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Aqueous extracts of marine invertebrates from Cuba coastline display neutral aminopeptidase inhibitory activities and effects on cancer cells and *Plasmodium falciparum* parasites

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Neutral aminopeptidases are enzymes distributed in all living organisms. By hydrolyzing biologically active peptides in tissues and biological fluids, they are involved in the control of many physiological processes. They became established targets for new therapeutic agents in cancer, but also in parasitic diseases like malaria. Marine organisms are promising sources for biomolecules but few examples of neutral aminopeptidase inhibitors are described. The goal of this work was to search in Cuban marine invertebrates, for inhibitory activities of neutral aminopeptidases of biomedical relevance, belonging to the M1 and M17 metallopeptidase families. The screening of inhibitory activities was performed using aqueous crude extracts and their 2.5 % TCA treatments. The treatments with 2.5 % TCA increased the recovery of inhibitory activities versus all enzymes tested and from all of marine species. These inhibitory activities were dose-dependent in all cases, with certain selectivity for PfA-M17 regarding hLAP, and good inhibition of hAPN. Interestingly, some TCA treated extracts displayed promising effect on either *Plasmodium* parasite growth as well as on PC3 and 3LL cells. This contribution is the first report identifying inhibitory activities from marine invertebrates, directed against human and malarial neutral aminopeptidases, suggesting a potential for biomedical applications for the corresponding marine species.

Keywords: Cancer, Inhibitors, Malaria, Marine invertebrates, Neutral aminopeptidases, Screening.

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Introduction

Neutral aminopeptidases are enzymes that catalyze the cleavage of neutral amino acids from the N-terminus of protein or peptide substrates. They have been classified in several metallopeptidase families (like M1 and M17 families)^{1,2}. These enzymes are present in all living organisms, but the diversity of the functions in which they are involved is far from being entirely deciphered.

In humans, neutral aminopeptidases are distributed in several tissues and found in various subcellular

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Supplementary figures are available online only.

organelles, in the cytoplasm, and as integral membrane proteins. In particular, neutral metalloaminopeptidase (APN, EC 3.4.11.2, M1 family) catalyzes the cleavage of neutral and basic amino acids from the N-terminus of protein or peptide substrates and is involved in the degradation of enkephalin, now a target for the development of new pain management strategies^{3,4}. This enzyme is also up-regulated in human pathologies such as various types of cancers, inflammation, and skin diseases⁵⁻⁷. Strategies for inhibition of APN⁸ have been developed primarily for the treatment of pain^{4,9} and less so for cancer and skin pathologies. Mammalian leucyl aminopeptidase (LAP, EC 3.4.11.21, M17 family) is cytosolic and involved in the breakdown of peptide products of intracellular proteinases^{1,2}. It is one of the enzymes that trim proteasome-produced peptides for class I antigen presentation and is also associated with tumor cell proliferation, invasion and/or angiogenesis^{10,11}.

Neutral aminopeptidases are also distributed in Apicomplexan protist parasites like Plasmodium falciparum, the main agent of malaria in humans. Malaria remains the most deadly human parasitic disease in many parts of the world, especially in the Sub Sahara Africa, and is responsible for over 438,000 deaths. In 2015, more than 214 million people had malaria. Malaria is caused by five Plasmodium species transmitted by the bite of female Anopheles mosquitoes, Plasmodium falciparum being by far the most lethal species of *Plasmodium*. The widespread appearance of drug-resistant parasites, even to newly-developed second and third generation therapeutics such as artemisinin and its derivatives, illustrates the need to design the next generation of anti-malarial drugs to inhibit biochemical pathways critical for parasite survival and/or transmission¹². The most important clinical stage of the complex P. falciparum life cycle¹³, which has attracted the highest attention for the development of antimalarials, takes place in the human erythrocyte where significant hemoglobin degradation occurs under the concerted action of endo and exo peptidases^{14,15}. These enzymes include PfA-M1 (M1 family) and PfA-M17 (M17 metallo-aminopeptidases family). two neutral involved in the final steps of hemoglobin digestion¹⁶. Both enzymes are promising chemotherapeutic targets because their inhibitors can kill parasites in vitro and *in vivo*¹⁷⁻²⁷.

While natural products are a major source of drug lead compounds, the tropical marine sources of aminopeptidases inhibitors, in particular from the Caribbean invertebrate fauna, remain largely unexplored and have the advantages of their tremendous richness and diversity. Two inhibitors of the M1 family enzymes have been isolated from marine organisms, PsammaplinA²⁸ and HcPI²⁹. However, there are no available reports in the literature of inhibitors isolated from marine sources that are targeting the M17 family of neutral aminopeptidases.

The aim of the present study was to investigate whether Cuban marine invertebrates represent valuable sources of promising inhibitors of M1 and M17 neutral aminopeptidases of medical relevance for either cancer or malaria.

Materials and Methods Materials

Human placenta from the Institute of Research on Placental Histotherapy was kindly donated by Institute Oncology and Radiobiology, Cuba. DEAE Sephacel was purchased from Amersham Biosciences. L-Leu-AMC, bestatin and amastatin were purchased from Sigma-Aldricht. Leu-pNA was purchased from Bachem. The rest of the reagents were of analytical grade. PC3 (CRL 1435) and 3LL (CRL-1642) tumor cell lines were from ATCC (American Type Culture Collection, EEUU) and were kindly donated by the Center for Molecular Immunology, Cuba. The media and supplements for tumor cell cultures were purchased from Gibco. Marine invertebrates used in the present study were identified by Aida Hernandez-Zanuy, PhD, from the National Institute of Oceanology (CITMA, Cuba) (Table 1, Plate1).

Aminopeptidase N and Leucyl aminopeptidase preparations from human placenta

Human Aminopeptidase N (hAPN) was prepared according to Byzia *et al.*³⁰ in the form of microsomes by differential centrifugation, but without Triton X-100. The microsomal fraction was resuspended in 50 mM Tris-HCl pH 7.5 (buffer A) for use in enzymes assays. Human Leucyl

Table 1 — Marine invertebrate's species from the Havana coastline studied in the present contribution.

Species	Phyla: Class: Family		
Cenchritis muricatus (Linnaeus,	Mollusca: Gastropoda:		
1758)	Littorinidae		
Nerita peloronta (Linnaeus, 1758)	Mollusca: Gastropoda:		
	Neritidae		
Nerita versicolor (Gmelin, 1791)	Mollusca: Gastropoda:		
	Neritidae		
Lissodendoryx (Lissodendoryx)	Porifera: Demospongiae:		
isodictyalis (Carter, 1882)	Coelosphaeridae		
Tripneustes ventricosus	Echinodermata: Echinoidea:		
(Lamarck, 1816)	Toxopneustidae		
Echinaster (Othilia) echinophorus	Echinodermata: Asteroidea:		
(Lamarck, 1816)	Echinasteridae		
Isostichopus badionotus (Selenka,	Echinodermata:		
1867)	Holothuroidea:		
	Stichopodidae		
Stichodactyla helianthus (Ellis,	Cnidaria: Anthozoa:		
1768)	Stichodactylidae		
Bunodosoma granuliferum(Le	Cnidaria: Anthozoa:		
Sueur, 1817)	Actiniidae		
Physalia physalis (Linnaeus,	Cnidaria: Hydrozoa:		
1758)	Physaliidae		

Species names were validated on the reference WORMS database (http://www.marinespecies.org/index.php).

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Plate 1 — Some of the marine invertebrates assayed in the present contribution: a) *Nerita peloronta*, b) *Nerita versicolor*, c) *Cenchritis muricatus*, d) *Isostichopus badionotus*, e) *Tripneustes ventricosus*, f) *Echinaster (Othilia) echinophorus*, g) *Stichodactyla helianthus*, and h) *Bunodosoma granuliferum*. Pictures were supplied by Professor Jose Espinosa, PhD from the National Institute of Oceanology, CITMA, Cuba.

aminopeptidase (hLAP) was partially purified from the soluble fraction of human placenta homogenates obtained from the first centrifugation at 10 000 \times g. This supernatant was extensively dialyzed in presence of buffer A, using dialysis membranes with a 10 000 Da cut off (Spectrapor). The dialyzed sample was applied onto a DEAE Sephacel column (12 \times 1.6 cm) previously equilibrated with the same buffer A. The column was washed with 3 column volumes of buffer A to wash out the non-retained material. Then, elution was carried out by a stepwise salt increase using 0.15, 0.3, 0.5, 0.75 and 1.0 mol/L NaCl concentrations in buffer A with 3 column volumes of buffer each step. Active fractions (0.15 mol/L NaCl) were collected, concentrated via ultrafiltration using a membrane of 10 000 Da of cutoff, and used for enzyme activity.

Preparation of recombinant forms of malarial PfA-M1 and PfA-M17 aminopeptidases

Both malarial enzymes were produced as recombinant, enzymatically active proteins, in Escherichia coli. For PfA-M1, a genomic DNA fragment encoding residues 192-1085 of native PfA-M1^{31,32}was amplified from genomic DNA of the FcB1 strain of P. falciparum, with primers appending a N-terminal hexahistidine tag followed by cleavage site for the tobacco etch virus (TEV) protease (ENLYFOS), and was cloned into the BamH1 and NotI sites of the pET45b vector (Novagen). For PfA-M17, a synthetic gene (Genecust, Luxembourg) encoding a cleavage site for the TEV protease fused to residues 84-605 of native P. falciparum leucyl aminopeptidase PfA-M17 (PlasmoDB PF3D7 1446200), was cloned into BamHI and SalI sites of the pET45b vector (Novagen), which appended an N-terminal hexahistidine tag. Both plasmids were transformed into Escherichia coli BL21(DE3) Rosetta 2 (Novagen) after appropriate validations by Sanger sequencing (Beckman Coulter Genomics). Bacterial cultures were grown in autoinduced LB medium (Merck) supplemented with appropriate antibiotics (carbenicillin 50 µg/mL, chloramphenicol 34 µg/mL) for 24 h at 25 °C under vigorous agitation. The clarified lysates were loaded onto Ni²⁻ charged HisTrap column (GE Healthcare) equilibrated in phosphate buffer supplemented with 20 mM imidazole, extensively washed in phosphate buffer supplemented with 20 mM imidazole, and bound recombinant proteins were eluted in respectively

80 mM imidazole in phosphate buffer (rPfA-M1) and 200 mM imidazole in phosphate buffer (rPfA-M17). Eluted fractions were extensively dialyzed at 4 °C into respectively 50 mM Tris-HCl, 200 mM NaCl, 10 μ M ZnCl₂, pH 7.4 (rPfA-M1)²¹ and 50 mM Tris-HCl, 200 mM NaCl, 10 μ M ZnCl₂ pH 8 (rPfA-M17)³³.

Preparation of crude extracts from marine invertebrates

Organisms were collected along the coast of Havana (Cuba), classified, transported alive on ice, and stored at -20 °C until used for aqueous extract preparation. Specimens for each species were thawed, washed with distilled water, dried on paper filter, weighted, cut into small pieces with sterile scissors and homogenized in distilled water 2:1 (mL/g) at 4 °C in a warring blender (10 sec for 3 repeated cycles) as described earlier^{29,34}. Distilled water was preferred as extracting solvent for extract preparation with the aim

to have a crude extract enriched in molecules soluble and stable in aqueous conditions. This selection was based on their potential future application in different biomedical models, all of them requiring physiological conditions. Homogenates were centrifuged at 10 000 g for 30 min at 4 °C. The supernatants were filtered on fiberglass to yield aqueous crude extracts. These extracts were lyophilized and kept at -20 °C until being suspended in the correspondent buffer to assay for neutral aminopeptidase inhibitory activities.

Trichloroacetic clarification treatment of marine crude extracts

About 200 mg of each lyophilized crude extract was suspended in 5 mL buffer A and treated with 2.5 % final concentration of the chaotropic agent trichloroacetic acid (TCA) for 15 min at 4 °C following procedure described earlier^{29,34}. The aim of this clarification step was to eliminate contaminants (mainly proteins susceptible to this condition), and to promote dissociation from endogenous inhibitor-target complexes that do not allow the detection of inhibitory components in crude extracts^{29,34}. The TCA clarified and dialyzed extracts were then kept at -20 °C until assayed for neutral aminopeptidase inhibitory activities.

Protein concentration determination for marine extracts

The protein concentration was determined at 280 nm as described by Scopes³⁵ using one unit as an arbitrary extinction coefficient (0.1 %, 1 cm) and bovine serum albumin as standard in a GENESYS 10S UV-Vis spectrophotometer (Thermo Scientific) similar to previously described^{29,34}.

Monitoring of human and malarial enzymatic activities and inhibitions

The activities of the neutral aminopeptidases used in the present work were measured using the substrate L-Leu-AMC in specific conditions established for each enzyme. The substrate concentration in each assay was equal or lower than one K_M value for each enzyme to avoid competition. In all cases kinetic assays were run at 37 °C during 40 min in 96 well plates and the AMC fluorescence was measured at 460 nm (bandwidth: 40) upon excitation at 360 nm (bandwidth: 40) in a BioTek FL600 fluoromark spectrofluorometer^{31,32}.

Human placental aminopeptidase N activity (0.25 mg/mL) was measured using buffer A, in presence of 0.030 mmol/L of L-Leu-AMC in a final assay volume of 200 μ L. Human placental Leucyl aminopeptidase activity (0.84 mg/mL) was measured

using buffer A, in presence of 0.200 mmol/L of L-Leu-AMC in a final assay volume of 200 μ L. rPfA-M1 activity (666 nmol/L) was measured using buffer A, in presence of 0.160 mmol/L of L-Leu-AMC in a final assay volume of 100 μ L. rPfA-M17 activity (280 nmol/L) was measured using buffer Tris HCL 50 mmol/L pH 8, 1 mmol/L Cobalt (Buffer B) in presence of 0.050 mmol/L of L-Leu-AMC in a final assay volume of 100 μ L. One unit of enzyme activity was defined as the amount of enzyme needed to produce one arbitrary unit of fluorescence (AUF) per minute in the specified conditions.

Inhibition by aqueous crude and 2.5 % TCA cleared extracts from marine invertebrates was determined by quantifying the decrease in activity of each enzyme (hAPN, hLAP and rPfA-M1, rPfA-M17) pre-incubated with the extract sample for 30 min at 37 °C prior to addition of L-Leu-AMC substrate, regarding the control assay. One unit of inhibitory activity was defined as the amount of the extract (in terms of protein amount) needed to inhibit one unit of enzyme activity and was calculated using the formula

sIA (U/mg)= $(v_o-v_i)*V_{assay}/_{mg prot sample}$

where sIA is specific inhibitory activity, v_o is initial rate of the control, v_i is initial rate in presence of the inhibitory sample, and V_{assay} is assay volume.

Based on the increment of the specific inhibitory activity detected for 2.5 % TCA cleared extracts, dose response studies were carried out in these cases in the range of 0.1-1000 μ g/mL of each extract (n=3). IC₅₀ (half maximal inhibitory concentration) values were determined by non-linear regression adjustment of the dose-response curves (residual activity vs protein concentration of each treated extract) to the IC_{50} equation included in the GRAFIT 6.0 software and was defined as the amount of the extract (in terms of protein concentration) needed to inhibit the control enzyme activity by 50 %. Dose-response studies in presence of bestatin (1-100 µM, equivalent to 0.34-34.40 µg/mL) and amastatin (4-400 µM, equivalent to 1.89-189.00 µg/mL) were done in parallel as positive control of inhibition for each target aminopeptidase.

Evaluation of the effects on tumor cell lines cultures growth Detection of the APN activity in the PC3 and 3LL tumor cell lines cultures

The APN activity present in the membrane of the tumor cells in study was measured spectrophotometrically

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according to the work done by Ashmun and Look³⁶ using Leu-pNA as substrate. A total of 5×10^4 cells per well were seeded, in flat end 96 well plates (Costar, High Binding, EEUU) and incubated in DMEM-F12 with 10 % of SFT, during 24 h at 37 °C at 5 % of CO₂. After this, the medium was removed and 90 µL of DMEM-F12 was added to the intact cells. To begin the enzymatic reaction, 10 µL of Leu-pNA (3 mmol/L in the assay) previously incubated at 37 °C, were added to each well. The reaction was carried out at 37 °C for 1 h. The formation of pNA was followed by absorbance at 405 nm in a 96 well plate kinetic spectrometer (iMark Biorad), using the "M6M" software supplied by the manufacturer. The enzyme activity was determined using the equation given below and expressed as $nmol/h/10^6$ cells. Assays were run in triplicates using in parallel a standard free of cell and a control free of substrate.

EA (nmol/h/10⁶ cells) = $(\Delta DO/\Delta t) \times (1/\xi) \times (Va/(No.Cel./10⁶ cells))(1)$

where, EA is enzymatic activity; $\Delta DO/\Delta t$ (λ = 405 nm) is variation in the absorbance at 405 nm after 1 h; Va is total volume assay (mL); No.Cel. is number of cell per well; ξ is extinction coefficient of pNA at 405 nm ([[8800 (mol/L)

Effects on mammalian cancer cell viability

The viability of each cell line in presence of each 2.5 % TCA cleared and dialyzed extracts were determined by a modification of the colorimetric assay based on MTT (3-[4,5-dimethylthiazol-2-yl]-2,5 diphenyl tetrazodium bromide) method previously described by Mosmann³⁷. Flat-bottomed 96-well microculture plates (Costar) were seeded with PC3 (human prostatic adenocarcinoma) or 3LL (mouse lung Lewis carcinoma) cell line in 90 μ L (10⁴ cell)/well and grown in DMEM-F12 with 10 % SFT and kept at 37 °C with 5 % CO₂, for 24 h before the addition of the inhibitor sample diluted in 10 µL of media. After 72 h (for PC3) and 48 h (for 3LL) of incubation at 5 % of CO₂, cells viability was measured by the addition of MTT at 1 mg/mL. All the tests were performed in triplicates. Cells without treatment were included as a maximum cell growth point. Bestatin was included as positive control of inhibition. Percentage of viable cells in test wells was calculated using the following formula:

Viable cells (%) = (O.D.540 nm-620 nm of treated cells/O.D.540 nm-620 nm of non-treated cells) \times 100

Effect on *P. falciparum* growth in cultures of the 2.5 % TCA cleared extracts displaying inhibitory activities on malarial rPfA-M17

The general procedure was similar to that previously described (Flipo et al.)¹⁸. Antimalarial assays were performed using the P. falciparum FcB1 strain, grown in vitro according to Trager and Jensen³⁸. Parasite culture medium contains RPMI 1640 medium (Life Technologies, Inc.), 25 mM Hepes, 27.5 mM NaHCO₃, and 11 mM glucose (pH 7.4) and is supplemented with 8 % (v/v)compatible heat-inactivated human serum. Human red blood cells (RBC) were added at hematocrit of 2 %, and the parasite culture was maintained at 37 °C under an atmosphere of 3 % CO₂, 6 % O₂, and 91 % N₂, with daily medium changes. The effect of selected TCA clarified extracts on the intraerythrocytic development of P. falciparum was tested on asynchronous cultures containing mixtures of all stages at a final parasitemia of ~1 % that were distributed in 96-well microplates. Serial dilutions of selected extracts were added to the wells, in parasite culture medium, in triplicates. Parasites were allowed to grow in a candle jar system and after 24 h in culture, ³H-hypoxanthine was added (0.5 μ Ci/well). After an additional 24 h incubation period at 37 °C in the candle jar system, parasites were harvested on filters after a freeze-thawing cycle. Dried filters were submerged in a liquid scintillation mixture (OptiScintHisafe, Perkin Elmer) and counted in a 1450 Microbeta counter (Wallac, Perkin Elmer). Parasite growth inhibition (IC₅₀) was calculated from the parasite-associated radioactivity (incorporated into nucleic acids) in treated cultures compared with control cultures performed in the presence of equivalent amounts of buffer A as described earlier^{18,39}. Bestatin was included as positive control of parasite growth inhibition.

Results

Inhibition of hAPN, hLAP, rPfA-M1 and rPfA-M17 by aqueous extracts from marine invertebrates

In the present study, initially, the inhibitory activity of crude extracts from 10 species of marine invertebrates from the Cuban coastline (Table 1), against four neutral aminopeptidases: hAPN, hLAP and the malarial rPfA-M1 and rPfA-M17 was evaluated. These species belong to the phyla Mollusca, Poriphera, Echinodermata, and Cnidaria. The studies were performed using the substrate L-Leu-AMC at a final concentration equivalent to their respective Michaelis constant (K_M) values, determined in the present work, which were: 25.53 μ mol/L (hAPN), 238.00 μ mol/L (hLAP), 69.40 μ mol/L (rPfA-M1), and 43.33 μ mol/L (rPfA-M17), to avoid substrate competition during inhibition assays.

The initial evaluations performed allowed detection of inhibitory activity of the target enzymes in the aqueous crude extracts as follow: hAPN: Cenchritis *Isostichopus* badionotus; muricatus and hLAP: C. muricatus, Lissodendoryx (Lissodendoryx) isodyctialis, I. badionotus and Stichodactyla helianthus; rPfA-M17: C. muricatus, Echinaster (Othilia) echinophorus, I. badionotus, Physalia physalis, S. helianthus, and Bunodosoma granuliferum (Specific Inhibitory activity values are given in Table 2). Screening performed against recombinant malarial PfA-M1 (rPfA-M1) failed to detect any inhibitory activity from either extract (data not shown). Interestingly, in some assays with aqueous crude extracts, increased L-Leu-AMC hydrolysis rates over the control value were found instead of inhibitory activities of the tested enzymes. These results suggest either the presence of an activator of the target enzymes used in the assays or the presence of neutral aminopeptidase-like enzymes hydrolyzing L-Leu-AMC, in the corresponding aqueous extract.

Clarification of all the aqueous crude extracts with a 2.5 % TCA treatment increased, in all cases, the recovery of specific inhibitory activities as compared to their detection in positive crude extracts. The treatment also allowed the identification of inhibitory activities from species that were negative after screening using aqueous crude extracts. This result indicates that this clarification step was useful in the elimination of contaminants and/or induced dissociation from endogenous inhibitor-target complexes that did not allow the detection of inhibitory components in crude extracts. These 2.5 % TCA treated extracts were used to continue the inhibition studies.

Additionally, the enhanced activities over the control detected in some extracts were lost, in all cases, after the 2.5 % TCA treatments, suggesting susceptibility to the chaotropic agent and/or to the acidic pH, of the molecule(s) responsible for these effects in the concerned extract. To test for the possible presence of potential neutral aminopeptidaselike enzymes in aqueous crude extracts (which could be responsible for these observations), enzymatic assavs were performed using different amounts of the (in absence of the initial samples target aminopeptidases) in presence of L-Leu-AMC, in buffer A. A linear dependence of the initial rate versus the amount of crude extract in the assays was detected for the species Nerita peloronta, N. versicolor, L. (L.) isodictyalis, Tripneustes ventricosus, E. (O.) echinophorus, S. helianthus, B. granuliferum, and P. physalis. This allowed the determination of the specific enzyme activity in each case (Table 2). These results suggest the presence of neutral aminopeptidase activities in the aqueous crude

Table 2 — Summary of the screening of inhibitory activity against hAPN, malarial rPfA-M17 and hLAP.							
Species	Crude extracts			2.5% TCA cleared extracts			
	NA-like activity (x 10 ⁴ U/mg)	rPfA-M17 sIA (U/mg)	hLAPsIA(U /mg)	hAPNsIA (U/mg)	rPfA-M17 sIA (U/mg)	hLAPsIA (U/mg)	hAPNsIA (U/mg)
Cenchritis muricatus	ND	0.40	0.16	0.46	7.56	4.30	5.20
Nerita peloronta	6.95±2.84	ND	ND	ND	10.61	6.48	5.06
Nerita versicolor	12.56±1.70	ND	ND	ND	7.17	5.58	2.21
Lissodendoryx (Lissodendoryx) isodictyalis	1.51±0.49	ND	0.27	ND	256.13	312.01	171.92
Tripneustes ventricosus	6.39±0.07	ND	ND	ND	41.85	43.31	56.86
Echinaster (Othilia) echinophorus	58.87±12.62	0.11	ND	ND	9.10	7.85	10.82
Isostichopus badionotus	ND	0.18	1.24	1.56	55.83	27.86	33.81
Stichodactyla helianthus	9.63±2.68	0.55	0.68	ND	38.15	18.67	32.81
Bunodosoma granuliferum	39.44±5.45	0.19	ND	ND	5.45	4.08	4.31
Physalia physalis	2.04±0.52	0.40	ND	ND	19.21	11.00	13.13

Inhibitory activities found in aqueous crude and 2.5 % TCA extracts are expressed as specific Inhibitory Activity (sIA) in U/mg. One unit of enzyme activity was defined as the amount of enzyme needed to produce one arbitrary unit of fluorescence (AUF) per minute in the specified conditions and inhibitory activities are expressed per mg of corresponding extracts. The first column indicates tested species, the second column neutral aminopeptidase-like activity (NA), detected in aqueous crude extracts only, using L-Leu-AMC as substrate in Buffer A and the remaining columns refer to sIA on corresponding enzymes. ND: not detected.

extracts of all of these species. Characterizing of these in future is planned.

Dose response studies of inhibition of hAPN by 2.5 % TCA clarified extracts: Preliminary effect on PC3 and 3LL cancer cell lines

Table 3 summarizes the results of the evaluation of the inhibitory potential displayed by the 2.5 % TCA clarified extracts onnative hAPN. Once the crude extracts were treated with TCA at 2.5 % and extensively dialyzed, inhibitory activity of hAPN were detected in all treated extracts. All the clarified extracts inhibited hAPN in a dose dependent manner. The inhibition was characterized by a concave behavior observed in all the dose-response curves (Supplementary Fig. 1), indicating the reversibility of the inhibition and corroborating the presence of inhibitory molecules in the samples (and not artifacts interfering with the enzyme activity). IC_{50} values were in the range of 11.7-567.6 µg/mL. Inhibition of hAPN with IC_{50} values around or less than 100 µg/mL was detected in four of the ten species tested (L. (L.))isodictyalis, T. ventricosus, I. badionotus, and S. helianthus). IC₅₀ values for bestatin and amastatin were obtained in parallel as positive controls of inhibition (6.70±1.90 and 63.45±7.61 μg/mL, respectively). The IC₅₀ values for the TCA extract from L. (L.) isodictyalis and T.ventricosus were similar to that of bestatin. Once the inhibition of hAPN was corroborated, the effect of each treated extract on the viability of two cancer cell lines PC3 and 3LL was evaluated. As first step, we determined the neutral aminopeptidase activity in these cells in presence of L-Leu-AMC. PC3 and 3LL were

characterized by an activity of 87.78±3.14 and 27.53 ± 2.27 nmol/h/10⁶ cells, respectively. All of the treated extracts and bestatin used as positive control displayed a dose-response effect on PC3 and 3LL cell viability, with the best effects i.e. lowest IC₅₀ values (below $100\mu g/mL$) obtained was for L. (L.) isodictyalis (strongest effect, IC₅₀ value below of 5µg/mL, similar to bestatin effect), T. ventricosus, I. badionotus and S. helianthus, on both cell lines. Globally, the IC₅₀ values of cell viability were in good agreement with the IC₅₀ values on hAPN inhibition, including bestatin results (IC₅₀ on 3LL and PC3: 0.54 ± 0.01 and 3.15 ± 0.72 µg/mL, respectively). The rest of the TCA clarified extracts displayed moderately interesting IC₅₀ values in the range of 234.5-711.3 μg/mL.

Dose response studies of inhibition of malarial rPfA-M17 and hLAP: Preliminary effects on the growth of *P. falciparum* FcB1 strain

Table 4 summarizes the results of the evaluation of the inhibitory potential of the 2.5 % TCA clarified extractson malarial rPfA-M17 and native human LAP. We were able to detect inhibitory activities of both tested M17 aminopeptidases, in all TCA clarified extracts. These activities were characterized by a concave dose-response behavior corroborating the presence of reversible inhibitory molecules in the samples (Supplementary Fig. 2) with IC₅₀ values in the range of 15.3-509.2 µg/mL and 66.3-12 429.6 µg/mL, for rPfA-M17 and hLAP, respectively (Table 4). Similar behavior was detected in the presence of bestatin and amastatin as positive controls of inhibition vs both enzymes with IC₅₀ values

Table 3 — Summary of the IC₅₀ determination for the 2.5 % TCA clarified extracts against hAPN. Preliminary effects on growth of cancer cell lines 3LL and PC3.

Species	IC ₅₀ value vs hAPN (µg/mL)	IC_{50} value vs 3LL (µg/mL)	IC_{50} value vs PC3 (µg/mL)			
Cenchritis muricatus	450.20±77.40	214.00±46.50	352.90±65.00			
Nerita peloronta	237.80±20.90	273.30±78.80	299.10±31.70			
Nerita versicolor	370.00 ± 50.00	358.80±70.20	289.70±39.70			
Lissodendoryx (Lissodendoryx) isodictyalis	11.70±2.70	< 5.00	< 5.00			
Tripneustes ventricosus	25.00±3.10	39.90±2.00	77.00±3.90			
Echinaster (Othilia) echinophorus	198.20±27.20	265.70±29.60	405.60±50.40			
Isostichopus badionotus	69.70±10.00	57.10±2.70	83.10±3.00			
Stichodactyla helianthus	103.60 ± 20.60	110.80±13.20	58.10±7.50			
Bunodosoma granuliferum	567.60±88.00	786.80±37.10	711.30±29.30			
Physalia physalis	123.10±21.30	257.30±6.70	234.50±5.00			
Bestatin (positive control)	6.70±1.90	0.54±0.01	3.15±0.72			
Amastatin (positive control)	63.45±7.61	ND	ND			
Bestatin and amastatin were tested in parallel as positive control. ND: not determined.						

Species	IC ₅₀ value vs rPfA-M17 (µg/mL)	IC ₅₀ vs hLAP (µg/mL)	IC ₅₀ hLAP/IC ₅₀ rPfA- M17	IC ₅₀ Pf FcB1 (µg/mL)	
Cenchritis muricatus	113.40±3.00	341.00±110.00	3.00	> 400	
Nerita peloronta	22.20±2.70	329.50±100.00	14.80	291.80±38.50	
Nerita versicolor	207.00±30.60	12 429.60±633.00	60.00	325.50±0.80	
Lissodendoryx (Lissodendoryx) isodictyalis	27.30±9.40	66.30±27.60	2.42	2.60±0.60	
Tripneustes ventricosus	84.80±7.30	607.70±300.50	7.16	0.24±0.01	
Echinaster (Othilia) echinophorus	127.50±82.10	308.70±100.00	2.42	201.60±162.10	
Isostichopus badionotus	86.70±32.60	272.70±63.50	3.13	183.70±155.70	
Stichodactyla helianthus	15.30±6.20	234.90±34.60	15.35	> 400	
Bunodosoma granuliferum	509.20±100.90	1171.00±92.10	2.29	> 400	
Physalia physalis	293.70±100.00	550.00 ± 85.00	1.87	206.00±84.00	
Bestatin (positive control)	0.15±0.02	11.83±2.61	78.86	1.14±0.27	
Amastatin (positive control)	60.70±19.84	158.05±28.44	2.60	ND	
Bestatin and amastatin were tested in parallel as positive control. ND: not determined.					

Table 4 — Summary of the IC₅₀ determination for the 2.5 % TCA clarified extracts against rPfA-M17 and hLAP. Preliminary inhibitory effects on the growth of *Plasmodium falciparum* FcB1 strain

of: rPfA-M17: 0.15±0.02 and 60.70±19.84 µg/mL, respectively and hLAP: 11.83±2.61 and 158.05± 28.44 μ g/mL, respectively. We detected inhibition of rPfA-M17 with IC₅₀ values up to ~100 μ g/mL for 6 of the 10 extracts (those of C. muricatus, N. peloronta, L. (L.) isodictyalis, T. ventricosus, I. badionotus and S. helianthus). Comparing the inhibitions on rPfA-M17 and hLAP, in all cases the plasmodial enzyme was more susceptible than its human counterpart, with ratios of selectivity between 1.87 and 60 times. The most selective extract was from N. versicolor, an attractive situation even if its IC_{50} value on the malarial enzyme is moderate. Once the inhibition of rPfA-M17 was corroborated, we evaluated the effect of each extract on P. falciparum growth in cultures. With the exception of C. muricatus, S. helianthus and B. granuliferum, which displayed an IC₅₀ value greater than 400 μ g/mL, all of the treated extracts displayed a dose-response effect on FcB1 cells (IC₅₀ values in the range of 0.24-325.5 µg/mL). Bestatin tested in parallel as positive control displayed an IC₅₀ value of 1.14±0.27 µg/mL. The best effects were obtained for T. ventricosus (IC₅₀ value of 0.24 μ g/mL, one order lower than bestatin effect indicating strongest effect for this extract) and L. (L.) isodictyalis (IC₅₀ value of $\sim 2.8 \,\mu g/mL$, similar to bestatin effect).

Discussion

Peptidases are one of the most abundant groups of enzymes in living organisms, and they are involved in a wide variety of biochemical and physiological processes such as digestion, fertilization, growth, differentiation, cell signaling/migration, immunological defense, wound healing, and apoptosis. Peptidases are also involved in pathological outcomes in many human diseases including propagation of infectious agents and resulting pathogenesis and infectious diseases. For these reasons specific inhibitors of target peptidases are emerging as promising therapeutics uses both in the treatment of metabolic disorders (like type II diabetes mellitus. hypertension. cancers. inflammation. immunological and respiratory disorders and parasitic diseases)⁴⁰⁻⁴². Metallo-aminopeptidases infectious catalyze the cleavage of amino acids from the N-terminus of protein or peptide substrates. They belong to different families, the M1 and M17 families being the best characterized^{1,2,6}. Among M1 family enzymes, APN has been extensively investigated in mammals, with recent important findings about its structure-function relationships^{43,44} and its implication tumorigenesis, immune system and in pain management^{4,8,9}. PfA-M17 from *P. falciparum* is one of the best characterized M17 family members and current target for the development of new antimalarials because of the importance of this enzyme for parasite survival^{25,27,45-49}

Marine invertebrates are an abundant source of bioactive molecules such as toxins^{50,51}, peptidases⁵² and peptidase inhibitors of different mechanistic classes. They are characterized by a great diversity of chemical structures, high potency and diverse specificity including metallo inhibitors^{28,29,53-56}. These biomolecules are involved in nutrition, reproduction, and communication processes. Additionally, most invertebrates (like sponges, bryozoans, tunicates,

among others) lack morphological defense structures. These biomolecules are also part of mechanisms related with protection against predator, infection, and competition⁵⁷. Currently, identification of APN inhibitors from marine sources remains scarce with only two reports of molecules behaving as inhibitors of the porcine enzyme^{28,29} and inhibitory activity reported from one ascidian species⁵⁸. However, marine peptidase inhibitors that may interact with human and plasmodial M1 and M17 enzymes have not yet been described.

As result of the screening of inhibitory activity of neutral aminopeptidases of biomedical relevance in the present contribution, the present findings are the first to show the presence of inhibitory activity of human APN in the 2.5 % TCA clarified and dialyzed extracts from С. muricatus, Ν. peloronta, N. versicolor, L. (L.) isodictyalis, T. ventricosus, E. (O.) echinophorus, I. badionotus, S. helianthus, B. granuliferum, and P. physalis. In all the cases, inhibition was highly dependent on the dose. The higher inhibitions characterized by promising IC₅₀ values (around 100 µg/mL or lesser) were obtained for L. isodictvalis, T. ventricosus, S. badionatus, and S. helianthus. The IC₅₀ values for almost all the ten species were lower than the data previously described for both TCA and heat clarified extracts from the ascidian Diplosoma listerianum vs pAPN in similar assay conditions but in the presence of susbtrate Leu-pnitroanilide⁵⁸. Additionally, for *L. isodictyalis* and S. helianthus inhibition of porcine pancreatic elastase (EPP) and human neutrophil elastase (HNE), by heat treated extracts have been recently reported⁵⁹. HNE is a serine peptidase that contributes to the onset and progression of many inflammatory diseases with severe impact on organ tissue integrity (like pulmonary emphysema, rheumatoid arthritis, and cystic fibrosis) that is now viewed as a promising target for the development of new therapeutics⁶⁰. The EPP inhibition was characterized by IC₅₀ values in the same order of hAPN inhibition identified in the present contribution for these two species. Findings show that that EPP/HNE and hAPN belong to different peptidases clans with completely different catalytic mechanisms^{1,2}. Further, the clarification treatments employed in each work were different. It is very likely that the molecules responsible for the hAPN and EPP inhibition in L. isodictyalis and S. helianthus are not the same molecular entities. The present results support these two species as potential new sources of inhibitors for biomedical relevant

human peptidases like hAPN and HNE. All the clarified extracts had effect on 3LL and PC3 cancer cell characterized by the presence of APN activity reported in the present study. The higher effects with IC_{50} values below 100 µg/mL were observed for the same species that were also displaying the strongest hAPN inhibition. In particular, the identification of an IC_{50} value under 5 µg/mL for *L. isodictyalis* extract vs both cancer cells lines, similar to the effect displayed by bestatin (a pure compound), indicates that we are in presence of a potentially very good extract for the isolation of hAPN inhibitors, as defined by Mambu and Grellier⁶¹.

A wide variety of anticancer activities and isolated compounds with a large diversity of chemical structures (alkaloids, steroids, macrolides, polyketides, depsipeptides among others) have been identified from a large diversity of marine organisms^{57,62,63}. Some are currently in clinical and late preclinical development⁶⁴. To the best of our knowledge, this is the first time concomitant indications of activity on hAPN target are also provided.

As a second part of the screening of inhibitory activity for neutral aminopeptidases performed in the present study, inhibition of rPfA-M17 in all the 2.5 % TCA cleared extracts assayed was identified. The inhibition was also dose-dependent, with six species with IC₅₀ values lower than 100 μ g/mL. All the species also inhibited the human LAP, but the inhibition was higher for the malarial enzyme indicating certain degree of selectivity of the entities responsible for rPfA-M17 inhibition. These results constitute the first report of inhibition of M17 (human and plasmodial) by marine enzymes invertebrates' species extracts. With the exception of C. muricatus, S. helianthus and B. granuliferum, all the treated extracts showing inhibition of rPfA-M17 also have an inhibitory effect on the growth of FcB1 strain of P. falciparum. The extracts from L. isodictyalis and T.ventricosus were characterized by the most promising activity in terms of IC_{50} value under 3 µg/mL. These results indicate that we are in presence of two very good extracts as defined Mambu and Grellier⁶¹. Particularly attractive are the T. ventricosus extracts that displayed an effect 300 times more potent on FcB1 strain of P. falciparum regarding the human cancer cells, indicating parasite effect specificity. Additionally, this effect of T. ventricosus extract on FcB1 strain was stronger than bestatin effect (a pure compound). Due to the fact that IC_{50} for *L. isodyctialis* and

T. ventricosus is lower on FcB1 strain of *P. falciparum* than on PfA-M17 enzymes, it is very likely that these extracts may contain not only PfA-M17 inhibitors but also other compounds active on other malarial targets. For example, *L. isodyctialis*, inhibition of subtilisin from *Bacillus licheniformes* with IC₅₀ value of 3 μ g/mL has been recently described⁵⁹. Another interesting result is the selectivity for rPfA-M17 regarding hLAP of *N. versicolor* extract showing effects on parasite growth with IC₅₀ in the same order of enzyme inhibition, indicating that this species is also attractive.

More than 50 natural compounds isolated from marine organisms have been described with antimalarial activity, most of them from sponges. The molecular mechanism of the effect has been identified for only two of these natural compounds^{65,66}. Sponges are an abundant source of compounds with activity against protozoa responsible for the so called "neglected diseases" (such as leishmaniasis, amoebiasis, trichomoniasis, African sleeping sickness and Chagas disease)⁶⁶. The present results are similar in that good antimalarial activity vs FcB1 strain of P. falciparum (IC₅₀ of 2.8 µg/mL) in the 2.5 % TCA extract from the Caribbean sponge L. (L.) isodictyalis, which also displayed a good inhibition of rPfA-M17 (IC₅₀ 27.3 µg/mL) was identified. For this species, Mendiola et al.⁶⁷ previously identified inhibition of F32/Tanzania P. falciparum strain in an aqueous crude extract prepared without any treatment, with a MIC of 50 μ g/mL. In the same work the researchers described inhibition of the same F32/Tanzania P. falciparum strain by aqueous extracts from three species of Ascidians (Phallusia nigra, Ascidia sidneiensis and Microcosmus goanus) and two echinoderms (Holothuria sp. and Ε. (O.)echinophorus). The last species was also tested in the present contribution but the aqueous extract was characterized by a high aminopeptidase activity in presence of L-Leu-AMC at pH 7.5 that did not allow the detection of inhibition of enzymes tested. The TCA treated extract showed that inhibition of rPfA-M17 and the effect of FcB1 P. falciparum cells (IC₅₀ 232.22 μ g/mL) was higher than the one reported for the aqueous crude extract of this species (500 μ g/mL) by Mendiola *et al.*⁶⁷, probably due to a more clarified extract in the present study. The results of the present study supports that sponges as well as marine invertebrates from other Phyla such as mollusks, echinoderms, and cnidarians are a good and unexplored source of potential anticancer and antimalarial compunds associated with the inhibition of neutral aminopeptidases from M1 and M17 families, involved in these human pathologies.

Conclusion

The present study constitutes the first report of inhibitory activity of human and plasmodial neutral aminopeptidases in aqueous extracts from marine species. The results strongly suggest that all of the species in this study have potential as sources of inhibitors of hAPN that could be applied in future studies on cancer models involving this enzyme. L. isodictyalis, T. ventricosus, S. badionatus and S. helianthus are the most promising in terms of IC_{50} vs both enzyme and cell models. Although all the species extracts inhibited rPfA-M17, the most promising of them for future applications in malaria models are L. isodictyalis and T. ventricosus characterized by the higher effects on growth of FcB1 P. falciparum strain cells and with effects on rPfA-M17. The potentialities of the remaining tested species in the malaria field will depend whether the molecules responsible for the inhibition detected vs hAPN and rPfA-M17 are different, and in all cases, whether the purified compounds are not toxic vs normal human cells i.e. vs host. The present results strongly support that marine fauna, in particular the invertebrates, are an important and unexplored source of inhibitors of metallo-aminopeptidase with promising applications in cancer and malaria treatment.

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