



Golden Sea Cucumber: Identification and the Antioxidant Activity of Its Collagen Hydrolysates

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Abstract

Golden sea cucumber or locally known as “teripang emas” is one of Indonesia’s most popular sea cucumber and widely processed as functional food or supplement due to its bioactivities. The sea cucumber is often misidentified due to its morphological similarities with other *Stichopus* spp. This study aimed to identify the golden sea cucumber obtained from West Nusa Tenggara, Indonesia, by a molecular method and study the antioxidant activities of its collagen hydrolysates. The hydrolysates were produced by hydrolyzing acid collagen extract using neutrase for 30, 60, 120, 180, and 240 mins. The products were then analyzed for their degree of hydrolysis, peptide content, molecular weight distribution and radical scavenging activity by the 2,2-Diphenyl-1-picrylhydrazyl (DPPH) method. Results showed that hydrolysis for 180 mins was optimal in producing the highest peptide content, 12.79 ± 0.44 mg/mL, with a degree of hydrolysis (DH) of $55.2 \pm 1.50\%$. However, the highest antioxidant activity (IC_{50} of 5.25 ± 0.15 mg/mL) was demonstrated after 60 mins hydrolysis with molecular weight (MW) ranged from less than 14.4 kDa to approximately 25 kDa. The hydrolysate might be categorized as a weak to moderate antioxidant. Based on the molecular identification, the golden sea cucumber had 99% similarities with *Stichopus horrens* and *S. monotuberculatus*.

Keywords: golden sea cucumber, identification, collagen hydrolysates, neutrase, antioxidant

1. Introduction

Sea cucumbers have long been known as a potential source of functional food and biomedicine products. This is because their various compounds contain therapeutic activities, such as saponin, chondroitin sulfates, fatty acids, vitamins, minerals, bioactive collagen peptides, amino acids, etc. (Bordbar, Anwar & Saari, 2011). Various Indonesian sea cucumbers have been explored for their potential bioactivities (Adriansyah & Sulastri, 2014; Dewi, Patantis, Fawzya, Irianto, & Sa’diah, 2020; Khirzin, Sukarno, Yuliana, Fawzya, & Chasanah, 2015; Pangestuti & Arifin, 2018). However, many other local Indonesian sea cucumbers need further exploration of their biological activities related to their health benefit

effects and possibilities as new sources of functional constituents.

Golden sea cucumber belongs to the genus *Stichopus*, which can be found in Sumatra, Nusa Tenggara and Sulawesi waters (Fad’ha, Arma, & Busman, 2019; Nursid, Marraskuranto, Kuswardini, & Winanto, 2019; Pangestuti et al., 2016; Safithri, Setyaningsih, Tarman, Yuhendri, & Meydia, 2018). To date, golden sea cucumbers are often called *Stichopus hermanni* or *S. variegatus* because of their similar morphological characteristics. For the same reason, misidentification often occurs to differentiate between *S. horrens*, *S. monotuberculatus*, *S. quadrifasciatus* and *S. naso* (Purcell, Samyn, & Conand, 2012). Identification using the mitochondrial

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cytochrome c oxidase subunit I (COI) gene could distinguish several species in a specific genus of dried sea cucumber (Patantis, Dewi, Fawzya, & Nursid, 2019). Based on the previous studies above, molecular identification should be done to confirm the species of golden sea cucumber.

Similar to other sea cucumbers of high economic values, most golden sea cucumbers are processed into *beche-de-mer*. They also have been widely used for traditional healing products, which are trending now, shown by the number of supplement products on the market. However, the utilization of golden sea cucumbers found in West Nusa Tenggara is still limited to the dried food products.

Research on the biological activity of sea cucumbers from Nusa Tenggara have been reported, including radical scavenging activity of ethanolic extract of *Holothuria leucospilota*, *H. atra*, *H. fuscocinerea*, and *H. excellens* (Pangestuti et al., 2016), α -glucosidase inhibition of methanolic extract of *S. hermanni* (Mariyanti, 2017), and tyrosinase inhibitor activity of *S. hermanni* collagen (Safithri et al., 2018). Other explorations of golden sea cucumber or gamat (*S. hermanni* or *S. variegatus*) from various locations that have been published were studies on collagen extraction (Alhana, Suptijah, & Tarman, 2015; Fawzya, Chasanah, Poernomo, & Khirzin, 2016), sulfated glycosaminoglycan content (Masre, Yip, Sirajudeen, & Ghazali, 2010), angiotensin-I-converting enzyme (ACE) inhibitor and antioxidant activity of collagen peptide (Khirzin et al., 2015), the potential of *S. hermanni* to improve the growth of stem cells (Arundina, Suardita, Setiabudi, & Ariani, 2016), the effect of sea cucumber extract on the lymphocytes numbers of Wistar rat's oral mucus during the healing process from a traumatic ulcer (Arundina, Yuliati, Soesilawati, Damaiyanti, & Maharani, 2015); the growth-promoting effect of *S. variegatus* its water extract to boost the proliferation of spinal astrocytes (Patar, Jamalullail, Jaafar, & Abdullah, 2012) and the ability of sulfated glycosaminoglycan from *S. hermanni* integument in the process of wound healing (Masre et al., 2010).

Sea cucumbers are rich in protein, of which 70% is collagen with broad applications in producing biomedical, pharmaceutical, body care, and food products (Siddiqui et al., 2013). In the last decade, marine collagens, including sea cucumber collagens and their peptides, have significantly been an interest for study related to their potential biological activity. Various proteases have been reported to produce collagen hydrolysates, including pepsin (Khirzin et al., 2015; Siddiqui et al., 2013), alcalase (Kusumaningtyas, Nurilmala, & Sibarani, 2019; Safari & Yaghoubzadeh, 2020); trypsin, papain, bromelain,

flavorzyme, and protamex (Forghani et al., 2012). The use of neutrase to hydrolyze sea cucumber collagen from *Acaudina molpadioides* produced the highest antioxidant activity of hydrolysate compared to those produced using papain, pepsin and trypsin (Jin, Xu, Li, Zhang, & Xie, 2019). Neutrase was also reported as the best enzyme that produced hydrolysates efficiently with beneficial functional and bioactive characteristics (Liu, Ma, Che, Wang, & Li, 2018; Ou et al., 2010).

This study was aimed to produce collagen hydrolysates from golden sea cucumber obtained from West Nusa Tenggara using neutrase. The peptide content, molecular weight distribution, and radical scavenging activity of hydrolysate were determined as the important essential figures on utilizing the sea cucumber for antioxidative nutraceuticals.

2. Material and Methods

2.1. Material and Chemicals

Fresh and dried sea cucumber samples were collected from Labuhan Terata, Sumbawa, West Nusa Tenggara, Indonesia, in February 2017, with 31.88 ± 6.62 cm in length, 9.42 ± 2.35 cm in width, and 796.67 ± 334.75 g in weight each. The samples were washed with freshwater, gutted, washed again and packed into plastic bags. They were then transported (about six hours) in a cool box filled with ice to the laboratory in Jakarta, and then stored at -20°C until used for collagen extraction (for about seven days). Due to the limited fresh samples, dried samples were employed for species identification. The dried samples were obtained from a sea cucumber processor at the same location as the fresh one.

Neutrase® 0.8 L (a protease from *Bacillus amyloliquefaciens*) was supplied by Novozymes (Denmark). SnakeSkin™ dialysis tubing 10 K MWCO was obtained from ThermoScientific (Denmark). DNeasy Blood Tissue Kit was bought from Qiagen (Netherlands) and KOD FX Neo master mix for the PCR was supplied by Toyobo (Japan). DPPH and tryptone casein were obtained from Sigma-Aldrich (Germany). All other chemicals and reagents were pro analysis.

2.2. Identification of golden sea cucumber

A dried sea cucumber sample was identified by a molecular method as described by Uthicke, Byrne, and Conand, 2010, which was based on PCR amplification of ± 675 bp nucleotides from the gene of mitochondrial (COI). COIeF: 5'-ATAATGATA GGAGGRTTTGG-3' and COIeR: 5'-GCTCGTG

TRTCTACRTCCAT-3' primers were used to amplify that region (Arndt, Marquez, Lambert, & Smith, 1996), whereas sequencing was conducted at 1st Base Laboratories (Malaysia). The COI Sequence data were analyzed based on the Basic Local Alignment Search Tool (BLAST). MEGA 7 was used to construct the phylogenetic tree (Kumar, Stecher, & Tamura, 2016).

2.3. Chemical Composition of Sea Cucumber

Chemical composition of sea cucumber was analyzed, including ash content using SNI: 01-2354.1-2006 (BSN, 2006a), moisture content by SNI 01-2354.2-2006 (BSN, 2006b), protein by SNI: 01-2354.4-2006 (BSN, 2006c), and fat content by SNI: 01-2354.3-2006 (BSN, 2006d).

2.4. Extraction of Collagen

The sea cucumber collagen extraction was conducted according to Alhana et al. (2015) with slight modification. All extraction steps were carried out in chilling conditions (approximately 4°C). Previously, samples were cut into small pieces (about 2 cm x 2 cm), immersed, and gently stirred in aquadest (1:5 w/v) for 30 mins, then the water was removed by filtration using a plankton net. Afterward, samples were soaked in 5 volumes (v/w) of 0.1 M NaOH solution to remove other proteins. This process was performed while stirred continuously for 48 h. The NaOH solution was replaced with a fresh one every 24 h. The mixture was then filtered and the sample was washed using aquadest until the remaining washed aquadest had a neutral pH. The resulting mixture was the crude collagen fiber.

The crude collagen was extracted using 0.1% acetic acid solution (1:5 w/v) for 48 h, followed by filtration. The collagen-containing filtrate was subsequently precipitated by a slow addition of NaCl powder until the NaCl final concentration reached 1 M, then kept at 4 °C for 24 h. The mixture was centrifuged at 15,344 x g, 4 °C for 15 mins to produce collagen as the residue material. The next step was dialysis of collagen solution (in 10 mL 0.5 M acetic acid) using a 10K-MWCO SnakeSkin™ dialysis tubing against 1 L 0.1 M acetic acid for 12 h. The dialysis solution was changed twice. Finally, the collagen was dialyzed against distilled water for an hour. The dialyzed collagen was centrifuged at 15,344 x g, 4 °C for 15 mins, and the pellet as the dialyzed collagen was ready to be hydrolyzed. The yield of crude and dialyzed collagen was calculated as the percentage of each collagen weight to initial fresh sea cucumber weight (% wet base or % wb).

2.5. Hydrolysis of Collagen

The collagen hydrolysates of sea cucumber were prepared according to Ou et al. (2010). Firstly, collagen was added with 0.01 M phosphate buffer (pH 6.7) with a ratio of 1:4 (w/v), then digested with 0.5% (v/v) neutrase (4416.8 U/mL). Hydrolysis was done on a hot plate with continuous stirring for 30, 60, 120, 180, and 240 mins. The hydrolysis process was stopped by heating the mixture in boiling water for 10 mins. The hydrolysates were then separated by centrifugation at 4 °C; 9,820 x g for 10 mins. Before the analyses, the hydrolysates were dried using a concentrator. The yield of dried collagen hydrolysates was expressed as percentage of dried collagen hydrolysates weight to crude or dialyzed collagen or fresh sea cucumber weight.

2.6. Determination of Degree of Hydrolysis (DH)

The degree of hydrolysis was measured using trichloroacetic acid (TCA) precipitation method (Baharuddin, Halim & Sarbon, 2016) with a slight modification. Approximately 500 µL of hydrolysate was added to 500 µL of 20% TCA, homogenized with vortex and kept for 30 mins at 4 °C to allow precipitation. Nitrogen content of both hydrolysates and supernatants obtained from the hydrolysate treated with TCA (by centrifugation at 4 °C; 7,800 x g for 15 mins) were determined using the Kjeldahl method. The DH was calculated using Equation 1.

$$DH (\%) = \frac{\text{Soluble N in 20\% TCA} \times 100\%}{\text{Total N in the hydrolysate}} \quad (1)$$

2.7. Determination of Peptide Content

The peptide content of collagen hydrolysates was determined by o-phthalaldehyde (OPA) method (Budiari, Chasanah, Suhartono, & Palupi, 2018). OPA was prepared by mixing 25 mL of 100 mM sodium tetraborate, 2.5 mL of 20% (w/v) sodium dodecyl sulfate (SDS), 1 mL OPA solution (40 mg of OPA dissolved in 1 mL of methanol), 100 µL α-mercaptoethanol, and 21.4 mL distilled water reaching to a final volume of 50 mL. This reagent was prepared daily. A mixture containing 50 µL of sample and 2 mL of OPA solution was kept for 2 mins at ambient temperature, then read the absorbance at 340 nm. Tryptone casein in phosphate buffer (pH 7.4) with the series of concentrations 0; 0.1; 0.5; 1 and 2 mg/mL was used as a standard curve to determine peptide content.

2.8. Molecular Weight Analysis

The molecular weight of collagen was described using sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) method (Bollag, Rozycki, & Edelstein, 1996). Collagen (15 μ L) was mixed with the sample loading buffer (5 μ L) and boiled before loading to 4% separating and 10% resolving gel. Low molecular weight markers (LMW) with a range of 14.4–97 kDa and high molecular weight markers (HMW) with a range of 53–220 kDa (GE Healthcare, USA) were used as a marker. The electrophoresis was conducted for approximately 2 h at a constant voltage of 125 V. The gel was stained with Coomassie blue staining.

2.9. Determination of Antioxidant Activity

The antioxidant activity of the collagen hydrolysates was analyzed based on DPPH radical scavenging activity, according to Khirzin et al. (2015) with modification. The hydrolysates were dried using a vacuum concentrator. Stock solution (20 mg/mL) was firstly prepared by dissolving 20 mg sample in one milliliter phosphate buffer (pH 7) because the sample did not completely dissolve in methanol. A series concentration of samples (1; 2; 4; and 8 mg/mL) was prepared by diluting the stock solution using methanol. Each sample (160 μ L) was placed into the 96-well microplate and added with 40 μ L of 0.78 mM DPPH solution (methanol as solvent). The mixture was incubated at ambient temperature for 30 mins in the dark room and measured their absorbance at 517 nm. The negative control used was methanol with DPPH addition, methanol solution was used as blank, whereas ascorbic acid with a concentration of 0.008 mg/mL was used as the positive control. The DPPH radical scavenging activity (%) was calculated using Equation 2.

$$\text{DPPH radical scavenging activity (\%)} = \left[\frac{((Ac)-(As-Ab))}{Ac} \right] \times 100\% \quad (2)$$

Where Ac is the absorbance of the control reaction, As is the absorbance of the sample, and Ab is the absorbance of the blank solution.

2.10. Statistical Analysis

All analyses of the collagen hydrolysates were carried out in duplicates. The data were presented as the mean \pm SD and analyzed by one-way ANOVA with a significance level at $p < 0.05$. The Duncan test was used to determine significant differences at the 5% probability level.

3. Results and Discussion

3.1. Identification of Sea Cucumber

The result of the phylogenetic analyses (Figure 1) showed that the golden sea cucumber used in this study, coded as *NF 1 Gab* and *NF 2 Gab*, had similarities with *S. horrens* and *S. monotuberculatus*. The COI similarity result generated by BLAST analysis indicated that the similarity was 99%. This result supported Purcell et al. (2012) findings that *S. horrens*, *S. monotuberculatus*, *S. quadrifasciatus*, and *S. naso* are often indistinguishable due to their similar appearances. The local name golden sea cucumber or “teripang emas” are commonly known as *S. variegatus* which are found in some publications (Adriansyah & Sulastris, 2014; Cahyati et al., 2018) or *S. hermanni* (Arundina et al., 2015; Damaiyanti, 2015; Safithri et al., 2018; Susanto, Savitrim, & Tarman, 2018). However, it was also reported that the *S. hermanni* was previously known as *S. variegatus* (Pangestuti & Arifin, 2018). The present study gave different results that the golden sea cucumber obtained from Lombok island, West Nusa Tenggara was closer to *S. horrens* and *S. monotuberculatus*. The previous study showed that a phylogenetic tree constructed by the same method has successfully separated *S. horrens*, *S. hermanni*, and *S. monotuberculatus* (Patantis et al., 2019), which proved that the phylogenetic tree of COI sequences could also differentiate *Stichopus* at the species level.

3.2. Chemical Composition of Sea Cucumber and Yield of Collagen

The sea cucumber chemical composition used in this study is presented in Table 1, along with the proximate composition of other *Stichopus* sp. that had been reported previously. The wet base (wb) data was converted into the dry base (db) for comparison purpose. It is shown that the sea cucumber moisture and protein contents used in this study are similar to those of other species of *Stichopus*. At the same time, the fat content of the sample was relatively lower (< 2% db) compared to that of other *S. horrens* as well as other *Stichopus* sp. reported. This result was used as a preliminary data for non-alcohol pre-treatment in the collagen extraction to remove fat instead of those conducted by Fawzya et al. (2016) and Nagai, Tanoue, Kai, and Suzuki (2015). The moisture, ash, protein and fat contents of *Stichopus* varied from 90.86 to 95.14; 31.69 to 56.12; 30.73 to 54.32 and 1.46 to 4.89%, respectively. The high ash content might be attributed to the high content of minerals and other

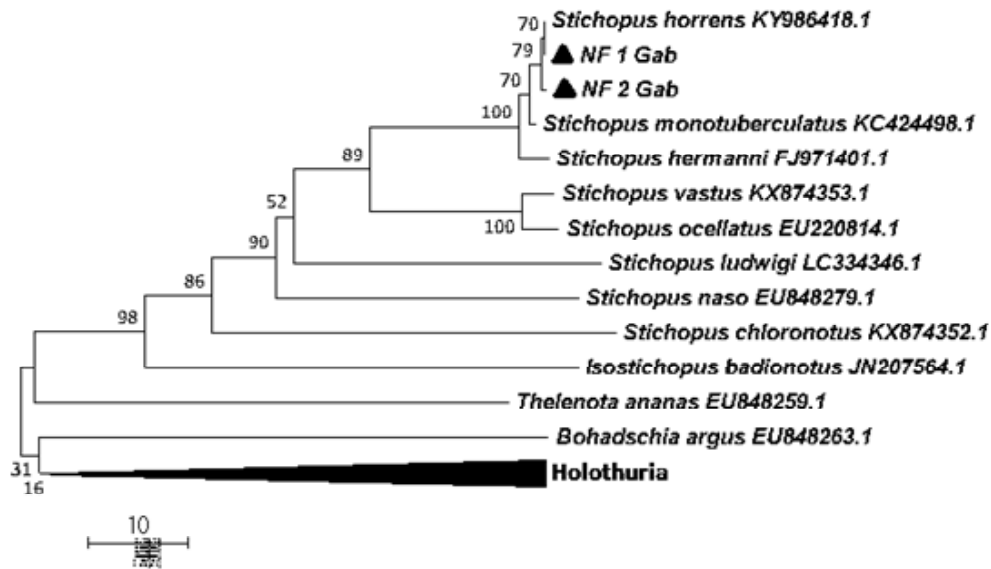


Figure.1. The phylogenetic tree of golden sea cucumber *beche-de-mer* from Lombok island, coded as *NF 1 Gab* and *NF 2 Gab*. The tree was created using the neighbor-joining method by testing bootstrap of 1000 replications. The numbers shown next to the branches showed bootstrap values.

Table 1. Chemical composition of golden sea cucumber

No	Sea cucumbers	Moisture (% wb)	Ash (% db)	Protein (% db)	Fat (% db)	Ref
1	<i>S. horrens</i>	94.51 ± 0.03	55.01 ± 0.08	34.97 ± 0.25	1.46 ± 0.03	Present study
2	<i>S. horrens</i>	93.10	39.13*	41.01*	3.04*	Forghani et al., 2012
3	<i>S. horrens</i>	95.14	31.69*	36.62*	4.11*	Gianto et al., 2017
4	<i>S. hermanni</i>	92.74 ± 0.36	40.94 ± 2.55	30.73 ± 2.40	4.89 ± 0.24	Mariyanti, 2017
5	<i>S. variegatus</i>	91.19 ± 0.07	40.18 ± 1.93	54.32 ± 0.68	2.72 ± 0.23	Fawzya et al., 2016
6	<i>S. variegatus</i>	93.84 ± 0.09	43.18 ± 1.95*	43.67 ± 0.49*	2.92 ± 0.10*	Alhana et al., 2015
7	<i>S. ocellatus</i>	93.80	56.12*	35.32*	3.06*	Rasyid & Ardiansyah, 2016
8	<i>Apostichopus japonicus</i>	90.86 – 91.19	32.22 – 33.90	45.12 – 46.10	3.87 – 5.11	Wu et al., 2015

*Converted into %db

inorganic matter in the sea cucumber's body wall. It might be related to skeletal elements, called ossicles, within cellular vesicles of the sea cucumber's body wall (Blowes et al., 2017). Ossicles are composed mainly of calcium carbonate (Bechtel, Oliveira, Demir, & Smiley, 2013). Variation of each species' proximate composition might be affected by several factors, such as seasonal variations, geographical location, and feeding system (Oh, Ko, Lee, Heo, & Jung, 2017; Santos et al., 2015). Procedures of handling also could influence sea cucumbers' chemical composition (Özer, Mol, & Varlýk, 2005).

Collagen extraction from the golden sea cucumber produced of 16.70% of crude collagen (% wb) or 7.20% of dialyzed collagen (% wb) (Table 2). After being hydrolyzed by neutrase, dried collagen hydrolysates' yield was in the range of 0.83-0.88% (% wb) or equivalent to 15.12-16.03% (% db). Other research showed that dried collagen extracted from *S. vastus* and *H. atra* produced yield of 0.92% and 0.88% wb (Yuniati & Sulardiono, 2019). Meanwhile, the dried collagen extracted from golden sea cucumber (*S. hermanni*) had a lower yield, i.e., 0.66% (Safithri et al., 2018). In this research, the dialyzed collagen was

Table 2. Yields of crude collagen, dialyzed collagen and collagen hydrolysates extracted from the body wall of golden sea cucumber*

Product	Yield (% wb) ¹	Yield (% wb) ^{2,3}
Crude collagen ¹	16.70 ± 4.24	
Dialyzed collagen ^{1,2}	7.20 ± 0.36	44.27 ± 9.12
Collagen hydrolysates from hydrolysis time (min) ^{1,3} :		
30	0.87	11.61
60	0.84 ± 0.08 ^a	11.67 ± 0.54 ^a
120	0.83 ± 0.02 ^a	11.29 ± 0.20 ^a
180	0.88 ± 0.05 ^a	12.24 ± 0.14 ^b
240	0.87 ± 0.06 ^a	12.02 ± 0.17 ^a

* Analyzed using one-way ANOVA

^{a,b}Different letters show significant differences in the same column ($p < 0.05$)

¹⁾ percentage of fresh sea cucumber

²⁾ percentage of crude collagen

³⁾ percentage of dialyzed collagen

directly hydrolyzed by neutrase without drying. The difference yield was most likely related to the variation method of collagen extraction and sea cucumber species as the source of collagen. In this work we omitted several pre-treatment steps to improve product yield and process efficiency. Collagen extraction consists of several steps, including pretreatment using certain chemicals intended to remove non-collagen protein, fats and other impurities. The difference of collagen yield was probably affected by the chemical concentration, ratio of the substance to the solvent and the extract separation method. The inappropriate concentration, ratio and separation method (such as centrifuge, pressing or filtering using a specific size of planktonet) will reduce the extraction effectiveness, which further produces a lower yield.

3.3. Degree of Hydrolysis (DH) and Peptide Content

The level of collagen cleavage of golden sea cucumber by neutrase was described on the degree of hydrolysis results and peptide content. The degree of hydrolysis (DH) is defined as the proportion of peptides bond digested to the total peptides bond. Figure 2 shows that DH increases rapidly in the first hour of hydrolysis, then the rate declined slowly in the following hours. DH's rise showed that a quantity of free amino groups of the protein increased and the

change of protein molecular weight occurred. The highest DH was achieved after 180 mins of hydrolysis, reaching 55.2% of DH value ($p < 0.05$). After three hours, the DH decreased because of the product inhibition and/or enzyme denaturation.

It was reported that the hydrolysis of *S. horrens*' tissue was more effective when using alkalase than other proteases (Forghani et al., 2012). This enzyme (2103 U/mL) produced the highest DH (39.8%) in 200 mins and the hydrolysate had the highest ACE inhibitory activity. The use of neutrase with the total activity of 2.01×10^5 U produced DH of Alaska pollock hydrolysates of approximately. 25% and reached equilibrium at 160 mins hydrolysis (Liu et al., 2018). They stated that neutrase was the most efficient enzyme to improve protein solubility compared to six food-grade proteases. It seems that DH resulting from the hydrolysis process was affected by the type and concentration of enzymes as well as type of substrates.

The increase of DH was followed by peptide increment (Figure 3) due to the release of peptides as a result of neutrase activity during the hydrolysis process. The correlation test exhibited that peptide content was strongly correlated to DH (R^2 0.79, data not shown). However, the increment of peptide content was insignificant after 180-mins of hydrolysis, reaching 12.79 ± 0.44 mg/mL of peptides. This might be related

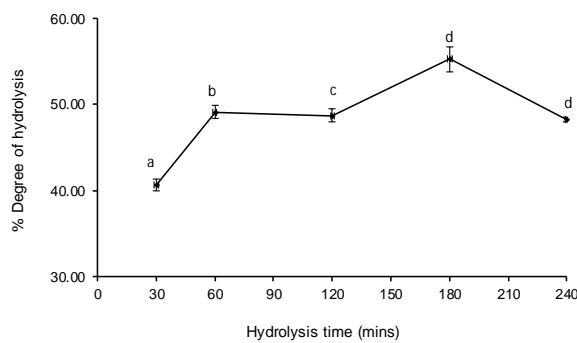


Figure 2. Degree of hydrolysis of collagen hydrolysates produced from golden sea cucumber

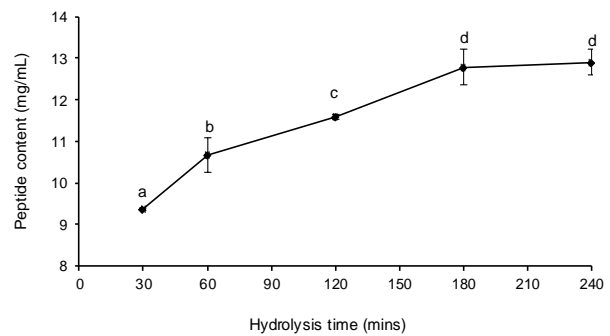


Figure 3. Peptide content of collagen hydrolysates produced from golden sea cucumber

to one (or more) of these reasons: the decrease of enzyme activity to hydrolyze substrate (collagen hydrolysate), the reduction in the amount of accessible peptide bonds as the substrate to be digested or inhibition of the hydrolyzed product to the remaining substrate (Guérard, Dufossé, De La Broise & Binet, 2001). Putalan, Munifah, Nurhayati and Chasanah (2018) reported the similar pattern that the highest DH and peptides on hydrolysis of striped trevally fish (*Selaroides leptolepis*) using *B. licheniformis* protease was reached after six hours of hydrolysis.

3.4. Molecular Weight Distribution

The SDS-PAGE profile generated from the hydrolysates is presented in Figure 4. It is indicated that the initial unhydrolyzed collagen (line 1) showed a thick and clear band with a size of approx. 220 kDa. The addition of neutrase hydrolyzed collagen rapidly resulted in hydrolysates with molecular weight ranging from less than 14.4 kDa to 53 kDa. Hydrolysis for 30 mins exhibited three visible thin bands with the size of approx. 53; 45 and 25 kDa, meanwhile 1 h hydrolysis produced hydrolysates with molecular weight of 25; 14.4 and under 14.4 kDa. The band almost completely disappeared after two hours of hydrolysis. This is probably related to the neutrase as a neutral endopeptidase which randomly digests peptide bonds of non-terminal amino acids (Anonymous, 2016) and aids other protein hydrolysis (Hong, Min, & Jo, 2019). The decrease of molecular weight occurred in accordance with the hydrolysis time.

This pattern was similar to the molecular weight distribution of porcine skin collagen hydrolysates treated with neutrase (Hong et al., 2019). They investigated hydrolysate produced from the hydrolysis process with the enzyme:substrate ratio of 1:100 had disappeared entirely after six hours of incubation. They

further mentioned that protease type, hydrolysis time and the ratio of enzyme/substrate affected the digestion pattern of protein into peptides and amino acids. Liu et al. (2018) observed that complete degradation of the Alaska pollock protein isolate by neutrase occurred at 40 to 240 mins of hydrolysis with the average molecular weights produced were less than 10 kDa. At the final hydrolysis time (240 mins), small peptides (500 Da) were produced. According to Hong et al. (2019), molecular weight exhibited a level of fractionation. A fraction with MW of < 3 kDa indicates a collagen hydrolysate; a fraction with MW greater than 50 kDa is gelatin; and a fraction with MW greater than 300 kDa is collagen. Thus, a lower molecular weight marker (less than 10 kDa) is necessary for SDS-PAGE or HPLC analysis for precise results.

3.5. Antioxidant Activity

The activity of collagen hydrolysate produced by the golden sea cucumber was assayed by the DPPH radical scavenging activity method with the concentration of 1; 2; 4 and 8 mg/mL. Ascorbic acid was used as a positive control with the concentration of 0.004; 0.006; 0.008 and 0.010 mg/mL. The results showed that the antioxidant activity of samples were increased in a concentration-dependent manner. Further determination of IC_{50} indicated that the hydrolysate produced from 60-min hydrolysis had the lowest IC_{50} (5.25 ± 0.15 mg/mL) those that of the other hydrolysates, which means that the antioxidant activity was the highest. However, the IC_{50} value was still much higher than that of ascorbic acid (0.0082 ± 0.0001 mg/mL) as control. The antioxidant activity of ascorbic acid, equivalent with $46.5 \mu\text{M}$ (MW of 176.12 Da), was 4.5 up to 8-fold than that of collagen hydrolysate produced from 60-min hydrolysis (210 and 375 μM , respectively) with an estimated molecular weight of 25 and 14 kDa (Figure 4). Thus, the collagen

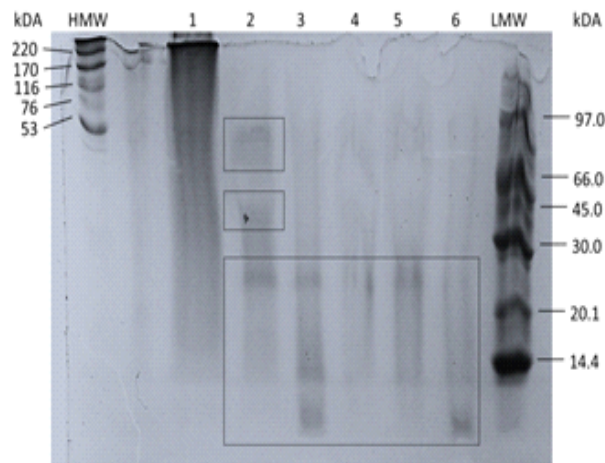


Figure 4. Molecular size distribution profile of golden sea cucumber collagen hydrolysates produced by neutrase hydrolysis. HMW: High Molecular Weight; 1: Unhydrolyzed collagen; 2: Hydrolysis 30 min; 3: Hydrolysis 60 min; 4: Hydrolysis 120 min; 5: Hydrolysis 180 min; 6: Hydrolysis 240 min; LMW: Low Molecular Weight. SDS-PAGE was done using a separating gel concentration of 15% at 125 Volt

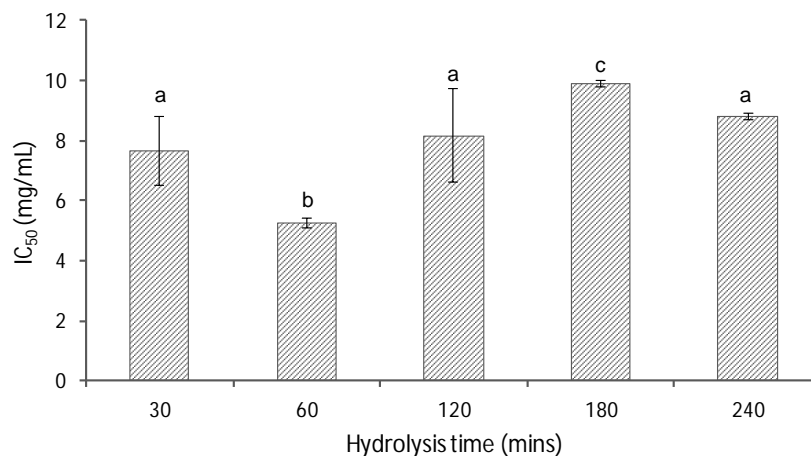


Figure 5. Antioxidant activity of collagen hydrolysates derived from golden sea cucumber (mean values with different letters indicate significant differences ($p < 0.05$))

hydrolysate may be classified as a weak to moderate antioxidant, even though there is no classification for IC_{50} values that can be referred.

To date, protein hydrolysates with strong antioxidant activity are rarely reported. However, low MW protein hydrolysates from yellowfin tuna skin (3-10 kDa) and the spotted Babylon snail (3-5 kDa) were reported as potent antioxidants with IC_{50} of 75.94 $\mu\text{g}/\text{mL}$ and 2.69 $\mu\text{g}/\text{mL}$, respectively (Nurilmala, Hizbullah, Karnia, Kusumaningtyas, & Ochiai, 2020; Petsantad et al., 2020). Liu et al. (2018) investigated that hydrolysis of the Alaska pollock protein isolate using neutrase significantly improved the DPPH free radical-scavenging activity. The inhibition of 5 mg/mL hydrolysate produced from 240-min incubation reached 85.22%. They further found that the longer the

hydrolysis time, the lower the molecular weight and the higher the antioxidant activity. In this study, the highest antioxidant activity exhibited by hydrolysate produced from 60-mins hydrolysis with the DH reached 49.11% and peptide content of 10.67 mg/mL. The probable reason was that the amino acid composition and sequence contributed to the antioxidant activity were present higher in the 60-min hydrolysate than those in the other counterparts.

According to You, Zhao, Regenstein, & Ren (2010), hydrophobic amino acids and residues of proline, cysteine, methionine, histidine, tyrosine, phenylalanine, and tryptophan could improve the antioxidant activity of peptides available in fish protein hydrolysates. Other factors that may affect the inhibitory activities of protein hydrolysates, including

size and structure of peptides, DH, molecular weight, etc. (Cheung, Cheung, Tan, & Li-Chan, 2012; Raghavan & Kristinsson, 2008). DPPH radical is a stable compound that can react with various antioxidants. The DPPH method is also simple and relatively inexpensive, making it used broadly to determine the antioxidant activity of many samples from various resources (Santos-Sánchez, Salas-Coronado, Villanueva-Cañongo, & Hernández-Carlos, 2019).

4. Conclusion

Golden sea cucumber obtained from West Nusa Tenggara was identified as *Stichopus horrens* or *S. monotuberculatus*. The sea cucumber collagen hydrolysis using neutrase produced hydrolysates with molecular weight less than 53 kDa and smaller peptide products in line with the increase of hydrolysis time. The hydrolysate produced after 60 mins of hydrolysis demonstrated the highest radical scavenging activity with IC_{50} of 5.25 ± 0.15 mg/mL that might be classified as a weak to moderate antioxidant. However the highest DH and peptides were produced after 180 mins of hydrolysis.

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