

Article

Phytochemical Characterization of *Schinus terebinthifolia* Leaf Phytobiotics and Their Potential as Natural Antibiotic Adjuvants Against Carbapenem-Resistant Bacteria

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Abstract

The emergence of carbapenem-resistant Gram-negative bacteria has intensified the search for natural compounds capable of restoring antibiotic efficacy by targeting β -lactamase-mediated resistance. This study investigated the antibacterial, β -lactamase inhibitory, and in silico potential of *Schinus terebinthifolia* leaf phytobiotics. Soxhlet (STS), maceration (STM), and essential oil (STEOL) extracts were evaluated against carbapenem-resistant *Escherichia coli*, *Klebsiella pneumoniae*, and *Pseudomonas aeruginosa* using agar diffusion, broth microdilution, checkerboard, and colorimetric β -lactamase inhibition assays. In parallel, the phytochemicals identified by GC–MS and LC–MS were virtually screened against the VIM and OXA-48-like β -lactamases using PyRx, followed by drug-likeness prediction using SwissADME. All phytobiotics exhibited significant antibacterial activity and synergistically enhanced the efficacy of cefotaxime against the majority of the tested bacterial strains ($\Sigma\text{FICI} \leq 0.5$). Moreover, they markedly inhibited β -lactamase activity, with complete inhibition achieved at MIC/2. Among the tested phytobiotics, STS exhibited the greatest β -lactamase inhibitory potency, with an IC_{50} of 49.8 $\mu\text{g/mL}$. Virtual screening identified diosmetin, catechin, quercetin, ellagic acid, and epicatechin gallate as the most promising candidate β -lactamase inhibitors, exhibiting strong binding affinities toward VIM and OXA-48-like β -lactamases. SwissADME analysis further indicated favorable drug-like properties for the top-ranked compounds. Collectively, these findings suggest that *S. terebinthifolia* phytobiotics have promising antibacterial and β -lactamase-inhibitory potential and may represent a source of candidate compounds for further investigation as antibiotic adjuvants against carbapenem-resistant Gram-negative pathogens.

Keywords: β -lactamase inhibition; cefotaxime; antibiotic potentiation; *Schinus terebinthifolia*; carbapenem resistance; molecular docking; virtual screening; drug-likeness prediction; phytochemicals



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1. Introduction

β -Lactams are among the most widely used and diverse classes of antibiotics. Their mode of action relies on interacting with and blocking penicillin-binding proteins (PBPs), which are implicated in the final steps of cell wall formation [1]. Carbapenems are a critical class of β -lactams, often employed as the ultimate therapeutic option against multidrug-resistant bacteria [2]. Carbapenems are structurally different from other β -lactams and have a broad range of antimicrobial activity and high strength against β -lactamases [2]. However, since the 2000s, the global rise in carbapenem-resistant microorganisms has become a major public health concern [3]. Resistance is particularly mediated by the production of β -lactamases, reduced membrane permeability, and efflux pumps, all of which significantly compromise therapeutic efficacy [4]. Therefore, the World Health Organization (WHO) recently identified ESKAPE bacteria (*Enterococcus faecium*, *Staphylococcus aureus*, *Klebsiella pneumoniae*, *Acinetobacter baumannii*, *Pseudomonas aeruginosa*, and *Enterobacter* species) as being in the critical group as one of the most significant pathogens driving the need for research, development, and strategic interventions to combat antimicrobial resistance [5,6]. In 2022, antimicrobial resistance led to an estimated five million deaths across the world, highlighting the grave impact of infections caused by *Enterobacterales* [7]. This growing concern highlights that carbapenem resistance constitutes a critical global health challenge, with particularly alarming implications in tropical regions [8].

In response to the rising threat of carbapenem resistance, it has become increasingly urgent to explore new antibiotics. Researchers have now shifted their focus from traditional microbial sources to alternative ones, particularly plants. Natural extracts have emerged as promising alternatives to conventional antimicrobials due to their low toxicity and multitargeting activity [9,10]. Phytobiotics are bioactive plant-derived substances, including plant extracts, essential oils, and phytochemicals such as flavonoids, phenolic acids, alkaloids, and terpenoids, that exhibit diverse biological activities, particularly antimicrobial effects. Their ability to inhibit microbial growth and potentially enhance antibiotic activity has stimulated interest in their investigation as natural antibiotic adjuvants in the context of antimicrobial resistance [11]. Their broad-spectrum activity stems from their wide diversity of secondary metabolites [12].

Hence, research on natural products is increasing, particularly in the identification and characterization of phytochemicals and plant species with antimicrobial activity, primarily against carbapenem-resistant bacteria. The flora of Brazil is particularly rich, with many species belonging to families such as *Anacardiaceae*, *Piperaceae*, *Meliaceae*, *Euphorbiaceae*, *Fabaceae*, *Amaranthaceae*, *Rubiaceae*, *Malvaceae*, *Malpighiaceae*, *Annonaceae*, *Sapindaceae*, and *Myrtaceae*. These botanical groups are known to be valuable sources of diverse phytochemicals with promising antibacterial properties [13]. However, it is essential to identify and characterize the species with antimicrobial efficacy within these families [13]. *Schinus terebinthifolia* Raddi is a Brazilian medicinal plant known for its diverse phytochemicals and antimicrobial properties [13]. Several studies have demonstrated the antimicrobial potential of *S. terebinthifolia*, supporting further investigation of this species as a source of bioactive agents [14]. Mu et al. reported that extracts from *Schinus terebinthifolia* exhibited strong activity against Gram-negative bacteria and inhibited the growth of Gram-positive food-associated microorganisms [15]. Similarly, the leaf essential oil from Zimbabwe showed inhibitory activity against a broad range of bacteria, including *Pseudomonas aeruginosa*, *Yersinia enterocolitica*, *Escherichia coli*, *Acinetobacter calcoaceticus*, *Klebsiella pneumoniae*, and *Bacillus subtilis*. This activity has been associated, at least in part, with monoterpenes, which can disrupt bacterial membrane integrity by increasing membrane permeability and fluidity, altering membrane-associated proteins, and disturbing cellular respiration and ion homeostasis [16].

Therefore, the objective of our research was to investigate the antimicrobial effects of the essential oil and two leaf extracts of *S. terebinthifolia* against reference and carbapenem-resistant Gram-negative bacteria, including *Klebsiella pneumoniae*, *Escherichia coli*, and *Pseudomonas aeruginosa*, particularly carbapenemase-producing isolates associated with human nosocomial infections. The synergistic effects of cefotaxime combined with *S. terebinthifolia* phytobiotics, together with their β -lactamase inhibitory activity, were evaluated to further investigate their potential as antibiotic adjuvants. Moreover, *in silico* analyses were performed to explore the potential interactions of phytochemical constituents with clinically relevant β -lactamases and provide mechanistic insights into the observed inhibitory activity. Although *S. terebinthifolia* has been traditionally used for medicinal purposes in several countries, in Morocco the plant is primarily encountered as an ornamental species, and its potential antimicrobial benefits remain largely unexplored. We therefore sought to investigate its potential as a source of natural antibiotic adjuvants against multidrug-resistant Gram-negative bacteria. Importantly, the novelty of this study does not lie in the identification of new phytochemical compounds but rather in providing integrated evidence linking the phytochemical profile of *S. terebinthifolia* with antibacterial activity, antibiotic potentiation, and β -lactamase inhibition against clinically relevant carbapenem-resistant pathogens. To our knowledge, this is the first study to comprehensively evaluate the potential of *S. terebinthifolia* leaf phytobiotics against carbapenemase-producing Gram-negative bacteria using a combined *in vitro* and *in silico* approach. This work provides new insights into the potential application of this traditionally underexplored Moroccan ornamental plant as a source of natural antibiotic adjuvants and contributes to the search for alternative therapeutic strategies in regions heavily affected by multidrug-resistant Gram-negative infections.

2. Materials and Methods

2.1. Botanical Material

The leaves of *S. terebinthifolia* Raddi were collected in November 2022 from the Faculty of Science Ain Chock, Hassan II University, Casablanca, Morocco (geographic coordinates: 33.5454° N, 7.6565° W). The plant samples were botanically identified based on similarity to the Herbarium of USAMV Cluj-Napoca CLA under sample code No. 30358. We rapidly washed the Brazilian pepper leaves with tap water and dried them on a paper towel. After air-drying in a dark environment for two weeks, the plant samples were finely ground with an electric blender. After grinding, the powdered leaves were stored in airtight containers at room temperature and protected from light until extraction.

2.2. Extraction of Leaves of *Schinus terebinthifolia*

Plant material was extracted using two methods: Soxhlet extraction and maceration. Approximately 62 g of the powder was weighed and extracted with ethanol for 6 h in Soxhlet equipment, and another roughly 62 g of leaf powder was extracted using maceration at room temperature on a magnetic stirrer for 72 h. Extracts were filtered through Whatman No.1 filter paper; the filtrate was concentrated, the solvent was removed, and it was then concentrated on a rotovap under reduced pressure at 79 °C. The extraction yield was expressed as the fraction of the extract mass over the dry mass of the plant material for the two ethanolic extracts. Yields were approximately 20.22% for the extract obtained by maceration (STM) and 26.82% for the one obtained by Soxhlet extraction (STS). The concentrated extract was stored at 4 °C until further analysis.

Essential oil (EO) was obtained from 200 g of *S. terebinthifolia* leaves by hydrodistillation using a Clevenger-type apparatus. The plant material was distilled with 1.5 L of water for 3 h. The EO yield was calculated on a dry-weight basis by dividing the mass

of recovered EO by the dry mass of plant material subjected to extraction. Following extraction, the EO was stored at 4 °C until chemical and biological analyses.

2.3. Chemical Composition of *Schinus terebinthifolia* Photobiotics

2.3.1. Total Phenolic Content

Total phenolic content (TPC) was assessed in the two ethanolic extracts using the Folin–Ciocalteu assay, following the procedure reported by Dirar et al. [17], with minor modifications. Briefly, each sample was reconstituted in 3 mL of methanol/water (50:50, *v/v*). The resulting suspensions were subjected to ultrasound-assisted extraction at 25 °C for 10 min at a frequency of 50 kHz. After obtaining homogeneous extracts, the supernatants were passed through 0.22 µm PTFE (polytetrafluoroethylene) membrane filters. A gallic acid (GA) calibration curve was used for quantification. TPC values were calculated from the corresponding GA calibration equation and reported as milligrams of gallic acid equivalents per gram of extract (mg GAE/g extract).

2.3.2. Liquid Chromatography–Tandem Mass Spectrometry Analysis

Polyphenolic compounds present in the *S. terebinthifolia* extracts were identified and quantified by liquid chromatography–tandem mass spectrometry (LC-MS/MS), according to the method described previously [18]. Analyses were conducted using a Thermo Scientific system (San José, CA, USA) consisting of a TSQ Quantum Ultra™ triple-quadrupole mass spectrometer fitted with a heated electrospray ionization source (HESI-II) and an Accela Open autosampler equipped with a 20 µL injection loop. Chromatographic separation was performed on a Kinetex C18 column (100 × 2.1 mm, 2.6 µm, 100 Å; Phenomenex) maintained at 50 °C. The mobile phase comprised water (solvent A) and methanol (solvent B), both containing 0.1% formic acid. The gradient program began at 5% B, followed by an increase to 90% B over 11 min. This composition was maintained for an additional 3 min before returning to the initial conditions over 9 min. The flow rate was 0.2 mL min^{−1}, and 10 µL of sample was injected for each analysis, giving a total chromatographic run time of 25 min.

The mass spectrometer was operated in both positive- and negative-ionization modes, with two or three MS/MS transitions monitored for each target compound. External calibration curves were prepared in water over the concentration range of 0.1–10 mg L^{−1} for quantitative determination. The calibration models showed good linearity, with coefficients of determination (*R*²) greater than 0.9952 for all analyzed compounds.

2.3.3. Gas Chromatography/Mass Spectrometry

The volatile constituents of the EO were characterized by gas chromatography–mass spectrometry (GC-MS). The analyses were performed on a Shimadzu GC-2010 Plus gas chromatograph equipped with a flame ionization detector (FID) and a BP-5 capillary column (30 m × 0.25 mm i.d.; film thickness, 0.25 µm; SGE Ltd, Melbourne, Australia), coupled to a Shimadzu QP2010 Plus mass spectrometer and operated using software version 2.50 SU1.

The chromatographic conditions were as follows: transfer-line temperature, 300 °C; ion-source temperature, 200 °C; helium as the carrier gas at a linear velocity of 36.5 cm s^{−1}; split ratio, 1:40; electron ionization energy, 70 eV; mass range, 40–400 u; and scan time, 1 s. Compound assignment was based primarily on retention indices calculated relative to C9–C20 *n*-alkanes analyzed under the same chromatographic conditions. Identification was further supported by matching the obtained mass spectra against the Shimadzu and NIST05 spectral databases, as well as an in-house library. The latter was established using reference essential oils, laboratory-synthesized compounds, commercially available standards, and previously reported spectral and chromatographic data [19].

2.4. Characterization of the Antibacterial Activity of *S. terebinthifolia* Phytobiotics

2.4.1. Bacterial Strains and Culture Media

Six bacterial isolates were employed in the biological effect assays: three ATCC bacteria (*Pseudomonas aeruginosa* ATCC27853 (*P. aeruginosa*), *Klebsiella pneumoniae* ATCC13883 (*K. pneumoniae*), and *E. coli* ATCC25922) and three clinical isolates resistant to carbapenems (*P. aeruginosa*, *K. pneumoniae*, and *E. coli*). They were obtained from urinary tract infections, characterized as carbapenemase-producing Gram-negative bacteria, preserved in glycerol, and stored at $-20\text{ }^{\circ}\text{C}$. All strains were routinely subcultured on various culture media, such as Cetrimide agar, MacConkey agar plates, and Luria–Bertani broth, and isolate identification was reconfirmed. Culture media were purchased from Oxoid (Basingstoke, UK).

2.4.2. Well Diffusion Method

The antibacterial effect of the three leaf phytobiotics (STEOL, STS and STM) against the six pathogenic bacteria, including the reference and the clinical strains, was determined using the agar diffusion test as outlined in previous studies [20]. Briefly, Petri plates containing Mueller–Hinton agar (MHA) were inoculated with a 10^6 CFU/mL fresh bacterial suspension. After the agar had solidified, 5 mm diameter wells were carefully formed. Then, 70 μL of each phytobiotic preparation was added to each well (10,000 $\mu\text{g/mL}$ for STEOL, 10,000 $\mu\text{g/mL}$ for STM, and 15,000 $\mu\text{g/mL}$ for STS). The STS and STM extracts were diluted in ethanol, whereas the essential oil (STEOL) was dissolved and diluted in DMSO to obtain the required concentrations. The corresponding solvent controls were included in the assay, and meropenem was included as the reference antibiotic for comparison. All plates were maintained at $4\text{ }^{\circ}\text{C}$ for 4 h and then incubated at $37\text{ }^{\circ}\text{C}$ for 24 h. The inhibition zone diameter (IZD) of bacterial growth around the well was measured at three different points in the circular zones. The experiments were repeated three times.

2.4.3. Assessment of Antibacterial Activity Using the AlamarBlue Assay

The broth microdilution method was used to determine the minimum inhibitory concentration (MIC) of the *S. terebinthifolia* phytobiotics against the six bacterial strains included in this study, according to the guidelines of the Clinical and Laboratory Standards Institute [21]. The assays were performed in sterile 96-well microtiter plates using two-fold serial dilutions of the phytobiotics at concentrations of 10, 5, 2.5, 1.25, 0.6, 0.3, 0.15, and 0.07 mg/mL. DMSO and ethanol were used as solvents, and the corresponding solvent controls were included to verify that the solvents themselves did not affect bacterial growth. A growth control containing the bacterial inoculum without phytobiotics was also included. In addition, an Alamar Blue (resazurin) assay was performed to assess the effect of *S. terebinthifolia* phytobiotics on the metabolic activity of the studied strains, based on a modification of a previously described method [22]. The MIC was considered the lowest phytobiotic concentration at which no visible bacterial growth was detected. Bacterial metabolic activity was monitored by measuring the fluorescence generated by the reduction of resazurin to resorufin. Fluorescence was recorded with a FLUOstar Omega microplate reader (BMG Labtech, Ortenberg, Germany) at 544 nm excitation and 590 nm emission wavelengths. MIC values were reported in $\mu\text{g/mL}$, taking. The concentration producing 50% inhibition of bacterial metabolic activity (IC_{50}) was estimated using GraphPad Prism version 8.0.2. Each experiment was conducted in triplicate.

2.4.4. Minimum Bactericidal Concentration Determination

The minimum bactericidal concentration (MBC) was determined by subculturing samples from MIC wells in which no visible bacterial growth was detected. Aliquots were transferred onto Mueller–Hinton agar (MHA) plates and incubated at $37\text{ }^{\circ}\text{C}$ for 24 h. The

MBC was recorded as the lowest extract concentration that resulted in a complete absence of visible colony growth following incubation [22].

2.5. Evaluation of the Interaction Between *S. terebinthifolia* Phytobiotics and Cefotaxime

The interaction between the *S. terebinthifolia* phytobiotics and cefotaxime (CTX) was investigated by the checkerboard microdilution method in 96-well plates, based on a previously described procedure [23] with minor modifications. Cefotaxime was selected as the β -lactam partner because the tested isolates exhibited resistance to this antibiotic, allowing us to assess whether the extracts and essential oil could enhance its antibacterial activity. We tested *S. terebinthifolia* extract and essential oil concentrations ($2\times$ MIC, MIC, MIC/2, MIC/4, MIC/8, and MIC/16) in combination with six CTX concentrations ranging from 0.003 to 0.1 mg/mL. The bacterial inoculum was adjusted to 1.5×10^6 CFU/mL, and the plates were incubated at 37 °C for 24 h. Following incubation, bacterial viability was assessed using resazurin. Optical density and fluorescence measurements were obtained at 544/590 nm with a FLUOstar Omega microplate reader (BMG Labtech). Three independent experiments were performed.

The fractional inhibitory concentration index (FICI) was determined according to

$$\Sigma \text{FICI} = \text{FIC(A)} + \text{FIC(B)}$$

where FIC(A) corresponds to the MIC of the *S. terebinthifolia* phytobiotic in combination divided by its MIC when tested alone, while FIC(B) represents the MIC of CTX in combination divided by the MIC of CTX alone. The interactions were interpreted as synergistic when $\Sigma \text{FICI} \leq 0.5$, additive when $0.5 < \Sigma \text{FICI} < 1.0$, non-interactive when $1 \leq \Sigma \text{FICI} < 4$, and antagonistic when $\Sigma \text{FICI} \geq 4.0$ [23].

2.6. Evaluation of the β -Lactamase Inhibitory Effect of *S. terebinthifolia* Phytobiotics

The ability of the *S. terebinthifolia* extracts to inhibit β -lactamase-mediated hydrolysis was evaluated using the Colorimetric β -Lactamase Activity Assay Kit (AAT Bioquest, Pleasanton, CA, USA), according to the manufacturer's protocol. The standardized enzyme provided with the kit was used as the β -lactamase source. Reactions were conducted in 96-well microplates and incubated at 37 °C, followed by measurement with a microplate reader.

The extent of β -lactam hydrolysis and enzyme inhibition was calculated from the optical density measurements using the following equations:

$$\% \text{ inhibition} = [(\text{OD}_{\text{control}} - \text{OD}_{\text{treated}}) / \text{OD}_{\text{control}}] \times 100$$

$$\% \text{ hydrolysis} = 100 - \% \text{ inhibition}$$

2.7. Molecular Docking Study

2.7.1. Protein Preparation

Crystal structures of the β -lactamase targets were retrieved from the RCSB Protein Data Bank in PDB format. VIM and OXA-48-like β -lactamases were selected as representative targets because of their importance in carbapenem resistance among Gram-negative bacteria and their relevance to the resistant species examined in this study (*K. pneumoniae*, *E. coli*, and *P. aeruginosa*) [24]. The corresponding PDB structures used for docking were VIM (PDB ID: 8HXO) and OXA-48-like (PDB ID: 7AW5).

Before molecular docking, the protein structures were prepared by removing crystallographic water molecules, co-crystallized ligands, and other non-protein heteroatoms. Polar hydrogen atoms and Kollman charges were added, and the receptors were converted into PDBQT format using PyRx. The active binding pockets and interacting amino acid residues

were identified using the DoGSiteScorer web server [25], and these residues were used to define the docking grid.

2.7.2. Ligand Preparation

Of the 39 phytochemicals identified in the ethanolic extracts (Soxhlet and maceration) and essential oil from *S. terebinthifolia* leaves, 38 were subjected to virtual screening to evaluate their potential as β -lactamase inhibitors. The three-dimensional chemical structures of the selected compounds were downloaded from the PubChem database (<https://pubchem.ncbi.nlm.nih.gov/>) in SDF format. The ligands were imported into PyRx 0.8, where their geometries were energy-minimized using the Open Babel module. The optimized structures were subsequently converted into PDBQT format for molecular docking.

2.7.3. Virtual Screening and Molecular Docking Simulations

Virtual screening and molecular docking calculations were carried out in PyRx v0.8 using the AutoDock Vina 1.1.2 engine. The prepared ligands were individually evaluated against the VIM and OXA-48-like β -lactamase structures. For each target, the docking region was defined according to the experimentally characterized binding site of the co-crystallized reference ligand or inhibitor. The catalytic pocket was further localized using DoGSiteScorer, and the resulting coordinates were used to establish the docking grid.

For VIM, the grid was centered at $X = 11.7385$, $Y = 1.7180$, and $Z = 32.3516$, with dimensions of $23.8230 \times 25.3451 \times 22.8063$ Å.

For OXA-48-like β -lactamase, the grid center was positioned at $X = 1.2402$, $Y = -5.7259$, and $Z = 58.3599$ Å, with dimensions of $22.3426 \times 27.1057 \times 29.0488$ Å.

Docking calculations were conducted using the standard AutoDock Vina settings available in PyRx. For each ligand, the resulting docking score was reported as the predicted binding affinity in kcal/mol. The highest-ranked pose was retained for subsequent interaction analysis. Ligand–protein interactions were examined with BIOVIA Discovery Studio Visualizer 2021, including hydrogen bonding, hydrophobic contacts, π -interactions, and the corresponding amino acid residues. The same software was used to generate the two-dimensional interaction and three-dimensional representations of the docked complexes.

2.7.4. Reference/Co-Crystallized Inhibitors and Validation of the Docking Protocol

The co-crystallized β -lactamase inhibitors, namely 2-amino-5-isobutylthiazole-4-carboxylic acid and 4-[(E)-[3-(4-chlorophenyl)-5-sulfanylidene-1H-1,2,4-triazol-4-yl]iminomethyl]benzoic acid, were used as reference ligands to benchmark the docking results. These compounds were used as structural benchmarks because they were co-crystallized with the corresponding β -lactamases in experimentally determined crystal structures.

The reliability of the docking protocol was validated by redocking the co-crystallized ligand into the active site of each enzyme. The predicted binding pose was compared with the corresponding crystallographic conformation, resulting in RMSD values of 0.703 Å for OXA-48-like β -lactamases and 1.013 Å for VIM. Both values were below the commonly accepted threshold of 2 Å, confirming the reliability and accuracy of the docking protocol.

2.8. Prediction of Drug-Likeness Properties

The physicochemical and drug-likeness characteristics of the compounds selected from the docking analysis were predicted using the SwissADME web server (<http://www.swissadme.ch/>) [26]. Canonical SMILES representations of the selected compounds were entered into the platform for computational assessment.

Drug-likeness was primarily assessed according to Lipinski's Rule of Five, considering molecular weight, lipophilicity (LogP), hydrogen-bond donor count, and hydrogen-bond acceptor count. The commonly applied thresholds were a molecular weight ≤ 500 Da, $\text{LogP} \leq 5$, no more than five hydrogen-bond donors, and no more than ten hydrogen-bond acceptors.

The Brain Or IntestinaL EstimateD permeation (BOILED-Egg) model available in SwissADME was additionally applied to estimate gastrointestinal absorption and blood–brain barrier permeability from the compounds' physicochemical characteristics, particularly lipophilicity and topological polar surface area (TPSA). The analysis also provided predictions regarding interaction with P-glycoprotein (P-gp), an efflux transporter that can influence the absorption and distribution of xenobiotic compounds.

2.9. Data Analysis

An Excel-based statistical tool was used to enter the data. Statistical analysis and visualization were performed using GraphPad Prism version 8.0.2 software. The outcomes were expressed as the average, and the measure of variability was the standard deviation derived from three independent experiments (means $\pm$ standard deviations). Data were analyzed using one-way analysis of variance (ANOVA) followed by Šídák's multiple-comparison test for MIC and MBC determinations and Tukey's multiple-comparison test for the β -lactamase inhibition assay. All experiments were performed in triplicate. Descriptive analyses, such as counts, frequencies, and percentages, were computed. p -values less than or equal to 0.05 were considered statistically significant.

3. Results

3.1. Characterization of Chemical Compounds of the Phytobiotics

3.1.1. Characterization of the Polyphenolic Compounds

The total phenolic content index (TPI) was analyzed using the Folin–Ciocalteu method. Results revealed that the extract obtained by Soxhlet had a value of $18,882 \pm 524$ mg GAE/L, while the extract obtained by maceration had a value of $14,803 \pm 496$ mg GAE/L. The LC-MS/MS analysis provided the characterization of polyphenolic profiles in the ST extracts (STS and STM), as shown in Table 1. A total of 15 compounds in the STM and 16 in the STS were reported in Table 1. Among them were flavonol constituents, mainly consisting of quercetin derivatives; the rest were represented by hydroxybenzoic acids, ellagitannins, and hydroxycinnamic acids. The major compounds were gallic acid (1142 ± 222 mg L⁻¹), followed by quercetin-3-glucoside (812 ± 51 mg L⁻¹) and quercetin (483 ± 21 mg L⁻¹) in the STS, whereas the STM contained as the main compounds quercetin (430 ± 21 mg L⁻¹), followed by quercetin-3-glucoside (422 ± 21 mg L⁻¹) and gallic acid (207 ± 53 mg L⁻¹).

Table 1. Concentrations (mg L⁻¹) of polyphenols detected in the analyzed *S. terebinthifolia* ethanolic extracts by LC–MS/MS.

Compound Class	Compound Name	Molecular Formula	STS (mg/L)	STM (mg/L)
Hydroxycinnamic acids	4-Hydroxycinnamic acid	C ₉ H ₈ O ₃	1.7 $\pm$ 0.1	1.2 $\pm$ 0.1
	Chlorogenic acid	C ₁₆ H ₁₈ O ₉	70 $\pm$ 4	18 $\pm$ 4
Hydroxybenzoic acids	Gallic acid	C ₇ H ₆ O ₅	1142 $\pm$ 222	207 $\pm$ 53
	3,4-Dihydroxybenzoic acid	C ₇ H ₆ O ₄	231 $\pm$ 13	119 $\pm$ 31

Table 1. Cont.

Compound Class	Compound Name	Molecular Formula	STS (mg/L)	STM (mg/L)
Flavan-3-ols	Catechin	C ₁₅ H ₁₄ O ₆	10 ± 2	47 ± 12
	Epicatechingallate	C ₂₂ H ₁₈ O ₁₀	62 ± 10	85 ± 7
Flavonols	Quercetin	C ₁₅ H ₁₀ O ₇	483 ± 21	430 ± 21
	Quercetin-3-rutinoside	C ₂₇ H ₃₀ O ₁₆	25 ± 2	12 ± 3
	Quercetin-3-glucoside	C ₂₁ H ₂₀ O ₁₂	812 ± 51	422 ± 12
	Quercetin-3-glucuronide	C ₂₁ H ₁₈ O ₁₃	59 ± 8	24 ± 9
	Astragalin	C ₂₁ H ₂₀ O ₁₁	5.7 ± 1.2	5.8 ± 0.7
	Isorhamnetin	C ₁₆ H ₁₂ O ₇	45 ± 8	10 ± 2
	Diosmetin	C ₁₆ H ₁₂ O ₆	4.8 ± 1.0	-
	Kaempferol	C ₁₅ H ₁₀ O ₆	15 ± 1	11 ± 2
	Myricetin	C ₁₅ H ₁₀ O ₈	203 ± 16	129 ± 11
	Ellagitannins	Ellagic acid	167 ± 23	169 ± 12

3.1.2. GC-MS Analysis of the Essential Oil of *S. terebinthifolia*

GC-MS analysis of the leaf essential oil of *S. terebinthifolia* (STEOL) led to the identification of 23 compounds. The major constituents were (+)-4-carene (64.88%), *p*-cymene (11.74%), and *trans*-2-carene-4-ol (6.75%). The obtained chromatograms clearly illustrated the distribution of the identified compounds and revealed a marked predominance of monoterpenes in the essential oil (Table 2).

Table 2. Chemical composition of the leaf essential oil of *S. terebinthifolia*.

Retention Time (Min)	Peak Area (%)	Name
4.595	0.07	2-Carene
4.934	0.34	Cyclohexene, 4-methyl-1-(1-methylethyl)-
5.423	1.15	Bicyclo[3.1.0]hex-2-ene, 2-methyl-5-(1-methylethyl)-
5.578	1.11	Cyclooctene
5.753	64.88	(+)-4-Carene
5.79	0.59	Pyrrolidine, 1,5-dimethyl-3,3-diphenyl-2-ethylidene-
5.95	11.74	<i>p</i> -Cymene
6.07	0.57	D-Limonene
6.129	4.14	Eucalyptol
8.452	0.81	Benzene, 1-methyl-3-(1-methylethyl)-
8.909	0.91	Cyclooctanone
9.403	0.25	3-Cyclohexen-1-ol, 1-methyl-4-(1-methylethyl)-
9.894	6.75	<i>trans</i> -2-Carene-4-ol
10.108	1.79	<i>trans</i> -Carveol
10.659	0.12	1-(1,2,3-Trimethyl-cyclopent-2-enyl)-ethanone

Table 2. Cont.

Retention Time (Min)	Peak Area (%)	Name
11.092	0.12	Carvenone
12.807	1.12	(1S,2R,4R,7R)-4-Isopropyl-7-methyl-3,8-dioxatricyclo[5.1.0.0 ^{2,4}]octane
13.243	0.3	(3E,5E)-2,6-Dimethylocta-3,5,7-trien-2-ol
14.488	0.77	1,4-Dihydroxy- <i>p</i> -menth-2-ene
15.197	0.6	1,4-Dihydroxy- <i>p</i> -menth-2-ene
15.855	0.15	1,3-Dioxolane, 2,2-dimethyl-4,5-bis(1-methylethenyl)-
16.563	0.3	1,3-Dioxolane, 2,2-dimethyl-4,5-di-1-propenyl-
18.256	1.44	7-Oxabicyclo[4.1.0]heptane, 1-methyl-4-(2-methyloxiranyl)-

3.2. Characterization of the Antibacterial Activity

Antibacterial activity assays using the well diffusion and microdilution methods showed that the three phytobiotics were active against both the reference and the clinical isolates. The well diffusion method was used to qualitatively assess the antibacterial effect of the ST extracts by measuring the zone of inhibition (mm). The three phytobiotics, STEOL, STS, and STM, inhibited all studied strains, particularly the STS extract, which showed strong potential with inhibition zone diameters ranging from 16.5 to 23 mm, whereas STM produced inhibition zones between 13.5 and 15 mm. The results are reported in Figure 1. Regarding the leaf essential oil (STEOL), it exhibited inhibition zones ranging from 13.5 to 16.5 mm against the tested bacterial strains.

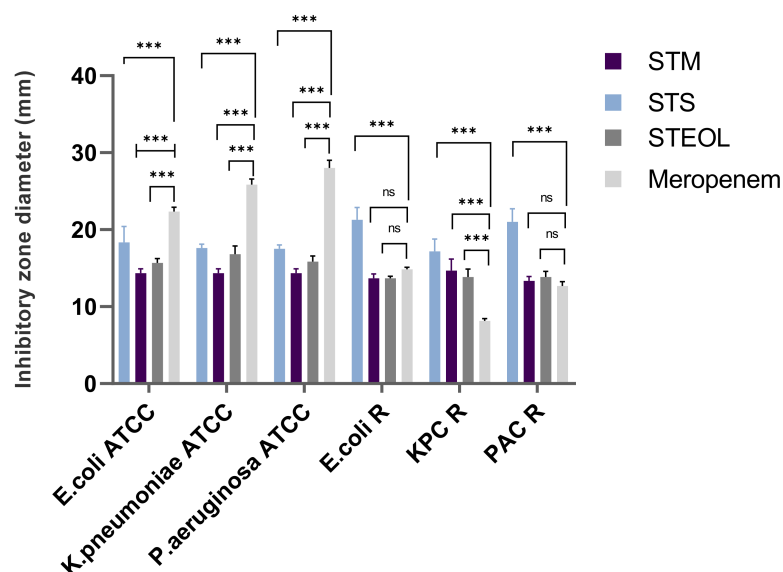


Figure 1. The antibacterial activity of *S. terebinthifolia* phytobiotics against *E. coli*, *K. pneumoniae*, and *P. aeruginosa* was determined by the well diffusion assay. *E. coli* R: carbapenem-resistant *E. coli*, KPC R: carbapenem-resistant *K. pneumoniae*, PAC R: carbapenem-resistant *P. aeruginosa*. STS: Soxhlet leaf extract, STM: leaf maceration extract, STEOL: leaf essential oil. *** indicates $p < 0.001$, while ns indicates $p > 0.05$.

The quantification assay of the antibacterial effect was conducted using the microdilution method. To confirm the impact of ST extracts on bacterial viability, a cell viability assay with Alamar Blue was performed. The results displayed in Figure 2 indicate that STS, STM, and STEOL produced a significant reduction in bacterial viability, especially at higher concentrations. Before outlining MIC and MBC values for each isolate, an

interesting antibacterial effect was noted for the STS extract, as already noted with the agar well diffusion test, which demonstrated the highest activity against the bacterial growth of the studied isolates. For STS, the MIC ranged from 310 to 766 $\mu\text{g/mL}$, and the MBC ranged from 600 to 2500 $\mu\text{g/mL}$. The MIC and MBC values of the STM extract ranged from 400 to 2000 $\mu\text{g/mL}$ and from 400 to 3300 $\mu\text{g/mL}$, respectively. In contrast, the leaf essential oil (STEOL) exhibited MIC values ranging from 925 to 1875 $\mu\text{g/mL}$ and MBC values ranging from 1250 to 5000 $\mu\text{g/mL}$.

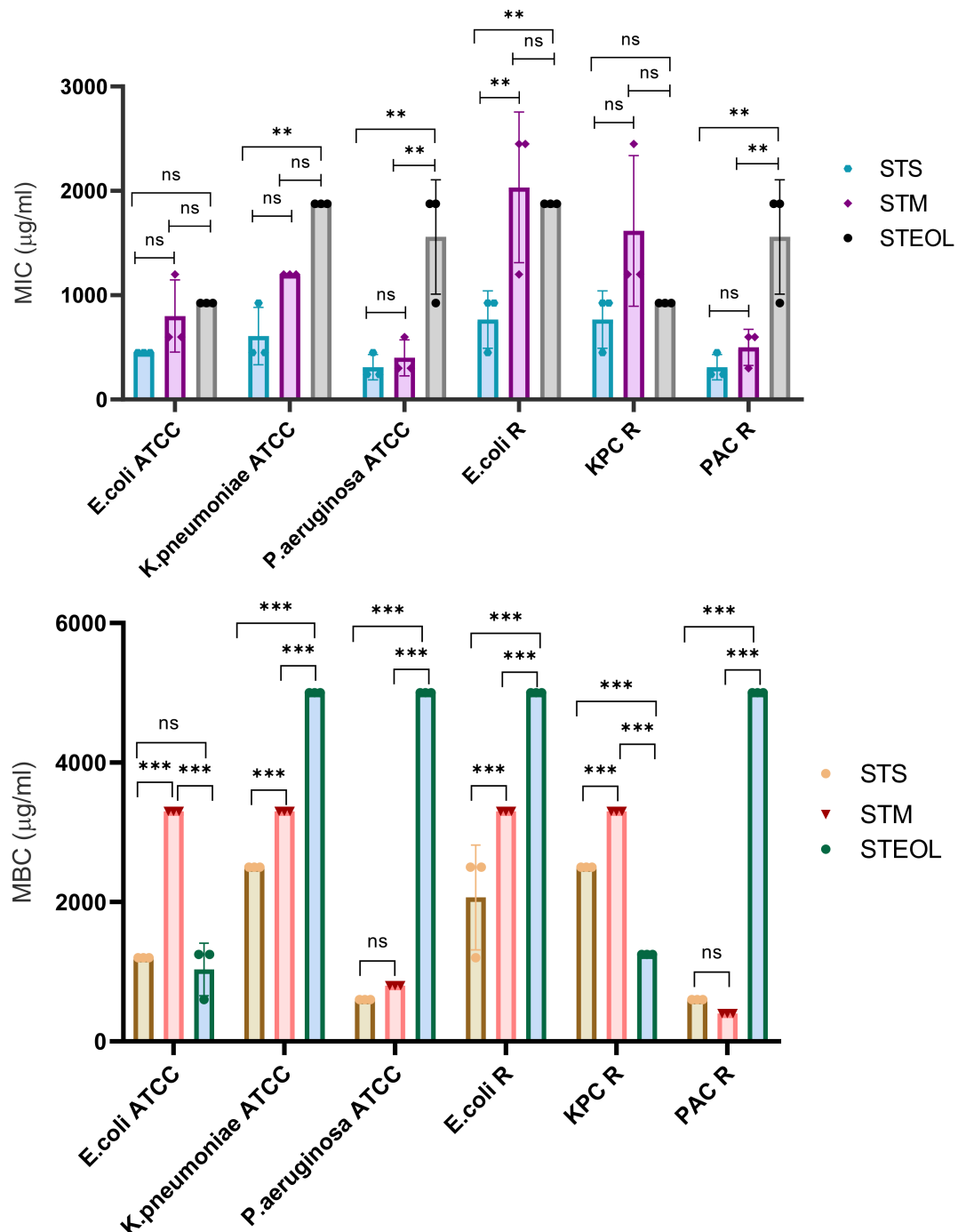


Figure 2. Minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC) of *S. terebinthifolia* phytobiotics against *E. coli*, *K. pneumoniae*, and *P. aeruginosa* strains. *E. coli* R: carbapenem-resistant *E. coli*, KPC R: carbapenem-resistant *K. pneumoniae*, PAC R: carbapenem-resistant *P. aeruginosa*. STS: Soxhlet leaf extract, STM: leaf maceration extract, STEOL: leaf essential oil. ** indicates $p < 0.01$, *** indicates $p < 0.001$, while ns indicates $p > 0.05$.

The MBC/MIC ratio was used to determine the antibacterial mode of action of the tested phytobiotics (Supplementary Materials). The STS extract exhibited a bactericidal effect ($\text{MBC/MIC} \leq 4$) against all tested strains, except for carbapenem-resistant *E. coli*, against which it showed a bacteriostatic effect. Similarly, STM exhibited bactericidal activity against all tested strains, except for *E. coli* ATCC, for which a bacteriostatic effect was observed. In contrast, the leaf essential oil (STEOL) exhibited bactericidal activity against all tested strains.

Moreover, the Alamar Blue assay showed that the three *S. terebinthifolia* phytobiotics reduced bacterial viability, with STS generally exhibiting the strongest effect, as indicated by its lower IC_{50} values (Table 3). The lowest IC_{50} value was observed for STS against *P. aeruginosa* CR ($4.74 \mu\text{g/mL} \pm 1.32$), followed by *P. aeruginosa* ATCC ($18.81 \mu\text{g/mL} \pm 0.20$) and *E. coli* ATCC ($24.66 \mu\text{g/mL} \pm 2.20$). In contrast, the highest values were observed mainly with STEOL, particularly against *K. pneumoniae* CR ($127.23 \mu\text{g/mL} \pm 0.013$) and *P. aeruginosa* CR ($114.2 \mu\text{g/mL} \pm 0.008$). Overall, the results indicate strain- and phytobiotic-dependent variation in antibacterial activity.

Table 3. The IC_{50} ($\mu\text{g/mL}$) values of *S. terebinthifolia* phytobiotics against *E. coli*, *K. pneumoniae*, and *P. aeruginosa*.

Bacteria	Extracts	Soxhlet Leaf Extract (STS)	Leaf Maceration Extract (STM)	Leaf Essential Oil (STEOL)
<i>E. coli</i> ATCC		24.66 ± 2.20	110.19 ± 3.79	69.33 ± 0.03
<i>E. coli</i> CR		83.72 ± 5.30	55.01 ± 3.47	60.03 ± 0.004
<i>K. pneumoniae</i> ATCC		92.57 ± 2.67	87.48 ± 2.35	84.12 ± 0.022
<i>K. pneumoniae</i> CR		107.66 ± 2.75	110.19 ± 3.79	127.23 ± 0.013
<i>P. aeruginosa</i> ATCC		18.81 ± 0.20	28.22 ± 1.76	81.1 ± 0.004
<i>P. aeruginosa</i> CR		4.74 ± 1.32	5.17 ± 0.79	114.2 ± 0.008

E. coli CR: carbapenem-resistant *E. coli*, KPC CR: carbapenem-resistant *K. pneumoniae*, PAC CR: carbapenem-resistant *P. aeruginosa*. Values are expressed as mean $\pm$ SD, $n = 3$.

3.3. Interaction Between *S. terebinthifolia* Phytobiotics and Cefotaxime

In our study, after confirming the growth inhibition activity exhibited by *S. terebinthifolia* phytobiotics toward the carbapenem-resistant strains, we explored the synergistic effect generated when combining the three phytobiotics with cefotaxime. Using the checkerboard method, we assessed the synergistic effect generated by the three extracts tested, which significantly decreased the MIC value of the antibiotic compared to cefotaxime alone (FIC index value was ≤ 0.5). The highest reduction in the MIC of CTX was obtained when it was combined with *S. terebinthifolia* phytobiotics against carbapenem-resistant *E. coli* and *K. pneumoniae* isolates, restoring the effect of CTX, which was ineffective alone against the two resistant bacteria. Also, the combination of STS with CTX modulated the action of CTX against carbapenem-resistant *K. pneumoniae*, carbapenem-resistant *P. aeruginosa*, and carbapenem-resistant *E. coli* strains, reducing its MIC values from $528 \mu\text{g/mL}$ to $50 \mu\text{g/mL}$, $278 \mu\text{g/mL}$ to $1.5 \mu\text{g/mL}$, and $235 \mu\text{g/mL}$ to $3.12 \mu\text{g/mL}$, respectively. The checkerboard method results are summarized in Tables 4 and 5.

Table 4. Evaluation of the synergistic effect of *S. terebinthifolia* leaf phytobiotics combined with cefotaxime against six bacterial strains.

		CTX (µg/mL)	CTX + Extract (µg/mL)	Extract (µg/mL)	Extract + CTX (µg/mL)	FIC
Soxhlet Leaf Extract (STS)	<i>E. coli</i> ATCC	12.5	0.78	450	75	0.22
	<i>E. coli</i> R	235	3.12	766	766	1.01
	<i>K. pneumoniae</i> ATCC	25	1.5	608	152	0.31
	KPC R	528	50	766	383	0.59
	<i>P. aeruginosa</i> ATCC	25	1.5	310	38.75	0.18
	PAC R	278	1.5	310	38.75	0.13
Leaf Maceration Extract (STM)	<i>E. coli</i> ATCC	12.5	0.78	800	200	0.31
	<i>E. coli</i> R	235	25	2033	2033	1.01
	<i>K. pneumoniae</i> ATCC	25	1.5	1616	269.3	0.22
	KPC R	528	5.25	1616	808	0.5
	<i>P. aeruginosa</i> ATCC	25	1.5	400	200	0.56
	PAC R	278	1.5	500	83.3	0.17
Leaf Essential Oil (STEOL)	<i>E. coli</i> ATCC	12.5	1.5	925	115.6	0.24
	<i>E. coli</i> R	235	6.25	1875	468.7	0.27
	<i>K. pneumoniae</i> ATCC	25	3.12	1875	312.5	0.29
	KPC R	528	3.12	925	231.2	0.25
	<i>P. aeruginosa</i> ATCC	25	0.78	1558	194.7	0.15
	PAC R	278	1.5	1558	194.7	0.13

E. coli CR: carbapenem-resistant *E. coli*, KPC R: carbapenem-resistant *K. pneumoniae*, PAC R: carbapenem-resistant *P. aeruginosa*, CTX: cefotaxime.

Table 5. Effect of *S. terebinthifolia* leaf phytobiotics in combination with cefotaxime against *P. aeruginosa*, *K. pneumoniae*, and *E. coli*.

	CTX/STS	CTX/STM	CTX/STEOL
<i>E. coli</i> ATCC	Synergy	Synergy	Synergy
<i>E. coli</i> CR	Indifference	Indifference	Synergy
<i>K. pneumoniae</i> ATCC	Synergy	Synergy	Synergy
KPC R	Additive	Additive	Synergy
<i>P. aeruginosa</i> ATCC	Synergy	Additive	Synergy
PAC R	Synergy	Synergy	Synergy

E. coli CR: carbapenem-resistant *E. coli*, KPC R: carbapenem-resistant *K. pneumoniae*, PAC R: carbapenem-resistant *P. aeruginosa*. STS: Soxhlet leaf extract, STM: leaf maceration extract, STEOL: leaf essential oil. CTX: cefotaxime.

3.4. β -Lactamase Inhibitory Effect of *S. terebinthifolia* Phytobiotics

The β -lactamase inhibitory activity of the three *S. terebinthifolia* leaf phytobiotics was evaluated. The inhibitory effects of the ethanolic extracts and the essential oil are presented in Figure 3. Remarkably, all three phytobiotics (STEOL, STS, and STM) exhibited strong β -lactamase inhibitory activity, achieving 100% inhibition at their MIC/2. Among them, the Soxhlet leaf extract (STS) demonstrated the highest inhibitory potency, with an IC₅₀ value of 49.8 µg/mL, indicating superior β -lactamase inhibition compared with the ethanolic extracts.

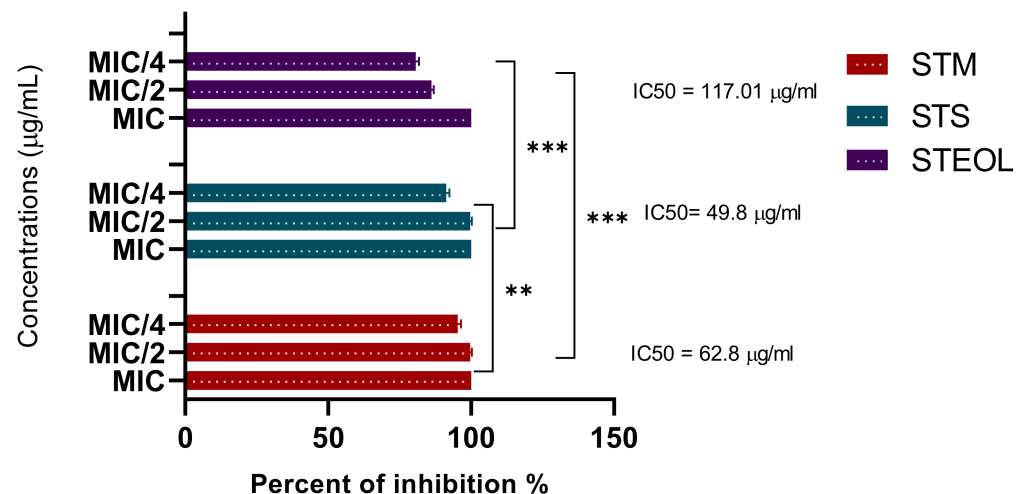


Figure 3. Percentage of β -lactamase activities inhibited by *S. terebinthifolia* leaf phytobiotics. ** indicates $p < 0.01$; *** $p < 0.001$.

3.5. In Silico Study

3.5.1. Molecular Docking Simulations

In this study, a total of 38 chemical compounds identified in the three *S. terebinthifolia* leaf phytobiotics were subjected to virtual screening against two clinically relevant β -lactamases, VIM and OXA-48-like, using PyRx software. The docking results revealed that the majority of the identified compounds exhibited higher binding affinities (more negative binding energies) than the reference antibiotic (Tables 6 and 7), suggesting a stronger interaction with the active sites of the target enzymes. Among the screened compounds, diosmetin, catechin, and quercetin exhibited the highest binding affinities for VIM, with docking scores of -9.3 , -8.9 , and -8.9 kcal/mol, respectively. Against OXA-48-like, ellagic acid, catechin, and epicatechin gallate were identified as the top-ranked compounds, with binding energies of -9.2 , -8.9 , and -8.7 kcal/mol, respectively. These findings suggest that the phytochemicals present in *S. terebinthifolia* may exhibit favorable predicted interactions within the catalytic pockets of both β -lactamases, highlighting these compounds as potential candidate inhibitors of VIM and OXA-48-like enzymes.

Table 6. Predicted binding energies (kcal/mol) of the phenolic compounds against VIM and OXA-48-like β -lactamases obtained using PyRx.

Compounds	Kcal/mol	
	OXA	VIM
4-Hydroxycinnamic acid	−5.9	−7
Chlorogenic acid	−7.8	−8.5
Gallic acid	−6	−5.6
3,4-Dihydroxybenzoic acid	−6.1	−6.4
Catechin	−8.9	−8.9
Epicatechingallate	−8.7	−7.2
Quercetine	−8.6	−8.9
Quercetin-3-rutinoside	−8.5	−8.1
Quercetin-3-glucoside	−8	−8.6

Table 6. *Cont.*

Compounds	Kcal/mol	
	OXA	VIM
Quercetin-3-glucuronide	−8.2	−8.6
Astragalin	−7.8	−8.3
Isorhamnetin	−8.4	−5.2
Diosmetin	−8.3	−9.3
Kaempferol	−8.5	−8.5
Myricetin	−8.4	−8.3
Ellagic acid	−9.2	−8.2
Reference compound	−5.9	−5.7

Table 7. Predicted binding energies (kcal/mol) of STEOL phytochemicals against VIM and OXA-48-like β -lactamases obtained using PyRx.

Compounds	Kcal/mol	
	OXA	VIM
2-Carene	−5.1	−6.9
Cyclohexene, 4-methyl-1-(1-methylethyl)-	−5.3	−6.3
Bicyclo[3.1.0]hex-2-ene, 2-methyl-5-(1-methylethyl)-	−5.1	−5.9
Cyclooctene	−4.5	−5.1
(+)-4-Carene	−5.1	−6.7
Pyrrolidine, 1,5-dimethyl-3,3-diphenyl-2-ethylidene-	−7.1	−6.3
p-Cymene	−5.3	−6.2
D-Limonene	−5.2	−6.3
Eucalyptol	−5	−5.9
Benzene, 1-methyl-3-(1-methylethyl)-	−5.2	−6.4
Cyclooctanone	−5.1	−5.5
3-Cyclohexen-1-ol, 1-methyl-4-(1-methylethyl)-	−5.1	−6.1
trans-2-Carene-4-ol	−5.4	−6.9
trans-Carveol	−5.6	−6.3
1-(1,2,3-Trimethyl-cyclopent-2-enyl)-ethanone	−5.7	−6.7
Carvenone	−6.1	−7
(1S,2R,4R,7R)-4-Isopropyl-7-methyl-3,8-dioxatricyclo[5.1.0.02,4]octane	−5.2	−6.1
(3E,5E)-2,6-Dimethylocta-3,5,7-trien-2-ol	−5.1	−6.5
1,4-Dihydroxy-p-menth-2-ene	−5.4	−6.5
1,3-Dioxolane, 2,2-dimethyl-4,5-bis(1-methylethenyl)-	−5.2	−6.1
1,3-Dioxolane, 2,2-dimethyl-4,5-di-1-propenyl-	−5.4	−5.6
7-Oxabicyclo[4.1.0]heptane, 1-methyl-4-(2-methyloxiranyl)-	−5.4	−5.8
Reference compound	−5.9	−5.7

Furthermore, to gain deeper insight into the binding mechanisms of the docked complexes, both 3D and 2D interaction analyses were performed using Discovery Studio Visualizer, as presented in Figures 4 and 5. The results showed that various intermolecular

interactions mediated the binding between the tested compounds and the three proteins, highlighting their potential to interact with key amino acid residues in the binding pocket. For example, epicatechin gallate interacts with Oxa-48 via various residues through significant interactions, including van der Waals forces with Asp101, Ser212, Thr213, Leu247, Ala69, Thr209, Ser118, Lys208, and Gly210; a Pi-Sigma interaction with Trp105; conventional hydrogen bonds (H-bonds) with Arg250 and Ser70; and Pi-Pi T-shaped interactions with Ile102, Val129, and Leu158.

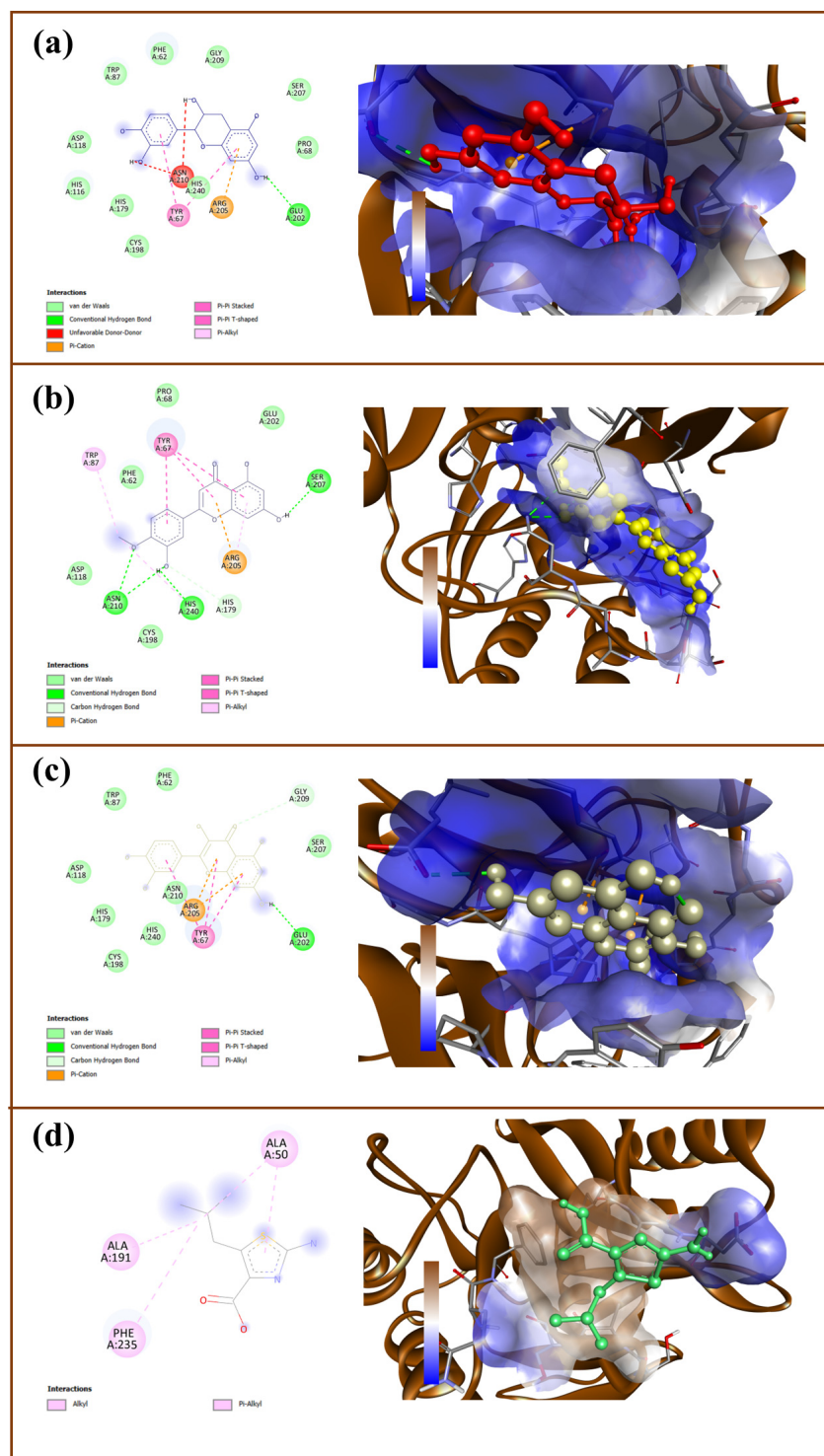


Figure 4. 2D and 3D visualization of the intermolecular interactions between VIM and the top-ranked phytochemicals (catechin (a), diosmetin (b), quercetin (c), and the reference inhibitor (d)).

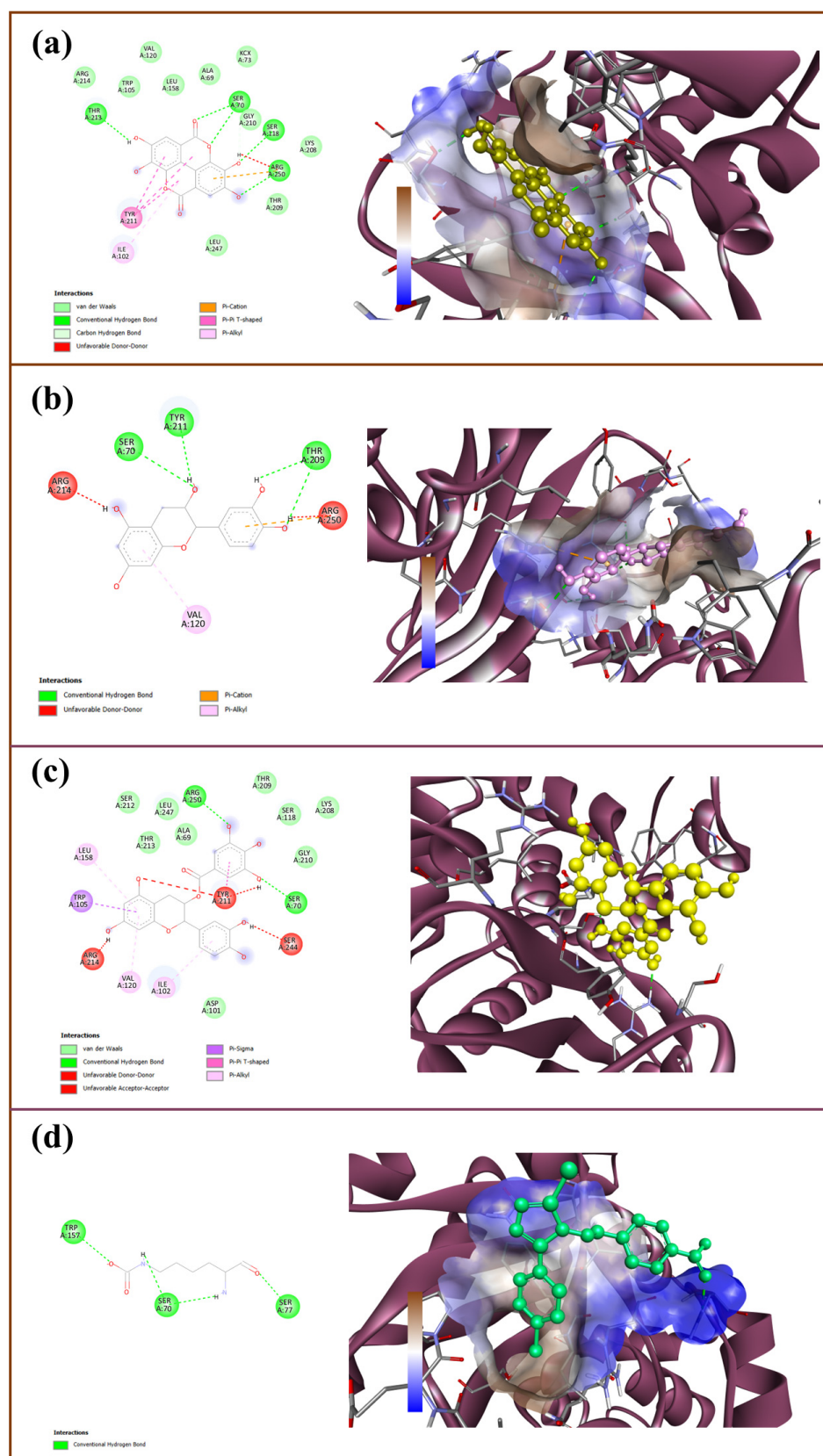


Figure 5. 2D and 3D visualization of the intermolecular interactions between OXA-48-like β -lactamases and the top-ranked phytochemicals (ellagic acid (a), catechin (b), epicatechingallate (c), and the reference inhibitor (d)).

3.5.2. Drug-Likeness Properties Prediction

To identify potential lead compounds with favorable drug-like properties, the top-ranked phytochemicals were evaluated using the SwissADME web server. Drug-likeness was assessed according to Lipinski's Rule of Five, the Ghose filter, and the Veber rule, based on key physicochemical descriptors including molecular weight, lipophilicity (LogP), hydrogen-bond donors, hydrogen-bond acceptors, topological polar surface area (TPSA), and the number of rotatable bonds (Table 8). Overall, all selected compounds complied with Lipinski's Rule of Five and fulfilled the Ghose criteria, indicating favorable drug-like characteristics. Furthermore, all compounds satisfied the Veber rule, except for epicatechin gallate, which exceeded the recommended TPSA threshold of 140 Å². Diosmetin and quercetin exhibited TPSA values below 130 Å², suggesting good membrane permeability.

Table 8. Physicochemical properties and drug-likeness prediction of the top-ranked phytochemicals evaluated using SwissADME.

Compound	MW (g/mol)	logP	Acceptors	H-Bond Donors	Lipinski Violation	Ghose Violation	Veber Violation	TPSA (Å ²)
Epicatechingallate	442.37	2.20	10	7	1	0	1	177.14
Catechin	290.27	1.22	6	5	0	0	0	110.38
Quercetine	302.24	1.99	7	5	0	0	0	131.36
Ellagic acid	302.1	1.31	8	4	0	0	0	141.34
Diosmetin	300.26	2.59	6	3	0	0	0	100.13

The BOILED-Egg model was further used to predict gastrointestinal (GI) absorption, blood–brain barrier (BBB) permeability, and P-glycoprotein (P-gp) substrate status (Figure 6). The analysis indicated that all compounds, except epicatechin gallate, were located within the white region, suggesting a high probability of passive gastrointestinal absorption. Epicatechin gallate was positioned outside the BOILED-Egg predictive regions, indicating poor passive absorption. Regarding P-gp substrate prediction, catechin was classified as a P-gp substrate (blue), whereas the remaining compounds were predicted to be non-substrates (red). None of the evaluated compounds were located in the yellow region of the BOILED-Egg plot, indicating a low probability of blood–brain barrier permeation.

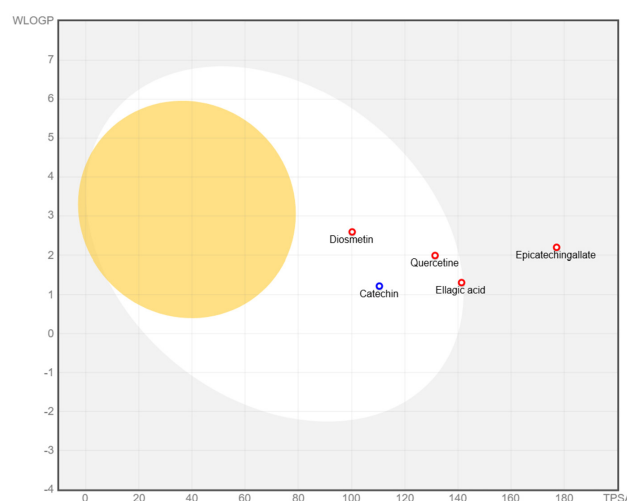


Figure 6. BOILED–Egg plot analysis of the top–ranked phytochemicals.

4. Discussion

The current study explored the potential antibacterial properties of extracts and essential oil from *S. terebinthifolia* leaves in fighting carbapenemase-producing Gram-negative bacteria. Results indicated that Soxhlet extraction was more efficient in extracting phenolic compounds than the maceration method. The difference in total phenolic content determined is linked to the extraction methods. The content of total phenolics detected in the aroeira leaves included in the current investigation was markedly superior to that presented in the literature for the same species of leaf extract: 384.64 mg GAE/g [27]. Soxhlet extraction varied between 5.44 and 309.03 mg GAE/g of Soxhlet extract, while extracts obtained by maceration ranged between 73.90 and 228.51 mg GAE/g [28]. A previous report showed that in the Moroccan *S. terebinthifolia* Raddi, the major compounds are represented by phenolic compounds, principally including galloyl derivatives; flavonoids represented the rest, particularly flavonols, namely myricetin, quercetin, and kaempferol derivatives [29]. Another study showed the presence of catechin, ellagic acid, gallic acid, and epicatechin in *S. terebinthifolia* extract [30]. Sereniki et al. (2016) confirmed that the *S. terebinthifolia* sample demonstrated the existence of the prevailing compounds, namely phenolic acids and flavonoids [31]. It exhibited a phenolic profile characteristic of the *Schinus* species, and phenolic acids and flavonoids were identified [32].

The volatile composition analysis of the leaf essential oil of *S. terebinthifolia* revealed a predominance of monoterpenes, with (+)-4-carene, p-cymene, and trans-2-carene-4-ol identified as the major constituents. Similar monoterpene-rich profiles have been reported for *S. terebinthifolia* essential oils from different geographical regions. For example, in Tunisia, the essential oil of *S. terebinthifolia* was also characterized by a predominance of monoterpenes, particularly α -pinene (14.85–15.18%) and limonene (6.62–8.79%) [33]. However, other studies have reported different chemical profiles. For example, Belhoussaine et al. (2022) found that the leaf essential oil of *S. terebinthifolia* collected in Rabat, Morocco, was dominated by globulol, γ -elemene, and β -elemene [29]. Such variations in the chemical composition of essential oils may be attributed to several factors, including geographical origin, climatic and environmental conditions, soil characteristics, plant genotype, developmental stage, harvesting season, and extraction method.

The in vitro biological investigations conducted on *S. terebinthifolia* extracts displayed their biological capabilities, including their antibacterial properties, for potential use against urinary and respiratory tract infections [34]. Our results demonstrated that *P. aeruginosa* was the most susceptible bacterial species to the *S. terebinthifolia* phytobiotics. This was evidenced by the lowest MIC values, particularly for the Soxhlet leaf extract (STS), which exhibited an MIC of 310 $\mu\text{g/mL}$. In contrast, *K. pneumoniae* was the least susceptible species, displaying the highest MIC values among the tested bacteria (Figure 2). Our findings agree with those previously published that describe the antimicrobial potential of *S. terebinthifolia* leaf phytobiotics against Gram-negative bacteria, including *P. aeruginosa* and *E. coli*, with an MIC value of 500 $\mu\text{g/mL}$ [35]. Additionally, prior research revealed that the leaves of the Anacardiaceae family plant had antibacterial activities against carbapenem-resistant Enterobacteriaceae, including *K. pneumoniae*, with MICs ranging from 512 to 1024 $\mu\text{g/mL}$ [36]. Similarly, antibacterial activities of *S. terebinthifolia* extracts reported that the IZD values were about 15.3 mm and 13 mm at concentrations of 2000 and 1000 $\mu\text{g/mL}$, respectively, against *E. coli*, and 18.3 mm at 2000 $\mu\text{g/mL}$ for *P. aeruginosa* [37]. These studies obtained MIC values close to those found in the present study.

In contrast to our findings, Giordani et al. (2021) suggested that *S. terebinthifolia* phytobiotics were active against Gram-negative bacteria at an MIC value of 3750 $\mu\text{g/mL}$ [38], an MIC value higher than what we found in our research. Another study reported that ST extracts exhibit their effect against *E. coli* with an MIC of 78 $\mu\text{g mL}^{-1}$ [39]. This greater

activity could be attributed to the richness of phenolic constituents because compounds of this class have demonstrated interesting antibacterial activity [40]. The variations in results may be linked to the variation in the chemical profile of each extract.

New antibacterial agents are being investigated to be used alone and in association with drugs to surmount antibiotic resistance. Plant metabolites have been used as a promising alternative because they can act via different mechanisms and have fewer side effects [41]. Natural products can induce an alteration in the effect of antibiotics, either by antagonizing or increasing the antibiotic effect [42]. The study of the application of natural compounds in association with antibiotics could be useful to treat infectious diseases caused by ESBL-producing Enterobacteriaceae [43]. In particular, *S. terebinthifolia* phytobiotics could be used as an excipient for β -lactam antibiotics, decreasing their dose. This alternative seems promising [44]. Based on a more in-depth analysis of activity, it is possible to observe that the combination with the three phytobiotics potentiated the action of cefotaxime against all bacteria included in this study by increasing the sensitivity of the tested strains toward cefotaxime (Table 4). In this context, similar to our findings, numerous studies have highlighted the synergistic effect of natural extracts with β -lactam antibiotics, which is a potential approach to combat β -lactamase-producing isolates. The *S. terebinthifolia* extracts potentiated the action of two conventional β -lactam antibiotics; exploring the association of *S. terebinthifolia* extracts with β -lactam antibiotics such as gentamicin induced a reduction of 50 to 96.87% in the MIC of antibiotics, indicating the capacity to enhance the antibiotic effect against MDR strains, with a significant effect against *E. coli*, where the MIC was decreased from 64 $\mu\text{g/mL}$ to 2 $\mu\text{g/mL}$ [45]. Furthermore, the lectin obtained from the leaf of *S. terebinthifolia* improved the action of ampicillin and presented a synergistic effect against MDR strains [46].

Indeed, it was shown that combining a natural compound with cefotaxime against metallo- β -lactamase-producing isolates presented an excellent synergistic effect [42]. Thus, Zhou et al. (2013) indicated that the plant extract had a synergistic impact on cefotaxime against ESBL-producing isolates, and they demonstrated that it exhibited enzymatic inhibition activity of β -lactamase against these isolates without any mutation in the ESBL gene [43]. However, other results evaluating the effects of the association of different antibiotics with different essential oils showed good activity against MDR strains producing extended-spectrum β -lactamase, such as *K. pneumoniae* and *E. coli* [47]. Additionally, it was confirmed that natural extracts against oxacillin-resistant isolates demonstrated synergistic action with cefotaxime against ESBL-producing *E. coli* strains [44]. In this case, the resistance mechanism involves the production of β -lactamase that stimulates the hydrolysis of the β -lactam ring, inducing its neutralization [48]. Interestingly, botanicals ameliorated the actions of several antibiotics and potentiated the effect of antibiotics against *K. pneumoniae* [37]. Thereby, the combination of an Anacardiaceae plant with imipenem decreased the MIC value of the antibiotic against MDR *P. aeruginosa* [43]. Another study by [45] indicated that some phytocompounds exhibited a synergistic activity when combined with imipenem. However, mechanisms of synergy are still poorly understood; nevertheless, various researchers propose that molecules disrupt the cell wall or increase the permeability of the cytoplasmic membrane and thereby produce efflux pump inhibitors, inhibit penicillin-binding proteins, or promote the influx of antibiotics [45].

Limited studies have been conducted to examine the impact of *terebinthifolia* extracts on inhibiting β -lactams hydrolysis. This is the first study reporting the inhibitory activity of β -lactamase using *S. terebinthifolia* extracts. Costa da Silva et al. (2022) attributed the effect of *S. terebinthifolia* essential oil on strains resistant to a β -lactam antibiotic to the inhibition of the β -lactam ring by the essential oil [46]. On the other hand, Winsou et al. (2022) observed strong β -lactamase inhibition activity with *Terminalia superba* and

Annona senegalensis [49]. Indeed, a hexane extract from *Anacardium occidentale* exhibited β -lactamase inhibitory activity on two strains resistant to penicillin by the synthesis of oxacillinase [50]. Moreover, [45] have reported that there is a relationship between the chemical compounds and the antibacterial effect of the plants [51].

In the present study, the extracts from *S. terebinthifolia* had potent antibacterial activity against carbapenemase-producing strains, targeting the hydrolysis of β -lactam antibiotics by the β -lactamase mechanism. Several natural constituents present in the plant employed in the current research could have contributed to the inhibition of β -lactamase in the included isolates. Furthermore, *in silico* analyses, including molecular docking, confirmed that phytochemicals of *S. terebinthifolia* exhibit a strong binding affinity within the pocket of carbapenemase enzymes, suggesting their effectiveness as potential inhibitors (Figure 7). Additionally, Spyarakis et al. [52] have suggested the β -lactamase inhibitory activity of natural compounds. The presence of the 4-oxo function in flavonoids, comparable to that in clavulanic acid and penicillin G, underlies this property [52]. Several *in silico* findings have highlighted that flavonoids can inhibit NDM-1, which is responsible for inducing resistance to standard antibiotics [53,54]. Moreover, Zhang et al. (2022) identified kaempferol, apigenin, fisetin, taxifolin, quercetin, and luteolin as effective inhibitors of OXA-48 [55].

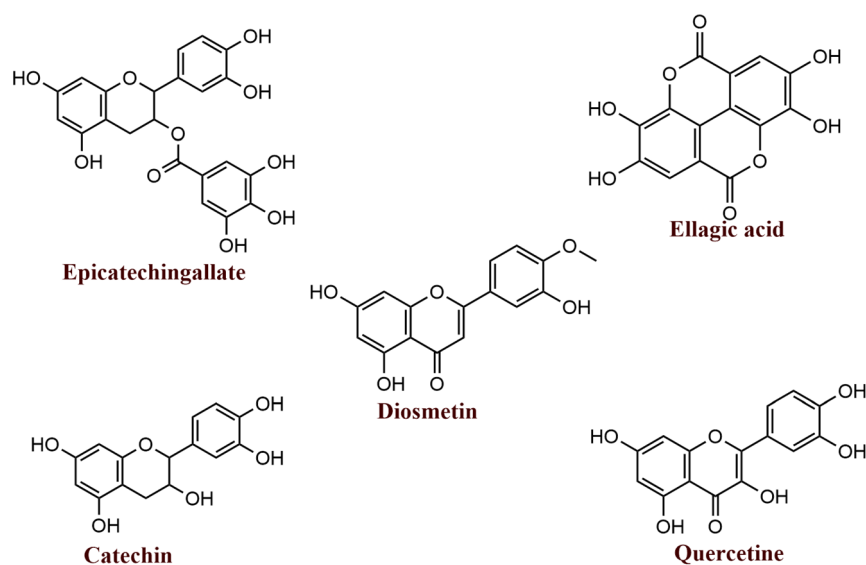


Figure 7. Chemical structures of the most promising compounds identified by virtual screening (ChemDraw v21.0.0).

Although the present study provides evidence for the antibacterial and β -lactamase inhibitory potential of *S. terebinthifolia* leaf extracts and essential oil, the observed antibacterial effects may involve multiple complementary mechanisms. In addition to β -lactamase inhibition, the antimicrobial effects of plant-derived preparations may result from interactions of their diverse constituents with bacterial cellular structures and physiological processes. The observed enhancement of cefotaxime activity, together with the β -lactamase inhibition results, suggests that interference with β -lactam resistance may contribute to the overall antibacterial effect. The molecular docking results further support the potential interaction of selected phytochemicals with β -lactamases.

5. Conclusions

In conclusion, *S. terebinthifolia* leaf extracts and essential oil exhibited antibacterial activity against the tested carbapenem-resistant Gram-negative isolates and enhanced the activity of cefotaxime in combination assays. The phytochemicals also showed β -lactamase

inhibitory activity, supporting their potential contribution to the observed antibiotic-potentiating effect. Molecular docking analyses further identified diosmetin, catechin, quercetin, ellagic acid, and epicatechin gallate as candidate β -lactamase-interacting compounds based on their predicted binding profiles. Overall, the findings support the potential of *S. terebinthifolia* phytobiotics as sources of natural antibiotic-adjuvant candidates. Further biochemical, toxicity, and in vivo studies are required to confirm the underlying mechanisms and assess their therapeutic relevance.

Supplementary Materials: The following supporting information can be downloaded at <https://www.mdpi.com/article/10.3390/app16189212/s1>, Table S1. The MIC, MBC, and MBC/MIC ($\mu\text{g/ml}$) values of Soxhlet Leaf Extract (STS) against *E. coli*, *K. pneumoniae*, and *P. aeruginosa*. Table S2. The MIC, MBC, and MBC/MIC ($\mu\text{g/ml}$) values of Leaf Maceration Extract (STM) against *E. coli*, *K. pneumoniae*, and *P. aeruginosa*. Table S3. The MIC, MBC, and MBC/MIC ($\mu\text{g/ml}$) values of Leaf Essential Oil (STEOL) against *E. coli*, *K. pneumoniae*, and *P. aeruginosa*.

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Abbreviations

The following abbreviations are used in this manuscript:

BBB	blood–brain barrier
BOILED-Egg	Brain Or IntestinaL EstimateD Permeation method
CTX	cefotaxime
<i>E. coli</i> R	carbapenem-resistant <i>E. coli</i>
EO	essential oil
ESKAPE	<i>Enterococcus faecium</i> , <i>Staphylococcus aureus</i> , <i>Klebsiella pneumoniae</i> , <i>Acinetobacter baumannii</i> , <i>Pseudomonas aeruginosa</i> , and <i>Enterobacter species</i>
FICI	fractional inhibitory concentration index
GA	gallic acid
GC/MS	gas chromatography–mass spectrometry
GI	gastrointestinal
H-bond	hydrogen bond
IZD	inhibition zone diameter
KPC R	carbapenem-resistant <i>K. pneumoniae</i>

LC-MS/MS	liquid chromatography–tandem mass spectrometry
MBC	minimum bactericidal concentration
MDR	multidrug resistance
MHA	Mueller–Hinton agar
MIC	minimum inhibitory concentration
MW	molecular weight
OD	optical density
PAC R	carbapenem-resistant <i>P. aeruginosa</i>
PBPs	penicillin-binding proteins
P-gp	P-glycoprotein
<i>S. terebinthifolia</i>	<i>Schinus terebinthifolia</i> Raddi
SMILES	Simplified Molecular Input Line Entry System
STEOL	leaf essential oil of <i>S. terebinthifolia</i>
STM	extract obtained by maceration
STS	extract obtained by Soxhlet extraction
TPC	total phenolic content
TPSA	topological polar surface area
WHO	World Health Organization

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