

**Research Article** 

# Antioxidant Activity and Potential Bioactive Peptides from Skin Protein Hydrolysate of Yellowfin Tuna (*Thunnus albacares*)

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# Abstract

The tuna fillet industry produces abundant skin by-products with high protein, which has the potential as a raw material for fish protein hydrolysate (FPH) for a source of bioactive peptides. Exploration of bioactive peptides from fish skin is generally from hydrolyzed gelatin and collagen. The study aimed to produce FPH directly from tuna skin as an antioxidant and identify potential bioactive peptides. The research began by defatting using multiple concentrations of NaH-CO<sub>2</sub> and immersion times. The defatted fish skin was produced as FPH by enzymatic hydrolysis method using different papain enzyme concentrations and hydrolysis times. The selected treatments were assessed for antioxidant activity and bioactive peptides. The results showed that the defatting process using a 0.50% NaHCO<sub>3</sub> for 30 minutes generated the highest reducing fat content in value was 80.53%. Using papain enzyme gave a significant effect on the DH of FPH, with value was 29.72-67.64%. Therefore, FPH obtained from different concentrations for 4 hours was chosen to characterize the antioxidant activity and bioactive peptide. Hydrolysis using 5% enzyme papain showed the highest antioxidant activity of DPPH, ABTS, and reducing power with values of 0.965 mg/mL, 0.495 mg/mL, and 0.415 absorbances, respectively. FPH possesses a molecular weight of 10.15-48.50 kDa. Functional groups detected were amides A, B, I, II, and III. Glycine, proline, and arginine became amino acids dominant of FPH. Based on the diversity of biopeptide compounds, several biological function candidates were detected, namely antioxidants, ACE inhibitors, and antifibrinolytics which have the potential to be used as nutraceutical products.

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## **1. Introduction**

The tuna industry is the most complex and dynamic industry; dominating the export of fishery products in Indonesia. The main export products for tuna are divided into four categories: canned tuna of 82,234 tons, frozen fillet of 58,871 tons, frozen tuna of 24,261 tons, and fresh tuna of 3,067 tons of a total export volume of 168,433 tons with a value of US\$ 713,9 million, generally originating from yellowfin tuna (KKP, 2020). The high total export volume of processed products will undoubtedly increase the volume of by-products. The tuna fillet industry has the highest export volume after canning products, which can be utilized because of the characteristics of fresh by-products compared to canned by-products that have undergone a heating process (Irianto and Akbarsyah, 2007; Gamarro et al., 2013). Several studies have shown that the most effective protein content in yellowfin tuna by-products is found in the skin 27.12-36.09% (Karunarathna and Attygalle, 2010; Nurilmala et al., 2019), in dark meat, 20.22-26.92% (Karunarathna and Attygalle, 2010; Sánchez-Zapata et al., 2011) and the minor protein content in the swim bladder is 17.52-19.17% (Hadinoto and Idrus, 2018; Hadinoto et al., 2019). The high protein content in tuna skin shows the potential to be utilized as a raw material for fish protein hydrolysate (FPH) products which can be value-added products.

Enzymatic hydrolysis is the best way to hydrolyze proteins with minimum losing the nutritional value of raw materials, the easiest to control in producing, and can improve the functional and biological properties of raw materials (Huang et al., 2016; Schmidt et al., 2020). The fat content in raw materials will affect the hydrolysis, requiring a defatting process to get the best FPH. In order to reduce the fat content is generally done by chemical and mechanical processes (Hua et al., 2020). FPH is used as an alternative protein source in commercial products (Lopera et al., 2018) and is also used as a source of additional nutrition (Kiewiet et al., 2018; Asmak et al., 2020), flavor enhancer (Vijaykrishnaraj et al., 2016; Witono et al., 2019) and to improve product characteristics (Jeewanthi et al., 2015). FPH is proven as a bioactive component source in the form of bioactive peptides released during hydrolysis. These bioactive peptides have positive effects on health due to biological activities, so they have the potential to be used as functional and nutraceutical food products (Li-Chan, 2015).

Bioactive peptides have very diverse functional properties and is determined by the constituent amino acid sequence (Baehaki *et al.*, 2020). Several studies report that FPH has functional properties as glucoregulatory and anti-inflammatory (Henaux et al., 2021), antihypertensive (Naik et al., 2021), anticancer (Yaghoubzadeh et al., 2020), antibacterial (Baco et al., 2022), antidiabetic (Rothwell et al., 2021), and antioxidants (Qiu et al., 2019; Baehaki et al., 2020; Agustin et al., 2021). Research related to the exploration of bioactive peptides from fish skin by-products is generally in the form of hydrolyzed protein products in gelatin products (Gonz et al., 2022) and collagen products (Prastyo et al., 2020; Tkaczewska et al., 2020). Exploration of bioactive peptides by direct hydrolysis fish skin is still limited so this research is essential to evaluate the potential of bioactive peptides in tuna skin FPH. Thus, this study has several objectives, first to get the best treatment in the defatting process. Second, to evaluate the effect of enzyme concentration and time on the degree of hydrolysis (DH) attribute as well as to assess the antioxidant activity and to identify bioactive peptides.

## 2. Materials and Methods

#### 2.1 Material

The main material in this study was tuna skin obtained from PT. Bumi Menara Internusa, Makassar, Indonesia. The material used for the defatting process was NaHCO<sub>3</sub> powder (Merck, Germany). The protein hydrolysate production was performed by the enzyme papain (EC3.4.22.2) commercial (Himedia, India) activity 1,000 units/mg. The chemicals used for analysis were 2,2-Diphenyl-1-Picrylhydrazyl (DPPH; Sigma-Aldrich, St. Louis, MO, USA), 2,2-Azinobis 3-ethyl benzothiazoline 6-sulphonic acid (ABTS; Sigma-Aldrich, St. Louis, MO, USA), ethanol (Merck, Germany), phosphate buffer (Sigma-Aldrich, St. Louis, MO, USA), K<sub>3</sub>Fe (CN)<sub>6</sub> (Merck, Germany) and FeCl<sub>3</sub> (Merck, Germany).

## 2.2 Tuna Skin Preparation and Defatting

Skin obtained from a by-product of the tuna industry was thawed and washed to remove dirt that sticks to the surface. Then it was cut into  $2x2 \text{ cm}^2$  pieces and crushed, then analyzed for proximate composition (AOAC, 2005). The mashed skin was then subjected to a defatting process according to Ramadhan *et al.* (2014), which was modified by immersion in NaHCO<sub>3</sub> solution with a concentration of 0%, 0.25%, 0.5%, and 0.75% (w/v) for 30, 60, and 90 minutes, respectively, using a 1:4 ratio of sample and NaHCO<sub>3</sub> solution at 4°C with periodic stirring. A fat content test was performed (AOAC, 2005) to determine the best NaHCO<sub>3</sub> concentration and immersion time.

## 2.3 Production of Protein Hydrolysate

Hydrolysate production based on Mutaminah et al. (2018) was modified with different concentrations of papain enzymes 4, 5, and 6% (v/v) at four, five, and six hours. The ratio of sample and distilled water 1:2 (w/v), pH 7 hydrolysis with a water bath shaker (BS-21, Jeio Tech, Korea) at 55°C, speed 100xg. The incubation process is complete followed by the inactivation process used, water bath (DWB-18, B-One, China) at 85°C and centrifuged (Sorval Lynx 6000, Thermo Scientific, Germany) at 10,000xg temperature 4°C, 20 minutes. Then the supernatant was collected and stored in a dark bottle at -15°C. The selected treatment was determined by testing the degree of hydrolysis (DH). The filtrate was mixed with 10% TCA with a ratio of 1:1 (v/v) and centrifuged at 7800xg for 15 minutes) (Hoyle and Merritt, 1994). The nitrogen content of the supernatant was measured using the Kjeldahl method (AOAC, 2005). This formula can calculate the degree of hydrolysis (%):

Degree of  
hydrolysis (%) = 
$$\frac{\text{TCA dissolved nitrogen 10 \%}}{\text{Sample total nitrogen}} \times 100\%... Eq 1$$

#### 2.4 Evaluation of Antioxidant Activity

The filtrate obtained from hydrolysis was dried using a freeze dryer (Missouri 64132, Labconco, USA) at -50°C, 0.5 Mbar pressure, and analyzed for antioxidant activity of DPPH, ABTS, and reducing power.

#### 2.4.1 DPPH radical scavenging activity

The 1.5 mL sample solution was added to 1.5 mL of 0.15 mM DPPH in 95% ethanol incubated in the dark for 30 minutes and the absorbance was measured at a wavelength of 517 nm (Wu *et al.*, 2003). Percent DPPH inhibition is calculated using the following formula:

DPPH  
inhibition (%) = 
$$\frac{A \text{ control} - A \text{ sample}}{A \text{ control}} \times 100 \dots \text{ Eq } 2$$

The sample used was dissolved with various concentrations to obtain the  $IC_{50}$  value representing the concentration of the sample solution required to reduce DPPH free radicals by 50%. The linear regression equation (y = a + bx) obtained from the inhibition value at each concentration is used to find the  $IC_{50}$  value.

## 2.4.2 ABTS radical scavenging activity

The ABTS radical scavenging activity was

performed according to the method by Stämpfli *et al.* (2007) with slight modifications. ABTS stock solution was prepared by reacting 7 mM ABTS solution and 2.45 mM potassium persulfate in 1:1 ratio in a dark room for 16 hours at room temperature. The solution was then added with ethanol to obtain an absorbance of  $\pm 0.785$  at a wavelength of 750 nm. A total of 1 mL of ABTS solution was added to 1 mL of the hydrolyzed sample. It was left for 10 minutes at room temperature. Absorbance was measured at a wavelength of 750 nm. The percentage of ABTS radical inhibition was calculated based on the formula:

ABTS inhibition (%) = 
$$\frac{A \text{ control} - A \text{ sample}}{A \text{ control}} \times 100... \text{ Eq } 3$$

The sample used was dissolved with various concentrations to obtain the  $IC_{50}$  value representing the concentration of the sample solution required to reduce ABTS free radicals by 50%. The linear regression equation (y = a + bx) obtained from the inhibition value at each concentration is used to find the  $IC_{50}$  value.

#### 2.4.3 Reducing power

Evaluation of reducing power was carried out using 1 mL sample (5 mg/mL) added to 2 mL of 0.2 M phosphate buffer (pH 6.6) and  $K_3Fe$  (CN)<sub>6</sub> and then incubated for 20 minutes at 50°C. After incubation, the sample was added with 1 mL of 10% TCA and centrifuged at 3500xg for 10 minutes. The 1 mL supernatant was added with 1 mL distilled water and 200 µL FeCl<sub>3</sub>. After 10 minutes of incubation, the absorbance was measured at a wavelength of 700 nm (Oyaizu, 1988).

## 2.5 Molecular Weight Analysis

The molecular weight of the samples was determined using the SDS-Polyacrylamide Gel electrophoresis method based on Laemmli (1970) with a concentration of 12% acrylamide gel as separator gel and 5% acrylamide gel as collector gel. The sample tested was tuna skin protein hydrolysate; 20 µL of the sample was mixed with 10 µL of sample buffer containing 2-mercaptoethanol, then heated at 85°C for five minutes, and then put into the well of the gel. Ten  $\mu$ L of the marker was included in the gel well. The gel was mounted in the electrophoresis apparatus by pouring the electrophoretic buffer into the chamber. The electrophoresis process lasted for 180 minutes at 180 volts and 50 mA in the electrophoresis apparatus, after which the gel was removed from the glass plate. The gel was soaked in Coomassie blue gel stain solution for two hours, followed by distaining until protein bands were obtained.

## 2.6 Functional Group Analysis

Functional group analysis was carried out using the *Fourier-transform infrared spectroscopy* (FTIR) principle based on Qiu *et al.* (2019) using an infrared spectrophotometer (Bruker Tensor 37, Ettlingen, Germany). A 2 mg sample was mixed with 100 mg KBr and pulverized. Then the sample was measured using the FTIR instrument at a wavelength of 500 - 4000 cm<sup>-1</sup>. FTIR instruments will detect absorption peaks at certain wave numbers. The wave numbers detected as peaks are identified based on the specific absorption region range of the protein functional groups.

## 2.7 Amino Acid Composition Analysis

The type of amino acid was characterized to determine the amino acid type in the sample. The instrumentation used was Ultra Performance Liquid Chromatography (UPLC) ACCQ-Tag Ultra C18 column, temperature 49°C, and mobile phase gradient composition system. The standard amino acids used in this analysis consist of 15 types of amino acids. A total of 100 mg of sample was weighed and put into a 20 mL headspace vial. Hydrolysis was carried out with an HCl solution. Then hydrolysate was removed and transferred into a 50 mL measuring flask, added distilled water the tera mark, and homogenized. The test solution was filtered with a 0.2 µm syringe filter to collect the filtrate. A total of 1 µL was included in the UPLC instrument (SIG, 2021). Calculation of amino acid levels in the sample using the ratio of the analyte area to the internal standard, with the following formula:

Standard or  
sample ratio = 
$$\frac{\text{The area of the amino acid analyte}}{\text{AABA standard internal area}}$$
 ... Eq 4

Amino acid  
content = 
$$\frac{\frac{\text{sample ratio}}{\text{standard ratio}} \times \frac{C_{\text{std}}}{1000000} \times \text{MW x Va x DF}}{W_{\text{spl}} \text{ or } V_{\text{spl}}} \dots \text{ Eq 5}$$
where:

MW = Amino acid molecular weight (g/mol)

- C<sub>std</sub> = Concentration of standard amino acid solutions (pmol/µL)
- Va = The final volume of the test solution  $(\mu L)$
- DF = Dilution factor

 $W_{Sol}$  = Test portion weight (g)

 $V_{Snl}$  = Volume of test portion (mL)

## 2.8 Diversity of bioactive peptides

The best protein hydrolysate was identified as a bioactive peptide using LC-MS, referring to the method

of Suetsuna et al. (2000). A 5 µL hydrolyzed tuna skin was injected into the LC-ESI-QTOF instrument. LC-MS analysis was performed using a UPLC-MS equipped with a binary pump. The LC was connected to a QTOF mass spectrometer with an ESI ionization source. The MS used was a Xeco G2-S TOF system with a positive ionization mode. The ESI parameters included a capillary temperature of  $120^{\circ}C$  and a 500 L/ hour fogging gas with a voltage source of 3 kV. The full scan mode from m/z 100-1500 was carried out with a source temperature of 110°C. The UPLC column used was Acquity UPLC HSS C18 1.8 µM×2.1×150 mm. The eluent was  $H_2O + 0.1\%$  formic acid (A) and 0.1%formic acid + acetonitrile (B). The eluent was set at a 0.3mL/min flow rate. The elution system was run isocratic at 0-1 minute with a ratio of 95:5, 0-5 minutes with a linear gradient elution of solvent A from 95% to 5%, 6-7 minutes isocratic elution with a ratio of 0:100, 6-7 minutes with a gradient elution linear solvent A from 0% to 100%, minutes 7.5-9 isocratic elution ratio 95:5, minutes 7.5-9 linear gradient elution solvent A from 95% to 5%. The resulting chromatogram data were converted to NetCDF format to make it easier to process data with Mzmine. The results obtained were in the form of mass array data from peptide hydrolysate chromatograms which included three variables, namely m/z, retention time, and peak intensity. Identification of the metabolite compounds contained in the samples was carried out by matching the m/z of the resulting chromatogram with the accurate mass database for the peptide compounds contained in tuna skin hydrolysate.

## 2.9 Data Analysis

The research design at the stages of defatting and protein hydrolysis used a Completely Randomized Design Factorial (CRDF), while for the determination of antioxidant activity a Completely Randomized Design (CRD) was used. Each result of the analysis of variance was significantly different; further testing was carried out with the Tukey test using Minitab software.

## 3. Results and Discussion

#### 3.1 Results

#### 3.1.1 The proximate composition of tuna skin

Yellowfin tuna skin as raw material for FPH is obtained from tuna fillet processing companies with an average fish weight of 15–70 kg/head. Several studies reported the proximate composition of scombroid fish skin (Table 1). Characteristics of raw materials can be determined by analyzing the proximate composition especially protein content, is of concern in this research.



**Figure 1**. Fat content with defatting treatment in the concentrations of NaHCO<sub>3</sub> and immersion time at 30 (**—**); 60 (**—**); 90 (**—**) minutes. Different superscripts show significant differences (p<0.05).



**Figure 2**. The degree of hydrolysis in the treatment of differences in the concentration of the enzyme papain and the duration of hydrolysis at 4 ( $\blacksquare$ ); 5 ( $\blacksquare$ ); 6 ( $\blacksquare$ ) hours. Different superscripts show significant differences (p <0.05).

	Yellowfin tuna		Mackerel tuna <sup>1</sup>		Skipjack <sup>2</sup>		Yellowfin tuna <sup>3</sup>	
Composition	(Thunnus albacares)		(Euthynnus affinis)		(Katsuwonus pelamis)		(Thunnus albacares)	
	(ww)	(dw)	(ww)	(dw)	(ww)	(dw)	(ww)	( <b>dw</b> )
	%	%	%	%	%	%	%	%
Moisture	$69.18\pm0.06$	-	71.66	-	59.6	-	57.42	-
Protein	$26.77 \pm 1.08$	81.5	24.63	87.43	28.31	70.07	37.45	87.95
Fat	$1.69\pm0.29$	5.2	2.72	9.65	7.45	18.44	3.8	8.92
Ash	$0.20\pm0.05$	0.59	0.17	0.6	1.33	3.29	0.49	1.15

Table 1. The proximate composition of scombroid fish skin

Description: <sup>1</sup>Komala et al. (2015), <sup>2</sup>Karunarathna and Attygalle (2010), <sup>3</sup>Nurjanah et al. (2021)

(ww=wet weight; dw=dry weight). Data represented as means  $\pm$  SD (n=3)

#### 3.1.2 Decreased fat content of tuna skin

The defatting method in this study was carried out by immersing the samples in various concentrations of NaHCO<sub>3</sub>, low temperature (4°C), and immersion time (Figure 1). The analysis of variance showed that the concentration of NaHCO<sub>3</sub>, immersion time and their interactions significantly affected the decrease in the lipid content of the tuna skin (p<0.05). The alkaline salt immersion treatment results were proven to remove 45-85% fat in the tuna skin.

**Table 2.** Antioxidant activity at different enzyme concentrations

Enzyme concen- tration (%)	DPPH IC <sub>50</sub> (mg/mL)	ABTS IC <sub>50</sub> (mg/mL)	<i>Reducing</i> <i>power</i> (Absor- bance 700 nm)
4	$1.277 \pm 0.016^{b}$	$0.647 \pm 0.016^{b}$	$0.415 \pm 0.007^{a}$
5	$0.964 \pm 0.002^{a}$	$0.495{\pm}0.002^a$	$0.415 {\pm} 0.001^{a}$
6	$0.963 \pm 0.001^{a}$	$0.491 \pm 0.001^{a}$	$0.416 \pm 0.004^{a}$

Data represented as means  $\pm$  SD (n=3). Different superscripts in the same column show significant differences (p < 0.05)

**Table 3.** Characteristics of tuna skin hydrolysate

 functional groups

Amide	Absorption area (cm <sup>-1</sup> )	Absorp- tion peak (cm <sup>-1</sup> )	Explanation
Amide A	3400 - 3440ª	3430	N-H stretching
Amide B	2922 – 2940 <sup>b</sup>	2934	C-Hasymmetrical stretching
Amide I	1625 – 1690°	1650	C=O stretching, C-N stretching
Amide II	1500 – 1600 <sup>d</sup>	1542	N-H bending, C-N stretching, C=H stretching
Amide III	1229 – 1301e	1244	C-N stretching, N-H bending

Description: <sup>a</sup>Kaewdang *et al.* (2014); <sup>b</sup>Devita *et al.* (2021); <sup>c</sup>Kaewdang *et al.* (2014); <sup>d</sup>Rasli and Sarbon (2019); and <sup>c</sup>Kong and Yu (2007)



**Figure 3**. Molecular weight of marker (M), tuna skin hydrolysate with papain enzyme concentration 4% (A); 5% (B) and 6% (C)

#### 3.1.3 Degree of hydrolysis

The level of enzymatic hydrolysis of proteins was measured by the degree of hydrolysis (DH) (Figure 2). The DH value in the hydrolysis process was affected by the addition of different concentration of papain enzyme (p<0.05), while time of hydrolysis and interaction had no significant effect (p>0.05).

#### 3.1.4 Antioxidant activity

The antioxidant activity was evaluated using three simple spectrophotometric methods (DPPH, ABTS, and reducing power) (Table 2). Analysis of variance showed a significant effect (p<0.05) with the addition of different concentration of papain enzyme on the value of antioxidant activity in the DPPH and ABTS methods. In the reducing power method, different concentration of papain enzyme did not have a significant effect (p>0.05).

## 3.1.5 Molecular weight profile

The molecular weight profile of tuna skin hydrolysate was determined using the SDS-PAGE (Figure 3). The results obtained were that the MW value tended to decrease with increasing the concentration of the papain enzyme in the hydrolysate in line with the increase in the DH value and the antioxidant activity obtained.

## 3.1.6 Function group of tuna skin hydrolysate

The polypeptide and protein units generally provided nine characteristic absorption peaks: amides A, B, and I–VII. The functional group spectrum analysis resulted from tuna skin hydrolysate samples using 5% enzyme concentration treatment for four hours (Table 3). The tuna skin hydrolysate functional groups showed absorption peaks in the amide A, B, I, II, and III absorption areas (Figure 4).

## 3.1.7 Amino acid composition of tuna skin hydrolysate

The amino acid composition was essential in determining the bioactivity of protein products. Amino acids of tuna skin, tuna skin hydrolysate, and collagen hydrolysate showed differences in total amino acids (Table 4). Tuna skin hydrolysate with the highest antioxidant activity had the highest amino acid composition in hydrophobic amino acids.

## 3.1.8 Bioactive peptide diversity

LC-MS is an integrated proteomics system for analyzing a component in a material. The results of LC-MS analysis related to peptides, molecular weight, and biological activity of tuna skin hydrolysate (Table 5). Based on various molecular and peptide formulas in tuna skin hydrolysate, various bioactive potentials are obtained as antibacterial, ACE inhibitor, antioxidant and antifibrinolytic.

## 3.2 Discussion

Fish are classified into two categories based on their protein and fat content. Two categories of fish, namely high protein with, moderate fat and high protein with, low fat. Yellowfin tuna is a type of fish with high protein content and low fat (Bahurmiz, 2019). The protein and fat content of the yellow fin tuna skin produced were 26.77% and 1.69%, respectively. The value of this protein content was lower than of mackerel tuna skin (Komala et al., 2015), and higher than skipjack tuna skin (Karunarathna and Attygalle, 2010d), and yellowfin tuna reported by Nurjanah et al. (2021). Differences in proximate composition can be influenced by species, habitat, body size, environmental conditions, type of feed, and preparation of ingredients (Songchotikunpan et al., 2008; Porto et al., 2016). The protein content of yellowfin tuna skin obtained in this study is relatively high, making it possible to use it as raw material for FPH.



**Figure 4**. FPH infrared spectrum of tuna skin using 5% papain enzyme concentration. Amide spectrum: A, B, I, II, and III

	<b>GUA</b> 1	Tuna skin	Collagen	
Types of amino acids	Skin <sup>1</sup>	hydrolysate	hydrolysate <sup>2</sup>	
Lysine <sup>a</sup>	2.3	3.06	4.04	
Valine <sup>ab</sup>	1.4	2.37	2.3	
Isoleucine <sup>ab</sup>	1.06	1.23	1.13	
Leucine <sup>ab</sup>	1.97	2.63	2.94	
Histidine <sup>ac</sup>	0.86	0.82	-	
Threonine <sup>ac</sup>	1.51	3.76	3.44	
Phenylalanine <sup>ab</sup>	1.23	2.11	2.63	
Glycine <sup>b</sup>	6.18	21.92	23.86	
Alanine <sup>b</sup>	3.66	8.46	8.79	
Arginine <sup>c</sup>	3.24	8.94	10.56	
Aspartate <sup>c</sup>	2.82	4.22	4.66	
Glutamate <sup>c</sup>	5.18	8.41	8.42	
Proline <sup>b</sup>	-	9.76	11.82	
Serine <sup>c</sup>	1.13	3.91	4.13	
Tyrosine	0.59	0.49	0.97	
AA Essential	10.33	15.98	16.48	
AA Hydrophobic	11.84	48.48	44.68	
AA Hydrophilic	14.74	30.06	31.21	
Total amino acids	33.13	82.09	89.69	

 Table 4. Amino acid composition of tuna skin, tuna skin hydrolysate and collagen hydrolysate

Description: <sup>1</sup>Suseno (2015) and <sup>2</sup>Prastyo (2021); <sup>a</sup>essential amino acids, <sup>b</sup> hydrophobic amino acids, and <sup>c</sup> hydrophilic amino acids

# Table 5. Bioactive peptide diversity in tuna skin protein hydrolysate

Peptide Confirmed <sup>1</sup>	Molecular formula <sup>2</sup> [M+H] <sup>-</sup>	Molecular weight <sup>1</sup> (Da)	Structure Molecule <sup>1</sup>	Biological activity of peptide <sup>1</sup>
Gly-Val-Arg	$C_{13}H_{26}N_6O_4$	330.383		Inhibitor ACE
Aminocaproic	C <sub>6</sub> H <sub>13</sub> NO <sub>2</sub>	131.173	о тын;	Antifibrinolytic
Val-Asn-Pro	$C_{14}H_{24}N_4O_5$	328.364		Antioxidant

Description: <sup>1</sup>Chemspider Database 2022; <sup>2</sup>MassLynMassLynx

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The chemical composition of tuna skin shows a low-fat content, but the presence of this fat can inhibit the effectiveness of the hydrolysis process and the results obtained. Several chemical and mechanical methods have been developed for degreasing (Hua et al., 2020). The application of the alkaline salt immersion method in this study intends to avoid economic, environmental, and food safety problems. The immersion treatment in NaHCO<sub>2</sub> at a low temperature of 4°C was effective in reducing the fat content, which can reduce the fat content of 85% in tuna skin before the defatting process. This condition is caused by the alkalinity of NaHCO<sub>3</sub>, which can remove fat and open the matrix bond structure in the skin so that fat and other compounds can be quickly released (Cho et al., 2005; Hemung, 2013). Immersion with 0.5% NaHCO<sub>2</sub> (w/v) and soaking time of 30 minutes resulted in 0.33% fat content with a reduction percentage of 80.53% from before the defatting process, which was the selected treatment for the next stage based on the efficient use of time in production. The decrease in fat content shows higher results when compared to research by Sae-leaw et al. (2016) on seabass skin gelatin with 30% isopropanol, decreased fat 50.32% and in Xu et al. (2017) on skin of southern catfish (Silurus meridionalis Chen) with 10% NaHCO<sub>3</sub>, the fat reduction was 69.37%. Lower results when compared with reported by Ramadhan et al. (2014) using catfish fillet with 0.75% NaHCO<sub>3</sub>, soaking time 10 minutes, which reduced fat 89.64%. Nilsuwan et al. (2021) on fat in collagen hydrolysate with a disk stack centrifugal separator (DSCS) two cycles and the addition of a solution 30% isopropanol which decreased fat 100%. Using different immersion media, tools, and raw materials in defatting, produces different fat reduction values. According to Hierro et al. (2021), low-fat insect extracts produce high antioxidant activity, so the low-fat content in the defatting process is also expected to produce high antioxidants from hydrolyzed tuna skin.

The DH value reflects changes in peptide bonds during enzymatic hydrolysis and is an essential parameter for measuring the enzymatic process of proteins (Shu *et al.*, 2018). Based on the results obtained, it was found that hydrolysis of tuna skin with papain enzymes 4-5% for 4-6 hours, DH 29.72–67.64%. These results indicate a higher DH when compared of reported by Prastyo *et al.* (2020) on tilapia collagen hydrolysis with papain enzyme for 15 minutes, DH 12.21-34.01% by Nurilmala *et al.* (2020) on skin collagen of yellowfin tuna with alcalase enzyme for three hours, DH 52.71%. However, these results are lower when compared to research reported by Karnila *et al.* (2020) on yellow pike conger (*Congresox talbon*) 0.15% papain enzyme for four hours, DH 99.06%. Several factors that affect the DH value are the enzyme concentration, hydrolysis time, and the type of protease used. Significantly different DH values can affect the molecular weight and amino acid composition, affecting the resulting peptides' antioxidant activity (Alahmad *et al.*, 2022). The multiple comparison follow-up test results showed that an increase in hydrolysis time above four hours did not show a significant increase in DH value. This condition is likely related to a decrease in the level of enzyme activation due to the autohydrolysis process at high temperatures for a long time (Fonseca *et al.*, 2016). Therefore, only treatments with enzyme concentrations of 4, 5, and 6% (v/v) and hydrolysis time of four hours were evaluated for their antioxidant activity.

The evaluation results showed that the antioxidant activity was higher in the ABTS method compared to the DPPH. Similar results were also reported by Kusumaningty as et al. (2019) and Nurilmala et al. (2020) on the hydrolysis of collagen from fish skin. The ABTS method tends to detect hydrophobic and hydrophilic compounds, while the DPPH method is suitable for hydrophobic compounds. This finding can be attributed to the raw material properties of tuna skin reported by Suseno (2015), which consists of 15.50% hydrophobic and 14.74% hydrophilic amino acids of the total detected amino acids. The differences of amino acids in the hydrolysate result in varying antioxidant activity. The type of proteases and different raw materials exhibit peptides with different amino acid lengths and compositions that hydrolyze proteins. The amino acid composition influences the hydrophobicity and charge of the peptide, which determines its bioactivity (Venuste et al., 2013). The absorbance value of the reducing power method showed a value of 0.415-0.416 at a sample concentration of 1 mg/mL. Samples with higher absorbance values show more ability to donate electrons to free radical elements. This research results showed an absorbance higher than that reported by Arlina et al. (2019) on eel myofibril hydrolysis at a sample concentration of 3 mg/mL with an absorbance of 0.305 nm and lower than that reported by Chen et al. (2014) on the protein hydrolysate sarcoplasm of tilapia at a sample concentration of 3 mg/mL with an absorbance of 0.629 nm. These results indicate that differences in raw materials and sample concentrations affect the samples' absorbance values and antioxidant activity. The antioxidant activity value of tuna skin hydrolysate was included in the very weak antioxidant category (>0.20 mg/mL). From the results obtained it was concluded that the concentration of hydrolysate used from tuna skin needs to be increased to obtain strong antioxidant activity.

The selected treatment from the antioxidant

test on tuna skin hydrolysate with an enzyme concentration of 5% (v/v) showed an MW value in the range of 10.15-48.50 kDa. The results obtained showed a higher MW value than that reported by Nurilmala et al. (2020) on collagen hydrolysate where the MW was in the range of 2.94-11.93 kDa and is lower than that reported by Mahmoodani et al. (2014) on the hydrolysate of catfish (Pangasius sutchi) skin gelatin, where the MW ranged from 45-66 kDa. The observed differences in MW from several hydrolysates were caused by the distribution of MW in different peptide of raw materials, the type of enzymes used Fonseca et al. (2016) and hydrolysis conditions. A lower MW value indicates a higher antioxidant activity value. Generally, bioactive peptides with high antioxidant activity have a molecular size of <10 kDa (Daliri et al., 2017).

The functional groups of tuna skin hydrolysate showed absorption peaks in the amide A, B, I, II and III absorption areas which showed similar results to those reported by Prastyo (2021) on tilapia skin collagen hydrolysate and Nurilmala et al. (2019) on tuna skin collagen hydrolysate. This is associated with the same type of raw material in the hydrolysate used. The absorption peaks obtained indicated that the tuna skin hydrolysate product had similar functional group characteristics to the collagen hydrolysate product. It is known that tuna skin hydrolysate shows a secondary protein structure in the form of  $\alpha$ -helix with amide I vibration at the absorption peak of 1651 cm<sup>-1</sup> (Table 3). The secondary structure of protein can vary depending on the type of material used, for example tilapia skin collagen hydrolysate reported by Prastyo (2021) shows an absorption peak at 1645 cm<sup>-1</sup> secondary protein structure is random coil while in tilapia skin collagen reported by Bi et al. (2019) at the absorption peak of 1628 cm<sup>-1</sup> the secondary protein structure is  $\beta$ -sheet.

The dominant types of amino acids in tuna skin hydrolysate are glycine, proline and arginine which are not different from the amino acid content reported by Prastyo (2021) in tilapia skin collagen hydrolysate by Blanco et al. (2017) in tuna skin collagen hydrolysate and collagen of mackerel (Scomberomorus guttatus) by Fatemi et al. (2021). Tuna skin hydrolysate contains the amino acids glycine, proline, and arginine, 21.92%, 9.76%, and 8.94%, respectively. These amino acids show similar results to collagen hydrolysate products, in which these amino acids are also reported to have a role in biological activity. The tuna skin hydrolysate obtained had the highest amino acid composition among hydrophobic amino acids. According to Chi et al. (2015), hydrophobic amino acids can prevent the oxidation of macromolecules by donating protons to reactive radicals. The arrangement of aromatic amino acid side chains (tyrosine, phenylalanine, and threonine) in hydrolysate also has radical scavenging and metal-chelating activities (Mutaminah *et al.*, 2018). The presence of these aromatic amino acid affects the evaluation of the antioxidant activity obtained previously.

Some of the peptides in the skin hydrolysate identified are multifunctional and exhibit bioactivity other than as antioxidant. The detected presence of Val, Met, Tyr, Leu, Phe, Lys, His, and Arg have been shown to contribute greatly to the potency of antioxidant peptides (Pezeshk et al., 2019). In addition, the hydrophobic nature of peptides can enhance their interaction with lipid targets or the entry of peptides into target organs through hydrophobic composition, which is desirable to achieve antioxidant effects (Wiriyaphan et al., 2012). Jiang et al. (2021) reported that the peptide component Val-Asn-Pro in baijiu drink showed potential antioxidant activity and was in line with the peptide component identified in tuna skin hydrolysate. This proves that tuna skin with an enzymatic hydrolysis process can be utilized as an antioxidant source in food and nutraceutical products, which was also supported by testing the antioxidant activity carried out in the previous stages. Findings by Miguel et al. (2006) showed that the presence of proline amino acid at the C-terminal end and hydrophobic amino acid residues at the N-terminal end (alanine, valine, leucine) had an impact on the good ACE ability of ovalbumin. The high percentage of hydrophobic amino acids in tuna skin hydrolysate is believed to increase ACE inhibitors. Manoharan et al. (2017) reported that the peptide components Gly-Val-Arg (GVR) from oyster mushroom extract (Pleurotus pulmonarius) produces antihypertensive effects that are competitive by analysis of ACE inhibitors in vitro and in vivo. The peptide components Gly-Val-Arg (GVR) was also detected in tuna skin hydrolysate with an integrated proteomic system. Identification was continued by matching the m/z of the resulting chromatogram with the accurate mass from the ChemSpider database. Confirmed peptides have the potential to be developed as nutraceutical products to reduce hypertension. Aminocaproic acid is a derivative and analogue of the amino acid lysine detected in tuna skin hydrolysate samples. Estcourt et al. (2016) reported the presence of aminocaproic acid as an antifibrinolytic for the prevention of bleeding in patients with hematological disorders.

# 4. Conclusion

*Pretreatment* of tuna skin with NaHCO<sub>3</sub> at a low temperature of 4°C effectively reduces fat content

by 80.53%. Tuna skin hydrolysate which was hydrolyzed for four hours with 5% (v/v) enzyme concentration had the best average percentage degree of hydrolysis and the best average  $IC_{50}$  value. The most effective antioxidant method for testing tuna skin hydrolysate is the ABTS method. FTIR results showed absorption peaks of amides A, B, I, II and III. The dominant amino acids are glycine, proline, and arginine. The detected potential bioactive peptides are ACE inhibitor, antioxidant and antifibrinolytic. This research should be continued with the fractionation and purification processes on protein hydrolysates to obtain higher antioxidant activity and specific bioactive peptides as antioxidants.

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# **Authors' Contributions**

All authors have contributed to the final manuscript. The contributions of each author are as follows, SHR; developed conceptual ideas and manuscript, prepared, ran, and controlled the course of the research, as well as collected and analyzed data. JS and SHS; developed conceptual ideas, monitored, and controlled the course of the research and critically revising articles. All authors discussed the results and contributed to the final manuscript.

# **Conflict of Interest**

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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