

ACE-Inhibitory Properties of Proteolytic Hydrolysates from Giant Jellyfish *Nemopilema nomurai*

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Abstract

This study aimed to determine the degree of hydrolysis and angiotensin-I-converting enzyme (ACE)-inhibitory activity of Giant Jellyfish *Nemopilema nomurai* (jellyfish) hydrolysates. The degree of hydrolysis using six proteolytic enzymes (Alcalase, Flavourzyme, Neutrase, papain, Protamex, and trypsin) ranged from 13.1-36.8% and the inhibitory activities from 20.46-79.58%. Using papain hydrolysate, we newly isolated and characterized ACE-inhibitory peptides with a molecular weight of 3,000-5,000 Da that originated from jellyfish collagen. The purified peptide (FII-b) was predicted to be produced from an alpha-2 fragment of the type IV collagen of jellyfish. The N-terminal sequence of FII-b was Asp-Pro-Gly-Leu-Glu-Gly-Ala-His-Gly- and showed 87% identity to the collagen type IV alpha-2 fragment of *Rattus norvegicus* and a predicted protein from *Nematostella vectensis*, indicating that the ACE-inhibitory peptide originated from the collagen hydrolysate and had an IC₅₀ value of 3.8 µg/mL. The primary structure of the fragment is now being studied; this peptide represents an interesting new type of ACE inhibitor and will provide knowledge of the potential applications of jellyfish components as therapies for hypertension.

Key words: ACE, Jellyfish, *Nemopilema nomurai*, Proteolytic hydrolysis, Papain

Introduction

Angiotensin-I-converting enzyme (ACE; EC 3.4.15.1) is a circulating enzyme that participates in the body's rennin-angiotensin system and plays an important physiological role in regulating blood pressure. ACE is known as peptidyl dipeptidase A and primarily cleaves a C-terminal dipeptide from substrates. It converts an inactive form of a decapeptide (angiotensin-I) to a potent vasoconstrictor, an octapeptide (angiotensin-II), and also inactivates the catalytic function of bradykinin, which has a depressor action (Ondetii et al., 1977; FitzGerald et al., 2004).

To date, many peptides with high ACE-inhibitory activity have been isolated and characterized from both natural products and processed foods. These include various protein hydrolysates, such as those of cheese (Smacchi and Gobetti, 1998), milk proteins (Gobetti et al., 2000), egg white

(Miguel et al., 2007), white and red wines (Pozo-Bayón et al., 2007), plant proteins (Dziuba et al., 1999; Wu et al., 2008), meat (Jang and Lee, 2005), and marine sources (Fujita and Yoshikawa, 1999; Je et al., 2005; Lo and Li-Chan, 2005; He et al., 2006). Some types of ACE-inhibitory peptide have also been isolated from fish proteins, such as sardines (Matsui et al., 1993), tuna (Kohama et al., 1988), cod (Kim et al., 2000), and bonito (Matsumura et al., 1993). These protein hydrolysates were usually produced using treatment with gastrointestinal (e.g., pepsin and pancreatin) or microbial (e.g., secreted by various microorganisms or supplied as commercial preparations, such as Protamex) enzymes under optimal conditions.

Nemopilema nomurai is a giant jellyfish species that has become abundant in offshore areas of Korea, China, and Japan in the last several years. Most fisheries dislike jellyfish because it

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is difficult to winnow the fishes from the bycaught jellyfishes, and because jellyfish toxins also decrease the quality of caught fish (Lee et al., 2007).

Salted jellyfish is a popular foodstuff because it contains about 97% moisture. Jellyfish have traditionally been used to treat asthma and hypertension in East Asian countries. Although jellyfish have been used for hypertensive therapy, information on the active compound(s) is lacking.

Thus, the purpose of this study was to determine the functional properties of the active compound of *N. nomurai* and its possible application as an antihypertensive therapy. We hydrolyzed *N. nomurai* jellyfish with six proteolytic enzymes that are widely used in industry (Alcalase, Flavozyme, Neutrase, papain, Protamex and trypsin), and then investigated the ACE-inhibitory activity of the purified hydrolysates.

Materials and Methods

Materials

The giant jellyfish, *Nemopilema nomurai*, was caught off the southern coast of Jeju Island, Korea. Only the umbrella (mesogloea) was collected, washed with deionized water, and then stored at -20°C until required. The proteases, trypsin, papain, and Flavozyme were purchased from Sigma Chemical Co. (St. Louis, MO, USA) and Protamex, Alcalase, and Neutrase were purchased from Novozymes (Bagsvaerd, Denmark). ACE from rabbit lung and a peptide substrate (Hip-His-Leu) were purchased from Sigma Chemical Co. The C₁₈ column used for the reverse-phase high-performance liquid chromatography (HPLC) was obtained from Phenomenex Inc. (Torrance, CA, USA).

Preparation of jellyfish hydrolysates

The influence of proteolytic enzymes on jellyfish hydrolysis was studied under the following conditions. First, jellyfish samples were lyophilized to remove excess water and then subjected to hydrolysis. Enzyme reactions were carried out for 1-4 h using 10% (w/v) lyophilized jellyfish and the following temperature and pH conditions: Alcalase, pH 7.0, 60°C; Flavozyme, pH 7.0, 50°C; Neutrase, pH 6.0, 50°C; papain, pH 6.0, 60°C; Protamex, pH 6.0, 40°C; and trypsin, pH 7.0, 40°C. To determine the optimal enzyme concentration, the ratio of reaction volume to enzyme quantity was varied from 1-4% (w/v). Reactions were stopped by heat treatment at 90°C for 15 min. The resultant slurries were centrifuged at 3,000 g for 10 min and the supernatants used as hydrolysates for further analysis.

In all experiments, the degree of hydrolysis was evaluated as the proportion (%) of α -amino nitrogen (N) with respect to the total N in the samples (Taylor, 1957).

Determination of ACE-inhibitory activity

The ACE-inhibitory activity assay was performed using a modified version of the method of Cushman and Cheung (1971). The standard reaction mixture contained 5 mM His-His-Leu as a substrate, 0.3 M NaCl, and 5 mU ACE in 50 mM sodium borate buffer (pH 8.3). A sample (50 μ L) was added to the enzyme solution (50 μ L) and then mixed with 8.3 mM Hip-His-Leu (150 μ L) containing 0.5 M NaCl to the same concentration as the standard reaction mixture. After incubation at 37°C for 30 min, the reaction was stopped by the addition of 1.0 N HCl (250 μ L). The resulting hippuric acid was extracted by the addition of 1.5 mL ethyl acetate. After centrifugation (800 g, 15 min), 1 mL of the upper layer was transferred into a new glass tube and evaporated at room temperature for 2 h in a vacuum. The extracted hippuric acid was dissolved in 3.0 mL distilled water, and absorbance was measured at 228 nm using a spectrophotometer (Model U-3210; Hitachi Co., Tokyo, Japan). The IC₅₀ value of the purified active fraction was determined by the standard method and defined as the concentration of inhibitor required to inhibit 50% of ACE activity.

Fractionation of jellyfish hydrolysates

To identify the ACE-inhibitory compound, the papain hydrolysate showing the highest inhibitory activity was selected and the resultant hydrolysate was fractionated through millipore membranes (Amicon Co., Beverly, MA, USA) with 5,000, 3,000, and 1,000 Da molecular weight cut-offs. Each fraction was lyophilized and then stored at -70°C until required.

Purification and peptide sequencing

Of the fractionated samples, fraction II showed the greatest inhibitory activity and was further purified by reverse-phase HPLC (5 μ M, 10 \times 250 mm; C₁₈ ODS 3100A; Phenomenex, Torrance, CA, USA) elution with a linear gradient of MeOH-H₂O (2 mL/min flow rate, ultraviolet detection at 214 nm). This procedure was repeated to increase purity. To determine the N-terminal sequence, the purified peptide showing the greatest ACE-inhibitory activity was subjected to automated Edman degradation on a protein sequencer (Perkin Elmer Model 470; Applied Biosystems, Foster City, CA, USA) according to the manufacturer's instructions. The amino acid sequence identity was aligned by searching database non-redundant protein sequences using the positron-specific iterative basic local alignment search tool (PSI-BLAST) of the National Center for Biotechnology Information (NCBI).

Results and Discussion

Degree of hydrolysis

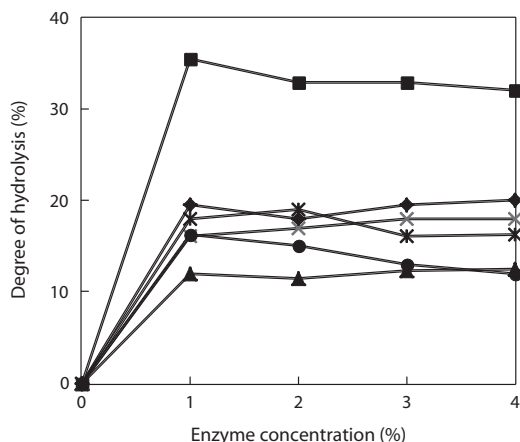


Fig. 1. Degree of hydrolysis of Giant jellyfish hydrolysates obtained by the action of proteolytic enzymes. Each enzyme reaction was varied in enzyme amounts (1-4% [w/v]). After 4 h reaction, all the treatments were followed by the same procedure. Alcalase, (◆); Flavozyme, (■); Neutrased, (▲); papain, (x); Protamex, (*); and trypsin, (●).

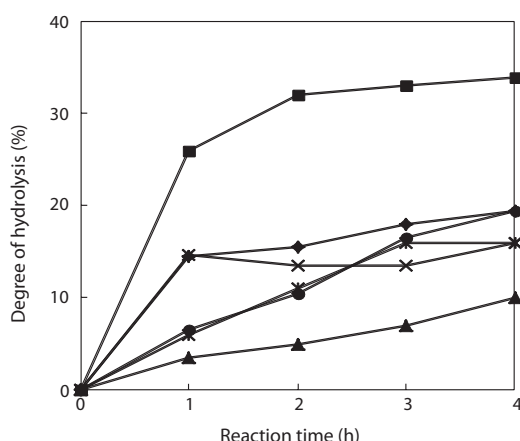


Fig. 2. Time dependence of the proteolytic enzyme reactions. Each enzyme reaction was conducted with 1% proteolytic enzymes (w/v) in the standard condition except for the reaction time (1-4 h) and all the treatments after the reaction were followed by the same procedure. Alcalase, (◆); Flavozyme, (■); Neutrased, (▲); papain, (x); Protamex, (*); and trypsin, (●).

The degree of hydrolysis of the jellyfish hydrolysates obtained using 1-4% enzyme and a 4-h reaction varied from 13.1-36.8%, and leveled off even when enzyme concentration was increased to 4% (Fig. 1). The degree of hydrolysis increased with reaction time; Flavozyme resulted in the greatest hydrolysis and Neutrased the lowest. Moreover, Alcalase and papain showed identical hydrolysis patterns and trypsin showed a pattern similar to that of Protamex (Fig. 2). These results indicate that a 4-h reaction time allowed for complete hydrolysis under the conditions mentioned above. Lee et al. (2001) determined the degree of hydrolysis and ACE-inhibitory activity of laver (*Porphyra tenera*) hydrolysates. Wide

ranges of hydrolysis (29.7-60.1%) and inhibitory activities (6.0-34.9%) were obtained, suggesting that the degree (%) of hydrolysis and inhibitory activities depended on the enzyme used.

ACE-inhibitory activities of enzymatic hydrolysates

To select the protease hydrolysate with the maximum ACE-inhibitory activity, lyophilized jellyfish were hydrolyzed with papain, trypsin, Flavozyme, Alcalase, Protamex, and Neutrased. Of these hydrolysates, that of papain revealed a high ACE-inhibitory activity in a 1-h reaction (76.73±2.10%) (Table 1). Increasing the enzyme concentration up to 3% or prolonging the reaction time up to 3 h had no effect on ACE-inhibitory activity, although the degree of hydrolysis increased (Fig. 2). Interestingly, Flavozyme-treated hydrolysate showed the highest degree of hydrolysis (Fig. 1) but the lowest inhibitory activity (Table 1). In contrast, the papain and Alcalase hydrolysates, which showed similar hydrolysis patterns, had differing ACE inhibition ratios (papain 76.73%, Flavozyme 36.01%), indicating that the primary structure of the hydrolysates obtained by these two enzyme reactions affected ACE inhibition (Fig. 3).

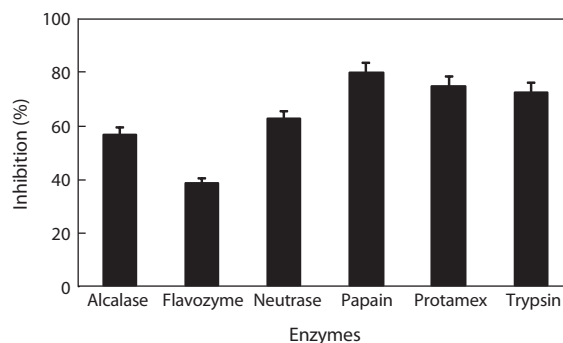


Fig. 3. Angiotensin-I-converting enzyme (ACE)-inhibitory activity of the Giant jellyfish hydrolysates. Same amount of the hydrolysate was added to the standard reaction mixture and the ACE-inhibitory activities of each hydrolysates were compared.

Table 1. ACE-inhibitory activities on incubation times and addition amounts of enzymes

Enzyme	ACE-inhibitory activity (%)		
	1%		3%
	1 h	3 h	3 h
Alcalase	39.61 ± 2.14	31.75 ± 1.63	20.46 ± 2.12
Flavozyme	36.46 ± 1.33	39.79 ± 1.52	47.60 ± 2.11
Neutrased	62.29 ± 1.24	58.47 ± 1.40	63.62 ± 1.53
Papain	76.73 ± 2.10	79.58 ± 1.31	75.23 ± 2.43
Protamex	70.01 ± 1.73	66.14 ± 2.23	72.74 ± 1.80
Trypsin	68.01 ± 1.82	65.85 ± 1.54	63.41 ± 2.22

ACE, angiotensin-I-converting enzyme.

Fujita and Yoshikawa (1999) isolated the prodrug-type ACE-inhibitory peptide LKPNM from the thermolysin digest of katsuobushi (*Katsuwonus pelamis*). He et al. (2006) reported that the molecular weight (Mw) < 1 kDa fraction of enzymatic hydrolysates of sea cucumbers possessed ACE-inhibitory activity. Here, we used six proteolytic enzymes; each hydrolysate showed differing degrees (%) of hydrolysis and inhibitory activities, suggesting that each contained several types of ACE-inhibitory peptide.

Characterization of the papain hydrolysate

Digestion with papain resulted in the greatest ACE-inhibitory activity. The papain hydrolysate was fractionated by ultrafiltration (5, 3, and 1 kDa Mw cut-offs) and then subjected to ACE-inhibitory activity assays. The ACE-inhibitory activities of fractions II and IV were 63.2 and 61.1%, respectively, whereas fractions I and III showed lower activities (30.2 and 27.8%, respectively) (Fig. 4). We assumed that most of the inhibitory compounds were within fractions II and IV, and that those activities were newly exposed by papain treatment, because no recognizable inhibitory activity was detected in a plain water extract (data not shown). The Mw of fractions II and IV were approximately 3,000-5,000 Da and less than 1,000 Da, respectively. The inhibitory activity of fraction II was similar to that of papain hydrolysate and fraction IV, suggesting that their specific activities were similar.

Byun and Kim (2001) revealed that activity increased markedly with decreasing Mw. A correlation was observed between hydrolysate Mw and the specificity of the ACE-inhibitory activity. Kinoshita et al. (1993) obtained two major fractions of high and low Mw from soy sauce, but reported that only the high-Mw fraction reduced blood pressure in hypertensive rats. In this study, we found no correlation between Mw and ACE-inhibitory activity.

Purification and characteristics of the inhibitory peptide

To purify the active compound, fraction II was subjected to reverse-phase column (ODS C₁₈) chromatography and the inhibitory activity of each eluate was determined. The FII-b fraction showed the highest inhibitory activity (59%), whereas the activities of FII-a, -b, and -c were less than 20% (Fig. 5). Therefore, the majority inhibitory activity was present in the FII-b fraction, although lower inhibitory activities were detected in the whole range (FII-a, -c, -d, -e, and -f). Treatment with papain thus produced a peptide of Mw 3,000-5,000 Da. To determine the primary structure, the FII-b fraction was again separated by chromatography to increase purity, and the N-terminal sequence was determined and the sequence identity compared. The N-terminal sequence of FII-b was Asp-Pro-Gly-Leu-Glu-Gly-Ala-His-Gly- and showed 87% identity to the collagen type IV alpha-2 fragment from *Rattus norvegicus*

and a predicted protein from *Nematostella vectensis*. These results suggested that the ACE-inhibitory peptide originated from collagen hydrolysate and had an IC₅₀ value of 3.8 µg/mL (Fig. 6). Although its entire primary structure is now be-

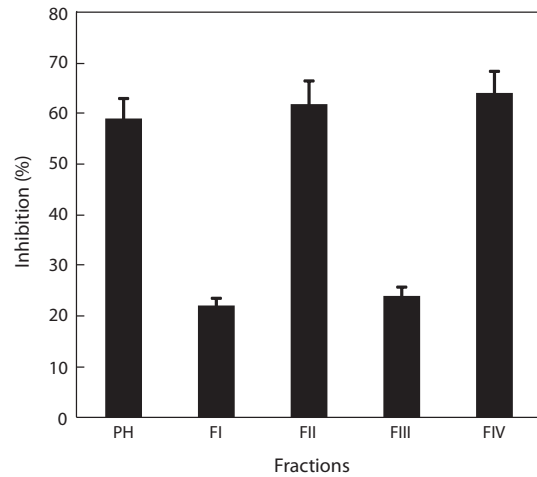


Fig. 4. Angiotensin-I-converting enzyme-inhibitory activity of papain hydrolysate fractionated using millipore membrane filters. The papain hydrolysate was fractionated periodically from smaller one. PH, papain hydrolysate; I, >5,000 Da; II, 5,000-3,000 Da; III, 3,000-1,000 Da; and IV, <1,000 Da.

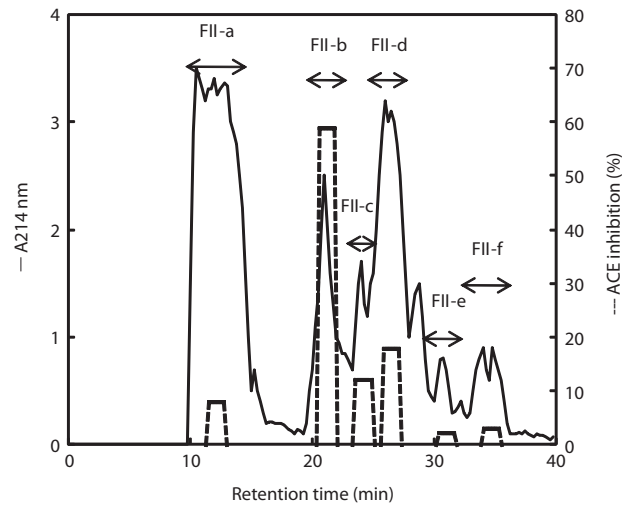


Fig. 5. The column chromatogram and angiotensin-I-converting enzyme (ACE)-inhibitory activity of each fraction. The fractions, FII-a, FII-b, FII-c, FII-d, FII-e and FII-f were collected separately and the inhibitory activities were conformed. —, Abs 214 nm; ---, ACE inhibition (%)

FII-b	1	DPGLQGAHG	9	FII-b	1	DPGLQGAHG	9
		DPG QGAHG				DPGLQGA G	
Sbjct1	350	DPGFQGAHG	358	Sbjct2	128	DPGFQGAHG	136

Fig. 6. N-terminal sequence alignment of FII-b by searching database non-redundant protein sequence using PSI-BLAST (NCBI).

ing determined, we believe that FII-b may represent a novel antihypertensive food additive.

Several types of ACE-inhibitory peptide have been reported, especially those originating from animal sources (Kohama et al., 1988; Matsui et al., 1993; Smacchi and Gobbetti, 1998; Jang and Lee, 2005). Most of these peptides have low Mw. Here, we isolated and characterized ACE-inhibitory peptides with a Mw of 3,000-5,000 Da originating from jellyfish collagen. FII-b was predicted to originate from an alpha-2 fragment of the type IV collagen of the jellyfish, *N. nomurai*. The precise size and sequence of the active compound remain unclear. The primary structure of the fragment is now being studied; this peptide represents an interesting new type of ACE inhibitor and will provide knowledge of the potential applications of jellyfish components as therapies for hypertension.

Acknowledgments

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