

Chemical and Biological Evaluation of the Aqueous Extract of *Peumus boldus* Molina (Monimiaceae) Leaves

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ABSTRACT

Background: Traditional medicine has inspired the scientific community to discover natural compounds with the ability to inhibit the enzyme acetylcholinesterase (AChE) and with significant antioxidant capacity. These themes are associated with therapeutic processes to decrease the symptoms of Alzheimer's disease (AD). Acetylcholine (ACh) is an important neurotransmitter in the central nervous system responsible for transmitting the nervous impulse and increasing intestinal motility and dilation of sphincters in the gastrointestinal tract. As the species *Peumus boldus* Molina is widely known and used in folk medicine to treat gastrointestinal discomfort, its constituents could have an affinity for AChE and could be used to treat AD. **Materials and Methods:** This study evaluated the chemical, inhibitory activity of AChE (using the Elmann method and *in silico*), antioxidant capacity (DPPH, ABTS, FRAP, and oxidation inhibition of the β -carotene/linoleic acid system), and quantification of total leaf phenols from a commercial sample of *P. boldus*. **Results:** The mass spectral analysis allowed identification of the alkaloids isoboldine, boldine, clocaurine, laurotetanine, *N*-methylurotetanine, reticuline, and *N*-methylcoclaurine. In the *in silico* prediction test, the alkaloid *N*-metillaurotetanine was the best acetylcholinesterase enzyme inhibitor, with energy of -10.1 kcal mol⁻¹. **Conclusion:** These results are of scientific relevance for the discovery of new drugs to treat dementia associated with Alzheimer's disease and stimulate the continuity of chemical and biological studies (anticholinesterase and antioxidant action) of prepurified fractions and substances isolated from *P. boldus* leaves.

Key words: Boldo, Acetylcholinesterase, Antioxidant activity, Acetylcholine, Alzheimer.

INTRODUCTION

Alzheimer's disease (AD) is a neurodegenerative pathology characterized by cognitive decline and mental degeneration, with massive and progressive loss of neurons in different regions of the brain. In 2015, approximately 46.8 million people were diagnosed with AD, and by 2050 this number could reach 131.5 million therefore, new approaches are needed for its treatment.^[1-3]

Since the discovery of the disease, researchers have recognized that its symptoms may be associated with the development of neuritic (senile) plaques, by the extracellular deposit of β -amyloid ($A\beta$) protein aggregates, and by neurofibrillary tangles, which occur through intracellular deposit of aggregates of phosphorylated tau protein.^[1] The $A\beta$ protein originates through the cleavage of a transmembrane protein called the amyloid precursor protein (APP), and this precursor has an $A\beta$ domain inserted in the membrane that is cleaved by two types of protease enzymes called secretases.^[2] When the cleavage is performed by the α -secretase enzyme, a non-amyloid soluble fragment (p3) is formed. If the protein is cleaved by

the β -secretase enzyme, the $A\beta$ amyloid fragment is formed, which is insoluble and rich in crossed β sheets, and deposit in brain tissue is followed by aggregation and polymerization in the form of senile plaques.^[1,3,4] A common hypothesis was that amyloid plaque formation was the main cause of dementia; however, this hypothesis has lost strength due to new studies that observed loss of synapses in regions of the brain where these plaques do not exist.^[5]

The TAU protein is an important component of the neuronal cytoskeleton, mainly in the axons, and is responsible for the stabilization of microtubules. Aggregates of this protein in the hyperphosphorylated state result in the formation of highly insoluble intracellular tangles with disruption of the cell cytoskeleton and consequent neuronal death.^[4,5] These, together with senile plaques, represent the pathological findings characteristic of Alzheimer's disease. When combined, these two events decrease the number of neuronal cells and consequently result in functional deficit of synapses and atrophy

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of the cerebral parenchyma, causing dementia.^[6] Many studies associate the accumulation of A β protein as the activator in the hyperphosphorylation of the tau protein, but the factors that lead to these events are still unclear.^[5]

Studies reveal that the accumulation of amyloid plaques can stimulate the production of reactive oxygen and nitrogen species (ROS and RNS),^[5,6] justifying the search for antioxidant compounds that can protect neurons from the oxidative processes generated by the accumulation of the amyloid plaques. Oxidative stress is characterized as an imbalance between the production of free radicals and the ability of the cell to defend against them.^[6-10] High oxygen consumption and relatively low levels of antioxidants result in the susceptibility of brain tissue to oxidative damage. Numerous oxidative stress biomarkers have been found in patients with Alzheimer's, including excessive lipid peroxidation, neuronal degeneration, protein oxidation, and DNA and RNA glycooxidation and oxidation.^[6]

Studies also found that oxidative stress can increase the production and accumulation of the β -amyloid peptide and facilitate the phosphorylation and polymerization of the Tau protein, forming a vicious cycle that promotes initiation and progression of AD.^[5] Clinical studies in humans have not yet obtained effective results,^[6] for the treatment of AD; therefore, interest in finding an effective therapy for AD remains high. One of the promising avenues for achieving this goal is to inhibit enzymes capable of degrading neurotransmitters that transmit the nerve impulse to the postsynaptic neuron through nicotinic and muscarinic receptors.

One approach to treating AD to inhibit the AChE enzyme, in which main objective is to minimize the deficit in neurotransmitter molecules and enable an increase in signal transmission in patients affected by AD.^[11,12] Some chemically synthesized compounds display an affinity with the enzyme, the most relevant being donepezil, rivastigmine, tacrine, and recently galantamine. However, these compounds have low antioxidant potential and adverse side effects such as nausea, vomiting, dizziness, anorexia, abdominal pain, and headaches. These conditions encourage the scientific community to continue research to discover new molecules with inhibitory capacity against the AChE enzyme.

One of the active sites of AChE is called the "aromatic throat", site presents itself as a deep and narrow channel about 20 Å deep. Its composition contains more than 50% amino acids with conserved aromatic residues (Phe120, Phe288, Phe290, Phe330, Phe331, Trp84, Trp233, Trp279, Trp432, Tyr70, Tyr121, Try130, Try334, and Tyr442).^[13] Other active sites consist of the catalytic triad (Ser200, Glu327, and His440), anionic catalytic site (Trp84, Glu199, and Phe330), oxyanion cavity (Gly118, Gly119, and Ala201), and acyl group cavity (Phe288 and Phe290).^[14]

Galantamine, recently discovered as an important enzyme inhibitor is an alkaloid found in species of the family *Amaryllidaceae*. Similar to this compound, other alkaloids also have important anticholinesterase activity,^[15-19] and are capable of reestablishing the neurotransmission system. Therefore, studies have explored these groups as promising natural inhibitors of acetylcholinesterase.

Every medicinal plant causes a therapeutic resource that is used to alleviate symptoms or cure diseases. One plant widely used for medicinal purposes is boldo, due to its extensive capacity to treat problems related to the stomach and liver, especially by people who commonly drink alcoholic beverages. One of the mechanisms of action of the ACh neurotransmitter is to increase intestinal motility; therefore, the chemical constituents present in boldo leaves could have the capacity to inhibit the catalytic action of AChE in the hydrolysis of ACh.

Boldo (*Peumus boldus*) is a plant native to the central and southern regions of Chile and belongs to the family Monimiaceae.^[18] This herb

is widely consumed in Brazil and other countries around the world due to its wide spectrum of pharmacological effects. It is mainly indicated for the treatment of various infections in the digestive and hepatobiliary system. In addition to its role as a liver protector and digestive stimulant, *P. boldus* is traditionally used as an anti-inflammatory, antispasmodic, and nervous sedative agent.^[18,20] The aforementioned pharmacological properties are obviously associated with the chemical composition of the boldo leaves.

The most important active principles contained in the boldo leaves are alkaloids and flavonoids. Among alkaloid compounds, boldine is the most important active principle of *P. boldus*, and some works indicate that the anti-inflammatory activity and antioxidant properties of the species are related to this substance.^[7,18,20,21]

Boldo leaves also have tannins, essential oil, flavonoids, and glycolipids.^[18] Its leaves are reported to contain sparteine; therefore, its use is not recommended during pregnancy, as this alkaloid can accelerate delivery (oxytocic activity).^[22]

Considering the presence of alkaloids is major constituents of the species, this group of compounds might be healthy alternatives to synthetic inhibitors.

In this context, this work aimed to discover potential AChE inhibitors in the aqueous extract of leaves from a commercial sample of *P. boldus* and to evaluate the antioxidant activity of these samples using the free radical DPPH and ABTS methods, auto-oxidation of the β -system carotene/linoleic acid, and iron reduction (FRAP), as well as quantifying the content of total phenolic compounds.

MATERIALS AND METHODS

Collection and identification of plant material

Boldo leaves were obtained from a store in the city of Itacoatiara, AM, Brazil, in October 2018.

Botanical identification was carried out based on the analysis of diagnostic characteristics such as shape, phyllotaxis, and leaf indumentum. To confirm authenticity, the samples were compared to reference collections in virtual herbaria.

Tea Preparation

The plant material was subjected to the infusion process for 10 min, according to the popular preparation method. The leaves of the boldo species were standardized at a concentration of 12 mg ml⁻¹, then the solutions were stored in vials, frozen, and lyophilized. The obtained extract was abbreviated AEBL - Aqueous Extract Boldo Leaves.

Mass Spectrometry Analysis

The stock solution was prepared by weighing 5 mg of the aqueous extract in a calibrated 5-ml volumetric flask using a calibrated analytical balance (\pm 0.1 mg) and MeOH as solvent, to obtain a final concentration of 1 mg ml⁻¹.

Then an intermediate solution was prepared to obtain the final solution with a concentration of 5 μ g ml⁻¹. These samples were analyzed by direct infusion in a mass spectrometer.

All mass spectra were acquired in monitoring mode (application Thermo LCQ Fleet Tune) using an LCQ Fleet ion-trap mass spectrometer (Thermo Frota LCQ, San Jose, CA, USA) with an electrospray ionization (ESI) interface and running in positive ion mode to perform ESI-MS and ESI-MSn analysis. Spectra were obtained from an average of at least 10 exams per spectrum. The samples were directly infused into the ion source through the syringe pump on the instrument (10 ml min⁻¹). The analytical conditions of MS were spray voltage, 5 kV; gas sheath, 10 arb; auxiliary gas, 5 arb; sweep gas, 0 arb; capillary temperature, 200; capillary

voltage, 40 V; tube lens, 115 V; mass range, m/z 200 to 400. Helium was used as a gas collision, and ESI-MSn spectra were obtained using collision energies ranging from 20 to 30%.

Antioxidant Activity

Antioxidant capacity analyzed by DPPH, ABTS, FRAP, and β -carotene/linoleic acid cooxidation methods

The tests were carried out at the Laboratory of Nanostructured Polymers (NANOPOL) at the UFAM. The samples were solubilized in MeOH P.A. at 1.0 mg ml⁻¹ and subjected to the following analyzes.

DPPH Assay

For this assay, 10 μ L of the sample was added to 190 μ L of the DPPH solution (100 μ M) and incubated in a dark environment for 30 min. Afterwards, absorbance readings were performed in a Microplate Reader at 515 nm (Epoch 2, Biotek). A Trolox standard curve from 100 to 1500 μ M ($y = -0.0004x + 0.7071$, $R^2 = 0.9931$) was built and the results were expressed in μ M of Trolox Equivalents.^[11,23]

ABTS Assay

For the ABTS assay, 2 μ L of the sample was added to 200 μ L of the ABTS⁺ solution (0.70 ABS) and incubated in a dark environment for 6 min. Then the absorbance reading was carried out in a Microplate Reader at 750 nm (Epoch 2, Biotek). A standard Trolox curve was made from 125 to 2000 μ M ($y = -0.0003x + 0.7351$, $R^2 = 0.9995$) and the results were expressed in μ M of Trolox Equivalents.^[23]

Iron reduction (FRAP)

The FRAP reagent was prepared beforehand by adding acetate buffer (0.3 mol L⁻¹, pH = 3.6), TPTZ solution (10 mM) and FeCl₃.6H₂O (20 mM), in the ratio 10:1:1. On a microplate, an aliquot of 9 μ L of the sample (500 μ g ml⁻¹) was mixed with 27 μ L of ultrapure water and added to 270 μ L of the FRAP reagent. The microplate was incubated for 30 min at 37°C and the absorbance measured at 595 nm using a microplate reader (Epoch 2, Biotek). The results were obtained by the regression equation of the curve obtained from the standard solution of ferrous sulfate (250 to 2000 μ M) and expressed in the Fe (II) [μ M Fe(II) g⁻¹] extract. The assay was performed in triplicate.^[23]

β -carotene/linoleic acid cooxidation

The Oxidation Inhibition assay of the β -carotene/linoleic acid system consists of the reaction mixture of linoleic acid, Tween 40, β -carotene, and water saturated with oxygen, with absorbance between 0.6 and 0.7 at 470 nm. First, 250 μ L of this reactive mixture and 10 μ L of samples (500 μ g ml⁻¹) were added to a microplate and incubated at 37°C, and then absorbance readings were performed in a Microplate Reader (Epoch 2, Biotek). Results were expressed as percentage of oxidation inhibition.^[24]

Determination of total phenols

For the assay of total phenols, the sample was added to a reaction mixture (1:1) of Folin Cicalteu reagent and sodium bicarbonate (6%), maintained in the dark for 90 min for further analysis in a Microplate Reader at 725 nm (Epoch 2, Biotek). The standard curve for gallic acid was made from 7.8 to 1000 μ g ml⁻¹ ($y = 0.0028x + 0.2847$, $R^2 = 0.9931$), and the results are expressed as milliequivalents of gallic acid per gram of sample.

Assessment of acetylcholinesterase inhibition *in vitro*

The enzymatic activity of acetylcholinesterase was determined on microplates using the Ellman spectrophotometric method.^[11] The enzyme

used was acetylcholinesterase (AChE) from *Electrophorus electricus* type VI (Sigma[®]), 5,5'-dithiobis (2-nitrobenzoic acid) (DTNB), iodinated acetylcholine substrate and eserine, Sigma standard. In each well of the microplates, 25 μ L of 15 mM iodinated acetylcholine, 125 μ L of DTNB, 50 μ L of 50 mM Tris-HCl buffer pH 8.0, and 25 μ L of the plant extracts at a concentration of 1 mg/ml were added. MeOH and tris-HCl were used as controls and eserine (10 μ M) as standard inhibitor. Absorbance was measured at 405 nm using a Thermoplate[®] microplate reader, model TP-reader, with 15-sec intervals. After these readings, 25 μ L of 0.22 U ml⁻¹ acetylcholinesterase were added to the wells. Absorbances were measured again at 405 nm, with 15-sec intervals. Increases in absorbance due to spontaneous substrate hydrolysis were corrected by subtracting the reaction rate before addition of the enzyme. Percentage of inhibition was calculated by comparing sample rates to controls. All assays were performed in triplicate. Enzymatic activity and percentage of AChE inhibition were calculated using the Microsoft[®] Excel program.

In silico evaluation against AChE enzyme by molecular docking

All structures of the major compounds identified in the samples were designed in ACD/Chemsketch and saved in sdf format. Furthermore, binder structures were previously verified by Marvin Srtch 15.4.207 for anchorage studies. Before the docking calculations, Autodock Tools 1.5.6 was used to prepare the protein and ligand structures according to the following steps: (1) definition of the grid for each target protein, considering each active site previously described by crystallographic studies; (2) verification for each torsional ligand structure and addition of polar hydrogens, Gaigster, and Kollman charges; and (3) saving all molecules pdbqt format.

Autodock Vina was used to calculate the affinity energies for each complex. After the structures were selected, to observe the results *in silico* by molecular docking, the Discovery Studio Visualizer program was used.

RESULTS AND DISCUSSION

Constituents identified in commercial *P. boldus* samples

The AEBL was analyzed in a direct infusion mass spectrometer in positive mode. The obtained spectra presented several similar peaks with [M+H]⁺ pair, but with different intensities. Peaks indicate masses that are compatible with alkaloids already reported for *P. boldus* as well as some alkaloids observed in studies about other species in the same family. The peaks at m/z 286, 300, 328, 330, 342, and 355 ([M+H]⁺) were detected in greater relative abundance (Figure S1 – Supporting information), the others were observed with low intensity.

Some of these peaks can be identified, such as the ion at m/z 286 ([M+H]⁺), which is compatible with the quasi-molecular ion ([M+H]⁺) of Coclaurine, an alkaloid already reported for this species.^[16,19] Its fragmentation allows detection of the daughter ion at m/z 269 with greater intensity, in addition to the ions at m/z 237, 178, 175, 145, 143, 137, and 107, which are compatible with fragmentation data for this substance.

The ion observed at m/z 300 ([M+H]⁺) is compatible with the quasi-molecular ion ([M+H]⁺) of *N*-methylcoclaurine, an alkaloid already reported in *P. boldus*.^[25] Fragmentation revealed daughter ions at m/z 269, 237, 192, 175, 145, 143, 137, and 107, which were compatible with the fragmentation data of *N*-methylcoclaurine.

Another ion observed was m/z 328 ([M+H]⁺), compatible with the quasi molecular ion ([M+H]⁺) of boldine, an aporphinic alkaloid reported in the species.^[26] When fragmented, daughter ions were detected at m/z 297

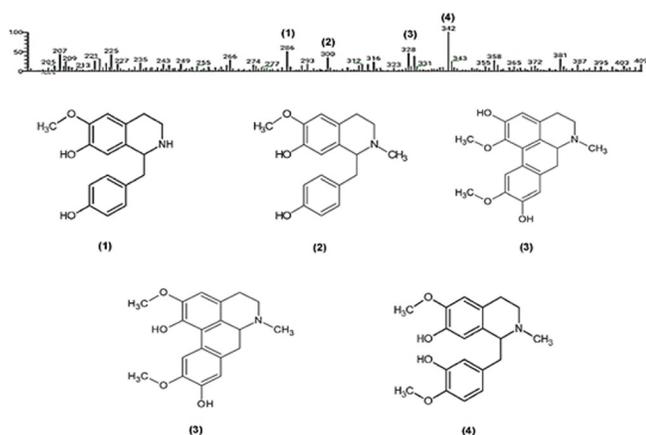


Figure S1: Total ion chromatogram of the AEBL sample and the main identified constituents: 1 - Coclaurine; 2 - *N*-methylcoclaurine; 3 - Boldine and Isoboldine; 4 - *N*-methyllaurotetanine.

and 268, in agreement with the literature data on the fragmentation of this alkaloid.

The mass analysis of the m/z 328 ion, when subjected to fragmentation into MS^n , exhibited an initial loss of 31 Da, and sequential losses of 32 Da and 28 Da, suggesting that it is an aporphine alkaloid containing an *N*-methyl and a methoxyl adjacent to a hydroxyl in the A ring. The alkaloids suggested for this spectrum are boldine and isoboldine, as they have the same ion mass of m/z 328.^[16,17,19,27]

The aporphine alkaloid laurotetanine was identified by fragmentation of the ion m/z 328 and through the formation of ions at m/z 311, 279, 264, and 236. The initial loss of 17 Da was attributed to the NH_3 group, which indicates that nitrogen present in the molecule is linked to a hydrogen and not to the methyl group.^[17,28]

The m/z 330 ($[M+H]^+$) ion is compatible with the reticuline alkaloid. When fragmented, the daughter ions were generated at m/z 299, 267, 192, 175, and 137, which are values noted in the literature.^[25]

The structure of the m/z 342 alkaloid was confirmed by comparing existing data in the literature.^[17,28] The m/z 342 ion, when submitted to fragmentation analysis, evidenced sequential losses of 31, 15, and 31 Da, suggesting an aporphinic structure containing an adjacent *N*-methyl and methoxyl in the A ring; such fragments are compatible with the *N*-methyllaurotetanine alkaloid.

The m/z 356 ion ($[M+H]^+$), which is compatible with the quasi-molecular ion ($[M+H]^+$) of glaucine, in its fragmentation, could detect the daughter ions of m/z 325, 294, 279, 251, and 236 (Table 1).

Evaluation of the mass spectra identified the presence of three phenolic compounds in the sample, Figure 1. The ion at m/z 727, when subjected to fragmentation into MS^n , showed an initial loss of 132 Da, corresponding to dehydrated pentose, and sequential losses of 162 Da and 146 Da, suggesting the presence of hexose and rhamnose, respectively. Such fragments, in comparison with literature data, are compatible with the flavonoid Luteolin-pentosyl-glucosylrhamnose. The ion at m/z 741 is compatible with the flavonoid kaempferol - 3-O-glycosyl-rhamnosyl-rhamnose, when sequential losses of 132 Da are observed, referring to one dehydrated pentose unit and two 146 Da, pertaining to losses of two rhamnose units.^[17]

The compound identified by the ion at m/z 633 showed a typical fragmentation pattern of chlorogenic acids. Through the loss of 341 Da,

Table 1: Alkaloids identified in the AEBL extract.

Precursor ion m/z	Fragmented ions m/z	Identification	MF	Refs.
286	269, 237, 178, 175, 145, 143, 137, and 107	Coclaurine	$C_{17}H_{19}NO_3$	[16,19]
300	269, 237, 192, 175, 145, 143, 137, and 107	<i>N</i> -methylcoclaurine	$C_{18}H_{21}NO_3$	[25]
314	297, 265	Lauroitsine	$C_{18}H_{19}NO_4$	[17]
316	299, 178	Norreticuline	$C_{18}H_{21}O_4$	[17]
328	297, 265, 237, and 205	Boldine	$C_{19}H_{21}NO_4$	[25,26]
328	297, 265, 237, 165	Isoboldine	$C_{19}H_{21}NO_4$	[16,27]
328	311, 279, 264, 236	Laurotetanine	$C_{19}H_{21}NO_4$	[17,28]
330	299, 267, 192, 175, and 137	Reticuline	$C_{20}H_{23}NO_4$	[17,25]
342	311, 296, 280, 265, 253, 237	<i>N</i> -methylaurotetanine	$C_{20}H_{23}NO_4$	[17,28]
342	296, 279, 264, 248	Isocoridine	$C_{20}H_{23}NO_4$	[17]
356	325, 294, 279, 251, and 236	Glaucine	$C_{21}H_{25}NO_4$	[25,28]

MF = Molecular Formula

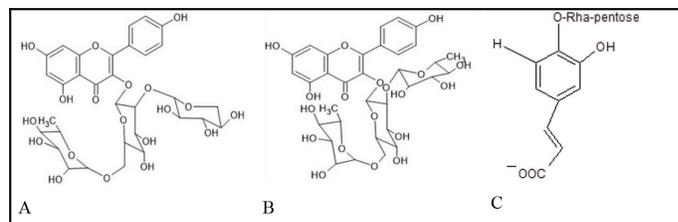


Figure 1: Phenolics identified in the AEBL extract: A- Luteolin-pentosyl-glycosyl-rhamnose; B - Kaempferol-3-O-glycosyl-rhamnosyl-rhamnose; and C - Caffeoylquinic phenolic acid glycoside.

the cinnamoyl glycoside part was identified and, through the loss of 323 Da, the caffeoyl glycoside unit was identified. These fragments suggest the glycoside caffeoylquinic phenolic acid.^[29]

Antioxidant activity and quantification of total phenolic compounds

Table 2 presents the results of the antioxidant activity of the AEBL extract evaluated by the DPPH, ABTS, FRAP methods and by the beta-carotene/linoleic acid system, in addition to the quantification results of the total phenolics.

DPPH is sequestered by phenolic compounds through two mechanisms: 1) possible donation of hydrogen to stabilize the radical compound; 2) donation of electrons from the phenoxide anion to DPPH. For these mechanisms, phenolics can participate together with flavonoids as an antioxidant by presenting free phenolic hydroxyl radicals and by the bioavailability of conjugation.^[23,30] However, this high free radical scavenging activity observed in the *P. bordo* extract is not just related

Table 2: Results of the antioxidant activity of the aqueous extract of *Peumus boldus*.

Sample	DPPH ($\mu\text{M ET}$)	ABTS ($\mu\text{M ET}$)	FRAP ($\mu\text{mol de Fe}$ (II) g^{-1}) ¹	Inhibition (%)	FT (mg EAG g^{-1})
AEBL	1513.6 \pm 1.2	1928.1 \pm 5.7	1512.7 \pm 4.2	81.2 \pm 1.3	82.4 \pm 0.9

¹ μmol of Fe (II) g^{-1} : power of reduction in μmol of Fe (II) per gram of sample; 2 mEAG g^{-1} : equivalent to gallic acid per gram of sample; ³ $\mu\text{M ET}$: antioxidant capacity equivalent to trolox.

to the presence of phenolic compounds because the extract exhibited a low concentration of phenolics (82.804 \pm 0.9 mg EAG g^{-1}). These results indicate that the observed oxidizing activity may be related to the presence of alkaloids. Previous studies have evidenced the antioxidant potential of alkaloids. Giordani *et al.*^[15] proved by the DPPH method the antioxidant activity of the alkaloids lycorine and boldine with IC_{50} values of 0.326 mM and 0.0552 mM, respectively. Other compounds, such as helindicine and lycopsamine, both alkaloids, containing a laconic ring in their structure,^[31] erisodine, hydroxyerisodine, melosmine, and O-Methyl-moschatoline, with the presence of the oxaporphine nucleus, have an expressive capacity to capture the DPPH radical.^[32] Capturing the activity of reactive oxygen species (ROS) in alkaloid groups may be related to the exceptional redox property conferred to nitrogen,^[16,17,20] present in the structures of these secondary metabolites.

Treatment of AD through antioxidant therapies can prevent or at least attenuate organic deterioration due to excessive oxidative stress. Significant evidence indicates oxidative stress in the progression of AD, leading to progressive and significant loss of specific neuronal cells, contributing to the pathogenesis of the disease.^[4] Thus, the discovery of constituents with antioxidant properties is very beneficial to delay the progression of the disease.

In the ABTS method, the extract also presented expressive antioxidant activity (1928.1 \pm 5.7 $\mu\text{M ET}$), which is related to the presence of alkaloids that act as good electron donors.

The methodology using β -carotene autooxidation measures the capacity of the extract to inhibit β -carotene oxidation against linoleic acid. The AEBL presented antioxidant activity with a percentage of 81.2 \pm 1.3%. This activity is directly related to the total phenolic content in the extract. The extract managed to reduce a significant amount of Fe^{3+} in Fe^{2+} (1512.7 μMFe^{2+} mg^{-1}). The antioxidant activity observed is due to the presence of phenolics and was complimented by the presence of alkaloid. These molecules can donate hydrogen radicals to pair with other available radicals, impeding oxidation and stabilizing the system.^[33]

Assessment of acetylcholinesterase inhibition *in vitro*

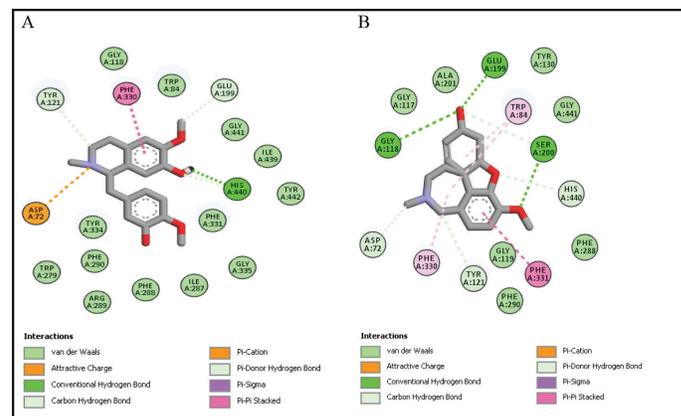
The AEBL did not exhibit inhibitory activity to acetylcholinesterase at the evaluated concentration of 1 mg/ml. For the AEBL, the literature reports a potent inhibitory action of aqueous extracts of *P. boldus* leaves at a concentration greater than 1 mg ml^{-1} ,^[18,20] suggesting that the *in vitro* test for the AEBL should be repeated at a higher concentration.

Molecular Docking

The results for the dockings of isoboldine, boldine, coclaurine, leurotetanine, *N*-methylleurotetanine, reticuline, and *N*-methylcoclaurine in AChE were practically located in the catalytic cavity (Trp84). The Trp84 residue is located next to the triad (Ser200 – His440 – Glu327) and is considered the main residue located at the end of the “aromatic throat”.^[13,14] Trp84 has been used in many other molecular docking

Table 3: Energy results of the alkaloids identified in the aqueous extract of *P. boldus* leaves.

Compounds	Binding energy (kcal/mol)
<i>N</i> -methylleurotetanine	-10.1
Isoboldine	-10.0
Boldine	-9.9
Lauroilsine	-9.7
<i>N</i> -methylcoclaurine	-9.6
Coclaurine	-9.4
Reticuline	-9.1

**Figure S2:** Result of molecular docking using the Autodock program for a) reticuline; b) galantamine.

studies as an important marker of successful enzyme inhibition. The isoquinoline alkaloid reticuline was the only compound that showed weaker interaction with the Trp84 residue, as this type of ligand recognition may justify a higher value of free binding energy in the enzyme. The higher the energy value, the lower the binding affinity and the higher the drug concentration required to inhibit enzyme activity^[11,34] Although reticuline does not present a docking energy value as satisfactory as the other alkaloids identified in the *P. boldus* sample (Table 3), it exhibits an inhibitory capacity due to the other electrostatic interactions observed with the amino acids of the enzyme.

The inhibitory capacity of reticuline is justified by Van der Waals interactions with Gly 118, Trp84, Try 334, Phe 290, and Phe 288 residues that belong to the catalytic triad, oxyanion cavity, peripheral anionic site, and acyl binding site as well as electrostatic interactions, for example the π -cation-type interaction with the Asp 72 residue pertaining to the peripheral anionic site (PAS), hydrogen bonding with the His 440 residue of the catalytic site and the hydroxyl group of the molecule, and the π - π type-stacking interaction with Phe 330 from the catalytic triad (Figure 2 and S2a).

The analysis of the AChE complex with galantamine, the reference molecule in this study, found several interactions in the active site of the enzyme, justifying its ability to stabilize the free energy of the drug-receptor complex (-9.6 kcal mol^{-1}) and the inhibitory potential of this substance. The interactions found include π - π stacking-type interaction with Trp 84 and Phe 330 residues of the catalytic triad, hydrogen bonding with residues Gly 118, Glu 199, and Ser 200 from the catalytic site, and the oxyanion cavity of the enzyme with the hydroxyl and methoxyl groups of galanthamine, π -sigma-type interaction with the Phe 331 residue with the aromatic ring of the molecule, Van der Waals interaction

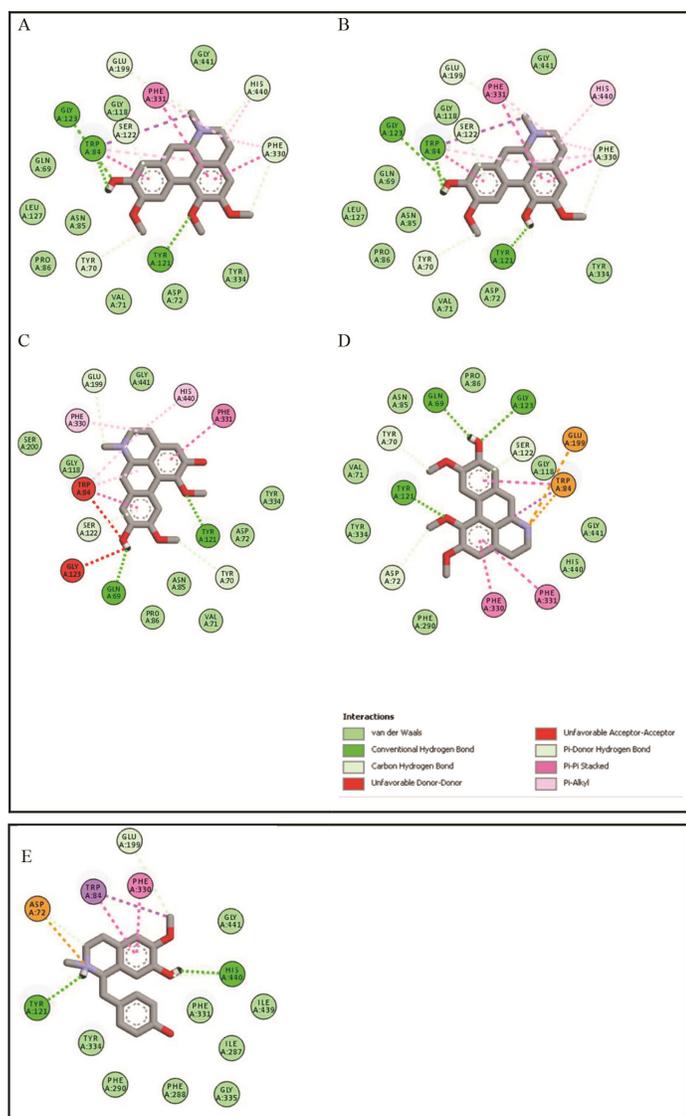


Figure S3: Molecular docking result for: A - *N*-methylaurotetanine; B – isoboldine; C-boldine, D – laurotetanine, and E - *N*-methylcoclaurine, using the Autodock program.

with the anionic subsite (oxygen channel) with the Ala 201 residue, and with the catalytic acylation site (CAS) formed by Phe 290 and Phe 288 (Figure S2b).

As mentioned above, the lower the molecular docking energy, the stronger the enzyme inhibition process for the evaluated substance. *N*-methylaurotetanine exhibited the best value, followed by isoboldine, boldine, laurotetanine, and *N*-methylcoclaurine, the latter displaying the same value as galantamine (an important enzyme inhibitor).

N-methylaurotetanine and isoboldine have a hydrogen bond interaction (stronger interaction) of the Trp 84 residue, which belongs to the catalytic triad, with the hydroxyl group of these alkaloids. This type of interaction is not observed in the other substances, justifying its higher energy values (Figures 3 and S3a-e). Hydrogen bonding was also observed at the Try 121 residue of the peripheral anionic site.

Boldine did not have any favorable interaction for the formation of the hydrogen bond of the hydroxyl group with the Trp 84 and Gly 123 residues; however, this type of bond was observed with the Try 121 residue and the methoxyl group. We verified π -alkyl-type interactions,

with the residue His 440 of the catalytic site and with the residue Phe 330 of the catalytic triad, and π - π stacking of the residue Phe 331, belonging to the oxyanion cavity, with the aromatic ring of the molecule.

Laurotetanine and *N*-methylcoclaurine exhibited similar interactions. The main differences observed include the presence of electrostatic interaction of the π -cation type between the residues Glu 199 and Trp 84 with the nitrogen of the aporphine alkaloid laurotetanine, and the presence of hydrogen bonding between the residue His 440 and the hydroxyl group of the alkaloid *N*-methylcoclaurine, as well as between the Try 121 residue and the proton.

CONCLUSION

The aqueous extract of *Peumus boldus* leaves exhibited important secondary metabolites, such as flavonoids, phenolic compounds, and alkaloids. These constituents were responsible for the antioxidant activity observed in the DPPH, ABTS, and FRAP methods and beta-carotene/linoleic acid system. Such results suggest that the studied sample can protect the neurons from the oxidative processes generated by the pathological processes of Alzheimer's disease. These activities correlate the promising results of the AChE enzyme inhibition. Despite the negative result of the extract for its *in vitro* inhibitory potential of the enzyme, the alkaloids identified in the sample were considered active for *in silico* anticholinesterase activity. The substances isoboldine, boldine, coclaurine, laurotetanine, *N*-methylaurotetanine, reticuline, and *N*-methylcoclaurine were able to inhibit the enzyme, with the alkaloid *N*-methylaurotetanine achieving the best inhibition. These data allow us to conclude that the aqueous extract of boldo leaves could be active at a concentration higher than 1 mg ml⁻¹ and future studies are needed to fractionate and purify the fractions to isolate the molecules responsible for the observed inhibition.

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CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

ABBREVIATIONS

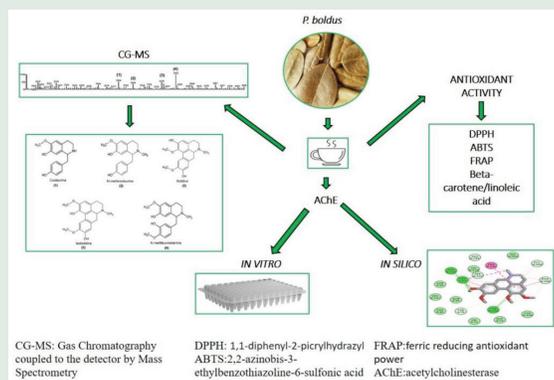
AD: Alzheimer's disease; **DPPH:** (2,2-diphenyl-1-picrylhydrazyl); **ABTS:** [2,2'-azino-bis(3-ethylbenzothiazoline)-6-sulfonic acid]; **FRAP:** Ferric Reducing Antioxidant Power; **ESI:** Electrospray Ionization; **AChE:** Acetylcholinesterase; **Ach:** Acetylcholine; **AEBl:** Aqueous Extract Boldo Leaves; **ROS:** Reactive Oxygen Species.

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GRAPHICAL ABSTRACT



SUMMARY

Medicinal plants have been traditionally used to treat all kinds of infirmities. Known in Brazil as boldo-do-chile, the species *Peumus boldus* Molina (Monimiaceae) is widely employed to relieve bowel discomfort. Increased gastrointestinal motility is associated to the action of the neurotransmitter acetylcholine (ACh), responsible for transmitting nerve impulses from one neuron to another for this purpose. To avoid excessive transmission of nerve impulses and its deleterious effects on biological functioning, our body produces an enzyme (acetylcholinesterase or AChE) that hydrolyzes ACh into choline (Ch), which is then reabsorbed by the first neuron to be reconverted to ACh during transmission of nerve impulses. AChE is involved in Alzheimer's Disease (AD) — a medical condition currently affecting about 35.6 million people worldwide — in that the latter causes degradation of neurons, thus decreasing the action of ACh. Although there is no cure to AD, it can be effectively treated by means of AChE-inhibiting drugs, which prevent ACh degradation. The traditional use of *P. boldus* suggests that the decoction of its leaves contains substances capable of inhibiting AChE, thereby promoting intestinal motility as more acetylcholine is available for inter-neuron communication. This study comprises the procedures for the preparation of the aqueous extract of *P. boldus* leaves (EAFPb), identification of its constituents by means of electrospray ionization mass spectrometry (ESI-MS), assessment of antioxidant activity (DPPH, ABTS, FRAP, and oxidation inhibition of the β -carotene/linoleic acid system), quantification of total phenols, and *in silico* determination of AChE inhibitory activity through molecular docking assays. Mass spectrometry identified the presence of the alkaloids isoboldine, boldine, clocaurin, laurotetanin, *N*-methyllaurotetanin, reticulon, and *N*-methylclocaurine in EAFPb. The *in silico* prediction assay indicated that the alkaloid *N*-methyllaurotetanin constitutes the best acetylcholinesterase enzyme inhibition ligand, with energy of -10.1 kcal/mol. The extract was shown to be potentially active in all analyses for antioxidant activity. The results obtained in this study suggests that EAFPb contains chemical constituents with potential for treating AD.

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