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Altered Ca²⁺ homeostasis in the skeletal muscle of DJ – 1 null mice

Alexander Shtifman^{a,*}, Nan Zhong^a, Jose R. Lopez^{b,c}, Jie Shen^d, and Jin Xu^a

^a Department of Neurology, Caritas St. Elizabeth's Medical Center, Tufts University School of Medicine, 736 Cambridge St., CBR-406, Boston, MA 02135, United States

^b Department of Anesthesia, Brigham & Women's Hospital, Harvard Medical School, 20 Shattuck St., Boston, MA 02115, United States

^c Centro de Biofisica y Bioquimica, Instituto Venezolano de Investigaciones Cientificas (IVIC), Caracas, Venezuela

^d Department of Neurology, Brigham & Women's Hospital, Harvard Medical School, 77 Avenue Louis Pasteur, Boston, MA 02115, United States

Abstract

Loss-of-function mutations in DJ – 1 are associated with early-onset of Parkinson's disease. Although DJ – 1 is ubiquitously expressed, the functional pathways affected by it remain unresolved. Here we demonstrate an involvement of DJ – 1 in the regulation of Ca²⁺ homeostasis in mouse skeletal muscle. Using enzymatically dissociated flexor digitorum brevis muscle fibers from wild-type (wt) and DJ – 1 null mice, we examined the effects of DJ – 1 protein on resting, cytoplasmic [Ca²⁺] ([Ca²⁺]_i) and depolarization-evoked Ca²⁺ release in the mouse skeletal muscle. The loss of DJ – 1 resulted in a more than two-fold increase in resting [Ca²⁺]_i. While there was no alteration in the resting membrane potential, there was a significant decrease in depolarization-evoked Ca²⁺ release from the sarcoplasmic reticulum in the DJ – 1 null muscle cells. Consistent with the role of DJ – 1 in oxidative stress regulation and mitochondrial functional maintenance, treatments of DJ – 1 null muscle cells with resveratrol, a mitochondrial activator, or glutathione, a potent antioxidant, reversed the effects of the loss of DJ – 1 on Ca²⁺ homeostasis. These results provide evidence of DJ – 1's association with Ca²⁺ regulatory pathways in mouse skeletal muscle, and suggest the potential benefit of resveratrol to functionally compensate for the loss of DJ – 1.

Keywords

DJ – 1; DJ – 1 null mice; Ca²⁺; Resveratrol; Excitation-contraction coupling; Ca²⁺ release

1. Introduction

DJ – 1 is a ubiquitously expressed protein, which has been linked to the early-onset autosomal-recessive Parkinsonism (Bonifati et al., 2003; Bandopadhyay et al., 2004). DJ – 1 is present in the cytoplasm as well as intracellular organelles, such as the nucleus and mitochondria (Canet-Aviles et al., 2004; Xu et al., 2005). Although the precise function of this protein remains to be determined, mounting evidence indicates that DJ – 1 is a regulator

*Corresponding author. Tel.: +1 617 789 2678; fax: +1 617 789 3111. alex.shtifman@caritaschristi.org (A. Shtifman).

Conflict of interest

There are no actual or potential conflicts of interest.

of cellular response to oxidative stress and can directly protect cells from oxidative damage by serving as an antioxidant protein (Kinumi et al., 2004; Taira et al., 2004; Andres-Mateos et al., 2007; Dodson and Guo, 2007). Additionally, DJ-1 is a transcriptional co-activator that promotes the activities of transcriptional factors including androgen receptor and PPAR γ co-activator-1 α (PGC-1 α) (Takahashi et al., 2001; Xu et al., 2005; Zhong & Xu, 2008). It also promotes the expression of a number of mitochondrial enzymes involved in ROS removal and stabilizes a potent transcriptional regulator of mitochondrial antioxidative proteins, nuclear factor erythroid 2-related factor (Nrf2) (Clements et al., 2006).

Consistent with DJ-1's ability to regulate oxidative stress response, DJ-1 null mice are more susceptible to mitochondrial toxin MPTP-induced neuronal cell loss (Kim et al., 2005). While the loss of DJ-1 does not replicate the symptoms and neuropathology associated with Parkinson disease (PD), it leads to severe locomotor irregularities and a substantial decrease in grip strength (Chandran et al., 2008), suggesting that loss of DJ-1 may directly affect muscle function. In support of the role of DJ-1 in muscle function, a recent study reported that oxidation-mediated inactivation of DJ-1 is associated with a muscle disorder, Inclusion Body Myositis (Terracciano et al., 2008).

A precise role of DJ-1 in the muscle function and other tissues remain to be investigated. Since locomotor properties appear to be affected in the DJ-1 null animals, it is feasible that cellular processes governing muscle contraction could be significantly altered by the loss of DJ-1. Skeletal muscle contraction is regulated by the excitation-contraction (EC) coupling process. EC coupling is initiated by an action potential that activates the voltage sensors of plasmalemmal ion channels, the dihydropyridine receptors (DHPRs). Activation of DHPRs leads to an initiation of rapid Ca²⁺ release from the sarcoplasmic reticulum (SR) through the ryanodine receptor Ca²⁺ release channels (RyRs), which subsequently activate the muscle contractile apparatus (Melzer et al., 1995). Conversely, muscle relaxation is achieved by sequestration of Ca²⁺ by the SR via the SR Ca²⁺ ATPase (SERCA). Because EC coupling can be affected by the oxidative state of the cell (Anzai et al., 2000; Sun et al., 2001), a reduced ability to regulate oxidative stress due to loss of DJ-1 could potentially affect force generation as well as locomotor properties. To assess the role of DJ-1 in muscle function we evaluated the effects of the loss of DJ-1 on Ca²⁺ homeostasis and depolarization-induced Ca²⁺ release in isolated muscle cells from the DJ-1 null mice. Using Ca²⁺ selective microelectrodes to monitor the resting intracellular [Ca²⁺] ([Ca²⁺]_i) and fluorescent Ca²⁺ indicator magFluo-4 AM to monitor whole-cell Ca²⁺ fluxes, we demonstrate that the loss of DJ-1 leads to substantial elevation of resting [Ca²⁺]_i and decreased Ca²⁺ release in response to electrical stimulation. Both of these characteristics can be partially reversed by the administration of a mitochondrial functional activator, resveratrol or a potent antioxidant, glutathione. These findings advance our understanding of the role DJ-1 plays in cellular function.

2. Materials and methods

2.1. Muscle fiber preparation

DJ-1 null and wild-type (wt) mice (Goldberg et al., 2005) were used in this study following a protocol approved by the Caritas St. Elizabeth's Medical Center Institutional Animal Care and Use Committee. Homozygous DJ-1 null and wt mice were generated by breeding of heterozygous DJ-1 null animals. Age-matched mice (8–12 months old) were euthanized by pentobarbital overdose. Flexor digitorum brevis (FDB) muscles were removed, placed in a Dulbecco's Modified Eagle Medium (Invitrogen, Carlsbad, CA) solution containing 2 mg/ml collagenase A (Roche, Nutley, NJ) and incubated for 3 h at 37 °C with gentle agitation. Thereafter, muscles were removed from the enzyme-containing media and rinsed twice in DMEM. Muscle bundles were then transferred to DMEM

supplemented with 10% bovine growth serum, 1% penicillin, 1% streptomycin and 1% glutamine and gently triturated with a polished glass pipette until a significant portion were dissociated to single cells. Myofibers were plated onto ECM-coated (Sigma, St. Louis, MO) glass-bottom 24-well dishes (MatTek, Ashland, MA) and allowed to settle to the bottom of the dish over night in an incubator at 5% CO₂ and 37 °C.

2.2. Fluorescence recording

All reagents, unless otherwise indicated, were purchased from Sigma–Aldrich (St. Louis, MO). Cells were bathed in a normal Ringer solution containing in [mM]: [125]NaCl, [5]KCl, [1.2]MgSO₄, [6]glucose, [25]HEPES, [2]CaCl₂, pH 7.4. Myofibers were loaded at room temperature for 30 min in Ringer solution supplemented with Ca²⁺ indicator dye (magFluo-4AM, 5 μM (Molecular Probes, Eugene OR)). Cells were later washed several times with Ringer solution to terminate further loading and placed in a 37 °C incubator for de-esterification of the dye. To eliminate the motion artifacts due to muscle contraction, *N*-benzyl-*p*-toluene sulphonamide (BTS, 50 μM), an inhibitor of the myosin II ATPase, was added to the bathing solution.

Whole-cell fluorescence changes were detected using an IonOptix fluorescence system (IonOptix, Milton, MA) interfaced with an inverted Zeiss Axiovert 200 microscope equipped with a Neofluar 40× oil-immersion objective. Changes in fluorescence were detected by a photo multiplier tube.

Ca²⁺ transients were elicited as described previously (Shtifman et al., 2008). Briefly, supra-threshold rectangular pulses (1 ms duration) were applied through two platinum electrodes placed on opposite sides of the experimental chamber. Changes in intracellular [Ca²⁺] were characterized as changes in Fluo-4 fluorescence intensity. All experiments were conducted at room temperature (22 °C). Detected changes in fluorescence within each cell were analyzed using IonOptix analysis software (IonOptix, Milton, MA). The resulting fluorescence changes were corrected for the background fluorescence within individual cells by dividing the magnitude of change in fluorescence (ΔF) by the mean baseline fluorescence intensity (F_0) to give the $\Delta F/F$ values.

2.3. Microelectrode preparation and recording

Microelectrode recordings were performed as described previously (Christensen et al., 2004). Briefly, resting, free Ca²⁺ concentration ([Ca²⁺]_i) and the plasma membrane potential (V_m) were recorded simultaneously using double-barreled Ca²⁺-selective microelectrodes that were prepared from thin-walled borosilicate glass capillaries (WPI PB150F-4, Sarasota, FL). Prior to pulling, all capillaries were washed with HCl followed by a rinse with distilled water and then dried at 150 °C for 3 h. Capillaries were pulled to make short-taper microelectrodes with an outside tip diameter of approximately 0.6 μm. The larger barrel (1.5 mm outside diameter) was silanized by exposure to dimethyldichlorosilane vapor. Twenty-four hours later, the tip was backfilled with the neutral carrier, ETH129 (Fluka, Ronkonkoma, NY). The remainder of the barrel was backfilled with pCa7 solution 24 h later. The smaller barrel (0.84 mm outside diameter) was backfilled with 3 M KCl (tip resistance 10–15 MΩ). All Ca²⁺-selective microelectrodes were calibrated at 22 °C in solutions of known [Ca²⁺] before and after the Ca²⁺ measurements (Alvarez-Leefmans et al., 1981). To better mimic the intracellular ionic conditions all calibration solutions were supplemented with 1 mM Mg²⁺. Only those Ca²⁺ microelectrodes that provided a Nernstian response between pCa3 and pCa7 (29.5 mV/pCa unit at 22 °C) were used in this study. The potentials from the 3 M KCl barrel (V_m) and the Ca²⁺ barrel (V_{CaE}) were recorded via high impedance amplifier (WPI FD-223, Sarasota, FL). The V_m potential was subtracted electronically from V_{CaE} potential, to produce a differential Ca²⁺-specific potential (V_{Ca})

that represents the $[Ca^{2+}]_i$ concentration. V_m and V_{Ca} were filtered (30–50 KHz) to improve the signal-to-noise ratio. Recordings were conducted at room temperature (21 °C).

2.4. Western blotting

The hamstring muscles were dissected and snap-frozen in liquid nitrogen. The tissues were ground into powder and lysed in RIPA buffer with protease inhibitor. Equal amount of proteins were resolved in either 6% for RyR or 4–20% for SERCA1 gels Tris–Glycine gels (Invitrogen, Carlsbad, CA). α -Actin was used as loading control. Antibodies for each protein were from Abcam, Inc. (Cambridge, MA).

2.5. Statistics

Statistical analysis was performed using Student's *t*-test for two independent populations. Differences were considered to be statistically significant at $p < 0.05$. All data are presented as mean $\pm$ SEM.

3. Results

3.1. Elevated resting $[Ca^{2+}]_i$ in DJ – 1 null muscle cells

To determine if the loss of DJ – 1 resulted in alteration in resting, myoplasmic Ca^{2+} concentration ($[Ca^{2+}]_i$), we performed simultaneous microelectrode measurements of $[Ca^{2+}]_i$ and membrane potential (V_m) in FDB muscle fibers from wt and DJ – 1 null mice. This method is well suited for accurate and consistent quantitative measurements of sub-sarcolemmal $[Ca^{2+}]_i$ in resting cells (Tsien and Rink, 1981) (Alvarez-Leefmans et al., 1981). Representative records of Ca^{2+} potential obtained from a wt and a DJ – 1 null myofiber are presented in Fig. 1A. As seen from these traces, DJ – 1 null muscle cell exhibited a less negative Ca^{2+} potential than the wt cells, corresponding to greater $[Ca^{2+}]_i$ after electrode calibration. Resting $[Ca^{2+}]_i$ in DJ – 1 null muscle fibers was 282.9 ± 8.1 nM ($n = 16$), which was more than two-fold greater than that observed in wt cells (113.3 ± 2.7 nM, $n = 12$; $p < 0.001$, *t*-test) (Fig. 1B). There was no difference in V_m between wt (81.9 ± 0.6 mV, $n = 12$) and DJ – 1 null cells (82.4 ± 0.6 mV, $n = 16$, $p > 0.5$ *t*-test) (Fig. 1C).

3.2. Effects of the loss of DJ – 1 on Ca^{2+} release

Depolarization-induced Ca^{2+} fluxes were monitored with a low-affinity Ca^{2+} indicator, magFluo-4AM, which is an appropriate indicator for examining kinetic parameters of rapid changes in $[Ca^{2+}]_i$. To assess the effect of DJ – 1 loss on SR Ca^{2+} release, individual, enzymatically dissociated FDB muscle fibers prepared from the wt and DJ – 1 null animals were field stimulated with single pulses (1 ms duration). Muscle fibers readily exhibited Ca^{2+} release activity in response to each stimulus. Fig. 2A presents the averaged Ca^{2+} transients from wt (gray) and DJ – 1 null muscle fibers (black). As demonstrated in Fig. 2B Ca^{2+} transients from the DJ – 1 null muscle fibers exhibited smaller peak amplitudes ($0.17 \pm 0.006 \Delta F/F$, $n = 22$) than those from the wt fibers ($0.23 \pm 0.02 \Delta F/F$, $n = 17$, $p < 0.05$).

The time course of decay in FDB muscle fibers from both animals could be well described by a single exponential function. Fig. 2C presents the comparison of decay time constants obtained from DJ – 1 null and wt Ca^{2+} transients. In wt fibers the decay time constant was 12.37 ± 2.6 ms ($n = 22$), whereas in DJ – 1 null fibers it was 10.2 ± 0.82 ms ($n = 17$, $p > 0.05$). Thus, it appears that Ca^{2+} clearance mechanisms were not significantly affected by the loss of DJ – 1.

3.3. Expression levels of RyR and SERCA1 in DJ – 1 null muscle

Given that DJ – 1 is a transcriptional regulator, it is feasible that some of the changes in resting $[Ca^{2+}]_i$ and Ca^{2+} release could be due to altered expression levels of key proteins involved in the maintenance of Ca^{2+} homeostasis. Immunoblots of muscle homogenates from the wt and DJ – 1 null hamstring muscles were used to quantify the expression levels of RyR1, and SERCA1 (Fig. 3A). A quantitative analysis confirmed that there were no significant alterations in the steady-state expression levels of these proteins in the DJ – 1 null muscle (Fig. 3B and C). These results argue against the possibility that altered Ca^{2+} handling observed in the DJ – 1 null muscle cells was caused by altered expression of the key proteins involved in excitation–contraction coupling.

3.4. Reversal of $[Ca^{2+}]_i$ and Ca^{2+} release alterations by resveratrol in DJ – 1 null muscle cells

Resveratrol (*trans*-3,5,4'-trihydroxy-*trans*-stilbene) is a phytoalexin produced by plants, and found in the skin of red grapes. It has been demonstrated to exert multifaceted anti-oxidant and anti-inflammatory effects in various disease models. Resveratrol treatment mimics the effect of caloric restriction and delay aging by activating sirtuin family of NAD^+ -dependent deacetylases (Wood et al., 2004; Baur et al., 2006). In addition, resveratrol stimulates mitochondrial biogenesis by activating an energy sensor AMP-activated kinase and increasing the expression of PGC-1 α (Baur et al., 2006; Dasgupta and Milbrandt, 2007). Given recent results that show DJ – 1-mediated enhancement of PGC-1 α transcriptional activity (Zhong and Xu, 2008) and the role of DJ – 1 in mitochondrial function, we sought to investigate whether treatment of cells with resveratrol could reverse some of the effects caused by loss of DJ – 1. For this purpose, dissociated DJ – 1 null and wt muscle fibers were treated for 18 h with resveratrol (10 μ M or 50 μ M). Microelectrode recording of resting $[Ca^{2+}]_i$ revealed that resveratrol treatment resulted in a dose-dependent reduction of resting $[Ca^{2+}]_i$ in DJ – 1 null muscle cells, with no observable effect on resting $[Ca^{2+}]_i$ in the wt cells (Fig. 4A). It must be noted that treatment with higher dosage of resveratrol (50 μ M) reversed $[Ca^{2+}]_i$ in DJ – 1 null cells to a near normal, physiological level (137.4 ± 3.8 nM). In addition to reversal of $[Ca^{2+}]_i$, resveratrol treatment also resulted in an increase in the peak amplitude of action potential-elicited Ca^{2+} release in the DJ – 1 null muscle fibers (Fig. 4B), without affecting peak amplitudes of Ca^{2+} transients in the wt cells (not shown).

To determine whether alterations in Ca^{2+} handling could be reversed by treatments with other antioxidants, experiments were conducted where cells were treated with glutathione (GSH), a potent antioxidant, prior to measuring resting $[Ca^{2+}]_i$. An overnight treatment with GSH (1 mM) did not have an effect on resting $[Ca^{2+}]_i$ in either wt or DJ – 1 null muscle fibers (not shown). However, increasing of GSH concentration to 5 mM lead to a substantial decrease in $[Ca^{2+}]_i$ in the DJ – 1 deficient cells (153.3 ± 3.7 nM) without affecting $[Ca^{2+}]_i$ in wt cells (Fig. 4C). Together, these results demonstrate that antioxidant treatments can reverse changes in Ca^{2+} handling in DJ – 1 deficient muscle cells.

4. Discussion

This study is the first report that describes a functional link between DJ – 1 and maintenance of Ca^{2+} homeostasis. Using primary skeletal muscle cells from DJ – 1 null mice, we have found that the loss of DJ – 1 leads to a substantial increase in resting cytoplasmic $[Ca^{2+}]_i$ and that DJ – 1 null cells consistently exhibit reduced Ca^{2+} release from the sarcoplasmic reticulum in response to electrical stimulation. The alteration in Ca^{2+} handling could be reversed by resveratrol or GSH treatments. These results show an important role for DJ – 1 in muscle function and suggest that dysregulation of Ca^{2+} homeostasis could be a seminal event in the pathogenic mechanism linked to a loss of DJ – 1.

DJ – 1 is recognized as a protein linked to selected cases of early-onset Parkinson disease (Bonifati et al., 2003). Given its ubiquitous expression, it is important to characterize the physiologic function of DJ – 1, and the consequences of its loss or inactivation. Our current study provides evidence that loss of DJ – 1 leads to the impairment of at least two physiologic processes in skeletal muscle cells, the maintenance of myoplasmic $[Ca^{2+}]_i$ and stimulated Ca^{2+} release from the sarcoplasmic reticulum. These results are consistent with motor deficits reported in another strain of DJ – 1-null mice (Chandran et al., 2008), and the possible involvement of DJ – 1 inactivation in a skeletal muscle disorder, Inclusion Body Myositis (Terracciano et al., 2008), where similar alterations in Ca^{2+} handling have been reported (Moussa et al., 2006; Shtifman et al., 2008).

One of the main findings of this study is that a loss of DJ – 1 leads to a substantial elevation of $[Ca^{2+}]_i$ in muscle cells. Although caused by different factors, similar alterations in Ca^{2+} handling were been reported in other skeletal muscle disorders, such as malignant hyperthermia (Lopez et al., 1986, 1992; Yang et al., 2006) and Inclusion Body Myositis (Moussa et al., 2006; Shtifman et al., 2008). Since Ca^{2+} ions are integral to the proper function of the cells, substantial alterations in basal Ca^{2+} concentration reported here could either trigger or exacerbate deleterious effects in a myriad of cellular processes. Another aspect of Ca^{2+} handling that was altered in the DJ – 1 deficient cells is depolarization-induced Ca^{2+} release. It will be of interest to determine whether an analogous disruption in Ca^{2+} homeostasis by a loss or inactivation DJ – 1 also occurs in the nervous system. Interestingly, a recent study linked a deficiency in another protein genetically associated with autosomal-recessive PD, PINK1, to mitochondrial Ca^{2+} overload in neural cells (Gandhi et al., 2009). Another study reported that interplay between α -synuclein, cytosolic dopamine metabolite and intracellular Ca^{2+} contributes to the selective neurotoxicity in dopaminergic neurons (Mosharov et al., 2009). Reduction of intracellular Ca^{2+} by channel blocking or buffering significantly reduced the levels of cytosolic dopamine, a key factor in the toxic mix. Based on our current finding and those by others, it is evident that a better understanding of the interplay between the loss of DJ – 1 and altered Ca^{2+} handling in dopaminergic neuron could lead to a better understanding of PD pathogenesis.

There is considerable evidence that DJ – 1 is involved in the oxidative stress response and in maintenance of mitochondrial function (Dodson and Guo, 2007). Thus, a decrease in functional DJ – 1 or its elimination could lead to an enhanced generation of ROS by the mitochondria. It remains to be determined whether changes in $[Ca^{2+}]_i$ and Ca^{2+} release in the DJ – 1 null cells could have been mediated by oxidative stress. All isoforms of RyRs contain regulatory thiols that are sensitive to RedOx modifications. Once these channels are oxidized, their responsiveness to modulators is altered and they become hyperactive. Hyperactivity of RyR under resting conditions (“ Ca^{2+} leak”) can account for the observed changes in cytoplasmic $[Ca^{2+}]_i$ in the DJ – 1 null cells. In addition, chronic oxidation of RyRs can also alter Ca^{2+} release parameters during EC coupling. The possibility that altered Ca^{2+} handling in the DJ – 1 null cells was potentially mediated by ROS is further supported by the rescue effect of resveratrol and glutathione. While tripeptide glutathione acts as a strong reducing agent to antagonize ROS directly, resveratrol stimulates the activity and expression of PGC-1 α , a key regulator of mitochondrial biogenesis and ROS removal (Wu et al., 1999; Baur et al., 2006; St-Pierre et al., 2006; Dasgupta and Milbrandt, 2007). Interestingly, DJ – 1 may prevent the ROS accumulation by resembling the effects of both glutathione and resveratrol. The DJ – 1 protein, abundant in cysteine and methionine, has been shown to act as antioxidant molecule to directly remove H_2O_2 (Kinumi et al., 2004; Taira et al., 2004; Andres-Mateos et al., 2007). Furthermore, oxidized DJ – 1 accumulates in the Alzheimer’s disease and PD brains (Choi et al., 2006). In addition to removing free radical directly, DJ – 1 is also a transcriptional regulator (Takahashi et al., 2001; Niki et al., 2003; Xu et al., 2005). We have recently reported a synergy between DJ – 1 and PGC-1 α in

the transcriptional regulation of the human MnSOD (Zhong and Xu, 2008). Thus, similar to resveratrol, DJ – 1 may enhance PGC-1 α activity and lead to the expression of genes involved in mitochondrial biogenesis and ROS removal. It is notable that resveratrol reversed the aberrant Ca²⁺ handling in DJ – 1 null cells without affecting the Ca²⁺ homeostasis in muscle cells from the wild-type mice. Given the potent anti-oxidative stress effect of DJ – 1, the resveratrol treatment, especially at the doses used in this study (10 and 50 μ M), may not generate additional benefit in the wild-type cells. Since increased oxidation of DJ – 1 has been associated with normal aging (Choi et al., 2006; Meulener et al., 2006), and may lead to attenuated DJ – 1 function (Zhong and Xu, 2008), it will be of interest to investigate whether resveratrol may affect the Ca²⁺ signaling in cells from wild-type, aged animals.

The aberrant mitochondrial function resulting from the loss of DJ – 1 and presumed subsequent reduction of ATP production could underlie the alterations in depolarization-induced Ca²⁺ release reported here. It has been well described that function of main components of the depolarization-induced Ca²⁺ release, the DHPRs and the RyRs, are strongly dependent on cytoplasmic ATP concentration. For instance, RyR1 is strongly stimulated by ATP binding to the cytoplasmic regulatory site and weakly inhibited by ATP hydrolysis products ADP and AMP (Meissner et al., 1986; Laver et al., 2001). Additionally, voltage sensor-dependent Ca²⁺ release has been shown to decrease by lowering of intracellular ATP concentration (Owen et al., 1996; Blazev and Lamb, 1999a, b; Dutka and Lamb, 2004). However, it must be noted that in addition to ATP, other metabolic and cellular factors may be involved in reducing Ca²⁺ release in DJ – 1 deficient cells.

Our results indicate that the steady-state expression of RyRs and SERCA, which are essential for maintenance of myoplasmic [Ca²⁺]_i and regulation of Ca²⁺ release, are not directly affected by the loss of DJ – 1. Nevertheless, abnormal transcriptional regulation due to the loss of DJ – 1 may still be indirectly involved in the Ca²⁺ dysregulation, given that DJ – 1 is involved in the expression or activity of a number of key proteins essential to mitochondrial function and oxidative stress response. And it is possible that these gene products could modulate the function of either RyRs or SERCA.

In summary, the present data demonstrate considerable alterations in SR Ca²⁺ release and resting [Ca²⁺]_i arising from the loss of DJ – 1. These results contribute to our understanding of the physiological role of DJ – 1 and provide new insights into molecular mechanisms underlying DJ – 1-associated pathology.

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Fig. 1.

Increased $[Ca^{2+}]_i$ in DJ-1 null muscle fibers. Recording of intracellular Ca^{2+} concentration ($[Ca^{2+}]_i$) and resting membrane potential (V_m) with double-barreled microelectrodes in enzymatically dissociated flexor digitorum muscle fibers. A. Representative Ca^{2+} potential records (V_{Ca}) from a wt (gray) and a DJ-1 null (black) muscle fibers. The initial portion of each trace was obtained prior to impalement of the cells. Penetration and removal of microelectrode was accompanied by immediate downward and upward deflection, respectively. B. Resting $[Ca^{2+}]_i$ in wt ($n = 12$, gray), DJ-1 null ($n = 16$, black) muscle cells determined from V_{Ca} after calibration of recording microelectrodes. DJ-1 null cells exhibited a greater than two-fold elevation in $[Ca^{2+}]_i$. C. Resting V_m recorded from the same cells in panel B. Cells from DJ-1 null animals exhibited V_m similar to those from the wt animals ($p > 0.5$, t -test). Muscle fibers were prepared from 8 to 12 months old mice. Asterisk indicates statistical significance ($p < 0.001$, t -test).

Fig. 2. Reduced Ca^{2+} release in DJ-1 null muscle fibers. A. Mean Ca^{2+} transient from wt (gray) ($n = 17$) and DJ-1 null (black) ($n = 22$) FDB muscle fibers measured in response to 1 ms action potential stimuli. B–C. Peak amplitude ($\Delta F/F$) and decay time constant (τ , ms) of Ca^{2+} transients from wt and DJ-1 null muscle fibers from the same cells presented in panel A, respectively. Asterisk indicates statistical significance ($p < 0.05$).

Fig. 3.

DJ-1 does not affect the protein expression of RyR1 and SERCA1. A. Immunoblots of RyR1 and SERCA1 protein levels from hamstring muscle of wt and DJ-1 null mice. The membrane was reprobbed for α -actin, which was used as loading control. Bottom panel demonstrates that there was no detectable DJ-1 protein in the DJ-1 null muscle. B-C. Quantitative analysis confirmed that there were no significant alterations in the steady-state expression levels of RyR1 (B) and SERCA1 (C) in the DJ-1 null muscle.

Fig. 4.

Resveratrol treatment reverses $[Ca^{2+}]_i$ changes and improves Ca^{2+} release in DJ-1 null muscle cells. A. Resting $[Ca^{2+}]_i$ measurements performed with Ca^{2+} selective microelectrodes in wt and DJ-1 null muscle cells treated for 18 h with 10 μ M ($n_{wt} = 5$, $n_{DJ-1 \text{ null}} = 17$) or 50 μ M ($n_{wt} = 5$, $n_{DJ-1 \text{ null}} = 8$) resveratrol (RSV). Resveratrol treatment did not have an effect on $[Ca^{2+}]_i$ in wt cells, but lowered $[Ca^{2+}]_i$ in DJ-1 null cells in dose-dependent manner. B. Mean Fluo-4 fluorescence transient from untreated DJ-1 null muscle fibers (left) ($n = 15$) and DJ-1 null muscle cells treated for 12 h with 50 μ M resveratrol (right) ($n = 16$). Ca^{2+} transients were elicited by 1 ms action potential stimuli. Resveratrol treatment increased peak amplitudes of Ca^{2+} transients in DJ-1 null cells. C. Resting $[Ca^{2+}]_i$ measurements performed with Ca^{2+} selective microelectrodes in wt and DJ-1 null muscle cells treated for 18 h with 5 mM glutathione (GSH). GSH treatment did not have an effect on $[Ca^{2+}]_i$ in wt cells ($n = 4$), but lowered $[Ca^{2+}]_i$ in DJ-1 null cells ($n_{DJ-1 \text{ null}} = 6$, $n_{DJ-1 \text{ null}+GSH} = 10$). Asterisk indicates statistical significance ($p < 0.05$).