Expedient Microwave-Assisted Synthesis of Bis(\textit{n})-lophine Analogues as Selective Butyrylcholinesterase Inhibitors: Cytotoxicity Evaluation and Molecular Modelling

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In the brain of patients with chronic Alzheimer’s disease (AD), the butyrylcholinesterase (BuChE) levels rise while the acetylcholinesterase (AChE) levels decrease. Therefore, development of new selective BuChE inhibitors is of vital importance. Here we present a series of bis(\textit{n})-lophine analogues, where two lophine derivatives are connected by a methylene chain. The bis(\textit{n})-lophine analogues were synthesized through one-pot four component reaction between pyridinecarboxaldehydes, \textit{n}-alkanediamines, benzil, and ammonium acetate. The reactions were performed in a microwave reactor in one step for symmetrical bis(\textit{n})-lophines, and in two steps for unsymmetrical bis(\textit{n})-lophines. The compounds are strongly selective to BuChE, since none of them inhibit AChE. All the compounds, except 7a, 7b and 7c, displayed potent inhibitory activity against BuChE at a micromolar and sub-micromolar range (half maximal inhibitory concentration (IC\textsubscript{50}) 32.25-0.03 µM). The enzyme kinetic and docking studies suggests that the inhibitor act as a dual binding site inhibitor, binding into the bottom of the gorge and in the peripheral anionic site (PAS) of BuChE cavity. Furthermore, \textit{in vitro} studies showed that compounds 5b and 12b had no cytotoxic effects in kidney Vero, hepatic HepG2 and C6 astroglial cell lines.

Keywords: cholinesterase inhibitor, butyrylcholinesterase, bis(\textit{n})-lophine analogues

Introduction

The serine hydrolases acetylcholinesterase (AChE) and butyrylcholinesterase (BuChE) are structurally related enzymes that co-regulate the metabolism of the neurotransmitter acetylcholine (ACh).

Cholinesterase inhibitors (ChEIs) are a class of drugs that have been used in the management of various human ailments, including Alzheimer’s disease (AD),\textsuperscript{1} Parkinson’s disease,\textsuperscript{2} glaucoma,\textsuperscript{3,4} myasthenia gravis,\textsuperscript{5} Lewy bodies’ disease,\textsuperscript{6} and chronic pain in elderly.\textsuperscript{7}

Alzheimer’s disease is known as a neurodegenerative disorder with major importance and the principal cause of dementia among the elderly. There is no cure for AD, but there are drugs that target the symptoms in order to improve the cognitive function of the patient.\textsuperscript{1,8,9}

Four drugs are currently available for AD treatment and all were approved more than a decade ago. Of these, the first-line agents are the ChEIs (donepezil, rivastigmine, and galantamine).\textsuperscript{1} Tacrine (1,2,3,4-tetrahydroacridine, THA), was the first ChEI approved by the U.S. Food and Drug Administration (FDA) in 1993 for the palliative treatment of AD.\textsuperscript{5} However, it soon exhibited hepatotoxicity and consequently was withdrawn from the market shortly after its approval.\textsuperscript{10}
Although the cause of AD is still not fully understood, dysregulation of the amyloid-beta (Aβ) protein level and neurofibrillary tangles appear to be the predominant contributing factors. Paradoxically, previous clinical failures of anti-Aβ antigens and γ-secretase inhibitors, and the recent clinical failures of β-secretase (BACE1) inhibitors and monoclonal anti-Aβ antibodies, have led researchers to suggest dropping such therapies proposals.11

Although the role of AChE in cholinergic transmission is well known, the role of BuChE has not been sufficiently elucidated and an increasing number of studies3 have provided evidence suggesting that BuChE may play a distinct role in AD patients. Depending on the AD stage, there is a decline in AChE levels in the brain and a progressive increase of BuChE levels, which becomes responsible for acetylcholine hydrolysis. In the AD patients’ brain, AChE levels are decreased by around 50%, whereas BuChE levels increase by as much as 900%.12,13 BuChE is associated with insulin resistance, a typical feature of patients with type II diabetes.14 According to a recent mouse study,15 BuChE regulates ghrelin levels to control social behavior, such as aggression. Thus, specific BuChE inhibitors may be considered as a new and promising therapeutic strategy for neurodegenerative diseases and others related to the cholinergic system.13,15

Different structural features of AChE and BuChE are related to their substrate specificity: AChE has higher selectivity for small molecules like ACh, while BuChE is less substrate-specific, accommodating the metabolism of several different molecules, including various neuroactive peptides.16 Accordingly, the structural characteristic of BuChE provides a reasonable thought to design selective BuChE inhibitors.17 Furthermore, BuChE is associated with other pathological manifestations of AD, including the formation of Aβ from the initially benign form seen in normal aging to malignant fibrillar Aβ deposition. Thus, BuChE’s inhibitors may serve the dual purpose of increasing acetylcholine levels and inhibiting fibrillar Aβ deposition.13,18-20

The active site gorge of BuChE is larger than that of AChE (500 Å against 300 Å)1 and forms a bowl rather than a deep narrow gorge.21 Also, it has been shown that the gorge of BuChE contains about 40% fewer aromatic residues than AChE, where they are substituted by smaller aliphatic or even polar residues. This accounts for most of the specific properties of BuChE. The acylation site, where catalytic reactions take place, is at the bottom of the gorge, about 20 Å from the protein surface. It consists of the catalytic triad Ser198, His438, and Glu325. The triad of cholinesterase-related enzymes differs from that of serine proteases by having glutamate instead of aspartate. One of the most important features of the BuChE and AChE catalytic process is the H-bond stabilization of the transition state by the oxyanion hole. The oxyanion hole is composed of three highly conserved N–H dipoles from the main chain of residues Gly116, Gly117 and Ala119.22

The first designed hybrid molecule for the potential treatment of AD dates back to 1996 when Pang et al.23 reported the synthesis of alkylene-linked bis-tacrine compounds as a dual binding site inhibitor (DBS, namely compounds that are able to occupy the two binding sites). They envisaged that tacrine dimer targets the high affinity catalytic anionic site (CAS) and the low-affinity peripheral binding site (PAS) of AChE. These simple bis-tacrine dimer exhibited potency in orders 1,000 folds higher, compared to tacrine.

The literature reveals a plethora of studies of ChEIs based on DBS. A newly arising strategy is the synthesis of hybrids molecules, where two pharmacological molecules are mixed in one single molecule. These hybrids act as multi-target compounds, usually combining its potent cholinesterase inhibition with other pharmacological properties.24-29 However, the number of hybrids obtained from the combination of scaffolds different from those well-known ChEI drugs are quite restricted, and ChEIs with novel structural diversity are urgently needed.

The imidazolic nucleus is incorporated in many bioactive molecules playing a vital role in treating various types of diseases.30,31 The imidazole derivative 2,4,5-triphenyl-1H-imidazole, also known as lophine, can be used as fluorescent labeling reagents for amines, phenols, and carboxylic acids.32 The synthesis of various imidazoles and their derivatives are important targets in current years, among them 2,4,5-tri- and 1,2,4,5-tetrasubstituted imidazoles have received much attention.33-35 Also, hybrids containing lophine and pyrimidine nuclei connected by a methylene chain showed photophysical features that were successfully used to explore their interaction with bovine serum albumin (BSA) protein and exhibited significant suppression mechanism.36 In previous work,37 our group synthesized a series of bis(n)-lophine, where two lophine core were linked by a methylene chain. In this series, the bis(8)-lophine was the only active compound, showing a potent inhibition against AChE (half maximal inhibitory concentration \( IC_{50} = 42.55 \) nM).

Among several methods reported in the literature for imidazoles synthesis, multicomponent reactions (MCRs) have received considerable attention and proves to be one of the most efficient methods for obtaining substituted imidazoles. In an MCR, a product is assembled according to a cascade of elementary chemical reactions. Thus, there is a network of reaction equilibria, which finally flow into an irreversible step yielding the product. The formation of a
particular product is dependent on the reaction conditions: solvent, temperature, catalyst, concentration, the type of starting materials, and functional groups.

As part of our search for new ChEIs drug candidates, we report herein a new one-pot synthesis of bis(n)-lophine analogues as selective dual binding inhibitors of BuChE along with their cytotoxicity evaluation and molecular modeling studies.

**Experimental**

**Materials**

Microwave-assisted reactions were carried out using a MARS6 microwave oven (CEM Company, Charlotte, NC, USA) and the MARSXPress vessels with 10 mL of capacity. The temperature program used consisted in a heating ramp of 15 min to reach 110 °C, followed by 1 h of reaction, with magnetic stirring. All melting points were determined in open glass capillaries using a Büchi M-565 (Essen, Germany) apparatus. Infrared (IR) spectra were recorded on a Varian 640-IR (Palo Alto, USA) spectrometer in KBr disks. 1 H nuclear magnetic resonance (NMR) and 13 C NMR spectra were recorded in CDCl3 (400 MHz, Varian) with hexane-d4 as internal standard in 1 H NMR or from the solvent peak of CDCl3 solution on a Bruker BioSpin 400 MHz (Billerica, USA) spectrometer. The relaxation time in both analyses was 1.0 s and the chemical shifts (δ) are given in part per million from the peak of tetramethylsilane (δ 0.00 ppm) as internal standard in 1 H NMR or from the solvent peak of CDCl3 (δ 77.16 ppm) in 13 C NMR APT (attached proton test). Multiplicities are given as s (singlet), d (doublet), dd (doublet of doublets), dt (doublet of triplets), t (triplet), td (triplet of doublets), m (multiplet); coupling constants (J) are given in Hz. High resolution mass spectrometry with electrospray ionization (HRMS-ESI) data on the positive mode was collected on a Micromass Q-Tof instrument from Waters (Manchester, UK). Samples were infused from a 100 mL Hamilton syringe at flow rate range from 5 to 10 mL min⁻¹, depending on the sample. The instrument settings were the following: capillary voltage 3000 V, cone voltage 33 V, extraction cone voltage 2.5 V, desolvation gas temperature 100 °C. Nitrogen was used as the desolvation gas. Methanol (high performance liquid chromatography (HPLC) grade, Tedia, Fairfield, OH, USA) was used as solvent for the analyzed samples and filtered prior to injection. Purification by column chromatography was carried out on silica gel 60 Å (70-230 mesh, Sigma-Aldrich, St. Louis, USA). Analytical thin layer chromatography (TLC) was conducted on aluminum plates with 0.2 mm of silica gel 60F-254 (Merck, Darmstadt, Germany). Dimethyl sulfoxide (DMSO), 5,5'-dithiobis-2-nitrobenzoic acid (DTNB), acetylthiocholine iodide, heparin, dipotassium phosphate dibasic (K2HPO4) and potassium phosphate monobasic (KH2PO4) were obtained from Sigma-Aldrich (St. Louis, MO, USA). Solvents were obtained from Tedia (Fairfield, OH, USA) and Nuclear (Diadema, SP, Brazil), and reagents were purchased from Sigma-Aldrich, Acros Organics (Thermo Fisher Scientific, Waltham, Massachusetts, USA) and TCI (Tokyo, Japan).

**Synthesis**

General procedure for the synthesis of bis(n)-lophine homodimers (5-7)

A mixture of 1, n-alkanediamine (0.45 mmol), ammonium acetate (0.90 mmol), benzil (0.90 mmol) and n-pyridinecarboxaldehyde (0.90 mmol) in absolute ethanol (3.5 mL) was added to a vial compatible with the use on the microwave reactor and the temperature program was started. This program was repeated three times in order to complete 4 h of reaction. At the end of the reaction time, the solvent was removed under reduced pressure. The crude product was purified by column chromatography (eluting with hexane:ethyl acetate, 90:10 with gradient elution until 0:100) to give the desired product as a solid.

1,6-Bis(4,5-diphenyl-2-(pyridin-2-yl)-1H-imidazol-1-yl)hexane (5a)

Yellow solid; 39% yield; mp 75-76 °C; IR (KBr) νmax / cm⁻¹ 3052, 2935, 2849, 1582, 1458, 788, 703; 1 H NMR (400 MHz, CDCl3) δ 8.55-8.50 (m, 2H), 8.41 (d, J 8.0 Hz, 2H), 8.71 (td, J 8.0, 1.8 Hz, 2H), 7.58-7.50 (m, 4H), 7.48-7.43 (m, 6H), 7.39-7.33 (m, 4H), 7.27-7.21 (m, 6H), 7.20-7.14 (m, 2H), 4.36 (t, J 7.6 Hz, 4H), 1.50-1.35 (m, 4H), 0.98-0.87 (m, 4H); 13 C NMR APT (100 MHz, CDCl3) δ 150.7, 148.3, 144.0, 137.7, 136.6, 134.0, 130.9, 130.8, 130.2, 129.2, 128.9, 128.2, 127.5, 126.7, 126.6, 123.6, 44.5, 30.6, 25.6; HRMS-ESI m/z, calcd. for [M + H]+: 677.3387, found: 677.3386.

1,6-Bis(4,5-diphenyl-2-(pyridin-3-yl)-1H-imidazol-1-yl)hexane (5b)

White solid; 37% yield; mp 194-195 °C; IR (KBr) νmax / cm⁻¹ 3043, 2927, 2857, 1473, 773, 696; 1 H NMR (400 MHz, CDCl3) δ 8.91-8.85 (m, 2H), 8.69 (dd, J 4.8, 1.7 Hz, 2H), 8.08 (dt, J 7.9, 1.7 Hz, 2H), 7.54-7.49 (m, 4H), 7.49-7.42 (m, 8H), 7.37-7.32 (m, 4H), 7.26-7.14 (m, 6H), 3.79 (t, J 7.6 Hz, 4H), 1.22-1.11 (m, 4H), 0.97-0.87 (m, 4H); 13 C NMR APT (100 MHz, CDCl3) δ 149.8, 149.3, 144.3, 138.4, 136.7, 134.0, 130.9, 130.8, 130.2, 129.2, 128.9, 128.2, 127.5, 126.7, 126.6, 123.6, 44.5, 30.1, 25.3; HRMS-ESI m/z, calcd. for [M + H]+: 677.3387, found: 677.3381.
1,6-Bis(4,5-diphenyl-2-(pyridin-4-yl)-1H-imidazol-1-yl)hexane (5c)

White solid; 33% yield; mp 200-201 °C; IR (KBr) νmax / cm⁻¹: 3043, 2927, 2857, 1597, 835, 765, 703; ¹H NMR (400 MHz, CDCl₃) δ 8.76-8.69 (m, 4H), 7.63-7.58 (m, 4H), 7.53-7.43 (m, 10H), 7.38-7.32 (m, 4H), 7.27-7.15 (m, 6H), 3.84 (t, J 7.6 Hz, 4H), 1.24-1.12 (m, 4H), 0.74-0.64 (m, 4H); ¹³C NMR APT (100 MHz, CDCl₃) δ 150.3, 144.5, 138.8, 133.9, 131.0, 130.7, 127.9, 129.0, 128.1, 126.7, 126.7, 44.6, 30.1, 25.3; HRMS-ESI m/z, calcd. for [M + H]+: 677.3387, found: 677.3382.

1,7-Bis(4,5-diphenyl-2-(pyridin-2-yl)-1H-imidazol-1-yl)heptane (6a)

Yellow solid; 45% yield; mp 149–150 °C; IR (KBr) νmax / cm⁻¹: 3055, 2946, 2860, 1586, 1469, 693; ¹H NMR (400 MHz, CDCl₃) δ 8.55-8.49 (m, 2H), 8.35-8.29 (m, 2H), 7.76 (td, J 7.8, 1.8 Hz, 2H), 7.53-7.47 (m, 4H), 7.46-7.32 (m, 11H), 7.23-7.16 (m, 6H), 7.16-7.11 (m, 2H), 4.37 (t, J 7.6 Hz, 4H), 1.51-1.35 (m, 4H), 1.00-0.81 (m, 6H); ¹³C NMR APT (75 MHz, CDCl₃) δ 151.1, 148.2, 144.1, 137.9, 136.5, 134.6, 131.7, 131.3, 131.2, 128.9, 128.7, 128.1, 126.7, 126.3, 123.4, 122.4, 45.2, 30.7, 28.0, 26.1; HRMS-ESI m/z, calcd. for [M + H]+: 691.5344, found: 691.5343.

1,7-Bis(4,5-diphenyl-2-(pyridin-3-yl)-1H-imidazol-1-yl)heptane (6b)

Yellow solid; 39% yield; mp 76-77 °C; IR (KBr) νmax / cm⁻¹: 3070, 2931, 2853, 1484, 1026, 770; ¹H NMR (400 MHz, CDCl₃) δ 8.97-8.84 (m, 2H), 8.72-8.60 (m, 2H), 8.04 (dt, J 7.9, 1.8 Hz, 2H), 7.55-7.31 (m, 16H), 7.24-7.11 (m, 6H), 3.80 (t, J 7.6 Hz, 4H), 1.27-1.14 (m, 4H), 0.77-0.59 (m, 6H); ¹³C NMR APT (100 MHz, CDCl₃) δ 149.9, 149.5, 144.5, 138.8, 136.8, 134.2, 131.2 (6), 131.2 (1), 130.4, 129.3, 129.0, 128.3, 127.9, 126.9, 126.8, 123.7, 44.8, 30.5, 27.8, 26.0; HRMS-ESI m/z, calcd. for [M + H]+: 691.5344, found: 691.5343.

1,7-Bis(4,5-diphenyl-2-(pyridin-4-yl)-1H-imidazol-1-yl)heptane (6c)

White solid; 31% yield; mp 164-165 °C; IR (KBr) νmax / cm⁻¹: 3055, 2922, 2853, 1601, 1415, 972; ¹H NMR (400 MHz, CDCl₃) δ 8.76-8.67 (m, 1H), 7.65-7.60 (m, 1H), 7.51-7.41 (m, 3H), 7.39-7.32 (m, 1H), 7.24-7.12 (m, 2H), 3.88 (t, J 7.6 Hz, 1H), 1.29-1.18 (m, 1H), 0.81-0.63 (m, 1H); ¹³C NMR APT (75 MHz, CDCl₃) δ 150.4, 144.6, 139.0, 134.1, 131.2, 131.1, 129.4, 129.2, 128.3, 126.9, 126.8, 44.9, 30.4, 27.9, 26.0; HRMS-ESI m/z, calcd. for [M + H]+: 691.5344, found: 691.5344.
The organic phase was dried with anhydrous Na₂SO₄ and the solvent was removed under reduced pressure. The crude product was purified by column chromatography (eluting with chloroform-methanol-ammonium hydroxide, 98:1:5:0.5 with gradient elution until 80:19:5:0.5) to give the desired product as a yellow oil (47% yield).

General procedure for the synthesis of lophine heterodimers (10-12)

A mixture of N-alkylaminolophine (0.45 mmol), ammonium acetate (0.45 mmol), benzil (0.45 mmol) and n-pyridinecarboxaldehyde (0.45 mmol) in absolute ethanol (3.5 mL) was added to a vial compatible with the use on the microwave reactor and the temperature program was started. This program was repeated three times in order to complete 4 h of reaction. At the end of the reaction time, the solvent was removed under reduced pressure. The crude product was purified by column chromatography (eluting with hexane:ethyl acetate, 90:10 with gradient elution until 0:100) to give the desired product as solid.

2-(4,5-Diphenyl-1-(6-(2,4,5-triphenyl-1H-imidazol-1-yl)hexyl)-1H-imidazol-2-yl)pyridine (10a)

Yellow solid; 37% yield; mp 78-79 °C; IR (KBr) ν max / cm⁻¹ 3059, 2926, 2857, 1586, 1464, 771, 691; ¹H NMR (400 MHz, CDCl₃) δ 8.52-8.49 (m, 1H), 8.35-8.31 (m, 1H), 7.77 (td, J 7.6, 2.0 Hz, 1H), 7.66-7.61 (m, 2H), 7.54-7.48 (m, 4H), 7.47-7.39 (m, 9H), 7.37-7.30 (m, 4H), 7.23-7.10 (m, 7H), 4.29 (t, J 7.6 Hz, 2H), 3.78 (t, J 7.6 Hz, 2H), 1.38-1.27 (m, 2H), 1.24-1.13 (m, 2H), 0.81-0.70 (m, 4H); ¹³C NMR APT (100 MHz, CDCl₃) δ 151.0, 148.2, 147.6, 144.0, 137.9, 137.7, 136.6, 134.5, 131.6, 131.5, 131.4, 131.2, 131.1, 130.9, 129.8, 129.5, 129.1, 129.0, 128.9, 128.8, 128.7, 128.6, 128.5, 128.1, 128.0, 127.7, 126.8, 126.7, 126.5, 126.3, 126.2, 123.4, 122.4, 45.0, 44.5, 30.5, 30.1, 25.5, 25.4; HRMS-ESI m/z, calcd. for [M + H]+: 676.3435, found: 676.3436.

3-(4,5-Diphenyl-1-(6-(2,4,5-triphenyl-1H-imidazol-1-yl)hexyl)-1H-imidazol-2-yl)pyridine (10b)

Yellow solid; 25% yield; mp 167-168 °C; IR (KBr) ν max / cm⁻¹ 3059, 2927, 2857, 1480, 773, 687; ¹H NMR (400 MHz, CDCl₃) δ 8.27-8.09 (m, 1H), 8.68 (dd, J 4.8, 1.6 Hz, 1H), 8.05-8.00 (m, 1H), 7.65-7.61 (m, 2H), 7.54-7.49 (m, 4H), 7.48-7.40 (m, 10H), 7.38-7.32 (m, 4H), 7.26-7.12 (m, 6H), 3.82-3.71 (m, 4H), 1.19-1.06 (m, 4H), 0.71-0.55 (m, 4H); ¹³C NMR APT (100 MHz, CDCl₃) δ 149.7, 149.3, 147.6, 144.3, 138.5, 137.7, 137.6, 136.3, 134.4, 134.1, 131.4, 131.3, 131.0, 130.9, 130.8, 130.2, 129.5, 129.1, 129.1, 129.0, 128.9, 128.8, 128.6, 128.5, 128.1, 128.0, 127.7, 126.8, 126.7, 126.5, 126.3, 126.2, 123.4, 45.0, 44.5, 30.5, 30.1, 25.5, 25.4; HRMS-ESI m/z, calcd. for [M + H]+: 676.3435, found: 676.3436.
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4-(4,5-Diphenyl-1-(7-(2,4,5-triphenyl-1H-imidazol-1-yl)heptyl)-1H-imidazol-2-yl)pyridine (11c)

Yellow solid; 26% yield; mp 152-153 °C; IR (KBr) νmax / cm⁻¹ 3043, 2919, 2857, 1597, 781, 687; ¹H NMR (400 MHz, CDCl₃) δ 8.75-8.71 (m, 2H), 7.68-7.62 (m, 4H), 7.55-7.42 (m, 13H), 7.41-7.34 (m, 4H), 7.26-7.12 (m, 6H), 3.92-3.79 (m, 4H), 1.31-1.14 (m, 4H), 0.80-0.61 (m, 6H); ¹³C NMR APT (75 MHz, CDCl₃) δ 150.2, 147.6, 144.4, 138.9, 138.8, 137.7, 134.5, 134.0, 131.5, 131.0, 130.9, 130.8, 129.5, 129.2, 129.1, 129.0, 128.9, 128.7, 128.6, 128.1, 128.0, 126.9, 126.8, 126.6, 126.3, 122.9, 44.7, 44.6, 30.3, 30.1, 28.2, 28.1, 25.9, 25.9; HRMS-ESI m/z, calcd. for [M + H]+: 704.3748, found: 704.3739.

Cholinesterase inhibition assay

Electric eel AChE and horse serum BuChE were used as sources of both cholinesterases. AChE and BuChE inhibitory activities were measured in vitro by the spectrophotometric method developed by Ellman, with slight modifications. The lyophilized enzymes, 500 U AChE and 300 U BuChE, were dissolved in buffer phosphate A (8 mM K₂HPO₄, 2.3 mM NaH₂PO₄) to obtain 5 and 3 U mL⁻¹ stock solution, respectively. Further enzymes dilution were carried out with buffer phosphate B (8 mM K₂HPO₄, 2.3 mM NaH₂PO₄, 0.15 M NaCl, 0.05% Tween 20, pH 7.6) to produce 0.5 and 0.3 U mL⁻¹ enzyme solution, respectively. Samples were dissolved in buffer phosphate B with 1.25% CHCl₃ and 13.75% of MeOH as a cosolvent mixture. 300 µL of enzyme solution and 300 µL of sample solution were mixed in a test tube and incubated for 60 or 120 min at room temperature. The reaction was started by adding 600 µL of the substrate solution (0.5 mM DTNB, 0.6 mM acetylthiocholine iodide/butyrylthiocholine iodide (ATCI/BTCI), 0.1 M Na₂HPO₄, pH 7.5), and the absorbance was read at 405 nm for 120 s at 25 °C. Enzymes activity was calculated by comparing reaction rates for the samples to the blank. All reactions were performed in triplicate, and the IC₅₀ values were determined with GraphPad Prism 5. The Lineweaver-Burk plot showed a pattern of lines with increasing slopes, characteristic of non-competitive type inhibition (inhibition constant (Ki) 21.29 nM). The nonlinear regression of these data fitted with non-competitive inhibition with a coefficient of determination R² = 0.9830.
Evaluation of cytotoxicity in in vitro models

The Vero (African green monkey kidney cells), HepG2 (liver hepatocellular carcinoma), and C6 (astroglial) cell lines were purchased from BCRJ (Rio de Janeiro Cell Bank). All cells were maintained in Dulbecco’s modified Eagle medium (DMEM) supplemented with 10% fetal bovine serum (FBS, Vero and HepG2 cells) or 5% FBS (C6 cells), 0.5% amphotericin B 250 μg mL⁻¹, 0.5% penicillin 100 IU mL⁻¹ and streptomycin 10 mg mL⁻¹, at 37 °C, in a humid atmosphere containing 5% of CO₂. The cells were seeded in 96-well plates (3 × 10⁴ cell well⁻¹ for C6, and 2 × 10⁴ cell well⁻¹ for Vero and HepG2) and incubated for 24 h. For cytotoxic studies, the selected compounds were dissolved in culture medium containing 0.5% DMSO, and the cells were exposed to them in concentrations up to 125 μM (compounds 5b and 12b) or 500 μM (tacrine), for 24 h. The cell viability was determined by the MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) assay. At least three independent experiments were performed in triplicate for each trial (n = 3). The results were expressed as IC₅₀ (50% inhibiting concentration) compared to the control (vehicle DMSO). The GraphPad Prism 5 program was used for statistical analysis.

Molecular docking

The three-dimensional structures of the compounds were prepared using the Maestro Suite and isomers, protonation states, and tautomers of the ligands were determined using LigPrep/Epik from Maestro at pH 7.4 ± 0.4. The AChE and BuChE structures were prepared with the Protein Preparation Wizard tool from Maestro. The protonation states of the amino acid residues were determined using PROPKA at pH 7.48. Interestingly, Glu199 located near the catalytic triad was predicted to be neutral due to a high pKᵢ value (ca. 10) for all the AChE and BuChE structures. Recently, Wan et al. proposed that the protonated form of Glu199 can interact with a conserved water and stabilize the catalytic triad in the molecular simulations of the BuChE-tacrine complex. The optimization of the hydrogen bond network between the protein and reference ligand was performed to adjust the hydrogen atoms’ orientation, followed by energy minimization with fixed heavy atoms. Due to the large size of the bis(n)-lophine derivatives, the water molecules were removed from the binding site.

In this work, the docking experiments were performed with the molecular docking program Glide under the standard precision (SP) mode. We redocked the reference ligands (i.e., the co-crystallized compounds) into their respective AChE and BuChE conformations to validate the docking protocol adopted herein. We selected the BuChE structure complexed with a dual-binding site tacrine-tryptophan hybrid (Protein Data Bank (PDB) ID 6IOC from Homo sapiens, solved at 2.67 Å). For AChE, we selected four representative conformations of AChE to consider the significant conformational changes mainly observed on the PAS using an ensemble docking strategy. This approach consists of docking the compounds into each representative conformation of the receptor, aiming to implicitly consider large-scale protein movements. We kept the conserved water molecules already reported in our previous work except for Wat720 (numbering from PDB code 6IOC for the BuChE structure), which is more exposed to the solvent and interacts with only one amino acid residue. For each ligand, the top-energy pose was selected according to the lowest Emodel value (i.e., the Emodel is the Glide scoring function recommended to evaluate different poses of the same ligand). Thus, the binding mode with the lowest GlideScore among the four AChE representative structures was selected for each compound. The protein conformations selected in this work for AChE were 1ZGC (Torpedo californica solved at 2.1 Å) (Haviv et al.), 2CKM (Torpedo californica solved at 2.1 Å), 1IQ8 (Mus musculus, solved at 2.4 Å), and 4EY7 (Homo sapiens, solved at 2.4 Å). All the inhibitors from the four representative conformations of AChE act as dual inhibitors interacting with both CAS and PAS.

Results and Discussion

Synthesis

The symmetrical bis(n)-lophine analogues were synthesized through a one-pot four component reaction of pyridinecarboxaldehyde (1a-1c), 1,ₙ-alkanediamines (2a-2c), benzil (3) and ammonium acetate (4, Scheme 1). The reaction was performed in a microwave reactor at 110 °C for 5 h. The reaction provides the bis(n)-lophines analogues (5-7), in 31-55% yield, in one reaction step.

For the synthesis of unsymmetrical bis(n)-lophine analogues, it was necessary synthesize the N-alkylaminolophine precursors (9a-9c) through the multicomponent reaction between benzaldehyde (8), 1,ₙ-alkanediamines (2a-2c), benzil (3) and ammonium acetate (4, Scheme 2). In order to optimize the synthesis of precursors 9a-9c using a microwave reactor, we chose hexanediane as substrate. The aim was to examine some of this reaction’s general features, such as Lewis acid catalysts, stoichiometry, reaction time, temperature and reaction yield.
As shown in Table 1 (entry 6), the use of 1 equiv. of NH₄OAc and 1 equiv. of hexanediamine, 110 ºC for 5 h and InCl₃ as catalyst, afforded 9a in 50% yield. Interestingly, we found that in this reaction conditions, the use of Lewis acid was not necessary, affording the precursor 9a in 47% yield (entry 10). In this way, the conditions of entry 10 were applied for the synthesis of N-alkylaminolophine precursors 9b and 9c. Next, the multicomponent reaction of 9a-9c with pyridinecarboxaldehyde (1a-1c), benzil (3) and ammonium acetate (4) provided the unsymmetrical bis(n)-lophine analogues in 23-49% yield (10-12, Scheme 3).

**In vitro** inhibition studies on AChE and BuChE

AChE and BuChE activities were measured *in vitro* by the spectrophotometric method developed by Ellman *et al.*,41 with slight modifications, with tacrine as the reference inhibitor. The inhibitors’ effectiveness is expressed as IC₅₀, representing the concentration of an inhibitor required for 50% inhibition of the enzyme. For this study, compounds with IC₅₀ values over 50 µM were considered to be inactive. All the compounds, except 7a, 7b and 7c, displayed potent inhibitory activity against BuChE at a micromolar and sub-micromolar range (IC₅₀ 32.25-0.03 µM). The inhibition was found to be highly selective since none of the bis(n)-lophine analogues were active against AChE, at the tested concentrations. BuChE inhibitory activity results are summarized in Table 2. These results indicate that BuChE inhibition is mainly influenced by two features: the R substituent nature and the length of the alkyl spacer. Among all the tested compounds, the asymmetric analogues of bis(n)-lophine with 2-pyridine and 3-pyridine moiety were better inhibitors of BuChE than those containing 4-pyridine moiety. Compound 11b, characterized by an unsubstituted
value of 0.034 µM. In the case of asymmetric analogues of bis(n)-lophine with an eight-carbon linker, the most active compound was 12b (IC\textsubscript{50} 0.071 µM), and for those with a spacer of six methylenes, compound 10b resulted in the most active one (IC\textsubscript{50} 0.134 µM). Both derivatives 12b and 10b share with 11b the presence of phenyl and 3-pyridine moieties as R\textsubscript{1} and R\textsubscript{2}. Since compound 11b was the most effective BuChE inhibitor of the series, it was selected for the kinetic study of enzyme inhibition.

**Kinetic characterization of BuChE inhibition**

Enzyme activity was evaluated at different fixed substrate concentrations and increasing inhibitor concentrations. The data were used to elucidate the enzyme inhibition mechanism, and the results are illustrated in the form of Lineweaver-Burk plots (Figure 1). The double reciprocal plots obtained using increasing inhibitors concentrations demonstrated that compound 11b behaves as a non-competitive inhibitor (see Figure 1). In fact, experimental data obtained using increasing inhibitor concentrations describe straight lines

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**Table 1.** One-pot four component reaction for the synthesis of 9a under different conditions

| entry | NH\textsubscript{4}OAc / mmol | Hexanediamine / mmol | Temperature / ºC | Catalyst (0.15 mol%) | time / h | Yield / % |
|-------|------------------|------------------|------------------|-------------------|----------|-----------|
| 1     | 1                | 1                | 78               | InCl\textsubscript{3} | 1        | 13        |
| 2     | 1                | 2                | 78               | InCl\textsubscript{3} | 1        | 8         |
| 3     | 2                | 1                | 78               | InCl\textsubscript{3} | 1        | 0         |
| 4     | 1                | 1                | 110              | InCl\textsubscript{3} | 3        | 40        |
| 5     | 1                | 2                | 110              | InCl\textsubscript{3} | 3        | 10        |
| 6     | 1                | 1                | 110              | InCl\textsubscript{3} | 5        | 50        |
| 7     | 1                | 1                | 110              | FeCl\textsubscript{3}.6H\textsubscript{2}O | 5        | 41        |
| 8     | 1                | 1                | 110              | CeCl\textsubscript{3} | 5        | 44        |
| 9     | 1                | 1                | 110              | SnCl\textsubscript{2}.2H\textsubscript{2}O | 5        | 34        |
| 10    | 1                | 1                | 110              | –                 | 5        | 47        |

**Table 2.** Inhibition of BuChE activity for bis(n)-lophine analogues

| entry | Compound | R\textsubscript{1} | R\textsubscript{2} | n   | IC\textsubscript{50} BuChE / µM | log IC\textsubscript{50} ± SD |
|-------|----------|------------------|------------------|-----|-------------------------------|-----------------------------|
| 1     | 5a       | 2-Py             | 2-Py             | 6   | 0.246                         | -0.608 ± 0.062              |
| 2     | 5b       | 3-Py             | 3-Py             | 6   | 0.208                         | -0.691 ± 0.198              |
| 3     | 5c       | 4-Py             | 4-Py             | 6   | 25.84                         | 1.412 ± 0.053               |
| 4     | 6a       | 2-Py             | 2-Py             | 7   | 32.25                         | 1.508 ± 0.111               |
| 5     | 6b       | 3-Py             | 3-Py             | 7   | 12.04                         | 1.081 ± 0.091               |
| 6     | 6c       | 4-Py             | 4-Py             | 7   | 23.27                         | 1.367 ± 0.125               |
| 7     | 7a       | 2-Py             | 2-Py             | 8   | n.a.                          | –                           |
| 8     | 7b       | 3-Py             | 3-Py             | 8   | n.a.                          | –                           |
| 9     | 7c       | 4-Py             | 4-Py             | 8   | n.a.                          | –                           |
| 10    | 10a      | Ph               | 2-Py             | 6   | 0.155                         | -0.808 ± 0.058              |
| 11    | 10b      | Ph               | 3-Py             | 6   | 0.134                         | -0.874 ± 0.075              |
| 12    | 10c      | Ph               | 4-Py             | 6   | 0.164                         | -0.784 ± 0.109              |
| 13    | 11a      | Ph               | 2-Py             | 7   | 0.091                         | -1.041 ± 0.039              |
| 14    | 11b      | Ph               | 3-Py             | 7   | 0.034                         | -1.463 ± 0.063              |
| 15    | 11c      | Ph               | 4-Py             | 7   | 1.294                         | 0.112 ± 0.164               |
| 16    | 12a      | Ph               | 2-Py             | 8   | 0.578                         | -0.238 ± 0.070              |
| 17    | 12b      | Ph               | 3-Py             | 8   | 0.071                         | -1.151 ± 0.100              |
| 18    | 12c      | Ph               | 4-Py             | 8   | 1.586                         | 0.200 ± 0.044               |
| 19    | tacrine  | –                | –                | 50  | 0.004                         | -2.35 ± 0.07                 |

n: number of methylenes in the linker; IC\textsubscript{50}: half maximal inhibitory concentration; BuChE: butyrylcholinesterase; SD: standard deviation; Py: pyridine; Ph: phenyl; n.a.: not active.

**Figure 1.** Lineweaver-Burk plots of the BuChE inhibition by compound 11b with butyrlthiocholine (S) as substrate.
intersecting each other at a point on the x-axis. Moreover, the tested compounds determine the decrease of the maximum reaction velocity ($V_{\text{max}}$), without affecting the Michaelis-Menten constant ($K_m$) values appreciably. Thus, the enzyme kinetic study suggests that the inhibitor binds to both the CAS and PAS sites of BuChE. The inhibition constant $K_i$ was equal to 0.021 μM. For comparison purpose, the reported $K_i$ of tacrine is about 0.017 μM.

**Evaluation of cytotoxicity in in vitro models**

The bis(n)-lophine analogues 5b and 12b were evaluated for their cytotoxicity against cellular models of Vero (kidney), HepG2 (hepatic), and C6 (astroglial) cell lines, which are widely used as experimental models for seeking information about nephrotoxicity, hepatotoxicity, and neurotoxicity, respectively.

According to the results (see Table 3, and Supplementary Information section, Figures S38-S43), the compounds 5b and 12b showed no or slightly cytotoxic effects in all cellular models at the tested concentrations, exhibiting an IC$_{50}$ higher than 125 μM. These results suggest that the compounds have a degree of safety because there is a large difference between the IC$_{50}$ for BuChE inhibition (IC$_{50}$ = 0.208 μM for 5b and 0.071 μM for 12b) and the concentration to which all cell lines were exposed (up to 125 μM) in the cytotoxic assay. For the 5b and 12b derivatives, concentrations 600 and 1760 times higher than those active for inhibiting BuChE were used, respectively, and small or no reduction in the cell viability was observed. Also, it can be observed that the bis(n)-lophine analogues are significantly less toxic than tacrine to the hepatic cell line (HepG2), once at 125 μM tacrine displayed a cell viability of 44%, while the derivatives 5b and 12b showed about 70 and 95% in cell viability at the same concentration.

**Table 3.** Cell viability after treatment with the selected compounds for 24 h using MTT assay (IC$_{50}$) in Vero, HepG2 and C6 cell lines.

| Compound | Vero (µM) | HepG2 (µM) | C6 (µM) |
|----------|-----------|------------|---------|
| 5b       | > 125     | > 125      | > 125   |
| 12b      | > 125     | > 125      | > 125   |
| Tacrine  | 168.9 ± 14.0 | 114.9 ± 17.9 | 450.7 ± 4.2 |

IC$_{50}$: half maximal inhibitory concentration; SEM: standard error of the mean.

**Molecular docking**

The redocking experiment with the BuChE conformation used in this work (PDB code 6I0C) was able to reproduce the experimental binding mode of the co-crystallized ligand as the top-scored pose (GlideScore = −11.022 kcal mol$^{-1}$) within a root mean square deviation (RMSD) value of 0.53 Å (Figure 2). The chlorotacrine moiety was perfectly predicted near CAS, whereas the indole group exhibited an inverted mode, making T-stacking interaction with W231, a residue involved in the selectivity against BuChE over AChE. Also, the ensemble docking strategy adopted for AChE was successfully validated with all the reference ligands docked with RMSD ≤ 2 Å in the respective protein conformation according to the lowest GlideScore.

The two most potent compounds 11b and 12b were ranked at the top-3 best scored ligands (Table 4, −11.28 and −11.052 kcal mol$^{-1}$, respectively) and predicted to interact with the BuChE cavity with similar binding modes, with the unsubstituted lophine moiety interacting at the bottom of the gorge, whereas the N-substituted lophine moiety interacts with the PAS region. At the bottom of the binding cavity, these ligands interact through lipophilic and T-stacking interactions, mainly with the side chains of key aromatic residues such as Trp82, His438, Trp321, Phe329, and Tyr332 (Figure 3). Despite these interactions, the conserved π-stacking or cation-π interactions commonly observed with Trp82 for potent inhibitors and tacrine were not observed in the docking experiments, probably due to the larger size and small flexibility of the lophine moiety leading to a suboptimal fit to the binding site. Notably, the bis(n)-lophine derivatives are able to interact more deeply on the BuChE binding site than the AChE cavity, being closer to Trp82 in the first. For instance, a phenyl ring of the compound 11b was predicted to be within a distance of 3.5 and 5.5 Å to the closest indole ring atom from Trp82 of BuChE (GlideScore = −11.28 kcal mol$^{-1}$) and AChE.
In addition, the remarkable selectivity of the lophine dimers for BuChE over AChE might also be due to the absence of some aromatic residues at the BuChE binding site, leading to an opened binding site that is able to best accommodate such large ligands and the formation of specific hydrophobic pockets frequently related with selectivity against BuChE. For example, according to the docking results, the bis(n)-lophine dimers interacted with the Trp231 side chain through a T-stacking interaction, a residue only accessible in BuChE and claimed to be a key residue for selectivity. 40,59

Despite the predicted binding affinities being quite similar for all compounds, probably the main difference in the experimental affinities observed between the series of homo and heterodimers is related to the desolvation penalty arising from burying the pyridine group in the homodimer series without compensation with polar interactions with the environment. According to the docking results, all heterodimer derivatives were predicted to have the unsubstituted lophine moiety interacting at the bottom of the cavity, except the compounds with the nitrogen atom at the ortho position. On the other hand, the N-substituted phenyl ring at meta-position of the heterodimer compounds, such as 10b and 11b, allowed the formation of a hydrogen bond with the Asn68 side chain from PAS (2.92 and 3.17 Å, respectively).

**Conclusions**

In summary, two new series of bis(n)-lophine analogues were synthesized through a one-pot four component reaction performed in a microwave reactor. Also, a new protocol for the synthesis of N-alkylaminolophine precursors was developed without the need of Lewis acid as catalyst. The bis(n)-lophine analogues were strongly selective to BuChE and present IC₅₀ at a micromolar and sub-micromolar range (IC₅₀ 32.25-0.03 µM). Compounds 11b (IC₅₀ = 0.034 µM) and 12b (IC₅₀ = 0.071 µM) were the most potent inhibitors of the asymmetric bis(n)-lophines series. Of the symmetric bis(n)-lophines, the most active compound was 5b (IC₅₀ = 0.208 µM). The kinetic characterization of BuChE inhibition by compound 11b suggests that the compound bind to both CAS and PAS sites, corroborated by the docking results. According to the predicted binding mode for the compound 11b, it is able to interact deeply into the BuChE binding site and a hydrophobic pocket involved with selectivity formed mainly by the residue Trp231, which is only available at the BuChE cavity. The dimers 5b and 12b showed no cytotoxic effects in Vero (renal), HepG2 (hepatic), and C6 (astroglial) cell lines and a wide

| entry | Compound | GlideScore / (kcal mol⁻¹) | IC₅₀ / µM |
|-------|----------|---------------------------|-----------|
| 1     | 5a       | -9.708                    | 0.246     |
| 2     | 5b       | -9.657                    | 0.208     |
| 3     | 5c       | -10.379                   | 25.84     |
| 4     | 6a       | -10.808                   | 32.25     |
| 5     | 6b       | -9.827                    | 12.04     |
| 6     | 6c       | -10.468                   | 23.27     |
| 7     | 7a       | -10.395                   | n.a.      |
| 8     | 7b       | -10.594                   | n.a.      |
| 9     | 7c       | -10.484                   | n.a.      |
| 10    | 10a      | -10.102                   | 0.155     |
| 11    | 10b      | -9.98                     | 0.134     |
| 12    | 10c      | -10.028                   | 0.164     |
| 13    | 11a      | -10.405                   | 0.091     |
| 14    | 11b      | -11.28                    | 0.034     |
| 15    | 11c      | -11.323                   | 1.294     |
| 16    | 12a      | -10.944                   | 0.578     |
| 17    | 12b      | -11.052                   | 0.071     |
| 18    | 12c      | -10.927                   | 1.586     |
| 19    | tacrine  | -7.578 (0.934)            | 0.004     |

*a ROOT mean square deviation from the experimental binding mode present in the 4BDS (PDB code for human BuChE in complex with tacrine) conformation of BuChE. IC₅₀: half maximal inhibitory concentration; n.a.: not active.*

**Figure 3.** Predicted binding modes of the compounds (a) 11b (carbon atoms colored cyan), (b) 12b (carbon atoms colored cyan), and (c) tacrine (carbon atoms colored green) superimposed to 11b against BuChE (PDB code 6I0C). (d) Predicted binding mode of the compound 11b against AChE (carbon atoms colored yellow, PDB code 1Q84) superimposed with its binding mode predicted against BuChE (transparent sticks with carbon atoms colored cyan, PDB code 1Q84). Hydrogen bonds are represented as yellow dashed lines.
range of safety considering the active concentration for BuChE inhibition.

**Supplementary Information**

Supplementary data (\(^1\)H and \(^{13}\)C NMR spectra and cytotoxicity results) are available free of charge at http://jbcs.sbq.org.br as PDF file.

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**Author Contributions**

Viktor S. Câmara was responsible for the data curation, investigation, methodology, visualization, writing original draft, review and editing; Ana Julia Soares for data curation and methodology; Brunella Biscusi for data curation, formal analysis, investigation, methodology, visualization, writing original draft; Ana Paula Murray for the data curation, formal analysis, funding acquisition, investigation, methodology, resources, supervision and writing original draft; Isabella A. Guedes for the conceptualization, data curation, formal analysis, investigation, software, writing original draft; Laurent E. Dardenne for the conceptualization, data curation, funding acquisition, investigation, methodology, resources, supervision and writing original draft; Aline R. Zimmer for the data curation, formal analysis, investigation, methodology, validation and writing original draft; Marco A. Ceschi for the conceptualization, data curation, formal analysis, funding acquisition, investigation, methodology, project administration, resources, supervision, writing original draft, review and editing.

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