



OPEN 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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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 the 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 squares analysis (PLS)

The Lamiaceae family (mint family) encompasses a broad range of aromatic plants that are widely distributed but particularly prevalent in the Mediterranean region¹. Lamiaceae plants have been used extensively as home remedies and to enhance the taste and aroma of foods since ancient times². The mint family has long been used for decades in cosmetics and conventional medicine as antiseptics, carminatives, expectorants, and sedatives², 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 activities⁵. In addition to essential oils, Lamiaceae plants are rich in non-volatile secondary

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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}. Owing 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 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 squares 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 the most and the 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.

Materials and methods

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).

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.

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 3000g (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.

Antibacterial activity evaluation

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 43,300 as Gram-positive bacteria, and the Gram-negative bacteria *Escherichia coli* ATCC 25,922. 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).

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.

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 µL 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 µL of sterile phosphate-buffered saline (PBS) pH 7.2, fixed by drying for 1 h at 60 °C, then stained with 200 µL 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 µL of 33% acetic acid, and transferred (150 µL) 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:

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

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

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'-CAATACTATTTTCGGGTGTCTTCAC TCT-3; 5'-CAAGAACTGCAATATCTTCGGTAATCAT-3'), respectively²¹. The relative expression values of each gene were normalized to the value of the housekeeping gene 16 S rRNA. The forward and reverse primers used for 16 S rRNA were 5'-TGT CGT GAG ATG TTG GG-3', and 5'-TGT CGT GAG ATG TTG GG-3', respectively²¹. $2^{-\Delta\Delta CT}$ method was used to calculate the results²². The results were reported as means \pm SD of triplicate measurements.

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:

2,2-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.

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 hydrochloric acid, and 20 mM ferric chloride at a ratio 10:1:1 v/v/v, respectively. Freshly prepared FRAP reagent (1.5 mL) was added to 50 μ L 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 μ mol/L $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$ equivalent/mg of each extract.

Anti-inflammatory activity evaluation

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 μ g/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.

Tumor necrosis factor- α (TNF- α) and nuclear factor kappa-B (NF- κ B) 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 μ g/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- α (TNF- α) and nuclear factor kappa-B (NF- κ B) 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- κ B p100/NFKB2 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- κ B). The equivalent concentrations were determined from standard curves, and all measurements were carried out in triplicate.

Statistical analysis

All experiments were performed in triplicate, and the results are expressed as mean \pm 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.

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* = 0.9–1.0²⁷.

Results and discussion

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 Fig. 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 Fig. 2. Details of the assigned metabolites are discussed in the following subsections.

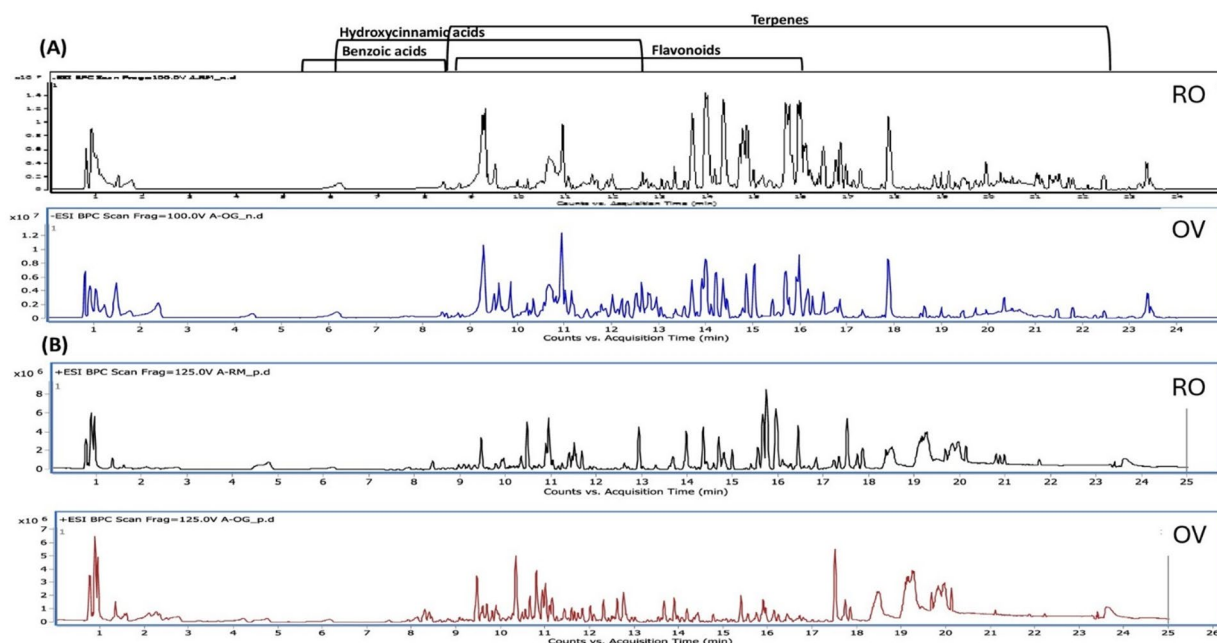


Fig. 1. Base peak chromatogram (BPC) of rosemary (RO) and oregano (OV) extracts analyzed by UPLC-QTOF-MS/MS in both negative (A) and positive (B) ion modes.

Hydroxycinnamic acid derivatives

Twenty-eight hydroxycinnamic acid derivatives of caffeic, coumaric, ferulic acids, and danshensu (salvianic acid A or α -hydroxy dihydrocaffeic acid $C_9H_{10}O_5$) 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 (Fig. 3).

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 Fig. 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 Fig. 2–4 S)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-OCH₃-CO₂][−]. A further significant ion at m/z 179 [C₉H₇O₄][−] confirms the presence of caffeic acid units within the structure (Supplementary Fig. 5S).

Caffeic acid trimers *vis.* salvianolic acids A and K were identified. Salvianolic acid K (cpd. 15, m/z 555.1129 [M-H][−], [C₂₇H₂₃O₁₃][−]) 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 Fig. 6S), respectively³³. The fragmentation pattern of this compound is presented as a representative example of the.

Salvianolic acid B and sagerinic acid are caffeic acid tetramers (rosmarinic acid dimers). Salvianolic acid B1 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][−], [C₃₆H₂₉O₁₆][−]), produced a base peak ion at m/z 359 [C₁₈H₁₅O₈][−] 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 Fig. 8S)³⁵.

Sagerinic acid (cpd. 21, m/z 719.1602 [M-H][−], [C₃₆H₃₁O₁₆][−]) 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 [C₉H₉O₅][−] corresponding to the danshensu moiety (Supplementary Fig. 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

No.	R _f (min.)	Metabolite name	Mol. Ion m/z		Δ mass (ppm)	Molecular formula	MS ² ions m/z (-)/(+)	Extract
			M-H	M+H				
Short-chain carboxylic acids								
1	0.99	Malic acid	133.0140		1.84	C ₄ H ₆ O ₅	115, 89, 71, 59	RO, OV
2	1.33	Citric acid	191.0192		2.74	C ₆ H ₈ O ₇	111, 87, 85	RO, OV
3	1.72	Succinic acid	117.0910		2.81	C ₄ H ₆ O ₄	99, 73	RO, OV
4	8.68	2-Isopropylmalic acid	175.0608		2.26	C ₇ H ₁₂ O ₅	131, 115, 85	RO, OV
5	11.43	Oxoadipic acid	159.0296		1.86	C ₆ H ₈ O ₅	113, 73, 68	OV
Hydroxycinnamic acids								
6	6.10	Danshensu (Salvianic acid A)	197.0450		2.26	C ₉ H ₁₀ O ₅	179, 151, 135, 123	RO, OV
7	6.20	Salvianolic acid D	417.0791		8.66	C ₂₀ H ₁₈ O ₁₀	219, 197, 179, 173	RO, OV
8	8.32	Hydroxyphenyllactic acid	181.0502		2.37	C ₉ H ₁₀ O ₄	163, 135, 119, 72	RO, OV
9	8.59	Coumaric acid-O-hexoside	325.0921		2.43	C ₁₅ H ₁₈ O ₈	163, 119	RO, OV
10	8.63	Caffeic acid-O-hexoside	341.0869		2.65	C ₁₅ H ₁₈ O ₉	179, 161, 135	RO, OV
11	9.04	Caffeic acid	179.0345	181.0495	2.68 /0.2	C ₉ H ₈ O ₄	135, 134, 117, 107 /163, 145, 135, 117	RO, OV
12	9.14	Neochlorogenic acid (O-caffeoylquinic acid)	353.0868	355.0998	2.84 /7.23	C ₁₆ H ₁₈ O ₉	235, 191, 179, 13,173, 161, 135 /325, 309, 193, 163	RO, OV
13	9.62	O-coumaroylquinic acid	337.0921	339.1049	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	519.1704		2.94	C ₂₁ H ₃₀ O ₁₂ (formate adduct)	473, 311, 149	RO
15	10.08	Salvianolic acid K	555.1129		2.72	C ₂₇ H ₂₄ O ₁₃	511, 357, 295, 197, 179, 135	RO, OV
16	10.21	Rosmarinic acid-O-hexoside (Salviaflaside)	521.1285		3	C ₂₄ H ₂₆ O ₁₃	359, 323, 197, 179, 161, 135	RO
17	10.23	Coumaric acid	163.0397	165.0546	2.24 /0.13	C ₉ H ₈ O ₃	119, 117, 103, 93 /147, 119, 91	RO, OV
18	10.41	Rosmarinic acid	359.0766	361.0919	1.78 /-0.29	C ₁₈ H ₁₆ O ₈	197, 179, 161, 135 /181, 163, 139, 135	RO, OV
19	10.51	Lithospermic acid	537.1021		3.25	C ₂₇ H ₂₂ O ₁₂	493, 359, 295, 179	OV
20	10.53	Salvianolic acid B	717.1437		3.35	C ₃₆ H ₃₀ O ₁₆	673, 519, 359, 321, 197	RO, OV
21	10.60	Sagerinic acid	719.1602		2.16	C ₃₆ H ₃₂ O ₁₆	539, 495, 359, 197, 179	RO, OV
22	10.80	Lithospermic acid derivative	897.1848		3.97	C ₄₅ H ₃₈ O ₂₀	537, 493, 471, 359	OV
23	11.31	Yunnaneic acid F	597.1231	599.1392	3.14 /0.56	C ₂₉ H ₂₆ O ₁₄	359, 197, 179, 161 /387, 359, 317, 181	RO
24	11.34	Ferulic acid	193.0501	195.0651	3.26 /0.44	C ₁₀ H ₁₀ O ₄	161, 135, 134, 133 /179, 161, 149, 133, 119, 117	RO, OV
25	11.49	Clinopodic acid A	343.0813		2.98	C ₁₈ H ₁₆ O ₇	197, 161, 145, 135	RO, OV
26	11.64	Schizotenuin C1	535.0865		3.17	C ₂₇ H ₂₀ O ₁₂	359, 197, 177, 161	OV
27	11.71	Salvianolic acid A	493.1127		2.67	C ₂₆ H ₂₂ O ₁₀	359, 295, 197, 179, 161	OV
28	11.99	Methyl melitrate A (Schizotenuin F)	551.1179		2.9	C ₂₈ H ₂₄ O ₁₂	519, 359, 179, 161, 135	OV
29	12.00	Coumaryl alcohol	149.0605		2.02	C ₉ H ₁₀ O ₂	131, 119, 108, 103	RO
30	12.09	Methylrosmarinic acid	373.0917		3.18	C ₁₉ H ₁₈ O ₈	193, 197, 179, 161, 135	RO, OV
31	12.68	Nepetoidin isomer	313.0709		2.74	C ₁₇ H ₁₄ O ₆	161, 151, 133	RO, OV
32	12.68	Vinyl caffeate	205.0503		1.61	C ₁₁ H ₁₀ O ₄	161, 133	RO, OV
33	12.97	Cleroden J	553.1339		2.26	C ₂₈ H ₂₆ O ₁₂	521, 477, 373, 179, 135	OV
Benzoic acid derivatives								
34	5.36	Protocatechuic acid hexoside	315.0712		3.02	C ₁₃ H ₁₆ O ₉	153, 109	RO, OV
35	7.98	Vanillic acid		169.0495	0.21	C ₈ H ₈ O ₄	151, 138, 125, 110, 65	RO, OV
36	7.99	Vanillic acid hexoside	329.0868		3.05	C ₁₄ H ₁₈ O ₉	167, 123, 121, 89	RO, OV
37	8.17	Hydroxytyrosol-O-hexoside	315.1077		2.66	C ₁₄ H ₂₀ O ₈	153, 137, 123	RO, OV
38	8.43	Syringic acid		199.0600	0.5	C ₉ H ₁₀ O ₅	181, 140, 125, 107	RO, OV
39	8.60	Protocatechuic acid		155.033	5.75	C ₇ H ₆ O ₄	137, 109, 81	RO, OV
40	8.79	Hydroxybenzoic acid-O-hexoside	299.0763		3.14	C ₁₃ H ₁₆ O ₈	137, 93	RO, OV
41	8.82	Hydroxybenzoic acid	137.0241	139.0390	2.3 /-0.21	C ₇ H ₆ O ₃	93 /121, 93, 65	RO, OV
Benzyl derivatives								
42	1.94	Hydroxyphenol hexoside	271.0814		3.4	C ₁₂ H ₁₆ O ₇	109, 108, 71	OV
43	2.26	Hydroxyphenol hexoside derivative	389.1075		3.68	C ₁₆ H ₂₂ O ₁₁	271, 161, 113, 101,73	OV
44	4.85	Calleryanin	301.0922		2.29	C ₁₃ H ₁₈ O ₈	225, 139, 121	OV
45	9.48	Benzyl alcohol-O-hexosyl-pentoside	401.1441	403.1575	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	813.1858		3.15	C ₃₅ H ₃₅ O ₂₀	769, 615, 571, 303	OV
47	12.16	Origanoside	435.1287		2.23 /5.56	C ₂₁ H ₂₄ O ₁₀	389, 227, 136, 92	OV
48	12.87	Orthosiphic acid A		523.1235	-0.02	C ₂₇ H ₂₂ O ₁₁	371, 325, 163, 153	OV
Continued								

No.	R _f (min.)	Metabolite name	Mol. Ion m/z		Δ mass (ppm)	Molecular formula	MS ² ions m/z (-)/(+)	Extract
			M-H	M + H				
Jasmonic acid derivatives								
49	9.21	Tuberonic acid hexoside	387.1651		2.46	C ₁₈ H ₂₈ O ₉	225, 207, 163, 59	RO, OV
50	9.87	Epi hydroxyjasmonic acid (Tuberonic acid)	225.1128	227.1278	1.91 /-0.06	C ₁₂ H ₁₈ O ₄	181, 163, 97, 59 /209, 191, 163, 149	RO, OV
51	14.81	Methyl jasmonate		225.1485	0.09	C ₁₃ H ₂₀ O ₃	207, 151, 133, 109	RO
Flavones								
52	9.45	Apigenin 6,8-di-C-hexoside (Vicenin 2)	593.1491	595.1657	3.52 /0.08	C ₂₇ H ₃₀ O ₁₅	503, 473, 383, 353 /541, 511, 475, 457	RO, OV
53	9.69	Luteolin-O-dihexuronide	637.1026		3.19	C ₂₇ H ₂₆ O ₁₈	351, 285	OV
54	9.85	Hydroxyluteolin-O-hexoside	463.0870	465.1029	2.58 /-0.32	C ₂₁ H ₂₀ O ₁₂	301, 191, 161 /303, 257, 163	RO
55	9.97	Luteolin-O-hexosyl-hexoside		611.1604	0.43	C ₂₇ H ₃₀ O ₁₆	449, 287	OV
56	10.16	Luteolin-O-rutinoside	593.1495	595.1655	2.85 /0.42	C ₂₇ H ₃₀ O ₁₅	285, 267, 217, 197 /449, 287, 147, 129	RO, OV
57	10.23	Vitexin		433.1118	2.6	C ₂₁ H ₂₀ O ₁₀	415, 397, 379, 337, 313, 283	OV
58	10.26	Luteolin-O-hexuronide	461.0712	463.0876	2.92 /-1.08	C ₂₁ H ₁₈ O ₁₂	357, 285 /287, 153	RO, OV
59	10.29	Luteolin-O-hexoside		449.1078	0.08	C ₂₁ H ₂₀ O ₁₁	287, 269	RO, OV
60	10.49	Nepitrin (6-Methoxyluteolin-O-hexoside)	477.1027	479.1183	2.4 /0.21	C ₂₂ H ₂₂ O ₁₂	315, 197, 161, 153 /317, 302, 285, 165	RO, OV
61	10.53	Luteolin-O-pentosyl-hexoside		581.1498	0.15	C ₂₆ H ₂₈ O ₁₅	435, 419, 287, 271	OV
62	10.54	Apigenin-O-rutinoside		579.1693	2.65	C ₂₇ H ₃₀ O ₁₄	433, 271	RO
63	10.67	Hispidulin-O-rutinoside		609.1808	0.98	C ₂₈ H ₃₂ O ₁₅	463, 301, 286	RO, OV
64	10.78	Luteolin-O-pentosyl-acetyl-hexoside		623.1605	0.26	C ₂₈ H ₃₀ O ₁₆	583, 449, 287	OV
65	10.78	Apigenin-O-hexoside		433.1128	0.29	C ₂₁ H ₂₀ O ₁₀	271, 185, 153, 119, 109	RO, OV
66	10.81	Apigenin-O-hexuronide	445.0762	447.0921	3.22 /0.2	C ₂₁ H ₁₈ O ₁₁	427, 341, 269, 225 /271, 203, 153, 109	RO, OV
67	11.18	Luteolin caffeoylhexoside		611.1390	0.87	C ₃₀ H ₂₆ O ₁₄	449, 287, 163, 153	OV
68	11.20	Apigenin-O-pentosyl-acetyl-hexoside		607.1657	0.08	C ₂₈ H ₃₀ O ₁₅	463, 287, 271, 199, 163	OV
69	11.53	Feruloylnepitrin	653.1489	655.1655	3.51 /0.38	C ₃₂ H ₃₀ O ₁₅	477, 315 /479, 317, 177	RO
70	11.60	Luteolin-acetyl-O- hexuronide	503.0818	505.0978	2.61 /-0.26	C ₂₃ H ₂₀ O ₁₃	443, 399, 285, 133 /463, 287, 201	RO
71	11.60	Acacetin-O-rutinoside (Linarin)	637.1752		3.46	C ₂₈ H ₃₂ O ₁₄ (formate adduct)	591, 283, 268	OV
72	11.86	Luteolin	285.0399	287.0549	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	11.92	Diosmetin-O-pentosyl-acetyl-pentoside		607.1657	0.08	C ₂₈ H ₃₀ O ₁₅	565, 302, 301, 286, 229	OV
74	12.00	Acacetin-O-hexuronide	459.0920	461.1063	2.79 /3.34	C ₂₂ H ₂₀ O ₁₁	283, 175, 113 /285, 270, 255, 153	OV
75	12.08	Hispidulin-O-hexoside (Homoplantaginin)	461.1076	463.1235	2.89 /-0.03	C ₂₂ H ₂₂ O ₁₁	299, 284 /301, 286	RO
76	12.39	Thymusin		331.0813	-0.21	C ₁₇ H ₁₄ O ₇	316, 301, 273, 181, 119	OV
77	12.56	Apigenin	269.0451	271.0603	1.66 /-0.74	C ₁₅ H ₁₀ O ₅	225, 201, 151, 149, 117, 107 /243, 153, 119, 109	RO, OV
78	12.74	Diosmetin	299.0554		2.04	C ₁₆ H ₁₂ O ₆	284, 256, 179, 151, 107	RO, OV
79	12.85	Hydroxyluteolin 7,3'-dimethyl ether		331.0813	-0.21	C ₁₇ H ₁₄ O ₇	316, 298, 270, 136	OV
80	13.34	Majoranin (Thymonin)	359.0763	361.0919	2.61 /-0.29	C ₁₈ H ₁₆ O ₈	344, 329, 314 /346, 331, 313	OV
81	13.51	Hispidulin	299.0555	301.0706	2.04 /0.22	C ₁₆ H ₁₂ O ₆	284, 256, 136.9, 117 /286, 121, 112	RO, OV
82	13.53	Hydroxygenkwanin (7-methyluteoline)	299.0554		2.37	C ₁₆ H ₁₂ O ₆	284, 256, 227, 151, 133	OV
83	13.69	Cirsimaritin	313.0710	315.0864	2.43 /-0.27	C ₁₇ H ₁₄ O ₆	297, 283, 163, 135, 117 /300, 282	RO, OV
84	13.90	Cirsilineol	343.0816	345.0970	2.11 /-0.35	C ₁₈ H ₁₆ O ₇	328, 313, 283, 147 /330, 312, 284	OV
85	14.18	Xanthomicrol	343.0816	345.0970	2.11 /-0.35	C ₁₈ H ₁₆ O ₇	328, 313, 298, 117 /330, 315, 297, 119	OV
86	14.32	Genkwanin (7-Methylapigenin)	283.0605	285.0758	2.45 /-0.18	C ₁₆ H ₁₂ O ₅	268, 211, 135, 117 /270, 167, 119	RO, OV
87	14.42	Pebrellin	373.0919	375.1074	2.65 /0.12	C ₁₉ H ₁₈ O ₈	358, 343, 161, 147 /360, 345, 327, 169	OV
88	14.61	Ladanein	313.0709	315.0864	2.74 /-0.27	C ₁₇ H ₁₄ O ₆	298, 283, 255, 148, 132 /300, 229, 133	RO, OV
89	14.93	Gardenin B		359.1123	0.64	C ₁₉ H ₁₈ O ₇	344, 326, 298, 209	OV
90	15.52	Salvigenin		329.1019	0.2	C ₁₈ H ₁₆ O ₆	314, 296, 268	RO, OV
91	16.25	Apigenin 7,4'-dimethyl ether		299.0914	0	C ₁₇ H ₁₄ O ₅	284, 256, 167, 133	RO, OV
Flavonols								
92	9.83	Rutin		611.1602	0.76	C ₂₇ H ₃₀ O ₁₆	465, 303, 287, 145, 129	RO
93	9.88	Quercetin-O-hexuronide	477.0661	479.0819	2.85 /0.24	C ₂₁ H ₁₈ O ₁₃	433, 343, 301, 113 /303, 285	RO, OV
94	10.30	Quercetin-O-hexoside (Isoquercetin)	463.0866	465.1027	3.45 /0.11	C ₂₁ H ₂₀ O ₁₂	301, 287, 175, 151 /303, 251, 145, 127	OV
95	10.31	Isorhamnetin-O-rutinoside	623.1388	625.1764	20 /-0.14	C ₂₈ H ₃₂ O ₁₆	477, 315 /479, 317	RO
96	11.06	Quercetin coumaroylhexoside	609.1235	611.1391	2.42 /0.71	C ₃₀ H ₂₆ O ₁₄	463, 301, 285 /465, 303, 287, 147	RO
97	11.97	Isorhamnetin	315.0506	317.0655	1.35 /0.25	C ₁₆ H ₁₂ O ₇	300, 271, 243 /302, 285, 121	RO, OV
98	12.87	Dimethylquercetin (Ombuin)	329.0660	331.0813	2.05 /-0.21	C ₁₇ H ₁₄ O ₇	314, 299, 271, 241, 199 /316, 121	RO, OV
Continued								

No.	R _f (min.)	Metabolite name	Mol. Ion m/z		Δ mass (ppm)	Molecular formula	MS ² ions m/z (-)/(+)	Extract
			M-H	M + H				
Flavanones								
99	8.77	Galocatechin	305.0693		-8.57	C ₁₅ H ₁₄ O ₇	225, 97, 59	RO, OV
100	10.48	Dihydroquercetin (Taxifolin)	303.0499	305.0655	3.7 /0.26	C ₁₅ H ₁₂ O ₇	285, 125, 109 /287, 259, 231, 153	OV
101	10.81	Hesperidin (Hesperetin-O-rutinoside)	609.1807		2.94	C ₂₈ H ₃₄ O ₁₅	301, 285, 251	RO
102	11.10	Aromadendrin	287.0553	289.0706	2.82 /0.22	C ₁₅ H ₁₂ O ₆	259, 243, 201, 177, 151, 125 /271, 243, 215, 153, 149	OV
103	11.77	Eriodictyol	287.0553	289.0706	2.82 /0.22	C ₁₅ H ₁₂ O ₆	151, 135, 125, 107 /271, 179, 163, 153	OV
104	12.49	Naringenin	271.0607	273.0751	1.83 /2.39	C ₁₅ H ₁₂ O ₅	253, 151, 119, 107 /255, 179, 153	OV
105	12.78	Hesperetin	301.0711	303.0854	2.19 /3.03	C ₁₆ H ₁₄ O ₆	286, 164, 151, 136 /177, 153, 137, 117	RO, OV
106	13.84	5,4'-Dihydroxy-6,7,8-trimethoxyflavanone	345.0970	347.1127	2.82 /-0.49	C ₁₈ H ₁₈ O ₇	330, 315, 301, 283, 119 /332, 227, 197	OV
107	14.26	Sakuranetin	285.0760	287.0915	2.96 /-0.35	C ₁₆ H ₁₄ O ₅	270, 165, 119 /167, 147, 119	OV
Monoterpenes								
108	8.45	Thymoquinol-O-dihexoside	535.2014		3.41	C ₂₂ H ₃₄ O ₁₂ ·CH ₂ O ₂	489, 327, 165, 161, 101	OV
109	9.62	Thymoquinol-O-hexoside	373.1495		2.42	C ₁₆ H ₂₄ O ₇ ·CH ₂ O ₂	327, 165, 164, 113, 101	OV
110	11.96	(+)-Camphor		153.1274	-0.06	C ₁₀ H ₁₆ O	109, 95, 81	RO, OV
111	12.98	Thymol		151.1117	0.28	C ₁₀ H ₁₄ O	123, 109, 91, 81	RO, OV
Sesquiterpenoids								
112	13.74	Spathulenol		221.1899	0.42	C ₁₅ H ₂₄ O	203, 147, 133, 119	OV
113	14.11	Caryophyllene epoxide		221.1900	-0.04	C ₁₅ H ₂₄ O	203, 147, 95	RO
114	18.11	Cubebene		205.1952	-0.6	C ₁₅ H ₂₄	165, 148, 121, 93	OV
Diterpenes								
115	13.51	Carnosic acid hexoside	493.2430		2.64	C ₂₆ H ₃₈ O ₉	331, 287	RO
116	13.53	Hydroxy-O-methylrosmanol	375.1803		2.69	C ₂₁ H ₂₈ O ₆	357, 345, 331, 316, 299	RO, OV
117	14.09	Rosmanol derivative	677.3676		2.81	C ₄₀ H ₅₄ O ₉	345, 331, 315, 301, 283	RO
118	14.14	Hydroxyrosmanol	361.1647		2.66	C ₂₀ H ₂₆ O ₆	317, 299	RO, OV
119	14.69	Rosmanol derivative	689.3312		2.78	C ₄₀ H ₅₀ O ₁₀	555, 345, 283	RO
120	14.73	Sageone	299.1646	301.1799	2.23 /-0.26	C ₁₉ H ₂₄ O ₃	243, 200 /283, 259, 241, 231	RO, OV
121	14.75	Hydroxy-O-methylrosmanol derivative	719.3424		1.79	C ₄₁ H ₅₂ O ₁₁	375, 360, 345, 313, 299	RO, OV
122	14.88	Hydroxyrosmadial	359.1489	361.1643	3.09 /0.74	C ₂₀ H ₂₄ O ₆	341, 331, 315, 287/333, 317, 289, 287	RO, OV
123	14.99	Carnosol	329.1750	331.1892	2.52 /3.59	C ₂₀ H ₂₆ O ₄	314, 285, 270, 201 /313, 289, 271, 247	RO, OV
124	15.44	Oridonin	363.1804		2.5	C ₂₀ H ₂₈ O ₆	345, 333, 315, 297	RO
125	15.60	Salvinorin F	373.1648		2.3	C ₂₁ H ₂₆ O ₆	358, 343, 329, 314, 299	RO, OV
126	15.63	Rosmic acid	389.1592	391.1752	3.53 /-0.18	C ₂₁ H ₂₆ O ₇	345, 313, 301, 285 /331, 303, 285	RO, OV
127	15.81	Methylrosmanol	359.1857	361.2012	1.94 /-0.69	C ₂₁ H ₂₈ O ₅	329, 315, 283 /329, 301, 273, 109	RO, OV
128	16.11	Sageone derivative	629.3466		2.82	C ₃₉ H ₅₀ O ₇	329, 299, 285	RO, OV
129	16.40	Dimethoxyrosmanol	389.1956		3.49	C ₂₂ H ₃₀ O ₆	374, 344, 330, 313, 298	RO, OV
130	16.74	Rosmadial	343.1544	345.1697	2.03 /-0.14	C ₂₀ H ₂₄ O ₅	328, 315, 299 /317, 299, 271, 231	RO, OV
131	16.77	Rosmaridiphenol	315.1960	317.2094	1.8 /5.44	C ₂₀ H ₂₈ O ₃	285, 270, 201, 179 /299, 281, 191	RO, OV
132	16.92	Rosmanol	345.1700	347.1854	2.16 /-0.29	C ₂₀ H ₂₆ O ₅	327, 301, 283, 268, 227 /301, 259, 241	RO, OV
133	16.94	Tetrahydro-hydroxyrosmariquinone	301.1802		2.38	C ₁₉ H ₂₆ O ₃	283, 273, 258	RO, OV
134	17.23	O-Methylcarnosol	343.1907		2.27	C ₂₁ H ₂₈ O ₄	299, 284	RO, OV
135	17.83	Pisiferal		301.2163	-0.31	C ₂₀ H ₂₈ O ₂	259, 245, 231, 219, 205, 163	RO, OV
136	17.91	Demethylsalvicanol	317.2114		2.57	C ₂₀ H ₃₀ O ₃	299, 243, 191, 179	RO, OV
137	18.05	Carnosic acid	331.1904		3.26	C ₂₀ H ₂₈ O ₄	313, 287, 244, 151	RO, OV
138	18.45	O-methylcarnosic acid	345.2063	347.2193	2.41 /6.89	C ₂₁ H ₃₀ O ₄	301, 286, 271 /305, 273, 121, 109	RO, OV
139	18.79	Sugiol	299.2010		2.18	C ₂₀ H ₂₈ O ₂	283, 243, 227, 215	RO, OV
140	20.34	Rosmadial derivative	631.3989		2.39	C ₄₀ H ₅₆ O ₆	567, 343, 299	RO
141	21.49	Hydroxyrosmadial derivative	631.3989		2.39	C ₄₀ H ₅₆ O ₆	359, 331, 315	RO, OV
142	22.42	Rosmadial derivative	615.4032		3.73	C ₄₀ H ₅₆ O ₅	572, 343, 299	RO, OV
Triterpenes								
143	14.85	Asiatic acid	487.3416		2.66	C ₃₀ H ₄₈ O ₅	469, 453, 441, 409	RO
144	17.09	Pomolic acid	471.3466	473.3622	2.93/0.71	C ₃₀ H ₄₈ O ₄	453, 407 /455, 409, 191	RO, OV
145	17.36	Micromeric acid	453.3363	455.3518	2.46 /0.38	C ₃₀ H ₄₆ O ₃	409, 407, 392 /437, 409, 205	RO, OV
146	19.22	Oleanolic acid	455.3517		3	C ₃₀ H ₄₈ O ₃	409, 391	RO, OV
147	19.48	Ursolic acid	455.3519		2.56	C ₃₀ H ₄₈ O ₃	407, 315	RO, OV
Continued								

No.	R_t (min.)	Metabolite name	Mol. Ion m/z		Δ mass (ppm)	Molecular formula	MS^2 ions m/z (-)/(+)	Extract
			M-H	M+H				
148	19.51	3-oxours-12-en-20,28-olide	451.3206		2.58	$C_{30}H_{44}O_3$	407, 389	RO
149	21.25	Corosolic acid	471.3466		2.93	$C_{30}H_{48}O_4$	453, 443, 427, 409, 393	RO
Fatty acids derivatives								
150	12.69	Corchorifatty acid F	327.2170		2.13	$C_{18}H_{32}O_5$	291, 229, 211, 171	RO, OV
151	13.29	Dihydroxyhexadecanoic acid	287.2221		2.37	$C_{16}H_{32}O_4$	269, 241, 141, 99	RO, OV
152	14.48	Hydroxy-octadecatrienoic acid		295.2266	0.58	$C_{18}H_{30}O_3$	277, 259, 133, 69	RO, OV
153	14.78	hydroperoxy-octadecadienoic acid	311.2220		2.51	$C_{18}H_{32}O_4$	293, 223, 195	RO, OV
154	16.81	Linolenic Acid	277.2164	279.2320	3.25 /-0.52	$C_{18}H_{30}O_2$	259, 233, 59 /123, 109, 95, 67	RO, OV
155	19.82	Hydroxystearic acid	299.2585		2.23	$C_{18}H_{36}O_3$	253, 59	RO
156	21.09	Camaryolic acid	581.3833		2.51	$C_{36}H_{54}O_6$	549, 497, 285	RO, OV
Others								
157	0.99	Quinic acid	191.0558		1.62	$C_7H_{12}O_6$	173, 171, 127, 85	RO, OV
158	1.00	Acetyl-maltose	383.1189		1.56	$C_{14}H_{24}O_{12}$	341, 191	RO
159	9.44	Hydroxybenzaldehyde	121.0293	123.0440	1.66 /0.46	$C_7H_6O_2$	92, 65 /95, 77, 51	OV
160	10.25	Amburoside A	421.1127		3.13	$C_{20}H_{32}O_{10}$	259, 153, 109	OV
161	10.82	Lolilide		197.1172	0.11	$C_{11}H_{16}O_3$	179, 161, 133, 107	RO, OV
162	11.62	Dihydroxyanthraquinone carboxylic acid	283.0240		2.86	$C_{15}H_8O_6$	239, 211, 135	OV
163	13.93	6-Oxocamphor	165.0917		2.43	$C_{10}H_{14}O_2$	150, 135	OV
164	14.45	Hotrienol		153.1274	-0.06	$C_{10}H_{16}O$	135, 107, 97	RO, OV

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.

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 caffeoylhexaside; cpd. **67**, and acacetin-*O*-rutinoside; cpd. **71**) and one compound in *S. rosmarinus* Spenn. (quercetin coumaroylhexaside; 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 $[M-H]^-$, $[C_{27}H_{29}O_{15}]^-$) 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 Fig. 10S)³⁶. Rutin, quercetin-*O*-rutinoside, (cpd. **92**, m/z 611.1602 $[M+H]^+$, $[C_{27}H_{31}O_{16}]^+$) was confirmed by the MS^2 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 Fig. 11S)³⁷. Luteolin-*O*-rutinoside (cpd. **56**, m/z 593.1495 $[M-H]^-$, $[C_{27}H_{29}O_{15}]^-$) 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 Fig. 12S) Luteolin-*O*-pentosyl-hexoside (cpd. **61**, m/z 581.1498 $[M+H]^+$, $[C_{26}H_{29}O_{15}]^+$), showed a base peak ion in the MS^2 spectra at m/z 287 $[M+H-162-132]^+$, suggesting the respective losses of hexose and pentose (Supplementary Fig. 13S). Apigenin-*O*-rutinoside (cpd. **62**, m/z 579.1693 $[M+H]^+$, $[C_{27}H_{31}O_{14}]^+$) 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 Fig. 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 $[M+H]^+$, $[C_{22}H_{23}O_{12}]^+$) displayed a prominent product ion in MS^2 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 Fig. 15S). Apigenin-*O*-hexoside (cpd. **65**, m/z 433.1128 $[M+H]^+$, $[C_{21}H_{21}O_{10}]^+$) and apigenin-*O*-hexuronide (cpd. **66**, m/z 447.0921 $[M+H]^+$, $[C_{21}H_{19}O_{11}]^+$) are monoglycoside flavones, both demonstrated a similar base peak ion at m/z 271 $[C_{15}H_{11}O_5]^+$, 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 Fig. 16S, 17 S). Homoplantagin (Hispidulin-*O*-hexoside) (cpd. **75**, m/z 463.1235 $[M+H]^+$, $[C_{22}H_{23}O_{11}]^+$) 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 Fig. 18S).

Flavonoid aglycones were detected in the chromatograms after R_t (10.48 min.). Eriodictyol (cpd. **103**, m/z 287.0553 $[M-H]^-$, $[C_{15}H_{11}O_6]^-$) 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 ($^{1,3}A^-CO_2$) and ($^{1,4}A^-$), respectively (Supplementary Fig. 19S)³⁹ Similarly, apigenin aglycone (cpd. **77**, m/z 269.0451 $[M-H]^-$, $[C_{15}H_9O_5]^-$), demonstrated the MS^2 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 ($^{1,3}A^-CO_2$) due to *C*-ring cleavage in addition

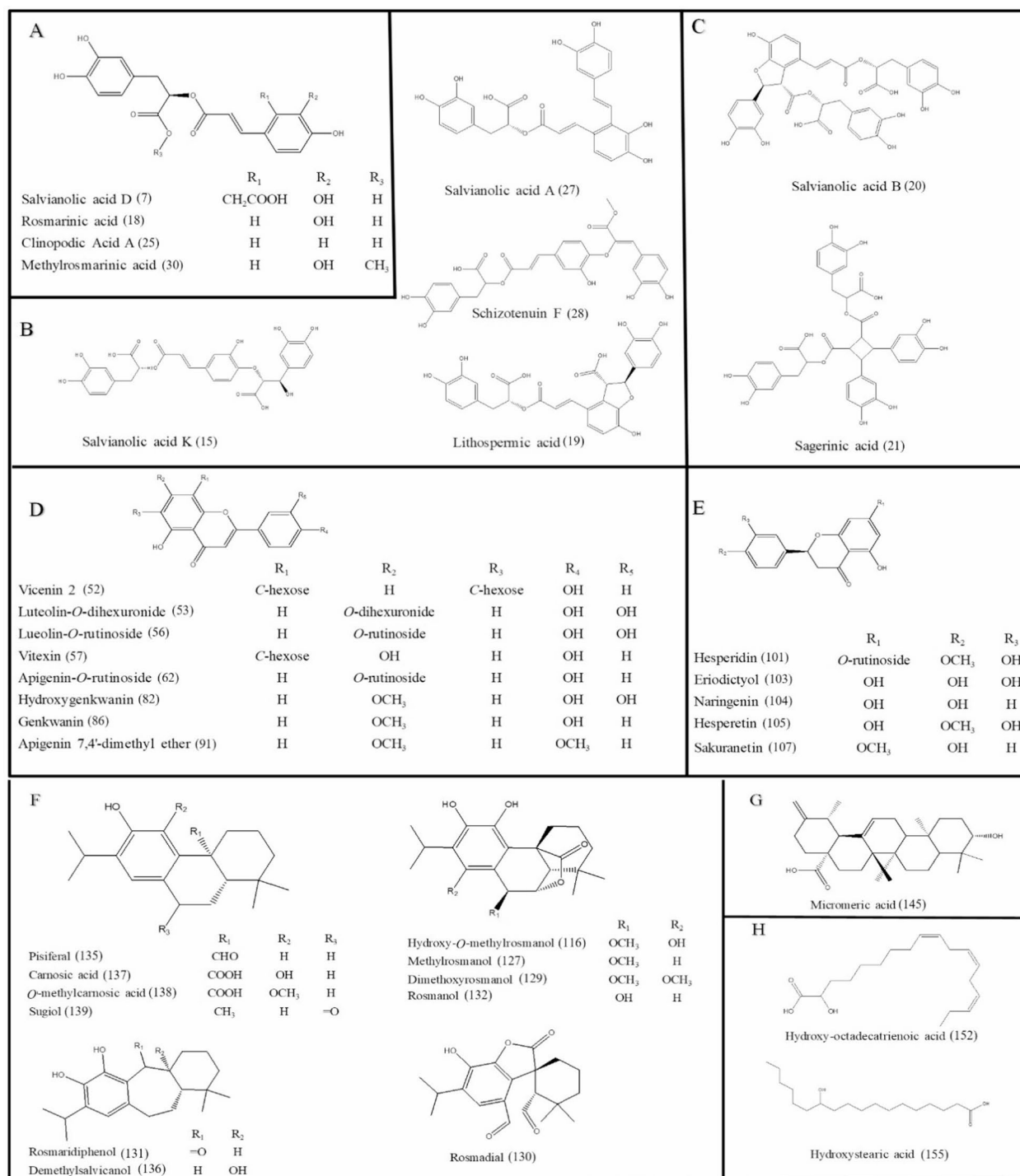


Fig. 2. Chemical structures of the major classes of metabolites identified in the rosemary & oregano extracts. The numbers listed refer to the identified metabolites listed in Table 3. **(A)** Caffeic acid dimers, **(B)** caffeic acid trimers, **(C)** caffeic acid tetramer, **(D)** flavones, **(E)** flavanones, **(F)** abietane diterpenoids, **(G)** triterpenoids, **(H)** fatty acids.

to other fragment ions resulting from small losses *vis.* m/z 225 and m/z 201 consecutive to CO_2 and C_3O_2 losses (Supplementary Fig. 20S)^{38,39}. Isorhamnetin; *O*-methyl-quercetin, (cpd. **97**, m/z 315.0506 $[\text{M-H}]^-$, $[\text{C}_{16}\text{H}_{11}\text{O}_7]^-$) was identified based on the distinctive MS^2 , 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 CO_2 and $\text{CO}_2 + \text{CO}$ groups, respectively (Supplementary Fig. 21S)⁴⁰. Hydroxygenkwanin (cpd. **82**, m/z 299.0554 $[\text{M-H}]^-$, $[\text{C}_{16}\text{H}_{11}\text{O}_6]^-$) was suggested as on dissociation produce fragment ions at m/z 284 $[\text{M-H-15}]^-$ denoting the loss of a methyl, m/z 256 $[\text{M-H-15-28}]^-$ indicating the losses of methyl and CO groups, m/z 227 $[\text{M-H-15-}$

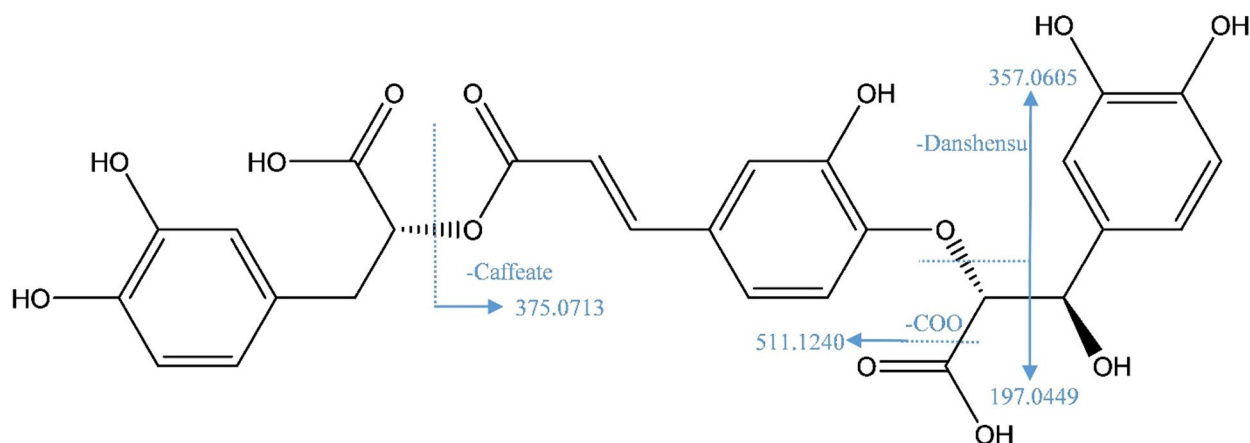


Fig. 3. Fragmentation pattern of Salvianolic acid K.

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 Fig. 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 Fig. 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 Fig. 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-methylrosmanol (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 Fig. 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 Fig. 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 Fig. 27S) and in accordance with previous reported data⁴⁷. 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 Fig. 28S). O-methylcarnosic 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 Fig. 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 properties⁴⁹. 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 Fig. 30S).

Antibacterial activity

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 ¼ MIC, reducing biofilm formation by 54.9 ± 2.3% and 61.4 ± 0.6%, respectively, and an unpaired t-test confirmed a statistically significant difference between them (p = 0.0091). Based on established criteria⁵⁰, a plant extract achieving a biofilm inhibition level greater than 50% is generally regarded as indicative of good antibiofilm activity. Accordingly,

Extract	MRSA ($\mu\text{g/mL}$)	<i>E. coli</i> ($\mu\text{g/mL}$)
RO	64	64
Aq.RO	256	128
OV	256	128
Aq.OV	64	64
Doxycycline	1	32

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.

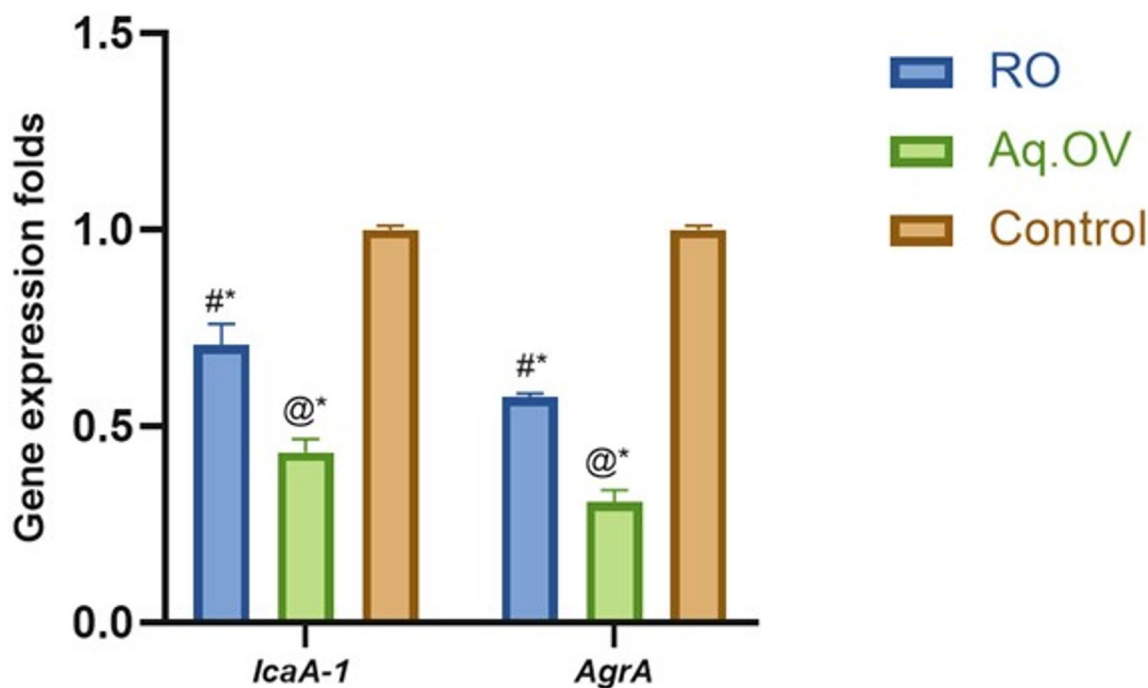


Fig. 4. Inhibition of *icaA-1* gene responsible for biofilm formation and *agrA* gene accountable for quorum sensing in MRSA. RO; *Salvia rosmarinus*, Aq.OV; Aqueous extract of *Origanum vulgare*. Data are represented as the mean \pm SD of three independent assays. Statistical analysis was performed 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 (*).

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 biofilm⁵¹.

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 MRSA⁵². 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 (Fig. 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.

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 ($\text{IC}_{50} = 6.56 \pm 0.035 \mu\text{g/mL}$) compared to the methanolic extract of

Antioxidant activity		
	RO	OV
DPPH "IC ₅₀ " (μg/mL)	6.56 ± 0.035 [*]	17.60 ± 0.333
FRAP (μmol/L)	122.78 ± 2.092	188.60 ± 10.058 [#]

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

	COX-II IC ₅₀ (μg/mL)	TNF-α (pg/mL)	NFκB (pg/mL)
RO	11.09 ± 0.46 ^{NS}	536.6 ± 13.4 ^{**}	616.2 ± 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

Table 4. Anti-inflammatory activity of *S. rosmarinus* Spenn. And *O. vulgare* L. methanolic extracts. Results are presented as (mean ± 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.

O. vulgare L. (IC₅₀ = 17.60 ± 0.333 μg/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 ± 24.06 μmol/L) compared to *S. rosmarinus* Spenn. (122.78 ± 2.09 μmol/L) (Unpaired Student's *t*-test, $p = 0.0004$) (Table 3).

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-κB 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-κB, using ibuprofen as a standard. *S. rosmarinus* Spenn. extract demonstrated strong anti-inflammatory potential, as evidenced by its COX-II inhibitory effect (IC₅₀ = 11.09 ± 0.46 μg/mL), which was comparable to ibuprofen (IC₅₀ = 8.79 ± 0.37 μg/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-κB 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-κB ($p = 0.0102$). 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).

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-κB, and 1/TNF-α.

The PLS Model was validated by the quality of fitness and prediction of Y. The autofit of the PLS model demonstrated excellent fit ($R^2_Y \text{ cum} = 0.996$) and predictive capability ($Q^2 \text{ cum} = 1$), 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 (Fig. 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, hydroxyrosmannol, rosmic acid, dimethoxyrosmannol, 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*

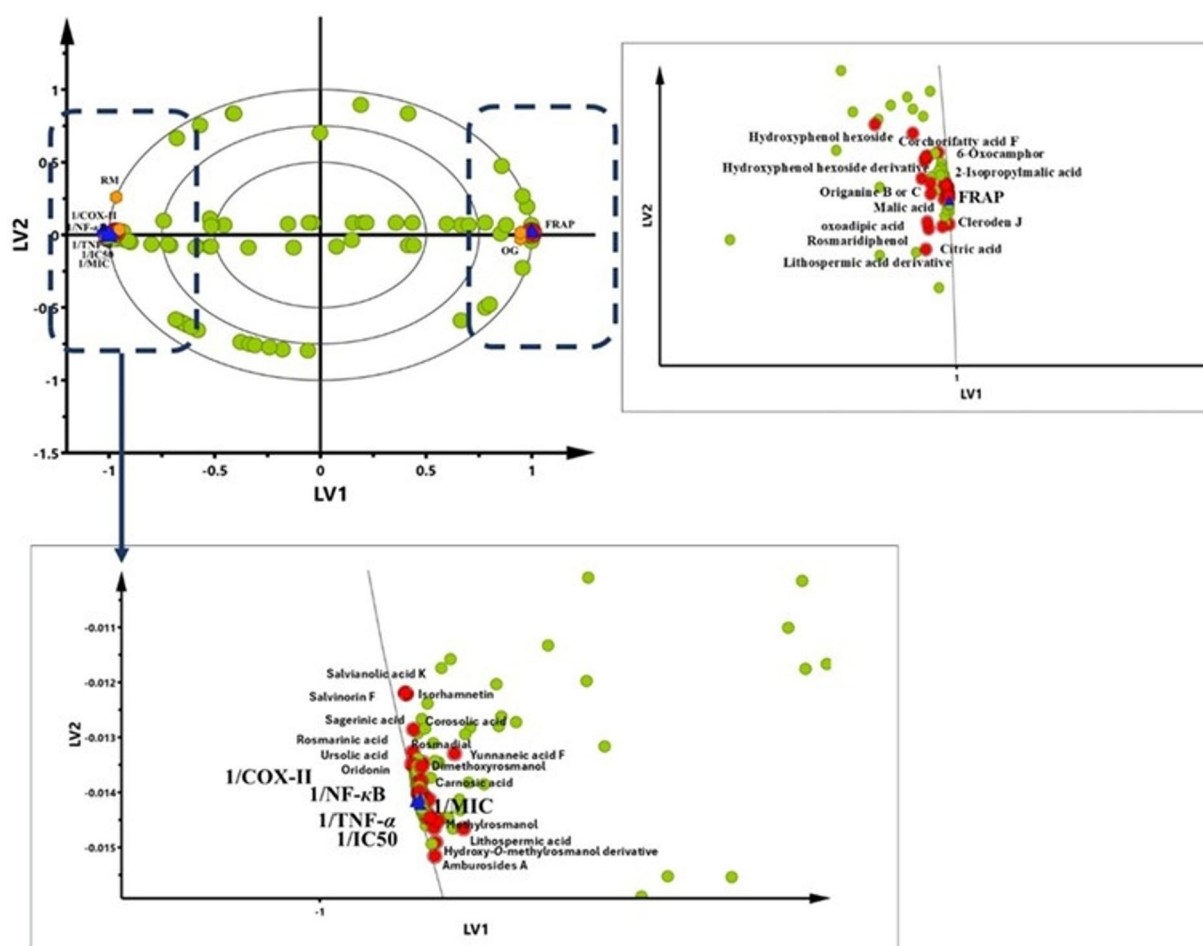


Fig. 5. PLS scores-loadings biplot describing the correlations of the identified metabolites in rosemary; RO and oregano; OV extracts and their studied bioactivities. In zoom, the detected compounds are annotated as metabolites.

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 groups⁵⁴. 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 potential⁸. 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-methylrosmanol, hydroxyrosmadial, and carnosol), which are well recognized for their anti-inflammatory effects⁵⁵. 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 cascades⁵⁶. 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 action⁵⁷.

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 (Fig. 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 (Fig. 7), leading to the selection of 66 annotated metabolites (Fig. 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. rosmarinus* Spenn. extract, namely, oridonin, carnosic acid, hydroxyrosmannol, rosmic acid, dimethoxyrosmannol, rosmadial, tetrahydro-7-hydroxyrosmariquinone, asiatic acid, corosolic acid, 3-oxours-12-en-20,28-olide, Yunnaneic acid F, salvianolic acid K, rosmarinic acid, nepitrin, isorhamnetin,

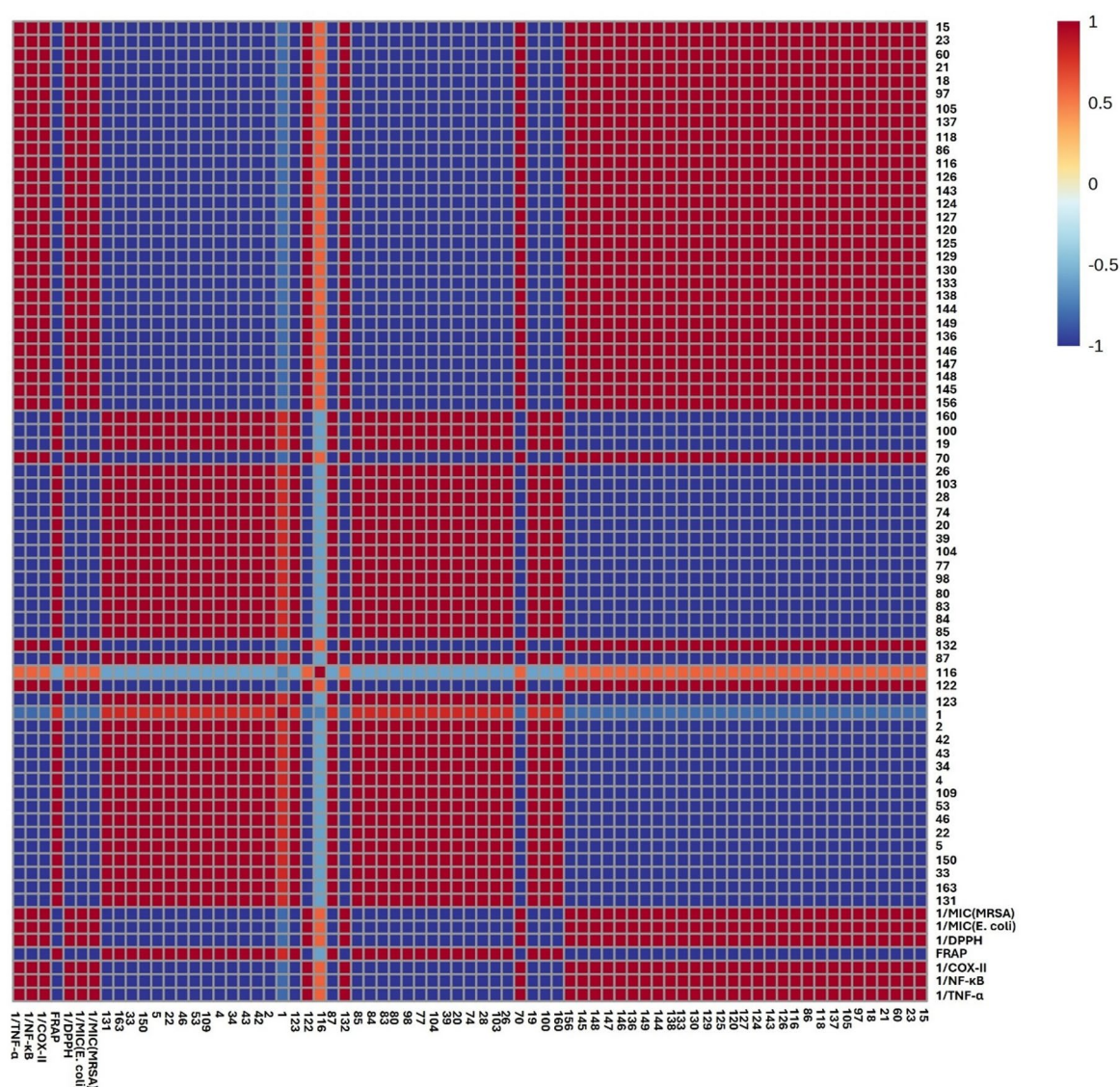


Fig. 6. Pearson's correlation between the metabolites and the antibacterial, antioxidant, and anti-inflammatory activities. Intensity of colors (blue and red) indicates correlation coefficients. The numbers listed on both axes refer to the identified metabolites listed in Table 1.

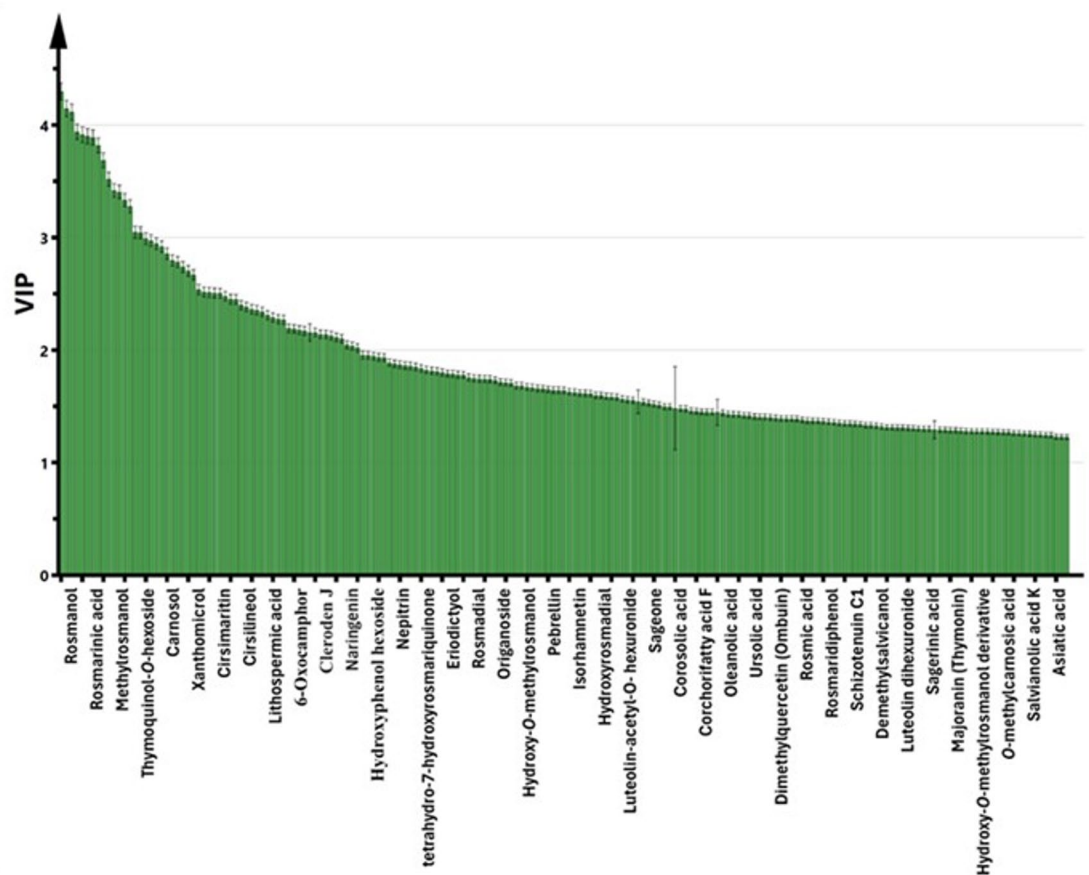


Fig. 7. Variable Importance in the Projection (VIP) plot of the PLS model for the top contributing metabolites to bioactivities (VIP ≥ 1).

and hesperetin. The antimicrobial activities of carnosic acid and rosmarinic acid are well-documented in the literature and appear to involve multiple complementary mechanisms. Carnosic acid has been reported to act as a potential quorum-sensing inhibitor, thereby suppressing bacterial virulence and biofilm formation. Additionally, its lipophilic nature allows it to incorporate into bacterial membranes, leading to membrane destabilization, increased permeability, and potential cell lysis⁵⁸. Similarly, rosmarinic acid exerts its antibacterial effects primarily through membrane disruption, inhibition of efflux pumps, interference with essential bacterial enzymes, and suppression of biofilm formation, with particularly pronounced activity against Gram-positive bacteria. These mechanisms collectively support the strong antimicrobial potential of these phenolic compounds, particularly in targeting persistent and resistant bacterial strains⁵⁹. Additionally, benzyl derivatives, organic acids & esters, and hydroxycinnamic acids are mainly present in *O. vulgare* L. extract exhibited strong positive correlations with the antioxidant activity investigated as FRAP, including hydroxyphenol hexoside, hydroxyphenol hexoside derivative, origanine B or C, lithospermic acid derivative, cleroden J, oxoadipic acid, and corchorifatty acid F. Lithospermic acid has been proven to possess significant antioxidant activity, primarily through its ability to scavenge free radicals. By neutralizing reactive oxygen species (ROS), it plays a crucial role in protecting cellular components from oxidative damage. These documented mechanisms highlight its potential as a therapeutic agent against oxidative stress-related diseases, reinforcing its value as a key natural antioxidant in plant-based systems⁶⁰. Also, the investigated anti-inflammatory activities 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 a 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.

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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Declarations

Competing interests

The authors declare no competing interests.

Additional information

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