

Article

Chemical-Structural Identification of Crude Gelatin from Jellyfish (*Stomolophus meleagris*) and Evaluation of Its Potential Biological Activity

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Abstract: The demand for jellyfish is growing worldwide, especially due to their high nutraceutical value. In this study, the extraction and characterization of crude gelatin from the brown cannonball jellyfish (*Stomolophus meleagris*), which is periodically found in large volumes on the American Pacific coasts, were carried out. The crude gelatin obtained by alkaline treatment, with subsequent heat and dialysis treatment, showed an ability to quench free radicals (via ABTS and ORAC methods), and protect human cells against oxidative damage (through inhibition of hemolysis by AAPH), and they protected against mutations caused by aflatoxin B₁ in the *Salmonella enterica* Typhimurium TA100 strain. Furthermore, it was established that these extracts were innocuous for eukaryotic cells (genotoxicity assay). The amino acid profiles indicate a high concentration of glycine and proline, as well as charged amino acids. Electrophoretic, FT-IR, and ¹H-NMR studies indicated that one of the main proteins present in this crude gelatin is collagen. The presence of collagen and other proteins was identified by proteomic studies. Alkaline crude gelatin from brown jellyfish could be considered as potential candidates to be evaluated as antioxidant agents in foods in future research.

Keywords: antioxidant; antimutagenic; jellyfish; proteomic

Key Contribution: This study shows that jellyfish *Stomolophus meleagris* proteins are a promising alternative source of antioxidant compounds. Furthermore, detecting different proteins with antioxidant capacities, such as collagen, tubulins, and histones, would further clarify, at least in part, jellyfish-related health benefits. This obtained information can be utilized for new protein findings or subsequent comparative investigations to enhance our comprehension of marine organisms' biological properties.



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1. Introduction

The demand for jellyfish suitable for human consumption is increasing worldwide, with uses as food that provide beneficial compounds for the consumer [1]. Some species of said edible jellyfish have been captured around the American continent, such as *Stomolophus* sp. [2] These marine species represent a low-cost, raw material for obtaining nutraceutical products [1]. Therefore, the identification and characterization of one of their main chemical components, proteins, are necessary.

Jellyfish are marine organisms that are found on the coasts of many countries, and some products have been obtained from them, but with a low commercial value. Given their reproduction rate, jellyfish are often regarded as plagues, mainly in coastal cities. One of the causes to which this massive increase in jellyfish has been attributed to is climate change. Previous research in various countries have been performed assessing the potential changes generated in the ecosystems by jellyfish [3].

Among the most common jellyfish species identified are the moon and cannonball jellyfish. The latter belongs to the genus *Stomolophus* spp. and is abundant on the coasts of the Pacific Ocean and the Gulf of Mexico. Different species of jellyfish can be found between the northern part of Sonora, in the Gulf of California, and Ecuador in South America [4]. Their high-water content and proportion of stromal proteins makes jellyfish a highly perishable species, so a conservation process must be performed immediately after capture [3]. The large amounts of jellyfish processed leads to the generation of waste, particularly comprising anatomical areas that are not used. These residues can be recovered and used efficiently, which can only reduce the negative impact on the environment, but also generate products with high commercial value, such as proteins.

It has been previously reported that in jellyfish species, in addition to their biochemical composition, their proteins have a high antioxidant activity [5–13], which makes these organisms a wholesome food and antioxidant compound supply.

In addition to the potential use of jellyfish for their bioactive compounds, these organisms have a high collagen content [14]. The collagen extracted from jellyfish via heat treatment has shown good functional properties, making it amenable to appliances in the food industry [13,15–17]. However, the information available on using crude gelatin samples from *S. meleagris* jellyfish in the foodstuff sector still needs to be improved.

The main aim of this work was to obtain a crude gelatin with bioactive properties using alkaline conditions from the brown cannonball jellyfish (*Stomolophus meleagris*). However, it has been widely documented that crude gelatin obtained under alkaline conditions represent a complex system of proteins with different structures and molecular masses. Therefore, to deepen our knowledge of this crude gelatin, proteomics can be utilized, which allows for the analysis of proteins on a large scale in a particular biological system [18]. Based on our information, no other reports exist whereby proteins obtained from the brown cannonball jellyfish under alkaline conditions have been successfully identified by proteomic analysis. In this investigation, the antioxidant and antimutagenic activities of crude gelatin obtained from the jellyfish under alkaline conditions and after heat treatment were established, and chemical-structural analyzes and further identification of the obtained crude gelatin using proteomic techniques were performed.

2. Materials and Methods

2.1. Sample Preparation

The jellyfish (*Stomolophus meleagris*) were obtained from the Gulf of California by local fishermen from Puerto Peñasco (31°18'24" N 113°32'24" W). The jellyfish were kept on ice during handling and transferred cut to the laboratory. They were rinsed and then stored at −20 °C inside polyethylene bags.

2.2. Protein Extraction and Gelatin Production

Protein extraction and gelatin production were executed, as explained in previous research [19], with certain adjustments. The jellyfish was cut into small pieces, soaked in alkali (0.1 N NaOH, 1:5 *w/v* for 24 h), and rinsed until the pH dropped to 7. After that, a thermic treatment was applied (water bath at 60 °C for 12 h). The protein solution was then filtered through the gauze and dialyzed at 4 °C in water, employing a 10 kDa molecular weight cut-off cellulose membrane. The gathered samples were frozen (−25 °C) and lyophilized.

2.3. Biological Activity

2.3.1. Antioxidant Activity

The in vitro antioxidant activities for the crude gelatin sample were evaluated by four spectrophotometric assays: ABTS [2,2'-Azino-Bis (3-ethylbenzothiazoline-6-sulfonic acid)], FRAP (ferric reducing-antioxidant power), ORAC (oxygen radical absorbance capacity), and the protective effect on human erythrocyte against AAPH [2,2'-azobis(2-amidinopropane) dihydrochloride].

The ABTS radical scavenging activity was evaluated as described previously [20]. Potassium persulfate (2.45 mM) was employed to dissolve ABTS. The 7 mM ABTS solution was incubated for 12 h in dark at room temperature to generate ABTS radicals. The ABTS solution was diluted (distilled water) until its absorbance (734 nm) reached 0.7 ± 0.02 . A crude gelatin sample dissolved in water (20 μ L) was taken and mixed with ABTS solution (270 μ L) and kept in the dark at 25 °C (30 min). Finally, the reduction in absorbance (734 nm) was determined. The sample concentration required to inhibit 50% ABTS (IC₅₀) was used to express ABTS findings. A curve standard, considering X-axis as the gelatin sample concentration and Y-axis as the % inhibition, was employed to establish the IC₅₀ value.

A FRAP assay was then carried out [21]. The method used was based on reducing a compound consisting of TPTZ (2,4,6-tripyridyl-s-triazine) and Fe⁺³ (ferric iron). The presence of an antioxidant in an acid medium induces the formation of a colorless complex, in contrast to Fe⁺², which exhibits a greenish-blue color [21]. For this determination, the sample (20 mL) was mixed with a solution (280 mL) containing acetic acid–sodium acetate (pH 3.4), TPTZ, and FeCl₃ (10:1:1). The mixture was placed on a microplate reader, and after 30 min incubation, the absorbance at 630 nm was read. The results were indicated as Trolox equivalents.

The ORAC assay was executed as described previously [22]. In brief, fluorescence loss in the presence of AAPH was assessed at 37 °C for 90 min. The reaction was executed into 75 mM sodium phosphate buffer (pH 7.4), with the reached total volume of 2000 μ L containing: 1700 μ L buffer, 100 μ L fluorescein, 100 μ L AAPH, and 100 μ L of the sample. The results were shown as Trolox equivalents.

The protective effect on human erythrocytes was determined via the method established previously [23]. A pure suspension of erythrocytes was achieved by washing them thrice with ten mM phosphate-buffered saline at pH 7.4. For the hemolysis assay, a 2% solution was produced by resuspending the erythrocytes in PBS (phosphate-buffered saline). The suspended erythrocytes solution (100 μ L) was mixed with crude gelatin sample (100 μ L) and AAPH (100 μ L); the mixture was then incubated at 37 °C and shaken at 30 rpm in the dark (3 h). Then, PBS (1 mL) was added to the solution and centrifuged at 1500 \times g (10 min). The supernatants were then transferred to 96-well microplates using a spectrophotometric microplate reader (Multiskan Go, Thermo Scientific, Waltham, MA, USA). The percentage inhibition of hemolysis was established by measuring the absorbance (540 nm).

2.3.2. Ames Assay

The antimutagenic potential of the jellyfish crude gelatin sample was determined by the Ames assay [24]. The bioactivated *Salmonella enterica* Typhimurium T100 strain (using the S9 enzymatic mix) was taken as the biological model. Briefly, the bacterial strain was mixed with the jellyfish crude gelatin sample with a top agar containing L-histidine and D-biotin, as well as an S9 enzyme mixture. The mixtures were then quickly disposed of over minimal glucose agar plates and incubated at 37 °C (48 h).

2.3.3. Genotoxicity Test

Onions (*Allium cepa*) were allowed to germinate by immersion in distilled water and were then stored in the dark at room temperature (25 ± 2 °C). When the roots were about 5 cm long, the onions were ready for testing. The roots of the onions were treated with the jellyfish crude gelatin sample in two concentrations of 50 and 100 ppm for 24 h. The control group was treated with distilled water. The root tips were dehydrated for 45 min in

a 3:1 (*v/v*) ethanol–acetic acid solution, and then fixed in 1N hydrochloric acid for 2 min at 60 °C, after which they were stained with orcein for 1 min, and finally squashed and observed with a microscope to count the mitotic cells [25].

2.4. Protein Analysis

2.4.1. Chemical Analysis

Moisture (oven-dried), crude protein (with the Kjeldahl method), and ash (muffle furnace) contents were determined using the official methods [26]. The nitrogen-free material was estimated by difference [100 – (ash + moisture + protein)].

2.4.2. Amino Acid Analysis

The amino acid analysis was performed by reverse-phase high-performance liquid chromatography (Model GmbH Hewlett-Packard RP- HPLC, Agilent Technologies Inc., Santa Clara, CA, USA) [27]. The crude gelatin sample (100 mg) was homogenized with performic acid (100 mL). Performic acid was prepared before being used by mixing 30% hydrogen peroxide (1 mL) with 97% formic acid (19 mL), which was maintained in a closed container at room temperature (2 h). Then, the prepared performic acid was cooled (0 °C) and added to the gelatin sample. After keeping the mixture at 0 °C for 2.5 h, cold water was added (0.9 mL). An aliquot of this mixture (200 mL) was lyophilized. After that, the crude gelatin sample was hydrolyzed at 110 °C (18 h) under pressure in a 6 M HCL and sodium thioglycolate (1:1, *v/v*). The hydrolyzed sample was neutralized with 4 N NaOH. An aliquot was carried out and mixed with an equal volume of 10 mg/mL of *s*-1- α -amine *n*-butyric acid as an internal standard. Then, an aliquot was acquired and vortexed (1 min), with four parts of potassium borate buffer (pH 10.4) and *O*-phthaldialdehyde (1:1, *v/v*). Immediately, 20 mL was injected into the reverse phase column (C18 octadecyl dimethylsilane, 100 mm \times 4.6 mm), coupled to a pre-column (30 mm \times 4.6 mm) packed with the same material. Amino acids were analyzed with a fluorescence detector, and the peaks produced were evaluated (Chem Station program, Agilent Technologies Inc., Santa Clara, CA, USA). Elution involved two buffers at a flow of 1.0 mL/min during 25 min. Peak areas and retention times were contrasted with the commercial amino acid's standards mixture.

2.4.3. Electrophoretic Profile (SDS-PAGE)

The electrophoretic profile was determined using sodium dodecyl sulfate (SDS)-polyacrylamide (PAGE) gels of 4% and 8% concentrations; separation systems were respectively prepared and injected into the sample (20 μ g). Coomassie R–250 blue was used to stain the gels, and a mixture of methanol, water, and acetic acid (5:4:1, *v/v/v*) was employed to decolorize [28].

2.4.4. Fourier Transformed–Infrared Spectroscopy (FT-IR)

The FT-IR spectrum of the jellyfish crude gelatin sample was recorded (PerkinElmer FT-IR/FIT spectrometer, Waltham, MA, USA) from 4000 to 500 cm^{-1} with a resolution of 4 cm^{-1} . The crude gelatin sample (1 mg) was mixed with dry potassium bromide (100 mg, KBr) pellet [29].

2.4.5. Nuclear Magnetic Resonance of Proton (^1H -NMR)

^1H -NMR spectrum of the jellyfish crude gelatin sample was recorded at 400 MHz on a Bruker Advance 400 NMR spectrometer (Billerica, MA, USA, EE. UU.). The crude gelatin sample (1 mg) was dissolved in 0.5 mL of a 1% (*v/v*) deuterated potassium hydroxide solution (KOD 40% in D_2O) with D_2O water. The reference employed was dimethylsilapentane-5-sulfonic acid (DSS). The spectral frame was 20 ppm at 24 ± 1 °C [29].

2.4.6. Protein Identification

The crude gelatin sample was dissolved, and then identified and quantified with the nano LCMS/MS platform (Ultimate 3000 nano UHPLC system, attached to an Orbitrap Q Exactive HF mass spectrometer with Nanospray Flex Ion Source, Thermo Scientific).

Proteins were precipitated from the protein solution using methanol and chloroform. Then, the protein pellet was dissolved in a 2 M urea aqueous solution. After that, the crude gelatin sample was denatured under the following conditions: 10 mM dithiothreitol and incubation for 60 min at 56 °C. The denatured sample was then alkylated with 50 mM indole-3-acetic acid. Then, the sample was incubated at room temperature in the dark (60 min). Afterward, 0.1% trifluoroacetic acid (TFA) was added into the solution to make a final concentration of 0.1% TFA with a pH of 1.0. Pepsin was used as the specific protease. The salt from the peptides produced was performed with a C18 SPE column (Thermo Scientific). Prior to LC-MS/MS analysis, the obtained peptides were resuspended in a solution of 20 mL of 0.1% formic acid. The crude gelatin sample (1 mg) was then loaded onto a Nanoflow UPLC by using two buffers as the mobile phase: 0.1% formic acid in water (buffer A), and 0.1% formic acid in acetonitrile (buffer B). The total flow rate was 250 nL/min. Buffer B was employed to obtain the LC linear gradient: 5 min from 2% to 8%, 60 min from 8% to 20%, 33 min from 20% to 40%, and 4 min from 40% to 90%. The full scan (300–1650 m/z) was acquired using a single high resolution of 60,000 (at 200 m/z), and the automatic gain control target was set to 3×10^6 . The MS/MS Top 20 mode scan resolution was 15,000 at 200 m/z, and the normalized collision energy was 28%. The automatic gain control target was then established to 1×10^5 , and the maximum injection time was 19 ms. The dynamic exclusion parameters were time window (1.4 Th) of 30 s, charge state of 1, and unassigned were rejected, while charge states >6 were not dismissed. All the MS/MS spectra were analyzed using Peaks Study 8.5 software (Bioinformatics Solutions Inc., Waterloo, ON, Canada) against the jellyfish protein database. The following restrictions were used: pepsin cleavage with up to 2 missed cleavage sites, tolerance of 10 ppm as the precursor, and 0.5 Da as the fragment ion mass. Carbamidomethylation of cysteine was accounted for as a fixed modification. The permissible variable was methionine oxidation.

2.5. Statistical Analyses

A randomized entirely design was employed in this study. Each trial was replicated at least threefold. A one-way analysis of variance was applied to analyze the obtained data from the antioxidant activity, Ames assay, and genotoxicity tests. The Tukey–Kramer method was used to carry out the comparison of the means. A significance level of 95% ($p < 0.05$) was applied. The obtained results were analyzed using the INFOSTAT statistical software. Data from electrophoretic analysis, FTIR, NMR, and proteomics were evaluated by descriptive statistics.

3. Results and Discussion

3.1. Biological Activity

The biological activity of the jellyfish crude gelatin sample was measured by different assays. The ABTS scavenging ability was used to establish antioxidant capacity. ABTS can become a stable molecule after accepting an electron atom from the antioxidants. The IC₅₀ values were 0.42 ± 0.10 mg/mL. The IC₅₀ value of the crude gelatin sample was lower than those reported for the amaranth proteins (*A. mantegazzianus*; IC₅₀ of 10.2 ± 0.8), suggesting that the jellyfish crude gelatin sample exhibits higher antioxidant effects than amaranth, but have comparable hydrolysates, with values of 1.16 ± 0.09 mg/mL [30]. These can be compared with the hydrolysate values from other sources, such as mutton ham or bacteria (*Spirulina platensis*), of 0.71 ± 0.23 and 1.5 ± 0.1 , respectively [31,32].

The crude gelatin sample's ability to quench the peroxy radicals generated by the azo compound AAPH was evaluated by the ORAC assay. The hydrogen-donating antioxidant obstructs the peroxy radical chain reactions. The antioxidant activity of the crude gelatin sample was assessed by comparing its fluorescence decay curve against a blank [33]. The

ORAC value for the jellyfish crude gelatin sample was measured at 479 ± 7.17 mmol TE/g; these values can be compared to those of other marine organisms, such as the lipid extracts from seaweeds, with values of 461.84 and 362.53 mmol TE/g for *M. pyrifera* and *E. radiate*, respectively, and krill protein (*Euphausia superba*), with values between 274 and 537 mmol TE/g [34,35].

Although the jellyfish crude gelatin samples showed the ability to quench radicals, they did not show the ability to reduce ferric-tripyridyltriazine (FeIII-TPTZ) and form ferrous-tripyridyltriazine (FeII-TPTZ).

The radical AAPH induces oxidation, causing the hemolysis of erythrocytes. Radical degradation can be inhibited by antioxidant agents. The jellyfish crude gelatin sample presented an IC_{50} of 6 mg/mL. IC_{50} values of 0.12 ± 0.01 mg/mL have been reported in peptides from dairy products [36]; however, an IC_{50} of 10 mg/mL is considered to yield high protection on the part of erythrocytes against radical damage [37]. Therefore, the jellyfish crude gelatin sample appears to be protective against the AAPH radical, preventing oxidative damage occurring in the erythrocytes.

The antioxidant activities yielded by all the utilized methods in this study were different since they all have other mechanisms of action. The ABTS evaluated the ability of antioxidants to scavenge the performed radicals by donating an electron. In contrast, the ORAC and antihemolytic tests measured the ability of antioxidants to act as hydrogen donors and thus inhibit radical initiation, including the scavenging of formed radicals in biological systems [33,37]. In addition to the mechanism of each methodology, the antioxidant activity of the proteins depends on the amino acid sequence, the structure, and the size of the peptides. The authors of [31,38] report that aromatic amino acid residues including Tyr, Trp, and Phe might have to repress the unpaired electron activity by forming a conjugated electron system that inhibits the reaction process of peroxidation mediated by the free radicals [39]. Nucleophilic sulfur-containing amino acids such as Met, and other amino acids such as Pro and Leu have been confirmed [40]. Peroxyl radicals are stabilized by adding hydrogen that can be donated by several amino acids including His, Tyr, Leu, and Met. Zheng L. et al. [41] suggested that Tyr- and Trp-Gly protect the erythrocytes against AAPH-induced hemolysis primarily by acting as direct scavengers of ROO. Therefore, in the present study, it was considered that the presence of Gly, Leu, Phe, Pro, and Tyr, and the absence of His in the protein (as will be shown in the amino acid results section) are amenable to the antioxidant activity detected.

Concerning the antimutagenic activity, the jellyfish crude gelatin sample showed the capacity to inhibit mutation induced by AFB₁ in the *S. enterica* Typhimurium TA100 strain (Figure 1). The percentage of inhibition was greater than 50% when 0.5 mg/plate was tested, representing moderate inhibition [42], whereas with 4.0 mg/plate, the inhibition was greater than 60%, representing strong inhibition [42]. A similar antimutagenic activity (40–50%) was derived when 0.5 mg/plate of ethanolic extracts from salmon (*Oncorhynchus masou*) was tested [43], but this activity was higher than that of the protein hydrolysate derived from giant squid skin, which showed 50% inhibition when 5 mg/plate was evaluated [44].

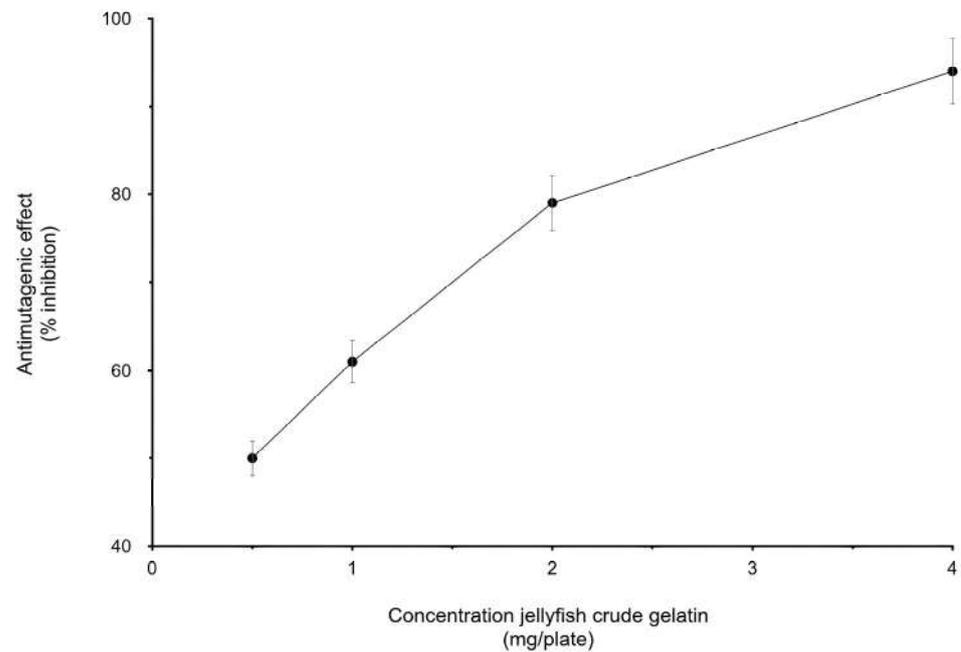


Figure 1. Effect of the gelatinous extract obtained from the cannonball jellyfish on mutagenicity induced by AFB₁, based on the *Salmonella enterica* Typhimurium TA100 assay.

3.2. Genotoxicity Test

Different phases of normal and abnormal mitosis are shown in Figure 2. Based on these images, we determined whether the crude gelatin sample had an adverse effect on the chromosomes. Table 1 shows the percentages of clastogenic cells; cells exposed to jellyfish crude gelatin sample at 50 ppm reached 3.44%, and those exposed to 100 ppm exceeded 5.17%. These values are low compared to those of other food additives, such as butylated hydrotoluene and butylated hydroxyanisole, at 28.67% and 28.60%, respectively [45].

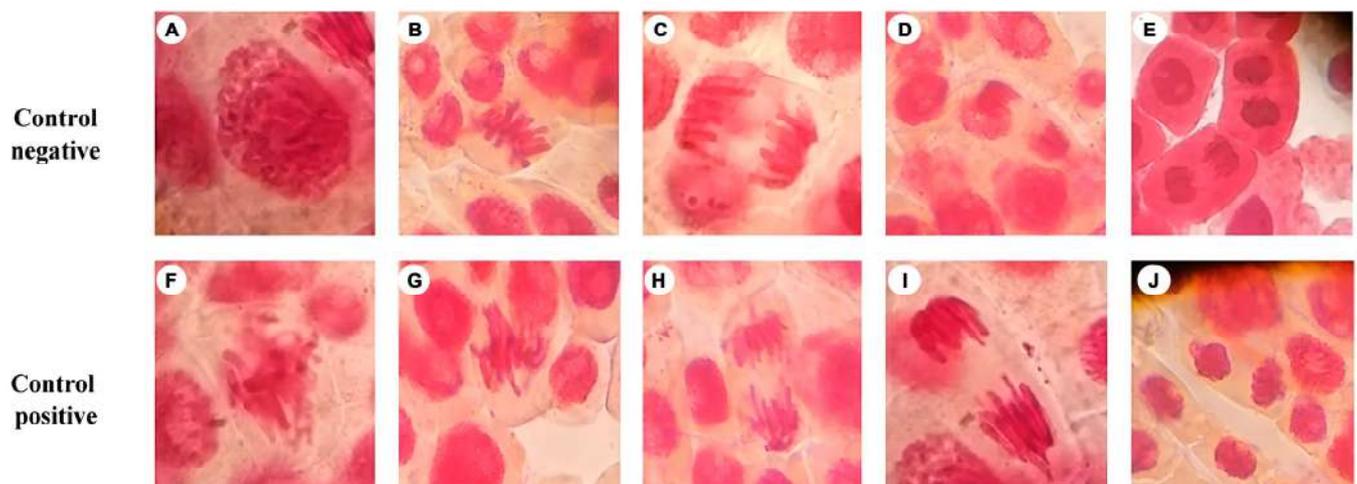


Figure 2. Phases of normal and abnormal mitoses exposed to water and sodium azide. Chromosomal aberrations observed in *Allium cepa* root tip cells: (A) normal prophase; (B) normal metaphase; (C) normal anaphase; (D) normal late anaphase; (E) normal telophase; (F) abnormal metaphase; (G) abnormal metaphase; (H) abnormal anaphase with lagging; (I) abnormal anaphase with lag; and (J) abnormal telophase.

Table 1. Clastogenic effects of the jellyfish crude gelatin sample (JCG) on the mitotic cells of *Allium cepa* *.

Treatment	Genotoxicity Percentage
Negative control	0 ^c
Sodium azide (10 ng)	88 ^a
JCG 50 ppm	3.44 ^b
JCG 100 ppm	5.17 ^b

* Values represent the average of three repetitions. Mean values with different letters indicate differences ($p < 0.05$) between the crude gelatin samples and the sodium azide control.

3.3. Protein Analysis

3.3.1. Yield

The yield of crude gelatin sample, expressed on a dry basis, and utilizing NaOH for extraction, was 10.49 ± 0.183 g lyophilized from 100 g of dry weight. This result is higher than other previously reported values, such as in *S. meleagris* (7.5 g/100 g) [17] and for *Cyanea nozakii* Kishinouye (5.5 g/100 g) [9].

3.3.2. Chemical Composition

The chemical composition of the jellyfish crude gelatin sample exhibited a moisture content of $10.67 \pm 0.74\%$, which is comparable to other marine organisms, such as red tilapia (gelatin—9.59%) [46] or commercial bovine (gelatin—9.7%) [47]. The crude protein result was $76.38 \pm 0.91\%$, which is higher than in dry-salted jellyfish (29.54%) [17], but comparable to commercial gelatin (87.52%) [47]. The ash result was $5.83 \pm 0.16\%$, which is low compared to dry-salted jellyfish gelatin ($56.61 \pm 0.13\%$). However, the value was much higher than commercial gelatin ($0.9 \pm 0.2\%$) [47]. This may be due to the marine nature of the crude gelatin sample, possibly containing high concentrations of salts [17]. A more effective method for salt removal should be developed since ash contents up to 2.5% are accepted for food products. The value of nitrogen-free material was high when compared to commercial gelatin (1.8%) [47].

3.3.3. Amino Acid Analysis

The amino acid composition of the jellyfish crude gelatin sample was determined (as shown in Table 2). The most abundant amino acid found was Gly. High concentrations of Pro and charged amino acids such as Arg, Asp, Glu, and Lys, and aromatic Phe and Tyr were all observed. This result was expected since Gly is the main constituent of collagen. It is typical to find repeating triplet Gly-X-Y sequence, X and Y, in most likely any amino acid. However, Pro and Hyp residues were the most frequent triplet in collagen [48]. The amino acid composition trends (high concentration of Gly, Arg, and Pro) are comparable to another marine organism, deep water rose shrimps [49].

Interestingly, histidine (His), considered a potent antioxidant amino acid, was not detected in the obtained crude gelatin sample. This result differs from other reports, such as those developed in *Rhizostoma pulmo* [50]. The absence of His could limit the crude gelatin sample's capacity to trap metallic ions, as was detected in this study. However, more studies will be required to confirm this.

Table 2. Amino acid composition of the crude gelatin sample obtained from cannonball jellyfish (*Stomolophus meleagris*) *.

Amino Acid	mg Amino Acid/100 g Protein
Alanine	269
Arginine	359
Aspartic acid	440
Glutamic acid	508
Glycine	1336
Histidine	n.d.

Table 2. Cont.

Amino Acid	mg Amino Acid/100 g Protein
Isoleucine	126
Lysine	175
Isoleucine	126
Leucine	175
Lysine	224
Phenylalanine	85
Proline	345
Serine	149
Threonine	190
Tyrosine	95
Valine	134

* Values represent the average of three repetitions. n.d. Not detected.

3.3.4. Electrophoretic Profile

The jellyfish crude gelatin sample's electrophoretic profile displayed two bands in continuous positions, with molecular weights of approximately 200 kDa and 97 kDa (Figure 3, Lane B). This profile suggests that the crude gelatin sample molecules comprise at least β and α collagen chains. Three polypeptide chains, named α chain, coiled around one another in a triple-helical shape, are present in all collagen molecules. Typical collagen gelatin displays one β band (200–210 kDa) and two α bands (around 100 kDa for α_1 and α_2), which in the triple helix are the unfolding chains [48,51]. The results were deemed to be similar to those of other marine organisms [52,53], wherein thermic treatment degrades the triple helix into the β and α chains.

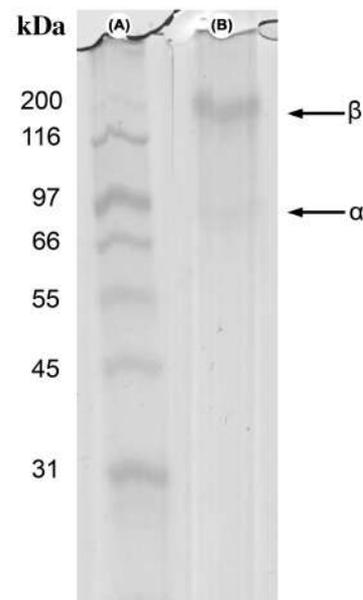


Figure 3. SDS–polyacrylamide protein pattern of the jellyfish (*Stomolophus meleagris*) crude gelatin sample. (A): molecular weight marker; (B): jellyfish crude gelatin sample.

3.3.5. Fourier Transformed–Infrared Spectroscopy (FT-IR)

The jellyfish crude yielded an FT-IR spectrum (Figure 4) similar to other jellyfish species, such as *Rhopilema esculentum* and *Rhizostoma pulmo* [50,54]. The FT-IR spectra exhibited peaks characteristic of amides A and B, and amides I, II, and III. Amide A was observed at 3290 cm^{-1} ; the NH stretching vibrations determined its position. The position of the amide B band was detected at 2920 cm^{-1} due to asymmetric CH stretching. At 1660 cm^{-1} the detection of the amide I band is due to C=O stretching. The amide II band

was observed at 1585 cm^{-1} , resulting from N-H and C-N torsional vibrations. Amide I and II bands' amplitudes are linked to the collagen structural spiral form, and linkages of α -helices, respectively [55,56]. The amide III band was observed at 1220 cm^{-1} due to the combined peaks of C-N stretching vibrations, and deformation of N-H from the amide linkages, and wagging vibration of the CH_2 groups of the Gly-backbone and Pro-side chains [57].

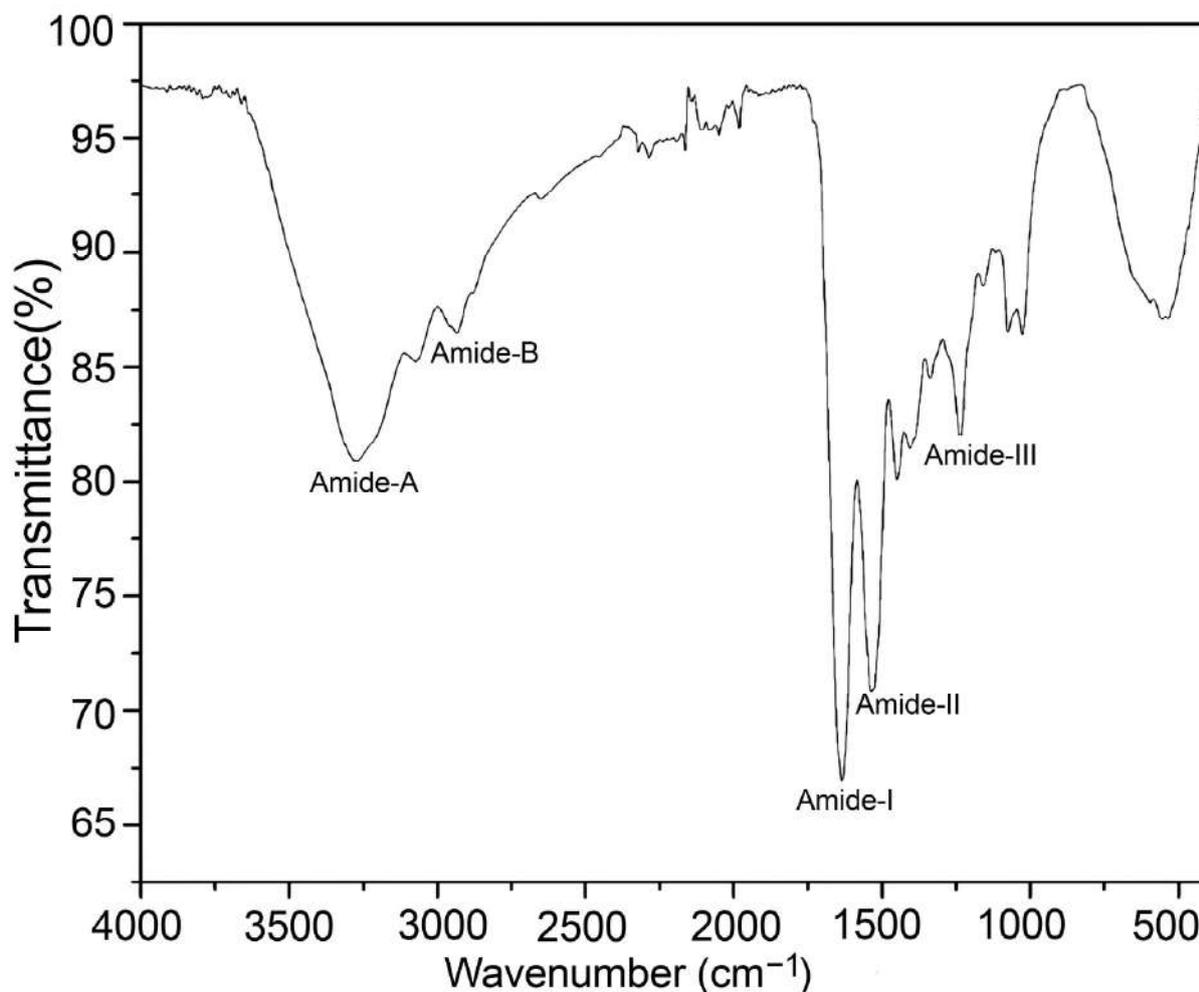


Figure 4. FT-IR spectra of the jellyfish crude gelatin sample.

3.3.6. Nuclear Magnetic Resonance of Proton ($^1\text{H-NMR}$)

The $^1\text{H-NMR}$ spectra of the jellyfish crude gelatin sample (Figure 5) were compared to those from previous studies and spectrum databases. Figure 5 indicates the signals corresponding to each amino acid, as well as the integration of the most relevant amino acids. The MestReNova software 8 (MestReNova v9.0.1-13254, Mestrelab Research S.L., Santiago de Compostela, Spain) was employed to analyze the $^1\text{H-NMR}$ spectra, and the areas of glycine, proline, and hydroxyproline were obtained by the integration of the different chemical shifts. The signal centered at 4.19 ppm, corresponding to the glycine signal (CH_2), while the signals at 1.19, 2.19, and 3.79 ppm correspond to proline and CH_2 , and the signals at positions 3.28 and 3.39 ppm correspond to hydroxyproline. These integrals suggest a 1:1 ratio of glycine to proline, while hydroxyproline is present at a smaller proportion. These observations are consistent with the amino acid content profiles discussed previously [13,58].

Amino Acid	δ ppm	Integral Values
Glycine	4.19	2.07
Proline	2.19	1.94
Hydroxyproline	3.28	0.25

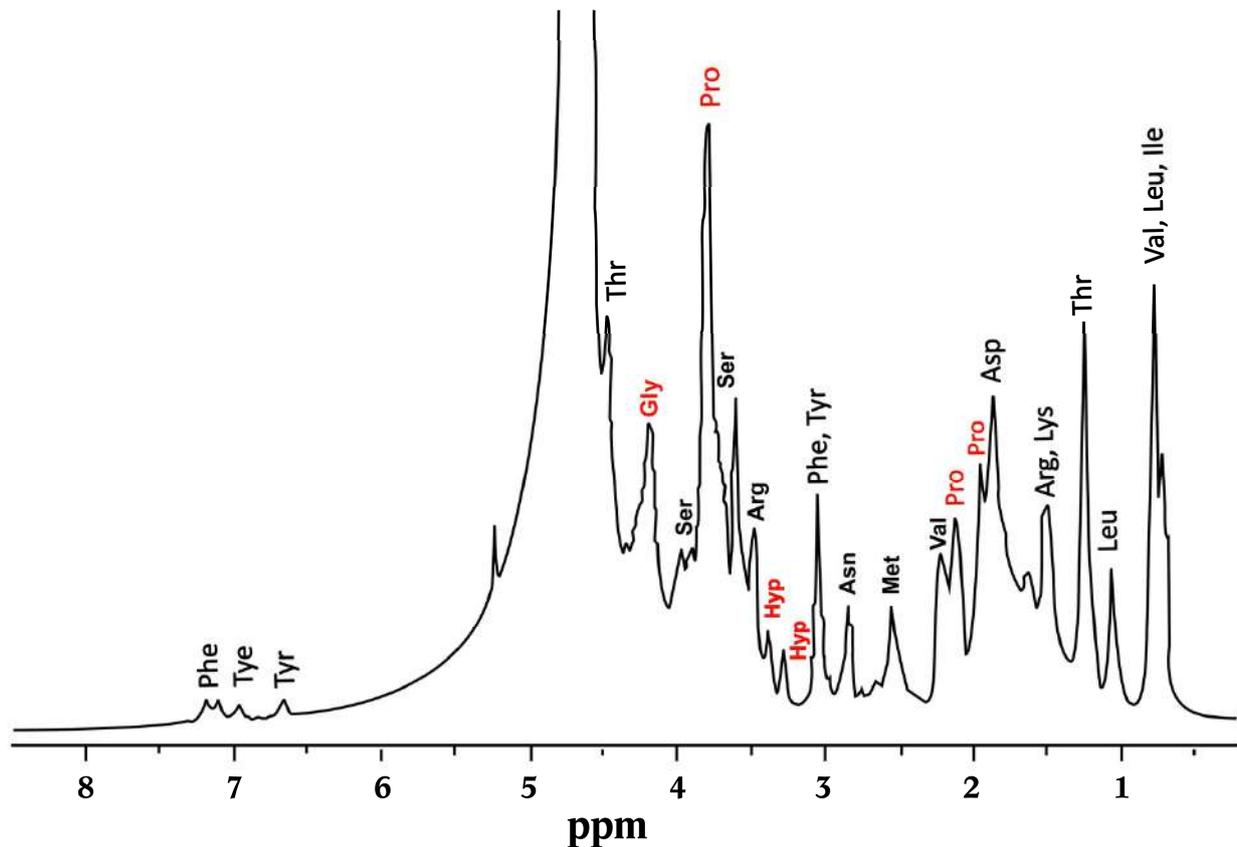


Figure 5. $^1\text{H-NMR}$ spectra of the jellyfish crude gelatin sample. Amino acids are indicated by their corresponding peaks.

3.3.7. Proteomic Identification

The protein identification of the crude gelatin sample from jellyfish is summarized in Table 3. These results confirm the presence of collagen, alongside other proteins, such as tubulins and histones. Tubulins carry out various cellular functions, such as in providing structural support and have been used as a drug for multiple diseases, including cancer [59]. Histone proteins, which were also identified in this work, are DNA-binding proteins present in many animal species, including the jellyfish *Stomolophus meleagris* [60]. Histones, such as peptides of the H3 class, are present in the obtained crude gelatin sample. Histones can act as antimicrobials against viruses, bacteria, fungi, and parasites [61]. The collagen in this study was characterized as belonging to type IV. Collagen extracted from jellyfish has been shown to reduce different human cancer lines [1]. Therefore, the bioactive properties identified in the present study could be associated with other proteins in the crude gelatin extracts alongside collagen. Still, these proteins need to be validated using further bioactivity assays.

Table 3. Proteins identified in the brown cannonball jellyfish crude gelatin sample with nanoLC-MS/MS and bioinformatic analysis.

Accession Number	Description	Uni. Pep.	PSM	Cov (%)	Score
tr Q7YZL5 Q7YZL5_AURAU	Tubulin alpha chain (Fragment) OS = <i>Aurelia aurita</i> OX = 6145 PE = 2 SV = 1	24	28	38	212.94
tr U3PBW0 U3PBW0_AURAU	Beta actin (Fragment) OS = <i>Aurelia aurita</i> OX = 6145 PE = 2 SV = 1	14	17	37	187.05
tr A0A2Z5WH75 A0A2Z5WH75_9CNID	Actin-2 OS = <i>Aurelia sp.</i> 2017-HT OX = 2136148 PE = 2 SV = 1	16	19	34	187.05
tr Q5VJP9 Q5VJP9_AURAU	Tubulin beta chain (Fragment) OS = <i>Aurelia aurita</i> OX = 6145 PE = 2 SV = 1	20	25	46	113.72
tr Q5VJP9 Q5VJP9_AURAU	Histone H3 (Fragment) OS = <i>Aurelia aurita</i> OX = 6145 PE = 3 SV = 1	2	3	9	56.75
tr G9IBZ5 G9IBZ5_CARAL	Cytochrome c oxidase subunit 1 OS = <i>Carybdea alata</i> OX = 1193083 GN = cox1 PE = 3 SV = 1	16	44	40	50.35
tr V9GWB0 V9GWB0_CRASO	Collagen type IV OS = <i>Craspedacusta sowerbii</i> OX = 128124 PE = 2 SV = 1	15	17	14	38.32

Uni.Pep., unique peptides; PSM, peptide spectrum matches; Cov, protein coverage; and Score reported after database search, with a score >32 indicating identity or extensive homology at $p < 0.01$.

4. Conclusions

Under the conditions of this study, the alkaline-extracted proteins obtained from the cannonball jellyfish showed in vitro antioxidant and antimutagenic capacities. Moreover, the extracts are not clastogenic. The bioactivities detected in this study can be attributed to the presence of proteins such as collagen, tubulins, and histones. All of these can contribute to free radical scavenging and antimutagenic activities. This information suggests that cannonball jellyfish proteins have potential uses in the food industry as antioxidant agents. Currently, work is in progress on applying jellyfish gelatin in the preservation of food products, and the results of this study further demonstrate the bioactivity of jellyfish proteins.

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Data Availability Statement: The data presented in this study are available in the article. Further information is available upon request from the corresponding author.

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