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The promising guide to LC–MS analysis and cholinesterase activity of *Luffa cylindrica* (L.) fruit using in vitro and in-silico analyses

S. Ankul Singh¹ and Chitra Vellapandian^{1*}

Abstract

Background *Luffa cylindrica* (L) is an annual climbing plant that produces fibrous fruit and can also be used as a vegetable in northern parts of India. Various studies have been carried out on the plant and found to have anti-inflammatory, antifungal, analgesic, anti-myocardial, anti-hyper triglyceride, immunostimulant, anti-allergic, and other properties. The ethanolic extract of the *Luffa cylindrica* (L) fruit has not yet been subjected to LC–MS analysis for several bioactive chemicals that target neurological diseases. Oxidative stress is an inevitable situation in AD mechanisms and is a key bridge connecting various AD pathways.

Results *Luffa cylindrica* contains various phytochemicals and showed highest alkaloid content of 21.39 ± 1.47 mg of AE/g. A total of 80 compounds were identified in the ethanolic extract from LC–MS analysis. The bioactive compounds were screened for eligibility by Lipinski's rule of five for docking with receptors responsible for causing oxidative stress-associated Alzheimer's disease. Perlolyrine was chosen to perform in-silico docking. An in vitro activity of cholinesterase showed highest inhibition at 500 µg/ml. In-silico docking of perlolyrine showed better binding affinity and score. Results revealed that out of 10 docked receptors, amyloid beta showed the highest binding affinity with an energy of - 46.1 kcal/mol showing promising drug for Alzheimer's disease.

Conclusion Based on current findings, the study reports the presence of a promising, bioactive compound (perlolyrine) and in turn provides an optimistic note in exploring its biological activity in vivo with oxidative stress-related Alzheimer's disease.

Keywords Luffa cylindrica, LC-MS, Alzheimer's disease, In silico, Amyloid beta

Background

Luffa cylindrica (L.) is an annual climbing plant that produces fibrous fruit and can also be used as a vegetable in northern parts of India. It belongs to the Cucurbitaceae family and is also called as loofa, bath sponge, sponge gourd, etc., and contains smooth and cylindrical-shaped fruits. The medicinal plant *Luffa cylindrica* has

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been used extensively to cure various ailments. Family Cucurbitaceae is economically beneficial as it is a significant food source and has pharmacological properties that include anticancer, antiulcer, anti-diabetic, analgesic, and nephroprotective [1]. In Bangladesh, *Luffa cylindrica* (L.) and *Luffa acutangula* are widely used as vegetables, meals, and folkloric medicines to treat various illnesses [2]. The plant is traditionally used to treat a variety of conditions, including asthma, intestinal worms, sinusitis, chronic bronchitis pain, carbuncles, abscesses, inflammation, heat rashes, bowel or bladder hemorrhage, hemorrhoids, jaundice, menorrhagia, hematuria, leprosy, splenopathy, and as an antiseptic, emmenagogue, and carminative [3, 4]. *Luffa cylindrica* (L.) fruit has been



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used extensively to clear phlegm along with antipyretic effect. Its fruit is also used to treat hyperglycemia [5, 6], cataract [7], anti-emetic, anti-inflammatory [8], and antimicrobial effect. [9] Its leaves are also studied extensively on cancer [10], hepatoprotective [11], and anthelminthic [12] activities. Similarly, its seed also has the potential in treating asthma-related effects [13], anti-fungal [14], abortifacient [15], and anti-HIV activity [16]. Various studies have been carried out on the plant and found to have anti-inflammatory, antifungal, analgesic, anti-myocardial, anti-hyper triglyceride, immunostimulant, antiallergic, and other properties [2, 17]. The components that make up plant antioxidants include terpenoids, polyphenolic chemicals, ascorbic acid, and tocopherols, which serve a variety of essential roles in both plants and humans [18]. Oxidative damage contributes significantly to the aging process and the etiology of various illnesses, including atherosclerosis, diabetes, cataracts, bronchial asthma, Alzheimer's disease, cancer, and rheumatoid arthritis. Chemical methods such as radical scavenging assays (DPPH, ABTS, hydroxyl assays), lipid peroxidation assays (carotene-linoleate model systems, thiobarbituric acid-reactive substances), and reduction power assays (FRAP, CUPRAC), as well as enzyme-based assays, can be used to investigate the antioxidant potential of plants [19]. These antioxidants can safeguard plant cells from reactive oxygen species damage, which frequently causes biochemical and physiological lesions, metabolic dysfunction, and ultimately cell death [20]. The endothelin receptor (ER) has a well-established significance in metabolic and cardiovascular problems because of its role and function in Ca²⁺ handling, protein synthesis/ folding, and secretory pathway control. Because ER redox state strongly affects protein folding, when disulfide bond production is regulated in response to ER stress, luminal oxidative stress rises, and ER function falls [21]. Chronic oxidative stress conditions speed up the neurodegenerative process by increasing the calcium influx into the cells caused by reactive oxygen species and altering the mitochondria by lowering succinate dehydrogenase (SDH) activity [22]. Additionally, it can disrupt ATP receptors by augmenting phosphorylation pathways and decreasing membrane depolarization. In conclusion, ozone exposure leads to oxidative stress, which stimulates the purinergic receptor and GSK-3 response, activating the nuclear factor kappa-light-chain-enhancer of activated B cells (NFkB) to produce the pro-inflammatory IL-1 and lowering IL-10, both of which give rise to cell death [23]. Luffa plants are a significant source of anti-retroviral medications since several species have ribosome-inactivating proteins (MAP30, luffin A & B) and anti-HIV activity [24]. In a study, ethanol extract proved the presence of various phytochemicals and antioxidant activity. It also demonstrated the beneficial effects of plant-based therapy on free radical scavenging effects [25].

The most common cause of dementia and cognitive decline in older people (over age 65) is Alzheimer's disease (AD) worldwide. Pathophysiological alterations include the build-up of toxic amyloid beta plaques, neurofibrillary tangles of hyperphosphorylated tau protein, and neurodegeneration carried by unregulated microglial activation in the brain, which secretes neurotoxins and inflammatory cytokines [26]. Populationbased European studies show that the age-standardized prevalence of dementia and AD is 6.4 percent and 4.4 percent among older adults of age 65. Vascular risk factors in middle-aged and older persons strongly contribute to the start and development of dementia and AD [27]. The ethanolic extract of the Luffa cylindrica (L.) fruit has not yet been subjected to LC-MS analysis for several bioactive chemicals that target neurological diseases. Cholinesterase inhibitors have been the best track record of effectiveness for several AD features. Newer potential therapeutic agents are still in the early stages of clinical research, and alternative therapy modalities have not undergone as much testing or demonstrated as much efficacy [28]. Traditional cholinesterase inhibitors are found to have cholinergic actions as well as activity against several AD molecular targets in several in vitro and in vivo investigations. However, their effects in slowing AD progression are still minimal [29]. Oxidative stress is an inevitable situation in AD mechanisms and is a key bridge connecting various AD pathways. Even though the majority of antioxidants have potent benefits in animal research, the outcomes of trials in people are insufficient [30]. For this instance, various oxidative stress receptors associated with Alzheimer's disease can be docked with identified compounds and understand the physiology to prevent the disease.

This work aims to provide a comprehensive view of the phytochemical screening of *L. cylindrica* fruit, identify various bioactive compounds from LC–MS analysis, in vitro cholinesterase activity, and perform in-silico docking of different receptors targeting Alzheimer's disease-associated oxidative stress.

Methods

Collection and identification of plant material

The entire fruit of *Luffa cylindrica* (L.) was collected in and around Uttar Pradesh from June to August 2021. The plant specimen, preserved in the Department of Pharmacognosy, was authenticated by Dr. P. Jayaraman, Director of the Plant Anatomy Research Center in Tambaram, Chennai (Registration No. PARC/2021/4528).

Processing and preservation of plant material

Luffa cylindrica (L.) unripe fruit was washed well in running tap water and then rinsed in distilled water. The unripe fruits were cut into small pieces and dried in the shade for two weeks to achieve total dryness. A mechanical grinder was used to ground the dried unripe fruits to a fine powder. For later usage, the powdered substance was stored in airtight containers.

Reagents and chemicals

All the organic solvents and acids used for these experiments were of analytical or HPLC grade and were procured from SISCO laboratories.

Preparation of ethanolic extract (Soxhlet extraction method)

Chemical constituent screening in medicinal plants is critical for obtaining helpful information for further research and in determining pharmacological effects. The dry powder of *Luffa cylindrica* (L.) fruit was defatted using petroleum ether and extracted with 2000 ml of ethanol using Soxhlet at 45 °C for 3 days [31]. Whatman filter paper (Grade 1- particle retention 11 μ m) was used to filter it, and the filtrate was concentrated in a water bath and then stored in an airtight sterile container for further research. Table 1 shows the obtained residue weights and extractive values of the extraction.

Phytochemical screening

Preliminary phytochemical results were carried out in order to identify the presence or absence of certain phytochemicals. The tests performed utilizing ethanolic extract were subjected to a variety of qualitative tests to identify the present phytoconstituents [25]. Table 2 lists the preliminary phytochemical screening results of ethanolic extract of *Luffa cylindrica* (L.) fruit. The ethanolic extract was subjected to variety of qualitative tests to identify the present phytoconstituents.

Determination of the total phenolics content

The total phenolic content of the *Luffa cylindrica* (L.) fruit's ethanolic extracts was determined using the spectrophotometric technique [32]. In a nutshell, 2.5 mL of the Folin-Ciocalteu's reagent (10 percent deionized water) and 4 mL of a NaHCO₃ solution (7.5 percent w/v) were combined with 0.5 mg of plant extract. The reaction mixture was then

Table 1 Extractive value of the plant ext	ract
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Content	Weight of the test substance taken	Extractive value
<i>Luffa cylindrica (L.)</i> fruit	1000 gm	5% w/w

 Table 2
 Preliminary
 phytochemical
 screening
 tests
 of
 Luffa
 cylindrica (Linn) fruit

S.no	Phytochemical tests	Result
1	<i>Carbohydrates</i> (a) Molisch test (b) Benedict test	Presence
2	<i>Alkaloids</i> (a) Mayer test (b) Wagner test	Presence
3	<i>Glycosides</i> (a) Modified borntrager test (b) Legal test	Presence
4	<i>Flavonoids</i> (a) Shinoda test (b) Lead acetate test	Presence
5	<i>Phytosterols</i> (a) Salkowski test (b) Liebermann-Burchard test	Absence
6	<i>Tannins</i> (a) Gelatin test (b) Ferric chloride test	Presence
7	Saponins-foam test	Presence
8	Phenols- Ferric chloride test	Presence
9	Triterpenoids-vanillin-sulfuric acid test	Presence

incubated for 30 min at room temperature with intermittent shaking for color development. The absorbance of the resulting reaction mixtures (blue color) was measured at 765 nm using a spectrophotometer (Jasco V-730). Gallic acid equivalent (mg of GAE/g dry weight) was used to express the overall amount of phenolics in the investigated samples.

Determination of the total flavonoids content

The total flavonoid content of the *Luffa cylindrica* (L.) fruit's ethanolic extracts was determined [33] using the spectrophotometric technique [21]. The extract was produced at the same concentration of 1 mg/mL for analysis. In a nutshell, 0.5 mg of the extract was combined with 0.1 mL of a 10% AlCl₃ solution, 0.1 mL of a 1 mol/L potassium acetate solution, and 4.3 mL of distilled water. The reaction mixture was then incubated at room temperature for 30 min. The spectrophotometer measured the absorbance at 510 nm (Jasco V-730). The standard calibration curve (10–100 mg/mL) was created using quercetin (Sigma-Aldrich). The quercetin equivalent (mg QE/g dry weight) was used to express the overall flavonoid content of the samples under investigation.

Determination of the total alkaloids content

The total alkaloid content of the *Luffa cylindrica* (L.) fruit's ethanolic extracts was determined by spectroscopic technique [34]. In a nutshell, 0.5 mg of *Luffa cylindrica* (L.) fruit extract was diluted in 20 mL of ethanol

solution (1:1) and filtered. The resulting mixture was poured into several funnels for separation. $\rm H_2SO_4$ was blended with the filtrates at a 60% ratio (5 ml). After the combination had been left for 5 min, 5 ml of 0.5 percent formaldehyde was added, and the mixture was then left for 3 h. The spectrophotometer measured the absorbance at 565 nm (Jasco V-730). The standard calibration curve (10–100 mg/mL) was created using the atropine standard solutions, which were manufactured similarly. The atropine equivalent (mg AE/g dry weight) was used to reflect the overall amount of alkaloids in the examined samples.

Determination of the total saponins content

The total saponin content of the *Luffa cylindrica* (L.) fruit's ethanolic extracts was determined by vanillin method using the spectrophotometric technique [35]. In a nutshell, 0.5 mg of *Luffa cylindrica* (L.) fruit extract was combined with 0.5 mL of an 8 percent vanillin solution and 5 mL of an H₂SO₄ solution (72%). The resulting liquids were carefully combined before being set aside to chill on ice. They were then incubated in a water bath at 60 °C for 15 min. On the ice, the mixtures were cooled once again, and the spectrophotometer measured their absorbance at 560 nm (Jasco V-730). Oleanolic acid (10–100 mg/mL) was utilized as a reference for creating the calibration curve. Oleanolic acid equivalent (mg OA/g dry weight) was used to express the overall amount of saponins.

Determination of the total triterpenoid content

The total triterpenoid content of the *Luffa cylindrica* (L.) fruit's ethanolic extracts was determined by using the spectrophotometric technique [36]. Briefly, 2 mL of H_2SO_4 and 0.5 mg of plant extract in acetic anhydride were combined in a microplate before being incubated for 10 min at room temperature. A calibration curve was created based on ginsenoside Re (10–100 mg/ml) as a standard. At 350 nm, the sample's absorbance was measured compared to a blank sample made up of acetic

anhydride-infused plant extract. Triterpenoid concentration was measured in milligrams of ginsenoside Re equivalents (GRE)/g of plant extract, and samples were conducted in triplicate. All the determination content is mentioned in Table 3.

Identification and quantification analysis of constituents by LC-MS analysis

LC-MS (liquid chromatography-mass spectroscopy) analysis was carried out using Column (Hypersil GOLD C18 100 × 2.1 mm-3 MICRON); 30min_+ESI_01112021_ MSMS.m method and detected with Q-TOF analyzer for the ethanolic extract of Luffa cylindrica (L.). MS Q-TOF (G6550A) component was deployed with 200 Ms Abs threshold and ion mode as Dual AJS ESI. Acquisition mode for AutoMS2 was set to Ms Min range to 120 (m/z) and max of 1100 m/z. Source parameters were set with gas temp $(250^{\circ}C)$; gas flow (13 l/min); nebulizer (35 psig). Injection model with needle wash of 5 µL volume was set to 3-secs wash time. Solvent composition was 0.1% formic acid in water (95%) for Channel A and 90% Acetonitrile +10% Water +0.1% Formic acid (5%) for channel B. Flow rate was kept constant to 0.300 mL/min, 1200 bar constant pressure, and 40 °C temperature. Nonetheless, to maximize the number of the monitored metabolites ions, MS analysis was carried out in a negative ionization mode. The resulting total ion chromatogram is a time vs. area plot that displays each constituent's overall response based on the abundance of its molecular ions. The identification of constituents by LC-MS analysis is mentioned in Table 4.

Cholinesterase activity

Ellman's approach was used to determine AChE inhibition. In a 1-mL cuvette, 640 L of Tris–HCl buffer with a pH of 8 and 0.12 U of AChE enzyme were introduced to the reaction mixture [37]. This reaction mixture was incubated for five minutes at room temperature. 100 mL of DTNB (Ellman's reagent of 7.5 mM) and 100 mL of

Table 3 Total content of Luffa cylindrica (Linn) fruit

Concentration (mg/mL)	Phenol content absorbance	Flavonoid content absorbance	Alkaloid content absorbance	Saponin content absorbance	Triterpenoid content absorbance
20	0.251	0.238	0.248	0.198	0.198
40	0.322	0.289	0.301	0.269	0.274
60	0.471	0.396	0.449	0.387	0.366
80	0.628	0.547	0.597	0.463	0.472
100	0.855	0.786	0.823	0.578	0.594
Unknown concentration	0.214	0.178	0.197	0.078	0.131

Unknown concentration is given in average of triplicates obtained from the result

Compounds	RT (min)	Molecular mass	Abundance	Molecular formula	Hits (DB)	BBB passage
Prolyl-arginine	1.428	271.1645	342,921	C ₁₁ H21N5O3	7	1
Mahaleboside	3.206	324.0833	145,936	C ₁₅ H ₁₆ O ₈	1	1
Feruloylputrescine	3.519	264.1456	105,908	C ₁₄ H ₂₀ N ₂ O ₃	8	1
Chlorogenic acid	4.033	354.0899	462	C ₁₆ H ₁₈ O ₉	9	1
Crotanecine	4.435	171.0889	140,956	C ₈ H ₁₃ N0 ₃	4	1
15-Hydroxymarasmen-3-one	4.727	264.1353	422,747	C ₁₅ H ₂₀ 0 ₄	10	0
Perlolyrine	6.362	264.279	81,774	C ₁₆ H ₁₂ N ₂ O ₂	9	0
Phrymarolin I	6.291	488.1269	144,012	C ₂₄ H ₂₄ O ₁₁	4	1
Diplodiatoxin	6.928	308.1979	47,386	C ₁₈ H ₂₈ 0 ₄	10	0
Dihydrocapsaicin	7.318	307.2122	234,849	C ₁₈ H ₂₉ N0 ₃	3	0
Morindone	8.488	270.052	235,899	C ₁₅ H ₁₀ O ₅	10	1
Corchorifatty acid	8.665	328.222	159,453	C ₁₈ H ₃₂ O ₅	5	1
Polyporusterone B	15.364	476.3098	113,332	C ₂₈ H ₄₄ O ₆	5	1
Camelledional	16.748	440.328	326,723	C ₂₉ H ₄₄ O ₃	10	1
2alpha-(Hydroxymethyl)-5alpha- androstane-3beta,17beta-diol	16.961	322.2495	194,509	$C_{20}H_{34}O_3$	7	1
Taxa-4(20),11(12)-dien-5α,13α-diol	18.336	304.2392	369,304	C ₂₀ H ₃₂ O ₂	10	1

Table 4 Identification of constituents by LCMS analysis

All the compounds obtained following Lipinski's rule of five with (a) not more than 5 hydrogen bond donors, (b) not more than 10 hydrogen bond acceptors, (c) molecular weight less than 500 g/mol, (d) partition coefficient log *P* less than 5. 1 in blood–brain barrier (BBB) passage indicates that it does not pass BBB and 0 indicates vice versa

 Table 5
 Cholinesterase activity

Tested sample concentration (µg/ml)	OD value a triplicates	Mean of OD value		
Control	0.76	0.84	0.92	0.84
10	0.543	0.746	0.854	0.714
50	0.387	0.573	0.623	0.528
100	0.311	0.484	0.478	0.424
250	0.289	0.267	0.381	0.312
500	0.146	0.123	0.198	0.156

acetylcholinesterase (AchEI of 1 mM) were combined, and a spectrophotometer was used to measure the reading at 405 nm. Triplicates of each experiment were performed. The percentage of inhibition was used to characterize the anti-cholinesterase activity and is mentioned in Tables 5 and 6.

In-silico docking analysis

Ligand and receptor preparations

Based on the Lipinski rule of five, the ADMET analysis [38] was performed for all the phytocompounds chosen from the LC–MS data fractions. Using the software ChemDraw Ultra 12.0, the structure of perlolyrine was made and converted to.pdb format using Molegro molecular viewer. The docking computation and techniques to replicate the pocket binding residues were validated using co-crystal ligands. Autodock v4.0 (version) software was employed to open macromolecule and
 Table 6
 Percentage inhibition of AchE in extract

Tested sample concentration (µg/ ml)	Percentage of inhibition (in triplicates)			Mean value (%)
Control	100	100	100	100
10	28.55	11.19	7.17	15.64
50	49.08	31.79	32.28	37.72
100	59.08	42.38	48.04	49.83
250	61.97	68.21	58.59	62.92
500	80.79	85.36	78.48	81.54

ligand to initiate docking and mentioned in Table 7. In the current investigation, we started the docking analysis with the standard settings such as ligand preparation and saving in.pdbqt format. Grid was prepared for both ligand and macromolecule to cover the area and run for autogrid 4.0. Further docking was initiated to set all the parameters required and saved as a Lamarckian genetic algorithm to further run in Autodock v4.0. All molecules were deleted and analyzed for clusterings which were further confirmed as rank by energy. The RMSD table was used to cluster all the transformations, and the most advantageous binding postures were chosen based on their low free energies and low inhibition constants. The targets chosen for this investigation were based on earlier studies, and protein data bank was used to retrieve all the 3D crystal structures in PDB format. One of the best molecules discovered using LC-MS analysis is

 Table 7
 Binding energy of periolyrine with selected targets

S. no	Receptors	PDB ID	Binding energy (kcal/ mol)
1	Amyloid beta	1IYT	- 46.1
2	Cox-2	1PXX	- 7.66
3	Butyrylcholine esterase	6IOC	- 7.59
4	Tau kinase	1J1C	- 7.59
5	Keap-1	2FLU	- 7.26
6	SOD	1CB4	- 6.77
7	Catalase	2CAG	- 6.54
8	Acetylcholinesterase	3LII	- 6.38
9	GSK-3β	1UV5	- 6.14
10	Glutathione peroxidase	2P31	- 5.75

perlolyrine. Perlolyrine showed improved idiopathic pulmonary fibrosis [39], osteoarthritis [40], and vascular dementia [41], respectively.

Purification and refinement of proteins and ligands

All the proteins utilized in the study were purified using Biovia Discovery studio to remove unwanted ligands, water molecules, and other contaminants. For better interactions, polar hydrogens were added to the protein during the preparation process, followed by Kollman calculations and determining Gasteiger charges. The PubChem database was used to visualize the 2D and 3D structures of perlolyrine and confirm with our structure (PubChem CID: 160179).

Root-mean-square validation of docked structures

Autodock v4.0 predicted 9 different docking sites for the studied ligand. All of them had different RMSD values. However, we have considered only those sites which gave RMSD value 0 for the betterment of our understanding.

In-silico toxicity predictions

Based on a proven Lipinski concept, compounds' potential drug-likeness was anticipated [42]. All pharmacologically significant substances changed their structures to their canonical, streamlined molecular-input line-entry scheme (SMILE). To identify organ toxicities and toxicological endpoints, obtained SMILE was entered to OSI-RIS Property explorer [43]. A drug score measure is used to select molecules as drug candidates. The likelihood of a molecule being regarded as a drug candidate increases with the drug score value as mentioned in Table 8 [44].

Statistical analysis

Each experiment was repeated in triplicate, and data are presented as mean \pm standard deviation (SD).

Results

Total phenolic content

Total phenolic content was determined to be 21.42 mg of GAE/g, respectively, when expressed in terms of gallic acid equivalent.

Total flavonoids content

The total flavonoid content expressed in terms of quercetin equivalent was found to be 19.57 ± 2.39 mg of QE/g, respectively.

Total alkaloids content

The total alkaloid content expressed in terms of atropine equivalent was found to be 20.39 ± 1.47 mg of AE/g, respectively.

Total saponin content

The total saponin content expressed in terms of oleanolic acid equivalent was found to be 14.39 ± 2.36 mg of OA/g, respectively.

Total triterpenoid content

The total triterpenoid content expressed in terms of ginsenoside Re equivalent was found to be 9.48 ± 1.64 mg of GRE/g, respectively (Fig. 1).

Liquid chromatography-mass spectrometer (LC-MS) analysis

The outcomes showed the presence of numerous medicinally significant substances, including Mahaleboside (Coumarin glycosides), Chlorogenic acid (Polyphenols), Crotanecine (Pyrrolizidine alkaloids), Perlolyrine (Harmala alkaloids), Phrymarolin I (Benzo-dioxoles), Diplodiatoxin (Gamma keto acids), Dihydrocapsaicin (Capsaicinoid), Morindone (Anthraquinone), Corchorifatty acid (Linoleic acid), and Polyporusterone B (triterpene carboxylic acid), which were all tentatively identified using database by comparing their molecular

Table 8 Predicted toxicity risks

logP	logS	H Donor	H acceptor	Mutagenic	Tumorigenic	Irritant	Reproductive effective
2.45	- 3.88	2	4	Orange	Green	Green	Green

Green: Low risk (Drug conform behavior); Orange: Medium risk



fragmentation patterns. The results of the LCMS analysis of the fruit's ethanolic extract, *Luffa cylindrica*, are shown in Table 4 and Fig. 2.

Liquid chromatography–mass spectrometer (LC–MS) chromatogram of Luffa cylindrica extract

All the compounds were subjected for screening based on Lipinski's rule of five, and perlolyrine compound (Fig. 3, Fig. 4) was able to cross blood–brain barrier which was observed through online server tool SwissADME (http:// www.swissadme.ch/). Perlolyrine compound was later subjected for in-silico molecular docking to see potential score of various receptors involving Alzheimer's diseaseassociated oxidative stress (Fig. 5).

Cholinesterases are specialized carboxylic ester hydrolases that break down esters of choline, and its activity with absorbance is mentioned in Figs. 6 and 7.

Molecular docking analysis

According to the LC–MS analysis, *Luffa cylindrica* (L.) extract included 80 bioactive compounds. Table 4 lists the top compounds with peak areas. These phytochemicals' effects on oxidative stress-related target proteins for Alzheimer's disease were examined. To clarify the binding affinities to the target proteins, phytoligand docking

studies were conducted using the Autodock v4.0 software. In general, the Lipinski rule of five was used to select the 80 phytocompounds that were discovered. All ten selected targets' corresponding protein structures were docked against perlolyrine; the best-screened ligand found using ADMET screening based on various criteria, including solubility, toxicity, absorption, molecular weight, and excretion. The best-docked complexes were then filtered based on their binding energy. Using Biovia discovery studio, the interaction between the ligand and the target protein was illustrated. Using Biovia discovery studio, distinct bonds between the ligand and the target protein were also displayed in 2D interaction diagrams. The binding conformations for perlolyrine ([5-(9H-pyrido[3,4-b] indol-1-yl) furan-2-yl] methanol) had the highest binding energy values for amyloid beta protein.

The perlolyrine docking outcomes with target proteins were rated based on binding energies. Perlolyrine docking results with proteins like catalase, glutathione peroxidase, superoxide dismutase, Cox-2, AchE, BuchE, A, GSK-3, tau kinase, and keap-1 indicated that the ligand has a stronger affinity for amyloid beta, which is shown in Fig. 8 and is a critical regulator in Alzheimer's disease. Figure 8 shows the docked structure depicting ligand



Fig. 2 Liquid chromatography-mass spectrometer (LC-MS) profile of metabolites from ethanolic fraction of Luffa cylindrica (L.) extract





6200 series TOF/6500 serie Q-TOF B.05.01 (B5125.3)

Sample Group

Version

Acquisition SW

interactions with essential amino acids through hydrogen bonds and Vander Waal forces. Table 7 shows the identified compound's affinity for proteins associated with Alzheimer's disease.

Info

Amyloid beta (1IYT) with a binding energy of -46.1 kcal/mol showed interactions with essential

amino acids, viz., ASN27, GLY29, GLY25, LYS28, SER26. Cox-2 (1PXX) with a binding energy of - 7.66 kcal/ mol showed interactions with essential amino acids, viz., THR-A206, ALA202, TRP-A387, LEU-A391, ALA-A199, HIS-A388, PHE-A210, HIS-A386, HIS-A207. Butyrylcholine esterase (6IOC) with a binding energy





Fig. 4 Structure of perlolyrine

of - 7.59 kcal/mol showed interactions with essential amino acids, viz., ASP-A170, GLY-A169, PRO-A171, GLU-A132, PRO-A133. Tau kinase (1J1C) with a binding energy of -7.59 kcal/mol showed interactions with essential amino acids, viz., GLN-A151, SER-A147, LEU-A252, THR-A152, GLY-A253, ALA-A143, LEU-A153, TYR-A146. Keap-1 (2FLU) with a binding energy of – 7.26 kcal/mol showed interactions with essential amino acids, viz., GLY-X367, VAL-X606, ALA-X366, ARG-X415. SOD (1CB4) with a binding energy of -6.77 kcal/ mol showed interactions with essential amino acids, viz., LEU-X557, GLY-X367, VAL-X606, ALA-X556, ARG-X415. Catalase (2CAG) with a binding energy of - 6.54 kcal/mol showed interactions with essential amino acids, viz., LYS-A85, LEU-A132, VAL-A110, CYS-A199, LEU-A188, ALA-A83, VAL-A70. Acetylcholine esterase (3LII) with a binding energy of -6.38 kcal/ mol showed interactions with essential amino acids, viz., TYR-B341, SER-B293, TRP-B286. GSK-3β (1UV5) with a binding energy of -6.14 kcal/mol showed interactions with essential amino acids, viz., LEU-A188, ILE-A109, VAL-A110, LEU-A198, CYS-A199, ASP-A200, LEU-A132, ALA-A83, VAL-A135. Glutathione peroxidase (2P31) with a binding energy of -5.75 kcal/mol showed interactions with essential amino acids, viz., HIS-B77, ARG-B177, PRO-B151, HIS-B78, VAL-B176, PRO-B76.

In-silico toxicity predictions

The toxicity of the compound was identified using SMILES in OSIRIS property explorer and identified as safe other than mutagenic response as displayed in Table 8.

Discussion

Luffa cylindrica (L.) is used in the food and cosmetic industries since it is a rich source of bioactive chemicals. A study observed that FT-IR analysis and physico-chemical characterization of luffa seed oil showed higher

content of unsaturated hydrocarbon and esters, enabling the oil to be suitable for the production of paint, pharmaceutical substances, and cosmetics [45]. Studies have shown that the fruit and seeds of the luffa plant contain phytochemicals that are good for human health. Gourd vegetables, especially bitter and bottle gourds, are preventative against gastric ulcers, bacterial and viral infections, arthritis, diabetes, hypertension, and cardiac conditions [46]. A study documented that vegetable peels have beneficial activities and antioxidant potential and promoted the utilization of agricultural wastes to achieve nutritional benefit at zero cost and maintain proper health [47]. Bioactive substances, particularly phytochemicals and antioxidants, are necessary for large-scale and significant bioactivities. Their presence necessitates adequate validation and verification of bioactive metabolites. According to this analysis, the fruits of L. cylindrica contained a reliable amount of phenolics, flavonoids, terpenoids, and alkaloids. As secondary metabolites are now shown to have multiple functions, a fresh wave of genetic and pharmacological studies has further blurred these boundaries [48]. The hydroxyl radical scavengers -carbolines and tetrahydrocarbolines have activity comparable to indole melatonin, a potent scavenger of hydroxyl radicals and antioxidants [49].

Several studies suggest various pharmacological actions, such as antioxidant, antibacterial, anti-inflammatory, and neuroprotective effects, present in harmala alkaloids [50]. Proximate analysis is a quantitative analysis performed to confirm the amount of identified phytochemicals to understand their role in the treatment of disease. Proximate analysis of various processing methods was seen in luffa gourds and confirmed the presence of several variables, including dry matter, crude protein, crude fiber ash, etc. [51, 52]. Alkaloids are renowned for their ability to scavenge free radicals, inhibit hydrolytic and oxidative enzymes, and have anti-inflammatory properties [53]. Alkaloids have been studied for their neuroprotective properties for a long time. They have disclosed their neuroprotective properties and can be useful for curing cognitive impairments and associated diseases such as Alzheimer's [54, 55]. Alkaloids are essential in the pharmaceutical and cosmetic industries. They have antiparasitic properties, anti-plasmodial, anti-corrosive, antioxidative, antibacterial, and insecticidal properties [56]. Among the most common diseases of the twenty-first century, dementia and depression are becoming more significant as the world's population ages [57].

The most recent therapeutic approach, which takes into account the primary pathobiochemical changes in the system caused by Alzheimer's disease, is an anti-amyloid therapy intended to slow down the progression of the disease [58]. Cucurbitan species are reported highly





MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
265.0953	1	81773.91	C16H12N2O2	(M+H)+
266.0987	1	15840.32	C16H12N2O2	(M+H)+

MSMS Spectrum



MS/MS Spectrum Peak List				
m/z	z	Abund		
163.0389		62.9		
167.0582	1	125.81		
196.964		61.79		
206.0818	1	1027.46		
219.091	1	235.59		
223.9756		117.12		
235.0842	1	117.65		
247.0848	1	602.53		
265.0948	1	2576.85		
271.0575	1	146.46		

Fig. 5 Ms zoomed spectrum of perlolyrine



Fig. 6 Percentage Inhibition of Ache in Luffa cylindrica extract



Fig. 7 Percentage inhibition of AchE in extract

on inflammation, cancer, and genotoxic activities [59–61]. Antineoplastic and anti-inflammatory action is the pharmacological activities that have received the greatest attention in the Cucurbitaceae family, where anti-inflammatory, antioxidant, and immunomodulatory potential has been studied recently. Mechanisms of the inhibition of COX-2, NOS, oxidative stress, pro-inflammatory cytokines, and modulation of acquired immunity proteins are highly focused and can also be studied in neuro-degenerative diseases like Alzheimer's disease, which also works on a similar mechanism [62].

Acetylcholine is one of the molecules responsible for communication between nerve cells and the brain. Alzheimer's disease symptoms begin when its level is lowered. The drugs known as acetylcholinesterase inhibitors stop the body from breaking down acetylcholine. Alzheimer's line of research lies on anticholinesterase and antioxidant mechanisms as it targets multiple pathways of dementia [63]. Acetylcholinesterase inhibitors (AChEIs) are primarily used to inhibit AChE since there is enough evidence of a deficit in cholinergic neurotransmission in AD to support therapy intended to repair this deficiency. AChE inhibitors also have antioxidant properties and control amyloid plaque development. Tacrine, donepezil, and rivastigmine are a few AChE inhibitors that have already received FDA approval. The majority of these inhibitors, however, are connected to limited availability and acute toxicity [64].

The extract of Luffa cylindrica (L.) showed inhibitory activity on the AchE enzyme. Similar reports have been reported by Patel SB et al. (2021) in Luffa echinata [1]. Liquid chromatography-high-resolution mass spectrometry (LC-HRMS) has become a robust analytical method for massive target screening. Its use can be expanded to assess the possibility of natural product poisoning in clinical situations [65]. An ultraliquid chromatography-quadrupoleperformance time-of-flight-mass spectrometry has been employed in several studies to characterize the potential components [66]. Additionally, LC-MS analysis revealed the presence of compounds like Mahaleboside (Coumarin glycosides), Chlorogenic acid (Polyphenols), Crotanecine (Pyrrolizidine alkaloids), Perlolyrine (Harmala alkaloids), Phrymarolin I (Benzodioxoles), Diplodiatoxin (Gamma keto acids), Dihydrocapsaicin (Capsaicinoid), Morindone (Anthraquinone), Corchorifatty acid (Linoleic acid), and Polyporusterone B (triterpene carboxylic acid) were known to possess anticancer, depression, hypertension, anti-inflammatory activities [67-70].

Compounds detected in high-resolution LC-MS analysis could help exhibit potent bioactivities in L. cylindrica. To find potential Alzheimer targeting compounds, affinity ultrafiltration LC-MS/MS was employed to screen and identify target compounds from the extract. A total of five AChE ligands were identified by LC-MS/ MS and confirmed the activity using molecular docking and in vitro acetylcholinesterase data [71]. Similarly, we employed HR-LCMS to screen extract and identify the potential components present and to confirm their activity by performing molecular docking and in vitro acetylcholinesterase activity. We performed LC-MS/MSbased molecular networking to ascertain their chemical composition and quantify existing metabolites using a pre-generated plant extract library. This provided a novel method for metabolite authentication and profiling and enabled the discovery of new, allegedly undiscovered metabolites for biological separation and evaluation in the future [72]. We identified various bioactive compounds, among which perlolyrine molecule showed less toxicity in comparison with other compounds by OSIRIS property explorer, and it had the potential to cross the blood-brain barrier. We planned for in-silico molecular docking of perlolyrine with several receptors, which are closely associated with Alzheimer's disease and oxidative stress, a pathological hallmark of Alzheimer's disease.



Fig. 8 Docking of perlolyrine with various receptors in Alzheimer's disease associated with oxidative stress



Fig. 8 continued

Conclusion

The current therapeutic strategies against AD include acetylcholinesterase inhibitor, anti-tau, N-methyl-Daspartate antagonist, etc., However, core pathogenesis of AD lies in oxidative stress mechanism and efficient compound with less toxicity is the need. Moreover, many researchers are focusing on oxidative stress-related mechanism in order to prevent extension of disease. This study's investigation of the extracts revealed that *Luffa cylindrica* (L.) extracts act on many target proteins of Alzheimer's disease and have sufficient AchE activity. Furthermore, this extract showed the highest concentrations



Fig. 8 continued

of various phytochemical components (alkaloids, phenolics, flavonoids, and saponins). Numerous investigations have confirmed that these phytochemical elements give the extract its bioactivity. The LCMS profile identified prominent metabolites that were reported to be good antioxidants and reported various other biological activities. The compound perlolyrine could be of interest against oxidative stress-related diseases like Alzheimer's and other pharmacological benefits. Based on current findings, the study reports the presence of a promising bioactive compound (perlolyrine) and, in turn, provides an optimistic note in exploring its biological activity (in vivo) in oxidative stress-related Alzheimer's disease.

Abbreviations

DPPH ABTS FRAP CUPRAC ER SDH ATP GSK-3 NF-KB Keap MAP30 AD LC-MS QE/g AE/g OA/g GRE/q	2,2-Diphenyl-1-picrylhydrazyl 2,2'-Azino-bis (3-ethylbenzothiazoline-6-sulfonic acid) Ferric-reducing antioxidant power assay Cupric ion-reducing antioxidant capacity Endothelin receptor Succinate dehydrogenase Adenosine triphosphate Glycogen synthase kinase-3 beta Nuclear factor kappa-light-chain-enhancer of activated B cells Kelch-like ECH-associated protein Mitogen-activated protein Alzheimer's disease Liquid chromatography-mass spectroscopy Quercetin equivalent per gram Atropine equivalent per gram Oleanolic acid equivalent per gram
GRE/g	Ginsenoside Re equivalents per gram

O-TOF	Quadrupole time-of-flight analyzer
	E E/ Dithiphis (2 pitrobanzois acid)
DIND	5,5°-Dithiodis-(2-hitrodenzoic acid)
AchEl	Acetylcholinesterase inhibitor
ADMET	Absorption, distribution, metabolism, excretion, toxicity
RMSD	Root-mean-square deviation
SMILE	Streamlined molecular-input line-entry scheme
BBB	Blood–brain barrier

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Author contributions

ASS helped in conceptualization, data curation, formal analysis, writing—original draft, writing—review & editing. CV was involved in conceptualization, data curation, formal analysis, writing—review & editing. All the authors critically reviewed the manuscript for intellectual content. All authors approved the final version of the manuscript for publication.

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Availability of data and materials

Data sharing is not applicable to this article as no datasets were generated or analyzed during the current study.

Declarations

Ethics approval and consent to participate Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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