



# LED therapy plus idebenone treatment targeting calcium and mitochondrial signaling pathways in dystrophic muscle cells

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## Abstract

Intracellular calcium dysregulation, oxidative stress, and mitochondrial dysfunction are some of the main pathway contributors towards disease progression in Duchenne muscular dystrophy (DMD). This study is aimed at investigating the effects of light emitting diode therapy (LEDT) and idebenone antioxidant treatment, applied alone or together in dystrophic primary muscle cells from *mdx* mice, the experimental model of DMD. *Mdx* primary muscle cells were submitted to LEDT and idebenone treatment and evaluated for cytotoxic effects and calcium and mitochondrial signaling pathways. LEDT and idebenone treatment showed no cytotoxic effects on the dystrophic muscle cells. Regarding the calcium pathways, after LEDT and idebenone treatment, a significant reduction in intracellular calcium content, calpain-1, caldesmon, and sarcolipin levels, was observed. In addition, a significant reduction in oxidative stress level markers, such as H<sub>2</sub>O<sub>2</sub>, and 4-HNE levels, was observed. Regarding mitochondrial signaling pathways, a significant increase in oxidative capacity (by OCR and OXPHOS levels) was observed. In addition, the PGC-1 $\alpha$ , SIRT-1, and PPAR $\delta$  levels were significantly higher in the LEDT plus idebenone treated-dystrophic muscle cells. Together, the findings suggest that LEDT and idebenone treatment, alone or in conjunction, can modulate the calcium and mitochondrial signaling pathways, such as SLN, SERCA 1, and PGC-1 $\alpha$ , contributing towards the improvement of the dystrophic phenotype in *mdx* muscle cells. In addition, data from the LEDT plus idebenone treatment showed slightly better results than those of each separate treatment in terms of SLN, OXPHOS, and SIRT-1.

**Keywords** Dystrophic muscle cells · Photobiomodulation · Calcium signaling · Oxidative stress · Mitochondrial parameters

## Introduction

Duchenne muscular dystrophy (DMD) is a genetic X-linked disease characterized by mutations of dystrophin protein, which leads to progressive muscle degeneration with a consequent loss of ambulation in adolescence and to death, at approximately 30 years of age, usually due to cardio-respiratory failure (Emery 2002; Landfeldt et al. 2018). There is currently no cure for DMD, and glucocorticoids are the main drug treatment used by dystrophic patients (Gloss et al.

2016). However, glucocorticoid continued use causes several adverse effects (Moxley et al. 2010), making it necessary to find new therapeutic approaches.

In addition to the primary genetic defect of DMD, the absence of dystrophin protein, several studies suggest that abnormal intracellular calcium concentration [Ca<sup>2+</sup>]<sub>i</sub>, reactive oxygen species (ROS), and mitochondrial dysfunction play a crucial role in the pathophysiology of this disease (Moore et al. 2020; González-Jamett et al. 2022).

Previous studies suggest that the sustained elevation of [Ca<sup>2+</sup>]<sub>i</sub> underlies muscle pathology and dysfunction in DMD. Research showed that the increase of [Ca<sup>2+</sup>]<sub>i</sub> enhances the expression and activity of calpains, the Ca<sup>2+</sup>-dependent proteases in dystrophic muscles (Spencer et al. 1995; Hussain et al. 2000; Sundaram et al. 2006; Voit et al. 2017). Furthermore, it was also observed that increased [Ca<sup>2+</sup>]<sub>i</sub> influences mitochondrial Ca<sup>2+</sup> uptake, which in turn

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leads to altered metabolism and increased ROS production (Mareedu et al. 2021).

Mitochondria are the main source of ROS in dystrophic muscles (Moore et al. 2020). A recent study showed abnormal mitochondrial structure and reduced mitochondrial function in the dystrophic muscle of *mdx* mice (the experimental model of DMD), prior to the onset of muscle dystrophic fiber damage, suggesting an early mitochondrial role in the pathophysiology of DMD (Moore et al. 2020).

In search of alternative treatments for DMD, previous studies have pointed to the use of photobiomodulation (PBM) as a potential therapy (Silva et al. 2015; Macedo et al. 2020). PBM has presented several benefits in dystrophic muscles, such as the mitigation of inflammatory response and oxidative stress (Silva et al. 2015; Macedo et al. 2020). The effects of PBM result in the absorption of the light through the mitochondria, leading to an increase in membrane potential, the electron transport, and the oxygen consumption (Karu 2008; Freitas and Hamblin 2016). In this study, seeking to increase the therapeutic efficiency of PBM, we associated the PBM therapy (PBMT) with idebenone, a potent antioxidant and electron carrier. Idebenone is a synthetic short-chain benzoquinone, which improves mitochondrial function, restores ATP production, and reduces ROS (Buyse et al. 2015). In addition, recently, our research group showed that idebenone protected the dystrophic muscle cells by reducing the  $[Ca^{2+}]_i$ , oxidative stress, and inflammatory process (Valduga et al. 2023).

Thus, in the present study, we evaluated the LED therapy plus idebenone treatment effects in dystrophic muscle cells from *mdx* mice, focusing on the calcium and mitochondrial signaling pathways.

## Materials and methods

### Mdx primary muscle cells

The development of primary skeletal muscle cell cultures from *mdx* (C57BL/10-Dmdmdx/PasUnib) mice was based on protocol previously described (Rando and Blau 1994; Mizobuti et al. 2019). All the experiments were performed under an approved protocol of the Committee on the Ethics of Animal Experiments of UNICAMP (#5603-1/2020) and were in accordance with the Brazilian College for Animal Experimentation guidelines.

### Experimental design (see Fig. 1)

#### LEDT

LEDT at 850 nm was applied on *mdx* primary muscle cells, using a ThorLabs Mounted High-Power equipment, and the parameters used (Table 1) are based on previous study (Rocha et al. 2022). The LEDT was applied to muscle cells at a perpendicular angle, by a single application, and irradiation was performed by one point at the center of each culture well. The cells were irradiated inside a laminar flow in a dark room without radiation. *Mdx* muscle cells not irradiated were used as control.

#### Idebenone treatment

The first experiment tried to establish the best dose of idebenone for *mdx* muscle cell proliferation by MTT assay. *Mdx* muscle cells received different doses of idebenone (0.5, 0.25, 0.12, 0.06, and 0.03  $\mu$ M) diluted in 0.5% carboxymethylcellulose sodium salt (CMC, Fluka,

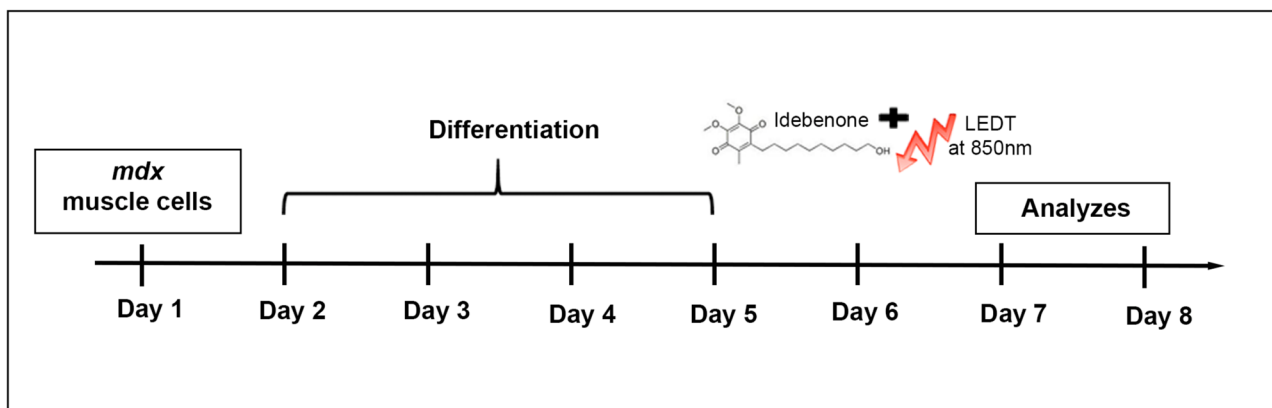


Fig. 1 Timescale of LEDT and idebenone treatment in *mdx* muscle cells

**Table 1** Specifications of LEDT parameters

Device information		
Manufacturer	ThorLabs Mounted High-Power equipment	
Irradiation parameters	96 weels	6 weels
Output power (mW)	110	38
Spot size (cm <sup>2</sup> )	0.32	9.5
Power density (w/cm <sup>2</sup> )	343	4
Energy density (J/cm <sup>2</sup> )	1.5	0.05
Time per point (s)	4.5	13.1
Wavelength (nm)	850	
Beam shape	Circular	
Operating mode	Continuous	
Number and frequency of treatment sessions	1	
Application technique	Applied directed to the cells through the bottom of each well	

Buchs, Switzerland) in water and were evaluated after 24 and 48 h of treatment. Subsequently, the *mdx* muscle cells received only two doses (0.06 and 0.03  $\mu\text{M}$ ; selected based on the results of MTT assay and IC50) and were evaluated after 24 and 48 h of treatment for cell viability (Neutral Red assay),  $[\text{Ca}^{2+}]_i$ , mitochondrial superoxide ( $\text{O}_2^{\bullet-}$ ), and  $\text{H}_2\text{O}_2$  production analyses. Lastly, the other analyses were performed in *mdx* muscle cells treated with idebenone, after 48 h of treatment, at the chosen dose (0.06  $\mu\text{M}$ ). For all experiments, the untreated *mdx* muscle cells were used as controls.

### Cell viability and proliferation

**MTT analysis** This colorimetric analysis was performed for quantification of cell metabolic activity. The protocol of this assay was previously described by Macedo et al. (2015) and Mizobuti et al. (2019). Briefly, after the MTT incubation phase, the crystals were dissolved with isopropanol acid, and the amount of formazan product was measured by a spectrophotometer (Synergy H1, Hybrid Reader, Biotek Instruments, Winooski, VT, USA) at 570 nm with a 655-nm reference wavelength. In addition, the IC50 was calculated based on the MTT data.

**Neutral Red** This colorimetric analysis was performed to quantify membrane permeability and lysosomal cell activity in response to different doses of idebenone (0.06 and 0.03  $\mu\text{M}$ ). The protocol of this assay was previously described by Borenfreund and Puerner (1985). Briefly, primary muscle cells were washed in PBS once, and NR medium was added (250  $\mu\text{l}$ ), and the cells were incubated (37 °C) for 3 h. After this period, the NR medium was removed, the muscle cells were washed in PBS, and desorption solution was added. A shaker was used to extract the NR from the cells, and the absorbance was measured in a spectrophotometer (Synergy

H1, Hybrid Reader, Biotek Instruments, Winooski, VT, USA) at 540 nm.

### Intracellular calcium content

$[\text{Ca}^{2+}]_i$  analysis was performed by Fluo-4 dye (Molecular Probes, Oregon, USA). The protocol was previously described (Macedo et al. 2015; Mizobuti et al. 2019). Briefly, after incubation with Fluo-4 AM, the calcium-sensitive dye intensities were observed on a fluorescent inverted microscope (Nikon, Eclipse TS100/TS100F) for qualitative data. Quantitative measurements were investigated using a spectrophotometer (Synergy H1, Hybrid Reader, Biotek Instruments, Winooski, VT, USA) at excitation and emission wavelengths of 494 and 516 nm, respectively.

### $\text{H}_2\text{O}_2$ production

$\text{H}_2\text{O}_2$  levels were determined by Amplex® Red assay kit (Molecular Probes, Life Technologies, California, EUA) according to the manufacturer's instructions. The Amplex UltraRed reagent (50  $\mu\text{M}$ ) and HRP (0.1 U/ml) were added for 60 min. The absorbance was determined at 530- (excitation) and 590-nm wavelength (emission). Measurements of ROS were previously calibrated using exogenous 10  $\mu\text{M}$   $\text{H}_2\text{O}_2$  (positive control). All measurements were performed in phenol red-free culture medium (1 ml), pH 7.4, at 37 °C.

### Mitochondrial superoxide ( $\text{O}_2^{\bullet-}$ ) production

The mitochondrial superoxide ( $\text{O}_2^{\bullet-}$ ) production was determined by fluorescent dye MitoSOX™ Red (M36008, ThermoFisher) according to the manufacturer's instructions. Briefly, the dystrophic primary muscle cells were incubated with MitoSOX™ Red for 15 min at 37 °C. MitoSOX™ is selectively accumulated in the mitochondria, and it emits red fluorescence when oxidized by the superoxide anion.

The intensities of MitoSOX<sup>TM</sup> fluorescence were monitored on a fluorescent inverted microscope (Nikon, Eclipse TS100/TS100F) for qualitative analyses. MitoSOX<sup>TM</sup> Red was excited at 514 nm with the fluorescent images been collected at 570–600 nm. Quantitative measurements were performed using a spectrophotometer (Synergy H1, Hybrid Reader, Biotek Instruments, Winooski, VT, USA) at 510-nm excitation and 580-nm emission wavelengths.

### Oxygen consumption

We evaluated oxygen consumption rates using an O2k-FluoRespirometer and used the DatLab software package (OROBOROS, Innsbruck, Austria) for data acquisition and analysis. After treatment, we trypsinized dystrophic muscle cells, centrifuged them (1250 RPM), and incubated them with 2 ml of air-saturated respiration medium in an Oxygraph-2k (O2k, OROBOROS Instruments). We used the following drugs in the assay: 1  $\mu$ M oligomycin (Oligo), 2  $\mu$ M carbonyl cyanate *m*-chlorophenyl hydrazone (CCCP), and 1  $\mu$ M antimycin (Ant). We determined the basal oxygen consumption rate (OCR) by subtracting OCR pre-oligomycin from OCR post-antimycin; ATP-linked OCR was calculated by subtracting OCR post-oligomycin from OCR pre-oligomycin; proton leak was obtained by subtracting OCR post-oligomycin from OCR post-antimycin; maximal OCR was determined by subtracting OCR post-CCCP from OCR post-antimycin; reserve capacity was determined by subtracting OCR post-CCCP from OCR pre-oligomycin; and non-mitochondrial was the value of OCR post-antimycin.

### Western blotting

The Western blotting protocol was previously described (Mizobuti et al. 2019; Rocha et al. 2022). Briefly, the Bradford method was used to determine the total protein content in cell extracts. Thirty micrograms of total protein homogenate was loaded on 6–15% SDS-polyacrylamide gels. After electrophoresis, the proteins were transferred onto nitrocellulose membranes, and the membranes were incubated with primary antibodies: 4-HNE (Bio-Rad AHP1251), Calpain 1 (Santa Cruz Biotechnology sc-7530), calsequestrin (Affinity BioReagents VIIIID12), sarcolipin (Merckmilipore ABT 13), serca 2a (Cell-Signaling 4388S), serca 1a (Cell-Signaling D54G12), Oxphos (Abcam STN-19467), PGC-1 $\alpha$  (Calbiochem 4C1.3), PPAR $\delta$  (Invitrogen PA1-823A), SIRT-1 (Cell-Signaling C14H4), and  $\beta$ -actin (Sigma-Aldrich A1978). Next, the membranes were incubated with the peroxidase-conjugated secondary antibodies: anti-rabbit (Promega Corporation W4011), anti-mouse (Promega Corporation W4021), and anti-goat (KPL14-13-06). To control protein loading,

Western blot transfer, and nonspecific changes in protein levels, the blots were stripped and re-probed for  $\beta$ -actin. The ImageJ software was used to determine the band intensities.

### Statistical analysis

All experiments were repeated independently at least three times. IC50 values of compounds were analyzed using GraphPad Prims 8 software package (GraphPad Software, CA, USA) using non-linear regression curve fitting with the normalized response. Statistical differences were analyzed using one-way ANOVA followed by Tukey test or Student's *t*-test using GraphPad Prism 8. Significant differences were defined as  $P < 0.05$ . All results are expressed as mean  $\pm$  standard deviation (SD).

## Results

### Cell proliferation and viability

To assess the dose-dependent toxicities of idebenone on dystrophic muscle cells, we used the MTT and Neutral Red assays.

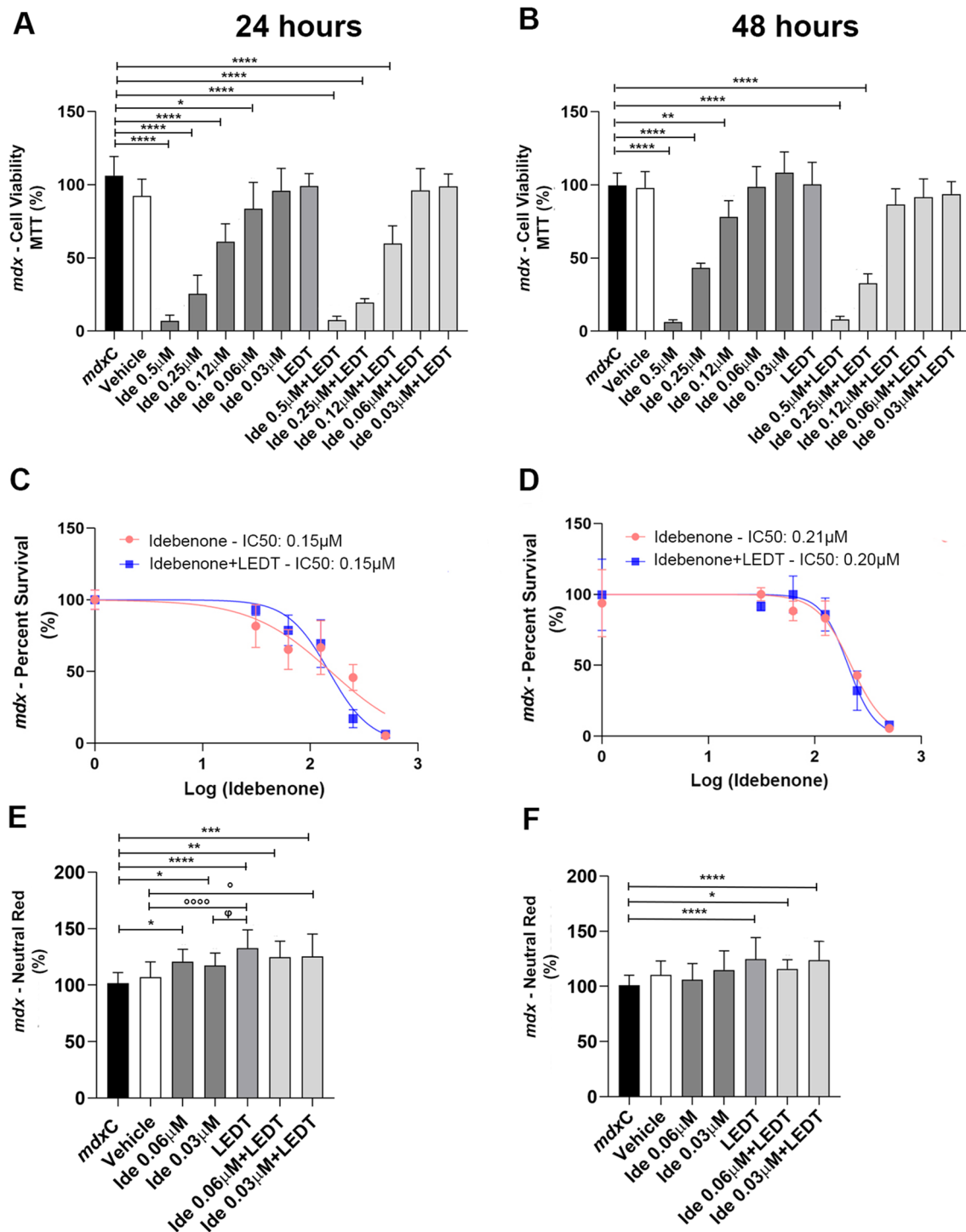
A significant reduction in cell proliferation in treated *mdx* muscle cells was observed in the MTT assay (Fig. 2A, B), when compared to untreated *mdx* muscle cells after 24-h treatment (93.0% for Ide 0.5  $\mu$ M; 76.0% for Ide 0.25  $\mu$ M; 42.4% for Ide 0.12  $\mu$ M; 21.3% for Ide 0.06  $\mu$ M; 93.0% for Ide 0.5  $\mu$ M + LEDT; 81.6% for Ide 0.25  $\mu$ M + LEDT; and 43.6% for Ide 0.12  $\mu$ M + LEDT) and after 48-h treatment (93.7% for Ide 0.5  $\mu$ M; 56.6% for Ide 0.25  $\mu$ M; 21.5% for Ide 0.12  $\mu$ M; 91.9% for Ide 0.5  $\mu$ M + LEDT; and 67.0% for Ide 0.25  $\mu$ M + LEDT).

The half maximal inhibitory concentrations (IC50) (Fig. 2C, D) at 24 h were 0.15  $\mu$ M for idebenone and idebenone plus LEDT and at 48 h were 0.21  $\mu$ M for idebenone and 0.20  $\mu$ M idebenone plus LEDT.

Through the analysis of Neutral Red assay (Fig. 2E, F), a significant increase in the viability of treated *mdx* muscle cells was observed when compared to untreated *mdx* muscle cells after 24-h treatment (18.8% for Ide 0.06  $\mu$ M; 15.3% for Ide 0.03  $\mu$ M; 30.6% for LEDT; 22.6% for Ide 0.06  $\mu$ M + LEDT; and 23.2% for Ide 0.03  $\mu$ M + LEDT) and after 48-h treatment (23.6% for LEDT; 14.8% for Ide 0.06  $\mu$ M + LEDT; and 22.7% for Ide 0.03  $\mu$ M + LEDT).

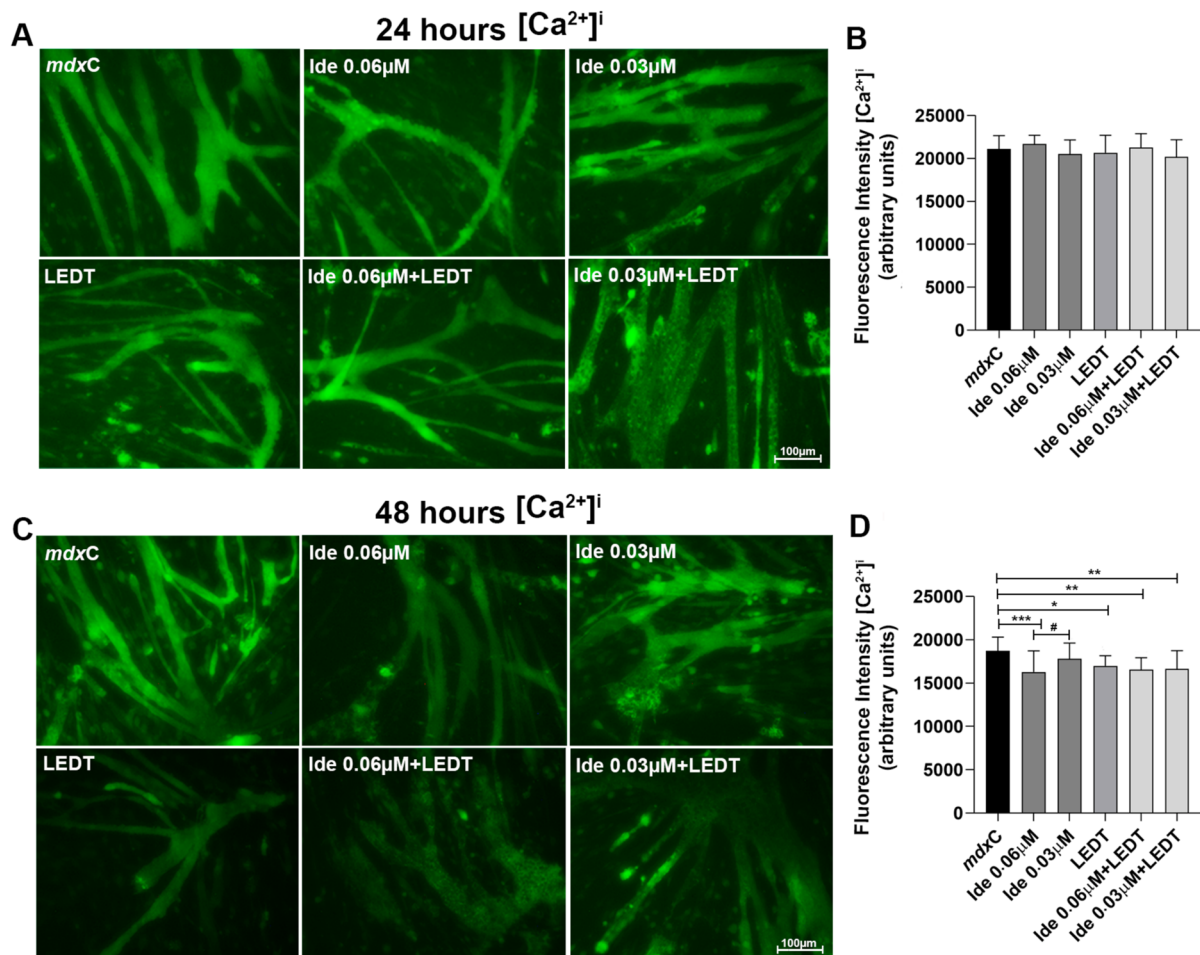
### Intracellular calcium content

Twenty-four hours after applying the treatments (idebenone, LEDT and/or idebenone plus LEDT), no significant



**Fig. 2** MTT assay in untreated *mdxC* muscle cells (*mdxC*); *mdxC* muscle cells treated with carboxymethylcellulose sodium salt (Vehicle); *mdxC* muscle cells treated with different doses of idebenone; *mdxC* muscle cells treated with LEDT (LEDT); and *mdxC* muscle cells treated with LEDT and different doses of idebenone, after 24 h (A) and 48 h (B). The half maximal inhibitory concentrations (IC<sub>50</sub>) in *mdxC* muscle cells treated with different doses of idebenone and *mdxC* muscle cells treated with LEDT and different doses of idebenone, after 24 h (C)

and 48 h (D). Neutral Red assay in *mdxC*; vehicle; *mdxC* muscle cells treated with 0.06  $\mu$ M and 0.03 $\mu$ M of idebenone, LEDT, and LEDT and 0.06 $\mu$ M and 0.03 $\mu$ M of idebenone, after 24 h (E) and 48 h (F). All data are expressed by mean  $\pm$  SD, and the experiments were carried out in triplicate. \* $P$  < 0.05 versus *mdxC*; \*\* $P$  < 0.01 versus *mdxC*; \*\*\* $P$  < 0.001 versus *mdxC*; \*\*\*\* $P$  < 0.00001 versus *mdxC*;  $\circ P$  < 0.05 versus vehicle;  $\circ\circ\circ P$  < 0.00001 versus vehicle;  $\phi P$  < 0.05 versus Ide 0.03  $\mu$ M



**Fig. 3** Intracellular calcium concentrations [Ca<sup>2+</sup>]<sub>i</sub> in untreated *mdx* muscle cells (*mdxC*); *mdx* muscle cells treated with 0.06 μM and 0.03 μM of idebenone; *mdx* muscle cells treated with LEDT; and *mdx* muscle cells treated with LEDT and 0.06 μM and 0.03 μM of idebenone, after 24 h (A) and 48 h (C). Scale bar: 100 μm, × 20. Graph

showing fluorescence intensity of [Ca<sup>2+</sup>]<sub>i</sub> in experimental groups, after 24 h (B) and 48 h (D) of LEDT and idebenone treatment. All data are expressed by mean ± SD, and the experiments were carried out in triplicate. \**P* < 0.05 versus *mdxC*; \*\**P* < 0.01 versus *mdxC*; \*\*\**P* < 0.001 versus *mdxC*; #*P* < 0.05 versus Ide 0.06 μM

difference in [Ca<sup>2+</sup>]<sub>i</sub> was observed between the experimental groups (Fig. 3A, B). On the other hand, 48 h after treatments, the treated-*mdx* muscle cells showed a significant reduction of [Ca<sup>2+</sup>]<sub>i</sub> (13.0% for Ide 0.06 μM; 9.1% for LEDT; 11.6% for Ide 0.06 μM + LEDT; and 11.0% for Ide 0.03 μM + LEDT) compared to the untreated *mdx* muscle cells (Fig. 3C, D).

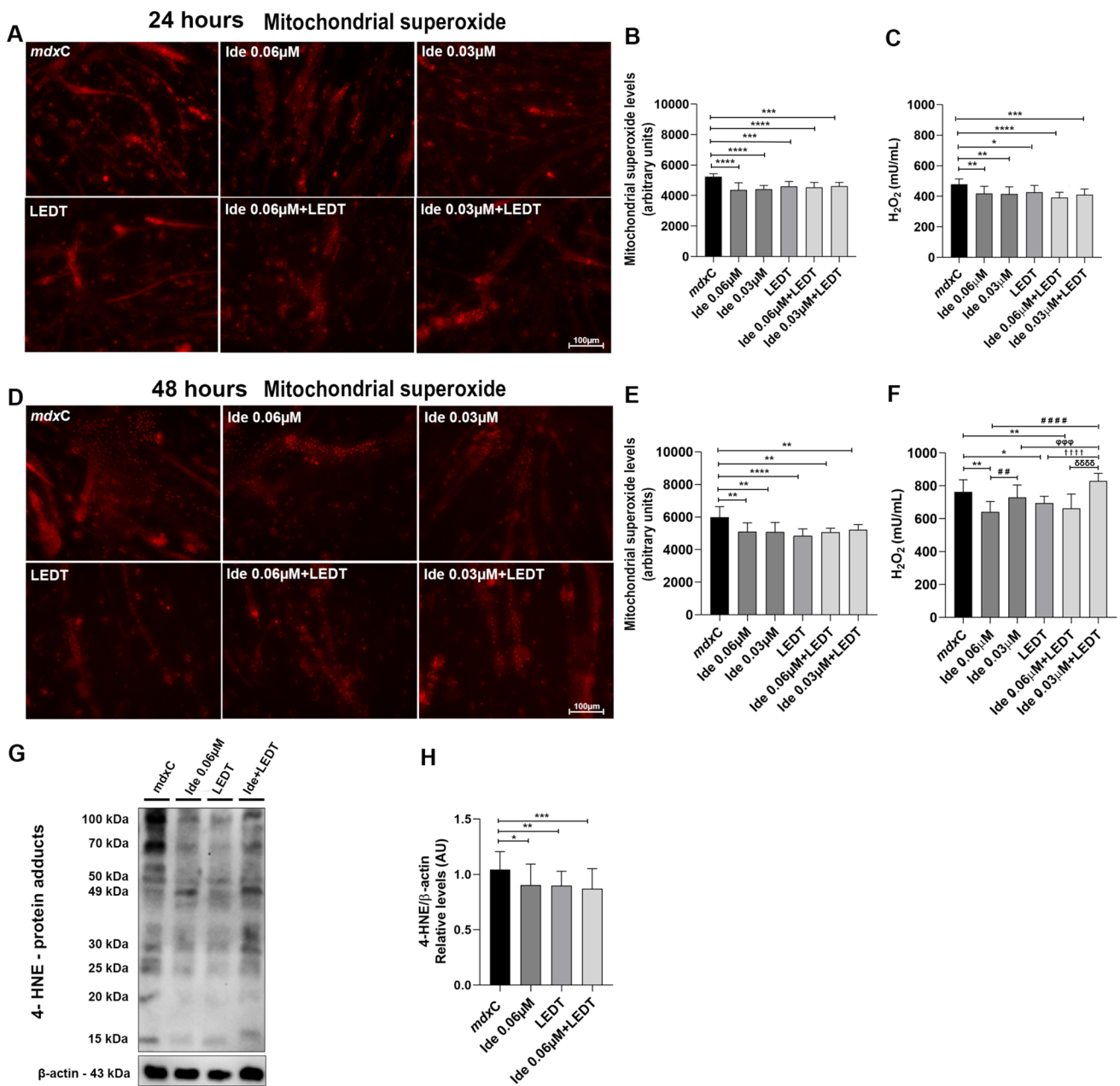
### Oxidative stress

Twenty-four hours after applying the treatments (idebenone, LEDT and/or idebenone plus LEDT), the treated-*mdx* muscle cells showed a significant reduction of O<sub>2</sub><sup>•-</sup> production (16.6% for Ide 0.06 μM; 15.6% for Ide 0.03 μM; 12.1% for LEDT; 13.6% for Ide 0.06 μM + LEDT; and 12.0% for Ide 0.03 μM + LEDT) compared to the untreated *mdx* muscle cells (Fig. 4A, B). Similar results were also observed

after 48 h of treatments (14.7% for Ide 0.06 μM; 15.1% for Ide 0.03 μM; 19.3% for LEDT; 15.3% for Ide 0.06 μM + LEDT; and 12.8% for Ide 0.03 μM + LEDT) compared to the untreated *mdx* muscle cells (Fig. 4D, E).

Regarding H<sub>2</sub>O<sub>2</sub> production, the treated-*mdx* muscle cells analyzed 24 h after all the treatments showed a significant reduction in its production (12.6% for Ide 0.06 μM; 12.9% for Ide 0.03 μM; 10.9% for LEDT; 18.0% for Ide 0.06 μM + LEDT; and 14.3% for Ide 0.03 μM + LEDT) compared to the untreated *mdx* muscle cells (Fig. 4C). However, 48 h after the treatments, only *mdx* muscle cells treated with Ide 0.06 μM, LEDT, and Ide 0.06 μM + LEDT showed a significant reduction in H<sub>2</sub>O<sub>2</sub> production (by 16.0%, 8.9%, and 13.0%, respectively) compared to the untreated *mdx* muscle cells (Fig. 4F).

Regarding the lipidic peroxidation marker, the *mdx* muscle cells treated with Ide 0.06 μM, LEDT, and Ide 0.06 μM +



**Fig. 4** MitoSOX<sup>TM</sup> fluorescence (red) images in untreated *mdxC* muscle cells (*mdxC*); *mdxC* muscle cells treated with 0.06  $\mu$ M and 0.03  $\mu$ M of idebenone; *mdxC* muscle cells treated with LEDT; and *mdxC* muscle cells treated with LEDT and 0.06  $\mu$ M and 0.03  $\mu$ M of idebenone, after 24 h (**A**) and 48 h (**D**). Scale bar 100  $\mu$ m,  $\times$  20. Graphs showing MitoSOX<sup>TM</sup> fluorescence intensity in experimental groups, after 24 h (**B**) and 48 h (**E**) of LEDT and idebenone treatments. The graphs show H<sub>2</sub>O<sub>2</sub> production in experimental groups, after 24 h (**C**) and 48 h (**F**) of LEDT and idebenone treatments. Western blotting analysis of 4-HNE protein adducts (**G**) in *mdxC*; *mdxC* muscle cells treated with 0.06  $\mu$ M of idebenone; LEDT; and *mdxC* muscle cells

treated with LEDT and 0.06  $\mu$ M of idebenone. The graphs (**H**) show 4-HNE protein adducts in experimental groups.  $\beta$ -Actin was used as a loading control. The relative value of the band intensity was quantified and normalized by the corresponding control. All data are expressed by mean  $\pm$  SD, and the experiments were carried out in triplicate. \* $P$  < 0.05 versus *mdxC*; \*\* $P$  < 0.01 versus *mdxC*; \*\*\* $P$  < 0.001 versus *mdxC*; \*\*\*\* $P$  < 0.00001 versus *mdxC*; ## $P$  < 0.01 versus Ide 0.06  $\mu$ M; ### $P$  < 0.00001 versus Ide 0.06  $\mu$ M;  $\phi\phi\phi\phi$  $P$  < 0.001 versus Ide 0.03  $\mu$ M;  $\dagger\dagger\dagger\dagger$  $P$  < 0.00001 versus LEDT;  $\delta\delta\delta\delta$  $P$  < 0.00001 versus Ide 0.06  $\mu$ M+LEDT

LEDT (analyzed 48 h after treatments) showed a significant reduction in 4-HNE protein adduct levels (by 10.0%, 11.0%, and 13%, respectively) compared to the untreated *mdx* muscle cells (Fig. 4G, H).

## Calcium pathways

Regarding calpain-1, the *mdx* muscle cells treated with Ide 0.06  $\mu$ M, LEDT, and Ide 0.06  $\mu$ M + LEDT showed a significant reduction in its levels (by 57.7%, 46.2%, and 64.7%, respectively) compared to the untreated *mdx* muscle cells (Fig. 5A, B).

The calsequestrin levels were significantly reduced in the *mdx* muscle cells treated with LEDT and Ide 0.06  $\mu$ M + LEDT (by 83.0% and 80.9%, respectively) compared to the untreated *mdx* muscle cells (Fig. 5A, B).

In relation to sarcolipin and serca 2a levels, only the *mdx* muscle cells treated with Ide 0.06  $\mu$ M + LEDT showed a significant reduction in their levels (by 49.2% and 30.0%, respectively) compared to the untreated *mdx* muscle cells (Fig. 5A, B).

The serca 1a levels were significantly increased in the *mdx* muscle cells treated with Ide 0.06  $\mu$ M, LEDT, and Ide 0.06  $\mu$ M + LEDT (by 227.4%, 201.9%, and 205.8%, respectively) compared to the untreated *mdx* muscle cells (Fig. 5A, B).

## Mitochondrial pathways

The OCR was evaluated, and it was found that the *mdx* muscle cells treated with Ide 0.06  $\mu$ M increased the basal, ATP-linked, and maximal capacity (by 98.8%, 100.0%, and 55.2%, respectively) compared to the *mdx* untreated muscle cells (Fig. 6A). In addition, the basal, ATP-linked, maximal and spare capacity were increased in the *mdx* muscle cells treated with LEDT (by 107.8%, 201.5%, 138.7%, and

511.7%, respectively) and/or Ide 0.06  $\mu$ M + LEDT (by 51.8%, 85.8%, 96.9%, and 198.6%, respectively) compared to the untreated *mdx* muscle cells (Fig. 6A).

The *mdx* muscle cells treated with Ide 0.06  $\mu$ M, LEDT, and Ide 0.06  $\mu$ M + LEDT presented a significant increase in the OXPHOS levels in complex V (by 125.0%, 94.1%, and 182.3%, respectively) compared to the untreated *mdx* muscle cells (Fig. 6B, C). In addition, the *mdx* muscle cells treated with Ide 0.06  $\mu$ M + LEDT, also showed a significant increase in complex II and I (by 39.1% and 55.4%, respectively) compared to the untreated *mdx* muscle cells (Fig. 6B, C).

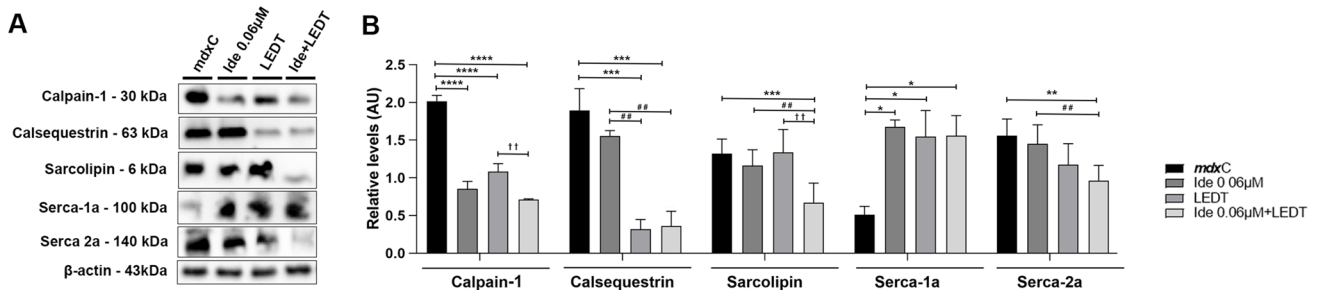
Regarding PGC-1 $\alpha$ , the *mdx* muscle cells treated with Ide 0.06  $\mu$ M, LEDT, and Ide 0.06  $\mu$ M + LEDT presented a significant increase in its levels (by 27.0%, 32.8%, and 27.0%, respectively) compared to the untreated *mdx* muscle cells (Fig. 6D, E). Similar results were also observed in PPAR $\delta$  where the *mdx* muscle cells treated with Ide 0.06  $\mu$ M, LEDT, and Ide 0.06  $\mu$ M + LEDT presented a significant increase in its levels (by 66.6%, 33.3%, and 38.8%, respectively) compared to the untreated *mdx* muscle cells (Fig. 6D, E).

On the other hand, only the *mdx* muscle cells treated with Ide 0.06  $\mu$ M + LEDT showed a significant increase in SIRT-1 levels (31.8%), compared to the untreated *mdx* muscle cells (Fig. 6D, E).

## Discussion

Here, we demonstrated beneficial effects in calcium and mitochondrial signaling pathways in dystrophic muscle cells subjected to LEDT and idebenone treatment applied separately and/or together.

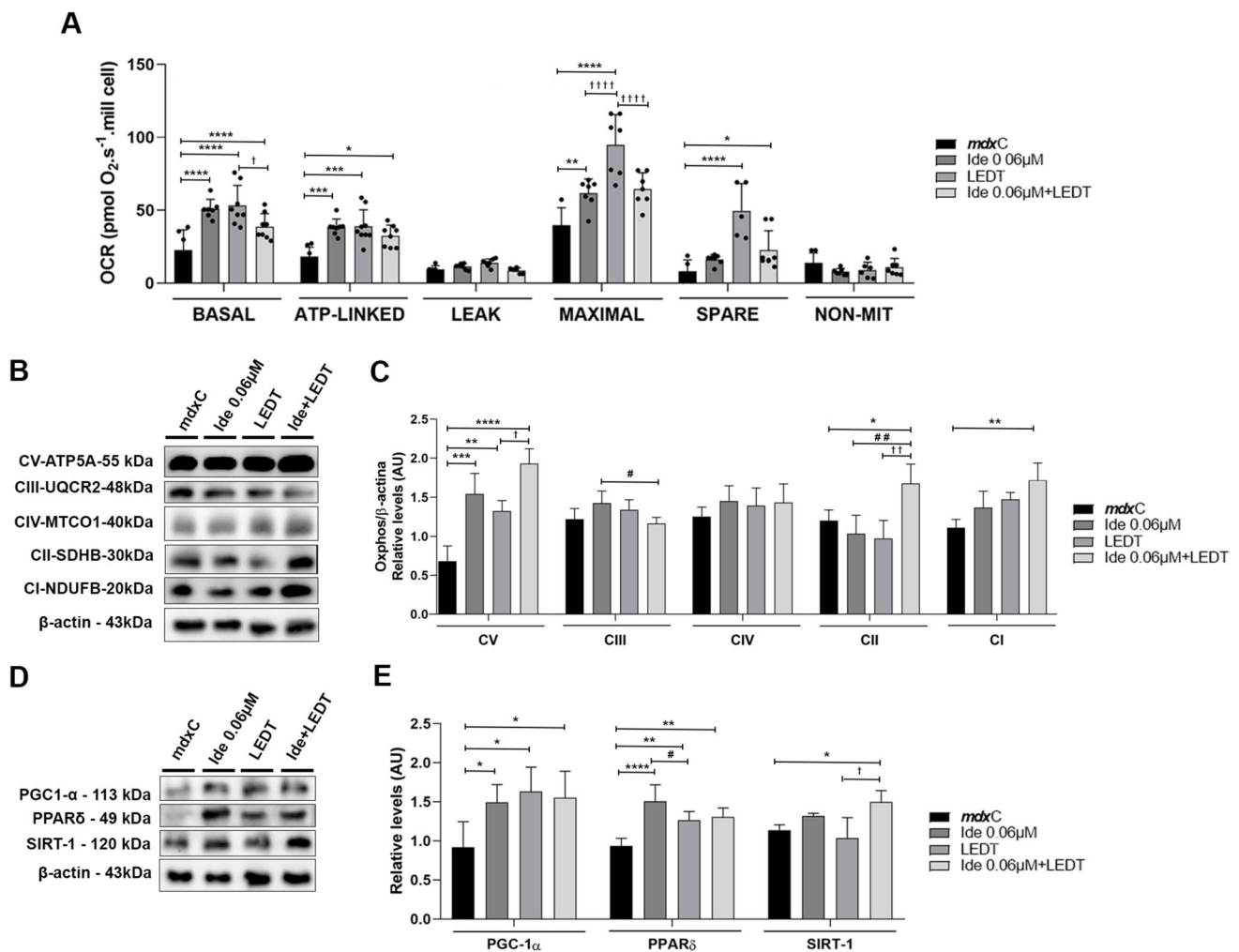
Although the mechanisms involved in DMD are complex and multifactorial, impaired calcium handling and calcium overload are key contributors to disease onset and progression (Mareedu et al. 2021). Increased intracellular



**Fig. 5** **A** Western blotting analysis of calpain-1, calsequestrin, sarcolipin, serca-1a, serca-2a in untreated *mdx* muscle cells (*mdxC*); *mdx* muscle cells treated with 0.06  $\mu$ M of idebenone; *mdx* muscle cells treated with LEDT; and *mdx* muscle cells treated with LEDT and 0.06  $\mu$ M of idebenone. **B** The graph shows the protein levels.  $\beta$ -Actin was used as an internal control. The relative value of the band inten-

sity was quantified and normalized by the corresponding control. All data are expressed by mean  $\pm$  SD, and the experiments were carried out in triplicate. \* $P$  < 0.05 versus *mdxC*; \*\* $P$  < 0.01 versus *mdxC*; \*\*\* $P$  < 0.001 versus *mdxC*; \*\*\*\* $P$  < 0.00001 versus *mdxC*; ## $P$  < 0.01 versus Ide 0.06  $\mu$ M; †† $P$  < 0.05 versus LEDT





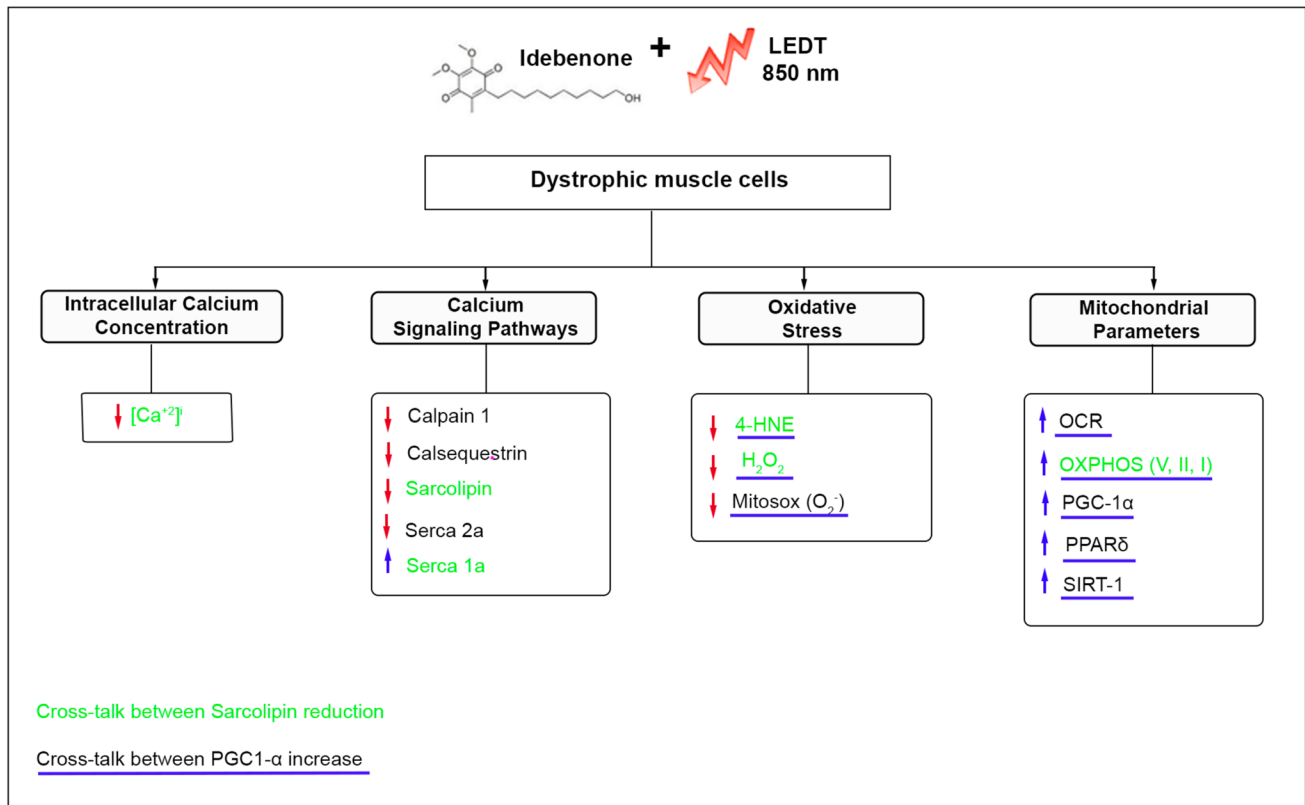
**Fig. 6** **A** OCR values were measured in untreated *mdx* muscle cells (*mdx*C), *mdx* muscle cells treated with 0.06 μM of idebenone, *mdx* muscle cells treated with LEDT, and *mdx* muscle cells treated with LEDT and 0.06 μM of idebenone. The measurements were taken for basal, ATP-linked, leak, maximal, and spare capacity. Non-mitochondrial OCR was subtracted from all data before being used for these analyses. **B** Western blotting analysis of OXPHOS and **D** of PGC-1α, PPARδ, and SIRT-1 in untreated *mdx* muscle cells (*mdx*C); *mdx* muscle cells treated with 0.06 μM of idebenone; *mdx* muscle cells treated

with LEDT; and *mdx* muscle cells treated with LEDT and 0.06 μM of idebenone. **C**, **E** The graph shows the protein levels. β-Actin was used as an internal control. The relative value of the band intensity was quantified and normalized by the corresponding control. All data are expressed by mean ± SD, and the experiments were carried out in triplicate. \**P* < 0.05 versus *mdx*C; \*\**P* < 0.01 versus *mdx*C; \*\*\**P* < 0.001 versus *mdx*C; \*\*\*\**P* < 0.00001 versus *mdx*C; #*P* < 0.05 versus Ide 0.06 μM; ##*P* < 0.01 versus Ide 0.06 μM; †*P* < 0.05 versus LEDT; ††*P* < 0.01 versus LEDT; ††††*P* < 0.00001 versus LEDT

calcium in the context of dystrophin deficiency is a main mediator in myocyte death and fibrotic development (Shirokova et al. 2013). Under our experimental conditions, we observed that LEDT and idebenone treatment decreased the [Ca<sup>2+</sup>]<sub>i</sub> in *mdx* muscle cells. Concomitant with the decrease in [Ca<sup>2+</sup>]<sub>i</sub>, a significant finding of this study was the decrease in sarcolipin (SLN) levels. In agreement with our results, a previous study found improvement in the cytosolic Ca<sup>2+</sup> homeostasis in *mdx* mice lacking SLN (Tanihata et al. 2018).

SLN is an important regulator of the SERCA pump, which is expressed exclusively in the striated muscle of all mammals (Vangheluwe et al. 2005). A previous study provides evidence that SLN upregulation is a molecular basis

for SERCA dysfunction in both DMD skeletal and cardiac muscles (Voit et al. 2017). In addition, SLN knockout mice display an increase in the SERCA function (Tupling et al. 2011; Bombardier et al. 2013). Enhancing the SERCA expression or activity appears to be a promising strategy to treat the dystrophic skeletal muscle, since SERCA accounts for ≥ 70% of Ca<sup>2+</sup> removal from the cytosol (Periasamy and Kalyanasundaram 2007; Tupling 2009). It is important to note that in the present study, we also showed an increase in SERCA 1 levels, which is the primary isoform expressed in the fast-twitch skeletal muscle (Wu and Lytton 1993). However, we also found a reduction in the SERCA 2 levels. In the skeletal muscle, there are some inhibitory proteins that alter



**Fig. 7** LEDT and idebenone treatment effects on mdx muscle cells. Upregulation: blue up arrow; downregulation: red down arrow; cross-talk between sarcolipin reduction: words in green; cross-talk between PGC1-α increase: word underlined in blue

the SERCA action, such as SLN (as mentioned before) and phospholamban (PLN) (Gamou et al. 2020). In vivo studies have consistently found that PLN preferentially binds with SERCA 2, which is highly expressed in the cardiac and slow-twitch muscle (Fujii et al. 1988; Briggs et al. 1992; Babu et al. 2007; Fajardo et al. 2013). On the other hand, SLN has been associated to SERCA 1 (MacLennan et al. 1973). Thus, these previous studies can collaborate with our findings about SERCA and SLN.

Still regarding the calcium signaling pathways, we found a reduction in the levels of calpain-1 and calsequestrin after LEDT and idebenone treatment. It is already well established that the increase in [Ca<sup>2+</sup>]<sub>i</sub> could activate calpains, thereby promoting muscle wasting through increased proteolysis in dystrophic muscles (Bodensteiner and Engel 1978; Spencer et al. 1995). In addition, it was reported that isolated mdx muscle fibers exhibit heightened calpain-mediated proteolysis when compared with normal muscle fibers (Gailly et al. 2007). Thus, targeting a reduction in calpain levels is a promising approach in DMD treatment. On the other hand, calsequestrin (the most abundant calcium-binding protein) showed higher levels in the spared muscles of mdx mice (Pertille et al. 2010). However, in our experimental

conditions, we observed a reduction in calsequestrin levels, suggesting that the decrease of [Ca<sup>2+</sup>]<sub>i</sub>, by other pathways, after LEDT and idebenone treatment, reduced the need for its upregulation.

Another potential target related to impaired Ca<sup>2+</sup> homeostasis in dystrophic muscles is the mitochondria (Robert et al. 2001). Reduced mitochondrial respiration and increased calcium deposits were reported in the fast-twitch muscle fibers of mdx mice (Gaglianone et al. 2019). In addition, mitochondrial Ca<sup>2+</sup> overload can cause mitochondrial ROS production contributing to oxidative stress in dystrophic muscles (Kyrychenko et al. 2015). In the present study, using LEDT and idebenone treatment, we observed improved oxidative capacity in dystrophic muscle cells, by OCR and OXPHOS analyzes. Concomitantly, we also found reduced levels of ROS and lipid peroxidation after treatment. A main mechanism of the beneficial effect of LEDT relies on the improvement of the mitochondrial function by the direct stimulation of the electron transfer chain (Salechpour et al. 2018b). Several studies showed that PBMT can improve mitochondrial functions by increasing ATP and the mitochondrial membrane potential and by decreasing ROS production (Salechpour et al. 2017, 2018a, 2018b). Regarding idebenone, its ability to modulate

the activity of the mitochondrial respiratory chain complexes, as well as its capacity to protect membranes against lipid peroxidation, is well known (Muscoli et al. 2002).

It is also important to highlight that the reduction in SLN levels correlates with the interplay between mitochondrial function and oxidative stress. Supporting what was said before, the reduction of SLN restored the complex proteins (OXPHOS) and their activities, as well as reducing the protein carbonyl content and lipid peroxidation, thus indicating reduced oxidative stress in the dystrophic muscles (Balakrishnan et al. 2022). Interestingly, in the present study, we also observed a reduction in SLN levels in the dystrophic muscle cells treated, suggesting that this signaling pathway may also be one of the mechanisms by which the LEDT and idebenone treatment showed antioxidant effects and improved mitochondrial function in the dystrophic muscle.

The signaling pathway between the silent mating type information regulation 2 homolog 1 (SIRT1), peroxisome proliferator-activated receptor (PPARs), and peroxisome proliferator-activated receptor- $\gamma$  coactivator-1 $\alpha$  (PGC-1 $\alpha$ ) could be another possible explanation for observed ROS reduction in the dystrophic muscle cells. PGC-1 $\alpha$  is a main regulator of mitochondrial biogenesis and function in several tissues (Handschin and Spiegelman 2006) and is regulated by PPARs and SIRT1 (Cantó and Auwerx 2009). LEDT and idebenone treatment increased SIRT1, PPAR $\delta$ , and PGC-1 $\alpha$  in the dystrophic muscle cells, in our study. In agreement with our results, previous studies have demonstrated that PBMT can induce mitochondrial biogenesis by the elevation of SIRT1 and PGC-1 $\alpha$  in the C<sub>2</sub>C<sub>12</sub> muscle cells (Nguyen et al. 2014) and in a transient cerebral ischemia mouse model (Salehpour et al. 2019). Recently, our research group also reported the correlation between elevated PGC-1 $\alpha$  levels and oxidative stress reduction in the dystrophic muscle treated with antioxidants (Silva et al. 2021; Mizobuti et al. 2022). Regarding idebenone, a previous study, collaborating with our results, showed that this antioxidant can positively influence mitochondrial biogenesis by increasing PPARGC1A gene expression in human-induced pluripotent stem cells (Augustyniak et al. 2017).

However, despite the novelty of this study, one limitation must be recognized. While the beneficial effects reported after LEDT and idebenone treatment are statistically significant, the relative change when compared to the control group is low. This fact may have occurred because a single application of LEDT and only one dose of antioxidant were evaluated. In addition, the analysis period after the treatments may also have interfered with the results obtained. In view of this, we intend to evaluate new variables (e.g., several LEDT applications and different exposure periods to the antioxidant treatment) in future experiments.

Summarizing, our findings suggest that LEDT and idebenone treatment, alone or together, can modulate the calcium

and mitochondrial signaling pathways, such as SLN, SERCA 1, and PGC-1 $\alpha$ , contributing to the improvement of the dystrophic phenotype in muscle cells (Fig. 7). It is also important to highlight that data from the LEDT plus idebenone treatment showed slightly better results than those of each separate treatment in terms of SLN, OXPHOS, and SIRT-1. In addition, this work contributes to the knowledge that the combining of new approaches holds promise for better and more specific DMD treatments in the future. The combined therapies usually show a synergistic effect and act on the treatment of the secondary consequences of the muscular dystrophy (e.g., oxidative stress and mitochondrial dysfunction) and are more effective than single therapies. Therefore, the future technical advances in combined therapeutic assays will help to lead to more effective treatment for DMD.

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## Declarations

**Conflict of interest** The authors declare no competing interests.

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