Bromelia karatas and Bromelia pinguin: sources of plant proteases used for obtaining antioxidant hydrolysates from chicken and fish by-products

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Abstract
In the present study, we evaluated new sources of plant proteases from fruits of Bromelia karatas (BK) and Bromelia pinguin (BP) to obtain antioxidant hydrolysates/bioactive peptides (BPs) derived from chicken by-products (CH) and fish by-products (FH). The profile of the peptides was identified by reverse-phase high-resolution liquid chromatography (RP-HPLC) and the size weight distribution by molecular exclusion chromatography (SEC). The hydrolysates obtained with BK in both sources of by-products showed greater antioxidant capacity compared to those obtained with BP, presenting similar or higher values when compared to a commercial plant enzyme. The use of new sources of plant proteases allowed to obtain hydrolysates of hydrophilic character with a high percentage (> 50%) of peptides with molecular weights < 17.5 kDa from chicken and fish by-products. Therefore, based on the results obtained in antioxidant capacity it is possible to consider the hydrolysates as potential ingredients, food additives, and pharmaceutical products.

Keywords: Antioxidant capacity, Bioactive peptides, By-products, Hydrolysates, Proteases

Introduction
Antioxidants are bioactive compounds of vital importance in food processing and human health [1], they can be obtained from different food matrices of plant and animal origin. Free radicals and oxidative stress are involved in the pathogenesis of a wide variety of conditions that include inflammatory diseases, cancer, atherosclerosis, diabetes mellitus, neurodegenerative diseases, HIV/AIDS as well as stimulating premature skin aging [2, 3]. Up to date, a large number of studies have been published about the antioxidant properties of hydrolysates or food-derived peptides [2], which are considered health-promoting biomolecules with medical applications, besides to possess a nutraceutical potential and application in the food industry [4]. BPs are the usual name of short amino acid sequences (AA), which could be generated from the protein hydrolysis obtained from different sources, but mainly of animal origin such as beef, pork, lamb, poultry, duck and various species of marine organisms [5]. The BPs are conformed by chains of 2–30 AA with molecular weights less than 10 kDa [6, 7]. BPs has a positive impact on the functions or conditions of the body and influences human health [8].

It is important to use proteins of animal origin from different species to obtain BPs [9, 10]. In the last few years, the interest for the production of these biomolecules
from by-products such as bird feathers, blood, viscera, skin, among others has increased [7, 11, 12], which represent a valuable source of protein and other essential nutrients [13]. In vitro studies have evaluated the antioxidant capacity and activity of different hydrolysates/peptides derived from poultry, chicken and ducks by-products [1–3, 14–16] and fish by-products [17–22].

Enzymatic hydrolysis by the action of different proteases of animal origin (i.e. trypsin and pepsin), microbial (i.e. alcalase and neutrase) and plants (i.e. bromelain and papain), has been the technological process commonly used for the production of hydrolyzates/peptides derived from food [4, 26–29]. The proteases of plant origin are relatively limited to the use of bromelain and papain, these proteases have been employed to obtain hydrolysates and bioactive peptides from fish by-products [17, 23, 24] and blood from livestock animals [23, 24]. In recent years, research on new sources of proteolytic enzymes has been encouraged, as well as their possible applications in industry. The obtaining of new sources of plant proteases is increasing due to their low production costs, relative abundance in some plants and attractive biochemical characteristics [25], such as their stability over a wide range of pH and temperature [26]. The fruits of *Bromelia pinguin* “guamara” and *Bromelia karatas* “cocuixtle” are presented as an alternative to obtaining plant proteases [4, 26–29]. However, little information exists on the application of these sources of proteases in different food matrices to obtain protein hydrolysates. In this regard, the objective of this work is to obtain and characterize peptides with antioxidant capacity from the enzymatic hydrolysis of chicken by-products and fish by-products using new plant sources of proteases from *B. karatas* “cocuixtle” and *B. pinguin* “guamara.”

**Materials and methods**

**Reagents**

ABTS (2,2’-azinobis (3-ethyl benzothiazoline-6-sulfonic)), TPTZ (2,4,6-tri (2-pyridyl)-s-triazine), DPPH (2,2-diphenyl-1-picrylhydrazyl), Trolox (6-hydroxy-2,5,7,8-tetramethylchroman-2-carboxylic acid), acetonitrile, trifluoroacetic acid was obtained from Sigma Aldrich (St. Louis, MO) and TNBS (2,4,6-trinitrobenzene sulfonic acid) was acquired from Thermo Fisher Scientific.

**Organic material**

The fruits of *B. karatas*, *B. pinguin*, chicken by-products (viscera 44.5% w/w, heart 44.5% w/w and blood 11% w/w) and fish by-products *Centropomus* ssp. (Scales 50% w/w and skin 50% w/w) were collected from a local market at Tepic, Nayarit, Mexico. The organic material was packed in polyethylene bags and transported to the integral food research laboratory of the Instituto Tecnológico de Tepic, Nayarit, Mexico. Organic material was frozen at −80 °C and then lyophilized. (70020, Labconco Corporation, Kansas City, MI, USA).

**Extraction, semi-purification and proteolytic activity of *B. pinguin* (BP) and *B. karatas* (BK) proteases**

Enzyme extraction and semi-purification of the fruits from *B. pinguin* and *B. karatas* was carried out using the methodology used by García-Magaña et al. [30]. The semi-purified enzyme extracts were centrifuged at 6000×g for 30 min at 4 °C, the recovered supernatant was lyophilized and then stored for further analysis.

**Proteolytic activity**

The proteolytic activity of the semi-purified protease extracts was determined by the methodology described by Natalucci et al. [31] with slight modifications by García-Magaña et al. [30], using egg albumin as a substrate. 1.1 mL of egg albumin solution (1 g/100 mL) was mixed with phosphate buffer (pH 6.0, 12.5 mM of L-cysteine) and 0.1 mL of protease extract (2.6 mg of the lyophilized enzyme in 1.5 mL of buffer) at 37 °C. The reaction was stopped by adding 1.8 mL of trichloroacetic acid (50 g/L). The solution was centrifuged (14,000×g for 10 min at 4 °C; Hettich, Mikro 200R, Tuttinglen, Germany) and the absorbance of the supernatant was measured at 280 nm (JENWAY, 6705, Bibby Scientific Limited, OSA, United Kingdom). A calibration curve with tyrosine (3 M) was used to determine the specific activity expressed as tyrosine units (UT), defined as the change in tyrosine concentration/mg protein/min at 37 °C. Protein concentration was determined by Bradford [32] and expressed in mg/mL. The specific activity was determined with the following equation:

\[
A_{U(37{\degree}C)} = \frac{\Delta \text{tyrosine concentration}}{(t^* V_{\text{enzyme}} * C_{\text{enzyme}})} \left( \frac{V_{\text{reaction}}}{V_{\text{enzyme}}} \right)
\]

where, Δ tyrosine concentration, is the difference in absorbances obtained between the test sample and the control; \(V_{\text{enzyme}}\) volume of the enzyme solution (0.1 mL); t, reaction time (20 min); \(V_{\text{reaction}}\) volume used in the reaction (3 mL); \(C_{\text{enzyme}}\) enzymatic concentration obtained by the Bradford method.

**Protein hydrolysates from chicken (CH) and fish by-products (FH)**

The lyophilized chicken and fish by-products were resuspended (1:20, w:v) in 0.2 M sodium phosphate buffer pH 6.5, homogenized for 2 min and subjected to heat treatment at 100 °C for 15 min. Subsequently, the protein substrates were individually hydrolyzed with the
enzymes extracted from *B. pinguin* (BP), *B. karatas* (BK) and the bromelain (BRO) was used as a control enzyme, adding 1 mL of enzyme solution (100 µg of lyophilized enzyme/mL) re-suspended in 0.1 M phosphate buffer, pH 6.5 containing, 5 mM l-Cysteine) at the conditions of 40 °C for 30 min at pH 6.5, 25 °C for 4 h and pH 6.5 and 37 °C for 4 h and pH 7.0, respectively. After this time, the samples were immediately heated at 100 °C for 15 min to stop the hydrolysis process. The hydrolysates were centrifuged at 14,000×g at 4 °C for 10 min and then filtered with 0.45 µm membranes (Millipore, MF-Membrane Filters HAWP).

**Soluble protein content**
The protein content was determined by the Bradford method [32], from a CH/FH solution of 30 mg/mL (w/v). 0.1 mL of hydrolysate solution was placed with 1 mL of Bradford reagent, allowed to stand for 10 min in darkness and the absorbance was measured on a UV–VIS spectrophotometer at 595 nm. Bovine serum albumin (BSA) was used as a standard. The results were expressed in mg/mL.

**Yield**
The hydrolysates were frozen at −80 °C and then lyophilized. The yield percentage of hydrolysates was calculated by the equation proposed by Chou et al. [2]:

\[
\text{Yield} \ (\%) = \left( \frac{\text{lyophilized hydrolysates} \ (g)}{\text{lyophilized by} - \text{product} \ (g)} \right) \times 100
\]

**Degree of hydrolysis**
The degree of hydrolysis (DH) was evaluated based on the quantification of the primary free amino groups liberated during hydrolysis by the method proposed by Adler-Nissen [33] modified by Tovar-Pérez et al. [34]. Briefly, we mixed 120 µL of hydrolysates, 1 mL of 0.2 M phosphate buffer at pH 8.2 and 1 mL solution of 0.1% 2,4,6-trinitrobenzene sulfonic acid (TNBS). The mixture was allowed to incubate for 60 min at 50 °C. The reaction was finished by adding 2 mL of 0.1 N HCl. The absorbance of the solution was measured at a wavelength of 340 nm. Blank was prepared in the same way. The degree of hydrolysis (%) was estimated with the following equation:

\[
\text{DH} (%) = \left( \frac{\text{NH}_{2t} - \text{NH}_0}{\text{NH}_{2\max} - \text{NH}_0} \right) \times 100
\]

where: \( \text{NH}_{2t} \) = concentration of free amino groups after hydrolysis with *B. pinguin*, *B. karatas*, and bromelain. \( \text{NH}_0 \) = concentration of free amino groups of chicken and fish by-products (without hydrolyzing). \( \text{NH}_{2\max} \) = concentration of free amino groups after total hydrolysis, which was estimated by hydrolysis with 6 N HCl at 100 °C for 24 h. The concentration of free amino groups (NH3) was expressed in mM eq l-Leucine/g protein with a previously elaboration of a standard curve of l-Leucine (0–2.5 mM).

**Antioxidant capacity (AoxC)**

**Abts**
The ability of the peptides to capture the ABTS+ radical (2,2’-(azinobis(3-ethyl benzothiazoline-6-sulfonic acid) was evaluated by the method proposed by Re et al. [35]. A solution of 7 mM ABTS was prepared in sodium phosphate buffer (0.1 M, pH 7.4) with potassium persulfate (2.45 mM). The reagent was adjusted to an absorbance of 0.8. Thereafter, 1800 µL of ABTS solution and 200 µL of hydrolysates (30 mg/mL) were placed and then allowed to stand for 7 min in the dark. Absorbances were read at 730 nm in a spectrophotometer. The phosphate buffer was used as a blank and Trolox (0.6 µM) as a standard solution. The results were expressed in mM equivalent to Trolox (ET)/mL.

**Dpph**
The ability of the peptides to capture the DPPH radical (2,2-diphenyl-1-picrylhydrazyl) was analyzed according to the methodology described by Park and Chin [36], with certain modifications according to Chou et al. [2]. The solution was obtained by mixing 0.8 mL of hydrolyzate (10 mg/mL) with 0.2 mL of 1 mM DPPH (D9132, Sigma Co.) dissolved in 95% ethanol (v/v), the mixture was kept in dark for 30 min at room temperature and then centrifuged at 2000×g for 5 min at 4 °C. The absorbance of the supernatant was measured in a SpectraMax M3 microplate reader at 517 nm. Distilled water was used as blank and Trolox (600 µM) as a standard solution. The results were expressed in µM ET/mL.

**Frap**
The ability of peptides to reduce ferric ion (Fe3+) was evaluated by the method of Benzie & Strain [37]. The FRAP reagent was prepared from 100 mL of sodium acetate buffer (300 mM, pH 3.6) with 10 mL of TPTZ solution (2,4,6-tris(2-pyridyl)-s-triazine) in 40 mM HCl) and 10 mL FeCl3 (20 mM) and then incubated at 37 °C for 30 min in the dark. Next, 100 µL of hydrolysates (30 mg/mL) reacted with 1000 µL of FRAP reagent, which was incubated at 37 °C for 30 min in the dark. The absorbance was read on a spectrophotometer at 595 nm. Distilled water was used as blank and Trolox as standard (260 µM). The results were expressed in µM ET/mL.
Determination of the peptide profile in the protein hydrolysates

Reversed-phase high-pressure liquid chromatography (RP-HPLC)

The peptide profile of the hydrolysates (CH and FH with BP, BK, and BRO) was measured by RP-HPLC following the methodology of Huang et al. [38], with some modifications. The analysis was performed with a reverse phase HPLC equipment (1100 series; Agilent Technologies Japan Ltd., Tokyo, Japan), the separation was carried out on a Discovery-C18 column (250 × 4.6 mm, 5 μm particle size, 180-Å pore size) of Supelco Inc. (Bellefonte, PA) with a solvent flow rate of 0.75 mL/min and an injection volume of 30 μL. Peptides were eluted from a non-isocratic gradient using solvent B (0.05% TFA in acetonitrile) and solvent D (0.05% TFA in Milli-Q water) for 40 min. 0–30 min (100% D), 30–35 min (60% B, 40% D), 35–40 min (100% B), 40 min (100% D). The peptide profile was monitored at 214 nm.

Size exclusion chromatography (SEC)

The molecular weight distribution (MW) of the CH and FH peptides was determined by Size exclusion chromatography (SEC) using a Varian HPLC system equipped with a diode array detector (DAD) (Varian™ ProStar) and Galaxy™ software (Varian, version 1.9.302.952) following the methodology used by Román-Gavilanes et al. [39]. 20 μL of the sample was injected at a concentration of 5 mg/mL. The mobile (isocratic) phase consisted of 150 mM phosphate buffer at pH 7 and a flow rate of 0.4 mL/min. A molecular exclusion column (Agilent BioSEC-™) of 4.6 mm internal diameter and 300 mm in length was used. The absorbance was monitored at 254 nm. The molecular size of the samples was determined based on their retention times using a peptide standard (BIO-RAD, 151–1901), consisting of thyroglobulin (670 kDa), gamma globulin (158 kDa), ovalbumin (44 kDa), myoglobin (17 kDa), and Vitamin B12 (1.35 kDa).

Statistical analysis

Statistical analysis of experimental data was performed using a one-way ANOVA. Differences between means were evaluated using the Tukey–Kramer multiple comparison test and were considered significant when P<0.05. Statistical analysis was performed using the NCSS 2007 software (NCSS LLC, Kaysville, UT).

Results

Proteolytic activity

The specific enzymatic activity of the semi-purified extracts of B. karatas and B. pinguin was 6.24 ± 0.01 AU/mg and 43.76 ± 1.60 AU/mg, respectively. These results confirmed that the extracted plant proteases are active.

Protein content, yield and degree of hydrolysis of CH and FH

In Table 1, we have shown the results of the protein content, yield and degree of hydrolysis of CH and FH with enzymes extracted from BP and BK, compared to BRO. The protein content of FH with BP and BK showed no significant differences (P>0.05) with those achieved with BRO, presenting a higher content (P<0.05) compared to those found with CH-BP and CH-BK. Regarding the yield (Table 1), the CH with BK and BRO presented significant differences (P<0.05) with a higher yield (65.4% and 65.1%, respectively) than the other hydrolysates.

The degree of hydrolysis (Table 1) of the CH-BP showed the highest value (P<0.05) with 22.9%, which may be related to the high enzymatic activity of BP.

Antioxidant capacity (AoxC) of the protein hydrolysates

The AoxC of CH and FH by means of the ABTS, DPPH and FRAP tests are presented in Fig. 1, where it is observed that the AoxC was affected by the type of substrate (chicken and fish by-products) and the type of plant proteases (BP, BK and BRO). Concerning the ABTS assay, the CH and FH with BK had the

Table 1  Protein content, yield and degree of hydrolysis of by-products hydrolysates

| Hydrolysates | Enzyme | Protein content (mg/mL) | Yield (%) | Degree of hydrolysis (%) |
|--------------|--------|------------------------|-----------|-------------------------|
| CH           | BP     | 0.59±0.04 a            | 46.8±0.4 a| 22.9±1.0 a              |
| FH           |        | 1.18±0.37 c            | 54.3±1.3 ab| 2.2±0.4 b               |
| CH           | BK     | 0.79±0.10 ab           | 65.4±4.7 d| 8.8±0.6 c               |
| FH           |        | 1.08±0.15 bc           | 49.7±2.6 ab| 2.2±0.3 b               |
| CH           | BRO    | 1.32±0.12 c           | 65.1±5.2 d| 5.1±0.3 d               |
| FH           |        | 1.34±0.16 c           | 54.8±1.4 bc| 2.8±0.6 b               |

Protein content, yield (%), degree of hydrolysis (%) of chicken by-products hydrolysates (CH) and fish by-products hydrolysates (FH) with enzymes of B. pinguin (BP) and B. karatas (BK). The means ± standard deviations (n = 3). The different letters mean significant differences (ANOVA, Tukey–Kramer, P<0.05) between treatments.
highest AoxC ($P < 0.05$) with 36.98 and 38.03 mM ET/mL respectively. The use of BP and BRO generate CH and FH with similar AoxC: ABTS ($P < 0.05$). The highest DPPH radical scavenging activity was obtained for the FH with BP and BRO (483.21 µM ET/mL and 454.84 µM ET/mL). No significant differences ($P < 0.05$) were observed with respect to the AoxC: DPPH of the CH (BP, BK and BRO). On the other hand, BRO increased the chelation capacity of ferrous ions only in the FH, presenting the highest AoxC: FRAP values ($P < 0.05$) of 1198.5 µM ET/mL, followed by CH-BK with 817.4 µM ET/mL. The use of plant proteases (BP and BK) increased the AoxC: FRAP values ($P < 0.05$) in the CH, showing significant differences ($P < 0.05$) with BRO (Fig. 1).

Profile of peptides from hydrolysates

Reversed phase-HPLC (RP-HPLC) can be used to separate peptides according to their hydrophobicity. HPLC profiles of the CH and FH peptides with BP, BK and BRO are shown in Figs. 2 and 3, respectively.

According to the chromatograms obtained, the use of plant proteases (BP, BK and BRO) generates of hydrophilic peptides. Peptides of CH with BP, BK and BRO started to elute from the column after the first 5 min, which indicates a lower net hydrophobicity and, therefore, a weak binding to the column. Most of the CH peptides eluted in the first 20 min when the mobile phase was formed 100% by Milli-Q water with 0.05% TFA (Fig. 2). Although, the peptides profile of CH predominates in hydrophilic peptides, it is also observed that CH-BP (Fig. 2a) generates a few hydrophobic peptides eluted in the 30–35 min.

Regarding the HPLC profiles of the FH with BP, BK and BRO, the peptides show mainly a hydrophilic character, this could be because most of the peptides eluted in the first 20 min (Fig. 3). FH-BK (Fig. 3b) and FH-BRO (Fig. 3c) present similar profiles, with higher peptide peaks compared to FH-BP (Fig. 3a) in the first 20 min. FH-BP and FH-BRO generates a few hydrophobic peptides eluted in the 30–35 min (Fig. 3).

Distribution of molecular weights (MW) of peptides from hydrolysates

The SEC process is based on the fractionation of peptides according to the retention time of the molecules in the particles of the stationary phases, in which the molecules separate from each other according to their molecular size [39]. The distribution of the MW of the CH and FH obtained with the proteases of BP and BK, are shown in Figs. 4 and 5.

According to the chromatograms obtained, the MW distribution of the peptides of CH and FH, is dependent on the plant protease used (BP, BK and BRO). The use of BRO in CH, showed a higher proportion of low MW (<1.35 kDa) eluted peptides compared to CH (Fig. 4c), compared to the chromatograms obtained with BP (Fig. 4a) and BK (Fig. 4b), which have a higher elution of peptides of 670–44 kDa. CH-BP and CH-BK have similar peptide profiles of 17–1.35 kDa and <1.35 kDa. Regarding FH, a similar MW distribution is observed with the use of BP (Fig. 5a), BK (Fig. 5b) and BRO (Fig. 5c), according to the chromatograms there is a greater peptide elution low MW (17–1.35 kDa; <1.35 kDa).

Respect to the percentage of the area of the chromatograms obtained in the CH, more than 50% of the peptides has MW below 17 kDa (Table 2). CH presents higher percentages of peptides 17–1.35 kDa compared to FH.
CH-BP (24.4%) and CH-BK (22.5%) have similar content of peptides < 1.35 kDa. Regarding the FH, the percentage of peptides < 1.35 kDa increases, compared with CH (Table 2). FH-BP presents the highest percentage of peptides < 1.35 kDa (51.3%), followed by FH-BRO (48.9%) and FH-BK (46.9%).

**Discussion**

The fruits of *B. karatas* and *B. pinguin* have been characterized as promising sources of plant proteases [25, 28–30, 40]. One of the important points of this research is the incorporation of these plant enzymes in food technology processes and the obtaining of protein hydrolysates from food by-products and their comparison with a commercial plant enzyme (bromelain). The use of enzymes for the production of hydrolysates has been the subject of research for years [41]. The enzymes extracted from BP and BK show differences between their enzymatic activities, which coincide with what was previously reported by García-Magaña et al. [30] (5.08 ± 0.70 for BK proteases and 43.58 ± 0.40 for BP proteases). BP proteases showed higher activity compared to BK proteases, this behavior is similar to the already reported by Aguilera-Aguirre et al. [40] and Meza-Espinoza et al. [25], and can be attributed to the presence of different types of cysteine proteases in both fruits [26, 30, 42]. In the same way, it has been reported that the different genetic

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**Fig. 2** Peptide fraction profiles of chicken by-products hydrolysates. Peptide fraction profiles corresponding to chicken by-products hydrolysates (CH) by BP (a), BK (b) and BRO (c) using reversed-phase HPLC at 214 nm.
characteristics of each fruit species synthesize endopeptidases with different structures and catalytic activities [25].

The application of BP and BK proteases in the hydrolysis of chicken by-products and fish by-products gave us variable results to their protein content, yield and degree of hydrolysis. Among the different biochemical parameters, the degree of hydrolysis is one of the most important characteristics, since it directly influences the length of the peptide, its nutritional and functional properties, as well as it is directly related to their solubility and digestibility [41]. Meza-Espinoza et al. [31] reported that BP proteases generate a greater degree of hydrolysis in different protein sources (milk, soy protein, and egg ovalbumin) compared to BK. Few references regarding the use of plant proteases to obtain hydrolysates from animal by-products can be found. It has been reported that the use of papain to hydrolyze by-products of bovine origin showed 6% of DH [43]. Likewise, the use of papain (EC 3.4.22.2) and bromelain (EC 3.4.22.32) with DH of 15% for obtaining hydrolysates of fish by-products with bioactive and functional properties [17].

The yield depends on the nature of the proteases used, the amount of proteins present in the substrate, the hydrolysis conditions, the concentration of the proteases and the hydrolysis time [4, 17]. According to the results obtained, we observed a greater affinity of the proteases of BP, BK and BRO to hydrolyze chicken by-product

![Fig. 3 Peptide fraction profiles of fish by-products hydrolysates. Peptide fraction profiles corresponding to fish by-products hydrolysates (FH) with BP (a), BK (b) and BRO (c) using by reversed-phase HPLC at 214 nm](image)
Fig. 4 Chromatographic profiles of chicken by-products hydrolysates. Chromatographic profiles of chicken by-products hydrolysates (CH) with BP (a) and BK (b) proteases and BRO (c). The graph shows the signal from the UV monitor at 254 nm.

Fig. 5 Chromatographic profiles of fish by-products hydrolysates. Chromatographic profiles of fish by-products hydrolysates (FH) with BP (a) and BK (b) proteases and BRO (c). The graph shows the signal from the UV monitor at 254 nm.
proteins compared to fish by-products presenting significant differences ($P < 0.05$) and resulting in the CH obtained a higher yield and DH compared to the FH.

A degree of hydrolysis greater than 10% generates peptides that have high solubility and superior bioavailability. On the other hand, a low degree of hydrolysis (1–10%) improves the functional properties of food products such as foaming power and emulsifier [41].

The biological effect of peptides isolated from by-products derived from the meat industry has been studied in vitro for their antihypertensive and antioxidant activity, which provides a basis for their use as bioactive peptides [5], and their activity is mainly related to structure and nature of the AA that conformed. An example is its AoxC, which is mainly attributed to the presence of aromatic AA (tyrosine, tryptophan, and phenylalanine) in its terminal chains, since they have the ability to efficiently neutralize free radicals, acting as electron donors/hydrogen due to the presence of phenolic, indole and imidazole groups; in the same way, they can reduce $\text{Fe}^{3+}$ ions to $\text{Fe}^{2+}$ and chelate $\text{Fe}^{2+}$ as well as $\text{Cu}^{2+}$ ions [43, 44]. It has been reported that the ferric ion (FRAP) can be oxidized by hydrophilic compounds, just like the ABTS radical [45]; although it has also been reported that ABTS may be neutralized by some compounds, poorly soluble or lipophilic [46]; while DPPH detects hydrophilic compounds in greater presence [47]. Therefore, the results of AoxC are directly related to the characteristics of hydrolysates/peptides derived from chicken by-products [1], antihypertensive character (Cys-Gly-Asp-Pro-Cys-Gly) [2], antioxidant capacity (Ser-Asn-Leu-Cys-Arg-Pro-Cys-Gly) [48], and neutral nature (His, Gly, Gln, Ser and Thr) in the C-terminal position. Of this AA, His, Lys and Arg are categorized in the AA group with strong AoxC together with hydrophobic AAs such as Met, Trp, and Tyr according to a study where they categorized 20 AA in relation to their AoxC [48]. Likewise, it has been suggested that a close relationship exists between polar peptides and the type of amino acid positioned in their side chains [49]. AA sequences of peptides derived from animal by-products having polar character AA in their side chains derived from poultry by-products with antioxidant capacity (Ser-Asn-Leu-Cys-Arg-Pro-Cys-Gly) [1], antihypertensive character (Cys-Gly-Lys-Pro, Ser-Gly-Arg, Cys-Thr-Ser-His, Val-Lys-Lys-Thr) [50] and (Ala-Ile-Tyr-His, Leu-Arg-Lys-Gly-Asn-Leu-Glu) [51] as well as by-products of marine origin with antioxidant capacity (Trp-Glu-Gly-Pro-Lys; Gly-Pro-Pro; Gly-Val-Pro-Leu-Thr) have been determined [52].

Peptides with MW < 10 kDa are considered bioactive which can generate multiple biological activities, thus the interest of incorporating these compounds in the food area, as well as in the pharmaceutical industry. Chicken liver hydrolysates with pepsin showed MW between 4.6 and 10 kDa [2], which have been shown to have multifunctional effects on in vivo systems with murine models induced to liver damage favoring (1) the regulation of lipid homeostasis, (2) decrease in lipid peroxidation through the improvement of antioxidant enzyme activities (3) improvement of gene expression related to the pathologies evaluated and (4) decrease in hepatic inflammatory responses [15, 53]. Based on the results obtained, the application of BP and BK proteases in chicken and fish by-products presenting significant differences ($P < 0.05$) and resulting in the CH obtained a higher yield and DH compared to the FH.

In the peptide profile presented in this study (RP-HPLC), a higher percentage of hydrophilic peptides with different polarity index is observed, which suggests the presence of hydrophilic AA (Asp, Glu, Pro, Lys, Asn, and Arg) and neutral nature (His, Gly, Gln, Ser and Thr) in the C-terminal position. Of this AA, His, Lys and Arg are categorized in the AA group with strong AoxC together with hydrophobic AAs such as Met, Trp, and Tyr according to a study where they categorized 20 AA in relation to their AoxC [48]. Likewise, it has been suggested that a close relationship exists between polar peptides and the type of amino acid positioned in their side chains [49]. AA sequences of peptides derived from animal by-products having polar character AA in their side chains derived from poultry by-products with antioxidant capacity (Ser-Asn-Leu-Cys-Arg-Pro-Cys-Gly) [1], antihypertensive character (Cys-Gly-Lys-Pro, Ser-Gly-Arg, Cys-Thr-Ser-His, Val-Lys-Lys-Thr) [50] and (Ala-Ile-Tyr-His, Leu-Arg-Lys-Gly-Asn-Leu-Glu) [51] as well as by-products of marine origin with antioxidant capacity (Trp-Glu-Gly-Pro-Lys; Gly-Pro-Pro; Gly-Val-Pro-Leu-Thr) have been determined [52].

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### Table 2 Chromatographic profiles of peptides of by-products hydrolysates

| Molecular weight (kDa) | CH (%) | FH (%) |
|------------------------|--------|--------|
|                        | BP     | BK     | BRO    | BP     | BK     | BRO    |
| > 670                  | 8.3    | 8.2    | 4.6    | 3.3    | 3.5    | 1.0    |
| 670–148                | 7.5    | 6.2    | 5.8    | 1.7    | 1.6    | 1.4    |
| 148–44                 | 8.3    | 7.3    | 5.9    | 2.5    | 2.3    | 3.0    |
| 44–17                  | 7.0    | 5.6    | 4.7    | 2.3    | 1.5    | 2.5    |
| 17–135                 | 44.6   | 50.2   | 51.1   | 38.8   | 44.3   | 43.2   |
| < 1.35                 | 24.4   | 22.5   | 27.9   | 51.3   | 46.9   | 48.9   |

Chromatographic profiles of peptides corresponding to chicken by-products hydrolysates (CH) and fish by-products hydrolysates (FH) obtained with BP protease, BK protease, and BRO.
fish by-products are useful for obtaining low MW peptides (17.5–1.35 kDa).

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Authors' contributions
MGRG Produced the data and drafted the original manuscript. EMM provided the equipment for the determination of exclusion molecular, analyzed and interpreted the data of size exclusion chromatography (SEC). AHM, BVC and AFGC provided the equipment for the determination of peptide profile in the protein hydrolysates and they analysed the data of peptide profile. EMG analysed data formally, supervised and revised the work substantially. MLGM interpreted the data of size exclusion chromatography (SEC). AHM, BVC and AFGC provided the equipment for the determination of exclusion molecular, analyzed and interpreted the data of SEC. MLGM formally analysed data in general, supervised, revised the work substantially and acquisition of resources for research development. All authors read and approved the final manuscript.

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Availability of data and materials
The datasets used and/or analysed during the current study are included in this article. All data analysed during this study are included in this article.

Competing interests
The authors declare that they have no competing interests.

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