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Biological Potential and Chemical Characterization of *Juniperus oxycedrus* Leaves and Seed Cones

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ABSTRACT: *Juniperus oxycedrus* (*J. oxycedrus*) is a traditional culinary spice and medicinal herb with a longstanding history of ethnopharmacological applications across diverse cultures. While prior research has explored the biological activities and phytochemical constituents of extracts derived from its leaves and seed cones, the present study systematically investigates their mineral and phenolic profiles alongside their multifunctional bioactive potential. Inductively coupled plasma-atomic emission spectroscopy (ICP-AES) analysis revealed a substantial abundance of essential macro- and microelements. Reversed-phase high-performance liquid chromatography (RP-HPLC) further identified high concentrations of phenolic acids (e.g., p-coumaric acid) and flavonoids (e.g., rutin and quercetin). The extracts exhibited potent radical scavenging activity against 2,2-diphenyl-1-picrylhydrazyl (DPPH), robust antioxidant capacity against hydrogen peroxide, and significant inhibition of xanthine oxidase (XO) activity. Notably, both extracts demonstrated marked antibacterial efficacy. *In silico* molecular docking studies suggested that the antimicrobial activity may stem from the phenolic constituents, which exhibited favorable binding affinities to the active site of bacterial target



proteins. These findings underscore *J. oxycedrus* as a promising reservoir of bioactive natural compounds, warranting further exploration for therapeutic and nutraceutical applications.

KEYWORDS: *Juniperus oxycedrus*; antioxidant; anti-inflammatory; antibacterial activity; chemical composition; computational biology

1 Introduction

The surge in research dedicated to phytochemistry and natural medicine has sparked a burgeoning interest in medicinal plants, owing to their potential applications in both the fields of medicine and food [1]. One such plant that has garnered attention is *Juniperus oxycedrus*, which thrives naturally in rocky regions of the Mediterranean and Near Eastern countries. Juniper seed cones, derived from this remarkable conifer, have long been celebrated in European and Scandinavian cuisines for imparting a distinctive flavor to meat dishes, particularly wild birds and game [2]. Moreover, they are an essential component of the beloved spirit, gin. Notably, the historical use of *J. oxycedrus* in traditional medicine has addressed a range of health issues, including obesity, hyperglycemia, tuberculosis, pneumonia, and bronchitis [3].

Emerging studies have further elucidated the therapeutic potential of *Juniperus oxycedrus*, with experimental evidence indicating that oral administration of its fruit and leaf extracts in streptozotocin-induced diabetic rat models elicited a statistically significant reduction in fasting blood glucose levels. Concurrently, these interventions demonstrated a marked attenuation of lipid peroxidation biomarkers, notably malondialdehyde (MDA), in both hepatic and renal tissues, suggesting antioxidative protective effects against oxidative stress-associated pathologies [4]. In Turkey, *J. oxycedrus* seed cones and leaves have been employed in the treatment of diabetes [4]. Intriguingly, polyphenols, the main class of natural antioxidants found in the compounds isolated from *J. oxycedrus*, have come to the forefront. Another study demonstrated that *J. oxycedrus* root bark exhibits substantial antioxidant activity when assessed *in vitro*, despite having fewer phenolic compounds than the needles [5]. Furthermore, extracts derived from the leaves and seed cones of *J. oxycedrus* have exhibited the ability to inhibit the growth of various microorganisms [6].

Notably, previous studies have indicated that polyphenol extracts generally exhibit greater antibacterial efficacy against Gram-positive bacteria compared to Gram-negative bacteria [7–9]. Previous *in silico* analyses, such as those by Lafraxo et al. [10], have demonstrated that the essential oil from *J. thurifera* bark exhibits inhibitory activity against the enzyme NADPH oxidase. Building on this, the present study explores the molecular docking of key compounds from *J. oxycedrus* to investigate their binding affinity and potential therapeutic relevance against *Staphylococcus aureus*. Indeed, this comprehensive study is designed to characterize aqueous extracts of *J. oxycedrus* leaves (JOL-AE) and seed cones (JOSC-AE) from Morocco. Also, the current study aimed to analyze the phenolic and flavonoid content, identify specific phenolic compounds using high-performance liquid chromatography (HPLC), and determine the mineral composition through inductively coupled plasma atomic emission spectrometry (ICP-AES). Antioxidant activities were evaluated against the DPPH• radical and hydrogen peroxide (H₂O₂), while the inhibitory potential against xanthine oxidase was assessed. Additionally, antibacterial activity against specific bacterial strains was examined. The findings of this study are expected to enhance the understanding of the medicinal and culinary applications of *Juniperus oxycedrus*, potentially supporting its broader utilization across various fields.

2 Materials and Methods

2.1 Chemicals and Reagents

The reagents and standards, including Quercetin, Ascorbic Acid, 2,2-diphenyl-1-picrylhydrazyl (DPPH[•]), Potassium dihydrogen phosphate (KH₂PO₄), Sodium hydrogen phosphate (Na₂HPO₄), Potassium ferricyanide (K₃[Fe(CN)₆]), Trichloroacetic acid (C₂HCl₃O₂), 2,3,5-triphenyltetrazolium chloride (TTC), and Iron(III) chloride hexahydrate (FeCl₃·6H₂O), were sourced from Sigma-Aldrich. Chloramphenicol was obtained from Thermo Scientific Chemicals. Mueller-Hinton agar, Mueller-Hinton broth, and Dimethyl sulfoxide (DMSO) were procured from Merck (Darmstadt, Germany). Methanol was supplied by Biosolve (Dieuze, France). Distilled water was produced using a Milli-Q water purification system (Millipore, Bedford, MA, USA).

2.2 Botanical Specimen

Leaves and seed cones of *Juniperus oxycedrus* were collected in May 2021 from the forest of the Middle Atlas Mountains in Taza City, Morocco (33°57'11.14" N, 4°3'4.68" O). Identification of the species level was performed by Professor Hamid Khamar of the Scientific Institute of Rabat, Morocco, and assigned the code number RAB108846. Following collection, the plant material was transported to the laboratory, washed with distilled water, and dried in the dark at room temperature for 60 days. The stems were then removed, and only the leaves and seed cones were retained for further analysis. The dried material was subsequently ground to a fine powder with a mean particle size of less than 910 µm. Also, voucher specimens were deposited at the Laboratory of Environmental and Health, Faculty of Sciences, Moulay Ismail University, Meknes, Morocco.

2.3 Preparation of Plant Aqueous Extract

Aqueous extracts of *J. oxycedrus* leaves (JOL-AE) and seed cones (JOSC-AE) were prepared by boiling approximately 10 g of the powdered material in 500 mL of distilled water for 30 min. The resulting extracts were filtered using a Büchner funnel and subsequently freeze-dried. The lyophilized extracts were stored in a desiccator, protected from light, until further experimentation. The extraction yields were 29.96% for the leaves and 39.21% for the seed cones.

2.4 Chemical Characterizations

2.4.1 Determination of Total Phenol Content

The total phenolic content of *J. oxycedrus* extracts was determined using the Folin–Ciocalteu method, as previously described by [11] with some modifications. In brief, 500 µL of various concentrations of each sample, solubilized in methanol, were mixed with 2.5 mL of Folin–Ciocalteu reagent (10%) dissolved in ultrapure water. After 1 min, 2.5 mL of Na₂CO₃ (7.5%) was added to the mixture, and the final solution was incubated in the dark at 25°C for 30 min. A blank was prepared by mixing 500 µL of methanol with 2.5 mL of Folin–Ciocalteu reagent and 2.5 mL of 7.5% Na₂CO₃. Absorbance was then measured at 765 nm. Gallic acid was used as a standard under the same conditions to construct the calibration curve. The total phenolic content was expressed as milligrams of gallic acid equivalent per gram of dried extract weight (mg GAE/g DW extract). Three independent experiments were performed in triplicate, and results were reported as the mean ± standard error.

2.4.2 Determination of Total Flavonoids Content

The total flavonoid content of *J. oxycedrus* extracts was assessed using a modified version of the method outlined by [11]. In this procedure, 2.5 mL of the plant extract, dissolved in ultrapure water, was

combined with 2.5 mL of a 2% aluminum chloride (AlCl_3) solution. The mixture was then incubated at room temperature in darkness for 1 h. The absorbance was recorded at 420 nm using a spectrophotometer. A calibration curve was constructed using quercetin as the reference standard under identical conditions. The flavonoid content was expressed in milligrams of quercetin equivalent per gram of dried extract weight (mg QE/g extract). Each experiment was conducted in triplicate, with results presented as the mean \pm standard error.

2.4.3 Qualitative Analysis by RP-HPLC

Chromatographic analysis was conducted following a modified version of the method reported by [12]. The separation was performed using an EC NUCLEOSIL column (5 μm , C18, 100-5, 250 mm \times 4.6 mm; Macherey-Nagel, Germany) under specific conditions. The mobile phase consisted of water containing 1% formic acid (A) and acetonitrile (B), with a gradient program starting at 7% B, increasing to 25% B at 40 min, 50% B at 85 min, and reaching 100% B at 120 min. The flow rate was set at 0.5 mL/min, and UV detection was carried out at 280 nm. The injection volume was 20 μL . ChromNAV 2.0-JASCO software was used for system control and data acquisition.

2.4.4 Quantification of Phenolic Acids and Flavonoids

The quantification of phenolic compounds was performed through high-performance liquid chromatography with ultraviolet detection (HPLC-UV) employing an EC NUCLEOSIL column (5 μm , C18, 100-5, 250 \times 4.6 mm; Macherey-Nagel, Germany) under chromatographic conditions consistent with those outlined in the identification protocol. Spectral data for all chromatographic peaks were acquired at a detection wavelength of $\lambda = 280$ nm. Data processing was conducted using the HPLC software ChromNAV 2.0 (JASCO). Concentrations of phenolic acids and flavonoids present in the aqueous plant extracts were determined through interpolation against calibration curves generated using corresponding reference standards.

2.4.5 Quantification of Mineral Contents

The quantification of minor and major mineral elements in digested plant material was conducted via inductively coupled plasma atomic emission spectroscopy (ICP-AES), following a protocol adapted from [13]. Briefly, 1 g of powdered *Juniperus oxycedrus* seed cones and leaves was subjected to digestion by heating at 110°C in a mixture of 5 mL concentrated nitric acid (HNO_3) and 15 mL hydrochloric acid (HCl). Following complete digestion, the samples were cooled to ambient temperature, and the final volume was adjusted to 100 mL using ultrapure water. Digestates were analyzed in duplicate, and trace metal concentrations were directly quantified using an Agilent 5110 ICP-OES Spectrometer (PerkinElmer, Inc., Waltham, MA, USA). Instrumental parameters were maintained as follows: plasma gas flow rate at 12 L/min, auxiliary gas flow rate at 1 L/min, and nebulizer gas flow rate at 0.7 L/min. The sample introduction rate was set to 1.5 mL/min, and radio frequency (RF) power was stabilized at 1200 W to ensure optimal plasma stability and analytical precision. Spectral data acquisition and calibration were performed under these standardized conditions to ensure reproducibility.

2.5 Antioxidant Activity

2.5.1 Assessment of DPPH Radical Scavenging Activity

The DPPH $^{\bullet}$ (2,2-diphenyl-1-picrylhydrazyl) radical scavenging activity was assessed using a protocol adapted with minor modifications from [14–17]. Briefly, 1.5 mL of methanolic sample solutions was combined

with 1.5 mL of DPPH methanolic solution (0.2 mM) and incubated under dark conditions at ambient temperature for 60 min. Absorbance measurements were subsequently recorded at a wavelength of 517 nm using a Synergy 168 HI spectrophotometer (BioTek, El Segundo, CA, USA). Three separate assays were conducted in triplicate, and the results were compared to those of ascorbic acid, which served as the positive control and was tested under identical conditions.

2.5.2 Hydrogen Peroxide (H_2O_2) Scavenging Activity of Antioxidants

Excessive concentrations of reactive oxygen species (ROS) in the human body have been associated with various pathological conditions. Among these, hydrogen peroxide (H_2O_2), when present at high levels, poses a significant threat to cellular integrity [18]. This molecule is primarily generated through the catalytic conversion of the superoxide radical ($O_2^{\bullet-}$) by the enzyme superoxide dismutase (SOD). To assess hydrogen peroxide (H_2O_2) scavenging activity, the method described by [19,20] was followed. A buffer solution (pH 7.4) containing 40 mmol/L hydrogen peroxide was prepared. Subsequently, 1 mL of the aqueous extract solution or standard (ascorbic acid) at varying concentrations was mixed with 0.6 mL of the H_2O_2 solution (40 mmol/L). After a 10-min incubation period, the absorbance of hydrogen peroxide was measured using spectrophotometry at a wavelength of $\lambda = 230$ nm. The experiment was conducted in three independent trials.

2.5.3 Assessment of Xanthine Oxidase Inhibitory Activity

Xanthine oxidase (XO) is recognized as a critical enzyme mediating the oxidation of hypoxanthine to xanthine, as well as the subsequent formation of uric acid. These catalytic processes are effectively suppressed in the presence of Allopurinol or phenolic compounds [21]. The inhibition of xanthine oxidase was evaluated using a modified protocol adapted from [22,23]. In this assay, a mixture was prepared by combining 1 mL of the test extract or Allopurinol (standard) with 2.9 mL of phosphate buffer (pH 7.5), followed by the addition of 0.1 mL of xanthine oxidase enzymatic solution (0.2 units/mL) and 1 mL of 0.5 mM xanthine solution. The resultant solution was incubated at 25°C for 15 min. To terminate the enzymatic reaction, 1 mL of HCl was introduced, after which the absorbance of the mixture was recorded at a wavelength of 295 nm. A control solution was prepared under identical conditions, excluding the enzymatic component. Allopurinol was employed as the positive control.

2.6 Antibacterial Activity

2.6.1 Bacteria Strains

Four bacterial strains, comprising two Gram-negative (*Escherichia coli* ATCC 25922 and *Salmonella Typhimurium* ATCC 700408) and two Gram-positive (*Staphylococcus aureus* ATCC 29213 and *Bacillus subtilis* ATCC 6633), were utilized in the bioassays. The bacterial strains were cultivated on Mueller Hinton Agar (Biokar, Beauvais, France) and incubated at 37°C for 24 h. Fresh 24 h cultures were prepared for each experimental trial.

2.6.2 Disc-Diffusion Assay

To evaluate the inhibition diameters of the studied extracts against the tested bacteria, we used the disc diffusion assay as previously described [24] with some modifications. Bacterial suspensions standardized to a 0.5 McFarland turbidity (approximately 10^8 CFU/mL) were prepared and inoculated via uniform swabbing onto Petri dishes containing Mueller-Hinton Agar (Biokar, Beauvais, France). Subsequently, 20 μ L of each test extract was applied to sterile 6 mm diameter filter paper discs (Whatman No. 4), which were then aseptically positioned onto the agar surface. Sterile distilled water and chloramphenicol (30 μ g/disc) served as negative

and positive controls, respectively. All plates were incubated aerobically at 37°C for 24 h, after which the diameters of the zones of inhibition (including disk diameter) were measured in millimeters.

2.6.3 Determination of MIC and MBC

The minimum inhibitory concentrations (MICs) and minimum bactericidal concentrations (MBCs) were assessed using a standardized microdilution technique in 96-well microtiter plates, as outlined in CLSI guidelines [17,25], with minor adaptations. Serial two-fold dilutions of the extract (40 µL) were prepared in sterile distilled water, followed by the addition of 40 µL of bacterial suspension (106 CFU/mL) and 120 µL of Müller Hinton broth (Biokar, Beauvais, France) to each well. A negative control, comprising sterile distilled water and extract, and a positive control, containing bacterial suspension and broth alone, were included. The plates were incubated at 37°C for 24 h. Subsequently, 40 µL of 0.2 g/mL 2,3,5-triphenyl tetrazolium chloride (TTC) was introduced into each well, followed by an additional 30 min incubation at 37°C. The MIC was defined as the lowest concentration at which no visible bacterial growth was observed. For MBC determination, 5 µL aliquots from non-turbid wells were aseptically transferred onto Müller Hinton Agar (Biokar, Beauvais, France), incubated at 37°C for 24 h, and the lowest concentration yielding no bacterial colony formation was recorded as the MBC.

2.7 In Silico Analysis

2.7.1 ADMET In Silico Pharmacokinetics

The implementation of computational drug discovery methodologies may significantly diminish the necessity for experimental investigations while concurrently enhancing the likelihood of successful therapeutic development [26,27].

To evaluate the pharmacokinetic profiles of quercetin and rutin for their prospective pharmaceutical applications, ADMET (absorption, distribution, metabolism, excretion, and toxicity) profiling was employed. The online pkCSM platform was utilized to predict ADMET characteristics and patient tolerability.

Based on established criteria, intestinal absorption was classified as optimal at values exceeding 90%, whereas values below 30% were indicative of suboptimal bioavailability. The volume of distribution (VDss) was categorized as elevated at thresholds above 0.45 L/kg, reflecting favorable systemic dispersion. For blood-brain barrier (BBB) permeation, log BB values surpassing 0.3 denoted efficient trans-barrier penetration, while values below −0.1 suggested limited cerebral distribution. Central nervous system (CNS) permeability was assessed using standardized thresholds, with log PS values greater than −2 signifying adequate CNS penetration and values below −3 correlating with restricted access. These parameters collectively serve as critical benchmarks for the systematic evaluation of drug candidate viability through ADMET profiling [28,29].

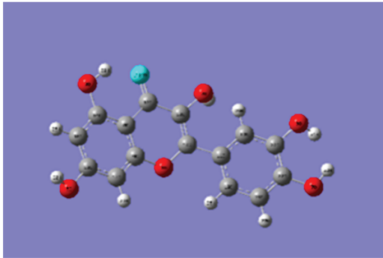
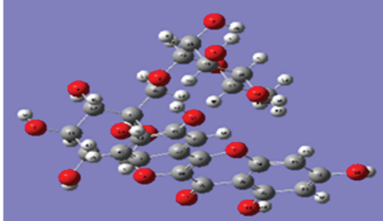
2.7.2 Molecular Docking

In this section, we focus on antibacterial activity and have chosen *S. aureus* because this bacteria strain is involved in a wide range of pathologies, with varying degrees of severity. These microorganisms have been identified as primary etiological agents of healthcare-associated infections (nosocomial infections), though their transmission is not restricted to clinical settings, with community-acquired infections being increasingly documented. Ubiquitously distributed across human and animal reservoirs, they constitute commensal components of cutaneous microbiota and demonstrate preferential colonization of external mucous membranes. Notably, environmental persistence has been well-documented in diverse ecological niches, including aqueous systems (non-potable water sources), terrestrial substrates, and contaminated fomites.

2.7.3 Preparation of Ligand

The most active compounds, rutin and quercetin, were analyzed by DFT using Gaussian 09 software (Frisch et al., 2004). The conformation was then optimized using the B3LYP/6-31G (d, p) method (# opt B3LYP/6-31G geom= connectivity) to obtain the most stable form, which was then saved under the extension (ligand.pdb) (Table 1), to be used for molecular anchoring with the 1mwt active site.

Table 1: Geometric shape of quercetin and rutin obtained using the DFT method

Compounds	Geometric structure obtained by gaussian	WM	Log P	EHOMO (ev)	ELUMO (ev)	Acceptors	Donors
Quercetin		302	1.988	-5.76	-1.43	7	5
Rutin		610	1.687	-5.23	-1.78	16	10

2.7.4 Preparation of the Active Site

The three-dimensional structural data of the target protein (PDB ID: 1MWT) were acquired from the RCSB Protein Data Bank. Structural refinement involved the removal of the co-crystallized penicillin ligand from the PLpro catalytic pocket, enabling topological analysis of ligand-interacting residues and subsequent delineation of the active site. Solvent molecules were computationally excised, and hydrogen atoms were parametrized using AutoDockTools [30] followed by molecular visualization via Discovery Studio Visualizer [31]. Ligand-protein interaction energies were quantified using the AUTOGRID37 algorithm [30]; to generate a three-dimensional affinity grid (Fig. 1). Binding free energies were estimated through empirical scoring functions, with conformational ensembles ranked based on thermodynamic stability. Computational parameters included a grid resolution of 0.35 Å and a cubic grid box spanning 60 × 60 × 60 Å³ to ensure comprehensive sampling of the catalytic domain.

2.8 Statistical Analysis

GraphPad Prism 6 software (San Diego, CA, USA) was utilized for the analysis of biological data. As the data demonstrated normal distribution and homogeneity of variance, an unpaired *t*-test with Welch's correction was performed to evaluate significant differences among the samples and relative to the control. A *p*-value of less than 0.05 was considered indicative of statistical significance.

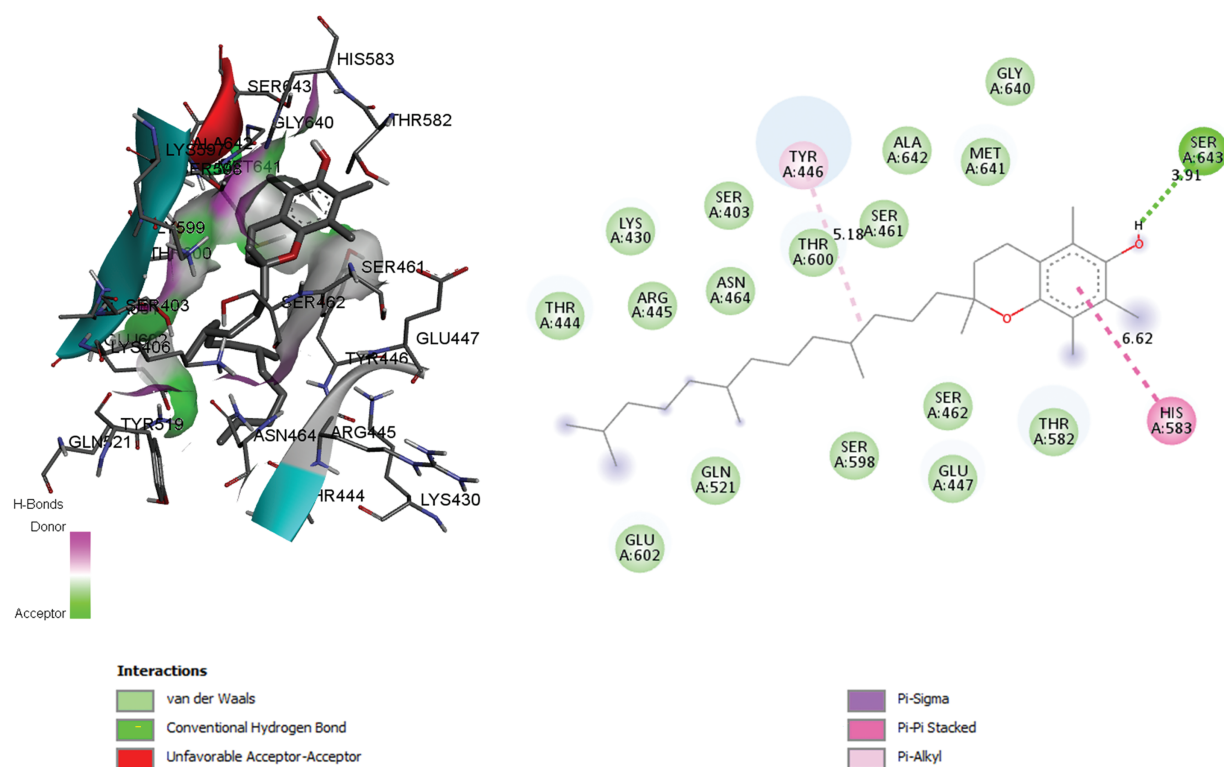


Figure 1: 2D and 3D anchor positions showing interactions of the co-crystallized compound (PENICILLIN G C₁₆ H₂₀ N₂ O₄ S) with penicillin G acyl and penicillin-binding protein 2a from *Staphylococcus aureus* strain 27r (binding energy 9.6 kcal/mol). This figure was created with Discovery Studio 3.5

3 Results and Discussion

3.1 Chemical Characterizations

3.1.1 Determination of Trace Elements Using ICP-AES

Macro-elements and microelements are the two categories of plants' mineral composition. These components play a crucial role in the cell's vital biological processes. The elemental composition of the leaves and seed cones of *J. oxycedrus* is presented in Table 2 below; the concentrations were expressed in mg/kg of the dry weight of the plant material. Macro-elements (S, K, Ca, Mg, P, Na, Fe), microelements (B, Cu, Mn, Zn, V), and heavy metals (Cd, Pb, Ni, Mo, Cr, As, Co) were determined in the studied organs of this plants. Our results show that sulfur (S) is the most abundant macro-element in both samples. The concentration of this element was 43.860 and 39.660 mg/kg respectively for leaves and seed cones. Calcium (Ca) and potassium (K) were the second most abundant elements. The concentration of potassium was 4530 and 7560 mg/kg, while that of calcium was 18820 and 5420 mg/kg respectively for leaves and seed cones. For the other macroelements, the concentrations varied between 730 and 860 mg/kg for magnesium (Mg), 584 and 710 mg/kg for phosphorus (P), 116.97 and 185.67 mg/kg for sodium (Na), and 39.51 and 217.85 mg/kg for iron (Fe).

Table 2: Mineral composition (mg/kg) of JOL-AE and JOSCAE

Minerals	JOL-AE	JOSCAE
Macro-elements		
S	43860	39660
Ca	18820	5420
K	4530	7560
Mg	860	730
P	710	584
Fe	217.85	39.51
Na	185.67	116.97
Micro-elements		
Mn	53.43	15.58
B	17.62	19.62
Zn	14.23	11.14
Cu	3.67	4.57
V	4.936	4.409
Heavy metals		
Ni	1.442	0.853
As	3.385	3.144
Mo	1.001	1.019
Cr	0.93	0.47
Pb	1.209	0.001
Cd	0.047	0.01
Co	0.001	0.001

The quantification of microelements showed that the most abundant element is manganese (Mn) with a concentration of 53.43 and 15.58 mg/kg respectively for the leaves and the seed cones of *J. oxycedrus*. Followed by boron (B) with a concentration of 17.62 and 19.62 mg/kg respectively for leaves and seed cones. The concentrations of the other microelements varied between 3.67 and 4.57 mg/kg for copper (Cu), 11.14 and 14.23 mg/kg for zinc (Zn), and 4.409 and 4.936 mg/kg for vanadium (V).

Both samples showed low concentrations of heavy metal residues in the organs studied (Table 2). Their concentrations varied between 0.047 and 0.01 mg/kg for Cadmium (Cd), 0.001 and 1.209 mg/kg for Lead (Pb), 0.853 and 1.442 mg/kg for Nickel (Ni), 1.001 and 1.019 mg/kg for molybdenum (Mo), 0.47 and 0.93 mg/kg for chromium (Cr), 3.144 and 3.385 mg/kg for arsenic (As) and 0.001 and 0.001 mg/kg for cobalt (Co).

Elevated concentrations of copper (Cu: 7.10 mg/kg), manganese (Mn: 27.79 mg/kg), chromium (Cr: 2.87 mg/kg), nickel (Ni: 10.43 mg/kg), and iron (Fe: 187.95 mg/kg) were reported in *Juniperus oxycedrus* seed cones from Turkey by [32], exceeding values quantified in the present investigation. In contrast, reduced zinc (Zn: 7.70 mg/kg) concentrations were documented relative to current findings. The seed cones and leaves of *J. oxycedrus* can be categorized as nutritious foods since they contain enough amounts of microelements and a high concentration of macronutrients. Additionally, the absence or presence of extremely low amounts of heavy metals in these plants is crucial for their therapeutic usage without toxicity.

3.1.2 Identification of Phenolic Compounds

The presence of phenolic acids and flavonoids in the aqueous extracts of *Juniperus oxycedrus* leaves and seed cones is identified through HPLC analysis. Six compounds were identified in JOL-AE, namely, para-hydroxybenzoic acid, caffeic acid, gallic acid, para-coumaric acid, caffeine, and rutin. While 10 were detected in JOSC-AE, which are Gallic Acid, Para-hydroxybenzoic Acid, Vanillic Acid, Para-coumaric Acid, Ferulic Acid, Quercetin, Catechin, Rutin, Caffeine and Vanillin (Table 3). To ensure the superposition of each standard with the corresponding molecule, we overloaded the extract with standards corresponding to the suspected molecules and we observed an increase in their concentrations.

Table 3: Phenolic profile of JOL-AE and JOSC-AE using RP-HPLC

Phenolic compounds	RT (min)	Concentrations (mg/g) of crude extract	
		JOL-AE	JOSC-AE
Caffeic acid	48.642	2.504	NI
Para-coumaric acid	54.80	0.975	0.060
Gallic acid	13.292	1.234	0.022
Para-hydroxybenzoic acid	23.91	2.937	0.204
Vanillic acid	47.308	NI	0.235
Ferulic acid	56.775	NI	NQ
Rutin	53.733	14.077	0.131
Vanillin	51.901	NI	0.072
Caffeine	48.00	8.414	0.134
Quercetin	85.133	NI	0.282
Catechin	44.092	NI	NQ

Note: NI: Not identified; NQ: Not quantified.

3.1.3 Quantification of Phenolic Compounds

The quantification of the identified compounds showed that the JOL-AE (Fig. 2) is more concentrated in phenolic compounds than that of the JOSC-AE (Fig. 3). The main compound in JOL-AE was rutin with a concentration of (14.007 mg/g of raw extract), followed by caffeine (8.414 mg/g), parahydroxybenzoic acid (2.937 mg/g), caffeic acid (2.504 mg/g), gallic acid (1.234 mg/g) and paracoumaric acid (0.957 mg/g) (Table 3). For JOSC-AE, the concentrations were (0.282 mg/g of extract) for quercetin, (0.235 mg/g) for vanillic acid, (0.204 mg/g) for para-hydroxybenzoic acid, (0.134 mg/g) for caffeine, (0.131 mg/g) for Rutin, (0.072 mg/g) for vanillin, (0.060 mg/g) for para-coumaric acid, (0.022 mg/g) for gallic acid (Table 3).

While some research has been conducted on this species, only a limited number focus on its typical consumption method (infusion). In a previous study by [33] who worked on *J. oxycedrus* collected in Amasya/Turkey. These authors noted that rutin was identified as the main compound of the methanolic extract of JOL (0.11 mg per g of extract), but its concentration was only 0.01 mg per g for seed cones extract. We also noted that the phenolic profile of the aqueous extracts of seed cones and leaves is close to that reported by [13] where they mentioned the presence of caffeic acid, para-coumaric acid, para-hydroxybenzoic acid, and rutin in the extracts of *J. oxycedrus* from Taza city, Morocco. These researchers also mentioned that rutin is the main compound of the leaf's aqueous extract with a concentration of 10.805 mg per gramme of dry extract and that the concentration of the identified compounds is very low in the seed cones extract.

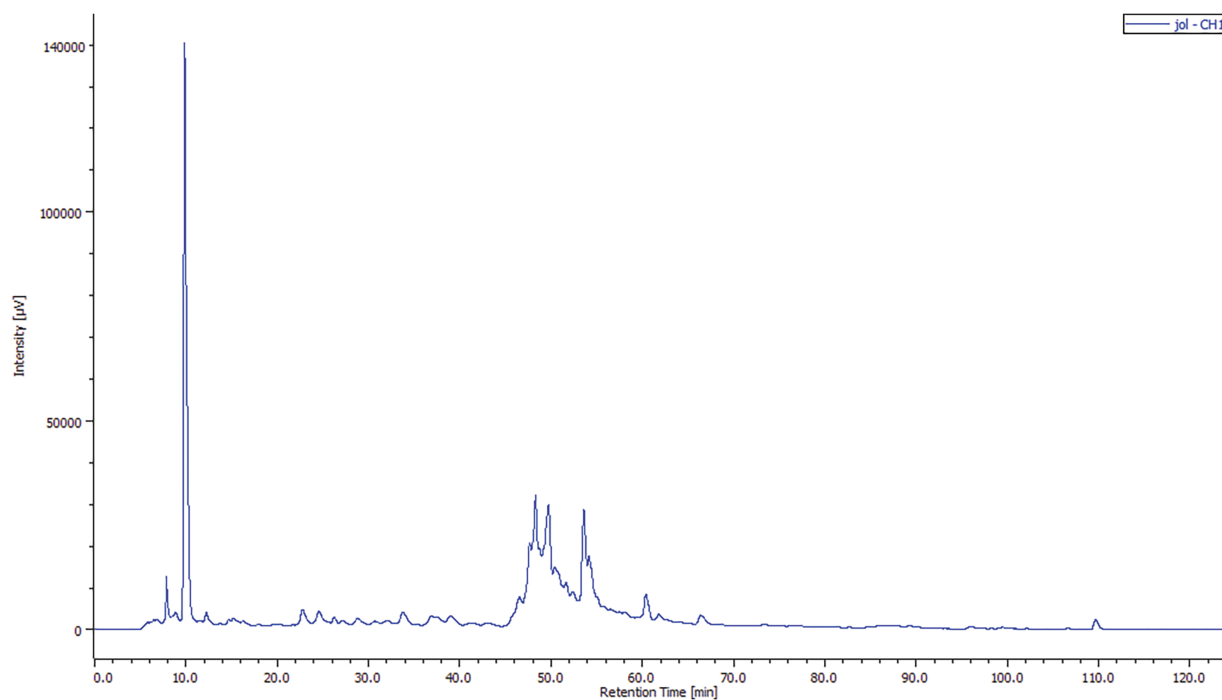


Figure 2: Chromatogram of the phenolic profile of JOL-AE by HPLC-UV

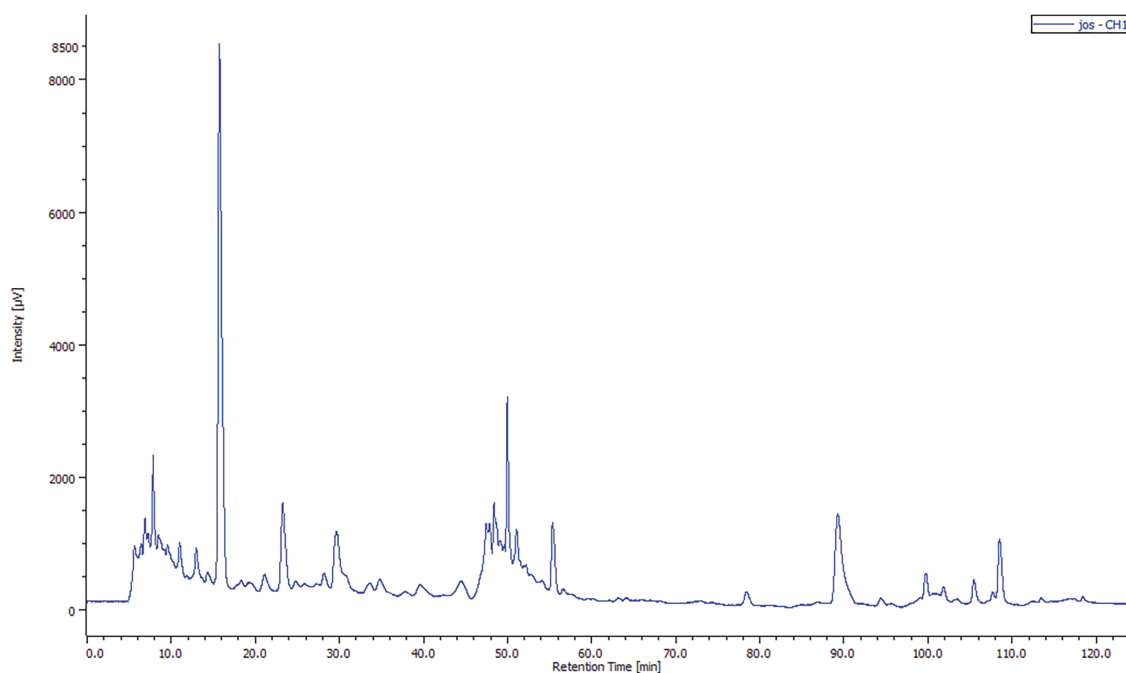


Figure 3: Chromatogram of the phenolic profile of JOSCAE-AE by HPLC-UV

3.1.4 Total Polyphenol and Flavonoid Quantification

Phenolic compounds are generally the best-known secondary metabolites in plants. These molecules are characterized by significant antioxidant activity and other beneficial biological effects for human health.

The phytochemical analysis of both extracts revealed varying levels of phenolic compounds (Table 4). Notably, JOL-AE exhibited the highest phenolic content (79.38 ± 0.82 mg GAE/g dw), whereas JOSCAE had the lowest (6.59 ± 0.27 mg GAE/g dw). A similar trend was observed for flavonoid content, with JOL-AE containing 29.47 ± 0.2 mg QE/g dw, compared to 2.69 ± 0.08 mg QE/g dw in JOSCAE.

Table 4: Total polyphenols and total flavonoid content; antioxidant activity; and xanthine oxidase inhibition ability of the aqueous extract of JOL-AE and JOSCAE

Samples	Phenolic content (mg EAG/g of dw)	Flavonoids content (mg QE/g of dw)	DPPH [•] (IC ₅₀ in µg/mL)	H ₂ O ₂ (IC ₅₀ in µg/mL)	Xanthine oxidase (IC ₅₀ in µg/mL)
JOL-AE	$79.38 \pm 0.82a$	$29.47 \pm 0.2a$	$16.05 \pm 0.15a$	$73.28 \pm 1.85b$	$46.70 \pm 1.93a$
JOSCAE	$6.59 \pm 0.27b$	$2.69 \pm 0.08b$	$9.39 \pm 0.09b$	$23.71 \pm 1.00b$	$11.66 \pm 1.30b$
Ascorbic acid	NT	NT	$52.77 \pm 0.11b$	$5.98 \pm 0.46c$	NT
Allopurinol	NT	NT	NT	NT	$51.14 \pm 0.47a$

Note: NT: Not tested. Different letters in the same row represent significant differences at $p < 0.05$.

Comparing our results with those obtained by other researchers. For *J. oxycedrus* leaves we observed that our sample showed a lower content of total phenolic compounds with a concentration of 79.38 ± 0.82 mg EAG/g of crude extract than that obtained by [13] who found a value of 147.29 mg EAG/g crude extract. On the other hand, for total flavonoids, our results showed higher content for needles extract with a content of 29.47 ± 0.2 mg EQ/g of crude extract than those reported by [34] who found a concentration of 1.68 ± 0.01 mg of EQ/g of crude extract.

For JOSCAE, few studies have focused on the identification and dosage of the phenolic compounds contained in the various aqueous extracts of this plant. Our results show that the concentration of total polyphenols is 6.59 ± 0.27 mg EAG/g of crude extract, while other authors found a higher value of 28.11 mg EAG/g of crude extract [13]. For flavonoids, our results show that this extract is poor in total flavonoids, the concentration obtained is 2.69 ± 0.08 mg of EQ/g of crude extract, this value closely approximates that obtained by [13] who found a concentration of 3.2 mg of EQ/g for the aqueous extract and of 8.3 mg of EQ/g for the methanolic extract. This difference may be caused by the use of another extraction method by these authors, as well as we cannot ignore that the subspecies were not the same, which influences the concentration of secondary metabolites of the studied samples.

3.2 Antioxidant Activity

3.2.1 DPPH[•] Radical Scavenging Activity

The evaluation of antioxidant activity in plant extracts necessitates the inclusion of the 2,2-diphenyl-1-picrylhydrazyl (DPPH[•]) radical scavenging assay as a fundamental screening method. By the protocols detailed in the methodology, *Juniperus oxycedrus* aqueous extracts (JOL-AE and JOSCAE) were subjected to DPPH[•] radical neutralization analysis. A concentration-dependent inhibitory effect was observed across experimental replicates. It was demonstrated that both extracts exhibited potent radical scavenging capacity, with half-maximal inhibitory concentrations (IC₅₀) quantified at 9.39 ± 0.09 µg/mL for JOSCAE and 16.05 ± 0.15 µg/mL for JOL-AE (Fig. 4; Table 4). Higher concentrations were excluded from the analysis due to solubility constraints. Notably, the scavenging efficacy of both extracts surpassed that of the reference antioxidant ascorbic acid, which yielded an IC₅₀ value of 52.77 ± 0.11 µg/mL.

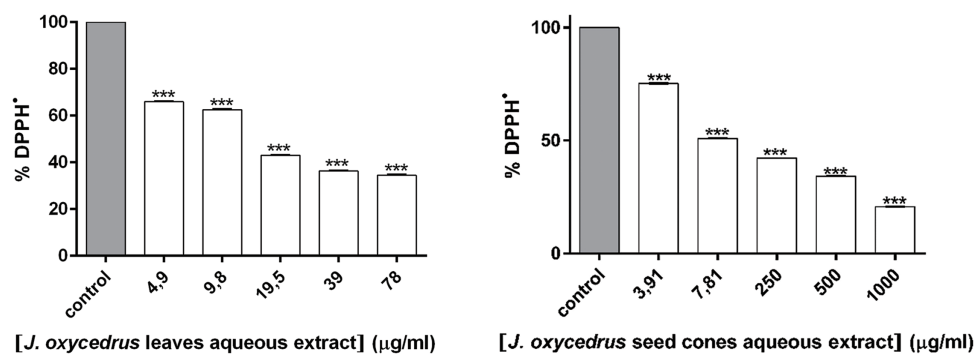


Figure 4: DPPH• scavenging activity of JOL-AE and JOSCAE. *** represent significant difference as determined by unpaired *t*-test with Welch's correction ($p < 0.001$)

For JOL-AE our results showed a better scavenging capacity against the DPPH• radical (IC_{50} value of 16.05 ± 0.15 µg extract/mL) than that reported by [35] who obtained an IC_{50} value of 140 ± 0.01 µg/mL, and similar activity to that obtained by [36] who reported an IC_{50} value of 17.91 ± 0.37 µg extract/mL, these authors used the same chemical systems and the same extraction method as well. On the other hand, [5] studied the hydro-methanolic extract (80%) and obtained a higher capacity to scavenge the radical DPPH• with an IC_{50} value of (10.95 ± 0.7) µg/mL. Conversely, the ability of JOSCAE was better than that reported by [13] who obtained an IC_{50} value of 960 ± 0.07 µg/mL. These authors used the same chemical systems and the same extraction methods. Similarly, reference [37] studied the ability of the ethanolic extract to scavenge the DPPH• radical and obtained a lower activity than ours with an IC_{50} value of 64.49 ± 0.23 µg of extract/mL. The observed variations in bioactivities may be partially attributed to discrepancies in the mass/solvent volume ratio, solvent polarity, and extraction duration employed across studies, as these parameters were not standardized. Furthermore, potential intersubspecific differences, which directly affect phytochemical constituents and their resultant pharmacological properties, cannot be disregarded as a contributory factor.

3.2.2 Antioxidant Activity against Hydrogen Peroxide (H_2O_2)

The antioxidant power of the JOL-AE and JOSCAE was studied against oxygen peroxide (H_2O_2), the obtained results show that both extracts have a strong activity against Hydrogen peroxide with IC_{50} values of 23.71 ± 1.00 µg/mL and 73.28 ± 1.85 µg/mL respectively for JOSCAE and JOL-AE. The ability of both extracts to scavenge the H_2O_2 was lower than that obtained by ascorbic acid which showed an IC_{50} value of 5.98 ± 0.46 µg/mL. The IC_{50} values obtained are presented in Table 4.

The chemical analysis showed the presence of many phenolic compounds, which are known for their antioxidant activity against Oxygen and nitrogen reactive species, which make sense of the obtained results. Other researchers [38] proved the inhibition of reactive oxygen species production in the presence of hydroethanolic extract of *J. oxycedrus* leaves using concentrations of 1.25, 2.5, and 5 µg/mL.

3.3 Inhibition of Xanthine Oxidase In-Vitro

The ability of JOL-AE and JOSCAE to inhibit xanthine oxidase activity was verified. Indeed, the seed cones extract has a good inhibitory activity against this enzyme with an IC_{50} value of 11.66 ± 1.3 µg/mL (Table 4), which was higher than that obtained by the leaf extract with an IC_{50} value of 46.7 ± 1.93 µg/mL. However, both extracts showed stronger activity than that obtained by the positive control (Allopurinol) which showed an IC_{50} value of 51.14 ± 0.47 (µg/mL).

To the best of our knowledge, the inhibition activity of JOL-AE and JOSCAE against xanthine oxidase has never performed before. Nevertheless, the activity of the studied extracts against XO is very interesting and may be due to the presence of flavonoids such as quercetin and rutin, especially *J. oxycedrus* leaves, which are known for this higher XO inhibition. The lowest concentrations of these flavonoids in *J. oxycedrus* seed cones proved that other bioactive compounds, such as terpenoids, phenolic acids, or other secondary metabolites, may also play a significant role in the observed activity (XO inhibition).

Several previous studies have mentioned that rutin and quercetin inhibit the activity of the XO enzyme [29,39]. Also, according to Cos et al. [29], a flavonoid compound with hydroxyl groups at C5 and C7 and a double bond between C2 and C3 is required for a strong inhibitory effect on xanthine oxidase. Moreover, Nagao et al. have stated that hydroxyl groups at the C5 and C7 of flavonoids were necessary for inhibiting the activity of xanthine oxidase but not a hydroxyl group at the C3 of quercetin's flavone structure [40]. Our results suggested that the amount and type of flavonoids and phenolic compounds in the extracts could influence their ability to inhibit XO activity. Our findings also agreed with those of previous reports suggesting that polyphenols and other minerals in aqueous extracts could interact synergically to increase their total antioxidant activity [41,42].

3.4 Antibacterial Activity

The antibacterial activity of the studied extracts was carried out by disc diffusion and broth micro-dilution assays, and results were presented in Table 5 and Fig. 5. Disc diffusion assay revealed that JOL-AE had the higher antibacterial activity with inhibition diameters ranging between 12.8 ± 0.3 mm and 17.2 ± 1.1 mm, while JOSCAE had inhibition diameters between 11.2 ± 0.3 mm and 15.2 ± 0.8 mm. Moreover, broth micro-dilution assay showed that JOL-AE has the highest antibacterial activity with MICs of 6.25, 12.5, 3.12, and 1.56 mg/mL against *E. coli*, *S. Typhimurium*, *S. aureus*, and *B. subtilis*, respectively. While JOSCAE showed MIC values of 12.5, 25.0, 6.25, and 6.25 mg/mL, respectively (Table 5). The ratio ($\text{MBC}/\text{MIC} \leq 4$) indicates that these extracts had bactericidal activity against the tested bacteria [43].

Table 5: MIC and MBC in mg/mL of the studied extracts against the tested bacteria

Bacteria	JOSCAE			JOL-AE			Chloramphenicol ($\mu\text{g/mL}$)		
	MIC	MBC	r	MIC	MBC	r	MIC	MBC	r
<i>E. coli</i>	12.5	50	4	6.25	25.0	4	4	16	4
<i>S. typhimurium</i>	25	100	4	12.5	50	4	64	128	2
<i>S. aureus</i>	6.25	25	4	3.12	6.25	2	4	16	4
<i>B. subtilis</i>	6.25	12.5	2	1.56	6.25	4	32	64	2

Note: Ratio (r): MBC/MIC ; (-): Inactive at the maximal tested concentration; JOSCAE: *Juniperus oxycedrus* seed cones aqueous extract; JOL-AE: *Juniperus oxycedrus* leaves aqueous extract.

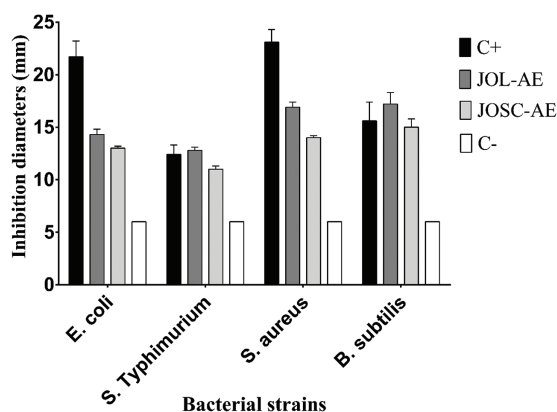


Figure 5: Inhibition diameters (mm) of the studied aqueous extracts against the tested bacteria. C+: positive control (Chloramphenicol; 30 µg); C-: negative control (sterile distilled water)

A previous Turkish study showed that the aqueous extract of *J. oxycedrus* displayed a bacteriostatic activity versus *S. aureus* (MIC of 78.12 µg/mL), while it was ineffective against Gram-negative [35]. However, Karaman and their colleagues showed that aqueous leaf extract of *J. oxycedrus* collected from Turkey was inactive against a panel of Gram-positive and Gram-negative bacteria when tested at a concentration of 300 µg/disc [44]. Taviano et al. [6] reported that methanolic extract of two *J. oxycedrus* subspecies (*oxycedrus* and *macrocarpa* (Sibth. & Sm)) from Turkey have a good antibacterial activity against *S. aureus* with a MIC value of 625.00 µg/mL [6]. These authors used a different solvent for extraction. Furthermore, we cannot ignore that the subspecies are unlikely to be identical, which also affects the chemical composition and thus the biological activity.

The antibacterial activity of plant extracts has been postulated to be associated with their distinct phytochemical profiles [45]. It is noteworthy that variations in the anatomical origin of the plant material and the quantitative variability of phenolic constituents are critical determinants of the antimicrobial efficacy observed in such extracts. Consequently, the antibacterial effects reported for *Juniperus oxycedrus* extracts may be mechanistically linked to the relative abundance of phenolic compounds, which are known to modulate bioactive properties through structure-dependent interactions [46]. In this regard, previous studies demonstrated that p-coumaric acid had strong antibacterial action against *S. aureus* and *E. coli* with MIC values of 20 and 80 µg/mL, respectively [47]. Additionally, previous studies have demonstrated the antibacterial capacity of caffeic acid, caffeine, and their derivatives; in addition to their synergic effect to enhance the antimicrobial activity of the conventional antibiotics against the drug-resistant bacteria [25,48,49]. Moreover, Rutin and its derivatives showed interesting antibacterial activity against bacteria and yeast (Al-Majmaie et al., 2019), and enhance the antibacterial activities of flavonoids against *Bacillus cereus* and *Salmonella Enteritidis* [50]. The observed variations in the phytochemicals and biological activities of *Juniperus* extracts are not only attributable to differences in geographic origin but could also stem from taxonomic distinctions between species. Prof. Adams' seminal work established that the eastern *J. oxycedrus*, commonly found in Turkey and surrounding regions, is a distinct species now recognized as *Juniperus deltoides* [51]. This species is taxonomically separate from the western *J. oxycedrus*, prevalent in the western Mediterranean, and from *J. macrocarpa*. The distinction between these species underscores the importance of precise taxonomic identification when interpreting comparative studies on *Juniperus* species [52]. The recognition of *J. deltoides* as a separate species highlights the potential for significant biochemical and ecological differences, which may influence the composition and activity of the extracts. Therefore, our study's findings should be viewed

within this context to avoid misleading interpretations, as these taxonomic distinctions may partially explain the differences observed.

3.5 *In Silico* Analysis

3.5.1 ADMET

The significant barrier-penetration potential of both compounds has been demonstrated in prior studies. Drug metabolism, defined as the biotransformation processes mediated by enzymatic activity within biological systems, was investigated (Table 6). Notably, inhibition of cytochrome P450 3A4 (CYP3A4) emerged as a critical pharmacological consideration, with the selected compounds exhibiting dual functionality as both substrates and competitive inhibitors of this enzyme. Pharmacokinetic evaluation further revealed reduced total clearance values, indicative of prolonged systemic retention, with one compound, Quercetin, demonstrating favorable pharmacokinetic persistence. Concurrent toxicity assessments confirmed the absence of adverse biological effects within the tested parameters. Collectively, these findings suggest that both compounds exhibit favorable pharmacokinetic profiles, characterized by efficient penetration, metabolic stability, and low toxicity, warranting further investigation for therapeutic applications.

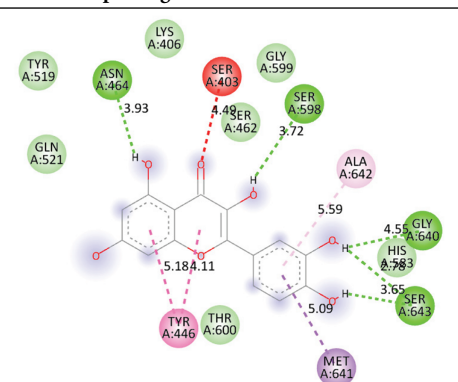
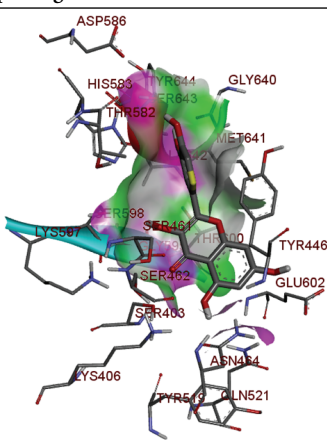
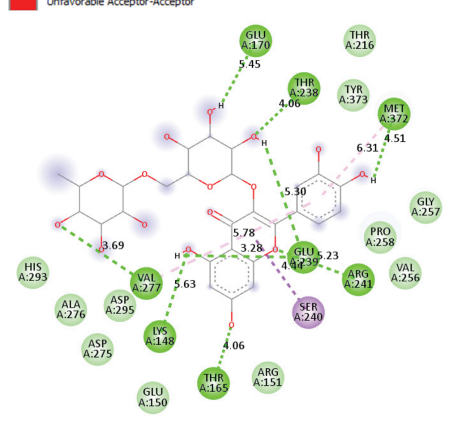
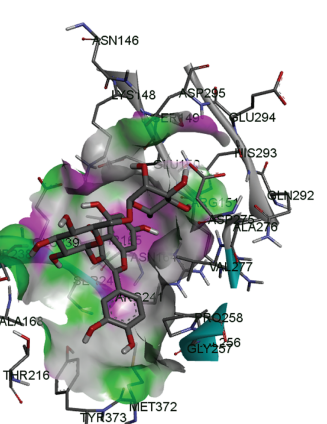
Table 6: ADMET *in silico* prediction of Quercetin and Rutin

Compounds	Absorption		Distribution		Metabolism							Excretion	Toxicity
	Intestinal absorption (human)	VDss (human)	BBB perme- ability	CNS perme- ability	CYP							Total clear- ance	AMES toxicity
					Substrate		Inhibitor						
					2D6	3A4	1A2	2C19	2C9	2D6	3A4		
					Categorical (yes/no)								
Numeric (%absorbed)	Numeric (log L/ kg)	Numeric (log BB)	Numeric (log PS)	Categorical (yes/no)							Numeric (log mL/min/kg)	Categorical (yes/no)	
Quercetin	73.048	0.223	-1.377	-3.475	No	Yes	No	yes	Yes	No	yes	0.663	No
Rutin	90.843	-0.34	-1.991	-5.89	No	No	Yes	No	No	No	yes	0.032	No

3.5.2 Docking Molecular

Quercetin exhibits a good binding pattern with the protein's active site (Table 7), which is ensured by three hydrogen bonds with Ser403, Ser598, Gly640 having a bond length of 4.11 Å, 3.44 Å, 4.00 Å, respectively, this type of interaction gives the molecule a high degree of stability in the active site, especially as it is oxygen-rich and has a higher electronegativity, making it more reactive and receptive to electrons, as can be seen with the bonds formed with the amino acid Serine. Interactions between residues Tyr446, His583, and Ala642, as well as other significant Pi-Alkyl interactions, are of great importance for the stability of the complex, particularly those with a distance less than 6 Å°. The other residues forming the complex are therefore mainly the result of Van der Waals interactions: Lys406, Tyr519, Gln521, Asn464, Glu602, Thr600, Thr582, Ser643, Gly599, and Ser462.

Table 7: 2D and 3D representation of the structure of quercetin and rutin, including active site

Compounds	Receptor-ligand interactions on a 2D	Receptor-ligand interactions on a 3D	Binding energy kcal/mol
Quercetin	 <p>Interactions</p> <ul style="list-style-type: none"> van der Waals Conventional Hydrogen Bond Unfavorable Acceptor-Acceptor <p>H-Bonds Donor (pink) Acceptor (green)</p>		-8.7
Rutin	 <p>Interactions</p> <ul style="list-style-type: none"> van der Waals Conventional Hydrogen Bond Unfavorable Acceptor-Acceptor <p>H-Bonds Donor (pink) Acceptor (green)</p> <p>Pr-Sigma, Pr-H Stacked, Pr-Alkyl</p>	 <p>H-Bonds Donor (pink) Acceptor (green)</p> <p>Pr-Sigma, Pr-H Stacked, Pr-Alkyl</p>	-7.6

In addition, the Rutin also binds well to the PLpro catalytic pocket of penicillin G acyl-Penicillin, involving eight hydrogen bonds with Glu170, Thr238, Met372, Glu239, Arg241, Thr165, Lys148 and Val277 with a bond length of 5.45 Å, 4.06 Å, 4.51 Å, 5.23 Å, 3.28 Å, 2.88 Å, 4.06 Å, 3.69 Å, respectively, and other important hydrophobic interactions via Ser240 with a distance of 4.44 Å°, other residues interact with the active site via Van der Waals interactions such as Asp164, Val165, Met208, Met 243, Ser 245, Ala246, Pro247, Pro248, Gly266, Thr308 (Table 7) This type of binding is called dipole-dipole interaction and is based on electrostatic attraction/repulsion between quercetin with fixed dipoles and the active site.

A comparison of the attacks of the two most active molecules shows that quercetin attacks virtually the same active site, while rutin attacks a different region of the protein, which explains the high antibacterial efficacy of both extracts. To sum up this theoretical section, the results obtained are very similar to the experimental results, leading us to propose these molecules as highly effective drugs against this type of bacteria.

4 Conclusion

The aim of the present study was to determine the antioxidant, anti-inflammatory, and antimicrobial potential, as well as the mineral composition of *J. oxycedrus* from the region of Taza Morocco. The aqueous extract showed strong antioxidant activity against DPPH[•] and H₂O₂ and a high potential to inhibit the catalyzing power of xanthine oxidase. The chemical composition showed the richness of this plant on phenolic acids, flavonoids, and mineralogical components with absence/minor concentrations of heavy metals. The presence of bioactivity appears to be linked in part to the major phenolic compounds, as these metabolites possess crucial structural features that influence biological activities. These findings collectively promote the use of this plant material. Furthermore, because of its abundance in bioactive natural compounds, its richness of benefit minerals in the absence of heavy metals incorporating this extract into dietary supplements and pharmaceutical formulations could prove beneficial in addressing diseases linked to inflammation, bacterial infections, and oxidative stress. *J. oxycedrus* extracts can serve as a valuable resource for the development of pharmaceutical products targeting oxidative stress and microbial infections. Furthermore, their natural origin and bioactive profile position them as excellent candidates for incorporation into nutraceutical formulations, offering health benefits through dietary supplements. Future studies should focus on scaling up the extraction process, validating the safety and efficacy through *in vivo* models, and exploring formulation strategies for clinical applications.

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Availability of Data and Materials: No datasets were generated or analyzed during this study.

Ethics Approval: Not applicable.

Conflicts of Interest: The authors declare no conflicts of interest to report regarding the present study.

Abbreviations

H ₂ O ₂	Hydrogen peroxide
XO	Xanthine oxidase
JOSC-AE	<i>Juniperus oxycedrus</i> seed cones aqueous extract
JOL-AE	<i>Juniperus oxycedrus</i> leaves aqueous extract
ICP-AES	Inductively coupled plasma atomic emission spectrometry
HPLC	High-performance liquid chromatography
TTC	2,3,5-triphenyltetrazolium chloride
FeCl ₃ ·6H ₂ O	Iron(III) chloride hexahydrate
RF	Radio frequency

MICs	Minimum inhibitory concentrations
MBCs	Minimum bactericidal concentrations
ADMET	Absorption, distribution, metabolism, excretion, and toxicity
BBB	Blood-brain barrier
CNS	Central nervous system

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