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Received: 3 September 2025

Accepted: 26 November 2025

Published online: 03 December 2025

Cite this article as: Zakarya A.Y., Rasheed D.M., Emad A.M. *et al.* Comparative metabolomic profiling and chemometric correlation of *Salvia rosmarinus* Spenn. and *Origanum vulgare* L. with antibacterial, antioxidant and anti-inflammatory activities. *Sci Rep* (2025). <https://doi.org/10.1038/s41598-025-30607-z>

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Comparative Metabolomic Profiling and Chemometric Correlation of *Salvia rosmarinus* Spenn. and *Origanum vulgare* L. with Antibacterial, Antioxidant and Anti-Inflammatory Activities

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Abstract

The Lamiaceae plants are recognized in folk medicine for their antibacterial and anti-inflammatory properties. This study reports the first MS-based metabolomics analysis, integrating with chemometrics to explore metabolome heterogeneity in *Salvia rosmarinus* Spenn. (rosemary) and *Origanum vulgare* L. (oregano), and to pinpoint the key metabolites driving their antibacterial, antioxidant, and anti-inflammatory activities. UPLC-QTOF-MS/MS facilitated the identification of 164 metabolites, including flavonoids and hydroxycinnamic acids, which were reported for the first time

in these species. For instance, salvianolic acid D and quercetin coumaroylhexoside were detected in rosemary, while salvianolic acid K, cleroden J, and flavonoids like nepitrin were newly reported in oregano.

In biological evaluation, rosemary strongly inhibited methicillin-resistant *Staphylococcus aureus* and *Escherichia coli*, exhibited the highest radical scavenging capacity in DPPH assay, and showed superior anti-inflammatory effects through COX-II inhibition and TNF- α and NF- κ B suppression. In contrast, oregano displayed the strongest reducing power in FRAP assay. Chemometric analyses revealed that flavonoids, hydroxycinnamic acids, and terpenes were principal discriminating classes. Partial least squares analysis correlated rosemary's antibacterial and radical scavenging activity with hydroxycinnamic acids, flavonoids, and terpenes, while its anti-inflammatory effects were linked to flavonoids and diterpenes. In oregano, FRAP reducing power correlated with benzyl derivatives, organic acids, and hydroxycinnamic acids.

Keywords: Antibacterial, Anti-inflammatory, Antioxidant, Lamiaceae (mint family), Metabolite profiling, Partial least square analysis (PLS)

1. Introduction

The Lamiaceae family (mint family) encompasses a broad range of aromatic plants that are widely distributed but particularly prevalent in the Mediterranean region ¹. Lamiaceae plants have been used extensively as home remedies and to enhance the taste and aroma of foods since ancient times ². The mint family has long been used for decades in cosmetics and conventional medicine as antiseptics, carminatives, expectorants, and sedatives ², in addition to its potential antioxidant and anti-inflammatory properties ^{3,4}. Herbs within Lamiaceae are recognized for their diverse chemical composition, particularly their essential oils, which contribute to their aroma and biological activities ⁵. In addition to essential oils, Lamiaceae plants are rich in non-volatile secondary metabolites *vis.* hydroxycinnamic

acids (e.g., rosmarinic acid, salvianolic acid isomers, caffeic acid), flavonoids (e.g., naringenin, apigenin, luteolin, quercetin), phenolic abietane diterpenoids (e.g., carnosol, carnosic acid, rosmanol), and other phenolics^{6,7}. Attributed to their rich and diverse phytochemical composition, Lamiaceae plants exhibit many biological activities, including antioxidant, antibacterial, antiviral, and antifungal effects^{8,9}. In particular, the antimicrobial activity is especially pronounced in their essential oils¹⁰. This generally highlights the broad biological potential of plants, as evidenced by continuous reporting of various biological activities of different medicinal plants and/or their derived metabolites including, but not limited to, antioxidant, anti-inflammatory, enzyme inhibitory, and cytotoxic effects^{11,12}.

Rosemary (*Salvia rosmarinus* Spenn., previously known as *Rosmarinus officinalis* L.) and oregano (*Origanum vulgare* L.) are mint family plants widely grown in many diverse regions worldwide¹³. Traditionally, these plants have long been used in relieving colds and coughs, and as immunomodulators owing to their pronounced antimicrobial and anti-inflammatory activities¹⁴. These traditional uses highlight their therapeutic importance, which has motivated the exploration of sustainable sources of their active constituents. In this regard, our earlier work underscored the potential of the different distillation by-products as promising antibacterial, antioxidant, and anti-inflammatory resources¹⁵.

This study aims to compare the MS-based phytochemical profiles of these Lamiaceae species and in correlation to their antimicrobial, antioxidant, and anti-inflammatory activities, given the interconnected relationship between oxidative stress, inflammation, and microbial infection, which collectively play a crucial role in the development of many diseases and often overlap in both cause and consequences¹⁶. Furthermore, this study seeks to underpin chemical constituents responsible for well-characterized biological effects using chemometric tools. To achieve such a goal, multivariate data analysis (MVA) represented by partial least square analysis (PLS) was employed to

identify relationships between metabolite profiles and associated biological activities for the first time in Lamiaceae and discern the metabolites contributing most and least to the studied biological activities. Given that *S. rosmarinus* Spenn. and *O. vulgare* L. are also widely used in traditional hygienic practices, mainly in aqueous preparations in the form of infusions or decoctions, antibacterial activity was additionally evaluated for the aqueous extracts to better represent their ethnopharmacological applications.

2. Materials and methods

2.1. Plant material

The aerial parts of *S. rosmarinus* Spenn. and *O. vulgare* L. were acquired from the Medicinal, Aromatic, and Poisonous Plants Experimental Station (MAPPES), Faculty of Pharmacy, Cairo University (Giza, Egypt) in June 2022 (early summer), before the flowering stage. Plants were authenticated supervisors and managers of the MAPPES. Voucher specimens were deposited at the Herbarium of the Pharmacognosy Department, Faculty of Pharmacy, Cairo University (specimens' numbers: *S. rosmarinus* Spenn.; 18.4.24-F, and *O. vulgare* L.; 17.4.24-F).

2.2. Preparation of plant extracts for biological evaluation

Shade-dried, pulverized plants (100 g, each) were extracted separately by maceration till exhaustion using 80% methanol in water to yield alcoholic extracts of *S. rosmarinus* Spenn. (RO) and *O. vulgare* L. (OV). After extraction, the solutions were filtered through Whatman No.1 filter paper and then concentrated under reduced pressure at 40°C using a rotary evaporator (R-210 evaporator, Büchi, Switzerland) to yield (19.5 g RO, 17.5 g OV). The samples were stored in airtight dark containers and kept in a refrigerator at 4°C till further assays.

Aqueous extracts of *S. rosmarinus* Spenn. (Aq.RO) and *O. vulgare* L. (Aq.OV) were prepared following the same procedure, using distilled water

instead of methanol. The resulting yields were 7 g for Aq.RO and 10.5 g for Aq.OV. These aqueous extracts were used exclusively for evaluating antimicrobial activity.

2.3. Preparation of extracts and UPLC-ESI-QTOF-MS analysis conditions

Dried pulverized plants (30 mg of each powder) were mixed separately in 2 mL 80 % HPLC grade methanol with 10 µg/mL umbelliferone (an internal standard), using a Turrax mixer (11000 RPM) for five 20-sec periods, then centrifuged at 3000*g* (4 °C, 15 min) to exclude plant debris, followed by filtration using a 22 µm pore-size filter (Agilent, USA).

An ACQUITY UPLC system (Waters, Milford, MA, USA) was used for UPLC-ESI-QTOF-MS analysis. Chromatographic separation was carried out by injection of alcoholic extracts (3.1 µL) on HSS T3 column (100 × 1.0 mm, particle size 1.8 µm; Waters) at a temperature of 40 °C, where mobile phase A was 0.1% formic acid in water, and mobile phase B was acetonitrile. The flow rate was maintained at 0.15 mL min⁻¹, with the following gradient program: 0-1 min, 5% B; 1-11 min, linear increase from 5% to 100% B; 11-19 min, 100% B; 19-20 min, decrease from 100% to 5% B; and finally, 20-25 min, 5% B. The analytical parameters of the instrument used were previously detailed¹⁷. The system was coupled to 6540 Agilent Ultra-High-Definition Accurate Mass Q-TOF LC/MS (Palo Alto, CA, USA) using an electrospray ionization (ESI) source in both positive and negative ion modes. The operating conditions were applied as described by Baky *et al.*¹⁷, with a fragmentation voltage of 100 V. The Mass Hunter Workstation software (Agilent Technologies) was used for handling data acquisition. Compounds were assigned by comparing retention times (*R_t*), exact masses, and characteristic fragmentation patterns (MS²), as well as the candidates' molecular formula (with 10 ppm mass accuracy limit), and data previously reported in other works of literature.

2.4. Antibacterial activity evaluation

2.4.1. Bacterial strains and culture conditions

Antibacterial activity of the aqueous and hydroalcoholic extracts of *S. rosmarinus* Spenn. and *O. vulgare* L. was carried out against two standard bacterial strains: Methicillin-resistant *Staphylococcus aureus* (MRSA) ATCC 43300 as Gram-positive bacteria, and the Gram-negative bacteria *Escherichia coli* ATCC 25922. The used bacterial strains were available in the stock culture in the Microbiology and Immunology Department, Faculty of Pharmacy (Girls), Al-Azhar University, Giza, Egypt. Nutrient agar, Mueller-Hinton agar, LB broth, and tryptic soy broth were purchased from Oxoid (Hampshire, UK). All reagents and chemicals for buffers were purchased from Sigma-Aldrich Co. (St. Louis, MO, USA).

2.4.2. Minimum inhibitory concentration (MIC) assessment using the broth microdilution method

For anti-microbial susceptibility testing, minimum inhibitory concentrations (MICs) were determined by the broth microdilution method according to the Clinical and Laboratory Standards Institute (CLSI) against standard strains of *E. coli* and MRSA¹⁸. The extracts were prepared in dimethyl sulfoxide (DMSO) at final concentrations ranging from 512 µg/mL to 16 µg/mL. MICs were carried out in triplicate (n = 3), and doxycycline was used as a reference drug control.

2.4.3. Biofilm inhibition assay

The biofilm formation inhibition assay was conducted against MRSA isolates to evaluate the biofilm formation inhibition potential of the tested samples. The biofilm formation inhibition was determined by measuring the absorbance of the adherent biofilms following treatment and comparing these values with those obtained from the untreated controls. A 100 µL of bacterial suspension (1.5×10^8 CFU) in trypticase soy broth supplemented with 1% glucose was added to each well of flat-bottom microtiter plates. Then

100 μ L of $\frac{1}{4}$ MIC of each extract was added to the corresponding well. In each microtiter plate, 6 wells were assigned for positive and negative controls. After incubation at 37°C for 24 h, the microtiter plates were decanted and washed three times with 250 μ L of sterile phosphate-buffered saline (PBS) pH 7.2, fixed by drying for 1 h at 60°C, then stained with 200 μ L of 0.1% w/v crystal violet, and kept at room temperature for 15 min. Finally, the microtiter plates were washed with distilled water, dried, filled with 200 μ L of 33% acetic acid, and transferred (150 μ L) to a new plate. A microplate reader (Tecan Elx800, USA) was used to measure the optical densities at 630 nm, as performed by Badawy *et al.* ¹⁹. The results were represented as a biofilm formation inhibition percentage, calculated using the following equation:

% Inhibition of biofilm formation

$$= 1 - \frac{\text{Optical density of sample}}{\text{Optical density of control (untreated)}} \times 100$$

The assay was performed in triplicate $\pm$ SD (n = 3).

2.4.4. RNA extraction and qRT-PCR-based relative gene expression analysis

The effectiveness of extracts (RO/Aq.OV) to inhibit expression of the *agrA*, *icaA* gene was assessed using quantitative real-time PCR.

MRSA was cultured overnight at 37 °C in LB broth with and without extracts at a concentration of $\frac{1}{4}$ MIC. The total RNA of the cultured MRSA was extracted and converted to DNA using the First High Pure RNA Isolation Kit (Roche Diagnostics GmbH, Germany) and QuantiTects Reverse Transcription Kit (Qiagen, USA), respectively, following the procedure of Saleh *et al.* ²⁰.

AgrA and *icaA* virulence genes were amplified using qRT-PCR in accordance with the instructions provided by One-Step Kit (Bioline, UK). The StepOne RT-PCR thermal cycler (Applied Biosystem, USA) was used to set up the qRT-PCR analysis.

The forward and reverse gene-specific PCR primers for *AgrA*, and *icaA* were (5'-GGA GTG ATT TCA ATG GCA CA-3'; 5'-ATC CAT TTT ACT AAG TCA CCG ATT-3'), and (5'-CAATACTATTTTCGGGTGTCTTCACTCT-3; 5'-CAAGAAACTGCAATATCTTTCGGTAATCAT-3'), respectively ²¹. The relative expression values of each gene were normalized to the value of the housekeeping gene 16S rRNA. The forward and reverse primers used for 16S rRNA were 5'-TGT CGT GAG ATG TTG GG-3', and 5'-TGT CGT GAG ATG TTG GG-3', respectively ²¹. $2^{-\Delta\Delta CT}$ method was used to calculate the results ²². The results were reported as means $\pm$ SD of triplicate measurements.

2.5. Antioxidant Activity Evaluation

In vitro antioxidant activity of methanolic extracts of *S. rosmarinus* Spenn. and *O. vulgare* L. was evaluated using 2,2-diphenyl-1-picrylhydrazyl (DPPH) and ferric-reducing antioxidant power (FRAP) assays. Color change of reaction mixtures resulting from the antioxidant activity of the extracts was monitored using a UV spectrophotometer (UV-1601 PC, Shimadzu, Kyoto, Japan), as follows:

2.5.1. 2,2-Diphenyl-1-picrylhydrazyl assay (DPPH) assay

The free radical scavenging effect of the samples was evaluated as per the method previously described by Karaçelik *et al.* ²³. The reaction was performed by mixing the sample with DPPH at room temperature (25°C) and incubating the mixture in the dark for 30 min. The absorbance was then recorded at 516 nm. Quenching of the color intensity of the DPPH indicates the scavenging activity of the extracts. The free radical scavenging activities were expressed as percent inhibition using the equation:

$$\text{Inhibition (\%)} = \frac{\text{Absorbance of control} - \text{Absorbance of sample}}{\text{Absorbance of control}} \times 100$$

The measurements were performed in triplicate, and the results are presented as mean $\pm$ SD.

2.5.2. FRAP Assay

The ferric ion-reducing capacity of alcoholic extracts was measured kinetically following the method described by Benzie and Strain²⁴. The FRAP reagent was prepared by mixing 300 mM sodium acetate trihydrate buffer (pH 3.6), 10 mM TPTZ (2,4,6-tripyridyl-*s*-triazine) dissolved in 40 mM of hydrochloric acid, and 20 mM of ferric chloride at a ratio 10:1:1 v/v/v; respectively. Freshly prepared FRAP reagent (1.5 mL) was added to 50 µL of each extract (concentration = 0.1 mg/mL). Over three minutes, the absorbance change (resulting from the production of a ferrous ion blue color) was monitored at 593 nm. The results were expressed as µmol/L FeSO₄·7H₂O equivalent/mg of each extract.

2.6. Anti-Inflammatory Activity Evaluation

2.6.1. COX-II inhibition assay

Cyclooxygenase II (COX-II) Inhibitor Assay Kit from Abcam (the USA): ab211097 was used to evaluate the samples' inhibitory activity. The reaction conditions and method comply with the manufacturer's guidelines, using 10-fold serial dilutions (100, 10, 1, 0.1, 0.01 µg/mL). The process is based on detecting prostaglandin G2, a metabolite from arachidonic acid developed by COX-II activity, fluorometrically. Spectrofluorometer Tecan Spark (Tecan Group Ltd., Switzerland) was used to measure the fluorescence of the samples (Ex/Em= 535/587 nm) kinetically for 5-10 min at 25°C. The IC₅₀ was determined by plotting % inhibition of enzyme activity against sample concentrations. All sample assays were performed in triplicate.

2.6.2. Tumor necrosis factor-alpha (TNF-α) and Nuclear Factor kappa-B (NFκb) quantification

The macrophage cell line RAW 264.7, used for the *in vitro* anti-inflammatory assay, was obtained from the American Type Culture Collection (Manassas, Virginia, USA). The cells were cultivated in Dulbecco's Modified

Eagle's medium (DMEM) with 10 µg/mL insulin, 1 % penicillin-streptomycin, and 10 % fetal bovine serum at 37°C. The reagents were all of molecular biology grade.

The levels of tumor necrosis factor-alpha (TNF-α) and nuclear factor kappa-B (NF-κB) were quantitatively assessed in lipopolysaccharide (LPS) stimulated RAW 264.7 macrophages using enzyme-linked immunosorbent assay (ELISA) kits (human TNF-α ELISA Kit, Abcam, USA: ab181421; human NF-κB p100/NFκB2 ELISA Kit, Abcam, USA: ab288581). All procedures were conducted according to the manufacturers' protocols. For both assays, absorbance was measured at 450 nm using a microplate reader (BIOLINE ELISA Microplate Reader for TNF-α; ROBONIK P2000 ELISA Reader for NF-κB). The equivalent concentrations were determined from standard curves, and all measurements were carried out in triplicate.

2.7. Statistical analysis

All experiments were performed in triplicate, and the results are expressed as mean ± standard deviation (SD). Data was analyzed using GraphPad Prism version 8.0.1 (GraphPad Software Inc., California, USA). Statistical differences between two independent groups were assessed using an unpaired *t*-test. For analyses involving more than two groups, one-way ANOVA was applied, followed by Tukey's post hoc test for multiple comparisons. A *p*-value < 0.05 was considered statistically significant.

2.8. Multivariate data analysis

The data processing software MZmine 3.3 (available at <https://github.com/mzmine/mzmine3>) was employed for peak detection, deconvolution, deisotoping, and alignment of the imported mzXML files²⁵. This workflow produced an aligned peak list, which served as the basis of a detailed data matrix incorporating information from all samples (in triplicate). The negative ESI mode demonstrated greater sensitivity for a broader range of expected metabolite classes compared to the positive ESI mode²⁶. The data matrix included columns detailing the scan number,

retention time (t_R), mass-to-charge ratio (m/z), and peak intensity of the eluted compounds. Subsequently, the dataset was exported to SIMCA-P (version 14.1, Umetrics, Ume, Sweden) for Pareto scaling before multivariate analysis (MVA).

Partial least squares (PLS) analysis was applied to establish associations between bioactivities and the UPLC-QTOF-MS/MS dataset of annotated metabolites. Significant metabolites contributing to bioactivity were identified using variable importance in projection (VIP) scores derived from the PLS model. Additionally, Pearson's correlation coefficient (r) was calculated for correlation analysis, and a correlogram was generated to visually represent the strength of the correlations between metabolites and the evaluated bioactivity. This visualization was created using the MetaboAnalyst 5.0 platform (<https://metaboanalyst.ca/>). The thresholds for interpreting correlation coefficients were defined as follows: negligible correlation for $r < 0.3$, weak correlation for $r = 0.3-0.5$, moderate correlation for $r = 0.5-0.7$, strong correlation for $r = 0.7-0.9$, and very strong correlation for $r = 0.9-1.0$ ²⁷.

3. Results and discussion

3.1. UPLC-QTOF-MS/MS metabolite profiling of *Salvia rosmarinus* Spenn. and *Origanum vulgare* L. extracts

To identify the metabolites likely to mediate the aforementioned antimicrobial, antioxidant, and anti-inflammatory effects, UPLC-QTOF-MS/MS was used in both ionization modes (positive and negative) to provide comprehensive detection of metabolites in *S. rosmarinus* Spenn. and *O. vulgare* L. extracts. Base peak chromatograms (BPC) of *S. rosmarinus* Spenn. and *O. vulgare* L. extracts are presented in **Figure 1**. The analysis resulted in the detection of 164 compounds within 25 min. UPLC run, of which 92 were detected in both extracts, including hydroxycinnamic acid derivatives, flavonoid derivatives, benzoic acid derivatives, terpenes, and organic acids,

as summarized in **Table 1**. The chemical structures of the major classes of metabolites are shown in **Figure 2**. Details of the assigned metabolites are discussed in the following subsections.

Hydroxycinnamic Acid Derivatives

Twenty-eight hydroxycinnamic acid derivatives of caffeic, coumaric, ferulic acids, and danshensu (salvianic acid A or α -hydroxy dihydrocaffeic acid C₉H₁₀O₅) were detected. These derivatives showed common major fragment ions at m/z 197⁻ for deprotonated danshensu, m/z 179⁻ for caffeate, m/z 161⁻ for caffeoyl, and m/z 135⁻ for decarboxylated caffeate in MS² spectra.

Depsides, *i.e.*, hydroxycinnamic acid dimers or oligomers linked *via* ester bonds, are characteristic chemomarkers of the Lamiaceae family²⁸. They are recognized as bioactive phenolics for their antioxidant, antiallergic, immunomodulatory, and antimicrobial activities²⁹. Twelve depsides were identified herein, with salvianolic acid D (cpd. **7**) detected for the first time in RO, and both salvianolic acid K (cpd. **15**) and cleroden J (cpd. **33**) in *O. vulgare*.

Salvianolic acid D and rosmarinic acid are caffeic acid dimers. Salvianolic acid D (cpd. **7**, m/z 417.0791 [M-H]⁻, [C₂₀H₁₇O₁₀]⁻) showed a base peak in MS² spectra at m/z 219 [M-H-198]⁻ signifying the loss of a danshensu moiety (**Supplementary figure 1S**)³⁰. Rosmarinic acid-*O*-hexoside (cpd. **16**, m/z 521.1285 [M-H]⁻, [C₂₄H₂₅O₁₃]⁻), rosmarinic acid (cpd. **18**, m/z 359.0766 [M-H]⁻, [C₁₈H₁₅O₈]⁻) and methylrosmarinic acid (cpd. **30**, m/z 373.0917 [M-H]⁻, [C₁₉H₁₇O₈]⁻) shared the same fragment ions at m/z 197 and 179 in MS² spectra, corresponding to deprotonated danshensu and caffeate moieties, respectively (**Supplementary figure 2S-4S**)³¹.

Cleroden J (cpd. **33**, m/z 553.1339 [M-H]⁻, [C₂₈H₂₅O₁₂]⁻) was detected in *O. vulgare* L. for the first time and has previously been isolated from other species within the same family³². The spectrum was dominated by a base peak at m/z 135, corresponding to the decarboxylated caffeic acid fragment [caffeic acid-H-CO₂]⁻. Additional product ions were observed at m/z 521 arising from the loss of a methoxy group [M-H-OCH₃]⁻, and at m/z 477,

corresponding to subsequent decarboxylation $[M-H-OCH_3-CO_2]^-$. A further significant ion at m/z 179 $[C_9H_7O_4]^-$ confirms the presence of caffeic acid units within the structure (**Supplementary figure 5S**).

Caffeic acid trimers *vis.* salvianolic acids A and K were identified. Salvianolic acid K (cpd. **15**, m/z 555.1129 $[M-H]^-$, $[C_{27}H_{23}O_{13}]^-$) revealed daughter ions at m/z 511 $[M-H-44]^-$ and 357 $[M-H-198]^-$ after the losses of carboxyl group and danshensu moieties (**Supplementary figure 6S**), respectively³³. The fragmentation pattern of this compound is presented as a representative example of the fragmentation pathway for this class and is illustrated in **Figure 3**. Salvianolic acid A (cpd. **27**, m/z 493.1127 $[M-H]^-$, $[C_{26}H_{21}O_{10}]^-$) yielded a major fragment ion at m/z 295 $[M-H-198]^-$, indicating the loss of a danshensu moiety (**Supplementary figure 7S**).

Salvianolic acid B and sagerinic acid are caffeic acid tetramers (rosmarinic acid dimers). Salvianolic acid B is formed by oxidative cyclization of two rosmarinic acid molecules, giving a 1,2-dihydronaphthalene ring structure, while sagerinic acid is formed by dimerization of 2 rosmarinic acid molecules and cyclobutane ring formation³⁴. Salvianolic acid B (cpd. **20**, m/z 717.1437 $[M-H]^-$, $[C_{36}H_{29}O_{16}]^-$), produced a base peak ion at m/z 359 $[C_{18}H_{15}O_8]^-$ representing rosmarinic acid, and the less abundant product ions that indicate sequential losses of two danshensu: at m/z 519 $[M-H-198]^-$, and m/z 321 $[M-H-198-198]^-$ (**Supplementary figure 8S**)³⁵.

Sagerinic acid (cpd. **21**, m/z 719.1602 $[M-H]^-$, $[C_{36}H_{31}O_{16}]^-$) generated a base peak fragment ion at m/z 359 $[M-H-360]^-$, resulting from molecular splitting $[M/2]^-$, corresponding to rosmarinic acid and another ion with lower intensity at m/z 197 $[C_9H_9O_5]^-$ corresponding to the danshensu moiety (**Supplementary figure 9S**)³⁴.

Flavonoids

Flavonoids comprised the major class of secondary metabolites in the alcoholic profiles of both *S. rosmarinus* Spenn. and *O. vulgare* L., where a total of 56 flavonoid derivatives were detected in this study. These included

representatives of flavanones, flavonols, and flavones, among which five compounds in *O. vulgare* L. (nepitrin; cpd. **60**, hispidulin-*O*-rutinoside; cpd. **63**, luteolin-*O*-pentosyl-acetyl-hexoside; cpd. **64**, luteolin caffeoylhexoside; cpd. **67**, and acacetin-*O*-rutinoside; cpd. **71**) and one compound in *S. rosmarinus* Spenn. (quercetin coumaroylhexoside; cpd. **96**) were identified for the first time in these species. The elution of flavonoids appeared to follow a decreasing polarity sequence in chromatograms. So firstly, at the R_t range (9.44-11.92 min.), flavonoid diglycosides were detected. Apigenin-6,8-di-*C*-hexoside (cpd. **52**, m/z 593.1491 [M-H]⁻, [C₂₇H₂₉O₁₅]⁻) was annotated after showing the losses of 120 amu (m/z 473) and 90 amu (m/z 503), indicative of a *C*-linked hexoside internal cleavage, and the losses of 240 (-2×120) amu (m/z 353) and 210 (-120-90) amu (m/z 383) that also indicate the internal cleavages that occur in both *C*-linked hexoside (**Supplementary figure 10S**)³⁶. Rutin, quercetin-*O*-rutinoside, (cpd. **92**, m/z 611.1602 [M+H]⁺, [C₂₇H₃₁O₁₆]⁺) was confirmed by the MS² spectra, which displayed fragment ions at m/z 465 and m/z 303, denoting the loss of a deoxyhexose sugar [M+H-146]⁺ followed by the loss of a hexose sugar [M+H-146-162]⁺, respectively (**Supplementary figure 11S**)³⁷. Luteolin-*O*-rutinoside (cpd. **56**, m/z 593.1495[M-H]⁻, [C₂₇H₂₉O₁₅]⁻) yielded a characteristic base peak of m/z 285 for the luteolin fragment after the loss of 308 (-162-146) amu of the rutinoside moiety (**Supplementary figure 12S**) Luteolin-*O*-pentosyl-hexoside (cpd. **61**, m/z 581.1498 [M+H]⁺, [C₂₆H₂₉O₁₅]⁺), showed a base peak ion in the MS² spectra at m/z 287 [M+H-162-132]⁺, suggesting the respective losses of hexose and pentose (**Supplementary figure 13S**). Apigenin-*O*-rutinoside (cpd. **62**, m/z 579.1693 [M+H]⁺, [C₂₇H₃₁O₁₄]⁺) represented major fragment ions at m/z 433 [M+H-146]⁺ and 271 [M+H-146-162]⁺, indicating the loss of a deoxyhexose sugar followed by a hexose sugar, respectively (**Supplementary figure 14S**)³⁸.

Flavonoid monoglycosides were identified in the following section of the chromatographic elution at R_t range of 9.85-12.08 min. Nepitrin (6-methoxyluteolin-*O*-hexoside) (cpd. **60**, m/z 479.1183 [M+H]⁺, [C₂₂H₂₃O₁₂]⁺)

displayed a prominent product ion in MS² at m/z 317 appeared after the loss of a hexose [M+H-162]⁺ moiety and another fragment at m/z 302 appeared after the losses of a hexose and methyl group [M+H-162-15]⁺ (**Supplementary figure 15S**). Apigenin-*O*-hexoside (cpd. **65**, m/z 433.1128 [M+H]⁺, [C₂₁H₂₁O₁₀]⁺) and apigenin-*O*-hexuronide (cpd. **66**, m/z 447.0921 [M+H]⁺, [C₂₁H₁₉O₁₁]⁺) are monoglycoside flavones, both demonstrated a similar base peak ion at m/z 271 [C₁₅H₁₁O₅]⁺, corresponding to apigenin aglycone after the losses of hexose and hexuronide moieties, respectively, and minor daughter ions at m/z 153 and 109 resulting from Retro Diels Alder (RDA) reactions (**Supplementary figure 16S, 17S**). Homoplantagin (Hispidulin-*O*-hexoside) (cpd. **75**, m/z 463.1235 [M+H]⁺, [C₂₂H₂₃O₁₁]⁺) showed a base peak ion at m/z 301 [M+H-162]⁺ after the loss of a hexose moiety and another fragment ion at m/z 286 [M+H-162-15]⁺ was produced due to the loss of a hexose followed by methyl moiety, that matched characteristic fragments of hispidulin (**Supplementary figure 18S**).

Flavonoid aglycones were detected in the chromatograms after R_t (10.48 min.). Eriodictyol (cpd. **103**, m/z 287.0553 [M-H]⁻, [C₁₅H₁₁O₆]⁻) gave fragment ions resulting from aglycone C-ring RDA cleavages, showing a base peak ion at m/z 135 (^{1,3}B⁻), a major fragment ion at m/z 151 (^{1,3}A⁻) and minor fragment ions at m/z 107 and m/z 125 for (^{1,3}A-CO₂) and (^{1,4}A⁻), respectively (**Supplementary figure 19S**)³⁹ Similarly, apigenin aglycone (cpd. **77**, m/z 269.0451 [M-H]⁻, [C₁₅H₉O₅]⁻), demonstrated the MS² data that illustrated the RDA fragment ions at m/z 151 (^{1,3}A⁻), m/z 149 (^{1,4}B⁻+2H), m/z 117 (^{1,3}B⁻) and 107 (^{1,3}A-CO₂) due to C-ring cleavage in addition to other fragment ions resulting from small losses *vis.* m/z 225 and m/z 201 consecutive to CO₂ and C₃O₂ losses (**Supplementary figure 20S**)^{38,39}. Isorhamnetin; *O*-methylquercetin, (cpd. **97**, m/z 315.0506 [M-H]⁻, [C₁₆H₁₁O₇]⁻) was identified based on the distinctive MS², where the loss of a methyl group (30 amu) was noted to produce a base peak ion at m/z 300 and minor fragment ions at m/z 271 and m/z 243 corresponding to the loss of CO₂ and CO₂+CO groups, respectively (**Supplementary figure 21S**)⁴⁰. Hydroxygenkwanin (cpd. **82**,

m/z 299.0554 [M-H]⁻, [C₁₆H₁₁O₆]⁻) was suggested as on dissociation produce fragment ions at m/z 284 [M-H-15]⁻ denoting the loss of a methyl, m/z 256 [M-H-15-28]⁻ indicating the losses of methyl and CO groups, m/z 227 [M-H-15-28-29]⁻ inferring the losses of methyl, CO and CHO groups and m/z 151 (^{1,3}A⁻-CH₃) and m/z 133 (^{1,3}B⁻) due to RDA cleavage of C-ring (**Supplementary figure 22S**)⁴¹. Sakuranetin (cpd. **107**, m/z 285.0760 [M-H]⁻, [C₁₆H₁₃O₅]⁻) had a characteristic MS² showing major fragment ions at m/z 165 and 119 resulting from RDA cleavage of the C-ring at (^{1,3}A⁻) and (^{1,3}B⁻), respectively (**Supplementary figure 23S**)⁷. Salvigenin (cpd. **90**, m/z 329.1019 [M+H]⁺, [C₁₈H₁₇O₆]⁺) showed fragment ions at m/z 314 [M+H-15]⁺ corresponding to the loss of a methyl group, m/z 296 [M+H-15-18]⁺ indicating the a further loss of water molecule and a minor fragment ion at m/z 268 [M+H-15-18-28]⁺ due to the collective loss of methyl, water and CO groups (**Supplementary figure 24S**)^{42,43}.

Terpenes

A total of 42 metabolites belonging to monoterpenes, diterpenes, sesquiterpenoids, and triterpenes were detected and mainly eluted at the middle and late sections of the chromatogram (R_t = 8.4-22.4 min.). Abietenes are phenolic diterpenes of limited distribution in some species of the Lamiaceae family⁴⁴. These compounds have various biological actions, including antioxidant, anti-inflammatory, and anti-microbial properties⁴⁵. Hydroxy-*O*-methylosmanol (cpd. **116**, m/z 375.1803 [M-H]⁻, [C₂₁H₂₇O₆]⁻) generated a major fragment ion at m/z 299 [M-H-44-31-1]⁻ after the losses of CO₂ and OCH₃ and molecular rearrangement, besides the daughter ions at m/z 345 [M-H-30]⁻ [C₂₀H₂₅O₅]⁻ for the rosmanol fragment, in addition to m/z 331 [M-H-44]⁻ and m/z 316 [M-H-44-15]⁻ attributed to the losses of CO₂ and CH₃, respectively (**Supplementary figure 25S**)⁴⁶. Rosmadial (cpd. **130**, m/z 343.1544 [M-H]⁻, [C₂₀H₂₃O₅]⁻) gave fragments at m/z 315 [M-H-28]⁻ for a loss of CO, and m/z 299 [M-H-44]⁻ suggesting a cleavage of CO₂ (**Supplementary figure 26S**)⁴⁷. Tetrahydro-hydroxyrosmariquinone (cpd. **133**, m/z 301.1802

[M-H]⁻, [C₁₉H₂₅O₃]⁻) was assigned after revealing product ions; *m/z* 283 [M-H-18]⁻, 273 [M-H-28]⁻, and 258 [M-H-43]⁻ following the losses of H₂O, CO, and (CH(CH₃)₂), respectively (**Supplementary figure 27S**) and in accordance with previous reported data ⁴⁷. Carnosic acid (cpd. **137**, *m/z* 331.1904 [M-H]⁻, [C₂₀H₂₇O₄]⁻) revealed a major fragment ion at *m/z* 287 [M-H-44]⁻ after decarboxylation that was followed by the loss of an isopropyl group (CH(CH₃)₂) yielding a fragment ion at *m/z* 244 [M-H-44-43]⁻ (**Supplementary figure 28S**). *O*-methylecarnosic acid (cpd. **138**, *m/z* 345.2063 [M-H]⁻, [C₂₁H₂₉O₄]⁻) on fragmentation showed two major ions at *m/z* 301 and *m/z* 286, corresponding to the losses of CO₂ [M-H-44]⁻ and a successive loss of methyl group [M-H-44-15]⁻, and fragment ion with low intensity at *m/z* 271 resulted from the further loss of a methyl group [M-H-44-15-15]⁻ (**Supplementary figure 29S**) ⁴⁸.

Six pentacyclic triterpenes were identified in alcoholic extracts of *S. rosmarinus* Spenn. and *O. vulgare* L., among them oleanolic acid, ursolic acid, and corosolic acid, which possess documented anti-inflammatory and antioxidant properties ⁴⁹. Asiatic acid (cpd. **143**, *m/z* 487.3416 [M-H]⁻, [C₃₀H₄₇O₅]⁻) showed characteristic fragment ions in its MS² spectra at *m/z* 469 [M-H-18]⁻ after dehydration, *m/z* 441 [M-H-46]⁻ corresponding to the loss of HCOOH, and *m/z* 409 [M-H-46-32]⁻ by the respective losses of HCOOH and CH₃OH (**Supplementary figure 30S**).

3.2. Antibacterial activity

3.2.1. Evaluation of minimum inhibitory concentrations (MIC) and biofilm formation inhibition

The extracts exhibited variable antibacterial activities against MRSA and *E. coli* **Table 2**. The methanolic extract of *S. rosmarinus* Spenn. and the aqueous extract of *O. vulgare* L. exhibited the strongest antibacterial activities (MIC = 64 µg/mL against both strains).

Biofilm development presents a significant challenge in microbial infections because of its role in increasing resistance. Therefore, the most

active extracts were tested for their ability to inhibit biofilm formation in MRSA. In our study, RO and Aq.OV, which showed the strongest antibacterial activity, were further examined for their antibiofilm formation activity. RO and Aq.OV extracts showed significant inhibition at $\frac{1}{4}$ MIC, reducing biofilm formation by $54.9 \pm 2.3\%$ and $61.4 \pm 0.6\%$, respectively, and an unpaired t-test confirmed a statistically significant difference between them ($p = 0.0091$). Based on established criteria ⁵⁰, a plant extract achieving a biofilm inhibition level greater than 50% is generally regarded as indicative of good antibiofilm activity. Accordingly, the inhibition levels observed in our study indicate that these extracts may represent promising alternatives to conventional antibiotics for managing MRSA infections, especially in biofilm-associated cases where standard therapies often fail due to the protective biofilm ⁵¹.

3.2.2. Reduction of *icaA* and *agrA* gene expression levels

The *icaA* and *agrA* are important genes that play crucial roles in virulence and biofilm development in MRSA; the *icaA* gene is responsible for biofilm formation and the *agrA* gene is responsible for quorum sensing of MRSA ⁵². The *icaA* was downregulated by 30% and 60% after treatment of MRSA with $\frac{1}{4}$ MIC of RO and Aq.OV extracts, respectively. Similarly, the *agrA* gene was also downregulated by 43% and 70% after treating MRSA with $\frac{1}{4}$ MIC of RO and Aq.OV extracts, respectively (**Figure 4**). For both *icaA-1* and *agrA* genes, Tukey's multiple comparisons test demonstrated statistically significant differences among all groups (RO extract, Aq.OV extract, and control) ($p < 0.0001$). These results are consistent with the observed antibiofilm activity, indicating that the extracts interfere with critical genetic regulators essential for biofilm formation.

3.3. Antioxidant activity

The antioxidant activity was assessed using complementary *in vitro* assays based on different mechanisms, including the DPPH radical scavenging assay and the FRAP reducing power assay.

The results revealed that the methanolic extract of *S. rosmarinus* Spenn. exhibited a significantly stronger radical scavenging capacity in the DPPH assay ($IC_{50} = 6.56 \pm 0.035 \mu\text{g/mL}$) compared to the methanolic extract of *O. vulgare* L. ($IC_{50} = 17.60 \pm 0.333 \mu\text{g/mL}$ (Unpaired Student's *t*-test, $p < 0.0001$) (**Table 3**). In contrast, *O. vulgare* L. exhibited a significantly higher reducing capacity in the FRAP assay ($188.61 \pm 24.06 \mu\text{mol/L}$) compared to *S. rosmarinus* Spenn. ($122.78 \pm 2.09 \mu\text{mol/L}$) (Unpaired Student's *t*-test, $p = 0.0004$) (**Table 3**).

3.4. Anti-inflammatory activity

Inflammation is a complex biological response that serves as both an indicator of pathological disturbances and a contributor to disease progression as a result of sustained disruption of inflammatory regulation⁵³. Pro-inflammatory cytokines, such as TNF- α , are released in response to inflammatory stimuli, which activate NF- κ B signaling and subsequently induce the expression of COX-II⁵³. In this study, the anti-inflammatory activity of the methanolic extracts of *S. rosmarinus* Spenn. and *O. vulgare* L. was evaluated through COX-II inhibition, as well as the suppression of TNF- α and NF- κ B, using ibuprofen as a standard. *S. rosmarinus* Spenn. extract demonstrated strong anti-inflammatory potential, as evidenced by its COX-II inhibitory effect ($IC_{50} = 11.09 \pm 0.46 \mu\text{g/mL}$), which was comparable to ibuprofen ($IC_{50} = 8.79 \pm 0.37 \mu\text{g/mL}$), with no statistically significant difference between them (Tukey's post hoc test, $p < 0.595$). In parallel, *S. rosmarinus* Spenn. extract significantly suppressed key inflammatory mediators, reducing TNF- α and NF- κ B levels to 0.385 and 0.31 fold, respectively, of the control values that were close to those achieved by ibuprofen, with significant differences observed for TNF- α ($p = 0.0013$) and NF- κ B ($p = 0.0102$). Conversely, *O. vulgare* L. extract exhibited noticeably weaker anti-inflammatory effects across all tested markers, showing significantly lower efficacy than ibuprofen ($p < 0.0001$) (**Table 4**).

3.5. Multivariate Data Analysis and Pearson's Correlation Study

The PLS model, as a supervised approach, was adopted to determine the relationship between the identified metabolites (**Table 1**) and the results of the biological investigations of *S. rosmarinus* Spenn. and *O. vulgare* L. extracts, utilizing the UPLC-QTOF-MS/MS dataset as X variables and the Y variables were the antibacterial activity represented as 1/MIC values against *E. coli* and MRSA strains, the antioxidant activity represented as 1/IC₅₀ values of DPPH radical scavenging and the values of the ferric ion reducing capacity of the extract (FRAP), and the anti-inflammatory activity represented by the down-regulation levels of the inflammation biomarkers calculated as 1/COX-II, 1/NF- κ B, and 1/TNF- α .

PLS Model was validated by the quality of fitness and prediction of Y. The autofit of the PLS model demonstrated excellent fit (R^2_Y cum = 0.996) and predictive capability (Q^2 cum = 1), suggestive of a strong model with no overfit. The PLS biplot, which combines score and loading charts, was used to visually represent the relation between the samples and variables contributing to differentiation across the extracts (**Figure 5**). The proximity of the X and Y variables to the sample clusters signifies their degree of contribution to the defining traits of each cluster. **Figure 5** illustrates that the *S. rosmarinus* Spenn. extract was segregated from the *O. vulgare* L. extract by the first latent variable (LV1). This may be attributed to differences in metabolite abundances.

Moreover, the bacterial inhibition activity and the DPPH free radical scavenging activity were positioned alongside the *S. rosmarinus* Spenn. extract, signifying their superior activities relative to the *O. vulgare* L. extract. Notably, diterpenes (*i.e.*, oridonin, carnosic acid, hydroxyrosmanol, rosmic acid, dimethoxyrosmanol, rosmadial, and tetrahydro-7-hydroxyrosmariquinone), triterpenes (*i.e.*, asiatic acid, corosolic acid, and 3-oxours-12-en-20,28-olide), hydroxycinnamic acids (*i.e.*, yunnaneic acid F, salvianolic acid K, and rosmarinic acid), and flavonoid derivatives (*i.e.*, nepitrin, isorhamnetin, and hesperetin) were found to be enriched in the *S.*

rosmarinus Spenn. extract. This demonstrates their beneficial effects on the DPPH free radical scavenging and bacterial inhibitory activities in *S. rosmarinus* Spenn. extract. The strong antibacterial and DPPH free radical scavenging actions of these metabolites can be attributed to the existence and quantity of phenolic hydroxyl groups ⁵⁴. In contrast, the *O. vulgare* L. extract had minimal projection towards those Y variables, positioned on the positive side of the latent variable (LV1). This observation indicated a lesser correlation of the *O. vulgare* L. extract with the examined bioactivities. These findings are aligned with results from the evaluation of the antibacterial activity study and evaluation of the free radical scavenging activity (DPPH assay), where *S. rosmarinus* Spenn. extract showed stronger bacterial inhibition than *O. vulgare* L. extract (**Table 2**). Also, *S. rosmarinus* Spenn. extract exhibited the highest scavenging capacity of DPPH• radical with the lowest IC₅₀ value, followed by *O. vulgare* L. extract, indicating strong antioxidant activity of the extract (**Table 3**).

However, antioxidant activity investigated as ferric ion reducing capacity (FRAP) was positioned alongside the *O. vulgare* L. extract, signifying their superior activities relative to *S. rosmarinus* Spenn. extract. These findings indicated that *O. vulgare* L. extract exhibited an antioxidant activity due to their enrichment in benzyl derivatives (*i.e.*, hydroxyphenol hexoside, hydroxyphenol hexoside derivative, and origanine B or C), hydroxycinnamic acids (*i.e.*, lithospermic acid derivative), and organic acids and esters (*i.e.*, oxoadipic acid, corchorifatty acid F, malic acid, and citric acid) were found to be enriched in the *O. vulgare* L. extract. The enrichment of these compounds may account for their notable ferric ion-reducing capacity and overall antioxidant activity. These constituents are known for their redox properties, free radical scavenging ability, and metal ion chelation, which collectively contribute to the observed antioxidant potential ⁸. Regarding the anti-inflammatory activity of both extracts assessed by inhibiting COX-II activity and modulating NF- κ B, and TNF- α pathways, those Y variables showed proximity to the extract of *S. rosmarinus* Spenn., indicating a direct

correlation. *O. vulgare* L. exhibited less projection, suggesting less correlation. *S. rosmarinus* Spenn. extract is richer in flavonoid derivatives (*i.e.*, luteolin-acetyl-*O*-hexuronide, apigenin, dimethylquercetin, and cirsimaritin), hydroxycinnamic acids (*i.e.*, salvianolic acid B and lithospermic acid), and diterpenes (*i.e.*, rosmanol, hydroxy-*O*-methylosmanol, hydroxyrosmadial, and carnosol), which are well recognized for their anti-inflammatory effects ⁵⁵. These findings are consistent with the anti-inflammatory activity of the extracts, where the *S. rosmarinus* Spenn. extract showed stronger anti-inflammatory activity than the *O. vulgare* L. extract (**Table 4**). Previous reports highlighted the importance of flavonoid derivatives and diterpenes as anti-inflammatory metabolites. For instance, carnosic acid and carnosol exert anti-inflammatory effects primarily by inhibiting the NF- κ B, MAPK, STAT3, and NLRP3 inflammasome pathways, leading to reduced expression of pro-inflammatory cytokines such as TNF- α , IL-1 β , and IL-6. They also activate SIRT1, which further suppresses inflammation by downregulating these signaling cascades ⁵⁶. Additionally, flavonoids such as luteolin and apigenin have been proven to exhibit anti-inflammatory effects mainly by inhibiting the transcriptional activity of NF- κ B without affecting its upstream signaling. They also slightly reduce JNK activation and suppress the expression of pro-inflammatory chemokines, contributing to their overall anti-inflammatory action ⁵⁷.

The correlation analysis was confirmed by calculating Pearson's correlation coefficient ($r \geq 0.7$), with statistical significance set at $p < 0.05$ and a false discovery rate (FDR) < 0.08 . A correlogram (**Figure 6**) was generated to visualize the strength of correlations among the variables. Notably, the analysis focused on metabolites with a PLS variable importance in projection (VIP) score of 1 or higher (**Figure 7**), leading to the selection of 66 annotated metabolites (**Figure 6**). The correlogram revealed strong positive correlations between the examined anti-microbial activity, DPPH radical scavenging activity, and hydroxycinnamic acid derivatives, diterpenes, triterpenes, and flavonoid derivatives, mainly present in *S.*

651 *rosmarinus* Spenn. extract, namely, oridonin, carnosic acid,
652 hydroxyrosmanol, rosmic acid, dimethoxyrosmanol, rosmadial, tetrahydro-7-
653 hydroxyrosmariquinone, asiatic acid, corosolic acid, 3-oxours-12-en-20,28-
654 olide, Yunnaneic acid F, salvianolic acid K, rosmarinic acid, nepitrin,
655 isorhamnetin, and hesperetin. The antimicrobial activities of carnosic acid
656 and rosmarinic acid are well-documented in the literature and appear to
657 involve multiple complementary mechanisms. Carnosic acid has been
658 reported to act as a potential quorum-sensing inhibitor, thereby suppressing
659 bacterial virulence and biofilm formation. Additionally, its lipophilic nature
660 allows it to incorporate into bacterial membranes, leading to membrane
661 destabilization, increased permeability, and potential cell lysis ⁵⁸. Similarly,
662 rosmarinic acid exerts its antibacterial effects primarily through membrane
663 disruption, inhibition of efflux pumps, interference with essential bacterial
664 enzymes, and suppression of biofilm formation, with particularly pronounced
665 activity against Gram-positive bacteria. These mechanisms collectively
666 support the strong antimicrobial potential of these phenolic compounds,
667 particularly in targeting persistent and resistant bacterial strains ⁵⁹.
668 Additionally, benzyl derivatives, organic acids & esters, and hydroxycinnamic
669 acids are mainly present in *O. vulgare* L. extract exhibited strong positive
670 correlations with the antioxidant activity investigated as FRAP, including
671 hydroxyphenol hexoside, hydroxyphenol hexoside derivative, origanine B or
672 C, lithospermic acid derivative, cleroden J, oxoadipic acid, and corchorifatty
673 acid F. Lithospermic acid has been proven to possess significant antioxidant
674 activity, primarily through its ability to scavenge free radicals. By
675 neutralizing reactive oxygen species (ROS), it plays a crucial role in
676 protecting cellular components from oxidative damage. These documented
677 mechanisms highlight its potential as a therapeutic agent against oxidative
678 stress-related diseases, reinforcing its value as a key natural antioxidant in
679 plant-based systems ⁶⁰. Also, the investigated anti-inflammatory activities
680 were strongly positively correlated with flavonoid derivatives, diterpenes,

and hydroxycinnamic acids. Pearson's correlation coefficient analysis strongly supported the PLS analysis findings.

Conclusion

This study provides the first evidence linking specific metabolites to some of the most important pharmacological effects of two Lamiaceae species, *Salvia rosmarinus* and *Origanum vulgare*. While *S. rosmarinus* Spenn. exhibited superior anti-inflammatory and radical scavenging activities, *O. vulgare* L. showed stronger reducing power. PLS and Pearson's correlation coefficients confirmed the strong positive correlation between different hydroxycinnamic acids derivatives (*i.e.*, Yunnaneic acid F, salvianolic acid K, and rosmarinic acid), diterpenes (*i.e.*, oridonin, carnosic acid, hydroxyrosmanol, rosmic acid, dimethoxyrosmanol, rosmadial, and tetrahydro-7-hydroxyrosmariquinone), triterpenes (*i.e.*, asiatic acid, corosolic acid, and 3-oxours-12-en-20,28-olide), and flavonoid derivatives (*i.e.*, nepitrin, isorhamnetin, and hesperetin) mainly present in *S. rosmarinus* Spenn. extract with strong antibacterial and DPPH radical scavenging activities. Moreover, the anti-inflammatory activity was positively correlated with flavonoid derivatives (*i.e.*, luteolin-acetyl-*O*-hexuronide, apigenin, dimethylquercetin, and cirsimaritin), diterpenes (*i.e.*, rosmanol, hydroxy-*O*-methylosmanol, hydroxyrosmadial, and carnosol), and hydroxycinnamic acids (*i.e.*, salvianolic acid B and lithospermic acids). The antioxidant activity evaluated as FRAP reducing power was positively correlated with benzyl derivatives (*i.e.*, hydroxyphenol hexoside, hydroxyphenol hexoside derivative, and origanine B or C), organic acids and esters (*i.e.*, oxoadipic acid, corchorifatty acid F, malic acid, and citric acid), and hydroxycinnamic acids (*i.e.*, lithospermic acid derivative and cleroden J). This study can be applied to a wide range of medicinal plants, driving a deeper exploration of the correlation between bioactivities and metabolite composition. In turn, this paves the way for a profound understanding of mechanisms of action, pharmacokinetics, and structure-activity relationships, and advances the development of herbal

medicine practice. Bioinformatic tools also enable this approach to be expanded toward other therapeutic targets.

Author contributions

Alaa Y. Zakarya: Formal analysis, Data interpretation, Methodology, Data curation, Investigation, Writing - original draft. **Dalia M. Rasheed, Ayat M. Emad:** Conceptualization, Supervision, Visualization, Writing, reviewing & editing. **Mohamed A. Farag:** Methodology, Visualization, Writing, reviewing & editing. **Zeinab M. Goda:** Software, Data curation, Formal analysis, Writing - original draft. **Omnia Karem M Riad, Sally T K Tohamy:** Methodology, Data curation, Formal analysis, Writing - original draft. **Ahmed H. Elbanna:** Conceptualization, Supervision, Visualization, Writing, reviewing & editing.

Conflicts of interest

The authors declare that there are no personal relationship or financial conflicts of interest, that could inappropriately influence the integrity of this work.

Funding

No funding was received to assist with conducting this study or preparation of this manuscript.

Data availability

All data generated or analyzed during this study are included in this published article and its supplementary information file

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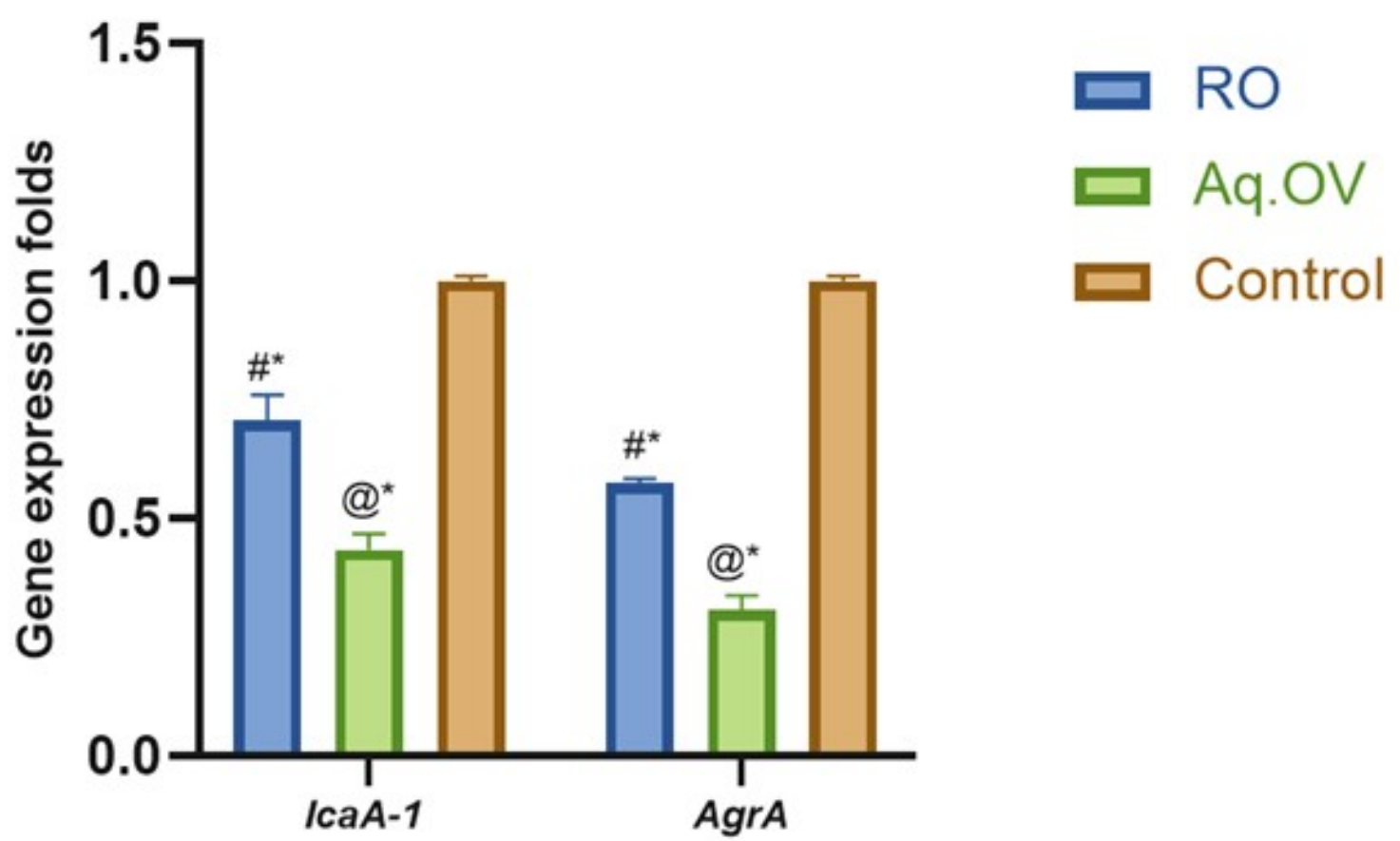
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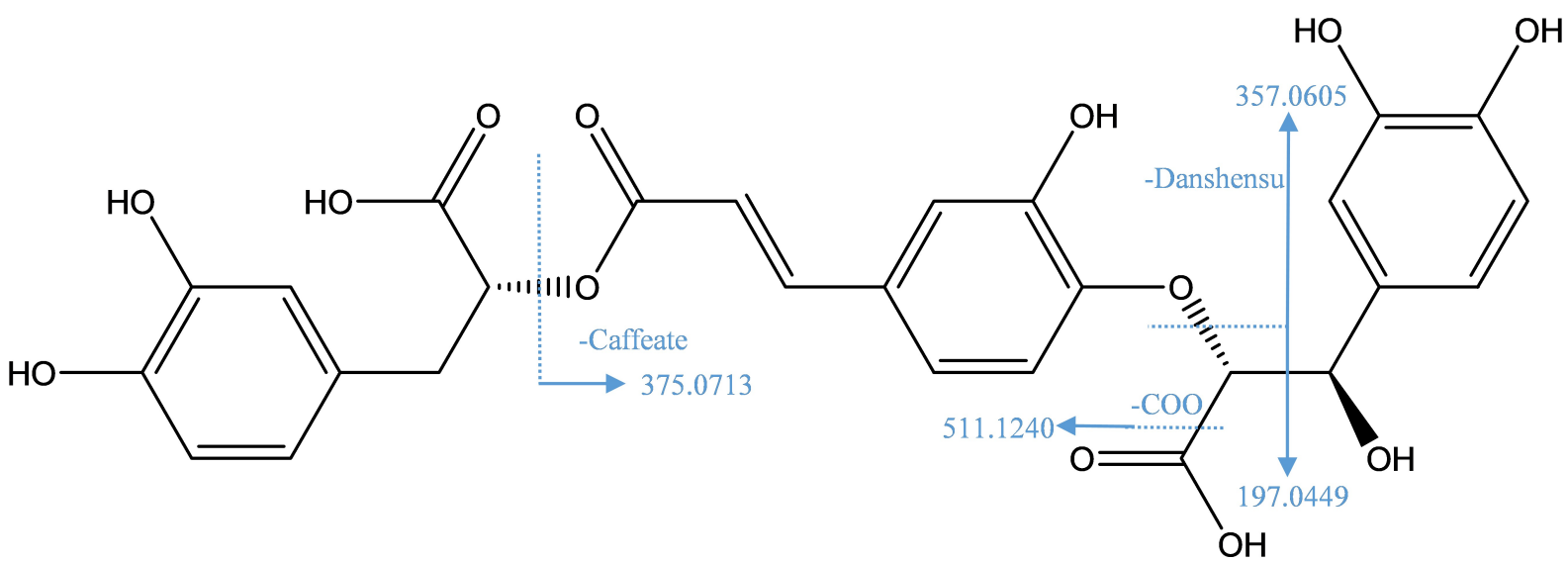
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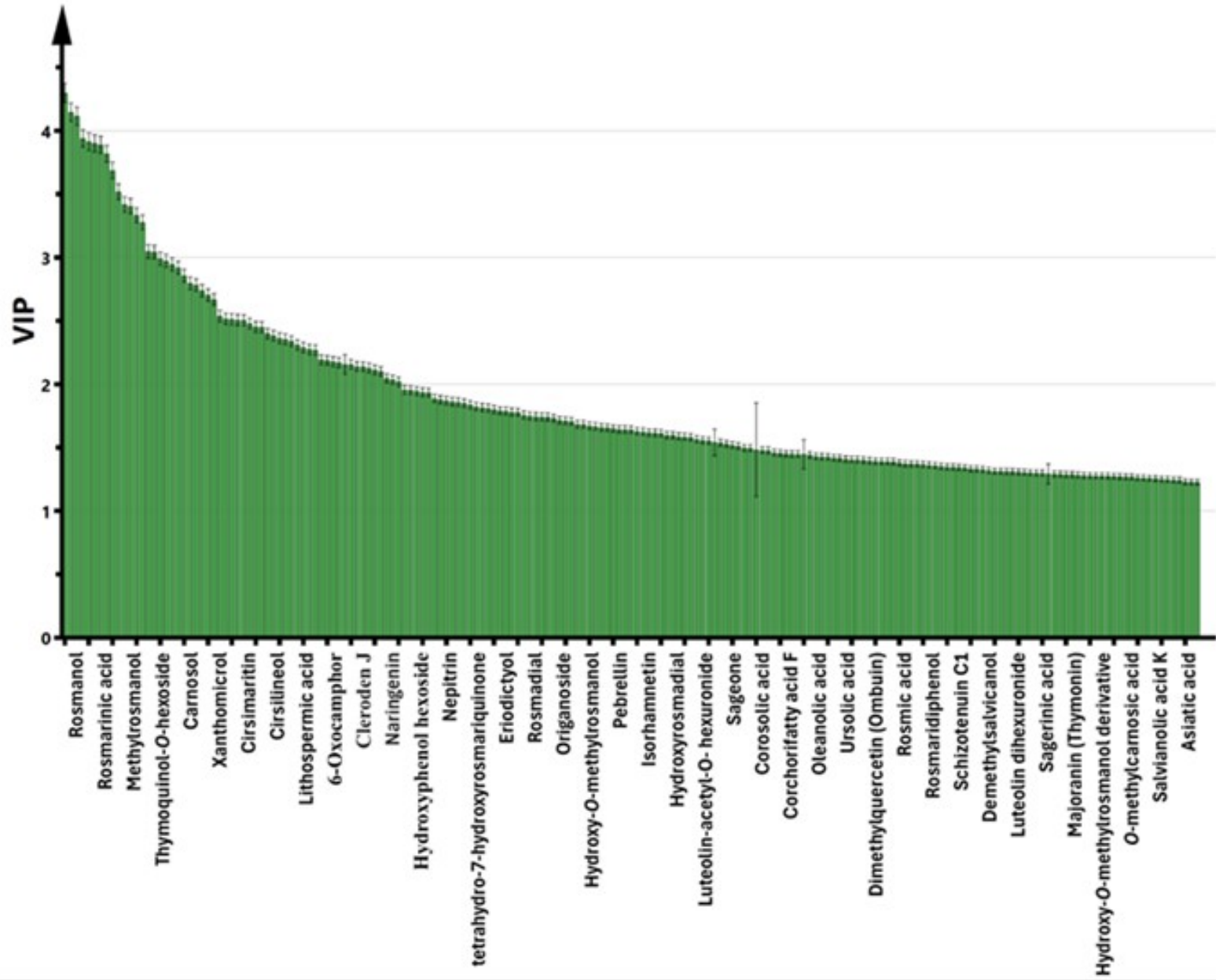
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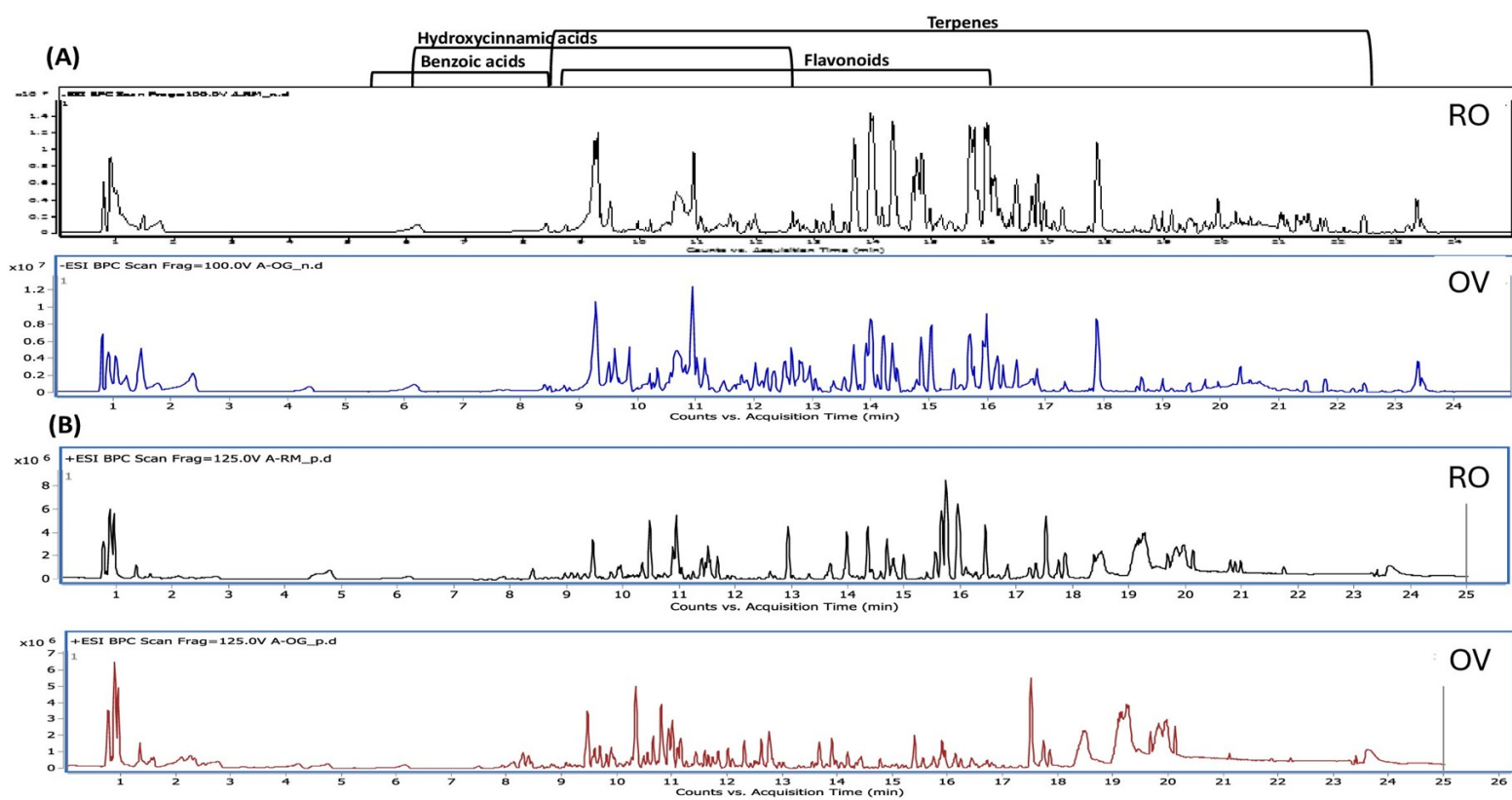
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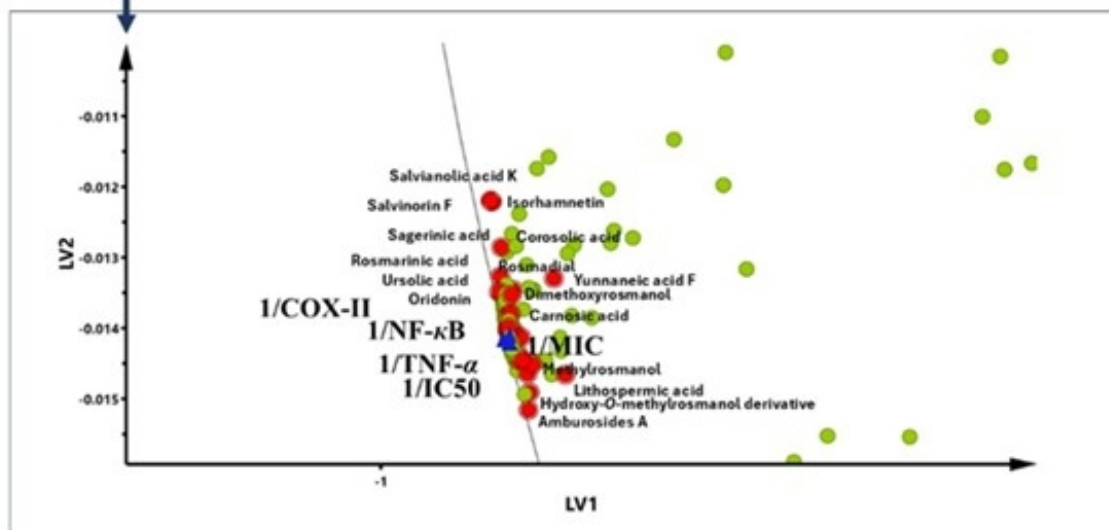
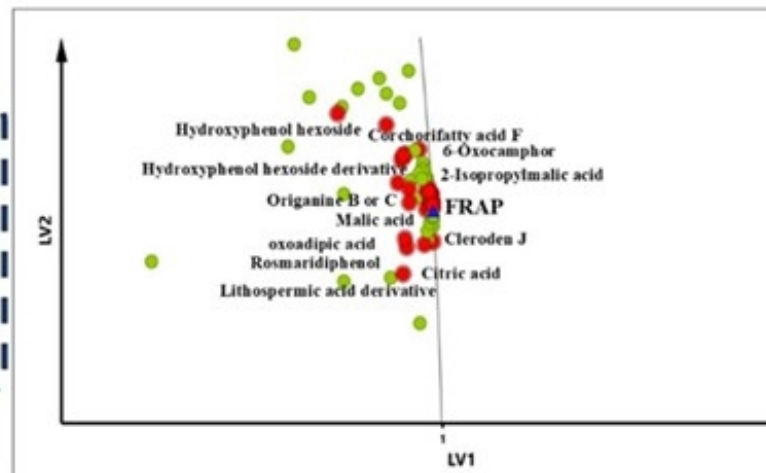
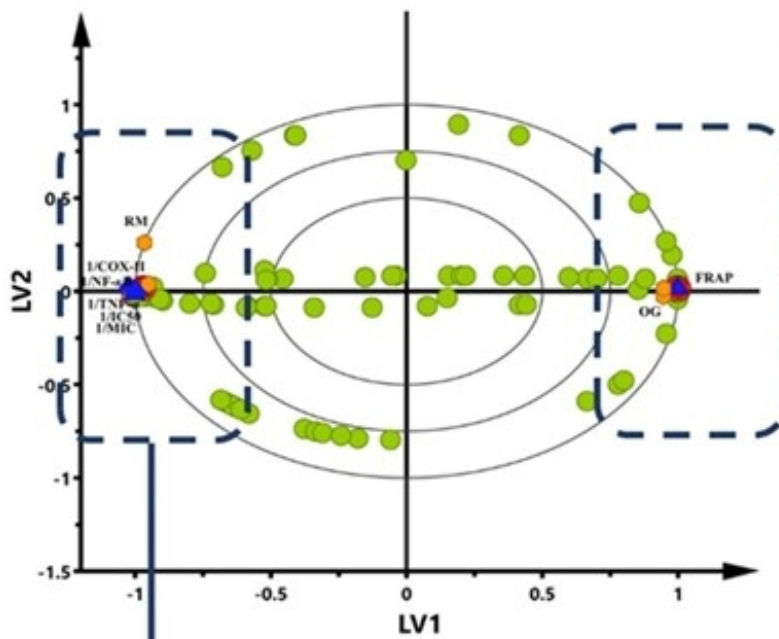
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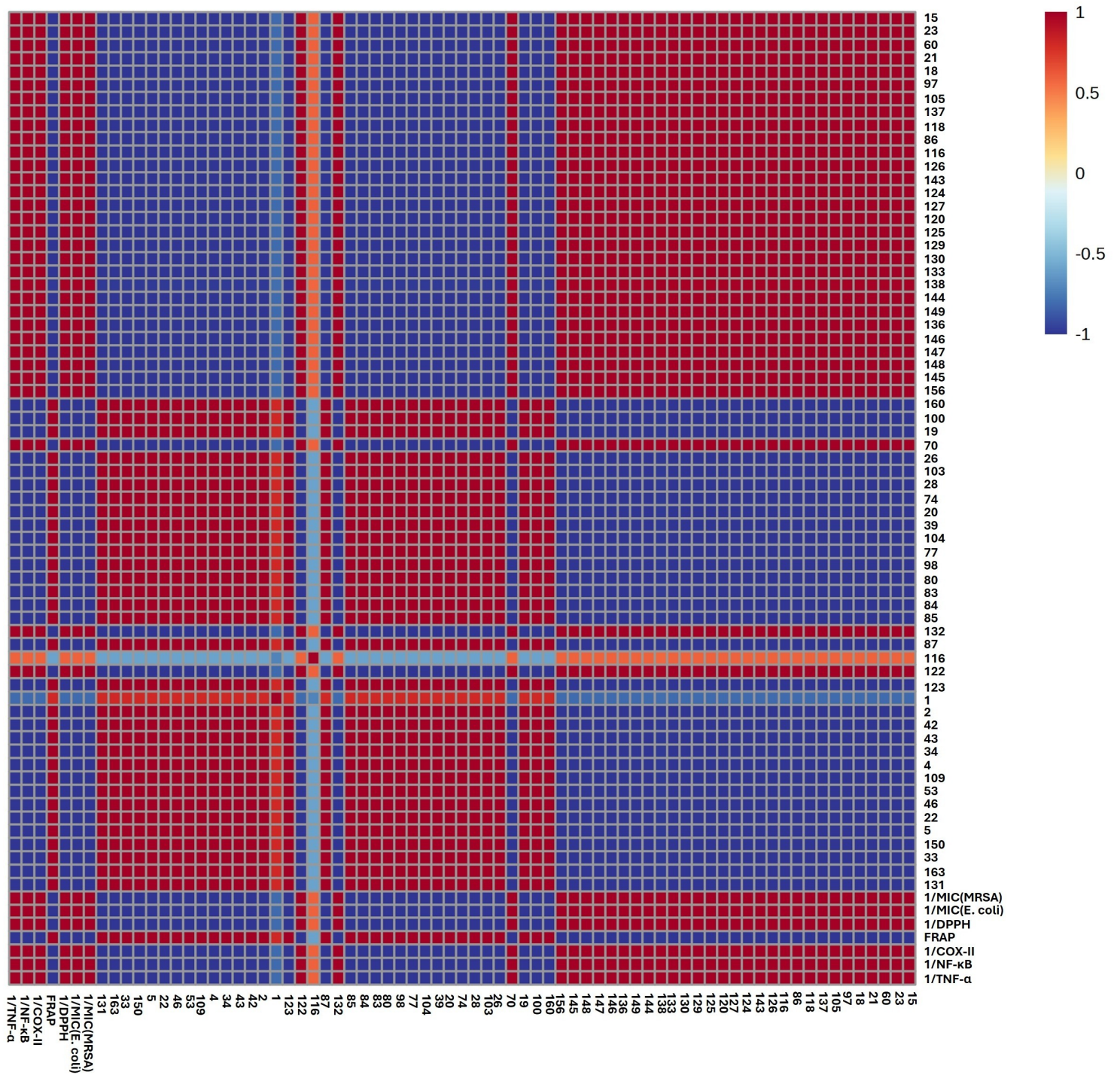


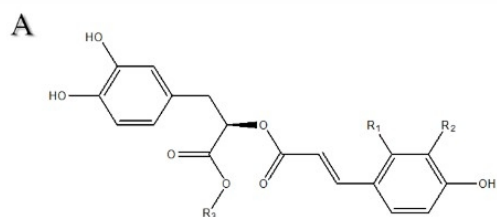




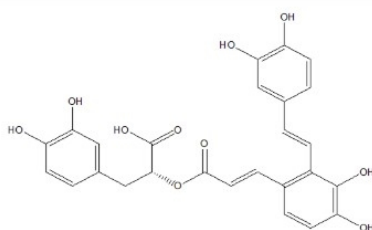




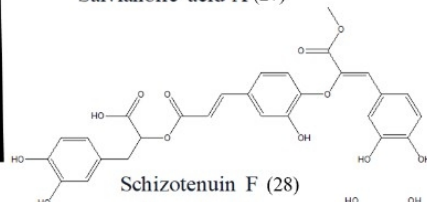




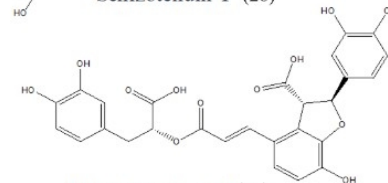
	R ₁	R ₂	R ₃
Salvianolic acid D (7)	CH ₂ COOH	OH	H
Rosmarinic acid (18)	H	OH	H
Clinopodic Acid A (25)	H	H	H
Methylrosmarinic acid (30)	H	OH	CH ₃



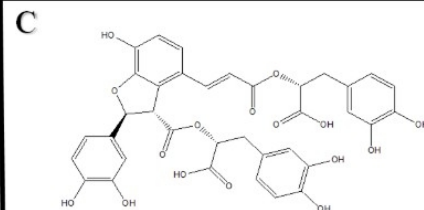
Salvianolic acid A (27)



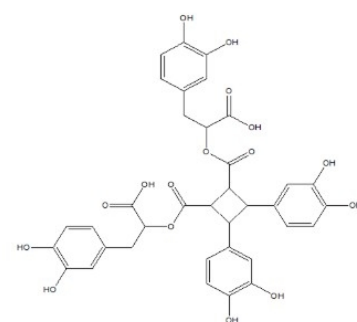
Schizotenuin F (28)



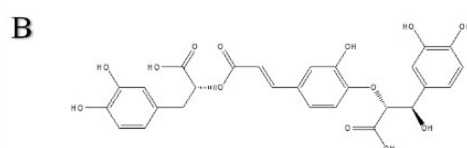
Lithospermic acid (19)



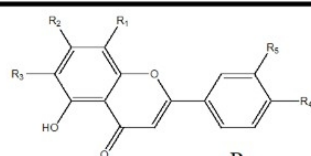
Salvianolic acid B (20)



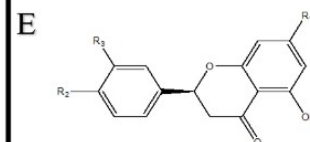
Sagerinic acid (21)



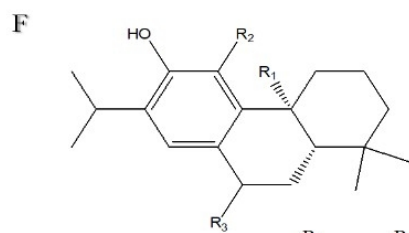
Salvianolic acid K (15)



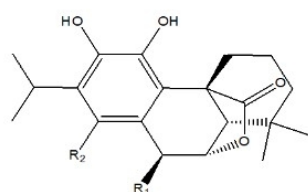
	R ₁	R ₂	R ₃	R ₄	R ₅
Vicianin 2 (52)	C-hexose	H	C-hexose	OH	H
Luteolin-O-dihexuronide (53)	H	O-dihexuronide	H	OH	OH
Luteolin-O-rutinoside (56)	H	O-rutinoside	H	OH	OH
Vitexin (57)	C-hexose	OH	H	OH	H
Apigenin-O-rutinoside (62)	H	O-rutinoside	H	OH	H
Hydroxygenkwanin (82)	H	OCH ₃	H	OH	OH
Genkwanin (86)	H	OCH ₃	H	OH	H
Apigenin 7,4'-dimethyl ether (91)	H	OCH ₃	H	OCH ₃	H



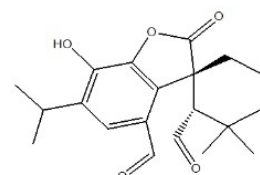
	R ₁	R ₂	R ₃
Hesperidin (101)	O-rutinoside	OCH ₃	OH
Eriodictyol (103)	OH	OH	OH
Naringenin (104)	OH	OH	H
Hesperetin (105)	OH	OCH ₃	OH
Sakuranetin (107)	OCH ₃	OH	H



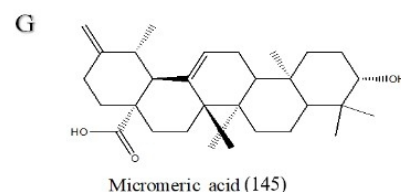
	R ₁	R ₂	R ₃
Pisiferal (135)	CHO	H	H
Carnosic acid (137)	COOH	OH	H
O-methylcarnosic acid (138)	COOH	OCH ₃	H
Sugiol (139)	CH ₃	H	=O



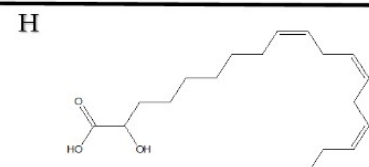
	R ₁	R ₂
Hydroxy-O-methylrosmanol (116)	OCH ₃	OH
Methylrosmanol (127)	OCH ₃	H
Dimethoxyrosmanol (129)	OCH ₃	OCH ₃
Rosmanol (132)	OH	H



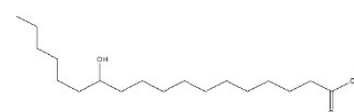
Rosmadial (130)



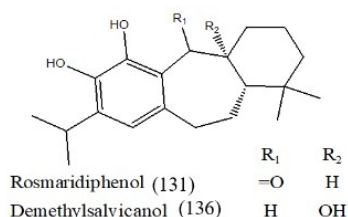
Micromeric acid (145)



Hydroxy-octadecatrienoic acid (152)



Hydroxystearic acid (155)



	R ₁	R ₂
Rosmaridiphenol (131)	=O	H
Demethylsalvicanol (136)	H	OH

Figure 1: Base peak chromatogram (BPC) of rosemary (RO) and oregano (OV) extracts analyzed by UPLC-QTOF-MS/MS in both negative (A) and positive (B) ion modes.

Figure 2: Chemical structures of the major classes of metabolites identified in the rosemary & oregano extracts. The numbers listed refer to the identified metabolites listed in **Table 13**. **A**; caffeic acid dimers, **B**; caffeic acid trimers, **C**; caffeic acid tetramer, **D**; flavones, **E**; flavanones, **F**; abietane diterpenoids, **G**; triterpenoids, **H**; fatty acids.

Figure 3: Fragmentation pattern of Salvianolic acid K.

Figure 4: Inhibition of *icaA-1* gene responsible for biofilm formation and *agrA* gene accountable for quorum sensing in MRSA. RO; *Salvia rosmarinus*, Aq.OV; Aqueous extract of *Origanum vulgare*. Data are represented as the mean $\pm$ SD of three independent assays. Statistical analysis was performed using one-way ANOVA, followed by Tukey's post hoc test was performed to calculate the statistical significance ($p < 0.0001$). As compared to RO (@), Aq.OV (#), and control (*).

Figure 5: PLS scores-loadings biplot describing the correlations of the identified metabolites in rosemary; RO and oregano; OV extracts and their studied bioactivities. In zoom, the detected compounds are annotated as metabolites.

Figure 6: Pearson's correlation between the metabolites and the antibacterial, antioxidant, and anti-inflammatory activities. Intensity of colors (blue and red) indicates correlation coefficients. The numbers listed on both axes refer to the identified metabolites listed in **Table 1**.

Figure 7: Variable Importance in the Projection (VIP) plot of the PLS model for the top contributing metabolites to bioactivities ($VIP \geq 1$).

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Table 1: Identified metabolites in *S. rosmarinus* Spenn. and *O. vulgare* L. extracts via UPLC-QTOF-MS/MS in negative/positive ionization modes. All molecular formulae were assigned with a mass accuracy limit of ± 10 ppm & the bolded fragments represent the base peak ions.

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
<i>Short-chain carboxylic acids</i>								
1	0.99	Malic acid	133.01 40		1.84	C ₄ H ₆ O ₅	115, 89, 71 , 59	RO, OV
2	1.33	Citric acid	191.01 92		2.74	C ₆ H ₈ O ₇	111, 87 , 85	RO, OV
3	1.72	Succinic acid	117.09 10		2.81	C ₄ H ₆ O ₄	99, 73	RO, OV
4	8.68	2-Isopropylmalic acid	175.06 08		2.26	C ₇ H ₁₂ O ₅	131, 115 , 85	RO, OV
5	11.4 3	Oxoadipic acid	159.02 96		1.86	C ₆ H ₈ O ₅	113, 73 , 68	OV
<i>Hydroxycinnamic acids</i>								
6	6.10	Danshensu (Salvianic acid A)	197.04 50		2.26	C ₉ H ₁₀ O ₅	179, 151, 135 , 123	RO, OV
7	6.20	Salvianolic acid D	417.07 91		8.66	C ₂₀ H ₁₈ O ₁₀	219 , 197, 179, 173	RO, OV
8	8.32	Hydroxyphenyllacti c acid	181.05 02		2.37	C ₉ H ₁₀ O ₄	163, 135 , 119, 72	RO, OV
9	8.59	Coumaric acid- <i>O</i> - hexoside	325.09 21		2.43	C ₁₅ H ₁₈ O ₈	163 , 119	RO, OV
10	8.63	Caffeic acid- <i>O</i> - hexoside	341.08 69		2.65	C ₁₅ H ₁₈ O ₉	179 , 161, 135	RO, OV
11	9.04	Caffeic acid	179.03 45	181.04 95	2.68 /0.2	C ₉ H ₈ O ₄	135 , 134, 117, 107 /163, 145, 135 , 117	RO, OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
12	9.14	Neochlorogenic acid (<i>O</i> -caffeoylquinic acid)	353.08 68	355.09 98	2.84 /7.23	C ₁₆ H ₁₈ O ₉	235, 191, 179, 173 , 161, 135 /325, 309, 193 , 163	RO, OV
13	9.62	<i>O</i> -coumaroylquinic acid	337.09 21	339.10 49	2.34 /7.52	C ₁₆ H ₁₈ O ₈	319, 293, 191, 163 /321, 221, 147, 175, 136	RO
14	9.77	Coumaryl alcohol dihexoside	519.17 04		2.94	C ₂₁ H ₃₀ O ₁₂ (formate adduct)	473, 311, 149	RO
15	10.0 8	Salvianolic acid K	555.11 29		2.72	C ₂₇ H ₂₄ O ₁₃	511, 357, 295, 197, 179, 135	RO, OV
16	10.2 1	Rosmarinic acid- <i>O</i> -hexoside (Salviaflaside)	521.12 85		3	C ₂₄ H ₂₆ O ₁₃	359, 323 , 197, 179, 161, 135	RO
17	10.2 3	Coumaric acid	163.03 97	165.05 46	2.24 /0.13	C ₉ H ₈ O ₃	119 , 117, 103, 93 /147, 119 , 91	RO, OV
18	10.4 1	Rosmarinic acid	359.07 66	361.09 19	1.78 /- 0.29	C ₁₈ H ₁₆ O ₈	197, 179, 161 , 135 /181, 163 , 139, 135	RO, OV
19	10.5 1	Lithospermic acid	537.10 21		3.25	C ₂₇ H ₂₂ O ₁₂	493, 359, 295 , 179	OV
20	10.5 3	Salvianolic acid B	717.14 37		3.35	C ₃₆ H ₃₀ O ₁₆	673, 519, 359, 321, 197	RO, OV
21	10.6 0	Sagerinic acid	719.16 02		2.16	C ₃₆ H ₃₂ O ₁₆	539, 495, 359 , 197, 179	RO, OV
22	10.8 0	Lithospermic acid derivative	897.18 48		3.97	C ₄₅ H ₃₈ O ₂₀	537, 493 , 471, 359	OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
23	11.3 1	Yunnaneic acid F	597.12 31	599.13 92	3.14 /0.56	C ₂₉ H ₂₆ O ₁₄	359, 197 , 179, 161 /387, 359, 317 , 181	RO
24	11.3 4	Ferulic acid	193.05 01	195.06 51	3.26 /0.44	C ₁₀ H ₁₀ O ₄	161, 135, 134 , 133 /179, 161, 149, 133, 119, 117	RO, OV
25	11.4 9	Clinopodic acid A	343.08 13		2.98	C ₁₈ H ₁₆ O ₇	197, 161 , 145, 135	RO, OV
26	11.6 4	Schizotenuin C1	535.08 65		3.17	C ₂₇ H ₂₀ O ₁₂	359, 197, 177 , 161	OV
27	11.7 1	Salvianolic acid A	493.11 27		2.67	C ₂₆ H ₂₂ O ₁₀	359, 295, 197, 179, 161	OV
28	11.9 9	Methyl melitrate A (Schizotenuin F)	551.11 79		2.9	C ₂₈ H ₂₄ O ₁₂	519, 359, 179, 161 , 135	OV
29	12.0 0	Coumaryl alcohol	149.06 05		2.02	C ₉ H ₁₀ O ₂	131, 119, 108 , 103	RO
30	12.0 9	Methylrosmarinic acid	373.09 17		3.18	C ₁₉ H ₁₈ O ₈	193, 197, 179 , 161, 135	RO, OV
31	12.6 8	Nepetoidin isomer	313.07 09		2.74	C ₁₇ H ₁₄ O ₆	161 , 151, 133	RO, OV
32	12.6 8	Vinyl caffeate	205.05 03		1.61	C ₁₁ H ₁₀ O ₄	161 , 133	RO, OV
33	12.9 7	Cleroden J	553.13 39		2.26	C ₂₈ H ₂₆ O ₁₂	521, 477, 373, 179, 135	OV
<i>Benzoic acid derivatives</i>								
34	5.36	Protocatechuic acid hexoside	315.07 12		3.02	C ₁₃ H ₁₆ O ₉	153 , 109	RO, OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
35	7.98	Vanillic acid		169.04 95	0.21	C ₈ H ₈ O ₄	151, 138, 125, 110, 65	RO, OV
36	7.99	Vanillic acid hexoside	329.08 68		3.05	C ₁₄ H ₁₈ O ₉	167, 123, 121, 89	RO, OV
37	8.17	Hydroxytyrosol- <i>O</i> - hexoside	315.10 77		2.66	C ₁₄ H ₂₀ O ₈	153 , 137, 123	RO, OV
38	8.43	Syringic acid		199.06 00	0.5	C ₉ H ₁₀ O ₅	181, 140 , 125, 107	RO, OV
39	8.60	Protocatechuic acid		155.03 3	5.75	C ₇ H ₆ O ₄	137 , 109, 81	RO, OV
40	8.79	Hydroxybenzoic acid- <i>O</i> -hexoside	299.07 63		3.14	C ₁₃ H ₁₆ O ₈	137 , 93	RO, OV
41	8.82	Hydroxybenzoic acid	137.02 41	139.03 90	2.3 /- 0.21	C ₇ H ₆ O ₃	93 /121 , 93, 65	RO, OV
<i>Benzyl derivatives</i>								
42	1.94	Hydroxyphenol hexoside	271.08 14		3.4	C ₁₂ H ₁₆ O ₇	109, 108 , 71	OV
43	2.26	Hydroxyphenol hexoside derivative	389.10 75		3.68	C ₁₆ H ₂₂ O ₁₁	271, 161, 113, 101, 73	OV
44	4.85	Calleryanin	301.09 22		2.29	C ₁₃ H ₁₈ O ₈	225, 139 , 121	OV
45	9.48	Benzyl alcohol- <i>O</i> - hexosyl-pentoside	401.14 41	403.15 75	3.04 /8.59	C ₁₈ H ₂₆ O ₁₀	269, 225, 161, 101, 73 /385, 333 , 271	RO, OV
46	9.63	Origanine B or C	813.18 58		3.15	C ₃₅ H ₃₅ O ₂₀	769, 615, 571 , 303	OV
47	12.1 6	Origanoside	435.12 87		2.23 /5.56	C ₂₁ H ₂₄ O ₁₀	389, 227 , 136, 92	OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
48	12.8 7	Orthosiphoic acid A		523.12 35	-0.02	C ₂₇ H ₂₂ O ₁₁	371 , 325, 163, 153	OV
<i>Jasmonic acid derivatives</i>								
49	9.21	Tuberonic acid hexoside	387.16 51		2.46	C ₁₈ H ₂₈ O ₉	225, 207, 163, 59	RO, OV
50	9.87	Epi hydroxyjasmonic acid (Tuberonic acid)	225.11 28	227.12 78	1.91 /- 0.06	C ₁₂ H ₁₈ O ₄	181, 163, 97, 59 /209, 191, 163, 149	RO, OV
51	14.8 1	Methyl jasmonate		225.14 85	0.09	C ₁₃ H ₂₀ O ₃	207, 151, 133 , 109	RO
<i>Flavones</i>								
52	9.45	Apigenin 6,8-di- <i>C</i> - hexoside (Vicenin 2)	593.14 91	595.16 57	3.52 /0.08	C ₂₇ H ₃₀ O ₁₅	503, 473 , 383, 353 /541, 511, 475, 457	RO, OV
53	9.69	Luteolin- <i>O</i> - dihexuronide	637.10 26		3.19	C ₂₇ H ₂₆ O ₁₈	351 , 285	OV
54	9.85	Hydroxyluteolin- <i>O</i> - hexoside	463.08 70	465.10 29	2.58 /- 0.32	C ₂₁ H ₂₀ O ₁₂	301 , 191, 161 / 303 , 257, 163	RO
55	9.97	Luteolin- <i>O</i> -hexosyl- hexoside		611.16 04	0.43	C ₂₇ H ₃₀ O ₁₆	449, 287	OV
56	10.1 6	Luteolin- <i>O</i> - rutinoside	593.14 95	595.16 55	2.85 /0.42	C ₂₇ H ₃₀ O ₁₅	285 , 267, 217, 197 /449, 287 , 147, 129	RO, OV
57	10.2 3	Vitexin		433.11 18	2.6	C ₂₁ H ₂₀ O ₁₀	415, 397, 379, 337, 313, 283	OV
58	10.2 6	Luteolin- <i>O</i> - hexuronide	461.07 12	463.08 76	2.92 /- 1.08	C ₂₁ H ₁₈ O ₁₂	357, 285 / 287 , 153	RO, OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
59	10.2 9	Luteolin- <i>O</i> -hexoside		449.10 78	0.08	C ₂₁ H ₂₀ O ₁₁	287 , 269	RO, OV
60	10.4 9	Nepitrin (6-Methoxyluteolin- <i>O</i> -hexoside)	477.10 27	479.11 83	2.4 /0.21	C ₂₂ H ₂₂ O ₁₂	315 , 197, 161, 153 / 317 , 302, 285, 165	RO, OV
61	10.5 3	Luteolin- <i>O</i> -pentosyl-hexoside		581.14 98	0.15	C ₂₆ H ₂₈ O ₁₅	435, 419, 287 , 271	OV
62	10.5 4	Apigenin- <i>O</i> -rutinoside		579.16 93	2.65	C ₂₇ H ₃₀ O ₁₄	433, 271	RO
63	10.6 7	Hispidulin- <i>O</i> -rutinoside		609.18 08	0.98	C ₂₈ H ₃₂ O ₁₅	463, 301 , 286	RO, OV
64	10.7 8	Luteolin- <i>O</i> -pentosyl-acetyl-hexoside		623.16 05	0.26	C ₂₈ H ₃₀ O ₁₆	583, 449, 287	OV
65	10.7 8	Apigenin- <i>O</i> -hexoside		433.11 28	0.29	C ₂₁ H ₂₀ O ₁₀	271 , 185, 153, 119, 109	RO, OV
66	10.8 1	Apigenin- <i>O</i> -hexuronide	445.07 62	447.09 21	3.22 /0.2	C ₂₁ H ₁₈ O ₁₁	427, 341, 269 , 225 / 271 , 203, 153, 109	RO, OV
67	11.1 8	Luteolin caffeoylhexoside		611.13 90	0.87	C ₃₀ H ₂₆ O ₁₄	449, 287 , 163, 153	OV
68	11.2 0	Apigenin- <i>O</i> -pentosyl-acetyl-hexoside		607.16 57	0.08	C ₂₈ H ₃₀ O ₁₅	463, 287, 271 , 199, 163	OV
69	11.5 3	Feruloylnepitrin	653.14 89	655.16 55	3.51 /0.38	C ₃₂ H ₃₀ O ₁₅	477, 315 /479, 317 , 177	RO
70	11.6 0	Luteolin-acetyl- <i>O</i> -hexuronide	503.08 18	505.09 78	2.61 /- 0.26	C ₂₃ H ₂₀ O ₁₃	443, 399, 285 , 133 /463, 287 , 201	RO

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
71	11.6 0	Acacetin- <i>O</i> - rutinoside (Linarin)	637.17 52		3.46	C ₂₈ H ₃₂ O ₁₄ (for mate adduct)	591, 283 , 268	OV
72	11.8 6	Luteolin	285.03 99	287.05 49	1.96 /0.4	C ₁₅ H ₁₀ O ₆	267, 257, 241, 217, 199, 175, 151, 133 /269, 219, 153 , 135	RO, OV
73	11.9 2	Diosmetin- <i>O</i> - pentosyl-acetyl- pentoside		607.16 57	0.08	C ₂₈ H ₃₀ O ₁₅	565, 302, 301 , 286, 229	OV
74	12.0 0	Acacetin- <i>O</i> - hexuronide	459.09 20	461.10 63	2.79 /3.34	C ₂₂ H ₂₀ O ₁₁	283 , 175, 113 / 285 , 270, 255, 153	OV
75	12.0 8	Hispidulin- <i>O</i> - hexoside (Homoplantaginin)	461.10 76	463.12 35	2.89 /- 0.03	C ₂₂ H ₂₂ O ₁₁	299 , 284 / 301 , 286	RO
76	12.3 9	Thymusin		331.08 13	-0.21	C ₁₇ H ₁₄ O ₇	316 , 301, 273, 181, 119	OV
77	12.5 6	Apigenin	269.04 51	271.06 03	1.66 /- 0.74	C ₁₅ H ₁₀ O ₅	225, 201, 151 , 149, 117, 107 /243, 153 , 119, 109	RO, OV
78	12.7 4	Diosmetin	299.05 54		2.04	C ₁₆ H ₁₂ O ₆	284 , 256, 179, 151, 107	RO, OV
79	12.8 5	Hydroxyluteolin 7,3'-dimethyl ether		331.08 13	-0.21	C ₁₇ H ₁₄ O ₇	316, 298 , 270, 136	OV
80	13.3 4	Majoranin (Thymonin)	359.07 63	361.09 19	2.61 /- 0.29	C ₁₈ H ₁₆ O ₈	344, 329 , 314 /346, 331 , 313	OV

No.	R _t . (mi n.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
81	13.5 1	Hispidulin	299.05 55	301.07 06	2.04 /0.22	C ₁₆ H ₁₂ O ₆	284 , 256, 136.9, 117 / 286 , 121, 112	RO, OV
82	13.5 3	Hydroxygenkwanin (7-methyluteoline)	299.05 54		2.37	C ₁₆ H ₁₂ O ₆	284 , 256, 227, 151, 133	OV
83	13.6 9	Cirsimaritin	313.07 10	315.08 64	2.43 /- 0.27	C ₁₇ H ₁₄ O ₆	297, 283 , 163, 135, 117 /300, 282	RO, OV
84	13.9 0	Cirsilineol	343.08 16	345.09 70	2.11 /- 0.35	C ₁₈ H ₁₆ O ₇	328 , 313, 283, 147 / 330 , 312, 284	OV
85	14.1 8	Xanthomicrol	343.08 16	345.09 70	2.11 /- 0.35	C ₁₈ H ₁₆ O ₇	328, 313 , 298, 117 /330, 315 , 297, 119	OV
86	14.3 2	Genkwanin (7-Methylapigenin)	283.06 05	285.07 58	2.45 /- 0.18	C ₁₆ H ₁₂ O ₅	268 , 211, 135, 117 / 270 , 167, 119	RO, OV
87	14.4 2	Pebrellin	373.09 19	375.10 74	2.65 /0.12	C ₁₉ H ₁₈ O ₈	358, 343 , 161, 147 /360, 345 , 327, 169	OV
88	14.6 1	Ladanein	313.07 09	315.08 64	2.74 /- 0.27	C ₁₇ H ₁₄ O ₆	298 , 283, 255, 148, 132 / 300 , 229, 133	RO, OV
89	14.9 3	Gardenin B		359.11 23	0.64	C ₁₉ H ₁₈ O ₇	344, 326 , 298, 209	OV
90	15.5 2	Salvigenin		329.10 19	0.2	C ₁₈ H ₁₆ O ₆	314, 296 , 268	RO, OV
91	16.2 5	Apigenin 7,4'- dimethyl ether		299.09 14	0	C ₁₇ H ₁₄ O ₅	284 , 256, 167, 133	RO, OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
<i>Flavonols</i>								
92	9.83	Rutin		611.16 02	0.76	C ₂₇ H ₃₀ O ₁₆	465, 303 , 287, 145, 129	RO
93	9.88	Quercetin- <i>O</i> - hexuronide	477.06 61	479.08 19	2.85 /0.24	C ₂₁ H ₁₈ O ₁₃	433, 343, 301 , 113 / 303 , 285	RO, OV
94	10.3 0	Quercetin- <i>O</i> - hexoside (Isoquercetin)	463.08 66	465.10 27	3.45 /0.11	C ₂₁ H ₂₀ O ₁₂	301, 287 , 175, 151 / 303 , 251, 145, 127	OV
95	10.3 1	Isorhamnetin- <i>O</i> - rutinoside	623.13 88	625.17 64	20 /- 0.14	C ₂₈ H ₃₂ O ₁₆	477, 315 /479, 317	RO
96	11.0 6	Quercetin coumaroylhexoside	609.12 35	611.13 91	2.42 /0.71	C ₃₀ H ₂₆ O ₁₄	463, 301 , 285 /465, 303 , 287, 147	RO
97	11.9 7	Isorhamnetin	315.05 06	317.06 55	1.35 /0.25	C ₁₆ H ₁₂ O ₇	300 , 271, 243 / 302 , 285, 121	RO, OV
98	12.8 7	Dimethylquercetin (Ombuin)	329.06 60	331.08 13	2.05 /- 0.21	C ₁₇ H ₁₄ O ₇	314, 299 , 271, 241, 199 / 316 , 121	RO, OV
<i>Flavanones</i>								
99	8.77	Gallocatechin	305.06 93		-8.57	C ₁₅ H ₁₄ O ₇	225, 97 , 59	RO, OV
100	10.4 8	Dihydroquercetin (Taxifolin)	303.04 99	305.06 55	3.7 /0.26	C ₁₅ H ₁₂ O ₇	285, 125 , 109 /287, 259, 231 , 153	OV
101	10.8 1	Hesperidin (Hesperetin- <i>O</i> - rutinoside)	609.18 07		2.94	C ₂₈ H ₃₄ O ₁₅	301 , 285, 251	RO
102	11.1 0	Aromadendrin	287.05 53	289.07 06	2.82 /0.22	C ₁₅ H ₁₂ O ₆	259, 243, 201, 177, 151, 125	OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extract
			M-H	M+H				
							/271, 243, 215 , 153, 149	
103	11.7 7	Eriodictyol	287.05 53	289.07 06	2.82 /0.22	C ₁₅ H ₁₂ O ₆	151, 135 , 125, 107 /271, 179, 163, 153	OV
104	12.4 9	Naringenin	271.06 07	273.07 51	1.83 /2.39	C ₁₅ H ₁₂ O ₅	253, 151 , 119, 107 /255, 179, 153	OV
105	12.7 8	Hesperetin	301.07 11	303.08 54	2.19 /3.03	C ₁₆ H ₁₄ O ₆	286, 164 , 151, 136 / 177 , 153, 137, 117	RO, OV
106	13.8 4	5,4'-Dihydroxy- 6,7,8- trimethoxyflavanone	345.09 70	347.11 27	2.82 /- 0.49	C ₁₈ H ₁₈ O ₇	330, 315, 301, 283 , 119 /332, 227, 197	OV
107	14.2 6	Sakuranetin	285.07 60	287.09 15	2.96 /- 0.35	C ₁₆ H ₁₄ O ₅	270, 165, 119 / 167 , 147, 119	OV
<i>Monoterpenes</i>								
108	8.45	Thymoquinol- <i>O</i> - dihexoside	535.20 14		3.41	C ₂₂ H ₃₄ O ₁₂ . CH ₂ O ₂	489, 327 , 165, 161, 101	OV
109	9.62	Thymoquinol- <i>O</i> - hexoside	373.14 95		2.42	C ₁₆ H ₂₄ O ₇ . CH ₂ O ₂	327, 165, 164 , 113, 101	OV
110	11.9 6	(+)-Camphor		153.12 74	-0.06	C ₁₀ H ₁₆ O	109 , 95, 81	RO, OV
111	12.9 8	Thymol		151.11 17	0.28	C ₁₀ H ₁₄ O	123, 109 , 91, 81	RO, OV
<i>Sesquiterpenoids</i>								
112	13.7 4	Spathulenol		221.18 99	0.42	C ₁₅ H ₂₄ O	203, 147 , 133, 119	OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
113	14.1 1	Caryophyllene epoxide		221.19 00	-0.04	C ₁₅ H ₂₄ O	203 , 147, 95	RO
114	18.1 1	Cubebene		205.19 52	-0.6	C ₁₅ H ₂₄	165, 148, 121 , 93	OV
<i>Diterpenes</i>								
115	13.5 1	Carnosic acid hexoside	493.24 30		2.64	C ₂₆ H ₃₈ O ₉	331 , 287	RO
116	13.5 3	Hydroxy- <i>O</i> - methylosmanol	375.18 03		2.69	C ₂₁ H ₂₈ O ₆	357, 345, 331, 316, 299	RO, OV
117	14.0 9	Rosmanol derivative	677.36 76		2.81	C ₄₀ H ₅₄ O ₉	345 , 331, 315, 301, 283	RO
118	14.1 4	Hydroxyrosmanol	361.16 47		2.66	C ₂₀ H ₂₆ O ₆	317, 299	RO, OV
119	14.6 9	Rosmanol derivative	689.33 12		2.78	C ₄₀ H ₅₀ O ₁₀	555, 345 , 283	RO
120	14.7 3	Sageone	299.16 46	301.17 99	2.23 +/- 0.26	C ₁₉ H ₂₄ O ₃	243 , 200 /283, 259, 241, 231	RO, OV
121	14.7 5	Hydroxy- <i>O</i> - methylosmanol derivative	719.34 24		1.79	C ₄₁ H ₅₂ O ₁₁	375 , 360, 345, 313, 299	RO, OV
122	14.8 8	Hydroxyrosmadial	359.14 89	361.16 43	3.09 /0.74	C ₂₀ H ₂₄ O ₆	341, 331, 315 , 287 / 333 , 317, 289, 287	RO, OV
123	14.9 9	Carnosol	329.17 50	331.18 92	2.52 /3.59	C ₂₀ H ₂₆ O ₄	314, 285 , 270, 201 /313, 289, 271, 247	RO, OV
124	15.4 4	Oridonin	363.18 04		2.5	C ₂₀ H ₂₈ O ₆	345, 333, 315 , 297	RO

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
125	15.6 0	Salvinorin F	373.16 48		2.3	C ₂₁ H ₂₆ O ₆	358, 343, 329, 314 , 299	RO, OV
126	15.6 3	Rosmic acid	389.15 92	391.17 52	3.53 /- 0.18	C ₂₁ H ₂₆ O ₇	345, 313, 301, 285 / 331 , 303, 285	RO, OV
127	15.8 1	Methylrosmanol	359.18 57	361.20 12	1.94 /- 0.69	C ₂₁ H ₂₈ O ₅	329, 315, 283 /329, 301, 273, 109	RO, OV
128	16.1 1	Sageone derivative	629.34 66		2.82	C ₃₉ H ₅₀ O ₇	329, 299 , 285	RO, OV
129	16.4 0	Dimethoxyrosmano l	389.19 56		3.49	C ₂₂ H ₃₀ O ₆	374 , 344, 330, 313, 298	RO, OV
130	16.7 4	Rosmadial	343.15 44	345.16 97	2.03 /- 0.14	C ₂₀ H ₂₄ O ₅	328, 315, 299 /317, 299, 271, 231	RO, OV
131	16.7 7	Rosmaridiphenol	315.19 60	317.20 94	1.8 /5.44	C ₂₀ H ₂₈ O ₃	285 , 270, 201, 179 /299, 281, 191	RO, OV
132	16.9 2	Rosmanol	345.17 00	347.18 54	2.16 /- 0.29	C ₂₀ H ₂₆ O ₅	327, 301, 283 , 268, 227 / 301 , 259, 241	RO, OV
133	16.9 4	Tetrahydro- hydroxy rosmariquinone	301.18 02		2.38	C ₁₉ H ₂₆ O ₃	283, 273, 258	RO, OV
134	17.2 3	<i>O</i> -Methylcarnosol	343.19 07		2.27	C ₂₁ H ₂₈ O ₄	299, 284	RO, OV
135	17.8 3	Pisiferal		301.21 63	-0.31	C ₂₀ H ₂₈ O ₂	259, 245, 231, 219, 205 , 163	RO, OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
136	17.9 1	Demethylsalvicanol	317.21 14		2.57	C ₂₀ H ₃₀ O ₃	299, 243, 191, 179	RO, OV
137	18.0 5	Carnosic acid	331.19 04		3.26	C ₂₀ H ₂₈ O ₄	313, 287, 244, 151	RO, OV
138	18.4 5	<i>O</i> -methylcarnosic acid	345.20 63	347.21 93	2.41 /6.89	C ₂₁ H ₃₀ O ₄	301, 286 , 271 /305, 273 , 121, 109	RO, OV
139	18.7 9	Sugiol	299.20 10		2.18	C ₂₀ H ₂₈ O ₂	283, 243 , 227, 215	RO, OV
140	20.3 4	Rosmadial derivative	631.39 89		2.39	C ₄₀ H ₅₆ O ₆	567, 343 , 299	RO
141	21.4 9	Hydroxyrosmadial derivative	631.39 89		2.39	C ₄₀ H ₅₆ O ₆	359 , 331, 315	RO, OV
142	22.4 2	Rosmadial derivative	615.40 32		3.73	C ₄₀ H ₅₆ O ₅	572, 343 , 299	RO, OV
<i>Triterpenes</i>								
143	14.8 5	Asiatic acid	487.34 16		2.66	C ₃₀ H ₄₈ O ₅	469, 453, 441, 409	RO
144	17.0 9	Pomolic acid	471.34 66	473.36 22	2.93/0. 71	C ₃₀ H ₄₈ O ₄	453, 407 /455, 409 , 191	RO, OV
145	17.3 6	Micromeric acid	453.33 63	455.35 18	2.46 /0.38	C ₃₀ H ₄₆ O ₃	409, 407 , 392 /437, 409 , 205	RO, OV
146	19.2 2	Oleanolic acid	455.35 17		3	C ₃₀ H ₄₈ O ₃	409 , 391	RO, OV
147	19.4 8	Ursolic acid	455.35 19		2.56	C ₃₀ H ₄₈ O ₃	407 , 315	RO, OV
148	19.5 1	3-oxours-12-en- 20,28-olide	451.32 06		2.58	C ₃₀ H ₄₄ O ₃	407 , 389	RO

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
149	21.2 5	Corosolic acid	471.34 66		2.93	C ₃₀ H ₄₈ O ₄	453, 443, 427, 409 , 393	RO
<i>Fatty acids derivatives</i>								
150	12.6 9	Corchorifatty acid F	327.21 70		2.13	C ₁₈ H ₃₂ O ₅	291, 229, 211 , 171	RO, OV
151	13.2 9	Dihydroxyhexadeca noic acid	287.22 21		2.37	C ₁₆ H ₃₂ O ₄	269, 241, 141 , 99	RO, OV
152	14.4 8	Hydroxy- octadecatrienoic acid		295.22 66	0.58	C ₁₈ H ₃₀ O ₃	277, 259, 133, 69	RO, OV
153	14.7 8	hydroperoxy- octadecadienoic acid	311.22 20		2.51	C ₁₈ H ₃₂ O ₄	293, 223 , 195	RO, OV
154	16.8 1	Linolenic Acid	277.21 64	279.23 20	3.25 +/- 0.52	C ₁₈ H ₃₀ O ₂	259, 233, 59 /123, 109, 95, 67	RO, OV
155	19.8 2	Hydroxystearic acid	299.25 85		2.23	C ₁₈ H ₃₆ O ₃	253, 59	RO
156	21.0 9	Camaryolic acid	581.38 33		2.51	C ₃₆ H ₅₄ O ₆	549 , 497, 285	RO, OV
<i>Others</i>								
157	0.99	Quinic acid	191.05 58		1.62	C ₇ H ₁₂ O ₆	173, 171, 127, 85	RO, OV
158	1.00	Acetyl-maltose	383.11 89		1.56	C ₁₄ H ₂₄ O ₁₂	341, 191	RO
159	9.44	Hydroxybenzaldehy de	121.02 93	123.04 40	1.66 /0.46	C ₇ H ₆ O ₂	92 , 65 /95, 77 , 51	OV
160	10.2 5	Amburosides A	421.11 27		3.13	C ₂₀ H ₂₂ O ₁₀	259, 153 , 109	OV

No.	R _t . (min.)	Metabolite name	Mol. Ion <i>m/z</i>		Δ mass (ppm)	Molecular formula	MS ² ions <i>m/z</i> (-)/(+)	Extra ct
			M-H	M+H				
161	10.8 2	Loliolide		197.11 72	0.11	C ₁₁ H ₁₆ O ₃	179, 161, 133 , 107	RO, OV
162	11.6 2	Dihydroxyanthraquinone carboxylic acid	283.02 40		2.86	C ₁₅ H ₈ O ₆	239, 211 , 135	OV
163	13.9 3	6-Oxocamphor	165.09 17		2.43	C ₁₀ H ₁₄ O ₂	150 , 135	OV
164	14.4 5	Hotrienol		153.12 74	-0.06	C ₁₀ H ₁₆ O	135, 107, 97	RO, OV

Table 2: MIC inhibitory test of methanolic and aqueous extracts of *S. rosmarinus* Spenn. and *O. vulgare* L. against MRSA and *E. coli*. **MIC**; minimum inhibitory concentration, **MRSA**; methicillin-resistant *Staphylococcus aureus*, ***E. coli***; *Escherichia coli*. MICs were assayed in triplicate.

Extract	MRSA (μg/mL)	<i>E. coli</i> (μg/mL)
RO	64	64
Aq.RO	256	128
OV	256	128
Aq.OV	64	64
Doxycycline	1	32

Table 3: Antioxidant activity of methanolic extracts of *S. rosmarinus* Spenn. and *O. vulgare* L. using 2,2-diphenyl-1-picrylhydrazyl (DPPH) and ferric-reducing antioxidant power (FRAP) assays. The results are

presented as means $\pm$ SD of three measurements ($n = 3$). Statistical analysis was performed using the Unpaired Student's t -test. * ($p < 0.0001$) and # ($p = 0.0004$).

Antioxidant activity		
	RO	OV
DPPH "IC₅₀" ($\mu\text{g/mL}$)	$6.56 \pm 0.035^*$	17.60 ± 0.333
FRAP ($\mu\text{mol/L}$)	122.78 ± 2.092	$188.60 \pm 10.058^\#$

Table 4: Anti-inflammatory activity of *S. rosmarinus* Spenn. and *O. vulgare* L. methanolic extracts. Results are presented as (mean $\pm$ SD) of three measurements ($n=3$). Statistical analysis was performed using one-way ANOVA, followed by Tukey's post hoc test, which was performed to calculate the statistical significance relative to the ibuprofen group indicated as follows: (*) at $p < 0.05$; (**) at $p < 0.005$; (****) at $p < 0.0001$; NS = not significant.

	COX-II IC₅₀ ($\mu\text{g/mL}$)	TNF-α (pg/mL)	NFκb (pg/mL)
RO	11.09 ± 0.46 NS	536.6 ± 13.4 **	$616.2 \pm 26.9^*$
OV	34.73 ± 1.44 ****	874.1 ± 35 ****	1025 ± 40.4 ****
Ibuprofen	8.791 ± 0.37	409.4 ± 14.7	496.2 ± 24.7
Control	--	1392 ± 21.3	1992 ± 48.7