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Antimicrobial, antioxidant and anti-inflammatory

Pooja G Singh^{1,¶}, Anisha S Jain^{1,¶}, Poojitha B. Sridhara Setty², Sushma BV¹, Sharanagouda S. Patil³, Ashwini P¹, Gopenath TS¹, Kuralayanapalya Puttahonnappa Suresh³, Guru Kumar Dugganaboyana¹, Karthikeyan Murugesan⁴, Ashok Gnanasekaran⁴, Chandan Shivamallu^{1,*}, Shiva Prasad Kollur⁵, Chandrashekar Srinivasa², Raghavendra HL⁶, Parthiban Rudrapathy⁷, Kanthesh M Basalingappa^{1,*}

¹JSS Academy of Higher Education and Research, Mysuru, Karnataka 570 015, India; ²Department of Studies in Biotechnology, Davangere University, Davangere, India; ³ICAR-National Institute of Veterinary Epidemiology and Disease Informatics, Yelahanka, Bengaluru, Karnataka 560 064, India; ⁴Department of Microbiology, Faculty of Medicine, Quest International University, Ipoh, Perak, Malaysia; ⁵Department of Sciences, Amrita School of Arts and Sciences, Amrita Vishwa Vidyapeetham, Mysuru Campus, Mysuru, Karnataka 570 026, India; 6Center for Biotechnology, Pravara Institute of Medical Sciences, Loni-413736, Rahata Taluk, Ahmednagar District, Maharashtra, India; 7Malabar Cancer Centre, Department of Clinical Laboratory Services & Translational Research, Thalassery, Kannur 670103, India; *Corresponding authors, 'Equal contribution & First authorship

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activities of seeds from emblica officinalis (Gaertn.)



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Research Article

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Author contacts:

Pooja G Singh - E-mail: singhpoojasingh824@gmail.com Anisha S Jain - E-mail: anishasjain@jssuni.edu.in Poojitha B. Sridhara Setty - E-mail: bspujitha@gmail.com Sushma BV - E-mail: sushmabv@jssuni.edu.in Sharanagouda S. Patil - E-mail: sharanspin13@gmail.com Ashwini P - E-mail: ashwinip@jssuni.edu.in Gopenath TS - E-mail: gopenath@jssuni.edu.in Kuralayanapalya Puttahonnappa Suresh - E-mail: sureshkp97@rediffmail.com Guru Kumar Dugganaboyana - E-mail: gurukumard@jssuni.edu.in Karthikeyan Murugesan - E-mail: murugesan.karthikeyan@qiup.edu.my Ashok Gnanasekaran - E-mail: gnanasekaran.ashok@qiup.edu.my Chandan Shivamallu - E-mail: chandans@jssuni.edu.in Shiva Prasad Kollur - E-mail: shivachemist@gmail.com Chandrashekar Srinivasa - E-mail: chandru.s@davangereuniversity.ac.in Raghavendra HL - E-mail: raghavendra@pmtpims.org Parthiban Rudrapathy - E-mail: parthi71975@gmail.com Kanthesh M Basalingappa - E-mail: kantheshmb@jssuni.edu.in

Abstract:

There is a shred of evidence to suggest that *Emblica officinalis* Gaertn, the botanical name for amla seeds, has greater medicinal potential than amla fruit. We conducted this work to assess the anti-inflammatory, antibacterial, and antioxidant capacities of *E. officinalis* seed extracts. The bioactive components from the seeds were fractionated using chloroform, hexane, methanol, and diethyl ether, according to the polarity of the solvents. The total amount of phenolic and flavonoid was estimated. Both the reducing power and antioxidant capacities of the extracts were evaluated using the DPPH (1,1-diphenyl-2-picryl-hydrazyl) technique. 15-lipoxygenase (LOX) was inhibited by seed extracts at doses ranging from 5 to 25 micrograms. *In silico* docking was employed to assess the results. Some human pathogenic microorganisms were tested for their antibacterial activity using the agar disc diffusion method. *Escherichia coli, Proteus vulgaris*, and *Klebsiella pneumonia* were inhibited by a methanolic extract with an IC50 value of 58g, making it the most common organic solvent extract. Methanolic extracts also showed good antioxidant and antibacterial activity. Our investigation led us to discover that amla seeds have anti-inflammatory, antioxidant, and antibacterial effects.

Keywords: Emblica officinalis; antioxidant; lipoxygenase inhibition; antimicrobial; LC-MS analysis

Background:

Most of the human diet is comprised of herbs and spices. Their nutritional, medicinal and preservative properties are wellknown in addition to their flavor [1]. Emblica officinalis' antimicrobial properties and lack of side effects have made it more popular in recent years, despite its long recovery time [2]. Medicinal plant E. officinalis Gaertn is widely used in Ayurvedic and Unani medicine systems [3]. The sour amla fruit is often soaked in saltwater and turmeric to make it more palatable in India or, in other words, pickled [4]. The amla fruit's chemical composition contains more than 80% water [5]. Protein, carbohydrate, fibre, and mineral content are also present, as well as gallic acid, which is a highly effective polyphenol [6]. When it comes to the physical and chemical properties of the fixed oil yield (16%), Iodine 139.5, unsaponifiable matter 381.8% and saturated fatty acids 72% are found in the linolenic acid (8.78%), the linoleic acid (44 %), and the oleic acid (28.10%) (0.95%) [7-8]. Amla is the most concentrated source of Vitamin C in the plant kingdom. It is easily absorbed by the human body when the entire fruit is used instead of an active ingredient. The tannins in the amla fruit bind to the vitamin C present in the fruit showing anti-diarrheal and antipyretic properties. E.officinalis contains several chemical constituents. Microbe-fighting constituents include quercetin and flavonoids (phyllantine, phyllantidine),

gallic acid, ascorbic acid and hydrolyzable tannins (emblicanin A and B). Therefore, it is of interest to evaluate the antioxidant, antiinflammatory and antibacterial activity of different solvent extracts of *E.officinalis* seeds.



Figure 1: (a) Three-dimensional structure of human 15-LOX (PDB ID: 4NRE) retrieved from the protein data bank. (b) Active pocket (red-colored) of 4NRE protein estimated from CASTp server surrounding the binding residues.

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Materials and Methods:

Reagents and Chemicals

Catechin, 1,1-diphenyl-2-picryl-hydrazyl (DPPH) and linoleic acid were procured from Sigma–Aldrich. Folin-Ciocalteu reagent, gallic acid, ascorbic acid, trichloroacetic acid, ferric chloride, aluminum trichloride, sodium carbonate and sodium hydroxide and other reagents and chemicals used were of AR grade.



Figure 2: The total phenol content was calculated using the standard gallic acid graph, and the findings were represented in GAE.



Figure 3: Total phenol content was calculated using a standard catechin graph, and the results were reported as catechin equivalents.



Figure 4: The DPPH scavenging capacity was calculated using the standard ascorbic acid graph, and the results were represented as IC50 values.



Figure 5: The DPPH scavenging capacity was calculated using the ascorbic acid standard graph, and the results were presented as IC50 values for hexane extract.



Figure 6: The DPPH scavenging capacity was calculated using the ascorbic acid standard graph, and the results were represented as IC50 values for diethyl ether extract.



Figure 7: The reducing power capacity of ascorbic acid was calculated using a standard graph, and the findings were reported as IC50 values for methanolic extract.

 Table 1: Total weight of E. officinalis seeds extracts and qualitative analysis of Tannins and Saponins

Parameters	Hexane	Chloroform	Diethyl Ether	Methanol
Extracts (g)	0.362	0.093	0.0512	0.4416
Tannins	-	-	-	+
Saponins	+	+	+	-

(+) present; (-) absent

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Figure 8: Methanolic extract of *Emblica officinalis* inhibits 15- LOX with an IC50 of 58 μ g.



Figure 9: A column bar graph determining the negative binding affinity (Kcal/mol) of each bioactive compound of *E.officinalis* with the active site of LOX protein.



Figure 10: An illustration of quercetin compound buried within the binding pocket of hydrophobicity surface of the target protein. The docked complex's interactions are highlighted in the 2D picture.

Plant Extract:

Seeds and rind:

Fresh fruits of amla (*Emblica officinalis*) were chosen from the local supermarket in Karnataka State, India. Seeds and rinds were segregated from the fruit and shade dried at 37°C for 3 days. The dried samples were coarsely ground using 500 ml of liquid nitrogen in a mortar and pestle. As soon as the amla seeds and rinds were ground to a coarse paste, they were stored in sealed glass bottles at 4°C until they were used **[9, 10]**.

Extraction of seeds and rinds:

On the basis of increasing polarity, hexane, chloroform, diethyl ether, and methanol solvents were used to dissolve 25 g of powdered amla seeds in 200 ml of solvent. The extraction was done for 48 hours at room temperature with a rotator shaker set to 30° C and 60 rpm. Filtration of the solvent via muslin cloth was used to separate it. The samples were then vacuum filtered and dried in glass petri plates at room temperature using a flash evaporator set to 44° C and 160 rpm. The final drying procedure was carried out for 3 hours at 40° C using a Speed Vac Concentrator (Savant SPD 2010), and the extracted material was kept in the dark at 4° C until further usage **[11]**.

Quantitative Analysis of Phytochemical Constituents: Determination of total phenolics

The Folin-Ciocalteu colorimetric technique was used to calculate the total phenolic content. The test sample (100 μ l) was incubated at 22°C for 5 minutes with 250 μ l of Folin-Ciocalteu reagent. The reaction was neutralized with 1500 μ l of saturated sodium carbonate and left to sit in the dark for 1.5 hours at 22° C. A Hitachi U-3900 UV/visible spectrophotometer were used to test the blue color's absorption at 765 nm. A calibration curve based on the absorbance of known amounts of Gallic acid standard (20 to 100 μ g) was used to calculate total phenolics. Gallic acid equivalence (GAE) in μ g was used to calculate the total phenolic content **[12]**.



Figure 11: A pictorial representation of gallocatechin compound engraved to the active pocket of hydrophobicity surface of the target protein. The 2D image highlights the interactions of the docked complex.

Table 3: Antimicrobial activity of E.officinalis methanolic extract against E. coli								
Solvent used	Test organism	Concentration	Zone of inhibition (mm)					
			T0	T12	T24	T48		
		Control	ND	0.15	0.32	0.37		
Methanol	E. coli	Drug	ND	0.74	4.0	4.32		
		1.00	ND	1.17	0.23	0.26		
		2.50	ND	0.37	0.40	0.47		
		5.00	ND	0.88	0.89	1.36		
		10.00	ND	1.4	1.96	2.55		

 Table 4: Antimicrobial activity of *E.officinalis* methanolic extract against *P. vulgaris*

 Solvent used
 Test organism
 Concentration
 Zone of Inhibition (mm)

rest organism	concentration	Lone of multitudi (mill)			
		T0	T12	T24	T48
	Control	ND	0.44	0.47	0.53
P. vulgaris	Drug	ND	0.76	1.03	1.34
	1	ND	0.19	0.28	0.33
	2.5	ND	0	0.28	0.45
	5	ND	0	0	2.73
	10	ND	0	0.66	1.17
	0	Control P. vulgaris Drug 1 2.5 5	T0ControlNDP. vulgarisDrugND1ND2.5ND5ND	T0 T12 Control ND 0.44 P. vulgaris Drug ND 0.76 1 ND 0.19 2.5 ND 0 5 ND 0	T0 T12 T24 Control ND 0.44 0.47 P. vulgaris Drug ND 0.76 1.03 1 ND 0.19 0.28 2.5 ND 0 0.28 5 ND 0 0



Figure 12: Inhibition of methanolic extracts against Escherichia coli at concentrations of 1 ug (a); 2.5 ug (b); 5 ug (c); and 10 ug (d) in comparison to the control (methanol; e) and the antibiotic, streptomycin (f).



Figure 13: Inhibition of methanolic extracts against Proteus vulgaris at concentrations of 1 ug (a); 2.5 ug (b); 5 ug (c); and 10 ug (d) in comparison to the control (methanol; e) and the antibiotic, streptomycin (f)

Estimation of total flavonoids:

The total flavonoid content was estimated using aluminum trichloride by the colorimetric method. Catechin (1-5 μ g) was utilized as a control. Extract (10-50 μ l) was made up to 1 ml using methanol, 4 ml distilled water, and 0.3 ml of 5% sodium nitrite solution. After 5 minutes of incubation, 0.3 ml of 10% aluminum trichloride solution was added and the combination was allowed to stand for 6 minutes. The combination was then raised to a final volume of 20ml with double-distilled water after the addition of 2ml of 1mol/L NaOH solution. The absorbance was measured at 510 nm after the mixture had been allowed to sit for 15 minutes. The total flavonoids content was determined using a calibration curve observed from measuring the absorbance of known concentrations of catechin as standard (10 to 50 μ g). The total flavonoid content was expressed as catechin equivalence (CE) in μ g **[13]**.

Test for tannins:

Around 0.1 g of amla seed extract was weighed and suspended in respective solvents. In a test tube, diluted extract (0.5 mL) was mixed with 10 mL distilled water and filtered using Whatman

No.1 filter paper. Three drops of ferric chloride reagent were applied to the filtrate. The existence of gallic or catechol tannins was established by the emergence of blue-black or green precipitation [14].

Test for Saponins:

01. g of amla seed extract was weighed and diluted in respective solvents. In test tubes, different solvent extracts (1 mL) were added to 5 mL of distilled water. The presence of saponins was shown by the formation of stable foam.

Estimation of antioxidant activity:

DPPH radical scavenging activity

An amla seed extract in hexane, diethyl ether, and methanol was tested for its ability to neutralize free radicals using the DPPH method. A solution of DPPH radicals was made in methanol **[15]**. This mixture comprises 5 μ l of test samples and 95 μ l of 50 μ g/ml DPPH. After 30 minutes at 37°C in the dark, a 96-well plate was used to perform the DPPH radical scavenging reaction. The radical scavenging activity was determined as a percentage using a solvent as a control **[16]**. The IC50 values were calculated, which show the concentration of extracts required to scavenge 50% of DPPH radicals. The standard was ascorbic acid (1–10 μ g). The following equation was used to calculate the percent scavenging effect:

Eq. (A.1).Inhibition (%) = [(Absorbance of control –Absorbance of the test sample)] / Absorbance of control × 100

Reducing power estimation

The reducing power assay of amla seed extract was determined by ferric chloride (FeCl3). Phosphate buffer (0.2 M, pH 6.6) and potassium ferricyanide were combined in equal parts with the test sample (0.1 g) (2.5 ml, 1%). It was incubated for 20 minutes at 50°C. The mixture was then centrifuged at 3000 rpm for 5 minutes with trichloroacetic acid (TCA) (10 percent, 2.5 ml). The supernatant (1 ml) was collected after centrifugation and combined with an equal volume of distilled water and FeCl3 in a test tube (0.5 ml, 0.1%). As a blank solution, phosphate buffer was utilized, and ascorbic acid was employed as a standard. Beckman-DU730 Coulter's life sciences UV/Visible spectrophotometer was used to detect absorbance at 700 nm. Stronger reducing power is shown by increased absorbance of the reaction mixture.

Anti-inflammatory Activity:

In vitro activity

The anti-inflammatory property determined was spectrophotometrically [17] with a slight modification in the procedure. Lipoxygenase (LOX) enzyme activity was assayed in a spectrophotometer. The 15-lipoxygenase (15-LOX) from soybeans was used in the test. Experimentally, the loss of soybean 15-LOX activity (5 g) was measured by using 0.2M linoleic acid (Sigma) as the substrate prepared in the solubilized state (dissolving in 10 µl methanol) in the presence of 0.2M borate buffer (pH 9.0) [18]. With a UV-Vis spectrophotometer, various extract concentrations (5, 10, 15, 20 and 25 µg) and a reference compound (ascorbic acid) was used to measure inhibition. We also calculated the percentage of enzyme activity that the extracts inhibited. IC50, or the concentration needed to inhibit 50% of LOX activity, was

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estimated as well **[19]**. The following equation was used to calculate hydrogen peroxide content and lipoxygenase activity:

Eq. (A.2). Specific activity (LOX)= Δ A.V/ ϵ .l.c

Where ΔA is the absorbance rise per minute, V is the incubation mixture volume, ϵ is the extinction coefficient for linoleic acid (25 x 10-3 mol/l/cm) **[20]**, l is the length of the cuvette (1 cm) and c is

the enzyme concentration in mg (0.005). The results represent the average of three separate tests.



Figure 14: UPLC-PDA-ESI/HDMS was used to assess the phytochemical profile in the methanol extract of Emblica officinalis seed.

Molecular docking:

A molecular docking approach is used to determine the interactions and binding affinity between three-dimensional structures of a ligand and protein. To carry out this, PyRx 0.8 virtual screening software was used. The 3D structure of the target protein, soybean 15-lipoxygenase (15-LOX) was retrieved from Protein Data Bank (PDB) having a resolution of 2.63 Å [21]. To prepare the target protein for docking, water molecules and bound ligands were detached from the structure, missing residues and polar hydrogens were added to it (Fig. 1). Among several phytocompounds present in the seed extract of E.officinalis [22, 23] we selected a few compounds, whose 3D structures were downloaded from the PubChem database. These structures were prepared for docking by converting the downloaded sdf files into pdb files using Open Babel GUI. The binding site of the target protein was estimated by CASTp online server. Along with the binding residues, the area and volume of the binding pocket are also calculated by this server. Finally, the protein and ligand files are loaded to the PyRx server where the docking analysis is carried out [24].

Antibacterial Activity:

Antibacterial activity of seed extracts was tested against gramnegative bacteria *Klebsiella pneumonia* and *Escherichia coli* and grampositive bacteria *Pseudomonas vulgaris* which were collected from the Department of Microbiology, JSS Academy of Higher Education and Research, Mysuru, India. All of these cultures were kept at 4°C on nutrient agar plates.

Agar disc diffusion:

To test for antibacterial activity, methanolic seed extract of *E.officinalis* was tested against a bacterial culture. Muller Hinton agar was inoculated evenly with 0.02 ml of a 24hr old culture using a sterile cotton swab. This created a lawn culture on the MHA. A seeded agar plate was used to test the antibacterial activity of methanolic extract of *E.officinalis*. The discs containing the extract were placed on the seeded agar plate **[25, 26]**. Discs carrying methanol solvent were used as controls. Streptomycin (10 μ g) antibiotic was used as a drug. Varying concentrations of methanol extract (1, 2.5, 5, 10 μ g) were used. The plates were incubated at room temperature for 48 hours before being examined for a zone of growth inhibition around the discs (Stereo Discovery V20 – Zeiss, Germany, Whitefield Microscope). The antibacterial activity of the extracts was determined by measuring the diameter of the zone of inhibition to the nearest millimeter for each bacterial species.

LC-MS analysis:

According to the manufacturer's protocol, the metabolites were analyzed qualitatively using Synapt G2 (UPLC separations with Quan Tof). The nitric oxide flow rate was set at 10 L/min at 350°C with a 60-psi nebulizer pressure. Positive ionization was used to acquire mass spectra from m/z 100 to 1000. For fragmentation of the isolated compounds, helium was used as a collision gas **[27]**. The detection conditions were as follows: skimmer voltage, -40 V; capillary voltage, 3500 V; trap drive level, 45.0; Oct RF, 150 Vpp; Lens 1, 5.0 V; Lens 2, 60 V; cap exit voltage, -158.5 V; Oct 1 DC, -12 V; Oct 2 DC, -2.45 V.

Table 2: List of phytocompounds selected for the docking study along with their structures and binding energy values estimated by PyRx software.

Phytocompounds Binding energy (Kcal/mol) Three-dimensional structure Caftaric acid -76 Cis-Resveratol -7.7 Gallic acid -6 Gallocatechin -9.1 Linoleic acid -59 Linolenic acid -61 Myristic acid -5.3 Naringenin -8.7 Oleic acid -5.7 Pyrogallol -5.7 Quercetin -9.7

Results:

Phytochemical constituents and active ingredients:

The physical properties of pH and moisture content of amla were found to be slightly acidic and 65% respectively. The Phytochemicals analysis of amla seed extract is summarized in **Table 1**. Amla seeds contain a lot of nutritive and anti-nutritive

compounds. Saponins, tannins, flavonoids and phenolic content were `found to be higher in methanolic extract than other solvent extraction. Smaller amounts of the biologically active compound were extracted by hexane, diethyl ether and chloroform. The amount of total phenol was observed as 120 μ g (**Figure 2**) of seeds as gallic acid equivalents and the amount of total flavonoids content was observed as 100 μ g (**Figure 3**) as catechin equivalents of amla seeds.

DPPH free radical scavenging activity and reducing power assay

The antioxidant activities of *E.officinalis* seed extracts were confirmed in our work using the DPPH and Reducing Power Assays. The hydrogen donating ability of test compounds is ascribed to the percentage of DPPH decolonization. The amla seed extract showed a wide range of DPPH activity. When compared to other extracts, Methanolic extract had higher antioxidant activity (IC5024 μ g), whereas Hexane (**Figure 5**) and Diethyl ether (**Figure 6**) had the lowest DPPH radical scavenging activity. At 42 μ g, the reference standard ascorbic acid demonstrated 50% inhibition.

Reducing power estimation

The IC50 of the methanolic extract is 40μ g in the reducing Power assay (**Figure 7**). The methanol extracts of amla seed had good activity, whereas hexane and diethyl ether exhibited modest activity. Ascorbic acid, the reference standard, demonstrated a 50% inhibition. Lower IC50 value implies higher antioxidant power. The presence of reductions capable of breaking free radical chains by donating hydrogen atoms and therefore converting them to a more stable non-reactive species is indicated by a higher reducing power.

Lipoxygenase inhibition

Hydroperoxylinoleic acid was detected as a rise in absorbance at 234 nm, indicating LOX activity. Concentration-dependent LOX inhibition was observed with *E.officinalis* seed extract. The methanolic extract had the strongest inhibitory effect, with an IC50 of 58 μ g. At a dose of 100 μ g, complete LOX inhibition was found (**Figure 8**).

Molecular docking analysis

The docking analysis was carried out between the target protein of 15-lipoxygenase and bioactive compounds of Emblica officinalis. Table 2 and Figure 9 represent the estimated binding affinity of the best-docked poses obtained from PyRx. An acceptable binding affinity between protein and ligand is indicated by low binding energy. Based on the graph, we can say that quercetin has shown the best binding affinity (-9.7 Kcal/mol) with the active site of the target protein followed by gallocatechin (-9.1 Kcal/mol). These docked complexes were further evaluated using visualization software. The docked complexes of 15-LOX with 2 two best compounds that are, quercetin and gallocatechin were further evaluated using Chimera 1.15 and the 2D images were procured from BIOVIA Discovery Studio Visualizer 2020. Both the ligands showed a high number of non-bonded interactions with the binding site residues of the target protein. In Figure 10, we can see that quercetin shared five hydrogen bonds with TYR A: 107, GLN A: 108, ARG A: 145, HIS A: 394, PHE A: 399 (bright green colored

discs and bonds), a pi-sigma bond with ILE A: 403 (purple colored disc and bond) and fourteen van der Waals interactions (light green discs). An unfavorable bond was also observed with THR A: 406 (red-colored disc). For the docked complex of gallocatechin (**Figure 11**), there were four hydrogen bonds with HIS A: 394, ARG A: 145, TYR A: 149, ARG A: 407, a single pi-sigma bond with ILE A: 403, an unfavorable bond with GLN A:108 and eleven van der Waals interactions.

Antibacterial activity:

As an antipyretic, analgesic and immunomodulatory agent, *E.officinalis* Gaertn has a long history of use. First, the antibacterial property of the extracts was determined using the standard agar disc diffusion method, as previously reported **[28]**. Extracted seed extracts were compared to streptomycin, a commonly used antibacterial drug, for their antibacterial efficacy *in vitro*. The methanolic solvent extract of the seed showed antibacterial property against the clinical cultures of *E. coli*, *P. vulgaris* and *K. pneumonia* and the results were recorded at different time intervals of 12h, 24 h and 48h and the results for *E. coli* is presented in **Figure 12**. The results for *P. vulgaris* are presented in **Figure 13**. Maximum antibacterial activity was obtained for *E. coli* (4.32 mm); (**Table 3**) and minimum response was recorded against *P. vulgaris* for methanolic extracts (**Table 4**). No activity was recorded for *K. pneumonia*.

LC-MS estimation:

The methanol seed extract of *E.officinalis* yielded 6 compounds. The mass spectral analysis was used to make a tentative identification of these compounds, which was then confirmed by comparing them to literature values. Complicated metabolite identification using chromatographic and internal MS data identified compound A as MS1-Digalloyl Hexose and Diaglloyl Glucose isomers. Compound B was identified as Patuletin glucoside and Patuletin Gallacatechin. The compound C is identified as Methyl 4,7,10,13,16 and 19-doco sahenaenoate (C23H34O2), Compound D was identified as MS3 cisresveratrol, caftaric acid and Quercitrin and Compound E were identified as MS4- (±) - naringenin by comparing retention time and MS/MS data (**Figure 14**).

Discussion:

A sophisticated traditional medicine system is based on plants, and natural products are excellent starting points for novel drug development programs **[29]**. Health programs in developing countries can benefit from the effective use of herbal medicine that the World Health Organization (WHO) encourages, promotes, and facilitates **[30]**. Studies have been conducted to determine the potential of natural products in inhibiting ROS and, as a result, attenuating health situations requiring ROS inhibition. It has been three decades since new antibiotics were created, yet bacteria have become more resistant to them **[31]**. Drug resistance is a common trait among bacteria. In the present study, a freshly prepared DPPH solution with maximum absorption at 517 nm is used in the DPPH method. Most of the time, if there is an anti-oxidant present in the medium, the purple color fade **[32]**. Antioxidant molecules provide hydrogen or electrons to DPPH free radicals, perhaps via a free

radical attack on the DPPH molecule resulting in an increase in absorbance at 517nm. As a result, the extract's antioxidant potency increases as the absorbance decreases. Indicators of potential antioxidant activity include a compound's reducing capacity [33-34]. Most of the time, the presence of reductions, which break the free radical chain by giving away a hydrogen atom, is linked to the reducing ability. The extract showed a reductive ability that grew when the concentration of the extract was raised [23]. An antimicrobial activity screening of plant extracts revealed that higher plants are a potential source of new anti-infectives. Although conventional antibiotics inhibit bacteria through different mechanisms, plant-derived antimicrobial compounds may have clinical value in treating infections caused by resistant microbes. Gram (-) bacteria, such as E. coli, and Gram (+) bacteria, such as P. vulgaris, were the most susceptible to methanol extract Differential susceptibility of Gram-positive or Gram-negative bacteria to phenolics was previously reported [35]. A major role is played by inhibiting the production of leukotrienes (LTs) and PGE2. Inflammation-inducing enzymes are inhibited in the initial screening of plants used for hydroxy peroxy linoleic acid is formed when the LOX activity increases. They provide valuable preliminary data for selecting seed extract with potential antiinflammatory activities that support their traditional use. Methanolic seed extract with potent LOX inhibition capacity [36]. Antioxidant activity has been related to plant phytochemicals, which humans cannot synthesize [13]. In the current study, UPLC-DAD identified cis-resveratrol, gallo catechin, quercetin among others. Gallocatechin and quercetin have also proved to be effective enough against the LOX protein via in silico analysis. According to Indian traditional literature, the presence of phytochemicals in the extract of amla seeds may also contribute to their medicinal qualities.

Conclusion:

With this study, we wanted to see how seed extracts of E. officinalis Gaertn worked as anti-oxidants, anti-inflammatory agents, and anti-microbial agents in vitro Hexane, chloroform, diethyl ether, and methanol solvents were used to separate the bioactive components from the seeds in increasing order of polarity. They were tested for antioxidant properties using the DPPH (1,1-diphenyl-2picrylhydrazyl) method, and for reducing power using the FeCl3 method, as well. Seed extracts were used to inhibit 15-lipoxygenase (LOX) at concentrations ranging from 5 to 25µl. A good antioxidant and antibacterial activity were observed in the methanol extracts, which were the most common organic solvent extract inhibition of Lox by methanolic extract with an IC50 value of 58µg have been reported. Total antioxidant capacities were found to be positively correlated with phenolic content, indicating that the phenolics in amla seeds were the dominant antioxidant constituents. We can confirm that amla seeds contain anti-oxidant, anti-inflammatory, and antibacterial activities as a result of our research.

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Conflict of interest:

The authors declare no conflict of interest.

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