ISSN: 2347-467X, Vol. 12, No. (1) 2024, Pg. 125-136



# **Current Research in Nutrition and Food Science**

www.foodandnutritionjournal.org

# Enzymatic Hydrolysis of Protein Hydrolysate from *Pangasius* sp. by-Product using Bromelain

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# Abstract

Fish protein hydrolysate (FPH) is a product resulting from the degradation of fish protein into simple peptides and amino acids through hydrolysis. This study aims to optimise the enzymatic hydrolysis conditions of Pangasius sp. by-products to produce high-quality fish protein hydrolysate. Bromelain enzyme was used as the catalyst for hydrolysis. The degree of hydrolysis (DH), pH and antioxidant activity of FPH were used as response parameters. The optimisation was done using response surface methodology (RSM) by applying two factors (enzyme concentration and incubation time) with a 3-level Central Composite Design (CCD) model. The result showed that the bromelain concentration and incubation time gave significantly different effects (p<0.05) on the response parameters of Pangasius protein hydrolysate. Hydrolysis of *Pangasius* protein with 0.04% bromelain enzyme and incubation time of 2.8 hours resulted in DH, pH and DPPH antioxidant activity of 35.88%, 7.07 and 29.86%, respectively. The response value of Pangasius protein hydrolysate was within the range of the predicted value of hydrolysate. Therefore, the optimum conditions suggested by RSM can be used in the future production of Pangasius FPH. In addition, amino acid profiles of Pangasius protein hydrolysate showed high concentrations of Glycine, L-glutamic acid and L-aspartic Acid.

# Introduction

*Pangasius* sp. is one of the popular fish commodities in Indonesia that has experienced an increase in

production in recent years, which was 11.53% in 2015.<sup>1</sup> The global production of *Pangasius* from 2015 to 2021 shows that catfish production is growing

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# Article History Received: 23 January 2024

Accepted: 06 April 2024

# Keywords

Bromelain; Fish Protein Hydrolysate; Optimisation; Pangasius By-Product. consistently and is expected to experience an increase in trade value by 2023.<sup>2</sup> *Pangasius* is mainly required in whole or fillet form.<sup>3</sup> Meanwhile, the amount of waste from *Pangasius* processing ranges from 20-60% of the raw material.<sup>4</sup> The rest of the processing in the form of trimming from *Pangasius* meat is still not well utilised, so it is necessary to use *Pangasius* by-products.<sup>5</sup> Fish protein can be converted into protein hydrolysates and bioactive peptides to increase its economic value, potentially as a high protein source and reduce possible negative environmental impacts.

Fish protein hydrolysate (FPH) results from breaking fish protein into simple peptides and amino acids through hydrolysis processes by enzymes, acids, or bases.<sup>6</sup> Fish protein hydrolysate contain bioactive peptides or short fragments of proteins with residues of 2-20 amino acids that have specific functions, including anti-inflammatory, anti-hypertensive, and anti-microbial.7 Fish protein hydrolysate can be used as an antioxidant source that captures free radicals and binds metal ions.8 Meanwhile, the benefits of protein hydrolysates in the food industry include fortifying ingredients into non-allergenic food formulations for infants and food supplements for diets, food emulsifiers, and ingredients that enhance the characteristics of various products.9 Several studies on fish protein hydrolysates have reported that they can be produced from various kinds of fish, such as milkfish,<sup>10</sup> tilapia,<sup>11</sup> salmon,<sup>12</sup> and catfish.<sup>5</sup> The process of making fish protein hydrolysate enzymatically is influenced by the incubation time factor and certain conditions, such as temperature, pH, enzyme concentration and others.<sup>13</sup> Incubation in the process of protein hydrolysate production utilises the ability of enzymes to help produce simple peptides in a controlled environment. The optimum conditions for producing protein hydrolysates can be identified by applying Response Surface Methodology (RSM) to attain the highest degree of hydrolysis. RSM is a useful method employing mathematical and statistical modelling techniques to enhance the optimisation of food production.14 Pangasius protein hydrolysate was optimised using the Design Expert v. 11 application in a Central Composite Design (CCD) design to determine the approximate optimal direction of the enzyme concentration and incubation duration treatment factors.

The production of FPH requires enzymes to help accelerate the reaction of the hydrolysis process. One of the enzymes used to obtain protein hydrolysates was proteases such as bromelain.<sup>15</sup> Bromelain works to catalyse the reaction of breaking peptide bonds in protein molecules by hydrolysis. Applying papain and different enzyme concentrations was conducted to develop fish protein hydrolysate from Pangasius.5 The use of bromelain in fish protein hydrolysate products has been investigated in independent studies with different bromelain concentrations.16,17 Thus, this study aims to optimise the concentration of bromelain and incubation time during the enzymatic hydrolysis of protein from trimming by-products to produce high-guality fish protein hydrolysate. Degree of hydrolysis (DH), pH and antioxidant activity were used as the responses in the response surface methodology (RSM).

## Materials and Methods Materials

The trimming of *Pangasius* sp. was obtained from a local processing factory in Sidoarjo, East Java. Samples were separated from fat and skin that was still attached. Then, the sample was ground using a chopper until homogenised. Bromelain enzyme was purchased from Nanning Pangbo Biological Engineering Co., Ltd, China. 2,2-diphenyl-1picrylhydrazyl (DPPH), NaOH 0.2 N, HCl, and other chemicals of analytical grade were procured from Merck (Germany) and HiMedia (India).

# Preparation of Fish Protein Hydrolysate (FPH)

The Pangasius protein hydrolysate was produced following the method outlined by He et al.18 The Pangasius trimmings (150 g) were homogenised with distilled water (1:2 w/v). Subsequently, pH adjustment was performed by adding 0.2 N NaOH to reach pH 7. The mixture was preheated at 55°C for 5 minutes. The hydrolysis of the sample was conducted under controlled temperature in a shaker incubator at 55 °C with various bromelain concentrations and incubation times. Subsequently, the mixture was heated to ~95 °C for 20 minutes to inactivate the enzyme and terminate reaction, followed by cooling to room temperature at ~28 °C. The samples were centrifuged at 4500 rpm for 30 minutes, and the resulting liquid and supernatant were collected.

# **Experimental Design**

Response surface methodology (RSM) was employed to predict the optimum fish protein hydrolysate (FPH) production using bromelain enzyme. Incubation duration (A, 0.6–3.4 hours) and enzyme concentration (B, 0.01–0.07%) were selected as independent variables. In our preliminary study, the control treatment (without the addition of bromelain) showed that hydrolysis did not occur even after 4 hours of incubation, as evidenced by the degree of hydrolysis remaining at 0%. The dependent (response) variables were DH, antioxidant activity, and pH. The range and midpoint values of the two independent variables were obtained based on the preliminary study. A Central Composite Design (CCD) was employed for twenty randomised experiments, with each variable coded at five levels (-1.41, -1, 0, +1, and +1.41) as outlined in Table 1. The randomisation of experiments aimed to mitigate the impact of unexpected variability in observed responses. The estimation of responses for both independent and dependent variables and the generation of response surface graphs was conducted using Design-Expert Version 11 software from Stat-Ease Inc. (Minneapolis, MN, USA). Threedimensional surface plots illustrate the relationship between the response and independent variables for each variable examined in this study.

Table 1: Parameter code settings for the *Pangasius* protein hydrolysate process using Central Composite Design

Independent Variable	Symbol	Code					
		-1,41	-1	0	+1	+1.41	
Incubation time (hours)	А	0.6	1	2	3	3.4	
Bromelain concentration (%)	В	0.01	0.02	0.04	0.06	0.07	

### **Degree of Hydrolysis**

The degree of hydrolysis of the hydrolysed soluble protein extract was determined according to Hoyle and Merritt.<sup>19</sup> Specifically, 2 mL of sample was mixed with 20% (v/v) trichloroacetic acid, followed by centrifugation at 10,000 rpm for 20 minutes using the Eppendorf Centrifuge 5415C (Crown Scientific Pty Ltd, Moorebank, NSW, Australia). The supernatant was collected and analysed for nitrogen content following the Kjeldahl method 20. The calculation of degree of hydrolysis was performed using the following formula:

$$Nitrogen (\%) = \frac{(T - B) \times N \times 14.008}{total \ weight \ (mg)} \times 100$$

$$DH (\%) = \frac{ICA - assolved httrogen}{total nitrogen in sample} \times 100$$

### pH Test

The pH test was conducted with a pH meter.<sup>20</sup> The cathode end of the pH meter indicator was washed with distilled water before use and cleaned with tissue. Furthermore, the pH meter was calibrated by dipping the cathode tip into the buffer solution. Then the cathode tip was dipped into 10 mL of *Pangasius* 

protein hydrolysate sample. The measurement results were read on the pH meter monitor.

## **Antioxidant Activity**

Antioxidant activity measurement was conducted based on the method outlined by Nurdiani *et al.*<sup>21</sup> Briefly, 100  $\mu$ L of protein hydrolysate sample solution was mixed with 3,900  $\mu$ L of 0.075 mM DPPH solution in 95% methanol, followed by a 30-minutes incubation. The absorbance values of samples and blanks were determined using a UV-Vis spectrophotometer at a wavelength of 517 nm. Antioxidant activity was calculated using the following formula:

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\label{eq:antioxidant activity (%) = } \frac{(Absorbance \ of \ blank - Absorbance \ of \ sample)}{Absorbance \ of \ blank} \ x \ 100
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# **Amino Acid Composition**

The amino acid profile was determined through ultra-performance liquid chromatography (UPLC) following the method outlined by SIG (2013) at Saraswanty Indo Genetech Laboratory, Bogor.<sup>22</sup> In this process, 0.1g of the sample was mixed with 5mL of 6N HCI thoroughly blended using an ionised vortex. Subsequently, the mixture underwent

hydrolysis at 110 °C for 22 hours, then was cooled and transferred to a 50-mL volumetric flask. After the addition of double-distilledwater, the mixture was filtered through a 0.45-µm filter. Further, 500 µL of the filtrate, 40 µm of alpha aminobutyric acid (AABA), and 460 µL of double-distilled water were combined. After adding 10 µL of the solution to 70 µL of AccQ-Fluor borate buffer, the mixture was vortexed and allowed to stand for 1 minute. Fluor A reagent (20 µL) was introduced, and the mixture was vortexed and left to stand for an additional 1 minute. Subsequently, the solution was incubated at 55 °C for 10 minutes, and 1 µL of the resulting solution was injected into the ultra-performance liquid chromatograph equipped with an ACCQ-Tag Ultra C18 column (Waters Co., Milford, MA, USA). Chromatographic separation was achieved at 49 °C, with the mobile phase consisting of two eluents: AccQ.Tagultra eluent A concentrate (5%, v/v) and water (95%, v/v) and AccQ Tagultra eluent B, each flowing at a rate of 0.7 µL/min. The PDA detector (Water, Massachusetts, USA) was set at a wavelength of 260 nm.

# **Data Analysis**

Experimental data from the different treatments were analysed using multiple regression analysis using Design Expert version 11 (Trial Version, Stat-Ease Inc., Minneapolis, MN, USA). Statistical analysis of the models was performed using Minitab 18 Statistical software (Minitab Pty Ltd., Sydney, NSW, Australia) to evaluate the analysis of variance (ANOVA).

# Results and and Discussion FPH Analysis using Response Surface Methodology (RSM)

Response surface methodology (RSM) was used to optimise conditions for enzymatic protein hydrolysis of *Pangasius* based on DH, DPPH antioxidant activity and pH parameters. The effect of incubation duration and enzyme concentration on DH, DPPH antioxidant activity and pH are shown in Table 2. The effect of two independent variables, A and B, on the response values are presented in Figure 1.

Run	Independent Va	)			
	Incubation Time (hours) (A)	Bromelain Concentration (%) (B)	Degree of Hydro- lysis (%) (Y1)	рН (Ү2)	Antioxidant Activity (%Inhibition) (Y3)
1	1	0.02	14.71	7.16	11.67
2	3	0.02	26.68	6.88	23.83
3	1	0.06	16.67	7.15	17.5
4	3	0.06	32.59	6.98	21.25
5	0.6	0.04	14.46	7.06	14.52
6	3.4	0.04	40.56	6.74	29.54
7	2	0.01	19.54	6.97	17.03
8	2	0.07	27.56	6.98	20.61
9	2	0.04	33.59	7.14	31.86
10	2	0.04	33.05	7.14	27.59
11	2	0.04	30.64	7.22	29.75

Table 2: Treatment design

Experimental data from the different treatments were analysed.A 3-dimensional (3D) response was developed to determine the effects between the two independent factors of incubation duration and enzyme concentration at 0.04% and 2 hours, respectively, and the dependent factors of DH, pH

and antioxidant activity, as suggested by design. The results of response surface plots related to the effect of enzyme concentration and incubation time of *Pangasius* FPH on DH, antioxidant activity, and pH can be seen in Figure 1.

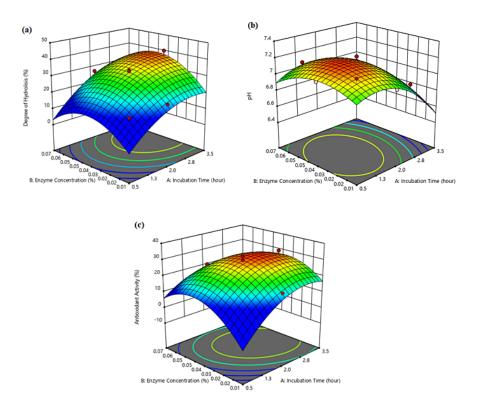


Fig. 1: Response surface plots of the effect of enzyme concentration and incubation time of Pangasius FPH on a. degree of hydrolysis, b. pH and c. DPPH

Source	Sum of Squares	df	Mean Square	F-value	Prob>F	
Model	742.02	5	148.40	22.45	0.0019	significant
А	524.90	1	524.90	79.39	0.0003	
В	46.14	1	46.14	6.98	0.0459	
AB	3.90	1	3.90	0.5900	0.4771	
A²	56.93	1	56.93	8.61	0.0325	
B²	150.08	1	150.08	22.70	0.0050	
Residual	33.06	5	6.61			
Lack of Fit	28.12	3	9.37	3.80	0.2153	not significant
Pure Error	4.93	2	2.47			-
Cor Total	775.08	10				

Table 3: Analysis of va	ariance (ANOVA)	hydrolysis dec	ree of FPH

R<sup>2</sup>=0.9573, A= Incubation Time (hours), B= Enzyme Concentration (%)

# Effect of Bromelain Concentration and Incubation Duration on Hydrolysis Degree (DH) of *Pangasius* FPH

The analysed DH response values ranged from 14.71% to 40.56% at design points 1 and 11 (Table 2). The analysis of variance (ANOVA) of the

Linear Response Surface model for FPH DH is shown in Table 3. Based on the results obtained, the model for DH of *Pangasius* FPH had a significant  $R^2$  value (p<0.05) at 0.9573. ANOVA results showed that enzyme concentration had a significant effect (p<0.05) on DH (0.0019). The same model has been reported in previous studies on enzymatic hydrolysis of different concentrations in different fish species such as herring,<sup>23</sup> salmon,<sup>24</sup> beluga,<sup>25</sup> and carp.<sup>26</sup>

In the quadratic coefficient,  $A_2$  and  $B_2$  significantly (p<0.05) affected the DH of *Pangasius* FPH. The lack of fit test was used to determine the fit of the model. The results showed that the p-value (0.2153) for the lack of fit test was not significant (p>0.05). Therefore, the model fits the experimental data and is selected to predict the conditions for enzymatic hydrolysis of *Pangasius* protein.

A 2D contour plot showing the interaction between incubation duration and bromelain concentration on the degree of hydrolysis response can be seen in Figure 1. Figure 1a shows the increase in the degree of hydrolysis at an incubation duration of about 2.8-3.4 hours and a bromelain concentration of about 0.04% - 0.05%, with the highest result of 40.56% (as shown in Table 2). Bromelain concentration of 1.5% and incubation time of 0.5h - 3h can increase the hydrolysis degree of *Pangasius* FPH, ranging from 10.38% to 29.36%.<sup>16</sup>

The length of incubation influences the degree of FPH hydrolysis. Witono *et al.*<sup>6</sup> stated that the longer the incubation time; the more thorough proteolysis activity is, the more it can increase protein degradation and produce a higher degree of hydrolysis. The degree of hydrolysis is also influenced by enzyme concentration. The longer hydrolysis duration with the addition of enzymes causes the breaking of peptide bonds with maximum speed so that it can produce a higher degree of hydrolysis.<sup>27</sup> However, the activity of cutting peptide bonds with bromelain can decrease due to the decreasing substrate available, which can cause inhibition of the active side of the enzyme. Thus, the higher enzyme concentration results in protein damage, and the degree of hydrolysis decreases. This is in accordance with the statement of Wijayanti et al.<sup>10</sup> that the addition of enzymes in the hydrolysis process has certain limitations where the addition of excess enzymes will result in a constant degree of hydrolysis because the addition of enzymes does not function on the available substrate.

The fat content can also influence the effect of bromelain performance on the degree of hydrolysis in *Pangasius*.<sup>28</sup> Fats are inhibitors in the protein breakdown process, thereby inhibiting the conversion of protein into peptides and simple amino acids. High fat content can limit substrate availability, reducing hydrolysis under particular conditions. Bromelain is one of the proteases capable of breaking lipoprotein bonds, thus allowing optimal bond breaking with the help of bromelain.<sup>29</sup>

Source	Sum of Squares	df	Mean Square	F-value	Prob>F	
Model	0.1799	5	0.0360	6.65	0.0289	significant
A	0.1018	1	0.1018	18.83	0.0074	
В	0.0014	1	0.0014	0.2507	0.6378	
AB	0.0030	1	0.0030	0.5594	0.4882	
A²	0.0648	1	0.0648	11.98	0.0180	
B²	0.0273	1	0.0273	5.06	0.0744	
Residual	0.0270	5	0.0054			
Lack of Fit	0.0228	3	0.0076	3.56	0.2271	not significant
Pure Error	0.0043	2	0.0021			-
Cor Total	0.2070	10				

R<sup>2</sup>=0.8694, A= Incubation Time (hours), B= Enzyme Concentration (%)

# Effect of Bromelain Concentration and Incubation Duration on the pH of *Pangasius* FPH

The pH response values analysed ranged from 6.74 to 7.22 at design points 1 and 11 (Table 2).

The analysis of variance (ANOVA) of the Response Surface Linear model for the pH of *Pangasius* FPH is presented in Table 4. Based on the results obtained, the model for DH of *Pangasius* FPH had a significant  $R^2$  value (p<0.05) at 0.8694. In the quadratic coefficient,  $A_2$  and  $B_2$  significantly (p<0.05) affected the antioxidant activity of *Pangasius* FPH. The lack of fit test was used to determine the fit of the model. The results showed that the p-value (0.2271) for the lack of fit test was not significant (p>0.05).

Figure 1b shows the 3D response surface plot of the effect of bromelain enzyme concentration and incubation time on the pH of *Pangasius* FPH. The lowest pH value is 6.74, and the highest pH value is pH 7.22. The lowest results in this study are not different from the research of,<sup>29</sup> with pH ranging from 6.58-6.81 resulting from hydrolysis with bromelain concentrations of 8% - 16% and incubation times of 2-6 hours. Meanwhile, the pH value of fish protein hydrolysate ranged from 7 to 9.<sup>31</sup>

The length of incubation for 6 hours showed the lowest pH.<sup>31</sup> This was due to the longer incubation duration, causing the enzyme to have a more extended hydrolysis process and releasing more carboxylic groups. A protein solution that undergoes hydrolysis results in a decrease in pH as proteases cleave peptide bonds, leading to the release of carboxylate groups and the liberation of hydrogen ions. Additionally, the addition of alkaline compounds (NaOH) in the protein hydrolysis process plays a role in achieving the optimum pH value and maintaining

its constancy during hydrolysis, facilitating the continued breaking of peptide bonds by enzymes.<sup>10</sup>

In this study, the neutral pH showed the best results for producing *Pangasius* protein hydrolysate. However, there is no standardised pH for fish protein hydrolysate. The pH optimise enzyme performance, ensuring maximum protein digestibility by measuring the degree of hydrolysis of fish protein hydrolysate. Easily digestible protein indicates that the protein can release hydrogen ions quickly, which is indicated by a faster decrease in pH within a certain period.<sup>31</sup>

# Effect of Bromelain Concentration and Incubation Duration on Antioxidant Activity of *Pangasius* FPH

The antioxidant activity response values analysed ranged from 11.67% to 31.86% at design points 1 and 11 (Table 2). The analysis of variance (ANOVA) of the Response Surface Linear model for FPH antioxidant activity is presented in Table 5. Based on the results, the model for DH FPH of *Pangasius* had a significant R<sup>2</sup> value (p<0.05) of 0.9563. In the quadratic coefficients, A<sub>2</sub> and B<sub>2</sub> significantly (p<0.05) affected the antioxidant activity of *Pangasius* FPH. The lack of fit test was used to determine the fit of the model. The results obtained showed that the p-value (0.5981) for the lack of fit test was not significant (p>0.05).

Source	Sum of Squares	df	Mean Square	F-value	Prob>F	
Model	437.91	5	87.58	21.88	0.0021	significant
А	172.53	1	172.53	43.09	0.0012	
В	8.64	1	8.64	2.16	0.2018	
AB	17.68	1	17.68	4.42	0.0896	
A²	105.26	1	105.26	26.29	0.0037	
B²	198.06	1	198.06	49.47	0.0009	
Residual	20.02	5	4.00			
Lack of Fit	10.90	3	3.63	0.7972	0.5981	not significant
Pure Error	9.12	2	4.56			5
Cor Total	457.93	10				

Table 5: Analysis of variance (ANOVA) for antioxidant activity of Pangasius FPH

R<sup>2</sup>=0.9563, A= Incubation Time (hours), B= Enzyme Concentration (%)

The increase in antioxidant activity was attributed to the length of incubation and the optimal enzyme concentration. A longer incubation time, correlating with the increased antioxidant activity of fish protein hydrolysate leads to a varied amino acid composition primarily composed of hydrophobic amino acids.<sup>21</sup> Hydrophobic amino acids have been shown to have strong radical scavenging activity due to the presence of imidazole rings as proton donors.<sup>8</sup> The antioxidant activity is highly dependent on the hydrophobic amino acid content rather than the peptide size.

During hydrolysis, enzymes influence in helping break peptide bonds produces amino acids with bioactive properties as antioxidants.<sup>32</sup> If the enzyme concentration is too high, it will affect the protein's free radicals inhibition activity.<sup>33</sup> The peptides produced from fish protein hydrolysates (which act as antioxidants) do not donate enough hydrogen, so the inhibition tends to decrease. In this study, too high enzyme concentration resulted in a lower inhibition value in the treatment with 0.04% bromelain concentration.

	Factor		Response			Desirability
-	Incubation Time (hours)	Bromelain Concentration (%)	Degree of Hydrolysis (%)	рН	Inhibition (%)	
Prediction	2.8	0.04	37.24	7.00	30.48	0.933
Confirmation/ Verification	2.8	0.04	35.88	7.07	29.86	
P-value			0.082	0.109	0.753	

# Table 6: Optimum point solution and verification

P-value > 0.05 = not significantly different

# **Optimum Value Prediction**

Prediction of the optimum value is conducted by setting criteria according to the Central Composite Design method analysis results. At the same time, the confirmation results are obtained by retesting the factor and response variables. The confirmation results predict the maximum value of the verified response parameters by checking reliability and repeatability. The optimum point solution and model verification results can be seen in Table 6.

Νο	Amino acid	Unit	Result	
1	L-Alanine	mg / kg	35765.4	
2	L-Arginine	mg / kg	31294.6	
3	L-Aspartic Acid	mg / kg	37680.2	
4	Glysine	mg / kg	58876.2	
5	L-Glutamic Acid	mg / kg	67879.3	
6	L-Histidine	mg / kg	7389.29	
7	L-Isoleucine	mg / kg	16492.2	
8	L-Cysteine	mg / kg	19098.4	
9	L-Leucie	mg / kg	37882.3	
10	L-Lysine	mg / kg	42737.2	
11	L-Methionine	mg / kg	4476.25	
12	L-Tryptophan	mg / kg	831.5	
13	L-Valin	mg / kg	19690.2	
14	L-Phenylalanine	mg / kg	22977.8	
15	L-Proline	mg / kg	23110.2	
16	L-Serine	mg / kg	19633.1	
17	L-Threonine	mg / kg	18715.1	
18	L-Tyrosine	mg / kg	6389.59	

# Table 7: Amino Acid Profiles of Pangasius FPH

The verification results in the design expert program with the Central Composite Design method showed that the verification value of the factor was not significantly different (p>0.05) from the predicted response value. The optimum Pangasius protein hydrolysate solution, according to the criteria, is an incubation time of 2.8 hours, bromelain concentration of 0.04% with a predicted response value of the degree of hydrolysis of 37.24%, pH response value of 7, antioxidant activity response value of 30.48%. The confirmation test results are still within the range of predicted values. The FPH optimisation process solution that is considered closest to the optimum is the solution that has a desirability value close to 1 on a scale of 0-1, with 0 indicating low desirability and 1 describing the highest desirability. The desirability value for the projected solution is 0.933. This value serves as a functional indicator for optimisation, indicating how well the program aligns with the set criteria for the final product.34 The optimal formula is identified as the one with the highest desirability value.

# Amino Acid Profiles of *Pangasius* Fish Protein Hydrolysates

The amino acid composition of fish protein hydrolysate obtained under optimal conditions was analysed using the UPLC method. The profile of 18 amino acids present in the fish protein hydrolysate is shown in Table 7.

In this study, variability in amino acid composition was observed in fish protein hydrolysates. Glycine, L-Glutamic Acid and L-Aspartic Acid showed the highest concentrations, about 58876.2 mg/ kg, 67879.3 mg/kg, 37680.2 mg/kg, respectively. Meanwhile, L-tryptophan, L-Methionine, and L-histidine showed lower concentrations, about 831.5 mg/kg, 4476.25 mg/kg, and 7389.29 mg/ kg, respectively. In most reported fish protein hydrolysates, aspartic acid and glutamic acid were identified as having higher concentrations than other amino acids.<sup>35</sup> The high glycine is due to collagen content derived from fish skin that is still attached to the epidermal fish skin. Humpback Grouper skin and scales have a more elevated glycine amino acid content.36

Fish protein hydrolysates have been identified as containing essential amino acids necessary for

maintaining good health. Fish protein hydrolysate is the breakdown product of the enzymatic conversion of fish protein into smaller peptides, which typically contain 2-20 amino acids.37 When employed as a nutraceutical supplement, the amino acid composition of protein hydrolysate is essential as a crucial determinant of its nutritional profile, functional characteristics, and antioxidative efficacy. Variations in the amino acid composition of fish protein hydrolysates mainly depend on various factors, including the origin of enzymes, the raw materials used to prepare the hydrolysates, and the specific hydrolysis conditions applied.<sup>38</sup> Processing conditions can influence the amino acid composition of hydrolysates.<sup>39</sup> This substantial amino acid diversity in Pangasius protein hydrolysate emphasises comprehensive analysis and consideration of nutritional implications.40

### Conclusion

The optimisation of *Pangasius* protein hydrolysate with the factors of incubation time and bromelain concentration presented a significantly different effect on the characteristics of *Pangasius* protein hydrolysate. Hydrolysis of *Pangasius* protein with 0.04% bromelain enzyme and incubation time of 2.8 hours resulted in DH, pH and antioxidant activity of 35.88, 7.07 and 29.86%, respectively, under optimised conditions. The response values were within the range of predicted hydrolysate values, so the optimum conditions suggested by RSM can be used in the production of *Pangasius* FPH.

### Acknowledgements

We would like to thank Jauharotul Afifah and Nada Itorul Umam for their contribution in reviewing and editing the manuscript.

# **Funding Sources**

The authors would like to thank the Ministry of Education, Culture, Research and Technology through the 2021 Basic Research Grant (Number: 438.1/UN10/C10/TU/2021) for funding the research and Hibah Dosen Asing 2023 (Number: 107.9/UN10/KS/2023) Universitas Brawijaya for financing the publication.

### **Conflict of Interest**

The author(s) declares no conflict of interest.

### **Data Availability Statement**

The manuscript incorporates all datasets produced or examined throughout this research study.

### **Ethics Statement**

The document accurately and thoroughly presents the authors' original research and analysis.

# Authors' Contribution

Rahmi Nurdiani designed the experiments, performed the experiments, authored or reviewed drafts of the paper, and approved the final draft. Muhamad Firdaus and Asep Awaludin Prihanto analysed the data, authored or reviewed drafts of the paper, and approved the final draft. Muhammad Rayhansyah Jati and Taufiq Rizki Abdurrahman designed the experiments, analysed the data, performed the experiments. Syaravina Ifilah and Elfriede Rositta Debataraja performed the experiments, analysed the data, prepared figures and/or tables. Abdul Aziz Jaziri and Nurul Huda authored, reviewed drafts of the paper and approved the final draft.

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