

Novel antioxidant and anti-inflammatory peptides from the Siamese crocodile (Crocodylus siamensis) hemoglobin hydrolysate

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2 (Crocodylus siamensis) hemoglobin hydrolysate

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31 Abstract Novel antioxidant and anti-inflammatory peptides were isolated from hydrolysates of Siamese 32 crocodile (Crocodylus siamensis) hemoglobin. Crocodylus siamensis hemoglobin hydrolysates (CHHs) were 33 obtained by pepsin digestion at different incubation times (2, 4, 6 and 8 H) at 37 °C and subjected to antioxidant 34 and anti-inflammatory activity assessment. CHH obtained by 2-H hydrolysis (2H-CHH) showed the highest 35 anti-inflammatory activity with respect to decreasing nitric oxide (NO) production, while the strongest 36 antioxidant activity was found for 6-H hydrolysis (6H-CHH) against nitric oxide radicals. To evaluate the anti-37 inflammatory and antioxidant activity of individual peptide components, 2H-CHH and 6H-CHH were purified 38 by semi-preparative HPLC. Peptide fraction P57 isolated from 6H-CHH was found to exhibit the highest nitric 39 oxide radical inhibition activity (32.0%). Moreover, purification of 2H-CHH yielded peptide fraction P16, 40 which displayed a high efficacy in decreasing NO production of macrophage RAW 264.7 cells (83.2%) and 41 significantly reduced pro-inflammatory cytokines and inflammatory mediators interleukin-6 (IL-6), interleukin-42 1 beta (IL-1β) and prostaglandin-E2 (PGE₂) production to about 2.0, 0.3 and 1.9 ng/mL, respectively. Using 43 LTO orbitrap XL mass spectrometry, active peptide sequences were identified as antioxidant KIYFPHF (KF7), 44 anti-inflammatory SAFNPHEKQ (SQ9) and IIHNEKVQAHGKKVL (IL15). Additionally, CHHs simulated 45 gastric and intestinal in vitro digestion positively contributed to antioxidant and anti-inflammatory activity. 46 Taken collectively, the results of this work demonstrate that CHHs contain several peptides with anti-47 inflammatory and antioxidant properties which may prove valuable as treatment or supplement against diseases 48 associated with inflammation and oxidative stress.

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50 *Keywords:* Crocodylus siamensis, enzymatic hydrolysis, hemoglobin hydrolysate, inflammation, oxidative stress,
 51 purification and identification

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Abbreviations: CHHs, Crocodylus siamensis hemoglobin hydrolysates; cHb, C. siamensis hemoglobin; COX-2,
cyclooxygenase-2; DH, degree of hydrolysis; IL-1β, Interleukin 1 beta; IL-6, Interleukin 6; NO, Nitric oxide;
PGE₂, Prostaglandin E2; TNF-α, Tumor necrosis factor alpha.

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61 **1. Introduction**

62 Inflammation, oxidative stress and free radical damage have recently been acknowledged as a global threat for 63 human health. Albeit being common by-products of a broad range of processes ensuring cellular homeostasis, 64 excessive generation of radical metabolites is unambiguously associated with a variety of serious disorders such 65 as cancer, heart disease, strokes, Alzheimer's, and premature aging [1]. Similarly, the inflammatory response 66 constitutes an important natural reaction of the host defense system to injury and the invasion of toxins or infectious particles. In the case of pathogenic infection, inflammation contributes to combatting the disease and 67 68 protecting vital parts of the body, while suspending the normal immune response and shunting certain metabolic 69 pathways at the same time. Specific pro-inflammatory mediators and cytokines, such as NO, PGE₂, TNF- α , IL-70 1β and IL-6, comprise vital parts of the inflammatory cascade and are generated to affect immune cell function 71 as well as proliferation activity [2]. In the long term, however, inflammatory processes are known to result in 72 progressive damage and are usually associated with excessive free radical release and oxidative stress [3].

73 The latter is defined as an inequality between the free radicals production and reactive metabolites resulting 74 in damage to cells and important biomolecules which severely impacts all organisms [4]. Free radicals, which 75 account for the majority of both reactive nitrogen species (RNS) and reactive oxygen species (ROS), are defined 76 as molecular species possessing one or more non-bonding paired electrons in spatially different orbitals of atoms 77 or molecules [5]. Also, they have high potential to either donate an electron to or accept an electron from other 78 molecules, consequently behaving as oxidants [6]. The latter process is capable of inflicting severe damage to 79 the membranes and nuclei of cells by oxidizing important biochemical molecules such as proteins, lipids, 80 carbohydrates, and DNA [7]. During the course of the inflammation high amounts of free radicals are produced within the inflamed tissue, in particular NO, which acts as a major pro-inflammatory mediator in the human 81 82 body and is generated from macrophage in response to the inflammation [8]. In this context NO is generally 83 regarded as the key mediator correlating inflammation and oxidative stress. The discovery and development of 84 substances capable of inhibiting either the activity or production of NO has therefore gained considerable 85 attention within the recent years as they may be utilized to reduce the detrimental effects of both inflammation 86 and oxidative damage.

Crocodylus siamensis, commonly called Siamese crocodile, is a small freshwater crocodilian populating
parts of Southeast Asia, including Thailand. Recently, several components of *C. siamensis* blood, i.e. plasma,
serum, white blood cells and hemoglobin have been reported to possess a broad spectrum of biological
properties, mainly attributed to abundance of a number of biologically active peptides and proteins. Among

91 these, hemoglobin constitutes the most abundant component and has been shown to exhibit antimicrobial [9],
92 antioxidant [10, 11] and anti-inflammatory activity [11, 12].

93 In addition to the direct extraction of specific proteins with desired biological activity, enzymatic hydrolysis 94 of protein sources has been established as a convenient method to generate novel protein fragments with 95 enhanced biological properties. A variety of animal proteins was shown to be applicable for producing bioactive 96 peptides via protein hydrolysis within recent decades, granting access with specific properties of interest such as 97 inhibitory activity on angiotensin I converting enzyme [13-16], antibacterial activity [17], antioxidant activity 98 [18, 19] and anti-inflammatory activity [20]. In this context, it is anticipated that the biological properties of C. 99 siamensis blood proteins and their potent antioxidant and anti-inflammatory activity may be further enhanced by 100 protein hydrolysis.

101 As a consequence, this study aimed to investigate the antioxidant and anti-inflammatory peptides from C. 102 siamensis hemoglobin hydrolysates (CHHs) derived by pepsin digestion. NO scavenging, linoleic peroxidation 103 and ferric reducing power assays were conducted to determine the antioxidant activity of the CHHs, whereas 104 MTT, NO (of macrophage RAW 264.7 cells), IL-6, IL-1 β and PGE₂ assays were used to investigate the anti-105 inflammatory activity. In vitro digestion and hemolytic activity were used to determine the application uses. 106 Furthermore, the identity of the antioxidant and anti-inflammatory peptides was elucidated via amino acid 107 sequence determination using ultra-high performance liquid chromatography-LTQ orbitrap XL mass 108 spectrometry.

109

110 **2. Materials and Methods**

111 2.1. Materials and reagents

112 RPMI 1640 medium, antibiotic/antimycotic (penicillin/streptomycin/amphotericin B), trypsin-EDTA and fetal 113 bovine serum (FBS) were purchased from Gibco (USA). Pepsin from porcine gastric mucosa, picrylsulfonic 114 acid solution (TNBS), sodium sulfite, L-leucine, ammonium thiocyanate, ferrous chloride, potassium 115 hexacyanoferrate, ferric chloride, Linoleic acid, 6-hydroxy-2,5,7,8-tetramethylchroman-2-carbonic acid 116 (Trolox), glutathione (GSH), 2,20-azobis-(2-amidinopropane)-dihydrochloride (ABAP), dimethyl sulfoxide 117 (DMSO), lipopolysaccharide (LPS), NaCl, HCl, PBS, bile salts solution, pancreatin solution, CaCl₂, Triton X-118 100 were purchased from Sigma-Aldrich (Germany). Sodium nitroprusside was purchased from Merck 119 (Germany). 3-(4,5-diamethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) was purchased from Eugene 120 (USA).

121 2.2. Crocodile hemoglobin preparation

122 Crocodile blood was collected from a slaughterhouse (Srirachamoda Co., Ltd.) in Thailand, and the extraction of 123 hemoglobin from red blood cells was performed following the method of Srihongthong et al. [9]. Blood was collected and transferred to 15 mL sterile tubes containing 0.08 g of EDTA. Blood samples were stored at 4 °C 124 125 overnight to allow blood cells to settle. Red blood cells (bottom laver) were collected in sterile tubes after 126 elimination of plasma (top layer) and white blood cells (middle layer). Isolated red blood cells were washed 127 three times with phosphate buffer saline (PBS), pH 7.0, and centrifuged at 3,000g for 5 min at 4 °C. Ice-cold 128 distilled water of five-fold volume was added to the RBC pellet, followed by vigorous mixing and allowing the 129 mixture to settle for 10 min. After centrifugation at 10,000g for 20 min at 4 °C, the supernatant was collected for 130 lyophilization and then stored at -70 °C.

131

132 2.3. Enzymatic hydrolysis

Enzymatic hydrolysis was performed according to the method of Yu et al. [21]. Shortly, the hemoglobin solution was digested by pepsin with a ratio of enzyme to substrate of 1:100 (w/w) at 37 °C for 2, 4, 6 and 8 H and boiled at 95 °C for 10 min to quench the reaction by inactivating the enzyme. The hydrolysis condition of hemoglobin by pepsin was performed at pH 2.0 (adjusted with 1 M HCl), followed by removal of insoluble components by centrifugation at 7,168*g* for 20 min. The supernatant was collected and adjusted to pH 7.0 by addition of 1 M HCl or 1 M NaOH. Finally, the supernatants of the crocodile hemoglobin hydrolysates (CHHs) were lyophilized and stored at -20 °C.

140

141 2.4. Degree of hydrolysis

142 The degree of hydrolysis was determined following the method of Benjakul et al. [22]. Briefly, 125 μ L of CHHs 143 were added to 2.0 mL of 0.21 M sodium phosphate buffer, pH 8.2, followed by addition of 1 mL of 0.01% 144 TNBS solution. The mixture was incubated in a water bath at 50 °C for 30 min in the dark, and 2 mL of 0.1 M 145 sodium sulfite was added to stop the reaction. The mixture was then allowed to cool for 15 min. The absorbance 146 was measured at 420 nm and the α -amino acid content expressed in terms of L-leucine. The percentage of the 147 degree hydrolysis was calculated using the formula:

148 $DH = [(L_t - L_0)/(L_{max} - L_0)] \times 100$

149 where L_0 determines the amount of α -amino acid expressed in the sample. L_t corresponds to the amount of α -150 amino acid released at time t. L_{max} determines the maximum amount of α -amino acid after hydrolysis by 5 M 151 HCl at 100 °C for 24 H.

- 152
- 153 **2.5.** Nitric oxide (NO) scavenging assay

154 Nitric oxide (NO) was generated from sodium nitroprusside and measured using Griess reagent. The assay was 155 conducted according to the method of Yen et al. [23] with modifications. For the experiment, 250 µL sodium 156 nitroprusside (10 mM) in PBS were mixed with 15, 31, 62, 125, 250 and 500 µg/mL of CHHs and incubated at 157 25 °C for 2 H. An aliquot of the incubated solution (100 µL) was added to 100 µL of Griess reagent (1% 158 sulfanilamide in 5% phosphoric acid and 0.1% naphthylethylenediamine dihydrochloride) and incubated in the 159 dark for 10 min before measuring the absorbance at 540 nm. Butylated hydroxytoluene (BHT) was used as a 160 positive control. All samples were analyzed in quadruplicate. The percentage of inhibition was calculated using 161 the formula:% inhibition = $[(Abs_{cont} - Abs_{test})/Abs_{cont}] \times 100.$

162

163 2.6. Linoleic peroxidation assay

This assay was performed following the method of Ledesma et al. [24] with modifications. Briefly, 50 µL of 164 CHHs (0.005, 0.05, 0.5, 5, 50 and 500 μ g/mL) were added to 50 μ L of linoleic acid (0.05% v/v) solution. 0.07 165 166 M ABAP (10 μ L) was added and the solution mixed for 10 min. After addition of 150 μ L 20% (v/v) acetic acid, the mixture was incubated at 70 °C for 1 H. Twenty microliter of each solution was then added to a 96-well 167 168 plate already containing 75% ethanol (160 μ L), 15% ammonium thiocyanate (10 μ L) and 10 mM ferrous 169 chloride (10 µL). The new solution was mixed and incubated at for 3 min. The absorbance was measured at 500 170 nm and Trolox used as positive control. All samples were analyzed in sextuplicate. The antioxidant activity was 171 analyzed and the percentage of antioxidant inhibition (% AI) of each sample was calculated using the following 172 formula:

173 % AI (antioxidant inhibition) =
$$100 \times [Abs_{cont} - Abs_{test}]/Abs_{cont}$$

174

175 2.7. Ferric reducing power assay

176 This assay was modified from the method of Girgih et al. [25]. Briefly, 250 μ L of CHHs (0.005, 0.05, 0.5, 5, 50 177 and 500 μ g/mL) and positive control (glutathione) were added to 250 μ L of 0.2 M phosphate buffer, pH 6.6. 250

- $178 \qquad \mu L \ of \ 1\% \ (w/v) \ potassium \ hexacyanoferrate \ solution \ were \ added, \ vortexed \ and \ the \ mixture \ incubated \ at \ 50 \ ^{\circ}C$

179 for 20 min. The reaction was quenched by adding 250 μ L of 10% TCA and incubated for 10 min before 180 centrifugation at 800*g* for 10 min. Aliquots of each solution (30 μ L) were then transferred to a 96-well plate 181 already containing double distilled water (160 μ L) and 0.1% (w/v) ferric chloride (10 μ L). The reaction mixture 182 was homogenized and incubated at 25 °C for 10 min, followed by measuring the absorbance at 700 nm. All 183 samples were analyzed in quadruplicate. The antioxidant activity for each sample was expressed as Trolox 184 (mM) equivalents and calculated using the equation of the standard curve of the positive control.

185

186 **2.8.** Cell culture

187 A murine macrophage cell line (RAW 264.7) was purchased from the American Type Culture Collection 188 (American Type Culture Collection [ATCC], USA) and cultured in RPMI 1640 medium supplemented with 189 10% heat-inactivated fetal bovine serum (FBS), 100 μ g/mL of streptomycin, 100 U/mL of penicillin, 25 μ g/mL 190 amphotericin B and incubated at 37 °C in a 5% CO₂ humidified atmosphere.

191

192 2.9. Measurement of nitric oxide

193 This assay was conducted according to the method of Phosri et al. [11]. CHHs (62, 125, 250 and 500 μ g/mL) 194 were mixed with LPS (100 ng/mL), and the resulting solution incubated with RAW 264.7 cells. Another 195 incubation was set up between LPS and RAW 264.7 cells at 37 °C in a 5% CO₂ humidified atmosphere for 24 196 H. 100 μ L of culture medium from each CHH sample were slightly mixed with 100 μ L of Griess reagent and 197 incubated at 25 °C for 10 min. The absorbance was measured at 540 nm using a microplate reader (BioRad, 198 Model 680, USA). Nitric oxide (NO) production was calculated as percentage of control. All samples were 199 analyzed in quadruplicate.

200

201 2.10. Cell viability

Cell viability determination was performed using the MTT assay according to the method of Phosri et al. [11]. RAW 264.7 cells (1×10^5 cells/mL) were cultured on a 96-well plate overnight. Lipopolysaccharides (LPS) were co-incubated with CHHs in defined concentrations (62, 125, 250 and 500 µg/mL) and 100 ng/mL of LPS at 37 °C in a 5% CO₂ humidified atmosphere. After incubation for 24 H the medium was discarded. MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) (0.5 mg/mL) was added to RAW 264.7 cells and incubated at 37 °C in a 5% CO₂ humidified atmosphere for 30 min before the medium was discarded. DMSO was added and the reaction mixture incubated at 25 °C for 30 min. The absorbance was measured at 570 nm and 209

the cell viability evaluated by comparing the absorbance with that of the control for each sample. All samples 210 were analyzed in quadruplicate.

211

212 2.11. Measurement of IL-6, IL-1β and PGE2

213 Aliquots of culture medium employed in the NO assay were further used for determination of IL-6, IL-1 β and 214 PGE₂ expression using the ELISA kit and following the instructions in the manufacturer's manual (R&D, 215 Minneapolis, MN, USA).

216

217 2.12. Peptide purification from crocodile hemoglobin hydrolysates

218 The purification of CHH was performed according to the method of Srihongthong et al. [9] with modifications. 219 A HPLC system connected to a C-18 reverse phase column (10×150 mm; SunfireTm prep 5 µm) was employed 220 in the purification process. Fractions were separated using a mobile phase system consisting of mobile phase A 221 (0.1% TFA in deionized water) and mobile phase B (60% acetonitrile in 0.1% TFA) with a flow rate of 1.0 mL/ min. A gradient of 0-25% (v/v) B over 15 min, then 25-100% (v/v) B over 95 min was applied. The elution 222 223 peaks were collected by monitoring the absorption at 220 nm. The antioxidant and anti-inflammatory activity of 224 all peaks were assayed, followed by determination of the amino acid sequences for each peak showing 225 significant biological activity.

226

227 2.13. Amino acid sequence analysis

228 The active fractions P16 of 2H-CHH and P57 of 6H-CHH were selected and the peptides identified using LTQ 229 orbitrap XL Mass spectrometry employing the following search parameters: non-specified enzymatic cleavage 230 with three possible missed cleavages, +/-0.8 Da mass tolerances for MS and MS/MS, a peptide mass tolerance 231 of +/-5 ppm, methionine oxidation and Gln->pyro-Glu (N-term Q) variable modification, and monoisotopic 232 mass. Data were additionally processed at the Mascot Server (http://www.matrixscience.com/) using MS/MS 233 ion searches against SwissProt (current release).

234

235 2.14. In vitro simulated gastric and intestinal digestion

236 In vitro digestion models were used to simulate the condition of human stomach and intestine for observing 237 remaining activity of CHHs as described by Cheong et al. [26] with modification. The ratio of sample: simulated 238 gastric: simulated intestinal was 1: 1.5: 2. Simulated gastric solution was prepared by dissolving 2 g of NaCl, 7

239 mL of HCl and 3.2 g pepsin with DI up to 1 L. The pH was adjusted to 1.6 using 1 M HCl and maintained at 37 240 °C in a temperature-controlled water bath (Julabo, Germany). Each CHHs were mixed and pH was adjusted to 2.5 using 1 M HCl then incubated at 37 °C for 1 H. The simulated intestinal solution was prepared by mixing 241 160 mL bile salts solution (5 mg/mL in PBS), 100 mL of pancreatin solution (4.8 mg/mL in PBS) and 40 mL of 242 243 CaCl2 solution (110 mg/mL in PBS). The pH was adjusted to 7 and maintained at 37 °C in a temperature-244 controlled water bath. After the gastric digestion, samples were immediately adjusted to pH 7 with 1 M NaOH. 245 Digested samples were added to simulated intestinal solution and incubated at 37 °C for 2 H. Samples were 246 adjusted to pH 9 to ensure enzyme inactivation.

247

248 2.15. Hemolytic activity

The hemolytic activity was determined according to the method of Pata et al. [27]. After isolation of the erythrocytes by centrifugation at 1,000*g* for 5 min, human red blood cells (hRBCs) were washed three times with PBS, pH 7.4 and then adjusted to 2% (v/v). Ten microliter CHHs were added separately to the 100 μ L reaction solution, incubated for 1 H at 37 °C and centrifuged at 1,000*g* for 5 min. 100 μ L of supernatant was then transferred into 96-well plate. The absorbance was measured at 415 nm. 1% (v/v) Triton X-100 and DI used as positive and negative control respectively. All samples were analyzed in quadruplicate.

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256 2.16. Statistical analysis

Statistical analysis was performed using ANOVA and followed by Dunnett's test (Prism 5.0, GraphPad Inc., San Diego, CA, USA). Data are presented as mean \pm SEM. A value of P < 0.05 was accepted to be significant (*P < 0.05, ** P < 0.01, ***P < 0.001).

260

261 **3. Results and Discussion**

262 3.1. Preparation of crocodile hemoglobin hydrolysates (CHHs) and degree of hydrolysis (DH)

In the present study, crocodile (*C. siamensis*) hemoglobin was extracted from red blood cells. To obtain active fragments, crocodile hemoglobin was hydrolyzed by pepsin digestion at different reaction times. The extent of enzymatic protein degradation was evaluated by degree of hydrolysis (DH) which was 20.4%, 23.1%, 25.0% and 35.9% for 2 H, 4 H, 6 H and 8 H of incubation, respectively (Fig. 1). Results indicated that cleavage of peptide bonds was higher in longer enzymatic hydrolysis.

269 3.2. Antioxidant activity, purification and amino acid sequence identification of CHHs

270 NO is a very unstable radical which produces highly reactive molecules [28]. Intact crocodile Hb and all 271 investigated Hb hydrolysates inhibited NO radicals (Fig. 2a). Notably, the hydrolysates had superior antioxidant 272 activity compared to intact Hb at all concentrations (15-500 µg/mL). The NO-scavenging activity of the intact 273 protein (500 µg/mL) was 75.7%, while the inhibitory ability of the hydrolysates, including 2H-CHH, 4H-CHH, 274 6H-CHH and 8H-CHH, showed inhibition percentages of 77.4%, 78.2%, 88.3% and 83.1%, respectively. 275 Moreover, BHT employed as positive standard displayed 77.8% NO radical inhibition. These results indicate 276 that CHHs obtained by pepsin digestion possess pronounced antioxidant activity against nitric oxide radicals. 277 From the report of Kabbua et al. [29] found that recombinant α -globin from cHb can act as a heme-nitric oxide 278 and/or oxygen binding (H-NOX) hemoprotein. Furthermore, the value of NO inhibition activity of CHH was 279 higher than yellowfin tuna hydrolysate [30].

280 Lipid peroxidation is considered to be a free radical process involving a source of secondary free radical, which 281 can act as secondary messenger or directly react with other biomolecules, enhancing biochemical lesions. The 282 results of linoleic peroxidation assay reveal that 2H-CHH, 4H-CHH, 6H-CHH and 8H-CHH (at concentrations 283 of 0.005-500 µg/mL) effected significant inhibition of linoleic peroxidation in a concentration dependent 284 manner when compared with Trolox. The inhibitory ability of all hydrolysates, including 2H-CHH, 4H-CHH, 285 6H-CHH and 8H-CHH were 94.2%, 79.0%, 97.0% and 82.2%, respectively. Moreover, Trolox (500 µg/mL) 286 showed 94.4% of linoleic peroxidation inhibition and intact Hb showed 89.1% inhibition (Fig. 2b). Furthermore, 287 the linoleic peroxidation inhibition of protein hydrolysates obtained from smooth hound has been investigated. 288 As shown by the results, the value of this activity was lower than that obtained in this study [31].

289 2H-CHH, 4H-CHh, 6H-CHH and 8H-CHH at the highest concentration (500 μ g/mL) displayed significant ferric 290 ion reducing power. 4H-CHH displayed the highest reduction activity equivalent to Trolox of about 2.1 mM, 291 while intact Hb, 2H-CHH, 6H-CHH and 8H-CHH displayed reduction equivalents of 0.8, 1.2, 1.4 and 0.6 mM 292 Trolox (Fig. 2c). Moreover, 500 μ g/mL glutathione (positive control) affected significant ferric ion reduction 293 equivalent to Trolox at 18.60 mM (data not shown). It has been reported that hydrolysates with high reducing 294 power show a great ability to donate electrons to form stable compounds and thereby interrupt the free radical 295 chain reactions and showing antioxidant activity [32].

296 Due to displaying the highest antioxidant activity in prior experiments, 6H-CHH was subjected to purification

by reverse phase C-18 column semi-preparative HPLC to identify the contained active peptide components. As

shown in Fig. 3a, 61 individual fractions were eluted. All elution peaks were normalized to a concentration of 78

μg/mL and screened with respect to antioxidant activity by nitric oxide scavenging assay. After the activity
screening, several active fragments including P2, P39, P40, P43, P45, P46 and P57 were associated with nitric
oxide radical inhibition rates of 19.5%, 21.2%, 17.3%, 16.1%, 18.7%, 18.1% and 32.0%, respectively. The
highest activity in inhibiting nitric oxide radicals was found for P57 (32.0%), while the positive control (BHT)
showed 68.1% nitric oxide radical inhibition (Fig. 3b).

304 The primary structures of the purified antioxidant peptides were elucidated using LTQ orbitrap XL mass 305 spectrometry (Fig. 4). Amino acid sequence of P57 was KIYFPHF (KF7) with molecular mass of 476.26 Da. As 306 shown in Table 1, antioxidant peptide KF7 showed 42% hydrophobicity and a +1 net charge. Alignment of the 307 amino acid sequences of the peptide fragments with cHb indicated the antioxidant peptides originated from the 308 α -subunit, this observation is in excellent agreement with results of Srihongthong et al. [9], who reported that 309 the α -subunit of cHb exhibited higher antioxidant activity than the β -subunit. Peptide fractions from acid-310 digested cHb with a molecular mass of 180 to 3,000 Da are reported to exhibit antioxidant activity [33]. 311 Notably, mass spectrometric analysis further revealed an apparent correlation of peptide length with biological 312 activity as antioxidant peptides were found to be smaller and shorter molecules. In a previous study, Peña-313 Ramos et al. [34] reported that the presence of particular amino acids including His, Tyr, Met, Lys, Trp and Pro 314 correlates with increased antioxidant potency of most food-derived peptides. Likewise, the presence of Leu, Ile, 315 His, Met, Tyr, Lys and Trp is assumed to contribute to the reducing power of protein hydrolysates [35]. In 316 addition, peptides containing His residues have been documented to exhibit protective effects against lipid 317 peroxidation. An imidazole ring of His has been implicated in the donation of hydrogen and trapping of lipid 318 radicals [36]. Trp, Tyr and Met exhibited the highest antioxidant properties, followed by Phe, Cys and His [37]. 319 Nitric oxide, however, is also scavenged by CHH, presumably due to the presence of reactive thiol groups [38].

320

321 3.3. Anti-inflammatory activity, purification and amino acid sequences identification of CHHs

The anti-inflammatory property of intact Hb and CHHs was evaluated on the basis of NO production and cell viability against macrophage RAW 264.7 cells (Fig. 5). After induction of inflammation in RAW 264.7 cells by LPS for 24 H, the percentage of nitric oxide production was defined as 100% when treated with LPS. Fig. 5a, 2H-CHH, 4H-CHH, 6H-CHH and 8H-CHH (at concentrations of 62-500 μ g/mL) show nitric oxide production decrease in a concentration dependent manner. Intact Hb, 2H-CHH, 4H-CHH, 6H-CHH and 8H-CHH at a concentration of 500 μ g/mL show nitric oxide production at 56.5%, 52.6%, 68.6%, 62.0% and 60.9%, respectively. In order to evaluate potential cytotoxic effects of CHHs, the viability of RAW 264.7 cells treated 329 with defined concentrations of CHHs was determined (Fig. 5b). The results showed that intact Hb, 2H-CHH, 330 4H-CHH, 6H-CHH and 8H-CHH at concentrations of 62-500 µg/mL had no observable effect on cell viability 331 while the viability percentage of cells treated with LPS only was defined as 100%. This result indicates that 332 CHHs show high efficacy to reduce nitric oxide production and are essentially non-toxic to macrophage RAW 333 264.7 cells, these results correlate with that from the report of Phosri et al. [11] who reported that cHb provides 334 anti-inflammatory activity via the suppression of nitric oxide synthase (NOS), which inhibits the NO 335 production. Jangpromma et al. [39] reported that cHb significantly decreased the production of NO in LPS-336 stimulated RAW 264.7 cells and decreased inducible nitric oxide synthase (iNOS). In addition, results showed 337 that CHHs have higher anti-inflammatory activity than gastropod (Harpa ventricosa) hydrolysate [40].

338 Since 2H-CHH was found to possess the highest anti-inflammatory activity, its peptide constituents were purified by reverse phase C-18 column semi-preparative HPLC. The resulting chromatogram depicted in Fig. 6a 339 340 consists of 59 fraction peaks which were diluted to a concentration of 31 µg/mL prior to anti-inflammatory 341 activity determination. The assay revealed 6 peaks (P16, P17, P28, P29, P30 and P38) which show nitric oxide 342 production at 83.2%, 86.0%, 87.9%, 88.8%, 86.3% and 83.6% (Fig. 6b) with no effect on the viability of RAW 343 264.7 cells (Fig. 6c). Moreover, these fractions were also selected to determine the influence on the expression 344 levels of IL 6, IL-1β and PGE₂ (Table 2). The results indicate that P16, P17, P28, P29, P30 and P38 are able to 345 significantly decrease IL-6 and PGE₂ production, whereas P16 and P17 further inhibited IL-1β production. 346 Notably, P16 showed the highest activity to decrease IL-6 levels (approximately 2.0 ng/mL) compared to LPS 347 and effectively decreased the IL-1 β level to about 0.3 ng/mL. P16 showed the highest activity to reduce PGE₂ 348 levels (approximately 1.9 ng/mL) when compared to LPS. Furthermore, P16 was found to reduce the production 349 of pro-inflammatory cytokines and inflammatory mediators IL-6, IL-1 β and PGE₂, which are associated with the 350 inflammation-related in neuropathological diseases, for example Alzheimer's, multiple sclerosis and cerebral 351 ischemia [41, 42]. IL-6 that acts as a multifunctional cytokine was up-regulated by lipopolysaccharide (LPS). In 352 inflammation, trauma and autoimmune diseases were found high expression levels of IL-1 β and IL-6 [43]. 353 Furthermore, IL-1 β induced IL-6 production is mediated predominantly through the p38 MAPK/NF- κ B 354 pathway [44]. cHb was found to exhibit anti-inflammatory effects in the cotton pellet model and RAW 264.7 355 cells by reducing expression levels of IL-1 β , TNF- α , COX-2 and IL-6 mRNA [39, 45]. The collected results in 356 this work indicate that the anti-inflammatory activity of sample might be related to an interaction with the 357 MAPK/NF- κ B pathway by their ability to decrease pro-inflammatory cytokine and inflammatory mediator 358 production.

359 The primary structures of the purified anti-inflammatory peptides were elucidated using LTQ orbitrap XL mass 360 spectrometry (Fig. 4). Amino acid sequences of P16 were determined as SAFNPHEKQ (SQ9) and 361 IIHNEKVQAHGKKVL (IL15) corresponding to molecular mass of 529.26 Da and 857.51 Da, respectively. As 362 shown in Table 1, anti-inflammatory peptides SQ9 and IL15 had hydrophobicity values of about 22% and 40%, 363 together with net charges of 0 and +2, respectively. The sequence of the anti-inflammatory peptide showed 364 strong correlation with the β -chain of cHb. Likewise, the anti-inflammatory activity of SQ9 and IL15 was traced 365 back to the presence of specific amino acids. According to reports, anti-inflammatory peptides can be found in a 366 wide molecular weight range [20, 46]. Hydrophobic amino acid side chains (e.g. Leu, Phe, Val, Ile, and Trp) as well as positively charged amino acids (Lys, Arg and His) were documented to have a major influence on the 367

- 368 anti-inflammatory activity of peptides [47-52].
- 369

370 3.4. In vitro simulated gastric and intestinal digestion of CHHs and its activity

371 Simulated gastric and intestinal digestion (in vitro digestion model) was used as the conditions of human 372 stomach and intestine. The degree of hydrolysis of 2H-CHH and 6H-CHH after intestinal digestion was 373 increased about 3.7% and 8.8%, respectively. From the results, it is inferred that CHH tended to be more 374 hydrolyzed by intestinal digestion. Gastrointestinal tract is known to be a major oxidation site where various 375 free radicals are generated in the digestion process [53]. As shown in Fig. 7a, 6H-CHH after intestinal digestion 376 was inhibited nitric oxide (42.5%, the concentration of 125 μ g/mL) and exhibited higher reducing power (2.8 377 mM Trolox equivalent, the concentration of 500 µg/mL) than 6H-CHH before intestinal digestion (Fig. 7b). 378 With excellent agreement with Moure et al. [36] who reported that the reducing power increased with higher 379 degree of hydrolysis. The anti-inflammatory activity of 2H-CHH after intestinal digestion showed NO 380 production (19.8%, the concentration of 125 µg/mL) without toxicity against macrophage RAW 264.7 cells 381 (Figs. 8a and 8b). During the course of the inflammation high amounts of free radicals are produced within the 382 inflamed tissue, in particular NO, which acts as a major pro-inflammatory mediator in the human body and is 383 generated from macrophage in response to the inflammation [8]. In this context, NO is generally regarded as the 384 key mediator correlating inflammation and oxidative stress. The capability of NO production has decreased 385 resulting in the reduction of detrimental effects of both inflammation and oxidative damage. The hemolytic 386 activity was performed to determine the toxicity against human red blood cells and the results showed that both 387 2H-CHH and 6H-CHH at all concentrations were not toxic against human red blood cells (Figs. 9a and 9b). So 388 far, only a few bioactive peptides including peptides in this study displaying activities in vitro that have been

proven effective *in vivo*. Results indicated that both 2H-CHH and 6H-CHH with intestinal digestion havepotential to be used as an alternative source of antioxidant and anti-inflammatory agents with safety.

391

4. Conclusions

393 A number of novel antioxidant and anti-inflammatory peptides were derived from pepsin hydrolysis of cHb. 394 Experimental evidence collected in this study indicates the antioxidant fragment is a comparatively small 395 peptide, consisting of only 7 amino acid residues with slightly positive charge and low hydrophobicity. 396 Similarly, the peptides with anti-inflammatory activity were found to comprise 9 and 15 amino acid residues, 397 respectively, and are also characterized by positive charge and low hydrophobicity. This is the first report to 398 identify antioxidant and anti-inflammatory peptides from pepsin digested cHb. Due to their remarkable 399 biological activity, ease of production and absence of cytotoxicity, these naturally-derived peptides are believed 400 to bear a great potential for a future application as health promoting supplements and therapeutic agents against 401 inflammation and oxidative stress-related conditions.

402

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411

412 6. References

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TABLE 1Structural characteristics of the anti-inflammatory peptides (P16) from 2H-CHH and
antioxidant peptide (P57) from 6H-CHH

	Properties Anti- inflammatory Anti- inflammatory		Peptide sequences		Hydrolysates	% hydropho- bicity	Net charge	Sequence alignment
			SAFNPHEKQ (SQ9)		2Н-СНН	22%	0	cHb β-subunit (position 2-10)
			IIHNEKVQAHGKKVL (IL15)		2Н-СНН	40%	+2	cHb β-subunit (position 55-69)
	Antioxid	lant	KIYFPHF (K	KF7)	6H-CHH	42%	+1	cHb α-subunit (position 41-47)
508								
509								
510	-							
511		TAB				ı in LPS-stimulat of 2H-CHH at a		
512		////	μg/m	-	<i></i>	.		
513		Fraction peaks (P)		The production concentration (ng/mL)				
514	_			IL-6		ΙL-1β	PC	GE_2
		I DC		10 (17 . 0	010	0.0.000		0 4 4 9
515		LPS		10.647 ± 0	.213	0.360 ± 0.002	6.4	486 ± 0.163
515 516		P16		10.64 / ± 0 2.019*** ±		0.360 ± 0.002 $0.315^{***} \pm 0.002$		486 ± 0.163 $371^{***} \pm 0.068$
					= 0.000		1.8	
516		P16		2.019*** ±	= 0.000 = 0.447	0.315*** ± 0.002	1.8	371*** ± 0.068
516 517		P16 P17		2.019*** ± 7.287*** ±	= 0.000 = 0.447 = 0.110	$0.315^{***} \pm 0.002$ 0.338 ± 0.001	1.8 1.8 3.0	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$
516 517 518		P16 P17 P28		2.019*** ± 7.287*** ± 7.967*** ±	= 0.000 = 0.447 = 0.110 .155	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$	1.8 1.8 3.0 3.5	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$
516 517 518 519		P16 P17 P28 P29		2.019*** ± 7.287*** ± 7.967*** ± 9.340* ± 0	= 0.000 = 0.447 = 0.110 .155 .116	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$	1.8 1.8 3.0 3.5 2.8	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$
516 517 518 519 520	*denotes	 P16 P17 P28 P29 P30 P38 	05 and *** der	$2.019^{***} \pm 10^{-1}$ $7.287^{***} \pm 10^{-1}$ $7.967^{***} \pm 10^{-1}$ $9.340^{*} \pm 10^{-1}$ $9.407^{*} \pm 10^{-1}$ $4.707^{***} \pm 10^{-1}$	= 0.000 = 0.447 = 0.110 .155 .116 = 0.173	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$ $0.430^{***} \pm 0.002$ $0.538^{***} \pm 0.011$	1.8 1.8 3.0 3.5 2.8 3.1	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$ $380^{***} \pm 0.205$
516 517 518 519 520 521		P16 P17 P28 P29 P30 P38 s P < 0.		2.019*** \pm 7.287*** \pm 7.967*** \pm 9.340* \pm 0 9.407* \pm 0 4.707*** \pm notes $P < 0.0$	= 0.000 = 0.447 = 0.110 .155 .116 = 0.173	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$ $0.430^{***} \pm 0.002$ $0.538^{***} \pm 0.011$ sed as a mean ± S.	1.8 1.8 3.0 3.5 2.8 3.1	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$ $380^{***} \pm 0.205$ $181^{***} \pm 0.377$
516 517 518 519 520 521 522		P16 P17 P28 P29 P30 P38 s P < 0.		2.019*** \pm 7.287*** \pm 7.967*** \pm 9.340* \pm 0 9.407* \pm 0 4.707*** \pm notes $P < 0.0$	= 0.000 = 0.447 = 0.110 .155 .116 = 0.173 001. Data expres	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$ $0.430^{***} \pm 0.002$ $0.538^{***} \pm 0.011$ sed as a mean ± S.	1.8 1.8 3.0 3.5 2.8 3.1	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$ $380^{***} \pm 0.205$ $181^{***} \pm 0.377$
516 517 518 519 520 521 522 523		P16 P17 P28 P29 P30 P38 s P < 0.		2.019*** \pm 7.287*** \pm 7.967*** \pm 9.340* \pm 0 9.407* \pm 0 4.707*** \pm notes $P < 0.0$	= 0.000 = 0.447 = 0.110 .155 .116 = 0.173 001. Data expres	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$ $0.430^{***} \pm 0.002$ $0.538^{***} \pm 0.011$ sed as a mean ± S.	1.8 1.8 3.0 3.5 2.8 3.1	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$ $380^{***} \pm 0.205$ $181^{***} \pm 0.377$
516 517 518 519 520 521 522 523 523		P16 P17 P28 P29 P30 P38 s P < 0.		2.019*** \pm 7.287*** \pm 7.967*** \pm 9.340* \pm 0 9.407* \pm 0 4.707*** \pm notes $P < 0.0$	= 0.000 = 0.447 = 0.110 .155 .116 = 0.173 001. Data expres	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$ $0.430^{***} \pm 0.002$ $0.538^{***} \pm 0.011$ sed as a mean ± S.	1.8 1.8 3.0 3.5 2.8 3.1	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$ $380^{***} \pm 0.205$ $181^{***} \pm 0.377$
516 517 518 519 520 521 522 523 524 525		P16 P17 P28 P29 P30 P38 s P < 0.		2.019*** \pm 7.287*** \pm 7.967*** \pm 9.340* \pm 0 9.407* \pm 0 4.707*** \pm notes $P < 0.0$	= 0.000 = 0.447 = 0.110 .155 .116 = 0.173 001. Data expres	$0.315^{***} \pm 0.002$ 0.338 ± 0.001 $0.420^{***} \pm 0.002$ $0.412^{***} \pm 0.001$ $0.430^{***} \pm 0.002$ $0.538^{***} \pm 0.011$ sed as a mean ± S.	1.8 1.8 3.0 3.5 2.8 3.1	$371^{***} \pm 0.068$ $377^{***} \pm 0.042$ $046^{***} \pm 0.220$ $578^{***} \pm 0.035$ $380^{***} \pm 0.205$ $181^{***} \pm 0.377$

529 Figure c	aptions
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530

FIG.1 Degree of hydrolysis (% DH) of CHH after pepsin digestion for 2, 4, 6 and 8 H. Hemoglobin
hydrolyzed enzymatically by pepsin displayed a direct correlation between the rate of hydrolysis (DH)
and the time of incubation (H).

534

535	FIG.2 (a) Nitric oxide scavenging activ	ity of CHHs at concentrations of 15-500 μ g/mL. Each bar displays the
536	mean ± SEM of four demonstrati	tons. (** $P < 0.01$ and *** $P < 0.001$) probability levels compared
537	with BHT. (b) Linoleic peroxidat	tion activity of CHHs at concentration of 0.005-500 μ g/mL. Each bar
538	displays the mean \pm SEM of six c	<i>lemonstrations.</i> (* $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$)
539	probability levels compared with	Trolox and (c) reducing power of CHHs at concentration of 0.005-
540	500 μg/mL expressed in Trolox e	quivalents. Each bar displays the mean \pm SEM of four demonstrations.
541	(***P < 0.001) probability level	s compared with glutathione.

542

543	FIG. 3	Reverse phase C-18 column semi-preparative HPLC profile of (a) 6H-CHH. 0.1% Trifluoroacetic acid
544		(TFA) in deionized water and 60% acetonitrile in 0.1% Trifluoroacetic acid (TFA) were chosen as
545		mobile phase A and B, respectively. CHH was filtered through 25 mm PES filters and 700 μ g (4 mL)
546		were injected at 1.0 mL/min of flow rate. The linear gradient was $0-25\%$ (v/v) mobile phase B over 15
547		min, then 25-100% (v/v) mobile phase B over 95 min. (b) The antioxidant activity of purified peaks
548		from 6H-CHH (78 μ g/mL) determined by the nitric oxide scavenging assay. Each bar displays the
549	//////	mean \pm SEM of four demonstrations (*** $P < 0.001$).

550

FIG. 4Mass spectrogram of anti-inflammatory peptides and antioxidant peptide from P16 (2H-CHH) and P57(6H-CHH), respectively, determined by LTQ orbitrap XL mass spectrometry. The following sequenceinterpretation was concluded: anti-inflammatory peptides were identified as (a) Ser-Ala-Phe-Asn-Pro-His-Glu-Lys-Gln (SAFNPHEKQ) and (b) Ile-Ile-His-Asn-Glu-Lys-Val-Gln-Ala-His-Gly-Lys-Lys-Val-Leu (IIHNEKVQAHGKKVL) and (c) the antioxidant peptide was identified as Lys-Ile-Tyr-Phe-Pro-His-Phe (KIYFPHF).

FIG. 5 The effect of CHHs on (a) NO production in LPS-activated macrophage RAW 264.7 cells and (b) the cytotoxicity (cell viability) of CHHs on macrophage RAW 264.7 cells determined by the MTT assay. CHHs (62-500 μ g/mL) were then incubated with macrophage RAW 264.7 cells and another overnight incubation was set between LPS and macrophage RAW 264.7 cells. The media were further used to measure the nitrite level (NO assay). Each bar displays the mean \pm SEM of four demonstrations (* P < 0.05, ** P < 0.01 and *** P < 0.001).

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FIG. 6 Reverse phase C-18 column semi-preparative HPLC profile of (a) 2H-CHH. 0.1% Trifluoroacetic acid 565 566 (TFA) in deionized water and 60% acetonitrile in 0.1% Trifluoroacetic acid (TFA) were chosen as 567 mobile phase A and B, respectively. CHH was filtered through 25 mm PES filters and 700 μ g (4 mL) 568 were injected at 1.0 mL/min of flow rate. The linear gradient was 0-25% (v/v) mobile phase B over 15 569 min, then 25-100% (v/v) mobile phase B over 95 min. (b) The anti-inflammatory property of purified 570 peaks from 2H-CHH (31 μ g/mL) against NO production determined in LPS-activated macrophage 571 RAW 264.7 cells. (c) The cytotoxicity (cell viability) of purified peaks from 2H-CHH (31 µg/mL) on 572 macrophage RAW 264.7 cells determined by the MTT assay. Each bar displays the mean \pm SEM of four 573 *demonstrations (* P < 0.05, ** P < 0.01 and *** P < 0.001).*

574

575**FIG. 7** (a) Nitric oxide scavenging activity of 6H-CHHs after intestinal digestion at concentrations of 15-125576 $\mu g/mL$. Each bar displays the mean \pm SEM of four demonstrations. (* P < 0.05 and *** P < 0.001)577probability levels compared with BHT. (b) Reducing power at concentration of 0.005-500 $\mu g/mL$ 578expressed in Trolox equivalents. Each bar displays the mean \pm SEM of four demonstrations. (***P <5790.001) probability levels compared with glutathione.

580

581FIG. 8The effect of 2H-CHHs after intestinal digestion on (a) NO production in LPS-activated macrophage582RAW 264.7 cells and (b) the cytotoxicity (cell viability) on macrophage RAW 264.7 cells determined by583the MTT assay. 2H-CHHs (31-125 µg/mL) were then incubated with macrophage RAW 264.7 cells and584another overnight incubation was set between LPS and macrophage RAW 264.7 cells. The media were585further used to measure the nitrite level (NO assay). Each bar displays the mean ± SEM of four586demonstrations (* P < 0.05, ** P < 0.01 and *** P < 0.001).</td>

588 FIG.9 The hemolytic activity of (a) 2H-CHH after intestinal digestion and (b) 6H-CHH after intestinal
589 digestion against human red blood cells. Each bar displays the mean ± SEM of four demonstrations
590 (*** P < 0.001).