by -90% , -85% and -65% in normal myotubes expressing V-Rad (-1.3 ± 0.5 pA/pF; $n = 9$; $P < 0.001$, unpaired t test; Fig. 3B), V-Gem (-1.9 ± 1.1 pA/pF; $n = 9$; $P < 0.001$, unpaired t test; Fig. 3C) or, as expected, V-Rem (-4.5 ± 1.7 pA/pF; $n = 11$; $P < 0.001$, unpaired t test; Fig. 3D). We also observed small, but significant, depolarizing shifts in $Ca_v1.1$ activation in myotubes expressing V-Rad, V-Gem and V-Rem (Table 1).

RGK proteins inhibit depolarization-dependent Ca^{2+} entry in myotubes

Previous work^{12,13} and the experiments described above have established that all RGK isoforms thought to be expressed in skeletal muscle are capable of inhibiting the ability of $Ca_v1.1$ to serve as the voltage-sensor for EC coupling and to conduct L-type Ca^{2+} current. Yet, it remains unknown whether RGK proteins affect depolarization-dependent Ca^{2+} entry into skeletal muscle. For this reason, we examined the effects of V-Rad, V-Rem and V-Gem expression on Ca^{2+} entry elicited by long, weak depolarizations. As expected, control myotubes expressing an mcherry transfection marker displayed substantial slowly activating Ca^{2+} transients in response to elevation of extracellular K^+ from 5 mM to 80 mM ($1.27 \pm 0.13 \Delta F/F$; $n = 7$; Fig. 4A–D). The transients were almost certainly indicative of Ca^{2+} entry because: (1) SR Ca^{2+} release was blocked by preincubation with 200 μ M ryanodine,^{10,11} and (2) no quantifiable transients were observed in nine naïve myotubes exposed to 100 μ M Gd^{3+} prior

to and during depolarization ($P < 0.001$, unpaired t test *vs.* control; Fig. 4D). In contrast, the entry mediated-transients were nearly ablated by overexpression of either V-Rad ($0.23 \pm 0.11 \Delta F/F$; $n = 6$; Fig. 4A), V-Gem ($0.19 \pm 0.08 \Delta F/F$; $n = 11$; Fig. 4B) or V-Rem ($0.02 \pm 0.01 \Delta F/F$; $n = 7$; all $P < 0.001$, ANOVA *vs.* control; Fig. 4C and D).

Discussion

The most significant finding of this study is that Rad, Rem and Gem each inhibit slow depolarization-dependent Ca^{2+} influx into skeletal muscle cells (Fig. 4) in such a way that parallels their abilities to dampen EC coupling and to reduce L-type Ca^{2+} current (Figs. 2 and 3, respectively). Thus, RGK proteins inhibit all three physiological processes that are dependent on the voltage-sensing ability of $Ca_v1.1$.

Depolarization-dependent Ca^{2+} entry is influenced by expression of various constituents of the EC coupling apparatus including the α_{1S} and β_{1a} subunits of $Ca_v1.1$,^{4,5,10} RyR1,^{4,5,9-11} JP-45,¹⁹ and calsequestrin1.¹⁹ In the present work, we present the first example of modulation of this mysterious mode of Ca^{2+} influx into skeletal muscle by dynamically-regulated cell signaling molecules (Fig. 4). In this regard, RGK proteins may regulate a variety of cellular processes that have been linked to depolarization-dependent Ca^{2+} entry such as maintenance of myoplasmic Ca^{2+} levels during activity,⁴ activation of transcription via NFAT translocation²⁰ and regulation of muscle strength¹⁹ and mass (see below). Moreover, altered RGK protein-mediated modulation of depolarization-dependent Ca^{2+} entry may potentially be involved in the pathophysiology of malignant hyperthermia,²¹⁻²³ central core disease,²⁰ and rippling muscle disease²⁴ since enhanced depolarization-dependent Ca^{2+} entry is characteristic of these muscle disorders.

The similar degree of inhibition between depolarization-dependent Ca^{2+} entry and L-type current represents yet another commonality between these 2 modes of Ca^{2+} flux into skeletal muscle (please see Table 4 of ref. 10). Such parallel inhibition provides further support for the idea that L-type current is, in fact, the predominant means of Ca^{2+} entry upon depolarization.^{10,11} However, the evidence presented here is still not conclusive because it is plausible that RGK proteins may stem depolarization-dependent Ca^{2+} entry through another channel which shares a common gating mechanism with $Ca_v1.1$.⁴ Even so, the mounting circumstantial evidence in favor of depolarization-dependent Ca^{2+} entry passing through the conventional $Ca_v1.1$ pore casts heavy doubt on the existence of a coupled 2-channel mechanism.

We also observed that overexpression of Rad, Rem or Gem had adverse effects on the morphology of cultured myotubes (Fig. 1A–C). Unfortunately, we were unable to quantify the impact of RGK protein overexpression on morphology because developing myotubes in a culture dish exhibit a broad range of morphological characteristics. Even so, our qualitative observations raise the possibility that RGK proteins may engage

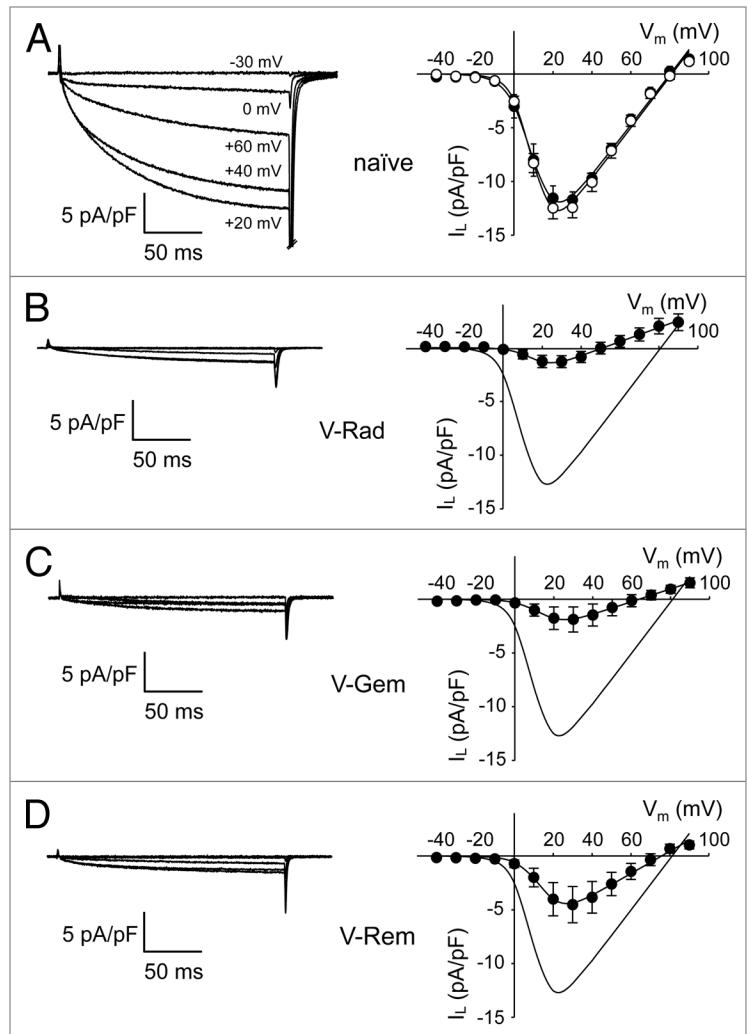


Figure 3. RGK proteins reduce L-type Ca^{2+} current amplitude in cultured myotubes. Representative recordings of skeletal muscle L-type currents elicited by 200 ms depolarizations from -50 mV to the indicated potentials are shown for a naïve normal myotube (A, left) and normal myotubes expressing V-Rad (B, left), V-Gem (C, left) or V-Rem (D, left). The peak I-V relationships corresponding to each current family are shown in the right panels. In panel (A, right), the I-V relationship for naïve normal myotubes expressing unfused YFP is also shown. In panels (B–D) the smooth line is the fit of I-V relationship for naïve myotubes. Currents were evoked at 0.1 Hz by test potentials ranging from -40 mV through $+90$ mV in 10 mV increments following a prepulse protocol (Adams et al., 1991; please see Materials and Methods). The smooth curves are plotted according to Eq. 1 with fit parameters displayed in Table 1. Throughout, error bars represent \pm SEM.

atrophic signaling through their ability to downregulate one or more of $Ca_v1.1$'s 3 functions. In this regard, *in vivo* siRNA-mediated knockdown of $Ca_v1.1$ expression is known to cause profound atrophy and fibrosis in mouse hindlimb muscle.²⁵ By the same token, one may speculate that pathological increases in RGK protein expression may have similar atrophic effects in a broad spectrum of disorders that affect skeletal muscle. In particular, Rad expression is substantially enhanced in muscle obtained from type II diabetics,^{14,16} amyotrophic lateral sclerosis patients,^{18,26} dystrophic (*mdx*) mice¹⁷ and denervated

Table 1. Conductance fit parameters

	G-V		
	G_{\max} (nS/nF)	$V_{1/2}$ (mV)	k_G (mV)
naïve	248 ± 15 (16)	10.3 ± 1.2	4.9 ± 0.4
YFP	241 ± 16 (16)	12.9 ± 1.4	4.8 ± 0.2
V-Rad	83 ± 17‡ (7)	18.9 ± 2.9*	4.6 ± 0.9
V-Gem	73 ± 23‡ (7)	17.5 ± 4.1*	6.8 ± 1.9
V-Rem	148 ± 31† (7)	16.7 ± 2.3*	5.5 ± 0.4

Data were fit by Eq. 1 and are given as mean ± SEM, with the numbers in parentheses indicating the number of myotubes tested. Since some individual current-voltage relationships for V-RGK protein expressing myotubes could not be fit with Eq. 1 because of the absence of inward Ca^{2+} current, these experiments have been omitted from the data presented in **Table 1**. However, these experiments were included in the calculation of peak current density presented in the Results section. For all the data given, the calculated average voltage error was < 5 mV. Asterisks indicate significant differences (* denotes $P < 0.05$; † denotes $P < 0.01$; ‡ denotes $P < 0.001$; † test vs. naïve control).

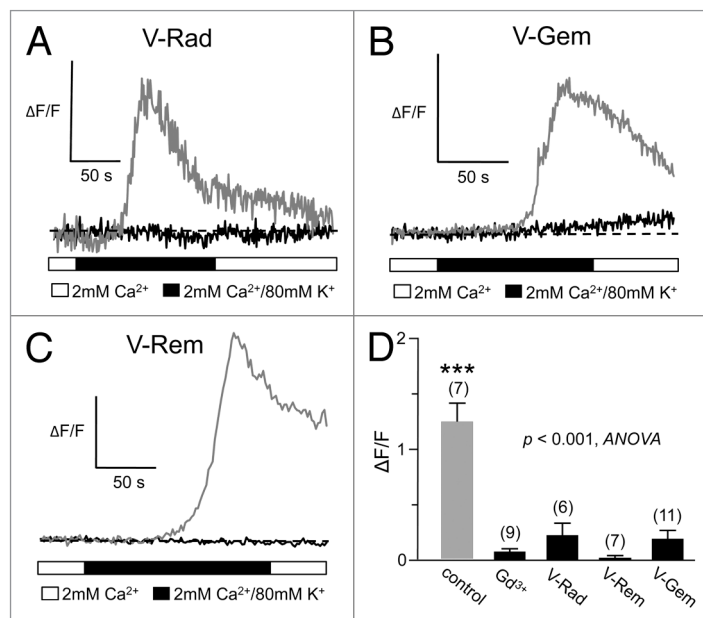


Figure 4. RGK proteins reduce slow depolarization-dependent Ca^{2+} entry into cultured myotubes. Representative Ca^{2+} transients evoked by global perfusion of 80 mM K^+ Ringer's solution for naïve normal myotubes V-Rad (A), V-Gem (B) or V-Rem (C). In panels (A–C), the fluorescence record for the RGK protein of interest (gray) is accompanied by a fluorescence record obtained from a presumably naïve (i.e., mcherry negative) normal myotube in the same field (black). The sampling rate for the experiments shown in panels (A and B) was 1.3 Hz and the sampling rate for the experiment shown in (C) was 0.45 Hz. Myotubes were exposed to 200 μM ryanodine for > 1 hour at 37 °C prior to experiments in order to block the EC coupling component of the Ca^{2+} transient. A summary is shown in panel (D). In one set of experiments, naïve myotubes were exposed to Gd^{3+} (100 μM) continuously in the bath to confirm that the Ca^{2+} signal was generated by extracellular Ca^{2+} influx, as opposed to SR Ca^{2+} release. The number of cells tested per experiment is indicated in parentheses. Asterisks indicate significant differences relative to control mcherry-expressing myotubes (***) denotes $P < 0.001$; ANOVA).

mice.¹⁸ Thus, the potential role for RGK proteins as agents of pathological muscle atrophy highlights the importance of understanding the mechanisms that these small GTP-binding proteins employ to regulate L-type channel activity in skeletal muscle.

Materials and Methods

Myotube culture and microinjection of cDNA

All procedures involving mice were approved by the University of Colorado Denver-Anschutz Medical Campus Institutional Animal Care and Use Committee. Primary cultures of normal (+/+ or +/*dysgenic*) myotubes were prepared from newborn mice as described previously.²⁷ For electrophysiological experiments, myoblasts were plated into 35 mm, plastic culture dishes (Falcon #353801) coated with ECL (Millipore #08–100). Myoblasts destined for imaging were plated into 35 mm culture dishes with laminin (Invitrogen #23017–015)-coated glass coverslip bottoms (MatTek #P35G-1.5–14-C).

Two days following differentiation, single nuclei were microinjected with a cDNA encoding a Venus fluorescent protein-RGK protein fusion construct (V-Rad, V-Gem or V-Rem, all generously provided by Drs. S.R. Ikeda and H.L. Puhl, III; 20 ng/ μl) or YFP only (Clontech; 5 ng/ μl). Myotubes to be used in Ca^{2+} imaging experiments were co-transfected with the plasmid pmCherry-C1 (Clontech; 5 ng/ μl) as a means to identify V-RGK positive cells following exposure to Fluo 3-AM dye (see below). N-Benzyl-P-toluensulfonamide (BTS; 20 μM ; Sigma-Aldrich #S949760) was added to the culture medium during microinjection in order to prevent contractions. Fluorescent myotubes were used in experiments 2 d following microinjection.

Imaging of V-RGK constructs

Live myotubes expressing YFP or V-RGK fusion constructs were examined in Rodent Ringer solution (in mM: 146 NaCl, 5

KCl, 2 CaCl₂, 1 MgCl₂, 10 HEPES, pH 7.4 with NaOH) using the confocal laser scanning microscope LSM 510 META with a Zeiss Plan-Apochromat 63X 1.4NA oil-immersion objective. In these experiments, YFP/Venus was excited with the 488-nm line of an argon laser (30-milliwatt maximum output, operated at 50% or 6.3A, attenuated to 12%), which was directed to the cell via a 488 nm dual dichroic mirror. The emitted YFP/Venus fluorescence was directed to a photomultiplier equipped with a 505 nm long-pass filter. Confocal fluorescence intensity data were recorded as the average of 8 line scans per pixel and digitized at 8 bits, with photomultiplier gain adjusted such that maximum pixel intensities were no more than ~70% saturated.

Myotube contractions

Myotube contractions were elicited by 10 ms, 100 V stimuli. The extracellular pipette contained 150 mM NaCl and the bath solution was Rodent Ringers solution.

Measurement of L-type Ca²⁺ currents

Pipettes were fabricated from borosilicate glass (~2.0 MΩ). The pipette solution consisted of (mM): 140 Cs-aspartate, 10 Cs₂-EGTA, 5 MgCl₂, and 10 HEPES, pH 7.4 with CsOH. The bath contained (mM): 145 TEA-Cl, 10 CaCl₂, 0.002 TTX, and 10 HEPES, pH 7.4 with TEA-OH. Linear capacitive and leakage currents were eliminated by -P/4 online subtraction. Electronic compensation was used to reduce the effective series resistance and the time constant for charging the linear cell capacitance. The corresponding mean values were: 2.6 ± 0.2 MΩ; 612 ± 50 μs and 311 ± 31 pF, respectively (n = 61 myotubes). Ca²⁺ currents were filtered at 2 kHz and digitized at 5 kHz. To measure macroscopic L-type current in isolation, a 1 s prepulse to -20 mV followed by a 100 ms repolarization to -50 mV was administered before the test pulse to inactivate voltage-gated Na⁺ and T-type Ca²⁺ channels (please see ref. 28). Cell capacitance was determined by integration of a transient from -80 to -70 mV using Clampex versions 9.3 and 10.3 (Molecular Devices) and was subsequently used to normalize current amplitudes (pA/pF). Current-voltage (I-V) curves were fitted using the following equation:

$$I = G_{\max} * (V - V_{\text{rev}}) / \{1 + \exp[-(V - V_{1/2}) / k_G]\}, \text{ [Eq. 1]}$$

where I is the normalized current for the test potential V, V_{rev} is the reversal potential, G_{max} is the maximum Ca²⁺ channel conductance, V_{1/2} is the half-maximal activation potential and k_G is the slope factor.

Assessment of depolarization-dependent Ca²⁺ entry

Myotubes were exposed to ryanodine (200 μM; Sigma-Aldrich #R6017) for > 1 h at 37 °C prior to experiments to eliminate voltage-dependent Ca²⁺ release from the SR. During the final 30 min of ryanodine exposure, myotubes were concurrently loaded with 5 μM Fluo 3-AM (Molecular Probes #F-1242) dissolved in Rodent Ringer's solution. Myotubes were then washed 3X in Rodent Ringer's solution with gentle agitation.

After, a 10 min de-esterification period, Fluo 3-AM-loaded cells bathed in Rodent Ringer solution were then placed on the stage of an LSM 510 META scanning laser confocal microscope and viewed with a Zeiss Plan-Neofluar 10X 0.3NA objective. Positively transfected myotubes were identified by mcherry fluorescence. Fluo 3-AM was excited with the 488-nm line of an argon laser (30-milliwatt maximum output, operated at 50% or 6.3A, attenuated to 5%). The emitted fluorescence was directed through a dual 488/543 dichroic mirror to a photomultiplier equipped with either a 505 nm long-pass filter or a 490–530 nm band-pass filter. Entry-mediated myoplasmic Ca²⁺ transients were elicited by application of high K⁺ Ringer solution (in mM: 71 NaCl, 80 KCl, 2 CaCl₂, 1 MgCl₂, 10 HEPES, pH 7.4 with NaOH) via a manually-operated, gravity-driven global perfusion system. Fluorescence amplitude data are expressed as ΔF/F, where F represents the baseline fluorescence prior to application of high K⁺ Ringer solution and ΔF represents the change in peak fluorescence during the application of high K⁺ Ringer solution. All experiments were performed at room temperature (~25 °C).

Analysis

All data are presented as mean ± SEM. Statistical comparisons were made by ANOVA (as appropriate) with P < 0.05 considered significant.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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Author contributions

C.F.R., D.B. and U.M. performed research, analyzed data, and wrote the paper. R.A.B. designed research, performed research, analyzed data, and wrote the paper.

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