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### **Attenuation of neuroinflammation in microglia cells by extracts with high content of rosmarinic acid from in vitro cultured Melissa**

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**Attenuation of neuroinflammation in microglia cells by extracts with high content of rosmarinic acid from *in vitro* cultured *Melissa officinalis* L. cells**

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**Abstract:**

Plant cell culture is a biotechnology cultivation method that permit to cultivate plants in a short period of time and to obtain extracts with a high degree of standardization and high safety profile. The aim of our study was to evaluate the anti-inflammatory and neuroprotective activity of a standardized *Melissa officinalis* L. phytocomplex extract (MD) obtained with an *in vitro* plant cell culture. The MD has been chemically characterized and the content of total polyphenols was  $5.17 \pm 0.1\%$  w/w, with a content

of rosmarinic acid (RA), its main constituent, of  $4.02 \pm 0.1\%$  w/w. MD was tested in an *in vitro* model of neuroinflammation, in which microglia cells (BV2) were stimulated with Lipopolysaccharides (LPS; 250 ng/mL) for 24 h and its pharmacological activity was compared with that of RA. MD (10  $\mu\text{g/mL}$ ) and RA (0.4  $\mu\text{g/ml}$ ) reduced pro-inflammatory factors (NF- $\kappa\text{B}$ , HDAC, IL-1 $\beta$ ) in LPS-stimulated BV2 cells and counteracted the toxic effect produced by activated microglia medium on neuronal cells. This work shows the efficacy of MD on reducing microglia-mediated neuroinflammation and promoting neuroprotection, highlighting the innovative use of *in vitro* plant cell cultures to obtain contaminant-free extracts endowed with marked activity and improved quali-quantitative ratio in the constituents' content.

**Keywords:** neuroinflammation; microglia; rosmarinic acid; *Melissa officinalis* L.

## 1. Introduction

Natural products have always had an enormous importance for healthcare, and still nowadays they represent a precious source of medicines, food supplements, cosmetics and so on. Plants produce a huge variety of metabolites, diversifying the chemical scaffolds to adapt, compete and communicate in the environment. Moreover, the coevolution of plants and animals, gives to plants metabolites high chance to be biologically active and also likely to be substrates for one or more of the many transporter systems that can deliver the compounds to their intracellular site of action[1]. Many natural compounds exhibit a wide range of pharmacophores and a high degree of stereochemistry, making them promising hits candidates, with very complex chemical structure often difficult to be obtained synthetically[2]. A review work by Newman & Cragg dealing with the new medicines approved by the US Food and Drug

Administration (FDA) from 1981 to 2010, highlighted that 34% of those medicines based on small molecules were natural products or direct derivatives of natural products[3]. Moreover, the documented use of plants in traditional medicine systems represents a further resource to supports natural product-based drug-discovery and to provide valuable information on plant safety profile. Plant cell culture technology is a technique for growing of plant cells under strictly controlled environmental conditions. Since plant cells are considered totipotent, they have the potential to express the full genetic machinery coded in the nucleus, and thus, they can produce the full spectrum of characteristic secondary metabolites, found in mother plants [4]. Plant cells are amenable to good manufacturing practice procedures and can be easily propagated by using large volume bioreactors independently on climate or soil or field management practices [4]. Moreover, in vitro cultured plant cells are characterized with fast growth, and the ability to accumulate large amount of uniform biomass for a short period of time [5]. Additionally, plant cell culture technology offers a reliable and powerful production platform for continuous supply of contamination-free, phytochemically uniform biomass from herbal, aromatic, medicinal, and even from rare and threatened plant species.

Plant cell cultures are very attractive platforms to produce high-biotech ingredients for cosmetic, nutraceutical and food purposes [6]. Their season-independent production and the great water saving compared to traditional agricultural practices make these systems suitable for industrial planning and more sustainable towards environment issues as water, carbon footprint and pesticide or herbicide uses. Moreover, when exploited to get non-food products, plant cell cultures are ethically more acceptable since they do not consume agricultural land[6].

The bioactivity of *Melissa officinalis* L. extracts is mainly attributed, as for the majority of plant formulation, to the qualitative and quantitative composition of secondary metabolites (i.e., phenolic acids, flavonoids and terpenoids). Rosmarinic acid (RA) is a caffeic acid ester with 3,4-dihydroxyphenyllactic acid and it is a main bioactive component of the *Melissa officinalis* L. extracts. The RA and phenolic acids content in the *Melissa officinalis* L. extract is highly variable [7]. The variability is associated with multiple factors, which are difficult to control: seasons, plant age, geographical growing areas and tissues used for the preparation of products [8]. Furthermore, the preparation of standardized *Melissa officinalis* L. derivatives with a reproducible content of metabolites is very difficult. The extreme variability in the content of phytoconstituents of plant preparations obtained directly from a plant, or parts thereof, by extraction negatively impacts the effectiveness of the same. An alternative method for obtaining contaminant-free standardized plant preparation in industrial quantities is to use *in vitro* plant cell cultures [9].

In traditional medicine, *Melissa officinalis* L. and its essential oil, have been used for the treatment of nervous, gastrointestinal, hepatic, and biliary disorders and in modern medicine, it has been shown to control symptoms associated to anxiety and insomnia [10]. In recent years *Melissa officinalis* L. was also associated with beneficial effects in several skin disorders and a protective activity from damages by UVB exposure was described [11]. This effect is related to the antioxidant and free radical scavenging properties of its main constituent, rosmarinic acid, indicating a potential antiaging activity of *Melissa officinalis* L. preparations [12].

The increasing life expectancy of populations in developed countries makes modulating aging an urgent medicinal priority. Recent evidence highlights chronic low-grade inflammation as a feature of aging, the so-called “inflammaging” [13]. In particular, neuroinflammation is an important physiological process responsible for the onset of numerous aging-degenerative diseases. The main cell type involved in this activity are microglia cells, devoted to the maintenance of homeostasis in the CNS. After an insult, microglia assume a pro-inflammatory phenotype (M1) and potentially release a large amount of pro-inflammatory cytokines, which are thought to induce tissue damage, demyelination, and neuronal death in the CNS [14].

On these bases, the aim of our study was to evaluate the pharmacological profile of an in vitro cell cultured *Melissa officinalis* L. cells extract (MD) with high content of RA. Our results showed that MD attenuated microglia-mediated neuroinflammation and promoted neuroprotectant effects, highlighting the use of extracts from in vitro plant cell cultures as an innovative and effective strategy.

## **2. Materials and Methods**

### **2.1 Phytocomplex preparation from selected *Melissa officinalis* L. cell line**

The stabilized and highly selected cell line specified on the synthesis of rosmarinic acid of *Melissa officinalis* L. was obtained as described in Pressi et al [9]. 10 days old cell cultures of *Melissa officinalis* L. were filtered by 50 µm mesh filter and the medium cultures were discarded. Cells were washed with a double volume of saline solution (0.9% w/v NaCl in sterile water) and added with 1% (w/w) of citric acid and then homogenized with ultraturrax at 15000 rpm for 20 minutes. The biomass of homogenate cells was dried to obtain a phytocomplex (MD). Drying of the homogenate cells was

performed using a Mini Spray Dryer (BUCHI-B290). The drying process was carried out by setting the following parameters in the Spray Dryer: 180°C of inlet air temperature, 12% of rate pump sample output, 55 mm of gas volume and 103°C-104°C of outlet air temperature.

## 2.2 NMR analysis

*Melissa officinalis* L. phytocomplex (MD) was extracted and subjected to  $^1\text{H}$  NMR analysis. Thirty mg of freeze-dried material were extracted using 1 mL of mixture (1:1) of phosphate buffer (90 mM; pH 6.0) in  $\text{H}_2\text{O}-d_2$  (containing 0.1% TMSP) and  $\text{MeOH}-d_4$  by ultrasonication (TransSonic TP 690, Elma, Germany) for 20 min. After this procedure, the sample was centrifuged for 10 min ( $17000 \times g$ ), then 700  $\mu\text{L}$  of supernatant were transferred into NMR tube.  $^1\text{H}$  NMR spectrum was recorded at 25°C on a Varian Inova instrument (equipped with a reverse triple resonance probe) operating at  $^1\text{H}$  NMR frequency of 600.13 MHz, and  $\text{H}_2\text{O}-d_2$  was used as internal lock. The  $^1\text{H}$  NMR spectrum consisted of 256 scans (corresponding to 16 min) with the relaxation delay (RD) of 2 s, acquisition time 0.707 s, and spectral width of 9595.8 Hz (corresponding to  $\delta$  16.0). A presaturation sequence (PRESAT) was used to suppress the residual water signal at  $\delta$  4.83 (power = -6 dB, presaturation delay 2 s).  $^1\text{H}$  NMR spectrum was manually phased and baseline corrected and calibrated to the internal standard trimethyl silyl propionic acid sodium salt (TMSP) at  $\delta$  0.0 using Mestrenova software (Mestrelab Research, Spain). The signals elucidation was performed also based on an in-house library and comparison with literature [15,16].

## 2.3 UPLC-DAD analysis



100 mg of powder of MD were weighed into a 15 ml test tube and 30 volumes of ethanol (Honeywell) and water 60:40 (v/v) were added. The suspension was mixed for 30 seconds with a vortex mixer and sonicated for 15 minutes in an ice bath; finally, it was centrifuged at 4000 rpm for 15 minutes at 6°C. At the end of centrifugation, the supernatant was recovered. 15 ml of supernatant were transferred into a new tube and preserved in ice until loading into the UPLC system. The sample was diluted 1:10 (first 1:5 in a solvent and then 1:2 in water). The diluted sample was filtered over 0.22 µm filters before being loaded into the UPLC system. Five independent replicates of the MD were extracted and measured. The chromatography system used for quantification of the RA consists in an Acquity UPLC BEH C18 1.7 µm column, size 2.1x100 mm, coupled to an Acquity UPLC BEH C18 1.7 µm VanGuard Pre-Column 3/Pk, size 2.1x5 mm. The platform used for the UPLC-DAD analysis comprises a UPLC system (Waters) consisting of an eluent management module, Binary Solvent Manager model I Class, and an auto-sampler, Sample Manager – FTN model I Class, coupled to a PDA  $\lambda$  diode array detector. Empower 3 (Waters) software was used to acquire and analyze the data. The chromatography method used was the following: solvent A: water, 0.1% formic acid; solvent B: 100% acetonitrile. The initial condition is 99% solvent A; moreover, the flow remains constant at 0.350 ml/min throughout the duration of the analysis. The chromatography column was temperature controlled at 30°C. Elution of the molecules was conducted by alternating gradient and isocratic phases, as indicated in Table 1.

**Table 1.** Elution of the molecules in UPLC-DAD analysis

| <b>Time from start of the analysis (min)</b> | <b>Percentage of solvent B</b> | <b>Slope</b> |
|--|--------------------------------|--------------|
| 0  | 1%                             |              |
| 1  | 1%                             | linear       |
| 11   | 40%                            | linear       |
| 12   | 100%                           | linear       |
| 13   | 100%                           | linear       |
| 13.10  | 1%                             | linear       |
| 15   | 1%                             | linear       |

Quantification of total polyphenols and RA in the samples were based on absorbance UV/VIS spectra measured at 330 nm. The amount of RA and the total polyphenols (expressed as equivalent of RA) were evaluated through the comparison with calibration curves obtained from serial dilution of the authentic commercial standard of RA (CAS 20283-92-5; purity $\geq$ 95%; Sigma Aldrich). The data analysis was carried out with Empower 3 software.

#### **2.4 Cell culture**

A microglial line BV-2 (microglial cells, Tema Ricerca, Genova, Italy) was used for this study. The cells were thawed and placed in a 75 cm<sup>2</sup> flask (Sarstedt, Nümbrecht, Germany) in a medium containing RPMI with the addition of 10% of heat-inactivated (56 °C, 30 min) fetal bovine serum (FBS, Gibco®, Milan, Italy) and 1% glutamine. Cells were grown at 37 °C and 5% CO<sub>2</sub> with daily medium change[17]. A human neuroblastoma cell line SH-SY5Y was kindly donated by Prof. Lorenzo Corsi

(University of Modena and Reggio Emilia, Italy), was cultured in DMEM and F12 Ham's nutrients mixture (Sigma-Aldrich, Milan, Italy), containing 10% heat-inactivated FBS (Sigma-Aldrich), 1% L-glutamine (Sigma-Aldrich), and 1% penicillin-streptomycin solution (Sigma-Aldrich) until confluence (70–80%). The cells were grown in a humidified atmosphere with 5% CO<sub>2</sub> at 37 °C. EDTA-trypsin solution (Sigma-Aldrich) was used for detaching the cells from flasks, and cell counting was performed using a hemocytometer by Trypan blue staining as previously reported [18].

### **2.5 Sulforhodamine B (SRB) assay**

Cell viability was assessed by SRB assay. BV2 were seeded in 96-well plates, each corresponding to a different treatment (2x10<sup>4</sup> cells in 200 µL). 100 µL of serum-free RPMI 1640 were added to the wells followed by 25 µL of 50% TCA trichloroacetic acid. The plate was then incubated at 4°C for 1 h. This was followed by 5 washes with double distilled water (100 µL/well). The plate was left overnight to dry at room temperature. The next day, 30 µL of SRB (4 mg/mL) in 1% acetic acid were added to each well and incubated for 30 min at room temperature. Four washes with 1% acetic acid (200 µL/well) were performed. Then 200 µL of TRIS HCl solution at pH 10 were added and incubated for 5 min. Finally, the absorbance was recorded at a wavelength of 570 nm. The treatments were performed in six replicates in three independent experiments, and cell viability was calculated by normalizing the values to the control's mean[19].

### **2.6 CCK-8 (cell counting kit) assay**

The neuroprotective effect of MD and its constituents was evaluated by assessing the cell viability of the SH-SY5Y cells treated with the LPS-conditioned BV2 medium for

24 h. Unstimulated BV2 medium was used as control. Cell viability was performed using the Cell Counting Kit (CCK-8, Sigma-Aldrich) according to the manufacturer's instructions. The absorbance was measured at 450 nm using a MP96 microplate reader spectrophotometer (Safas, Monte Carlo, Principality of Monaco). The treatments were performed in six replicates in three independent experiments, and cell viability was calculated by normalizing the values to the control's mean [20].

## **2.7 Preparation of cell lysate**

BV2 cells were seeded in 6-well plates ( $3 \times 10^5$  cells/well), pre-treated with MD 10  $\mu\text{g}/\text{mL}$  and RA 0.4  $\mu\text{g}/\text{mL}$  for 4 h, and then stimulated with LPS 250  $\text{ng}/\text{mL}$  for 24 h. Then, the conditioned BV2 medium was collected and centrifuged (1000  $\times$  g for 10 min, 37 °C). The pellet was discarded, and the supernatant was stored at -80 °C for BDNF dosages or used to treat SH-SY5Y cells ( $5 \times 10^5$  cells/well in 96-well plates). Proteins from cells were extracted using RIPA buffer (50 mM Tris-HCl pH 7.4, 150 mM NaCl 1% sodium deoxycolate, 1% Tryton X-100, 2 mM PMSF) (Sigma-Aldrich) and the insoluble pellet was separated by centrifugation (12000  $\times$  g for 30 min, 4 °C). The total protein concentration in the supernatant was measured using Bradford colorimetric method (Sigma-Aldrich) [18].

## **2.8 Supernatant protein precipitation**

TCA (25  $\mu\text{L}$ ) was added to 50  $\mu\text{L}$  of medium and incubated at 4 °C for 10 min. This was followed by centrifugation at 14000 rpm for 5 min. The supernatant was then removed, the pellet was washed three times by resuspending in cold acetone, and

centrifuging again at 14000 rpm for 5 min. At this point the microtubes were placed in the stove, favoring the evaporation of the acetone and the drying of the pellets. Finally, the pellet was resuspended in loading buffer 4x and the samples were stored at -20 °C.

## 2.9 Western Blotting

Protein samples (40 µg/sample) were separated by 10% SDS-PAGE. Proteins were then blotted onto nitrocellulose membranes (120 min at 100 V) using standard procedures. Membranes were blocked in PBST containing 5% non-fat dry milk for 120 min and incubated overnight at 4 °C with primary antibodies anti-IKBA $\alpha$  (1:1000, sc-1643, RRID:AB\_627772) (Santa Cruz Biotechnology, CA, USA), anti-pp38, phosphorylated on Tyr182 (1:250, sc-166182, RRID:AB\_2141746), anti-HDAC1 (1:1000, sc-81598), anti-HDAC2 (1:1000, sc-9959) BDNF (1:500, sc-65514) , anti-IL-1 $\beta$ (1:1000, bs6319R) (Bioss Antibodies, MA, USA). The blots were rinsed three times with PBST and incubated for 2 h at room temperature with HRP-conjugated mouse anti-rabbit (1:3000, sc-235,) (Santa Cruz Biotechnology) and goat anti-mouse (1:5000, bs-0296G) (Bioss Antibodies) and then detected by chemiluminescence detection system (Pierce, Milan, Italy). Signal intensity (pixels/mm<sup>2</sup>) was quantified using ImageJ (NIH). The signal intensity was normalized to that of GAPDH (1:5000, sc-32233) (Santa Cruz Biotechnology)

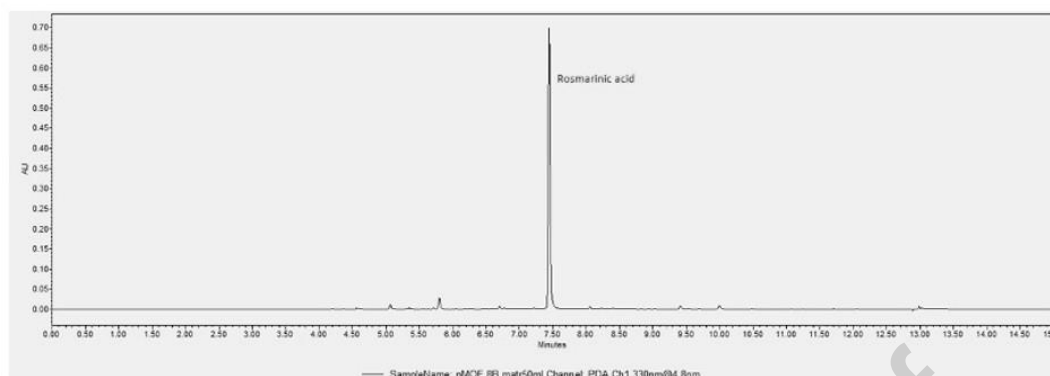
## 2.10 Statistical analysis

Data are expressed as the mean  $\pm$  SEM of six experiments and assessed by One-way or two-way ANOVA, followed by Tukey post hoc test. For each test a value of  $p < 0.05$  was considered significant. The software GraphPad Prism (version 5.0, San Diego, CA, USA) was used in all statistical analysis.

### 3. Results

#### 3.1 UPLC-DAD analysis

The UPLC-DAD analysis was used to estimate the content of total polyphenols and the content of RA in the MD. Total polyphenols expressed as equivalent of RA were quantified by compare of peak areas measured at 330 nm wavelength against the areas of the calibration curve of the reference standard: rosmarinic acid. The RA content was quantified by compare of peak area at retention time 7.5 minutes measured at 330 nm wavelength against the area of the calibration curve of the reference standard: rosmarinic acid. A calibration curve was obtained by the external standard method, using nine different concentrations of the reference standard with two injections per amount. The peak areas measured at 330 nm wavelength were plotted against the known concentrations of the standard solutions to establish the calibration regression; linear regression equation was calculated via the least square method. The chromatographic profile of the MD extract at the wavelength of 330 nm is shown in Fig 1. The content of total polyphenols identifies by their characteristic spectrum with  $\lambda_{\max}$  at 330 nm and expressed as equivalent of RA was  $5.17 \pm 0.1\%$  w/w. The content of RA, calculated measuring the peak area at retention time 7.5, was  $4.02 \pm 0.1\%$  w/w. Thus, the main component of total polyphenols in the MD is represented by RA.



**Fig 1. Chromatographic profile of the MD extract.** The main peak at retention time 7.5 min corresponds to RA.

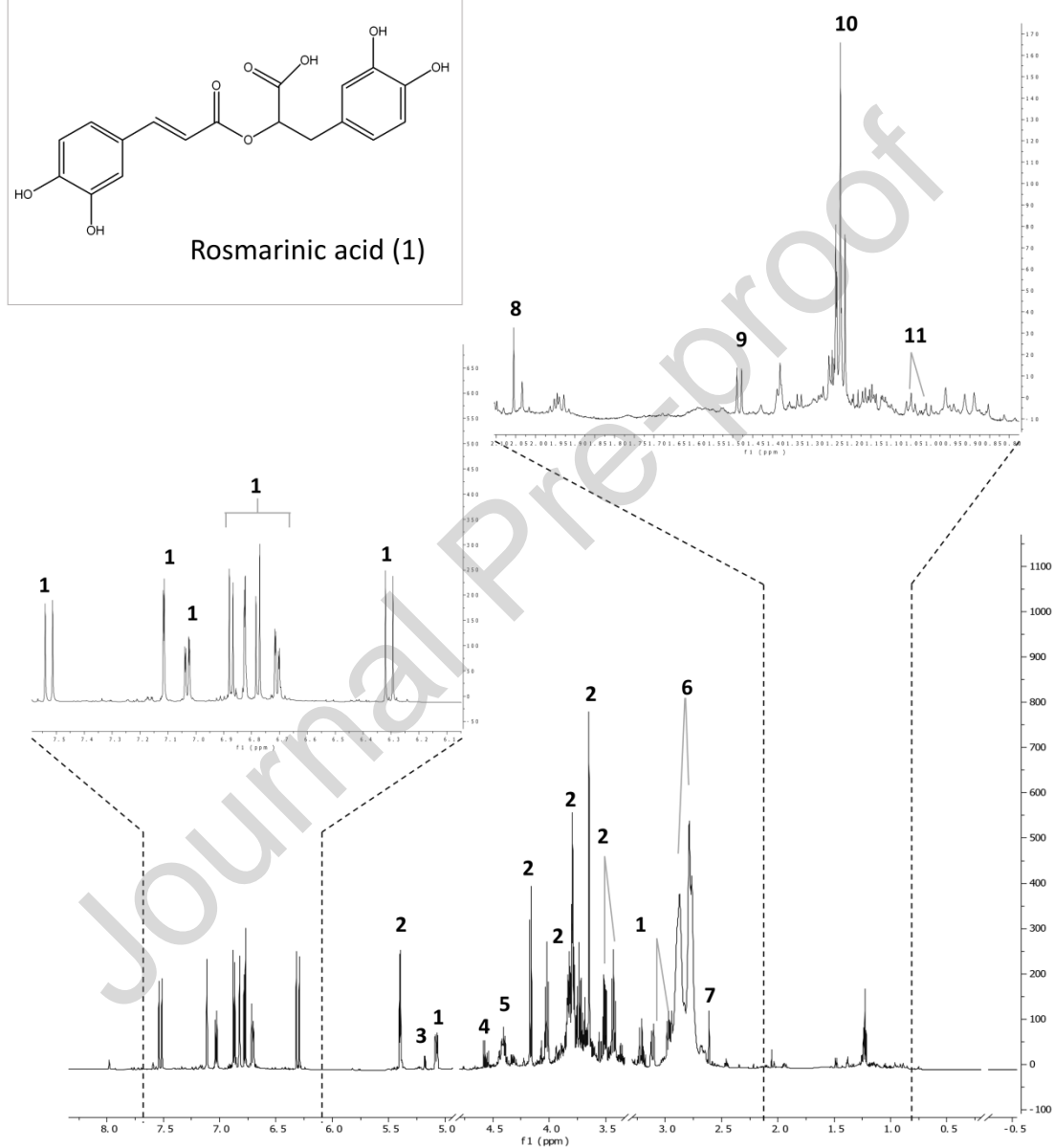
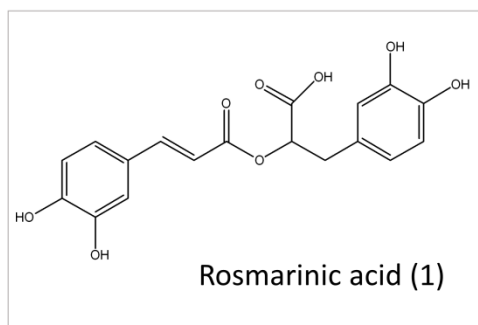
### 3.2 NMR analysis

The  $^1\text{H}$  NMR profiling of MD extract (Fig. 2) showed the presence of both primary and secondary metabolites in MD. In particular, sugars, amino acids and organic acids were detected, namely:  $\alpha$ -glucose,  $\beta$ -glucose, sucrose, alanine, valine, citrate, succinate and acetate. Sugars were identified on the basis of the diagnostic signals generated by their anomeric protons. In particular, sucrose was recognized by the doublet at  $\delta$  5.4 ( $J=3.9$  Hz) which is generated by the anomeric proton of the glucose moiety, while the doublets at  $\delta$  5.2 ( $J=3.8$  Hz) and 4.6 ( $J=7.9$  Hz) are typical of  $\alpha$ -glucose and  $\beta$ -glucose, respectively. Both alanine and valine were detected on the basis of the side chain, namely a methyl group in alanine, which diagnostic signal is a doublet at  $\delta$  1.49 ( $J=7.2$  Hz), and two methyl groups in valine, that generate two doublets at  $\delta$  1.05 and 1.00 ( $J=6.8$  Hz) [15,21]. Broad signals, most likely two doublets, were detected at  $\delta$  2.88 and  $\delta$  2.77 and are ascribable to citric acid; while the singlets at  $\delta$  2.61 and  $\delta$  2.05 are diagnostic of succinic acid and acetic acid, respectively [22]. Moreover, the aromatic

region of the spectrum showed quite intense signals ascribable to RA, whose chemical characterization by NMR and ESI-MS was recently reported by Mandrone, Marincich et al. 2021. In MD extract were evident all the signals of RA, namely: six signals between  $\delta$  6.68 and 7.12 generated by the protons of the two aromatic rings 3,4 hydroxylated, two doublets at  $\delta$  7.52 and 6.31 ( $J=15.9$  Hz) due to the protons of the double bond, the double doublet at  $\delta$  5.08 ( $J_1=3.6$  Hz,  $J_2=9.4$  Hz) generated by the proton in alpha position to the carboxylic group, and finally the two double doublets at  $\delta$  3.11 ( $J_1=3.6$  Hz,  $J_2=14.3$  Hz) and  $\delta$  2.96 ( $J_1=9.4$  Hz,  $J_2=14.3$  Hz) due to the protons of the aliphatic chain.

Moreover, the aromatic region of the spectrum showed quite intense signals ascribable to RA.



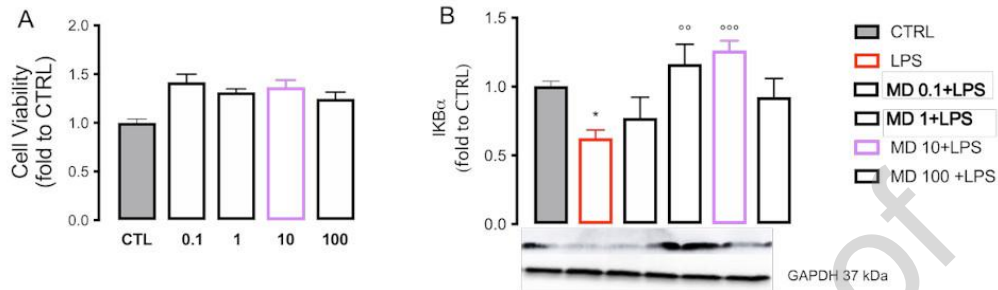


**Fig. 2.**  $^1\text{H}$  NMR profile of MD and chemical structure of RA. The regions of  $\delta$  4.94–4.78,  $\delta$  3.34–3.30, 0.20 – -0.20 were excluded from the analysis because of the residual signals of solvents and standard. 1= RA, 2= sucrose, 3=  $\alpha$ -glucose, 4=  $\beta$ -glucose, 5= unknown sugar, 6= citrate, 7= succinate 8= acetate, 9= alanine, 10= ethanol, 11= valine.

### 3.3 Concentration-response curve of MD extract in BV2 cells

Before directly evaluating the anti-neuroinflammatory of MD extract, we excluded any cytotoxic effect of the extract by treating BV-2 cells with increasing concentrations (0.1, 1, 10 and 100  $\mu\text{g}/\text{mL}$ ) of MD for 24 h. Using the SRB test, cellular toxicity was excluded at all tested concentrations (Fig 3A). To optimize the concentration to be used for investigating the anti-inflammatory effect of MD extract, we tested the extract at different concentration (0.1, 1, 10 and 100  $\mu\text{g}/\text{mL}$ ) in an in vitro model of neuroinflammation. In this model BV-2 cells were treated with LPS 250  $\text{ng}/\text{mL}$  for 24 h. NF- $\kappa\text{B}$ p65 represents a key modulator in microglia activation, indeed it is known to induce an increase of gene transcription involved in inflammation pathway [23].

The levels of NF-  $\kappa\text{B}$ p65 were evaluated indirectly by measuring the I $\kappa\text{B}$  $\alpha$  inhibitor. Figure 3B shows the reduction of I $\kappa\text{B}$  $\alpha$  levels in LPS-stimulated BV2, compared to untreated cell (CTRL). MD extract 1  $\mu\text{g}/\text{mL}$  was already able to reduce the activation of NF- $\kappa\text{B}$ p65, but the peak of the effect was observed at the concentration of 10  $\mu\text{g}/\text{mL}$ , while MD extract 100  $\mu\text{g}/\text{mL}$  started to lose the efficacy. Hence, the concentration of 10  $\mu\text{g}/\text{mL}$  which possessed the maximal activity and did not produce alteration on cell viability, was used for further experiments (Fig 3B).

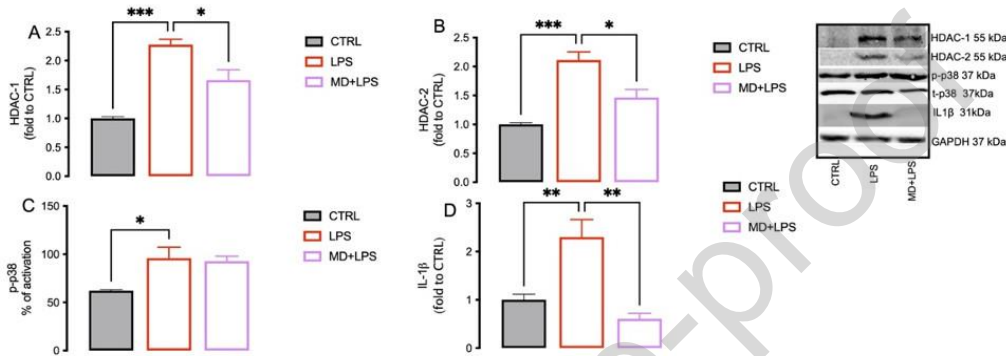


**Fig. 3 Dose-response curve for MD.** A) SRB assay test to evaluate the maximum non-toxic concentration of MD extract ( $\mu\text{g/ml}$ ). B) Effect of MD extract on the expression of IκB $\alpha$ . B) Effect of MD extract at the concentration of 0.1, 1, 10 and 100  $\mu\text{g/ml}$  on IκB $\alpha$  protein expression in LPS-stimulated BV2. One-way ANOVA \* $p < 0.05$  vs CTRL  $^{\circ\circ}p < 0.001$   $^{\circ}p < 0.01$  vs LPS.

### 3.4 MD extract reduced the expression of proinflammatory factors in LPS-stimulated BV2

We evaluated the effect of MD extract on different markers of neuroinflammation, namely HDAC1, HDAC2, and p-p38, which are involved in various signaling pathways related to the inflammatory phenotype of microglia. Indeed, we recently reported that HDAC1 and 2 can epigenetically control the expression of NF- $\kappa$ B, and, consequently, the activation of microglia to the proinflammatory phenotype [24]. LPS increased the expression of HDAC1 (Fig. 4A) and HDAC-2 (Fig. 4B) after 24 h of stimulation. MD extract 10  $\mu\text{g/mL}$  significantly reduced the expression of both HDAC1 and HDAC2. No

effect was observed regarding the activation of p38 (Fig. 4C). The anti-inflammatory effects were evaluated through the dosage of IL1 $\beta$ , a pro-inflammatory cytokine which resulted overexpressed in LPS-stimulated BV2, in the supernatants. We found that the pretreatment with MD completely prevented this effect (Fig. 4D).



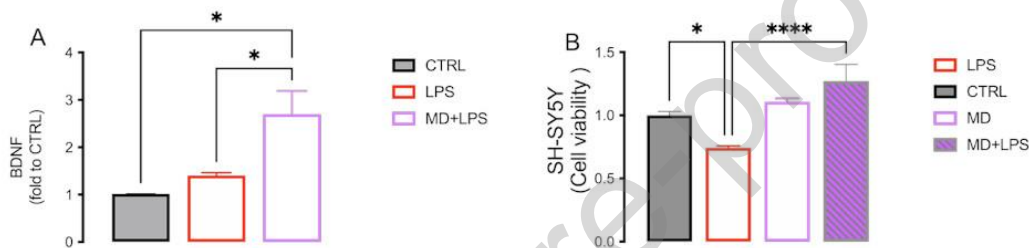
**Fig. 4 MD reduction of proinflammatory factors.** Effect of MD extract 10  $\mu$ g/ml on protein expression of HDAC-1 (A), HDAC-2 (B), p-P38 (C) and IL-1 $\beta$  in LPS-stimulated BV2. One-way ANOVA \*\*\*p<0.001 \*\*p<0.01 \*p<0.05

### 3.5 Neuroprotective effect of MD extract on inflammation-induced neurotoxicity in SH-SY5Y cells

Microglial cells are essential for maintaining neuronal homeostasis. Therefore, their shift towards a proinflammatory phenotype can cause the release of numerous factors that destabilize normal neuronal activity[25].

For this reason, we evaluated the ability of MD extract to modify the release of the neurotrophin BDNF from proinflammatory microglia, to protect neuronal development and its functionality [26]. As reported in Figure 5A, MD extract 10  $\mu$ g/mL increased

BDNF release compared to LPS-stimulated BV2, suggesting a neuroprotective effect of this extract. To better investigate the neuroprotective effect of the MD extract, we treated SH-SY5Y neuronal cells with the BV2 LPS-conditioned medium and evaluated the effect on cell viability. The results obtained (Figure 5B) indicate that the viability of cells treated with LPS alone is reduced compared to controls, while treatment with MD extract reduces the cytotoxicity induced by the model.

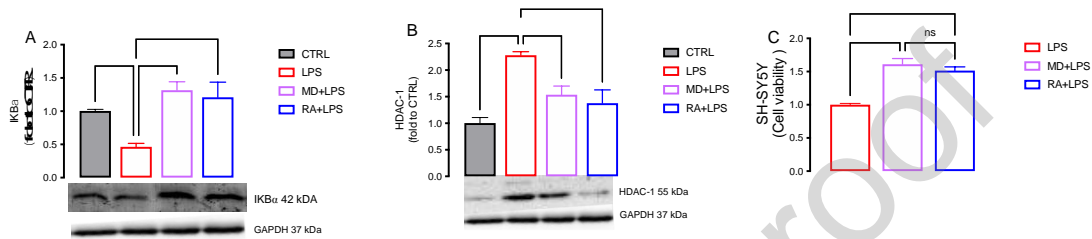


**Fig. 5 Neuroprotective effect of MD.** (A) Increase of BDNF released from BV2 cells stimulated with LPS by MD extract 10  $\mu\text{g}/\text{ml}$ . One way ANOVA \* $P < 0.05$ . (B) Evaluation of neuroprotective effect of MD extract 10  $\mu\text{g}/\text{ml}$  using the CCK8 to evaluate the viability of SH-SY5Y treated with LPS-BV2 conditioned medium. Two - way ANOVA \*\*\*\* $p < 0.00001$  \* $p < 0.05$

### 3.6 RA was the main responsible for the anti-inflammatory and neuroprotective activity of MD extract

To understand if RA was responsible for the activity of the MD extract, we tested the effect of RA in the *in vitro* model of neuroinflammation, using the concentration present in the active dose of MD (0.4  $\mu\text{g}/\text{mL}$ ). The anti-inflammatory effect of RA was

comparable to that produced by MD extract. Indeed, it induced an increase of IKB $\alpha$  in LPS-stimulated BV2 cells (Fig. 6A), while reducing HDAC1 expression (Fig. 6B). In SHSY5Y, RA reduced the toxicity produced by the LPS-conditioned BV2 medium, with an efficacy comparable to that of MD extract (Fig. 6C).



**Fig. 6 RA anti-inflammatory and neuroprotective activity.** Effect of RA (0.4 μg/ml) on IKB $\alpha$  (A) and HDAC-1 (B) in BV2 stimulated with LPS in comparison with MD extract 10 μg/ml. (C) Evaluation of the neuroprotective effect of RA (0.4 μg/ml) on the inflammation-induced neurotoxicity model on SH-SY5Y performed with CCK-8 compared to MD extract (10 μg/ml). One-way ANOVA \*\*\*\*p<0.0001, \*\*\*p<0.001, \*\*p<0.01, \*p<0.05

#### 4. Discussion

Traditional plant extracts are characterized by extreme variability in the extract composition, which depends on several factors, such as climate, soil and cultivation techniques. This variability cannot guarantee the efficacy and reproducibility of the effect of the extract in health care applications. Exploiting *in vitro* cell cultures, we obtained a standardized phytocomplex (MD) with high content of RA, free from pesticides, contaminants, and residual solvents, maintaining the same biological efficacy

in all batches[27]. The aim of our study was to evaluate the anti-inflammatory and neuroprotective activity MD extract (standardized to contain 5.17% w/w of total polyphenols, equivalent to 4.02% RA) obtained with an innovative process through plant cell cultures. The high and standardized content of RA in *Melissa officinalis* L. selected cell line is due to the high plasticity of the de-differentiated cells culture in optimized medium and under controlled conditions.

The cell plasticity of the *in vitro* plant cell culture allows to induce rapid stimuli to the biosynthesis of secondary metabolites by modifying the final liquid medium composition. As described in Pressi et al. 2021 [9] the final liquid medium of *Melissa officinalis* L. cell line with higher content of sucrose and lower content of plant growth hormones induced the biosynthesis of a high content of RA, in comparison to maintenance liquid medium.

*Melissa officinalis* L. is known to have anti-inflammatory activity mainly due to the presence of RA[28]. To confirm these properties and to extend to the evaluation of the activity of MD extract also as anti-neuroinflammatory agent, a microglial BV2-based cell model of neuroinflammation was developed. Under normal conditions, microglia cells are activated in response to a stimulus or insult to the body, developing a physiological inflammatory response to maintain the central nervous system homeostasis[29]. However, the inflammatory process tends to become chronic due to excessive microglial activation, with accelerated aging and loss of function. Thus, present method that involves the use of immortalized microglial cells appears ideal to investigate a protectant activity toward neuroinflammation. Cells were stimulated with LPS for 24 h, with the main consequence being the overexpression of inflammation-

related genes [30]. In response to a damaging stimulus, microglia assume a pro-inflammatory phenotype, characterized by morphological changes and overexpression of cytokines and chemokines such as IL-1, IL-6, and TNF- $\alpha$  [30]. Treating LPs-stimulated BV2 with MD resulted in a reduction of the inflammatory process, with decreased levels of NF- $\kappa$ B, HDAC1, HDAC2, and IL-1 $\beta$ . HDAC is an epigenetic enzyme that regulates the acetylation of histones, modulating the interaction of transcription factors with DNA [31,32]. We previously identified HDAC1 and HDAC2 as inflammatory markers, as a decrease in levels corresponds to a reduction in microglial activation and neuroinflammation [33]. NF- $\kappa$ B represents a transcription factor, the activation of which induces a signalling pathway that ends in the expression of inflammatory cytokines. The main inhibitor is represented by I $\kappa$ B- $\alpha$ , which regulates the expression of NF- $\kappa$ B and the inflammatory process through a negative feedback mechanism. A close correlation between HDAC1 and NF- $\kappa$ B p65 during microglia activation is well documented in the literature[23].

The maintenance of neuronal homeostasis, made possible by continuous communication between cells, is among the main activities of microglia, and BDNF is responsible for this interaction. BDNF has a neuroprotective role and modulates synaptic plasticity.

Activation of microglia towards a pro-inflammatory phenotype compromises neuronal integrity, due to overexpression of inflammatory cytokines and ROS, and decreased levels of neuroprotective factor. There is considerable scientific evidence supporting the neuroprotective effect of *M. officinalis*. For example, a reduction in ROS levels and cellular toxicity was observed *in vitro*[34]. To evaluate the effect of MD on neurotoxicity, we optimized an *in vitro* model, by treating SHSY5Y neuronal cells with



LPS-conditioned BV2 medium for 24 h [18]. The results obtained confirm the neuroprotective effect of MD, with a reduction in cellular toxicity, associated with decreased levels of inflammatory cytokines such as IL-1 $\beta$  and increased BDNF levels. MD has a high RA content which is known to be an effective antioxidant, antibacterial, antiviral, anti-inflammatory, analgesic, neuroprotective, and cardioprotective agent[12]. Particularly, the anti-inflammatory activity of RA was found to be related to the inhibition of the activation of the complement system, as well as to the reduction of the expression of the gene encoding for cyclooxygenase 2, and of several cytokines and pro-inflammatory mediators [35].

RA was tested at the concentration present in the active dose of MD in the neuroinflammation model, leading to a reduction in NF- $\kappa$ B and HDAC1 levels, with an efficacy comparable to that of MD. We thus confirmed that the anti-neuroinflammatory of MD extract is mainly due to its content in RA.

## 5. Conclusion

In conclusion, this work highlights how an extract obtained by in vitro cultured *Melissa officinalis* L. cells (MD) containing high concentration of RA can reduce neuroinflammation at the level of microglial cells, which contributes to the neuroprotectant effects. Indeed, this study indicates in vitro cultured *Melissa officinalis* L. cells MD as an innovative anti-neuroinflammatory approach with potential antiaging activities. Modulating aging is one of the most popular and challenging themes in modern medicine and present findings lay the groundwork to further investigations in more complex models.

## 6. Patents

Patent ITA102019000004113-PCT/IB2020/052589: Phytocomplex and selected extract of a meristematic cell line of a plant belonging to the genus *Melissa*

**Author Contributions:** “Conceptualization, N.G. and G.P.; methodology, N.G., V.B., G.P.; software, V.B validation, V.B., G.P.; O.B.; C.G.; M.M. and I.C.; formal analysis, N.G., V.B.; O.B.; C.G.; M.M. and I.C.; investigation, V.B., G.P; data curation, V.B.; O.B.; C.G.; M.M. and I.C.; writing—original draft preparation, N.G., V.B., G.P.; writing—review and editing, N.G., .G.P.; supervision, N.G.; project administration, N.G., V.B.; funding acquisition, N.G.. All authors have read and agreed to the published version of the manuscript.” Please turn to the CRediT taxonomy for the term explanation. Authorship must be limited to those who have contributed substantially to the work reported.

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**Conflicts of Interest:** The authors declare no conflict of interest.

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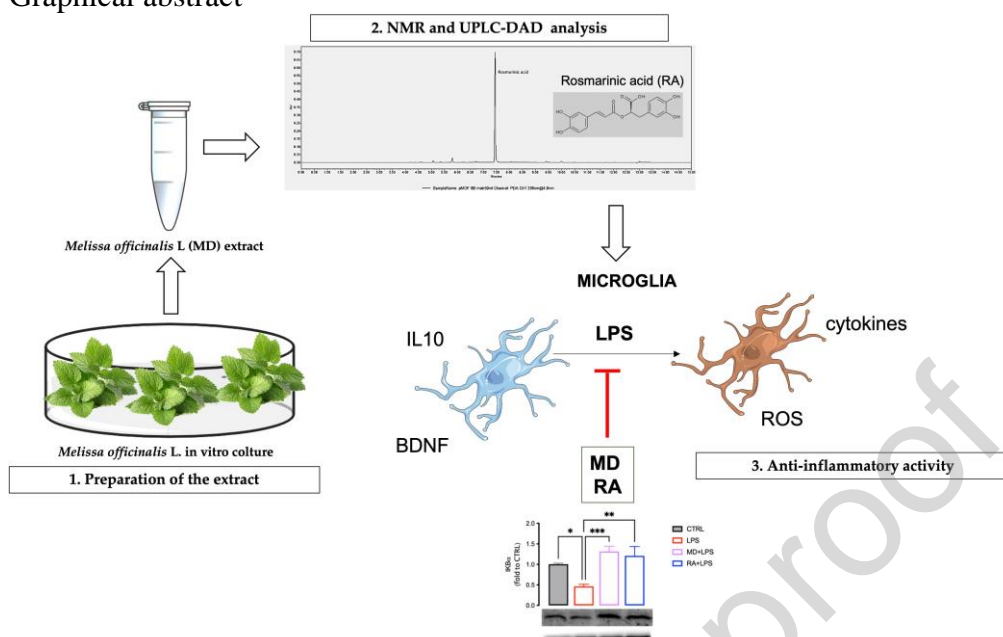
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#### Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

## Graphical abstract



## Highlights:

- Extracts obtained from plant cell culture are standardized and with safety
- Exploiting *in vitro* cell cultures as a standardized phytocomplex of *Melissa officinalis* L.
- *Melissa officinalis* L. extract possessed anti-inflammatory and neuroprotective effects
- These properties are related to the rosmarinic acid concentration.