

Protein Hydrolysate from Splendid Squid (*Loligo formosana*) Fins: Antioxidant, Functional Properties, and Flavoring Profile

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Abstract

Protein hydrolysates (PH) from splendid squid fins (*Loligo formosana*) (SFPH) with different degrees of hydrolysis (DH) (10-40%) were prepared using alcalase and characterized. The yield of SFPH was increased with increasing DH ($p < 0.05$). All SFPHs showed high protein content but low fat and ash contents. The lightness of SFPH was reduced with augmenting DH as advocated by increasing redness and yellowness ($p < 0.05$). All SFPH samples having DH above 10% exhibited relative solubility higher than 90% ($p < 0.05$). Foaming properties of SFPH were also improved with increasing DH. In general, antioxidant activities upsurged with increasing DH, except metal chelating activity and DPPH radical scavenging activity, which were decreased at higher DHs. Molecular weight (MW) of peptides was lowered with increasing DH as measured by size exclusion chromatography and SFPH having DH 40% (SFPH-40) possessed a higher proportion of peptides with smaller MW. SFPH-40 was rich in aspartic acid/asparagine and glutamic acid/glutamine. SFPH-40 had bitterness, especially at high concentrations. SFPH-40 contained a high number of volatile compounds associated with squid flavor. Thus, hydrolysates from squid fin could serve as an alternative nutrient with flavoring characteristics and antioxidant properties for food applications.

Introduction

Splendid squid (*Loligo formosana*) is an economically important and widely consumed seafood in Southeast Asian countries including Thailand (Singh, Mittal, & Benjakul, 2020). During processing, low-valued by-products such as fins, heads, and guts are produced, which can create serious environmental pollution, if not disposed or managed properly. The by-products from the squid processing account for nearly 75% of the total catch weight (Shahidi, 2006). In general, those by-products are known to contain high-quality functional and nutritive components such as protein, polyunsaturated fatty acids (PUFAs), etc. (Singh, Benjakul, & Kishimura, 2017). Squid processing

by-products such as skin, fin, ovary, etc. are the excellent sources of proteins (Singh et al., 2017; Zlatanov et al., 2006), which can be used for protein hydrolysate production. Squid skin has been mostly exploited for collagen or its derivatives such as gelatin or its hydrolysates (Singh, Mittal, & Benjakul, 2020). Nevertheless, less information has been available for the use of fin to produce protein hydrolysates (PH).

Enzymatic hydrolysis of food proteins is an effective method to recover potent bioactive peptides as compared to the chemical and physical methods (Thiansilakul, Benjakul, & Shahidi, 2007). Several commercial proteases including papain, alcalase, etc. have been employed to produce PH. Among them, alcalase from *Bacillus licheniformis* is one of the most

efficient and widely used proteases for hydrolysis of seafood proteins. It yielded hydrolysates with high bioactivities such as antioxidant activity, etc. as compared to other proteases (Liu et al., 2010). In addition to the bioactivities, hydrolysis is also known to enhance the nutritional, physicochemical, and functional properties of the parent proteins (Singh & Benjakul, 2019). Normally, interfacial properties (emulsifying and foaming properties) of hydrolysates are associated with molecular weight (MW), in which low-MW hydrolysate possessed the decreased viscosity with high solubility and could migrate to interface easily (Zhang et al., 2017).

Nowadays, ready-to-eat foods including healthy snacks are popular around the globe due to their availability and long storage time. However, during the processing of those snacks, the flavor and taste of the raw material can be depleted. Therefore, several kinds of flavorants have been introduced into those products to impart the desired flavor. Among them, food products with squid flavor produced by Bento, Lays, Pringles, etc. are widely popular in Southeast Asian countries. Squid flavor has been produced from the dried squid head, ink as well as mantle (Sukkhown et al., 2018). Squid fin can serve as potential raw material to produce flavorant, especially in the form of PH. Currently, PH has been of great interest associated with their health-promotion due to the presence of bioactive peptides (Cao et al., 2009). Those peptides are well known for excellent antioxidant activities including free radical scavenging, metal chelating, etc., in which oxidative damage toward proteins and lipids could be limited (Singh, Idowu, et al., 2020; Sinthusamran et al., 2018). Moreover, oxidative stress, associated with free radicals generated in human-body caused several diseases, such as Alzheimer, hypertension, diabetes cancer, aging, etc. (Rahman, 2007). Therefore, the objective of the current study was to produce and characterize squid fin protein hydrolysate (SFPH) using alcalase. Moreover, amino acid (AAs) profile and volatile compounds in SFPH responsible for squid flavor were also determined.

Materials and Methods

Raw Material, Enzyme and Chemicals

Splendid squid (*Loligo formosana*) fins were obtained from Sea Wealth Frozen Food Co., Ltd., Songkhla, Thailand and transported to International Center of Excellence in Seafood Science and Innovation, Prince of Songkla University, Thailand in ice within 1 h. The squid fin was then cut into small pieces ($1 \times 1 \text{ cm}^2$), placed in polyethylene bags, and stored at $-20 \text{ }^\circ\text{C}$ until use. The sample was stored for not more than 2 months.

Alcalase from *Bacillus licheniformis* (15 unit/mL) was obtained from Novozyme (Bagsvaerd, Denmark). All chemicals were of analytical grade and were purchased from Sigma Chemical Co. (St. Louis, MO, USA).

Sephadex™ G-25 and blue dextran were obtained from GE Healthcare (Uppsala, Sweden).

Preparation of Squid Fin Protein Hydrolysate (SFPH)

Pre-treatment of Squid Fin

Frozen splendid squid fin was thawed with running water and then bleached with 2% H_2O_2 (w/v) using a sample/solution ratio of 1:10 (w/v) for 6 h with mild stirring using an overhead stirrer in a $4 \text{ }^\circ\text{C}$ walk-in room. H_2O_2 solution was changed twice after 3 h. The bleached sample was rinsed three times with chilled water, followed by blending to obtain uniform homogenate.

Hydrolysis of Bleached Squid Fin

Blended bleached squid fin (10 g) was homogenized with 100 mL of distilled water for 2 min at 9,000 rpm using an IKA homogenizer (Labortechnik, Selangor, Malaysia). The mixture (pH 8) was pre-incubated at $60 \text{ }^\circ\text{C}$ for 15 min and then added with alcalase at 5, 10, 20, and 30 units/g squid fin. At designated hydrolysis time (5, 10, 15, 20, 30, 40, 60, 90, 120, and 180 min), 1 mL of sample was taken and incubated in a water bath at $90 \text{ }^\circ\text{C}$ for 10 min to inactivate enzyme activity. The sample was used for the degree of hydrolysis (DH) determination using 2,4,6-trinitrobenzene sulfonic acid (TNBS) solution (Benjakul & Morrissey, 1997). In brief, hydrolysate samples with the appropriate dilution (62.5 μL) were added with 1 mL of 0.2 M phosphate buffer (pH 8.2) and 0.5 mL of 0.01% freshly prepared TNBS solution. The solution was mixed thoroughly and incubated at $50 \text{ }^\circ\text{C}$ in a water bath for 30 min in the dark. The reaction was terminated by adding 1 mL of 0.1 M sodium sulphite. The mixtures were cooled at room temperature for 10 min in the dark. The absorbance was read at 420 nm using a UV-1601 spectrophotometer and α -amino group was expressed in terms of L-leucine. The DH was computed as follows:

$$\text{DH} = (L - L_0) / (L_{\text{max}} - L_0) \times 100$$

where L is the amount of α -amino groups of hydrolysate sample. L_0 is the amount of α -amino groups in the original sample. L_{max} is the total α -amino groups in the sample obtained after acid hydrolysis (6 M HCl at $130 \text{ }^\circ\text{C}$ for 24 h).

To prepare hydrolysates with several DHs (10-40%), the amount of alcalase was calculated based on a linear relationship between DH and \log_{10} (enzyme concentration). The hydrolysis was carried out as described previously at selected time of 60 min. The obtained hydrolysates were centrifuged at 8000 $\times\text{g}$ for 15 min at $4 \text{ }^\circ\text{C}$. Then, the supernatants were freeze-dried using a freeze-dryer (CoolSafe 55, ScanLaf A/S, Lyngø, Denmark) for 72 h. All PH were further subjected to analyses.

Analyses

Yield

The yield of SFPH with varying DHs was calculated using the following equations:

$$\text{Yield (\%, wb)} = \frac{\text{weight of hydrolysate (g)}}{\text{weight of initial sample (g)}} \times 100$$

$$\text{Yield (\%, db)} = \frac{\text{weight of dry hydrolysate (g)}}{\text{weight of initial dry sample (g)}} \times 100$$

where wb is wet weight basis and db is dry weight basis.

Proximate Compositions

Squid fin and SFPH having different DHs were analyzed for moisture, protein, fat, and ash contents using the AOAC method (AOAC, 2002).

Color

Color parameters such as lightness/brightness (L^*), redness/greenness (a^*), yellowness/blueness (b^*) and color difference (ΔE^*) were calculated according to the method of Temdee et al. (2021).

Molecular Weight (MW) Distribution

MW distribution of SFPHs was determined using a Sephadex G25 gel filtration column (2.5 x 50 cm) following the method given by Sinthusamran et al. (2018).

Solubility

The percentage of solubility was determined by measuring the protein content with respect to the total proteins in the samples as described by Sinthusamran et al. (2018).

Foaming Capacity (FC) and Foaming Stability (FS)

FC of solutions were estimated by whipping 20 mL of the sample solutions (2%, w/v) following the method of Phadtare et al. (2021). FS was measured by calculating the foam destruction within 60 min after whipping at room temperature.

Antioxidant Activities

ABTS and DPPH radical scavenging activities (RSA), ferric reducing antioxidant power (FRAP) and metal chelating activity of SFPH samples were determined as per the method of Sae-Leaw et al. (2016).

Amino Acid Profile, Bitterness Intensity, and Volatile Compounds in SFPH-40

Amino Acid Profile

The AAs composition of SFPH-40 powder was analyzed using AAs analyzer as per the procedure of Nilsuwan et al. (2021).

Volatile Compounds

The isolation and concentration of volatiles were accomplished using the headspace solid-phase microextraction (HS-SPME) method, followed by GC-MS analysis (Nilsuwan et al., 2021).

Bitterness

The bitterness of SFPH-40 at various levels (0, 1, 2, 3, and 4%; w/v) was examined by 11 trained panellists (6 male and 5 female) following the method of Idowu et al. (2019) with slight modifications.

Statistical Analysis

A completely randomized design (CRD) was used for the whole study and data were subjected to one-way analysis of variance (ANOVA). The differences between means were evaluated by Duncan's multiple range test using the SPSS statistic program (SPSS 11.0 for Windows, SPSS Inc., Chicago, IL, USA).

Results and Discussion

Degree of Hydrolysis (DH)

The change in DH of SFPH prepared using alcalase at different concentrations as function of hydrolysis time is shown in Figure 1A. DH was augmented with increasing enzyme concentrations and hydrolysis time. The highest DH was obtained when alcalase at 30 unit/g sample was used. The DH was increased rapidly within the first 20 min. Thereafter, gradual increase in DH was noticeable with further augmenting time up to 180 min. Similar results were reported during the hydrolysis of proteins obtained from yellow stripe (Klompong et al., 2007) and salmon (Gbogouri et al., 2004). This was associated with diminishing substrate, product inhibition or autodigestion of alcalase during the prolonged period of hydrolysis (Klompong et al., 2007). Alcalase at 2.83, 9.59, 32.40, and 109.47 units/g sample was used to achieve DH of 10, 20, 30, and 40% as determined by the linear relationship between DH and alcalase concentrations (data not shown) (Klompong et al., 2007). Based on preliminary work, hydrolysis using a higher concentration of alcalase resulted in a darker hydrolysate, possibly associated with dark color of alcalase used. Thus, only SFPHs with 10, 20, 30, and 40% DH were prepared.

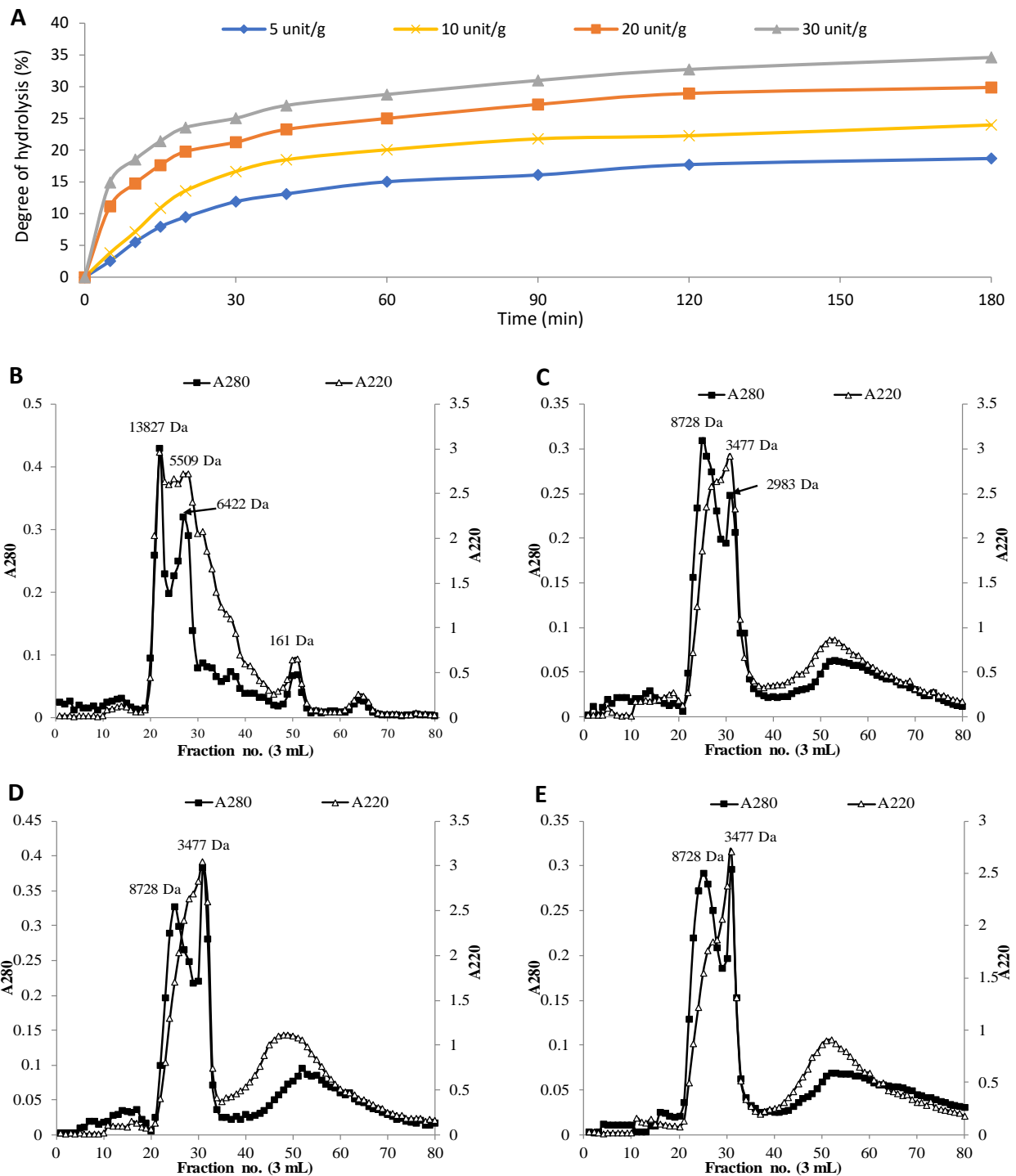


Figure 1. Degree of hydrolysis (DH) of splendid squid fin hydrolysates (SFPH) produced using alcalase at different concentrations as a function of time (A) and molecular weight distribution profile of SFPH with different DHs (B-E; 10-40%, respectively).

Proximate Compositions

Proximate compositions of fresh splendid squid fin and SFPH having various DHs are shown in Table 1. The raw squid fin had high content of proteins (80.13%) followed by fat (8.02%) and ash (6.68%) (dry weight basis). On a wet weight basis, squid fin contained 85.25% water, 11.82% protein, 1.05% fat, and 0.98% ash. Similar chemical composition was found in the squid meat from the mantle, which contained 75-84% water, 13-22% protein, 0.1-2.7% fat, and 0-9-1.9%

minerals (ash) (wet weight basis) (Sikorski & Kołodziejska, 1986). For SFPH, no differences in protein and fat contents were noticed, irrespective of DH ($p < 0.05$). Nevertheless, ash content was reduced with increasing DH ($p < 0.05$). With dry weight basis, protein, fat, and ash contents of all SFPHs were in the range of 93.72-94.33%, 2.65-3.16%, and 1.10-1.73%, respectively. All the SFPHs had high protein content, which might be associated with the solubilization of protein from the fins. SFPHs also showed high protein content than squid fin. Moreover, hydrolysis also

removed insoluble substances, lipid, and inorganic components (metal ions) associated with the proteins. The lipid removal was also supported by the lower fat content in SFPH as compared to raw squid fin. Overall, DH did not affect the chemical composition of hydrolysates prepared by alcalase.

Yield

The yield of SFPH was increased with augmenting DH, in which a higher amount of alcalase was used for hydrolysis (Table 1). The yields of 7.67, 8.27, 9.61, and 11.42% (wet weight basis) were found for SFPH-10, -20, -30, and -40 samples, respectively. These results showed the positive relation between amount of alcalase used and amount of SFPH obtained. Based on dry matter, yields of 48.49, 52.33, 60.76, and 72.21% were found for the aforementioned samples, respectively. Idowu et al. (2019) also reported increasing yield when DH of protein hydrolysate from salmon frames was increased. Alcalase was able to cleave peptide bonds inside the protein chain of squid fin during the hydrolysis process. A higher number of smaller peptides was generated and liberated into SFPH when a high amount of alcalase was used.

Color

Change in color values of SFPH with respect to the DH is shown in Table 2. The L^* value was decreased with increasing DH, indicating a reduction in the lightness of SFPH ($p < 0.05$), which was also supported by the

augmenting a^* and b^* values, which represent redness and yellowness, respectively ($p < 0.05$). In general, hydrolysate powder having higher DH was yellowish brown in color. Similarly, Klompong et al. (2007) detected brownish color at higher DH (60%) of hydrolysate from round scad protein produced using flavourzyme. The reduction in lightness of SFPH powders was more likely due to the non-enzymatic browning (Maillard) reaction. This reaction occurs between amino group of peptides and carbonyl compounds. Moreover, pigments present in the alcalase as well as those remaining in squid fin might have contributed to the color of hydrolysates. The ΔE^* value was also increased with an upsurge in DH, which suggested the color change. Hence, the varying color of SFPH was attributed to the composition and pigments of the raw material, type and amount of enzyme as well as and the hydrolysis condition.

Molecular Weight (MW) Distribution

Elution profiles of SFPH with various DHs are illustrated in Figure 1B-E. In general, A_{220} has been used to monitor peptides, while A_{280} indicates the peptides/proteins, primarily comprising aromatic amino acids (AAs) (Sinthusamran et al., 2018). All the samples had three major peaks at A_{280} , representing peptides comprising aromatic AAs, which had MW in the range of 161-13827 Da. On the other hand, A_{220} had two major peaks (MW ranging from 161 to 5509 Da) for all samples except SFPH-10, which had an extra peak having MW of 13827 Da (Figure 1B). When DH was higher than 10%,

Table 1. Proximate composition and yield of splendid squid fins and squid fin protein hydrolysate (SFPH) produced using alcalase with different degrees of hydrolysis.

	Splendid squid fins	SFPH-10	SFPH-20	SFPH-30	SFPH-40
Moisture	85.25±0.16	1.72±0.16 ^b	2.01±0.78 ^b	5.33±0.18 ^a	5.37±0.35 ^a
Protein	11.82±0.17 (80.13±1.12)	91.61±1.55 ^a (93.72±0.56 ^a)	91.72±3.26 ^a (94.12±3.33 ^a)	89.00±1.98 ^a (94.01±1.14 ^a)	89.26±4.14 ^a (94.33±4.37 ^a)
Fat	1.05±0.27 (8.02±0.23)	2.91±0.87 ^a (2.96±0.89 ^a)	2.80±0.84 ^a (2.65±1.06 ^a)	2.99±0.92 ^a (3.16±0.97 ^a)	2.98±0.33 ^a (2.83±0.35 ^a)
Ash	0.98±0.07 (6.68±0.49)	1.65±0.05 ^a (1.68±0.05 ^a)	1.50±0.31 ^{ab} (1.63±0.32 ^a)	1.30±0.13 ^b (1.37±0.14 ^b)	1.05±0.27 ^b (1.10±0.29 ^b)
Yield	-	7.67±0.12 ^a (48.49±0.77 ^a)	8.27±0.12 ^b (52.33±0.73 ^b)	9.61±0.24 ^c (60.76±1.50 ^c)	11.42±0.13 ^d (72.21±0.82 ^d)

Values are presented as mean ± SD ($n = 3$). Values are on wet weight basis, except in brackets, which is represented as dry weight basis. Different lowercase superscripts within the same row indicate significant difference ($p < 0.05$).

SFPH-10, SFPH-20, SFPH-30, SFPH-40: Squid fin protein hydrolysate (SFPH) powder with degree of hydrolysis of 10, 20, 30, and 40%, respectively.

Table 2. Color, solubility, and foaming properties of squid fin protein hydrolysate (SFPH) produced using alcalase with different degrees of hydrolysis.

Samples	Color				Solubility (%)	Foaming capacity (%)	Foaming stability (%)
	L^*	a^*	b^*	ΔE^*			
SFPH-10	89.79±0.11 ^a	0.88±0.04 ^c	10.62±0.13 ^c	10.43±0.13 ^c	89.80±1.67 ^c	72.46±2.45 ^d	31.78±2.52 ^c
SFPH-20	88.84±0.09 ^b	1.04±0.21 ^c	10.57±0.52 ^c	10.48±0.53 ^c	93.65±0.43 ^b	80.95±2.70 ^c	34.92±3.26 ^c
SFPH-30	87.30±0.21 ^c	1.32±0.10 ^b	13.41±0.35 ^b	13.61±0.34 ^b	96.49±1.43 ^a	85.98±1.36 ^b	50.40±2.87 ^b
SFPH-40	81.29±0.21 ^d	4.20±0.24 ^a	21.15±0.45 ^a	23.37±0.54 ^a	97.05±2.26 ^a	105.29±1.95 ^a	81.68±2.04 ^a

Values are presented as mean ± SD ($n = 3$).

Different lowercase superscripts within the same column indicate significant difference ($p < 0.05$). Caption: see Table 1

the peak (A_{280}) of peptides having MW of 13827 Da disappeared. This coincided with the formation of another peak with MW of 8728 Da. Similarly, A_{220} showed a shift in the peak of MW 5509 to 3477 Da. The result suggested the hydrolysis of high MW proteins/peptide with increasing DH, thus resulting in the formation of smaller proteins as well as peptides. This was advocated by increasing surface area of the peak of MW 161 Da monitored using both A_{220} and A_{280} . The increase in DH (Figure 1A) of SFPH was in line with the upsurging peak area of fractions with low MW peptides. Sinthusamran et al. (2018) also noticed the varying MW profiles as influenced by DH, which could be affected by the amount and type of enzyme as well as hydrolysis time.

Solubility

All samples showed 85-97% of solubility at neutral pH (Table 2). The lowest solubility was noticed in SFPH-10, followed by SFPH-20 sample ($p < 0.05$). The maximum solubility was noticed for SFPH-30 and SFPH-40 samples ($p < 0.05$), however both showed similar solubility ($p > 0.05$). PH from yellow stripe trevally meat produced using alcalase and flavourzyme with different DHs (5-25%) also exhibited solubility higher than 85% (Klompong et al., 2007). Higher solubility of SFPH samples with higher DH is associated with the presence of smaller peptides having polar residues, which were capable of forming H-bonds with water (Klompong et al., 2007). Hydrolysis has been known to alter molecular size as well as hydrophobicity, which directly influence the functional properties of resulting peptides (Singh & Benjakul, 2019). Moreover, hydrophobicity/hydrophilicity ratio or charged AAs also affect the solubility of PH (Gbogouri et al., 2004; Klompong et al., 2007). Hence, solubility of SFPH having different DHs is influenced by peptides size, hydrophobic-hydrophilic ratio, and charged AAs in the peptides.

Foaming Properties

Foaming capacity (FC) and foaming stability (FS) of SFPH powders with different DHs are shown in Table 2. FC was increased (72-105%) with an upsurge of DH ($p < 0.05$), in which the highest FC was noticed for SFPH-40 (105.29%) ($p < 0.05$). In general, foaming ability of proteins is related to their ability to form film at the interface of air and water (Singh et al., 2017). During whipping, protein, that rapidly adsorbs at the interface, undergoes unfolding as well as molecular

reorganization, thus improving foaming properties more effectively than protein that adsorbs slowly and resists unfolding at the interface of air and water (Zayas, 1997). Bao et al. (2017) also reported increasing FC of egg yolk hydrolysates with increasing DH. With increasing DH, peptides, which had smaller size, were able to relocate to the interfacial film instantly and occupy the interface more potentially than other having larger size. Similarly, FS of SFPH was upsurged with increasing DH ($p < 0.05$). Nevertheless, SFPH-10 and SFPH-20 had similar FS ($p > 0.05$). Generally, enzymatic hydrolysis reduced MW (Figure 1 B-E), thereby enhancing the solubility and exposing hydrophobic domains. Since peptides in SFPH-40 rapidly migrated to interface and could arrange themselves, in the way which stronger film could entrap air bubble potentially. As a consequence, the foam was stable as indicated by the highest foaming stability. On contrary, foaming properties of hydrolysate obtained from yellow stripe trevally were reduced with increasing DH (Klompong et al., 2007). This was more likely due to the difference in size and charge of peptides, which could be affected by the hydrolyzing enzymes, time, substrate, etc. (Klompong et al., 2007; Zayas, 1997).

Antioxidant Activities

Antioxidant activities of SFPH samples were measured by ABTS, DPPH, FRAP, and metal chelating assays as shown in Table 3. ABTS-RSA of SFPH was increased with augmenting DH and the highest activity was found for SFPH-30 and SFPH-40 samples ($p < 0.05$), but both samples showed similar activity ($p > 0.05$). Alcalase with a broad specificity for peptide bonds produced short-chain peptides, which could scavenge ABTS radicals. Similarly, upsurged DPPH-RSA was observed as DH was increased up to 30% ($p > 0.05$). When DH reached 40%, lower activity was noticed ($p < 0.05$). Therefore, peptides in hydrolysates with varying DHs might scavenge ABTS and DPPH radicals with different modes of action. Usually, peptides/proteins present in hydrolysate can act as H-donors, which convert radicals to more stable products and inhibit radical chain reaction (Khantaphant & Benjakul, 2008; Sinthusamran et al., 2018). ABTS assay determines the H-donating ability of compounds in aqueous phase (Singh, Idowu, et al., 2020). With high ABTS-RSA, it was postulated that SFPH mainly contained hydrophilic components. For DPPH-RSA, it is generally used to determine the H-donating ability of a compound in a lipophilic system (Wu, Chen, & Shiau, 2003). The result suggested that the hydrolysates at higher DH might contain more electron

Table 3. Antioxidant activities of squid fin protein hydrolysate (SFPH) produced using alcalase with different degrees of hydrolysis.

Antioxidant assays	SFPH-10	SFPH-20	SFPH-30	SFPH-40
ABTS ($\mu\text{mol TE/g sample}$)	862.15 \pm 93.91 ^c	1,074.64 \pm 98.31 ^b	1,219.10 \pm 83.58 ^a	1,213.59 \pm 37.79 ^a
DPPH($\mu\text{mol TE/g sample}$)	3.22 \pm 0.11 ^a	3.33 \pm 0.20 ^b	3.57 \pm 0.13 ^a	2.79 \pm 0.07 ^b
FRAP ($\mu\text{mol TE/g sample}$)	22.54 \pm 1.74 ^d	26.63 \pm 2.46 ^c	29.19 \pm 0.75 ^b	32.98 \pm 1.35 ^a
Metal chelating activity ($\mu\text{mol EDTA/g sample}$)	1.05 \pm 0.01 ^a	1.08 \pm 0.03 ^a	0.85 \pm 0.08 ^b	0.61 \pm 0.03 ^c

Values are presented as mean \pm SD ($n = 3$).

Different lowercase superscripts within the same row indicate significant difference ($p < 0.05$). Caption: see table 1.

donors in the form of hydrophilic AAs or peptides, which resulted in a lower capacity to scavenge lipophilic DPPH radicals. Similarly, Sinthusamran et al. (2018) observed lower DPPH-RSA of PH from Pacific white shrimp cephalothorax at higher DH.

FRAP activity was increased by augmenting DH ($p < 0.05$) (Table 3). Generally, reducing capacity is determined by its potential to donate an electron to a free radical (Wang et al., 2008). FRAP is commonly used to measure the capacity of a substance to reduce TPTZ-Fe(III) complex to TPTZ-Fe(II) complex (Sae-Leaw et al., 2016). The higher FRAP activity, especially at high DH was more likely due to the electron donation to the free radical, which prevents or retards propagation (Sinthusamran et al., 2018). The result of the study agreed with hydrolysates produced from gelatin and protein isolate extracted from splendid squid prepared using proteases from shrimp hepatopancreas (Hamzeh, Benjakul, & Senphan, 2016). Conversely, the lower FRAP activity of PH from yellow stripe trevally produced using alcalase was obtained at higher DH (Klompong et al., 2007). This might be governed by different peptides produced.

Based on metal chelating activity, SFPH-10 and SFPH-20 exhibited the highest activity ($p < 0.05$). However, both samples had similar values ($p > 0.05$) (Table 3). On the other hand, with further increase in DH to 40%, lower activity was noticed ($p < 0.05$). In general, complex formed between ferrozine and Fe^{2+} ion is disrupted in the presence of the chelating agent,

resulting in the decrease in color formation (Thiansilakul et al., 2007). The lower metal chelating activity at higher DH was mainly due to the alternation in structure and sequence of peptide chains. Hamzeh et al. (2016) also reported decreasing metal chelating activity of splendid squid gelatin hydrolysate with an increase in DH to 30%. However, no difference in metal chelating activity was noticed for hydrolysates from splendid squid mantle protein isolate having DH of 10-30% (Hamzeh et al., 2016).

Therefore, DH, enzymes used, raw material, etc. influenced the antioxidant activities of PH (Klompong et al., 2008). Based on yield, solubility, foaming properties, and antioxidant activities, SFPH-40 sample was selected for further characterization.

Amino Acid Profile, Bitterness Intensity, and Volatile Compounds of SFPH-40

Amino Acid Profile

AAs profile of SFPH-40 sample is shown in Table 4. Glutamic acid/glutamine (10.60 g/100 g sample) was the most abundant AAs in SFPH-40 followed by lysine (7.58 g/100 g sample) and aspartic acid/asparagine (7.26 g/100 g). Leucine (6.07 g/100 g sample) was also found at a high concentration along with glycine (5.94 g/100 g sample). The SFPH-40 had a favorable EAA/NEAA ratio (0.70), which was higher than that of muscle proteins (0.60) as reported by Iwasaki and

Table 4. Amino acid composition of squid fin protein hydrolysate (SFPH) produced using alcalase having 40% degree of hydrolysis.

Amino acids (g/100 g sample)	SFPH-40
L-Alanine	4.01±0.11
L-Arginine	5.94±0.21
Aspartic acid/asparagine	7.26±0.32
Cystine	ND
Glutamic acid/glutamine	10.60±0.47
Glycine	5.94±0.36
Histidine	3.60±0.10
Isoleucine	2.58±0.08
Leucine	6.07±0.36
Lysine	7.58±0.35
Hydroxylysine	0.28±0.00
Methionine	1.26±0.01
Phenylalanine	2.50±0.02
Hydroxyproline	1.24±0.01
Proline	3.73±0.05
Serine	3.35±0.06
Threonine	3.68±0.04
Tryptophan	0.20±0.00
Tyrosine	1.55±0.00
Valine	2.78±0.01
Essential amino acids (EAA)	30.53±2.03
Non-essential amino acids (NEAA)	43.62±3.63
EAA/NEAA ratio	0.70
Polar to non-polar ratio (P-value)	1.13±0.00
Hydrophobic amino acids	29.07±2.96
Average hydrophobicity	1064.14±8.06
Frequency of nonpolar side chains (NPS)	0.35±0.00

Values are presented as mean ± SD (n = 3). SFPH-40: squid fin protein hydrolysates having 40% degree of hydrolysis.

Harada (1985). The total hydrophobic AAs in SFPH-40 were 29.07 g/100 g. Normally, hydrophobic AAs are responsible for the bitterness of PH (Sinthusamran et al., 2020).

Polar to non-polar side chain ratio (P-value), frequency of non-polar side chains (NPS), and average hydrophobicity (H Φ) of SFPH-40 were measured from the AAs composition following the calculations given by Bigelow (1967). In general, the fibrous proteins such as collagen, elastin, tropomyosin, cilia, etc. from various sources have NPS and P-value, and H Φ in the range of 0.018-41, 0.09-2.14, and 440-1120, respectively (Bigelow, 1967; Tristram & Smith, 1963). Patil and Benjakul (2017) reported that the albumin from coconut milk proteins with high P-value (2.28) had lower NPS (0.21) and H Φ (780.23) values. On the other hand, globulin with low P-value (1.69) showed higher NPS (0.29) and H Φ (879.42). In current study, P-value, NPS, and H Φ of SFPH-40 were 1.13, 0.35, and 1064.14, respectively, which were higher than the average values of various fibrous proteins as well as albumin and globulin fractions of the coconut milk proteins (except P-value). Using 150 proteins, Bigelow (1967) observed that P-value did not always show relationship with NPS and H Φ values. Therefore, balance between hydrophobic and hydrophilic parts of SFPH-40 determined solubility and foaming properties, which were related with P-value, NPS, and H Φ values.

Bitterness

The bitterness score was directly proportional to the concentration of SFPH-40 sample, in which SFPH-40 sample at 1% (w/v) showed the lowest bitterness ($p < 0.05$). The bitterness scores of SFPH-40 sample at concentrations of 1, 2, 3, and 4% (w/v) were 4.36, 8.55, 13.64, 14.91, respectively ($p < 0.05$). In general, hydrolysis exposes hidden hydrophobic AAs located inside globular protein, thus causing bitterness (Singh, Idowu, et al., 2020). In general, bitterness of PH is affected by the structure of peptides (Kim & Li-Chan, 2006). The bulky hydrophobic AAs at the both C- and N-terminal in di- and tri-peptides caused bitterness, whereas bulky-hydrophobic groups either in the presence or absence basic-AAAs at the C-terminal and bulky basic-AAAs at the N-terminal in peptides larger than tetrapeptides are known for bitterness (Kim & Li-Chan, 2006). The bulky-hydrophobic groups of hydrophobic AAs and tyrosine at C-terminal in peptides were able to bind with taste bud and cause bitterness (Idowu et al., 2019). The hydrolysate with higher hydrophobic AAs had enhanced sensation of the bitter taste. Moreover, proline localized internally in the peptide chain contributed to bitterness (Idowu et al., 2019). The result was supported by high P-value and H Φ (Table 4). In addition, bitterness scores are affected by DH, number of carbons inside the chain, especially for branched-chain AAs, and their concentration (Yarnpakdee et al., 2015). Therefore, an appropriate concentration of SFPH-

40 must be considered for the supplementation in various food products to avoid bitterness perceived by consumers.

Volatile Compounds

Volatile compounds in SFPH-40 sample are listed in Table 5. Various classes of chemicals were identified as follows: 7 aldehydes, 2 alcohols, 2 amides, 2 alkanes, 1 alkene, 3 aromatic, 7 carboxylic acids, and 7 ketones. Aldehydes are considered as most predominant volatile secondary oxidation products (SOPs) in SFPH-40 sample. The formation of aldehydes indicates lipid oxidation taken place in foods. Due to their low odor threshold value, they might result in off-odor and off-flavor (Nilswan et al., 2021). Various kinds of aldehydes such as hexanal, octanal, nonanal, pentanal, etc. were generated during oxidation (Ross & Smith, 2006). Among all aldehyde compounds in SFPH-40 sample, butanal, 3-methyl- was highest in abundance followed by propanal, butanal, hexanal, benzaldehyde, and 2-propenal, respectively (Table 5), which might be oxidation products of unsaturated fatty acid. The 3-methyl-butanal showed green, fruity, nutty, cheese or sweat odors, which is found in Malaysian fish sauce (Mohamed et al., 2012). Cui et al. (2020) found 14 major aldehydes of low threshold value in cooked squid, which contributed to the flavor in cooked squid. During the enzymatic process, some lipids were oxidized and Strecker degradation of AAs occurred (Tan et al., 2018). The branched alcohol compounds namely, 2-penten-1-ol and 1,5-octadien-3-ol were found in SFPH-40 sample and might produce aroma in SFPH-40 owing to their low threshold value as compared to short and long straight chain alcohols. 8-carbon alcohols are known to be present in all species of fish (Nagarajan et al., 2015). The unsaturated alcohol might be generated from n-3 and n-6 polyunsaturated fatty acids. Alcohols in SFPH-40 could serve as flavoring agent in various products such as soups and sauces. Two furan compounds, namely 3-methyl-furan and 2-ethyl-furan were decomposed from 12-, 14-, and 16-hydroperoxide of linolenate, eicosapentaenoate and docosahexaenoate, respectively. Those compounds further produced conjugated diene radical through β -cleavage and reacted with oxygen to produce a vinyl hydroperoxide (Yarnpakdee et al., 2012). Moreover, breakdown of vinyl hydroperoxide through loss of hydroxyl radicals formed alkoxy radicals, which were crystallized to furan (Maqsood & Benjakul, 2011). Nevertheless, their contributions to SFPH-40 odor/flavor may be insignificant due to their low concentrations and high threshold values. Carboxylic acids were identified in SFPH-40 and might be formed from chemical and enzymatic oxidation of aldehydes. Moreover, N-terminal of peptide might consist of several alkyl carboxylic acids, which act as a blocking group, thus regulating various biological activities (Moscarello et al., 1992). Ketones significantly contributed to flavor of fish

products due to their low threshold value. Seven types of ketones were identified and quantified in SFPH-40 (Table 5), in which 1-Penten-3-one was most abundant, which is a fish, garlic, and mustard tasting compound. The 2,3-pentanedione was also present in sufficient amount, which is known to render the intense odorant in squid (Shahidi, 1998). Moreover, other compounds such as aliphatic hydrocarbons including, 2-methyl-1,3-butadiene, oxybis-methane, and 3-methyl pentane along with toluene and amides were also identified. Hydrocarbons present in SFPH-40 had less impact on the overall flavor of foods because of their high aroma thresholds (Wang et al., 2010). Since there were several volatiles in SFPH-40, those compounds could synergistically contribute to the flavor. Some of volatile compounds associated with specific flavors might be lost during the preparation of hydrolysate. Nevertheless, the major contributors to unique squid flavor should be further investigated.

Conclusions

Splendid squid fin protein hydrolysate (SFPH) prepared using alcalase with 40% DH (SFPH-40) had the highest relative solubility, foaming capacity, and foaming stability. All SFPH possessed antioxidant

activities, which varied, depending on the DH. The SFPH-40 was rich in essential AAs as well as hydrophobic AAs. The bitterness of the SFPH-40 was increased when its concentration was increased. Aldehydes, alcohols, carboxylic acids, and ketones were the abundant volatile compounds in the SFPH-40 sample. Overall, SFPH-40 at appropriate concentration could be used to improve the nutritional, functional, and squid flavor of various food products, in which further study is required.

Ethical Statement

Ethical approval is not required for this study.

Author Contribution

Conceptualization, A.S. and S.B.; methodology, A.S. and G.A.U.P.; software, A.S.; validation, A.S., S.B. and A.J.; formal analysis, A.S.; investigation, A.S., A.J. and G.A.U.P.; resources, S.B.; data curation, S.B.; writing—original draft preparation, A.S.; writing—review and editing, S.B., A.F.Y., and H.H.; visualization, A.S.; H.H., and A.F.Y.; supervision, S.B.; project administration, S.B.; funding acquisition, S.B. All authors have read and agreed to the published version of the manuscript.

Table 5. Volatile compounds of squid fin protein hydrolysate (SFPH) produced using alcalase having 40% degree of hydrolysis

Classes	Compounds	SFPH-40
Aldehyde	Benzaldehyde	1.02
	Butanal	1.31
	Butanal, 3-methyl-	2.65
	Hexanal	1.06
	(E)-4-Oxohex-2-enal	0.78
	Propanal	1.64
	2-Propenal	0.81
Alcohol	2-Penten-1-ol, (Z)-	1.59
	1,5-Octadien-3-ol, (Z)-	1.19
Heterocyclic	Furan, 3-methyl-	0.70
	Furan, 2-ethyl-	1.18
Carboxylic acid	Acetic acid	4.87
	Benzoic acid	0.24
	Butanoic acid	0.32
	Hexanoic acid	0.61
	Pentanoic acid	0.62
	Propanoic acid	6.11
	Sorbic Acid	0.34
Ketone	3-Buten-2-one	0.37
	2-Butanone	1.50
	2-Nonanone	0.46
	2-Pentanone	1.89
	1-Penten-3-one	4.69
	2,3-Pentanedione	1.17
	1-Phenyl-ethanone	1.49
Other compounds	Toluene	0.25
	1,3-Butadiene, 2-methyl-	0.24
	Acetamide	0.42
	Propanamide	0.59
	Methane, oxybis-	0.38
	Pentane, 3-methyl-	0.30

Values are expressed as abundance ($\times 10^8$). SFPH-40: squid fin protein hydrolysates of degree of hydrolysis of 40%.

Conflict of Interest

All authors declare that there is no conflict of interest.

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