



Melatonin action in tumor skeletal muscle cells: an ultrastructural study



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ABSTRACT

Melatonin (Mel), or N-acetyl-5-methoxytryptamine, is a circadian hormone that can diffuse through all the biological membranes thanks to its amphiphilic structure, also overcoming the blood–brain barrier and placenta. Although Mel has been reported to exhibit strong antioxidant properties in healthy tissues, studies carried out on tumor cultures gave a different picture of its action, often describing Mel as effective to trigger the cell death of tumor cells by enhancing oxidative stress.

Based on this premise, here Mel effect was investigated using a tumor cell line representative of the human alveolar rhabdomyosarcoma (ARMS), the most frequent soft tissue sarcoma affecting childhood. For this purpose, Mel was given either dissolved in ethanol (EtOH) or dimethyl sulfoxide (DMSO) at different concentrations and time exposures. Cell viability assays and ultrastructural observations demonstrated that Mel was able to induce a dose- and time-dependent cell death independently on the dissolution solvent. Microscopy analyses highlighted the presence of various apoptotic and necrotic patterns correlating with the increasing Mel dose and time of exposure. These findings suggest that Mel, triggering apoptosis in ARMS cells, could be considered as a promising drug for future multitargeted therapies.

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1. Introduction

Human rhabdomyosarcoma (RMS) is the most common pediatric soft tissue sarcoma that arises from mesenchymal precursors with the potential to differentiate into skeletal muscle cells, but failing to complete myogenesis because of the presence of chromosomal aberrations (Zanola et al., 2012). RMS are aggressive tumors whose current chemotherapy is often unsuccessful (Dantonello et al., 2015), especially for the alveolar rhabdomyosarcoma (ARMS) that is characterized by a chromosomal translocation that generating the Pax3-Foxo1 fusion protein confers a more aggressive tumor phenotype and resistance to radiation therapy (Jothi et al., 2013). Currently, the employment of a multimodal therapy has increased the survival rate for patients with localized disease to 70% albeit with significant toxicity (Soundararajan et al., 2012). Despite that,

the therapy success for high risk patients is about 20–30% (Li et al., 2013), suggesting the need of more effective drugs.

Over the past years, Mel, beside its strong and documented antioxidant effects (Salucci et al., 2014b; Ganie et al., 2015), has been shown to act as an anticancer drug in different tumors (Sánchez-Hidalgo et al., 2012; Batista et al., 2014; Codenotti et al., 2015), for example via induction of the mitochondrial apoptosis (Bejarano et al., 2009) or attenuation of telomerase activity both in vivo and in vitro (Leon-Blanco et al., 2003).

Numerous studies have described its anticancer action in solid tumors and in leukemia through induction of the apoptotic cell death, leading to replacement of neoplastic cells with healthy cells (Sánchez-Hidalgo et al., 2012; Chuffa et al., 2015). Additionally, Kubatka et al. (2001) demonstrated the anti-tumor action of Mel in breast cancer in vivo, and this effect was then confirmed in clinical studies showing its protective effect from mammary carcinogenesis (Nowfar et al., 2002) and in other neoplastic diseases (Hong et al., 2014; Wu et al., 2015).

Here we have evaluated the ultrastructural changes occurring in the human alveolar RH30 cells upon administration of Mel dissolved either in EtOH (Salucci et al., 2014a) or DMSO (Jardim-Perassi et al., 2014) at different doses and exposure times.

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2. Materials and methods

2.1. Cell culture

Human RH30 cell line, purchased from the European Collection of Cell Cultures (ECACC; Salisbury, UK), was routinely tested for the expression of the Pax3-Foxo1 chimeric protein by both gene sequencing and western blot analysis using polyclonal anti-Foxo1 (H-128) antibody (Rabbit, sc-11350, Santa Cruz Biotechnology, Dallas, USA). Cells were also routinely tested for mycoplasma contamination using the LookOut Mycoplasma PCR Detection Kit (MP0035 Lookout, Sigma), according to the manufacturer's instructions. RH30 cells were cultured in RPMI 1640, supplemented with 10% heat-inactivated fetal bovine serum, 2 mM glutamine, 1% antibiotics (Penicillin and Streptomycin) and maintained at 37 °C in humidified air with 5% CO₂ (Codenotti et al., 2015).

Mel (Sigma, St. Louis, MO, USA) was dissolved in 1% DMSO or 1% EtOH at the concentration of 100 mM. After 1 or 2 mM Mel treatment, the cell behavior was monitored by means of the inverted microscopy. Trypan Blue exclusion assay was employed to quantify cell viability. The relative number of dead and living cells was obtained counting the number of stained (dead) and unstained (live) cells using a Neubauer chamber at reverted microscope. Student's *t*-test was applied for statistical analyses to compare results obtained in control versus treated samples; *P*<0.05 was considered as significance threshold.

2.2. Scanning electron microscopy (SEM)

RH30 cells were cultured on coverslips in Petri dishes. After washing with 0.1 M phosphate buffer, adherent and suspended cells were fixed with 2.5% glutaraldehyde in 0.1 M phosphate buffer for 1 h. The suspended cell population was deposited on polylysine-coated coverslips and the samples were post-fixed with 1% osmium tetroxide (OsO₄) in 0.1 M phosphate buffer solution (PBS) for 1 h. After alcohol dehydration, the samples were critical point dried, gold sputtered and observed using a Philips 515 scanning electron microscope (Codenotti et al., 2015).

2.3. Transmission electron microscopy (TEM)

RH30-treated cells were washed and fixed with 2.5% glutaraldehyde in 0.1 M PBS for 15 min. The cells were scraped and centrifuged at 300 × g for 10 min. The pellets were fixed in 2.5% glutaraldehyde for an additional 30 min. The suspended cells were collected in Eppendorf, centrifuged and fixed for 45 min in glutaraldehyde. The samples were post-fixed in 1% OsO₄ for 1 h, alcohol dehydrated and embedded in araldite. Thin sections were stained with uranyl acetate and lead citrate and analyzed using a Philips CM10 transmission electron microscope (Salucci et al., 2014b).

2.4. Acridine orange (AO) and propidium iodide (PI) nuclei staining confocal laser scanning microscope (CLSM)

Cells were fixed with 4% paraformaldehyde in PBS (pH 7.4) for 30 min, and suspended cells were deposited on poly-lysinated coverslips in Petri dishes. Adherent and suspendend samples were washed twice using PBS and then pre-treated with RNase A 10 µg/mL in PBS for 30 min. After PBS washing samples were exposed to an equal mixture of PI (1 µg/mL, Life Technologies) and AO (1 µg/mL, Life Technologies) diluted in PBS at room temperature (protected from light) for 10 min and then observed at CLSM (Leica Microsystems CMS GmbH); FICT and PI excitation were at 488 and 500 nm, respectively, and their emission signals were detected at

617 and 525 nm. CLSM images are presented as maximum intensity projection or single-plane images (Salucci et al., 2014b).

2.5. Western blotting

Protein homogenates were obtained by harvesting cells in a cold lysis buffer, composed by 20 mM Tris-HCl (pH 7.6), 1% Nonidet P40, 0.5% sodium deoxycholate, 0.1% SDS, 50 mM NaCl and a cocktail of protease inhibitors (Roche) plus phosphatase inhibitors (1 mM Na₃VO₄ and 4 mM NaF).

Bradford reagent assay has been used to calculate protein concentration. Equal amount of protein samples were separated by SDS-PAGE under reducing conditions and transferred to polyvinylidene fluoride membranes. Sample were incubated with specific primary antibodies, Rabbit anti-Bax, polyclonal (Rabbit, sc-526, Santa Cruz Biotechnology, Dallas, USA) or Rabbit anti-Bcl-2, polyclonal (Rabbit, sc-492, Santa Cruz Biotechnology, Dallas, USA) followed by peroxidase-conjugated secondary antibodies (anti-mouse IgG from Santa Cruz Biotechnology, Dallas, USA; anti-rabbit IgG from Thermo Scientific, Erembodegem, Belgium). The immunocomplexes were visualized using the enhanced chemiluminescence reagent (GeneSpin, Milan, Italy) and immune-reactive bands were quantified using densitometry analyses (Software Gel Pro Analyzer, version 4) (Codenotti et al., 2015).

2.6. In silico analysis

The primary mouse tumor cultures from RMS-bearing mice were previously obtained by Keller and colleagues (Rubin et al., 2011) and utilized for gene expression analysis by employing the microarray approach (Illumina Mouse Ref-8 BeadChip, v1.1). Data sets were deposited in the NCBI Gene Expression Omnibus database with the accession number GSE22520. To analyze Mel receptor expression, data were processed using the Partek® Genomics Suite software version 6.6 (Partek, St. Louis, USA). Briefly, the microarray raw dataset were reprocessed by the background correction, normalization and summarization of probe intensities, using the robust multiarray average analysis to determine the specific hybridizing signal for each probe set. After background correction, the data were transformed in base-2 logarithm. Quality control was also performed to identify the presence of arrays with poor quality and evaluate whether batch effect significantly affected the data.

A one-way analysis of variance (ANOVA) was performed to determine differentially expressed genes in mouse alveolar RMS in comparison to normal muscle samples. Fold change >2 and *p*-values <0.05 were used as criteria to evaluate significant difference in gene expression.

3. Results

To analyze whether Mel might affect cell survival in human alveolar RH30 cells, Trypan Blue exclusion assay was performed, revealing that Mel dissolved either in EtOH or DMSO significantly reduced the viability of RH30 cells in a time and dose dependent manner (Fig. 1A).

In support of this evidence, SEM (Fig. 1) and TEM (Figs. 2 and 3) analyses confirmed that Mel induced cell death of RH30 cells in a dose- and time-dependent manner and regardless of the dissolution solvent. In particular, while control adherent cells appeared to be fusiform or star-shaped (1B), 1 mM Mel-treated cells showed a round shape morphology already after 24 h of treatment (1C, 1H). This effect was even increased after 48 h (1D, 1I) or 72 h (1E, 1L) of Mel treatment and accompanied by the membrane blebbing process. After 2 mM Mel treatment in EtOH (1F, 1G) or DMSO (1M-O),

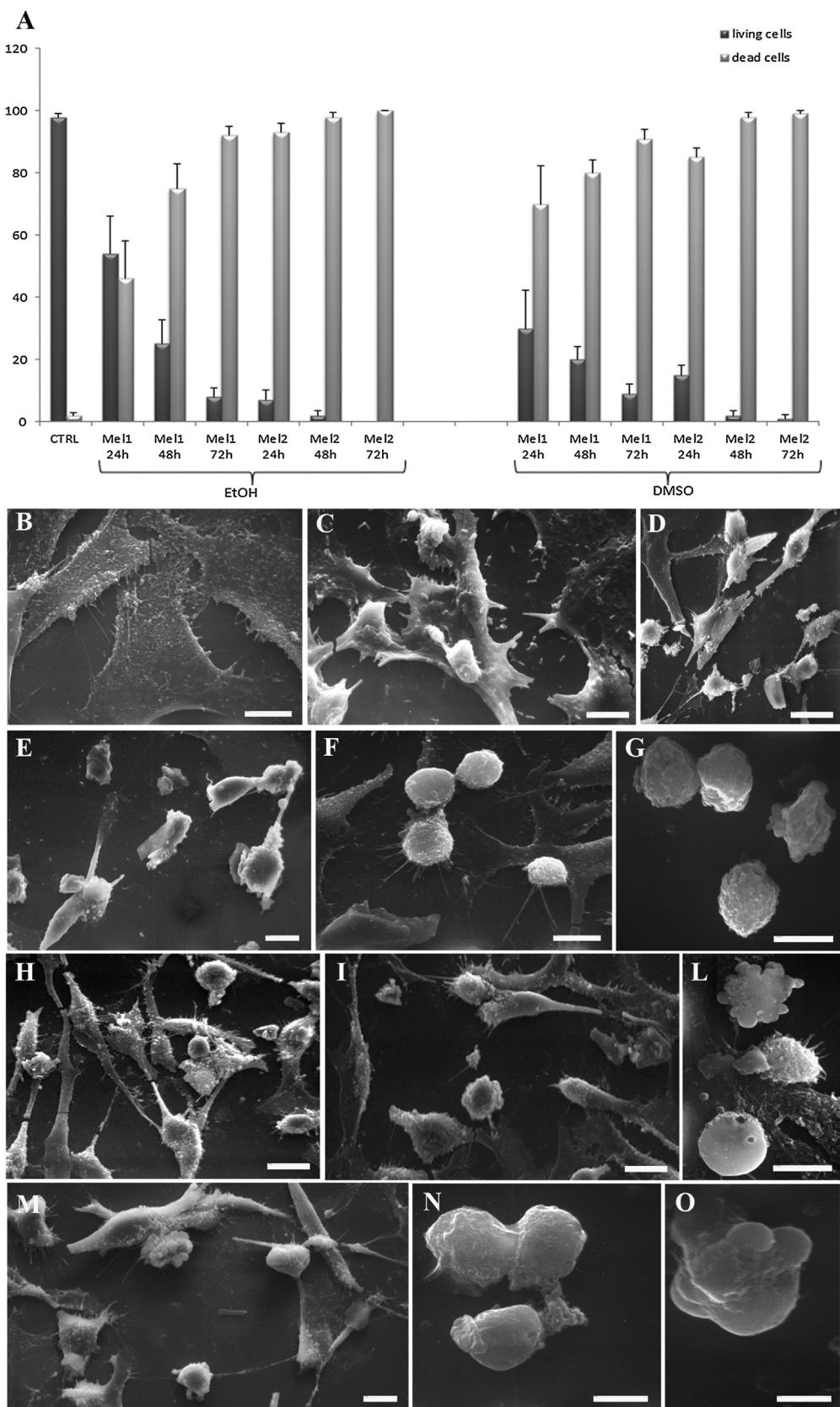


Fig. 1. Trypan blue exclusion assay. Histogram showing the percentages of living (dark grey column) and dead (light grey column) cells, under the different treatment conditions (A). Data are from three independent experiments and are shown as mean of value percentage \pm standard deviation. All Mel treatments are statistically significant versus control condition (T -test, $p < 0.05$).

SEM analysis of control untreated cells (B) versus Mel-treated cells (C–O); in detail, representative pictures showing cells treated with 1 mM Mel (in EtOH) for 24–48–72 h (C, D, E respectively) and 2 mM Mel (in EtOH) for 24–48 (F, G respectively) or, alternatively, with 1 mM Mel (in DMSO) for 24–48–72 h (H, I, L, respectively) and 2 mM Mel (in DMSO) for 24 (M)–48 h (N, O). Bars: 10 μ m B–O.

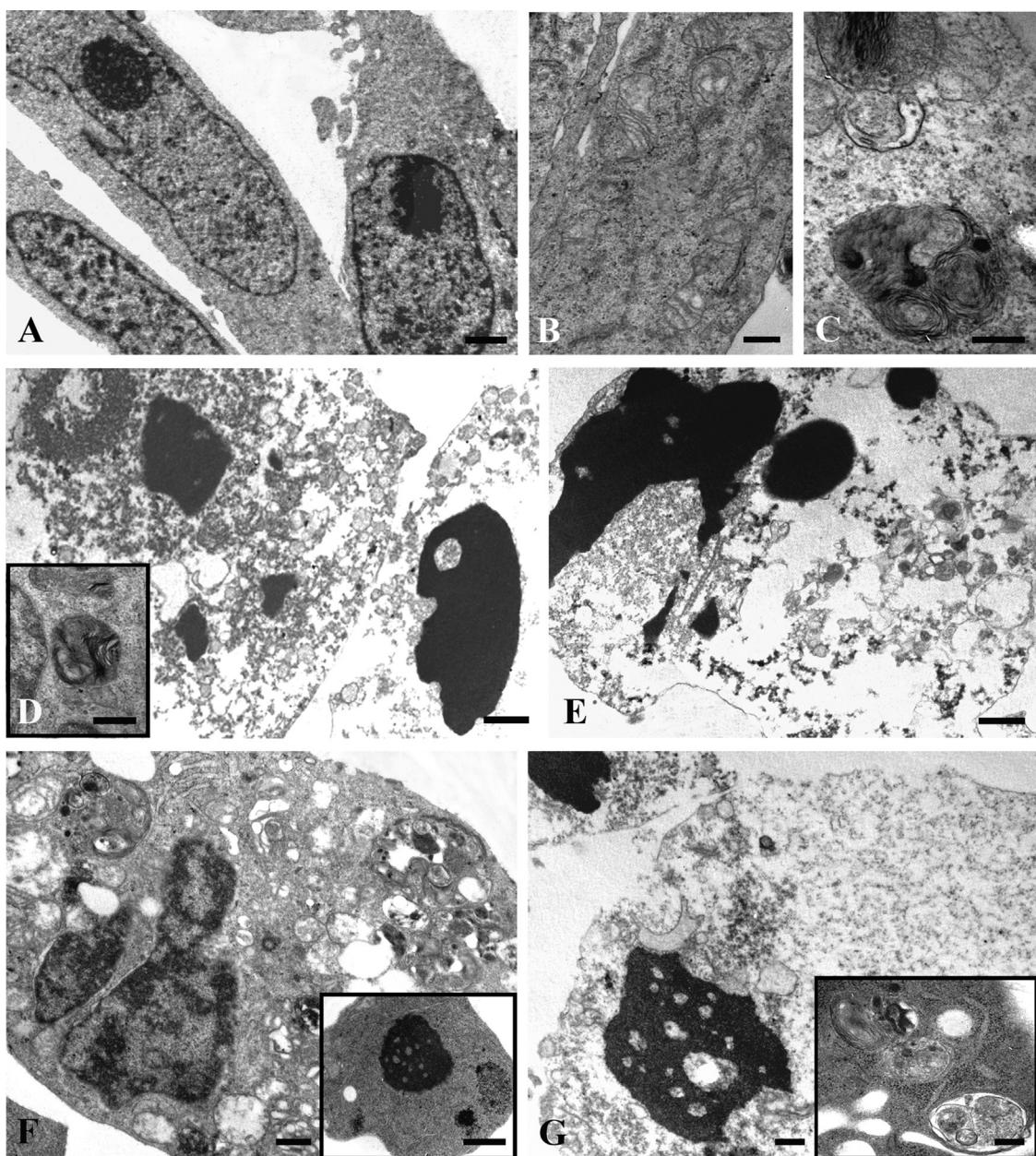


Fig. 2. TEM analysis of control untreated cells (A) versus Mel-treated RH30 cells (in EtOH) (B–G); in detail, representative pictures showing cells treated with 1 mM Mel for 24 h (B, C), 48 h (D, inset D), 72 h (E) and 2 mM Mel for 24–48 h (F, G respectively) Bars: A, B, C, F, inset D, 0.5 μ m; C, inset G, 0.25 μ m; D, E, inset F, 1 μ m.

the diffuse cell damage occurring after 24 h (1F, 1M) and 48 h (1G, 1N, 1O) was characterized by rounding and blebbled cells.

TEM observations (Fig. 2) showed control cells having an elongated shape and intact subcellular structures (2A). Instead, after 1 mM Mel administration for 24 h, cells showed altered mitochondria (2B) and autophagic vacuoles (2C). Occasionally, early apoptotic nuclear features were observed (data not shown). After 48 and 72 h, most of the treated cells appeared to be in late apoptosis (2D, 2E), whereas some of them showed a high number of autophagic-like structures (inset 2D).

Increasing Mel concentration up to 2 mM for 24 and 48 h triggered different cellular alterations, including a diffuse cytoplasmic vacuolization (2F), chromatin condensation (inset F) and autophagic features (inset G). In particular, cells in secondary necrosis were observed after 2 mM exposure for 48 h (2G). Consistent with these results, the effects produced on RH30 cells by Mel dissolved in DMSO were similar (Fig. 3). Early and late apop-

totic patterns could be observed after 1 mM Mel at 24 h (3A), 48 h (3B) and 72 h (3C). Secondary necrosis as well as autophagic features appeared after 2 mM Mel treatment at 24 h (3D, inset D, 3E) and 48 h (3F).

CLSM evaluation of RH30 cells exposed to Mel dissolved in DMSO was carried out by means of AO/PI double staining (Fig. 4). In Fig 4A, the uniform green labeling indicative of cellular healthy structures was visible in control condition. After AO and PI nuclear staining, treated cells (1 mM Mel for 24 h, Fig. 4B) showed a green stain similar to controls, although displaying a more rounded morphology compared to untreated cells. After 48 and 72 h, the number of apoptotic cells increased (Fig. 4C, inset C and D) and some cells with orange areas due to PI permeability were observed (Fig. 4C). Cells treated with 2 mM Mel for 24 h (Fig. 4E, inset E) showed orange nuclei due to late apoptotic events and necrosis.

In accordance with the observed apoptotic response after 1 mM Mel for 48 and 72 h, Western blot analysis showed Mel promoting

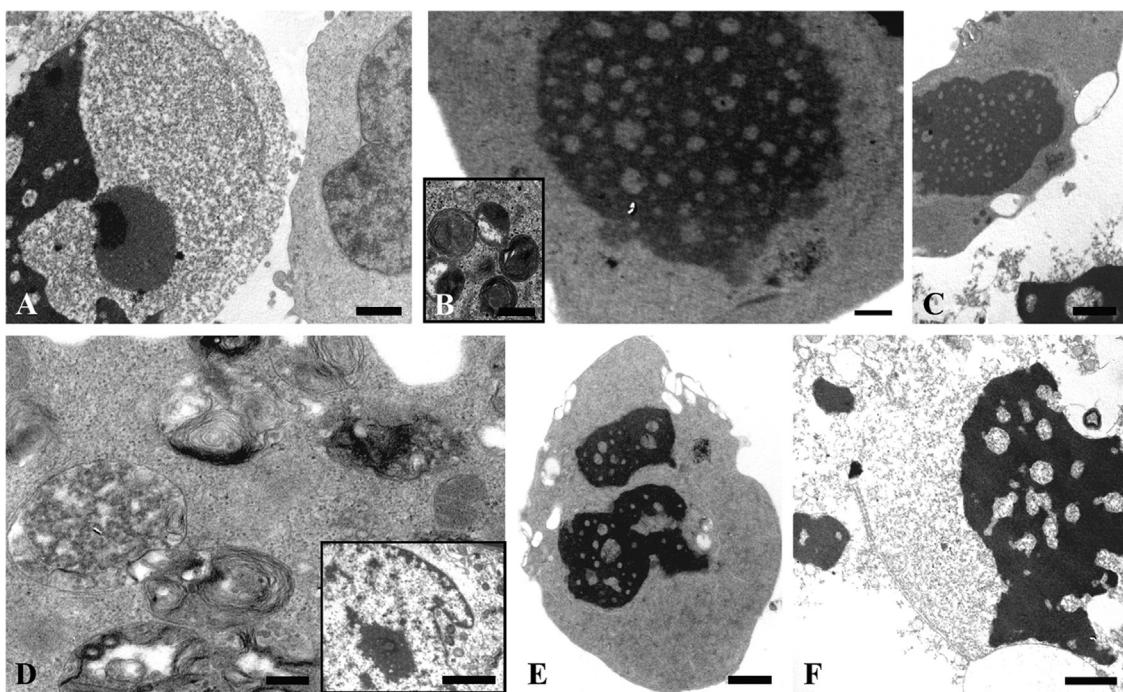


Fig. 3. TEM analysis of Mel-treated RH30 cells (in DMSO) (A–F); in detail, representative pictures showing cells treated with 1 mM Mel for 24–48–72 h (A, B, C respectively) and 2 mM Mel for 24 (D,E) and 48 h (F) Bars: A, C, E, 1 μ m; B, F, 0.5 μ m; inset B, D, 0.25 μ m; inset D, 2 μ m.

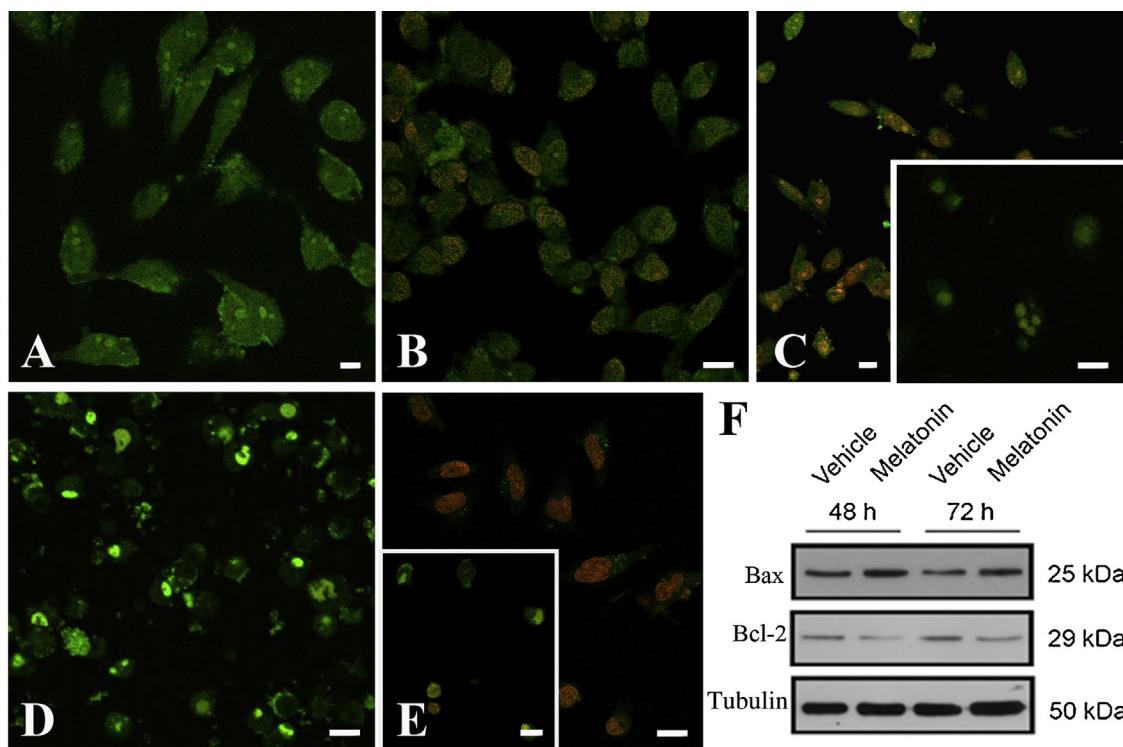


Fig. 4. CLSM analysis of control untreated (A) versus Mel-treated RH30 cells (in DMSO) after AO/PI double staining (B–E). In detail, representative pictures showing cells treated with 1 mM Mel for 24–48–72 h (B, C, D respectively) and 2 mM Mel for 24 h (E). Bars: A–E, 10 μ m. (For interpretation of the references to color in the text, the reader is referred to the web version of this article.)

Western Blot analysis showing expression levels of Bax and Bcl-2 in vehicle-treated versus 1 mM Mel-treated RH30 cells at the indicated time-points (F). Tubulin was used as protein loading control.

both an increase in the pro-apoptotic Bax expression and a decrease in the levels of anti-apoptotic Bcl-2 in comparison to untreated cells (Fig. 4F), therefore confirming that Mel induced apoptosis in RH30 cells.

Finally, an in silico analysis of microarrays data generated from primary mouse RMS tumor cultures (Keller et al., 2004; Keller and Capecchi, 2005) revealed that ARMS tumors could be sensitive to a receptor-dependent Mel action, as the transcript levels of Mel

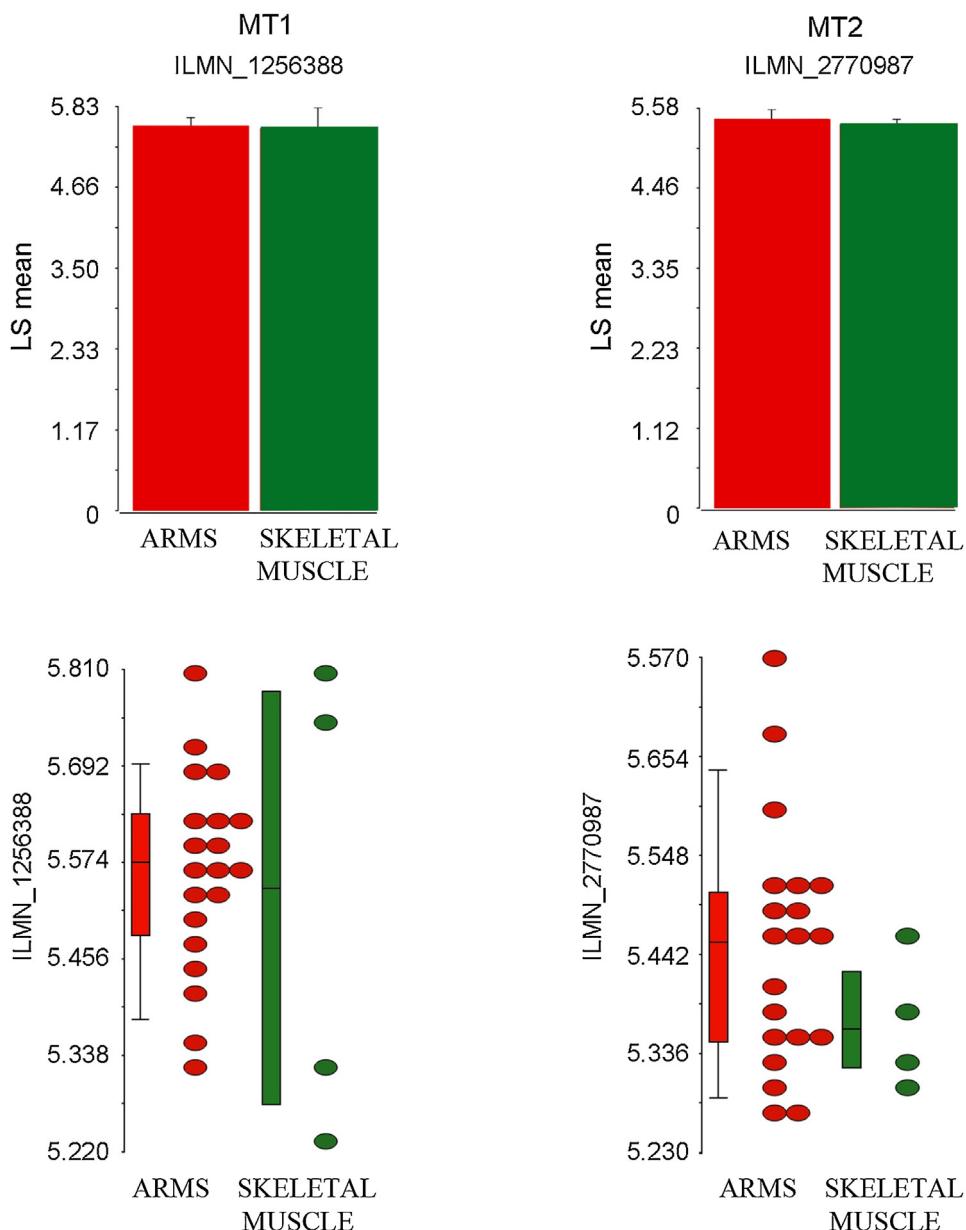


Fig. 5. Dot plot graphs and relative mean histograms representing the transcriptional levels of MT1 (ILMN_1256388 probe number) and MT2 (ILMN_2770987 probe number), as detected in silico by performing a comparison between twenty ARMS samples and four skeletal muscle samples. *p* value was not significant.

receptors MT1 and MT2 were detectable in ARMS subsets with expression levels comparable to those observed in skeletal muscle (Fig. 5).

4. Discussion

Epidemiological studies carried out over the past years have indicated that patients affected by some types of cancer have reduced bloody levels of Mel compared with healthy people (Mills et al., 2005; Lee, 2006). Since then, this pineal neurohormone Mel has been repeatedly described as a potential anti-cancer molecular tool given its demonstrated pro-apoptotic effect and oncostatic action towards different cancer cell lines (Bizzarri et al., 2013; Perdomo et al., 2013; Rodriguez et al., 2013; Hong et al., 2014), therefore contributing to cancer protection (Hrushesky et al., 2009; Mao et al., 2010; Bukowska, 2011; Santoro et al., 2012).

Based on this premise, here we focused on the ultrastructural effects induced by Mel on a human ARMS cell line carrying a unique

molecular signature found among sarcomas, the Pax3-Foxo1 fusion protein arising from a specific chromosomal translocation (Jothi et al., 2013). RMS neoplasm is the most common soft tissue sarcoma of childhood. Although aggressive treatment of RMS could provide long-term benefit, resistance to prolonged therapies is a recurrent problem (Faye et al., 2015). The last decade has seen an extraordinary increase of studies on RMS for understanding how apoptotic mechanisms may contribute on cancer eradication. In this regard, the integrity of apoptosis pathways in RMS constitutes a critical determinant of the sensitivity to most current treatment strategies (Fulda, 2013).

In this work we have first confirmed previous findings showing the ability of Mel administered at pharmacological concentrations (Rodriguez et al., 2013) in triggering cell death of RMS cells (Codenotti et al., 2015); moreover, since the dissolution solvent represents an important variable potentially influencing the Mel action, typically the absolute EtOH (Salucci et al., 2014a; Salucci et al., 2014b) or DMSO (Martín-Renedo et al., 2008; Jardim-Perassi

et al., 2014), here we have established that both these solvents did not influence the efficacy of neurohormone activity as death inducer in RH30 cell line.

Increasing both Mel doses and exposure time lead to late apoptosis and secondary necrosis a consistent number of RH30 cells. In this regard, our ultrastructural analyses revealed that 1 mM Mel dose for up to 48 and 72 h represented the ideal conditions for triggering classical apoptosis-related morphological and biochemical changes, such as surface blebbing, chromatin margination and condensation, mitochondria alteration, cytoplasmic vacuolization, autophagic vacuoles accompanied by Bax activation and Bcl-2 down regulation; in this case, RH30 cells maintained a morphological plasma membrane integrity and necrotic cells were detected only occasionally. On the other side, secondary necrosis and necrotic cell death appeared to be the major death response at very high Mel dose (2 mM). From a therapeutic point of view, killing cancer cells via apoptosis rather than necrosis could be considered a more preferable way, given that the appearance of necrosis *in vivo* might be accompanied by an inflammatory response typically associated to a more unfavorable outcome. Hence, the optimization of dosages for *in vivo* trials should take into account these preliminary *in vitro* observations.

The mechanisms by which Mel influences apoptosis have not been yet fully clarified (Sainz et al., 2003; Wölfle et al., 2001; Gong et al., 2003), although several reports have put forward the ability of Mel to elicit apoptotic effects through changes in the oxidative status. In this regard, a correlation between the increase in ROS production and the induction of Mel-driven apoptosis has been described in different cell lines (Büyükkavci et al., 2006).

Mel action is classically mediated by the interaction with its membrane receptors (Radogna et al., 2009), i.e., MT1 and MT2, which seemed to be expressed also in a number of mouse ARMS tumors, as inferred through an *in silico* analysis that still awaits further *in vitro* and *in vivo* confirmation. Hence, the pro-apoptotic Mel effect on RH30 cells could be receptor dependent, although it is also possible that high doses of Mel may favor its simple diffusion across the cell membrane in a receptor-independent manner. On the whole, understanding whether Mel action in ARMS cells may be mediated or not (or at least partially) by its receptors will be important for identifying the pathways underlying its pro-apoptotic action; in addition, the synthesis and selection of mimetic compounds with increased activity could provide a further pharmacological tool for overcoming the growth of this solid tumor.

5. Conclusions

The data reported in this work indicate that Mel, when administered at pharmacological doses, can profoundly affect cell survival in RMS, the most frequent myogenic soft tissue sarcoma affecting children and adolescents (Ognjanovic et al., 2009; Meyer and Spunt, 2004). In particular, Mel was able to trigger an apoptotic program, independently on the dissolution solvent, therefore representing a promising drug for counteracting RMS tumor progression.

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