

ฤทธิ์ด้านการอักเสบของเปปไทด์จากขนไก่ป่นในเซลล์เพาะเลี้ยงแมคโครฟาจ RAW 264.7



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จุฬาลงกรณ์มหาวิทยาลัย

CHULALONGKORN UNIVERSITY

บทคัดย่อและแฟ้มข้อมูลฉบับเต็มของวิทยานิพนธ์ตั้งแต่ปีการศึกษา 2554 ที่ให้บริการในคลังปัญญาจุฬาฯ (CUIR)

เป็นแฟ้มข้อมูลของนิสิตเจ้าของวิทยานิพนธ์ ที่ส่งผ่านทางบัณฑิตวิทยาลัย

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วิทยานิพนธ์นี้เป็นส่วนหนึ่งของการศึกษาตามหลักสูตรปริญญาวิทยาศาสตรมหาบัณฑิต

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ปีการศึกษา 2559

ลิขสิทธิ์ของจุฬาลงกรณ์มหาวิทยาลัย

ANTI-INFLAMMATORY ACTIVITY OF PEPTIDE FROM CHICKEN FEATHER
MEAL IN MACROPHAGE RAW 264.7

Miss Romteera Sukaboon



A Thesis Submitted in Partial Fulfillment of the Requirements
for the Degree of Master of Science Program in Biotechnology

Faculty of Science

Chulalongkorn University

Academic Year 2016

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Thesis Title ANTI-INFLAMMATORY ACTIVITY OF PEPTIDE
FROM CHICKEN FEATHER MEAL IN
MACROPHAGE RAW 264.7

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CHULALONGKORN UNIVERSITY

5772114823 : MAJOR BIOTECHNOLOGY

KEYWORDS: ANTI-INFLAMMATORY ACTIVITY / BIOACTIVE PEPTIDE / COX-2 / INOS

ROMTEERA SUKABOON: ANTI-INFLAMMATORY ACTIVITY OF PEPTIDE FROM CHICKEN FEATHER MEAL IN MACROPHAGE RAW 264.7. ADVISOR: ASSOC. PROF. APHICHART KARNCHANATAT, Ph.D., 69 pp.

Inflammation is a defense mechanism in the body induced by foreign chemicals or pathogens. Thus, inflammation is one of the most critical factors implicated in vasodilation, carcinogenesis and other degenerative disorders. However, the study of the anti-inflammatory mechanism of peptides has been limited. So studies on anti-inflammatory peptides from natural sources are increasing in interest. The objective of this study, to investigate the optimal condition of enzymatic hydrolysis to prepare an anti-inflammatory peptide from chicken feather meal by using 3 types of microbial protease (Alcalase[®], Flavourzyme[®], and Neutrase[®]) in various concentrations (1% 2.5% and 5%). Peptide hydrolysate with 1% Flavourzyme[®] produced the best results for inhibition of NO production (IC₅₀ 5.48±1.0 µg/ml) when compared with other proteases. These peptide fractions were fractionated by ultrafiltration with 10, 5, 3 and 0.65 kDa membranes and further analyzed for NO scavenging activity. Among the fractions, MW ≤ 0.65 kDa exhibited a high level of antioxidant activity toward the NO radical scavenging assay. The ≤ 0.65 kDa fractions were selected for further fractionation by gel filtration chromatography. The F2 fraction was the strongest in inhibiting nitric oxide production. After that, F2 fraction was selected to study the NO production and cytotoxicity test by MTT assay in LPS-stimulated macrophage RAW 264.7. The result shows that the peptide also inhibited NO production and wasn't toxic to macrophage RAW 264.7. The peptide fraction was further purified by HPLC and identification by mass spectrometry with amino acid sequence characterized into 4 peptides. In addition, the peptides are suppressed in the expression mRNA of iNOS, COX-2, IL-6 and TNF-α which are responsible for pro-inflammatory cytokine production. The results indicate that bioactive peptide from chicken feather meal can be developed to medical, pharmaceutical and cosmetic products.

Field of Study: Biotechnology

Academic Year: 2016

Student's Signature

Advisor's Signature

ACKNOWLEDGEMENTS

First of all, I would like to express my sincere gratitude and appreciation to my advisor Associate Professor Dr. Aphichart Karnchanatat for his precious advice, encouragement, guidance and supporting throughout the course of my thesis. His kindness and cheerfulness are also deeply appreciated.

Moreover, I also gratefully acknowledge the member of my thesis committee: Associate Professor Dr. Nattaya Ngamrojanavanich, Associate Professor Dr. Chanpen Chanchao and Dr. Chantragan Phiphobmongkol for discussion, advice and dedicating time for thesis examination.

I express special thanks to Assistant Professor Dr. Ornrapak Reamtong, Department of Molecular Tropical Medicine and Genetics, Faculty of Tropical Medicine, Mahidol University for her assistance in analysis and identification of amino acid sequence in this study.

I would like to thank Dr. Tanatorn Saisavoey and Ms. Papassara sangtanoo for their help and suggestion in analytical work in this research. I am thankful all members of room 704 and 705 for their help and kind friendship especially Ms. Piroonporn Srimongkol for helpful advice and assistance in RNA extraction and PCR method.

The author thanks Program in Biotechnology, Faculty of Science, and Institute of Biotechnology and Genetic Engineering, Chulalongkorn University for facilities and all supporting.

Finally, I would like to express the highest gratitude to my parents, everyone in my family for everything in my life, their understanding, encouragement and support for a long course of my education.

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LIST OF ABBREVIATIONS

| | |
|------------------|--------------------------------------------------------------|
| Abs | Absorbance |
| bp | Base pair |
| cDNA | Complementary DNA |
| CO ₂ | Carbon dioxide |
| DMSO | Dimethylsulfoxide |
| DMEM | Dulbecco's Modified Eagle Medium |
| eNOS | Endothelial nitric oxide synthase |
| <i>et al</i> | and others |
| FBS | Fetal bovine serum |
| h | Hour |
| IC ₅₀ | Median inhibitory concentration, 50% maximum inhibition |
| IL-6 | Interleukin-6 |
| iNOS | Inducible nitric oxide synthase |
| kDa | Kilodalton |
| LPS | Lipopolysaccharide |
| mg | Milligram |
| min | Minutes |
| ml | Milliliter |
| mM | Millimolar |
| MTT | 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide |
| NFκB | Nuclear factor-kappa B |
| ng | Nanogram |

| | |
|-----------------------------|------------------------------|
| nm | Nanometer |
| nNOS | Neuron nitric oxide synthase |
| NO· | Nitric oxide |
| O ₂ ⁻ | Superoxide anion |
| PBS | Phosphate buffer saline |
| PCR | Polymerase chain reaction |
| PGE ₂ | Prostaglandin E ₂ |
| psi | Pound per square inch |
| RNA | Ribonucleic acid |
| RNS | Reactive nitrogen species |
| ROS | Reactive oxygen species |
| rpm | Round per minute |
| RT | Reverse transcription |
| s | second |
| TAE | Tris-acetate-EDTA |
| TE | Tris-EDTA |
| TNF- α | Tumor necrosis factor alpha |
| UV | Ultraviolet |
| α | Alpha |
| β | Beta |
| °C | Degree Celsius |
| γ | Gamma |
| μ g | Microgram |
| μ l | Microliter |
| μ M | Micromolar |
| / | Per |

% Percentage

: Ratio



CHAPTER I

INTRODUCTION

Inflammation is a host defense mechanisms which involves physiological and pathological process within an organism and which is induce by the invasion of pathogens or tissue injury caused by biological, chemical or physical damage (Je J. Y. and S. K. Kim, 2012). The activation of several immune cells (monocytes and macrophages) produces inflammation mediators such as nitric oxide (NO), cyclooxygenase-2 (COX-2), prostaglandins E₂ (PGE₂) and other pro-inflammatory cytokines including TNF- α , IL-6 and IL-1 β (Berenbaum, 2000; Fujiwara and Kobayashi, 2005; Je J.-Y. and S.-K. Kim, 2012; Lawrence *et al.*, 2002). The macrophage is a major participant which play an important role in the inflammation mechanism and which is initiated by bacterial lipopolysaccharide (LPS) that is thought to originate from outer membrane of gram-negative bacteria, IFN- γ and pro-inflammatory cytokines (Guha and Mackman, 2001). In addition, oxidative stress is involved in the pathogenesis of several health problems. The physical and biochemical process in human body may generate free radical as by-products which can help trigger a wide range of diseases (Reddy and Reddanna, 2009). The interaction between reactive nitrogen species (RNS) such as nitric oxide and reactive oxygen species (ROS) interaction toxic agents and is strongly linked with inflammation and cancer (Lee *et al.*, 2013). However, excessive of inflammatory mediators contribute to acute and chronic disease characterized by the uncontrolled production of pro-inflammatory cytokines such as cancer, coronary heart disease, rheumatoid arthritis, asthma and Alzheimer's (Dia *et al.*, 2009). During the inflammation process, inflammatory mediators such as NO and PGE₂ are produced via the oxidation of L-arginine by inducible nitric oxide synthase (iNOS) and the conversion of arachidonic acid by cyclooxygenase (COX-2). Moreover, NO is a key signaling biological molecule involved in the vasodilation and regulation of blood pressure neurotransmission and host immune defense (de Mejia and Dia, 2009). The production of pro-inflammatory cytokine and nitric oxide are closely correlated

to inflammation. Inflammation can be regulated by suppressing pro-inflammatory cytokine and nitric oxide production (Lee S.-J. *et al.*, 2012).

Protein hydrolysate may be derived from a purified protein source through heating with chemical or proteolytic enzyme into peptides. This study focused on enzymatic hydrolysis to prepare protein hydrolysate from chicken feather meal. The production of protein hydrolysate is a sequential action between endopeptidase and exopeptidase (Clemente, 2000; Manninen, 2009). The benefits of enzymatic hydrolysis include that it is not consumed during reaction, it has a high conversion yield, high selectivity, high specificity, soft condition, is nontoxic, and can be produced a large-scale production (Kristinsson and Rasco, 2000). There are several factors that influence in enzymatic hydrolysate such as pH, incubation time, temperature, enzyme and substrate (Salwanee *et al.*, 2013). In general, peptides consist of a complex mixture of free amino acid around 3-20 residues in each sequence and bioactivity is characteristic of the amino acid composition and sequence (Pihlanto-Leppälä, 2000). Moreover, the sequences of amino acid within peptide of their native proteins usually take on an inactive bioactive role while specific enzymatic hydrolysis under controlled condition can be enhance the potential bioactivity of peptide (Sarmadi and Ismail, 2010).

Currently, natural compounds with biological activity are increasing. The requirement of functional food or dietary supplements has increased significantly in the health improvements of the human body (Cian *et al.*, 2012). Peptides are known for being highly selective with excellent safety and efficacy among humans (Fosgerau and Hoffmann, 2015). Recently, there has been much pharmaceutical research addressing the utilization and focus on peptides from plants or animals due to their different health benefits and biological activities as sources of bioactive peptides for drug development (Vo *et al.*, 2013). Bioactive peptide has been found to possess anticancer, antimicrobial, hypocholesterolemic, antihyperterolemic, antihypertensive, antithrombotic and anti-inflammatory properties (Ahn *et al.*, 2012).

Poultry processing plants and the related industry is expanding in many countries, especially in Thailand. Arising from this, chicken feather meal is a major waste by-product and produced in large amounts annually (Gang *et al.*, 2013). Chicken feather meal represents around 5-7% of chicken body weight. However, the substantial accumulation of chicken feather meal may cause environmental problems and pollutions (Nomura *et al.*, 2005). Chicken feather meal is also an alternative value-added product used as animal feed, organic fertilizer and feed supplement due to its content of approximately 90% protein, as well as its disulphide bonds and cross-linkages (Tiwary and Gupta, 2013). The structure is mainly composed of pure keratin and important amino acids like cysteine arginine glycine and phenylalanine (Kumar *et al.*, 2012). Therefore, the objective of this study was to prepare peptides from chicken feather meal by enzymatic hydrolysis using microbial protease and to determine anti-inflammatory activity in the macrophage MACROPHAGE RAW 264.7 . Therefore, the bioactive peptide from chicken feather meal can also be further formulated for peptide drug in the pharmaceutical industry or as ingredient in cosmetic products in the global market.



CHAPTER II

LITURATURE REVIEW

2.1 Free radicals

Free radicals are any chemical species such as atom or molecule which contain one or several unpaired electron in its outer shell. This unpaired electron also pulse radicals unstable and highly reactive. The radicals can react with both of radical and non-radical molecules to create chain reaction by donate its unpaired electron to a non-radical molecule or seize an electron from another molecule in order to form a paired electron and more damaging species (Lopez-Torres *et al.*, 1991). There are several types of free radicals in living systems. Mostly, radicals are mainly derived from oxygen (reactive oxygen species/ROs) such as superoxide anion $O_2^{\cdot-}$ and hydroxyl radicals $OH\cdot$. Moreover, they are also derived from nitrogen (reactive nitrogen species/RNSs) such as nitric oxide $NO\cdot$, nitrogen dioxide $NO_2\cdot$ and nitrate radical $NO_3\cdot$ (Table 2.1) (Halliwell, 2001). Free radicals are very short lived, with half-life in milli-, micro-, or nanoseconds, extremely unstable. The presences of free radicals are lead to overpower the radicals scavenging mechanism, oxidative stress and available to react with organic substrate including lipid, DNA and proteins to generate harmful products such as lipid peroxidation and other lipid adducts (Dhawan, 2014). The resulting in protein damage may cause loss of enzyme activity, while DNA damage can lead to mutagenesis and carcinogenesis (Davalos *et al.*, 2004; Wiseman and Halliwell, 1996). There are numerous studies suggested that free radicals are involved in the several human diseases, as well as in ageing, cancer and inflammation. Both of ROS and RNS also generate from either endogenous or exogenous source. For endogenous source, free radicals are generated from immune cell activation, inflammation, mental stress and infection. In contrast, exogenous of free radical mostly from air, water, pollution, cigarette smoke, alcohol and transition metals (Pham-Huy *et al.*, 2008).

Table 2.1 Different type of reactive oxygen species (ROS) and reactive nitrogen species (RNS) produced in cell (Dhawan, 2014)

| Reactive species | Free radicals | Formula |
|---------------------------------|------------------|----------------|
| Reactive oxygen species (ROS) | Superoxide | $O_2^{\cdot-}$ |
| | Hydroxyl | OH^{\cdot} |
| | Peroxyl | RO_2^{\cdot} |
| | Alkoxyl | RO^{\cdot} |
| | Hydroperoxyl | HO_2^{\cdot} |
| Reactive nitrogen species (RNS) | Nitric oxide | NO^{\cdot} |
| | Nitrogen dioxide | NO_2^{\cdot} |

However, the link between free radical and inflammation are investigated. One of the chronic inflammatory diseases which involved in oxidative stress is rheumatoid arthritis. The pathological process of this disease is caused from reproduction of ROS and RNS in human body (Kavazis and Powers, 2013).

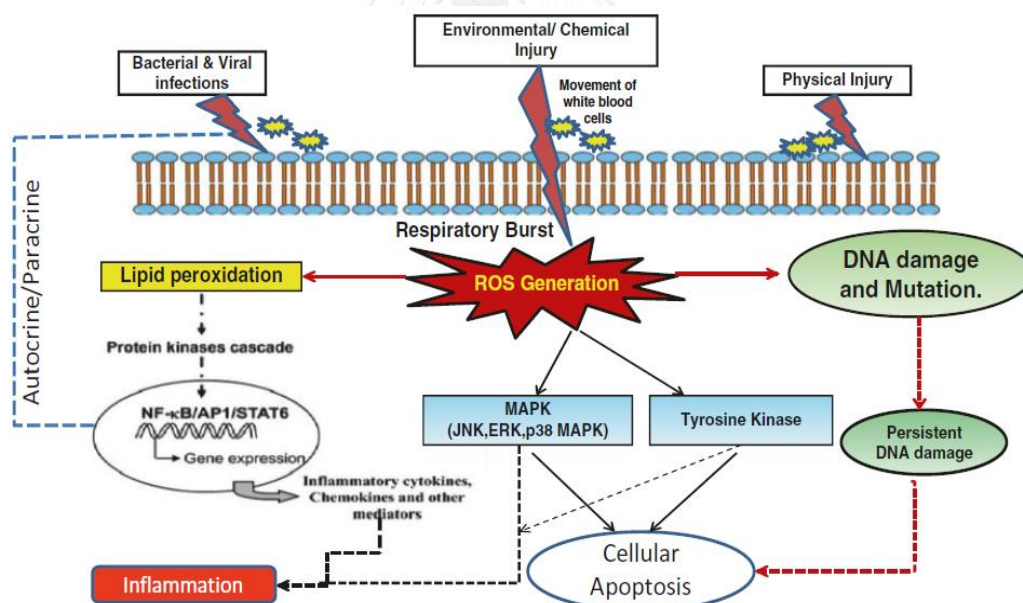


Figure 2.1 Role of ROS and RNS in tissue damage (Dhawan, 2014)

2.2 Inflammation

Inflammation is the complex process of the protective response of the body which is stimulated by microbial infection, chemical and physical damage (Abarikwu, 2014). The causes of inflammation are many and varied but most the common is that from the immune response to an infectious microorganism (Hoebe *et al.*, 2004). Inflammation can lead to several cardinal sign of pain, heat, redness, swelling and loss of function. The purpose of inflammation response is to eliminate the pathogenic insult and remove the injured tissue components. Excessive inflammation processes also lead to various diseases such as cancer, rheumatoid arthritis, atherosclerosis and asthma (Wang A.-Y. *et al.*, 2011). The inflammation mechanism can be classified into acute and chronic inflammation.

Acute inflammation is the immediate and early response to injury designed to deliver leukocytes to the sites of injury. Thus, invading microbes have been removed and necrotic tissues also broken down. During this process, the following events will occur:

2.2.1. Vascular change: revision in blood vessels resulting in increased blood flow (vasodilation) and structural change that plasma and protein leak (increased vascular permeability).

2.2.2 Cellular event: migration of the leukocyte from the microcirculation and accumulation in the focus of injury (cellular recruitment and activation).

Chronic inflammation is immortalized and prolongs the duration (weeks, months and years) in which the active inflammation tissue injury and healing proceed simultaneously. The causes of chronic inflammation are mostly from viral infection, persistent microbial infection, exposure to potentially toxic or chemical agents and autoimmune diseases. There is also tissue destruction which is directed by the inflammatory cell. Chronic inflammation also occurs following infiltration of the mononuclear cell (macrophage, lymphocytes and plasma cells), new vessel proliferation (angiogenesis) and fibrosis (Yu H. *et al.*, 2009).

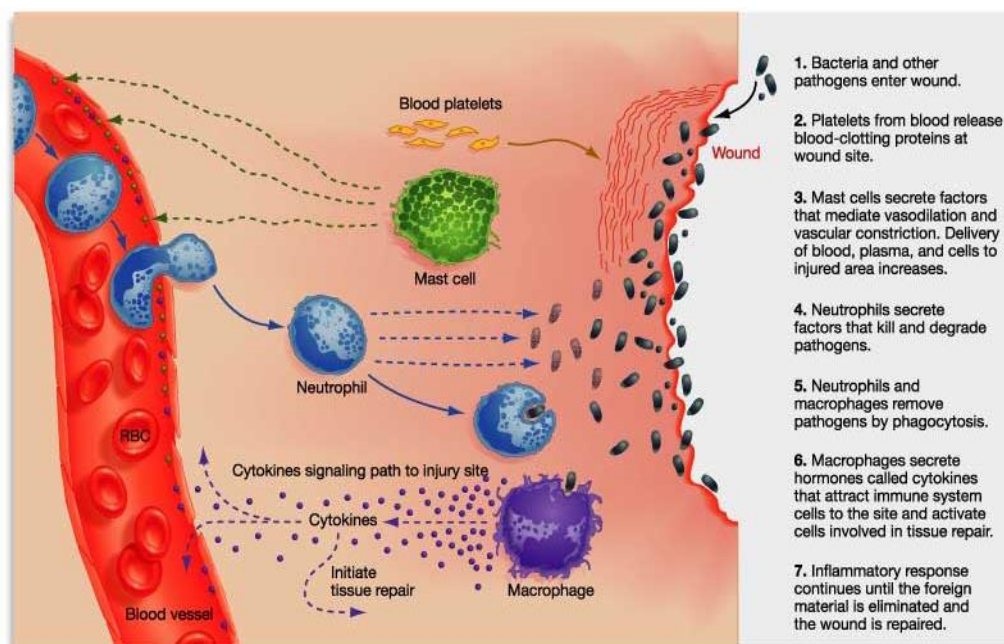
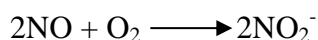


Figure 2.2 Step of inflammatory response against pathogens

2.3 Biological significance of nitric oxide

Nitric oxide is catalyzed by enzyme as nitric oxide synthase (NOS) which are present in the body. The synthesis of NO oxide is generated by conversion of L-arginine to L-citrulline in a highly control process. There are two cofactors have been found to be involved in this process which includes oxygen and NADPH (Omer *et al.*, 2012). In addition, there three isoforms of NOS have been identified as names which basis of their activities, following by neuronal NOS (nNOS), inducible NOS (iNOS) and endothelial NOS (eNOS)(Table 2.2) (Nathan and Xie, 1994). There are many chemical reactions which occur under biological conditions. The chemical reaction involving NO production is described below (Blough and Zafiriou, 1985):

1. Reactivity with oxygen to yield NO_2 gas or NO_2^- in solution:



2. Reactivity with superoxide anion (O_2^-) to yield the unstable intermediate peroxonitrite anion ($^-\text{OONO}$), which rearranges to form NO_3^-

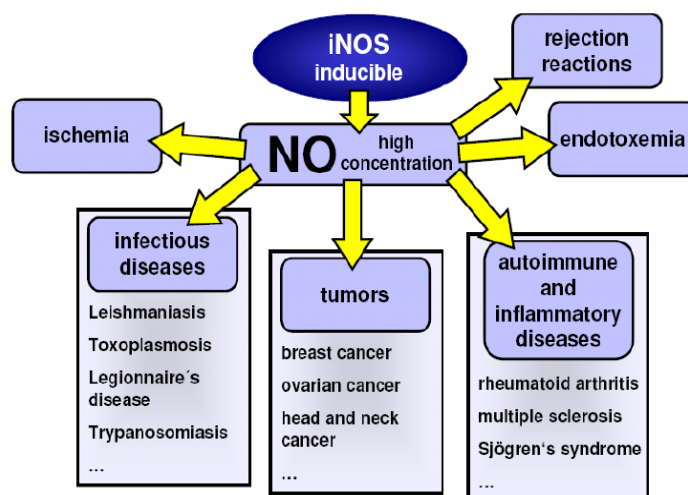
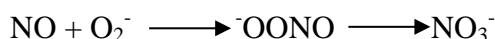


Figure 2.3 Assumption of iNOS to produce NO in various human diseases (Hemmrich *et al.*, 2003).

NO is diatomic free radical and chemically unstable, with extremely short life around 3-5 s in aqueous solution under physiological conditions of concentration, temperature, pH and oxygen (Bryan and Grisham, 2007). NO is one of the chemical mediators in inflammation of the body which significant for its role in regulating vascular tone and in signaling neurotransmission (Moncada *et al.*, 1991). The high level production of nitric oxide is dependent on induction and expression of the inducible nitric oxide synthase (iNOS) expressed in macrophages. However, overproduction of nitric oxide may lead to circulatory shock, chronic inflammation and carcinogenesis (Fang, 1997). NO also has been implicated in different mechanisms of diseases such as atherosclerosis, asthma, neurologic disorder and septic shock (Barton *et al.*, 1998). The isoform of nitric oxide synthase and their physical functions are summarized in Figure 2.4.

Table 2.2 Classification of mammalian nitric oxide synthase

| Isoform ^a | Typical localization | Cellular localization | Monomer molecular mass (kDa) |
|----------------------|----------------------|-------------------------|------------------------------|
| nNOS | Neuron | Soluble and particulate | -160 |
| eNOS | Endothelial cells | Particulate | -135 |
| iNOS | Macrophage | Soluble | -130 |

^anNOS, neuronal nitric oxide synthase; eNOS, endothelial NOS; iNOS, inducible NOS

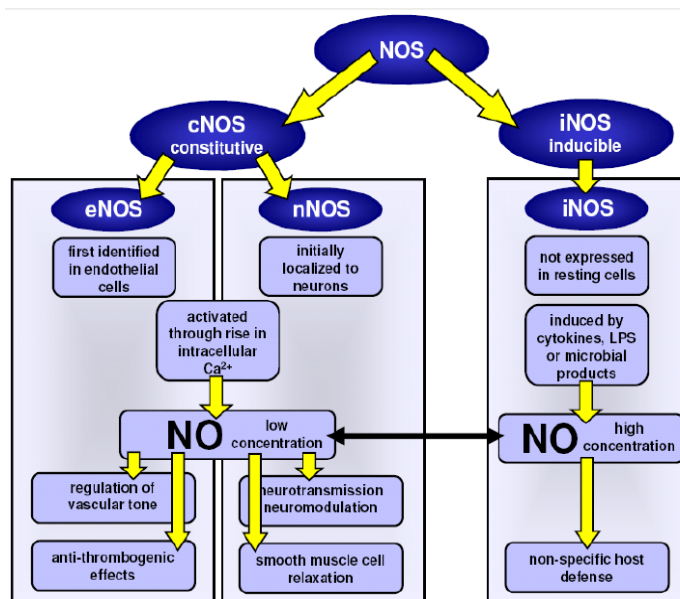


Figure 2.4 The isoform of nitric oxide synthase. Two cNOS enzyme (eNOS, nNOS) are contrasted by third, inducible NOS (iNOS) (Hemmerich *et al.*, 2003).

2.4 Role of nitric oxide physiological function and inflammation

All cells produce both reactive oxygen and nitrogen species (ROS/RNS) such as superoxide anion (O_2^-), hydrogen peroxide (H_2O_2), NO and peroxynitrite ($ONOO^-$) during normal respiration or activated macrophages. Moreover, ROS/RNS are primary members of cellular signaling pathways, particularly those leading to the inducible activation of nuclear factor kappa B (NF- κ B). NF- κ B is a transcription factor associated with the expression of various inflammatory genes including COX-2 and iNOS. Nitric oxide (NO) is a free radical involved in inflammation that is classified as a reactive nitrogen species. The role of nitric oxide is that of also regulating a variety of diseases and physiological functions of the body including blood pressure, neural signal transmission, and platelet function as well as playing a critical role in the anti-pathogen and tumoricidal response of the immune system (Liu *et al.*, 2008). However, during the inflammation process, in vivo formation of NO is a normal physiological function in the infection of tissues. Increased NO production levels may cause some adverse effects of NO such as the formation of the carcinogenic N-nitroso compound, deamination of DNA bases and mutagenesis, oxidation of DNA bases and tissue injury caused by potent oxidative agents (Liu *et al.*, 2008). Thus, production of NO via iNOS expression is also implicated as a deleterious agent in various pathophysiological conditions and tissue damage (Moncada *et al.*, 1991). Figure 2.9 shows the physiological and pathological roles of NO.

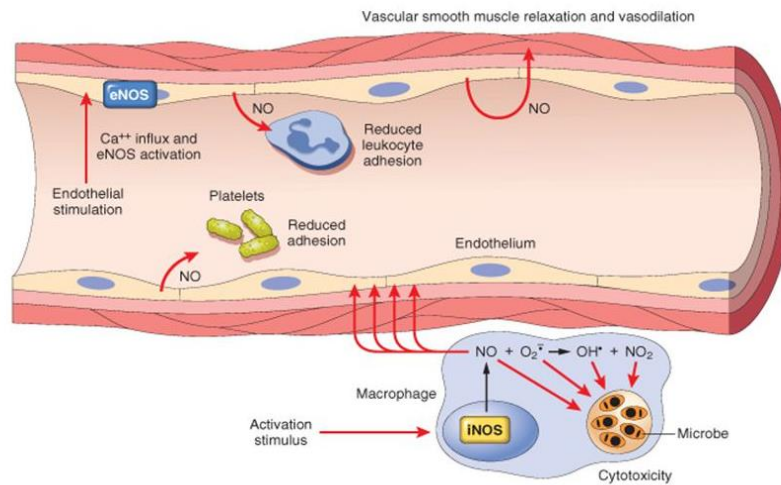


Figure 2.5 Function of nitric oxide (NO) in blood vessel and macrophage, produced by two NO synthase enzyme (eNOS and iNOS). (Kumar *et al.*, 2012)

2.5 Role of cytokines in inflammation process

Inflammation is mediated by a variety of soluble factors such as cytokines. Cytokines play a key role in cellular infiltrate, the state of cellular activation and the systemic response to inflammation. In addition, cytokines are associated with the extensive networks that correlate to synergistic as well as antagonistic interactions. There are several cytokines involved in the inflammatory process such as tumor necrosis factor-alpha ($\text{TNF-}\alpha$) and interleukin-6 (IL-6).

2.5.1 Tumor necrosis factor alpha (TNF- α)

Tumor necrosis factor (also known as TNF- α) is a polypeptide cytokine related to systemic inflammation and is a member group of cytokines that stimulate the acute phase of inflammation reaction. TNF- α has been implicated in common biological properties and a diverse range of inflammatory infectious. The importance of TNF- α in inflammation is controlling disease activity in rheumatoid arthritis and other inflammatory conditions. TNF- α also induces apoptotic cell death, inhibits tumorigenesis and viral replication. Overproduction of TNF- α has been implicated in human diseases such as cancer (Sprague and Khalil, 2009).

TNF- α is one of the products produced by activated monocyte/macrophage lineage, with T lymphocytes, neutrophils, mast cells and endothelium. Regulation of the transcription factor NF- κ B is a key component of TNF- α signal transduction. Physical stimuli such as ultraviolet light, X-radiation and heat are also responsible for TNF- α production and release. Normally, TNF- α is released from preformed storage by cleavage of the membrane TNF- α on the macrophage, neutrophils and activated T cells by a TNF- α converting enzyme (TACE/ADAM17) and the release of cytoplasmic granules from a mast cell and eosinophils (Feldmann and Maini, 2001). Large amounts of TNF- α are released in response to lipopolysaccharides and other bacterial products. Moreover, TNF- α can be induced by the expression of COX-2 which leads to vasodilation and thus increased blood flow.

2.5.2 Interleukin-6 (IL-6)

Interleukin-6 (IL-6) is produced by important immune cells macrophages and monocytes at the inflammation site. IL-6 is a cytokine with redundancy and pleiotropic activity that contributes to the host defense against acute environmental stress. Moreover, the down regulation of IL-6 production leads to a pathological role in various auto immune and chronic inflammatory diseases. When acting on hepatocytes, IL-6 strongly induces broad spectrum acute phase proteins such as C-reactive proteins (CRP) in addition to reducing albumin, cytochrome P450, fibronectin and transferrin (Figure 2.6) CRP is the best biomarker of inflammation and its expression mainly depends on IL-6. IL-6 also promotes the differentiation of

osteoclasts and angiogenesis, the proliferation of keratinocytes and mesangial cells, and the growth of myeloma and plasmacytoma cells (Tanaka *et al.*, 2014)

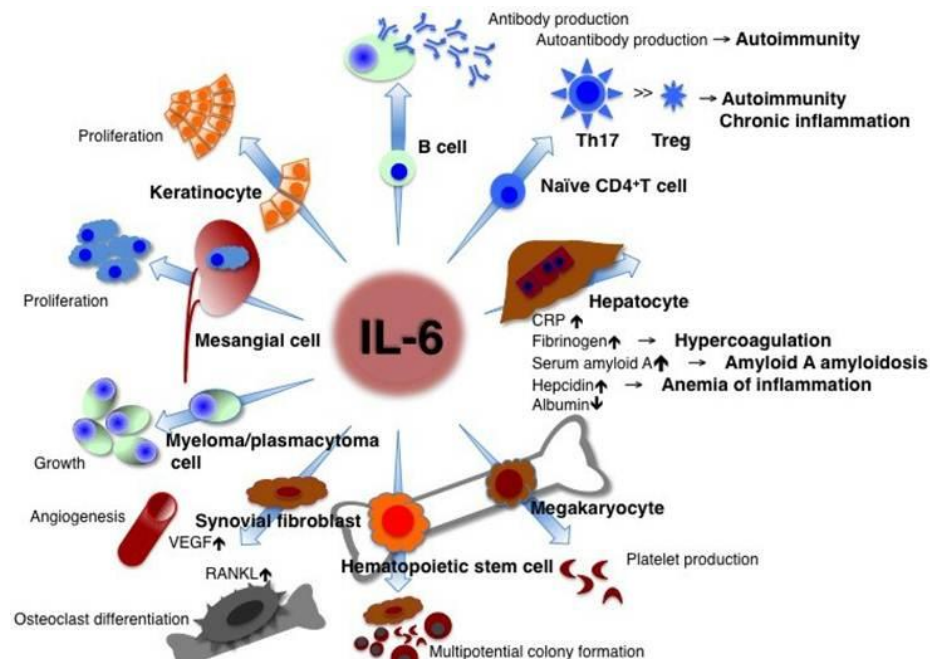


Figure 2.6 IL-6 has a pleiotropic effect which dysregulates persistent production and causes the onset and development of various autoimmune and chronic inflammatory diseases (Tanaka *et al.*, 2014).

2.6 Prostaglandins

Prostaglandin is a product of the COX-2 pathway, generating inflammation by increasing vascular permeability, vasodilation and release of proinflammatory cytokines to the site of injury (Issekutz and Movat, 1982). Moreover, the level of prostaglandins is connected to the setup of chronic inflammation and cancer. Cyclooxygenase production is the first step in the formation of prostaglandins and thromboxane from free arachidonic acid (Kim G.-H., 2008). There are two isoforms: COX-1 and COX-2. COX-1 also constitutively expressed in all cell types and involved in normal kidney, gastrointestinal and reproductive functions (Smith and Langenbach, 2001). COX-2 may be expressed in some tissue, but is largely inducible by a wide variety of mitogens, hormones, cytokines and other stimuli. COX-2 is also

associated with inflammation and disease. In addition, macrophages secrete prostaglandins following stimulation with LPS, primarily due to the induced transcription of the COX-2 gene and production of COX-2 enzyme (Ishimura *et al.*, 2004). The gene encoding for the expression of COX-2 protein is controlled by NF-KB, which is a transcription factor influenced by oxidant status.

2.6.1 Cyclooxygenase-2 (COX-2)

Cyclooxygenase (COX) or prostaglandin H₂ synthase (PGHS) is a key enzyme in the synthesis of prostaglandins (PGs) from arachidonic acid (AA) (Abarikwu, 2015). However, COX-2 expression is basically absent in normal cell function. In contrast, COX-2 expression is widely induced by inflammatory stimuli in cells such as macrophages, endothelial cells and synoviocytes. Induction of COX-2 by stimuli is also associated with cell activation and inflammation disease. COX-2 plays an important role in angiogenesis, colon cancer and Alzheimer's disease. Investigating the mechanism of the COX-2 expression could lead to the discovery of a new drug for reducing inflammation without removing the protective PGs in the stomach and kidneys. Therefore, COX-2 inhibitors also have great and innovative potential in reducing inflammation.

2.7 Macrophage

One major type of macrophages is the mononuclear phagocyte of the immune system which are important cells that originate from bone marrow including blood monocytes and tissue macrophage. Macrophages play a critical role in the initiation, maintenance and resolution of inflammation. During the inflammation process and infection, macrophages act as phagocytic cells capable of targeting and ingesting invading organisms activated by lymphokines and different bacterial products. Lipopolysaccharides (LPS) are the residues of the outer membrane of gram-negative bacteria, which are the most effective activators of monocytes. LPS can activate several signal transduction pathways through the induction of G-proteins, cAMP dependent kinase, protein kinase C and JNK mitogen activated kinase (MAPK) families (Furukawa *et al.*, 1988).

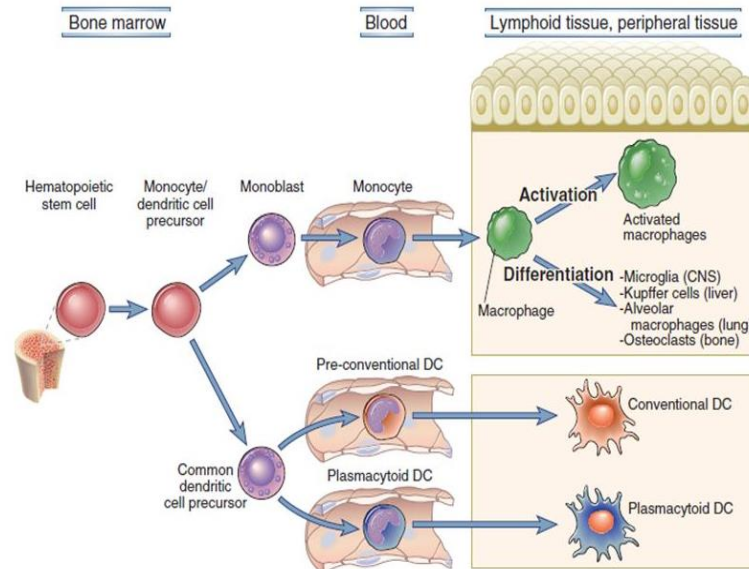


Figure 2.7 Maturation of mononuclear phagocyte develop from bone marrow (Abbas *et al.*, 2011).

Macrophages also produce a wide range of biologically active molecules that participate in the inflammatory process. Their function in inflammatory process is one of innate and adaptive immunity, playing a key role in the host defense mechanism. In innate immunity, the macrophage induces chemical mediators by releasing cytokine, which then leads to inflammation (Condeelis and Pollard, 2006).

2.8 Nitric oxide radical scavenging activity assay

Nitric oxide is one of the free radicals classified as reactive nitrogen species (RNS). Nitric oxide (NO) is produced from the conversion of amino acid L-arginine by endothelial cells, macrophages and neurons (Parul *et al.*, 2013). NO is an important signaling molecule in vasodilation, neurotransmission, and the host immune defense (Pacher *et al.*, 2007). There are three isoforms of the nitric oxide synthase (NOS): endothelial nitric oxide synthase (eNOS), neuronal nitric oxide synthase (nNOS) and inducible nitric oxide synthase (iNOS). These are constitutively expressed and essential for maintaining tissue homeostasis (Jung *et al.*, 2007). iNOS is influenced in response to pro-inflammatory cytokine production including interferon- γ (IFN- γ), tumor necrosis factor- α (TNF- α) and interleukin-6 (IL-6), and

several inflammatory responses (Ndiaye *et al.*, 2012). Determination of NO production can be carried out by various experimental methods. Mostly, in vitro nitric oxide radical scavenging activity is measured spectrophotometrically. When sodium nitroprusside is mixed with aqueous solution at physiological pH (7.2), nitric oxide (NO) is generated under aerobic conditions to produce nitrite and nitrate ion that can be estimated by the Griess reagent. The absorbance of the chromophore formed during diazotization of the nitrite with sulfanilamide and subsequent coupling with naphthylethylenediamine dihydrochloride is immediately measured at 540 nm (Menaga *et al.*, 2013).

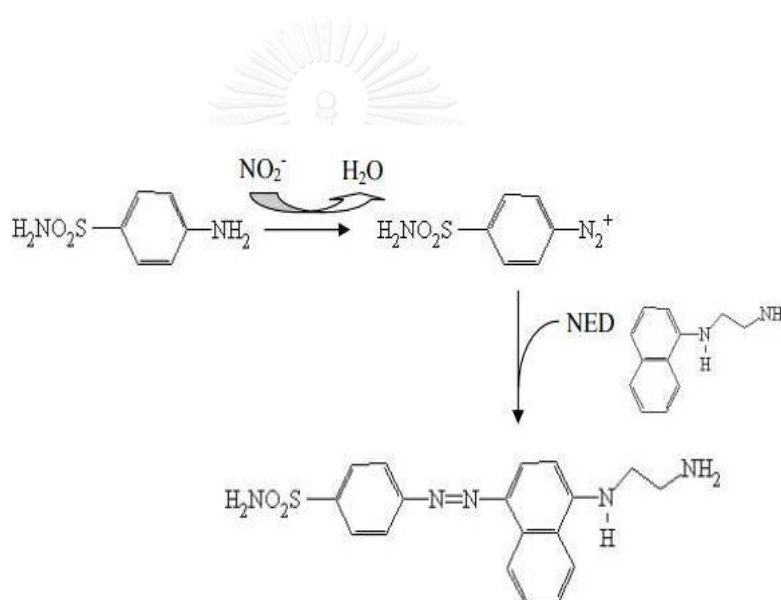


Figure 2.8 Chemical reactions involved in the determination of NO₂ using the Griess Reagent System.

2.9 Cytotoxic activity

Cytotoxic activity is an in vitro cytotoxicity test of the chemicals or unknown compounds affecting the normal functions of cells which are common to all cells. The toxicity of the compound can be evaluated by assessing the cellular damage. The normal experiment for the evaluation of cell death is that of MTT cytotoxicity assay (Figure 2.9). Other methods also widely use sensitive, quantitative and reliable colorimetric assays. MTT cytotoxicity assay is the measure of the viability of the proliferation and activation of cells (Senthilraja and Kathiresan, 2015). The assay is based on the capacity of the mitochondria dehydrogenase enzyme in the living cells to convert the yellow water-soluble substrate 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) into a dark blue formazan product which is insoluble in water. Living cells can reduce the yellow MTT under tetrazolium ring cleavage to a water-insoluble purple blue formation which precipitates in the cellular cytosol and can be dissolved after cell lysis, whereas cells that have died following toxic damage cannot transform MTT. This formation production is proportional to the viable cell number and inversely proportional to the degree of cytotoxicity. The reaction is mediated by the dehydrogenase enzyme involved with the endoplasmic reticulum and the mitochondria (Fotakis and Timbrell, 2006).

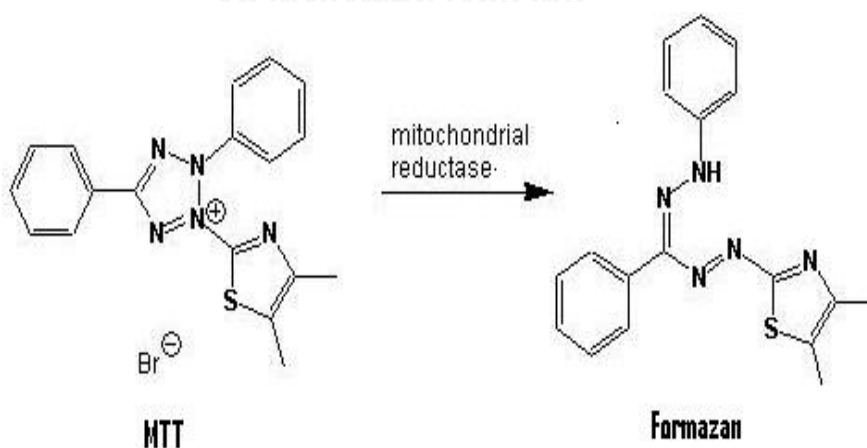


Figure 2.9 MTT assay (3-(4, 5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide a yellow tetrazole), is reduced to purple formazan in living cells

2.10 Bioactive peptide

Currently, there is increasing utilization of natural proteins as sources of biologically active peptides with diverse health benefits and these peptide can be generated by proteolytic enzyme treatment under specific conditions. Peptides are short polymers of amino acids linked by peptide bonds. Protein is the essential components of tissues in biological organisms and participates in a large number of physiological processes within the cells. In general, bioactive peptides also contain around 3-20 amino acid units (Pihlanto-Leppälä, 2000). Biologically active peptides, once liberated as dependent entities, act as potential metabolism modulators and control compounds with hormone-like activities. However, the bioactivities that have been reported for peptide include antihypertensive, antioxidant, anticancer, antimicrobial and opioid activities as well as immunomodulatory and cholesterol-lowering effects. In addition, peptides with anti-inflammatory effects have been less investigated (Montoya-Rodríguez *et al.*, 2014). The activity of bioactive peptides is dependent on their amino acid composition and sequence.

2.10.1 Anti-inflammatory peptide

The anti-inflammatory effect of peptides has been identified in a broad variety of natural sources. Tripeptide from salmon pectoral fin produced with low molecular weight, an anti-inflammatory peptide prepared from pepsin hydrolysate, has shown anti-inflammatory activity by suppressing the protein expression of inducible nitric oxide synthase, cyclooxygenase-2 and production of cytokines including tumor necrosis factor- α , interleukin-6 and -1 β (Ahn *et al.*, 2012). Another anti-inflammatory peptide, protein hydrolysate from yellow field pea seeds was reported to significantly inhibit nitric oxide production by activating macrophages and inhibiting the secretion of pro-inflammatory cytokines, TNF- α and IL-6 respectively (Ndiaye *et al.*, 2012). The *Mytilus edulis* hydrolysates fraction was able to inhibit LPS-stimulated inducible nitric oxide synthase (iNOS) and cyclooxygenase-2 (COX-2) protein and gene expression. These peptide fractions also inhibited the translocation of NF-KB (nuclear factor-kappa B) through the prevention of I κ B α (inhibitory factor kappa B alpha) phosphorylation (Kim Y.-S. *et al.*, 2016).

Table 2.3 Anti-inflammatory peptide from natural source

| Source of bioactive peptide | Enzymatic hydrolysis | Result | Amino acid sequence | Reference |
|-----------------------------------------------------|-------------------------------------------------------------------|---------------------------------------------------------------------------|---------------------------------------------------------|-----------------------------------------|
| <i>Pisum sativum</i> L. (Yellow field pea seeds) | Thermolysin | Inhibited NO production and secretion of pro-inflammatory cytokines | - | (Ndiaye <i>et al.</i> , 2012) |
| <i>Salmo salar</i> (Salmon byproduct) | Pepsin | Inhibited nitric oxide production by 63.08% and Prostaglandins by 45.33%. | Pro-Ala- Try | (Ahn <i>et al.</i> , 2012) |
| <i>Mytilus coruscus</i> (Korean mussel) | Alcalase, Flavpurzyme, Neutrase, Protamex, and Papain | Flavourzyme hydrolysate inhibited nitric oxide production on macrophage | Gly-Val- Ser-Leu- Leu-Gln- Gln-Phe- Phe-Leu | (Kim E.- K. <i>et al.</i> , 2013) |

Table 2.3 Anti-inflammatory peptide from natural source (continue)

| Source of bioactive peptide | Enzymatic hydrolysis | Result | Amino acid sequence | Reference |
|--------------------------------------------------------|-----------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------|-----------------------------------------------------------------|----------------------------------|
| <i>Ruditapes philippinarum</i> (Japanese carpet shell) | Alcalase, Flavpurzyme, Neutrase, Protamex, Papain, Pepsin, Trypsin and α -chymotrypsin | Alcalase hydrolysate inhibited LPS induced-NO production in RAW264.7 | Gln-Cys- Gln-Gln- Ala-Val- Gln-Ser- Ala-Val | (Kim E.-K. <i>et al.</i> , 2013) |
| Tuna cooking juice | Orientase, flavourzyme and alcalase | Alcalase hydrolysate inhibited nitric oxide production on murine macrophage RAW264.7 | Pro-Arg- Arg-Thr- Arg-Met- Met-Asn- Gly-Gly- Arg | (Cheng <i>et al.</i> , 2015) |

2.10.2 Enzymatic hydrolysis

Enzymatic hydrolysis is an alternative method to the conversion and utilization of biomass into bioactive protein. Hydrolysis of the protein involves the digestion of peptide bonds to generate peptides of various sizes and amino acid composition. There are three types of hydrolysis: enzymatic, acid and alkali hydrolysis. Chemical hydrolysis is a complex process to regulate and reduce the functional quality of products (Radha *et al.*, 2008). Enzymatic hydrolysis is used to produce protein hydrolysate without disruption of amino acid without more severe treatment such as chemical and physical treatment (Clemente, 2000). However, much research has shown that pH and temperature also affect enzymatic hydrolysis. The products obtained from enzymatic hydrolysis treatment also have high functionality and nutritive values. Production of protein hydrolysate by use of digestive proteolytic enzymes from animals or from plants and microorganisms have been reported safe for human consumption (McCarthy *et al.*, 2013).

Table 2.4 Comparison of the chemical hydrolysis and enzymatic hydrolysis to produce protein hydrolysate (Kristinsson and Rasco, 2000; Sanmartín *et al.*, 2009).

| Processing method | Advantage | Disadvantage |
|------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Chemical hydrolysis (acid and alkali) | <ul style="list-style-type: none"> -High protein recovery -Short processing time -Low processing cost | <ul style="list-style-type: none"> - Absence of homogeneous hydrolysates -Disturb functionality - Metal corrosion of equipment - Reaction is hard to control -Form toxic substance |
| Enzymatic hydrolysis (protease) | <ul style="list-style-type: none"> -Maintain functions and nutritive value of final hydrolysates -Produce homogeneous hydrolysate -Mild condition -Nontoxic and biodegradable | <ul style="list-style-type: none"> - High cost processing - Long processing time |

2.10.3 Proteolytic enzyme

Proteolytic enzymes (Proteinases, peptidase or protease) are enzymes involved in proteolysis. This process is the digestion of protein into small polypeptides or free amino acids by reacting with peptide bonds. Proteolytic enzymes can be split into endopeptidase and exopeptidase. Exoprotease hydrolyzes the peptide bonds at the terminal peptide bond and releases amino acids from either the N or C terminal side. Endopeptidase breaks down the peptide bond with protein molecules at a random side (Zhuang *et al.*, 2009). Protease has been used in the production of diverse bioactive peptides from native protein. (Sun *et al.*, 2016) reported that egg ovomucin hydrolyzed by alcalase might exert anti-inflammatory activity. Flavourzyme hydrolysate from *Ruditapes philippinarum* exhibited a strong anti-inflammatory activity (Lee S. J. *et al.*, 2012). Many bioactive peptides have been experimentally produced under specific conditions by using different commercial proteases as shown in Table 2.5.

2.10.3.1 Alcalase

Alcalase (E.C. 3.4.21.62) is an enzyme obtained from *Bacillus licheniformis* which contains several proteinases with various specificities. Its optimum pH for catalysis ranges from 6.5-8.5. Alcalase has been used widely to prepare soluble hydrolysates with the nutritional and functional properties of natural protein maintained, such as soy protein and fish protein (REBECA *et al.*, 1991) as well as to produce bioactive peptides.

2.10.3.2 Flavourzyme

Flavourzyme (E.C. 3.4.11.1) is a kind of compound enzyme composed of the incision enzyme and the circumscribed enzyme from *Aspergillus oryzae*. Flavourzyme also acts as exopeptidase and endoprotease. Enzyme preparation is numerous and used for protein hydrolysis in diverse industrial and research applications. (Doucet *et al.*, 2003; Klompong *et al.*, 2008).

2.10.3.3 Neutrase

Neutrase (E.C. 3.4.24.28) is a protease produced by *Bacillus amyloliquefaciens* which is extensively used to break protein down into peptide. The optimal working conditions are 45-55 °C and pH 5.5-7.5. Neutrase is formulated in a way that provides the highest degree of safety during handling (Lassoued *et al.*, 2015).

Table 2.5 Proteolytic enzymes used to preparation of protein hydrolysate and peptide (Najafian and Babji, 2012).

| Enzyme | Type of protease | Source of origin | Condition | |
|-------------|----------------------------------|-----------------------------------|-----------|------------------|
| | | | pH | Temperature (°C) |
| Alcalase | Endo-protease | <i>Bacillus licheniformis</i> | 6.5-8.5 | 45-65 |
| Flavourzyme | Endo-exo-peptidase | <i>Aspergillus oryzae</i> | 5-7 | 90 |
| Neutrase | Metallo-endo-proteinase (Zn) | <i>Bacillus amyloliquefaciens</i> | 5.5-7.5 | 45-55 |
| Papain | Cysteine protease, endopeptidase | Carica papaya (papaya latex) | 6-7 | 60-70 |
| Pepsin | Acid protease and endo-peptidase | Porcine gastric mucosa | 1-4 | 37-42 |

2.11 Chicken feather meal and feather meal

Enlargement of global poultry production has been accompanied by an increase in the amount of by products such as feather meal (Chaturvedi *et al.*, 2014). Chicken feather meals are bio-resources with high protein content of more than 750 g/kg of crude protein. In the past, chicken feather meal was generally used as animal feed because of its high protein content. In addition, keratin is the main component of feathers, comprising nearly 90% of the feather's weight. Feather keratin shows an elevated content of the amino acids such as glycine, alanine, serine, cysteine and valine (Grazziotin *et al.*, 2006). Accumulations of high amounts of chicken feather meal lead to environmental pollution. Today, environmental regulations and safety are of greater concern, restricting the methods in terms of processing feathers for animal meal or waste landfilling (Kwiatkowski *et al.*, 2013). Several pretreatment methods including chemical, enzymatic or biological treatment, have therefore been investigated in the past few years to improve the digestibility of feathers (Coward-Kelly *et al.*, 2006). (Taskin *et al.*, 2011) reported that protein hydrolysate from chicken feather was rich in protein, mineral and amino acids. However, the biological activity of chicken feather meal has been less investigated and evaluated. In contrast, there has been much research on the utilization of chicken feather meal (Table 2.6). So, chicken feather meal might act as a source of protein hydrolysate with novel anti-inflammatory activity.

Table 2.6 Utilization of chicken feather meal in many researched fields

| Source | Biological activity | Result | References |
|----------------------|----------------------------------|------------------------------------------------------------------------------------------------------------------------|------------------------------------|
| Chicken feather meal | Antioxidant and antihypertensive | Feather hydrolysates displayed <i>in vitro</i> antioxidant properties and inhibit the angiotensin I-converting enzyme. | (Fontoura <i>et al.</i> , 2014) |
| Chicken feather meal | Bioenergy | Bioenergy from chicken feather meal are successfully replaces the formerly used coal burner. | (Kwiatkowski <i>et al.</i> , 2013) |
| Chicken feather meal | Biological resource | The by-product can be produce useful biotechnological products. | (Lasekan <i>et al.</i> , 2013) |
| Chicken feather meal | Keratinase production | chicken feather as a potential renewable raw material to produce keratinase | (Demir <i>et al.</i> , 2015) |

CHAPTER III EXPERIMENTAL

3.1 Biological materials

Chicken feather meal (CFM) in this study was supplied by Betagro Co., Ltd. (Thailand) and amino acid composition of chicken feather meal was determined according to the Association of Official Analytical Chemist method (AOAC 994.12).

3.2 Chemicals

Alcalase and Flavourzyme were purchased from Brentag (Mülheim, Germany). Neutrase was purchased from Novozymes (Bagsvaerd, Denmark). Acetic acid, ethanol and phosphoric acid were purchased from Merck (Gibbstown, NJ). Bovine serum albumin (BSA), curcumin from *Curcuma longa* (Turmeric), 3-(4, 5-dimethylthiazol-2-yl)-2,5-diphenyl-2H-tetrazolium bromide (MTT), Dulbecco's modified Eagle medium (DMEM), fetal bovine serum (FBS), interferon-g from mouse (IFN-g), Lipopolysaccharide (LPS) from *Escherichia coli*, potassium persulfate, sodium nitrite, (1-Naphthyl)ethylenediamine (NED), sodium nitroprusside (SNP), sodium pyruvate, sulfanilamide, streptomycin sulfate were purchased from Sigma-Aldrich (St. Louis, MO).

3.3 Determination of amino acid content

Amino acid content of chicken feather meal was determined based on the standard AOAC 994.12 acid hydrolysis method. 1 g of chicken feather meal was dissolved in 5 ml of 6 N HCl in test tube, and placed in heating block at 110 °C for 24 h. to liberate the amino acid. Then, the internal standard (10 ml 2.5 mM L- α -amino-n-butyric acid in 0.1 M HCl) was added, diluted with deionized water to 250 ml and placed in a heating block at 55 °C for 10 min. Amino acid were determined by reverse-phase high performance liquid chromatography (RP-HPLC) analysis on a Hypersil GOLD column C₁₈ (4.6 mm X 150 mm, 3 μ M), elute with sodium acetate buffer pH 4.90 and 60% acetonitrile, at flow rate of 0.3 ml/min.

3.4 Preparation of enzymatic hydrolysate from chicken feather meal

CFM was dried at 60 °C overnight and ground pass through a 150 micron sieve. 5 g of CFM was mixed with 100 ml of phosphate buffer saline (20 mM phosphate buffer with 0.15 M NaCl pH 7.2). Then the suspensions were stirred overnight at 4 °C overnight. Three type of microbial protease; Alcalase[®], Flavourzyme[®] and Neutrase[®] were hydrolyzed in various concentrations (0, 1, 2.5 and 5%). The optimum pH, temperature and characterization of various enzymes are summarized in Table 3.1. The reactions were carried out for 4 h with shaking (150 rpm) respectively. After hydrolysis, the mixtures were heated at 90 °C for 10 min to inactive enzyme, and then centrifuged at 10,000 rpm for 15 min and collected the supernatant. The hydrolysates were stored in -20 °C until use.

Table 3.1 Condition for enzymatic hydrolysate from chicken feather meal

| Conditions | Alcalase | Flavourzyme | Neutrase |
|-------------------------|----------|-------------|----------|
| pH | 8.0 | 7.0 | 7.0 |
| Temperature (°C) | 50 | 50 | 50 |
| Time (h) | 4 | 4 | 4 |

3.5 Protein Content Determination

To determine the protein concentration of CFM protein hydrolysate was measured according to Bradford's procedure (Bradford, 1976), using bovine serum albumin (BSA) as the standard curve with various concentrations to construct the calibration curve. 20 µL of each sample were mixed with 200 µL Bradford working buffer into each of three well of a 96-well plate, and then incubate for 20 min. The absorbance was measured at 540 nm with a microplate reader spectrophotometer. The absorbance value was calculated to protein concentration using linear equation computed from standard curve.

3.6 Nitric oxide radical scavenging assay

Nitric oxide (NO) radical scavenging assay was slightly modified method from Chantaranothai et al., (2013). The peptide hydrolysates (5 μ L) was mixed with 10 mM of sodium nitroprusside in a phosphate buffer saline pH 7.2 and incubate at room temperature for 150 min. Then 100 μ L Griess reagent (0.33% sulfanilamide in 5% phosphoric acid) was added and incubated for 5 min. After that, 100 μ L of *N*-naphthyl-ethylidiamine (NED) was added and further incubated for 30 min at room temperature. The absorbance was determined at 540 nm spectrometer. The percentage of scavenging activity (%) and IC₅₀ were calculated. Curcumin was used as positive control. The percentage of scavenging activity was calculated using the following this equation.

$$\% \text{ inhibition} = \left[\frac{(\text{Abs control} - \text{Abs blank}) - (\text{Abs sample} - \text{Abs background})}{(\text{Abs control} - \text{Abs blank})} \right] \times 100$$

3.7 Purification of protein hydrolysate from CFM

3.6.1 Molecular weight cut-off by ultrafiltration

The peptide hydrolysate from CFM were fractionated through ultrafiltration (UF) membranes with a range of molecular weight cut-off (MWCO) of 10, 5, 3 and 0.65 kDa. Peptide hydrolysates were divided into five fraction including >10 kDa, 10-5 kDa, 5-3 kDa, 3-0.65 kDa and < 0.65 kDa. All fractions were collected to determine nitric oxide radical scavenging activity.

3.7.2 Gel filtration chromatography

The fraction was shown highest inhibit nitric oxide radical scavenging from molecular weight cut-off (\leq 0.65 kDa) by ultrafiltration was loaded onto a preparative Sephacryl S-200 gel filtration column chromatography (AKTA™ prime with Hitrap™, Amersham Biosciences), column (2.6 x 80 cm.) at flow rate 0.5 ml/min. Each fraction of sample was detected at 280 nm and assayed for nitric oxide radical scavenging.

3.7.3 Reverse-phase high-performance liquid chromatography (RP-HPLC)

The most effective fraction from gel filtration chromatography was further purified by reverse-phase high performance liquid chromatography on C₁₈ column (250×4.6 mm, Luna 5U, Phenomenex, Torrance, CA, USA). The peptide was eluted with rational gradient of mobile phase A (0.1% v/v trifluoroacetic acid) and mobile phase B (70% (v/v) acetonitrile in 0.05% (v/v) trifluoroacetic acid) at flow rate 0.7 ml/min. The elution peak was detected at 280 nm. Chromatographic analyses were completed with ChromQuest Software (Thermo Scientetific).

3.8 Identification of anti-inflammatory peptides

Amino acid sequences of the purified peptides were determined by LC-MS/MS Q-TOF mass spectrometer. The system consisted of liquid chromatography in combination with a quadrupole time to flight mass spectrometer (Q-TOF). LC-MS/MS data in mass range from 25 to 20,000 m/z. All of data were processed and analyzed by de novo sequencing which is to use the mass difference between two fragment ions to calculate the mass of an amino acid residue on the peptide chain.

3.9 Cell culture

The macrophage RAW 264.7 were grown in Dulbecco's Modified Eagle's Medium (DMEM) supplemented with 10% (v/v) FBS, 100 U/ml penicillin-G, 0.4 mg/ml streptomycin sulfate, 1% (w/v) sodium pyruvate, and 1% (w/v) of HEPES at 37 °C in a humidified atmosphere with 5% (v/v) CO₂. For routine maintenance in culture (passage) (Saisavoey *et al.*, 2016), cells were seed in non-tissue culture treated dish at approximately 10% confluency and grown to approximately 80% confluency, which typically took 2 days. Old medium was aspirated, cells were gently rinsed with PBS, then dislodged by gently scraping with a rubber spatula and then harvest by centrifugation 15,000 x g for 5 min. PBS was removed, the cell pellet was resuspended in CM and the in CM and the viable and total cell density was evaluated by using Trypan blue dye exclusion on an improved Neubauer hemacytometer. The following equation was then used to calculate the viable cell numbers.

Total cell count (cells/ml) = the number of cells counted in 16-large squares $\times 2 \times 10^4$

For storage, cells were harvested and resuspended in cold freezing media (CM plus 10% (v/v) dimethyl sulfoxide) at 2×10^6 cells/ml, aliquoted at 1 ml per cryogenic tubes and transferred to -80°C immediately overnight and then into liquid nitrogen. To initiate cultures from frozen stocks the cells were thawed at 37°C , transferred into 10 ml of serum free DMEM media and centrifuged at $15,000 \times g$ for 5 min. The cell pellet was resuspended in CM, plated and cultured as above except that after the cells had adhere (approximately 6 h), the old medium was replaced.

3.10 Pretreatment of macrophage RAW 264.7 Cells

The macrophage RAW 264.7 cells were seeded in 96-well plates at a density of 1×10^4 cells/well in 100 μl CM and incubated overnight at 37°C in 5% (v/v) CO_2 . After that the medium was replaced with a fresh supplemented without sample (negative control), various concentration of protein hydrolysate from CFM or parthenolide at 2.5 $\mu\text{g/ml}$ was utilized as a positive control and incubated for 1 h. Then, NO production was stimulated by the addition of 100 ng/ml of lipopolysaccharide (LPS) and incubated for 12 h.

3.11 MTT assay for measuring of cell proliferation

The cytotoxicity activity of CFM protein hydrolysate was determined by MTT assay with slightly modified according to Yang *et al.* (2014); (Chantaranothai *et al.* (2013)). The macrophage RAW 264.7 cells were plated at density of 1×10^4 in 96-well plate overnight. After overnight, cells were treated with sample in different concentration and LPS. Then 100 μl of 5 mg/ml MTT solution (in PBS) was added per well. After incubated at 37°C , 5% CO_2 incubator for 4 h. Finally, dimethyl sulfoxide (DMSO) was added 100 μl /well in 96-well plates due to DMSO solubilizes the formed formazan crystals. The volume of formazan crystals was determined by measuring the absorbance at 540 nm.

3.12 Determination of nitric oxide (NO) production from macrophage RAW 264.7 cells

Nitric oxide production was determined by measure nitrite production in the culture medium according to the Griess reaction. The macrophage RAW 264.7 cells were incubated with samples in various concentrations and without sample. Fifty-microliter of sulfanilamide was added to 50 μ l of the culture supernatant in a 96-well plate and incubate at room temperature for 10 min. after that, Fifty-microliter of NED solution was added and further incubate for 10 min, then absorbance at 540 was measured using microplate reader.

3.13 Total RNA Isolation from macrophage RAW 264.7 Cells

To study about the mechanism of action on cytokine release, the assay for mRNA expression were carried out. Total RNA was isolated from macrophage RAW 264.7 cells which treated with solvent (negative control), CFM protein hydrolysate in various concentrations and 2.5 μ g/ml of parthenolide (positive control). Cells were stimulated with 100 ng/ml of LPS. After 12 h incubation, to harvest the total RNA by using the MaterPure™ Complete DNA & RNA Purification Kit (Epicentre) were used according to the manufacturer's instruction. The RNA concentration was measured by using Nanodrop 2000 UV-vis Spectrometer (Thermo Fisher Scientific, Inc., USA).

3.14 Detection of iNOS, IL-6, TNF- α , COX-1 and COX-2 mRNA by Reverse transcription polymerase chain reaction (RT-PCR) analysis

A content of 1 μ g of total RNA from each condition was reverse-transcription became single-stranded complementary DNA (cDNA) was synthesis by using Precision™ nanoscript 2 Reverse Transcription kit (PrimerDesign Co., Ltd. UK) according to manufacturer's protocol. Total volume of cDNA synthesis were consisting of up to 5 μ g of total RNA, 1 μ l of RT primer and RNase/DNase free water were adjusted to final volume (10 μ l). After that incubate at 65 °C for 5 min, then immediately cool in ice water bath for 1 min. After that, nanoscript2 4X buffer 5 μ l, dNTP mix 10 mM, RNase/DNase free water 3 μ l and 1 μ l of nanoscript2 enzyme were added. The mixture was gently mixed and reaction was performed at 42 °C for 20 min, 75 °C for 10 min. cDNA was used as a template for PCR.

Then PCR analysed were performed on aliquots of the cDNA preparation to detect inducible nitric oxide synthase (iNOS), Interleukin 6 (IL-6), tumor necrosis factor alpha (TNF- α), Cyclooxygenase-1 (COX-1), Cyclooxygenase-2 (COX-2) and beta-actin (β -actin). The PCR primer of each gene in this study are listed below in Table 3.2 denaturation step at 95 °C for 3 min, 30 s at 95 °C for denaturation, annealing step of each primer were shown in table 3.2 for 1 min, and extension at 72 °C for 1 min. After amplification, the RT-PCR reaction product were separated with 1.2% (w/v) agarose gel electrophoresis and stained with ethidium bromide for 10 min. Then, visualized under a UV light using gel documentation system (Biorad, Hercules, CA).

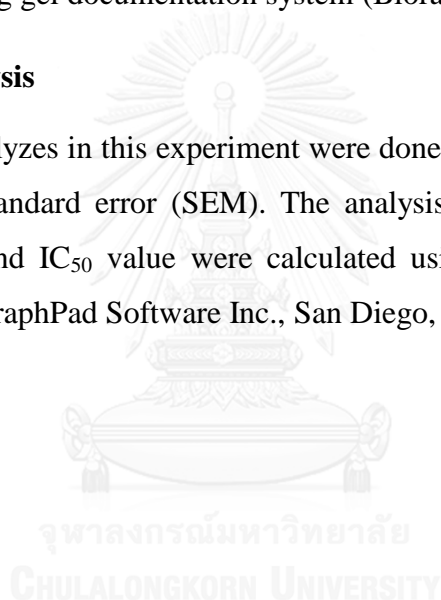
Table 3.2 The PCR primer in this study

| Primer | Sequence (5' to 3') | Annealing temperature (°C) | Base pair (bp) |
|------------------------|----------------------------|-----------------------------------|-----------------------|
| β -actin forward | ACCAACTGGGACGACATGGAGGA | 55 | 380 |
| β -actin reward | GTGGTGGTGAAGCTGTAGCC | | |
| iNOS forward | CCATCATGGACCACCACACA | 65 | 423 |
| iNOS reward | CCATGCAGACAACCTTGGTG | | |
| IL-6 forward | CATGTTCTCTGGGAAATCGTGG | 50 | 417 |
| IL-6 reward | AACGCACTAGGTTTGCCGAGTA | | |
| TNF- α forward | CCTGTAGCCCACGTCGTAGC | 50 | 375 |
| TNF- α reward | TTGACCTCAGCGCTGAGTTG | | |
| COX-1 forward | AGTGCGGTCCAACCTTATCC | 63 | 382 |
| COX-1 reward | CCGCAGGTGATACTGTCGTT | | |
| COX-2 forward | GGAGAGACTATCAAGATAGT | 50 | 861 |
| COX-2 reward | ATGGTCAGTAGACTTTTACA | | |

The PCR mixture consisted of cDNA 1 μ l, RNase free water 10.5 μ l, PCR reagent (Pcrbiosystem, United Kingdom) 12.5 μ l, 10 μ M forward primer 1 μ l and 10 μ M reverse primer 1 μ l. The PCR Reaction was carried out in a volume of 25 μ l (final volume) containing. Amplification was completed by using a PCR thermal cycler (Bio-Rad, Laboratories, Inc., USA) for PCR with following condition: initial denaturation step at 95 °C for 3 min, 30 s at 95 °C for denaturation, annealing step of each primer were shown in table 3.2 for 1 min, and extension at 72 °C for 1 min. After amplification, the RT-PCR reaction product were separated with 1.2% (w/v) agarose gel electrophoresis and stained with ethidium bromide for 10 min. Then, visualized under a UV light using gel documentation system (Biorad, Hercules, CA).

3.15 Statistical analysis

Statistical analyzes in this experiment were done for triplicate and results were shown as mean \pm standard error (SEM). The analysis was performed using SPSS statistical software and IC₅₀ value were calculated using GraphPad Prism Version 6.01 for Windows (GraphPad Software Inc., San Diego, CA, USA).



CHAPTER IV

RESULTS AND DISCUSSION

4.1 Amino acid composition of chicken feather meal

The amino acid composition of chicken feather meal is shown in Table 4.1. The amino acid composition of protein material plays a significant role in various physiological and biological activities. According to the result, chicken feather meal was rich in 18 types of amino acid including alanine, arginine, glycine, aspartic acid, valine, cysteine, glutamic acid, leucine, isoleucine, histidine, threonine, proline, lysine, methionine, serine, phenylalanine, tyrosine and tryptophan. In addition, anti-inflammatory peptide from *Mytilus coruscus* which consists of glycine, valine, serine, leucine, glutamine and phenylalanine has shown anti-inflammatory activity (Kim E.-K. *et al.*, 2013). In addition, Lee S. J. *et al.* (2012) reported that peptide consisting of glycine, cysteine, alanine, valine and serine can exhibit nitric oxide production. There is a previous study reporting that glycine, cysteine and histidine exhibited anti-inflammatory activity by inhibiting the NF-KB signaling pathway (Hasegawa *et al.*, 2012).

Table 4. 1 Amino acid profile of chicken feather meal (mg/100 mg)

| Amino acids | Results (%) |
|----------------------|--------------------|
| Alanine (Ala) | 3.80 |
| Arginine (Arg) | 6.30 |
| Glycine (Gly) | 6.82 |
| Aspartic acid (Asp) | 5.70 |
| Valine (Val) | 5.85 |
| Cysteine (Cys) | 2.90 |
| Glutamic acid (Glu) | 10.6 |
| Leucine (Leu) | 6.46 |
| Isoleucine (Ile) | 3.94 |
| Histidine (His) | 0.59 |
| Threonine (Thr) | 3.96 |
| Proline (Pro) | 8.37 |
| Lysine (Lys) | 1.45 |
| Methionine (Met) | 0.67 |
| Hydroxyproline (Hyp) | Not detected |
| Serine (Ser) | 7.84 |
| Phenylalanine (Phe) | 4.03 |
| Hydroxylysine (Hyl) | Not detected |
| Tyrosine (Tyr) | 1.10 |
| Tryptophan (Trp) | 0.20 |

4.2 In vitro nitric oxide radical scavenging assay

Nitric oxide is a reactive radical involved in the inflammatory process in living organisms. The NO scavenging activity may help prevent the chain reaction that promotes excessive generation of NO causing damage to human health (Yu M. *et al.*, 2014). The crude protein hydrolysate of chicken feather meal from various types of protease were subjected to testing of their anti-inflammatory activity by using nitric oxide radical scavenging assay. The inhibition results in Table 4.2 suggest that the 1% flavourzyme hydrolysate ($5.49 \pm 0.97 \mu\text{g/ml}$) exhibited the highest nitric oxide radical scavenging activity when compared with half maximal inhibitory concentration (IC_{50}). The 1% neutrase hydrolysate had shown weak NO scavenging activity ($175.70 \pm 3.96 \mu\text{g/ml}$). These results were comparable with curcumin as the standard for inhibiting this reaction (positive control). Therefore, 1% flavourzyme hydrolysate was more effective than curcumin. In addition, there has been some research about protein hydrolysate which has anti-inflammatory activity that has reported that flavourzyme hydrolysate from *Mytilus coruscus* has a high capability of inhibiting nitric oxide production in macrophage RAW 264.7 cells (Kim E.-K. *et al.*, 2013). The results for 1% flavourzyme hydrolysate concurred with previous study. Therefore, the 1% flavourzyme hydrolysate responsible for anti-inflammatory was purified for further study.

Table 4. 2 Nitric oxide radical scavenging activity of crude peptide from chicken feather meal

| Enzymes | NO radical scavenging activity IC ₅₀ (µg/mL) | | | |
|-------------|---------------------------------------------------------|---------------------------|--------------------------|--------------------------|
| | Control | 1% | 2.5% | 5% |
| Alcalase | 134±4.72 ^e | 80.59±5.40 ^d | 29.97±1.81 ^a | 40.23±0.78 ^b |
| Flavourzyme | 158.37±1.50 ^E | 5.49±0.97 ^A | 82.07±6.96 ^D | 68.68±3.57 ^C |
| Neutrase | 175.70±3.96 ^{ee} | 133.03±5.40 ^{dd} | 36.99±0.43 ^{aa} | 78.71±3.91 ^{cc} |

*All data are shown as mean±standard deviation of triplicate (the result show significant difference of $p \leq 0.05$)

** curcumin as a positive control with IC₅₀ = 60.53±3.18 µg/ml

4.3 Fractionation of 1% Flavourzyme hydrolysate by Ultrafiltration

For further study, 1% Flavourzyme hydrolysate was filtered through different molecular weight cutoff (MWCO) membranes: 10, 5, 3 and 0.65 kDa membrane filters (Pellicon XL filter, Merck). Then, each fraction was subjected to examination for nitric oxide radical scavenging activity. The results of each fraction are shown in Table 4.3. The anti-inflammatory activity of the fraction ≤ 0.65 kDa size molecular weight exhibited the most effective nitric oxide radical scavenging activity compared with the others (3.64±0.27 µg/ml). It was reported that the biological activity of hydrolysates depend on the molecular weight of each peptide (Wang A.-Y. *et al.*, 2011). An array of anti-inflammatory peptides with low molecular weights fractionated by ultrafiltration were effective in interaction with the internal barrier to promote the biological effect (Ahn *et al.*, 2015). So, low molecular weight peptides are more effective than high molecular weight peptides. Thus, this fraction was selected for further study.

Table 4. 3 Nitric oxide radical scavenging activity of peptide fraction from ultrafiltration

| Molecular weight (kDa) | NO radical scavenging activity IC ₅₀ (µg/mL) |
|------------------------|---------------------------------------------------------|
| > 10 kDa | 21.87±0.70 ^d |
| 5-10 kDa | 10.42±0.67 ^c |
| 3-5 kDa | 7.17±0.47 ^b |
| 0.65-3 kDa | 5.82±0.56 ^{ab} |
| <0.65 kDa | 3.64±0.27 ^a |

*All data are shown as mean±standard deviation of triplicate (the result show significant difference of $p \leq 0.05$)

** curcumin as a positive control with IC₅₀ = 60.53±3.18 µg/ml

4.4 Sephacryl S-200 gel filtration chromatography of 1% flavourzyme hydrolysate

The anti-inflammatory peptide fraction ≤ 0.65 kDa was subjected to the Sephacryl S-200 column, and the active peaks were separated into four peaks (F1, F2, F3 and F4), as shown in Figure 4.1. The four fractions were evaluated for nitric oxide radical scavenging activity. The results of each fraction are shown in Table 4.4. The F2 fractions exhibited the highest nitric oxide radical scavenging activity among all fractions with IC₅₀ values of 34.60±3.45 µg/ml. The IC₅₀ of nitric oxide radical scavenging activity for peptide fractions F3 and F4 could not be calculated due to their low concentrations of protein. Then the fraction F2 was further purified and studied.

Table 4. 4 Nitric oxide radical scavenging activity of peptide fraction from Sephacryl S-200 gel filtration chromatography

| Nitric oxide radical scavenging activity IC ₅₀ (µg/ml) | | | |
|-------------------------------------------------------------------|-------------------------|------------|------------|
| Fraction 1 | Fraction 2 | Fraction 3 | Fraction 4 |
| 73.49±2.77 ^b | 34.60±3.45 ^a | >51.47 | >33.71 |

*All data are shown as mean±standard deviation of triplicate (the result show significant difference of $p \leq 0.05$)

** curcumin as a positive control with IC₅₀ = 60.53±3.18 µg/ml

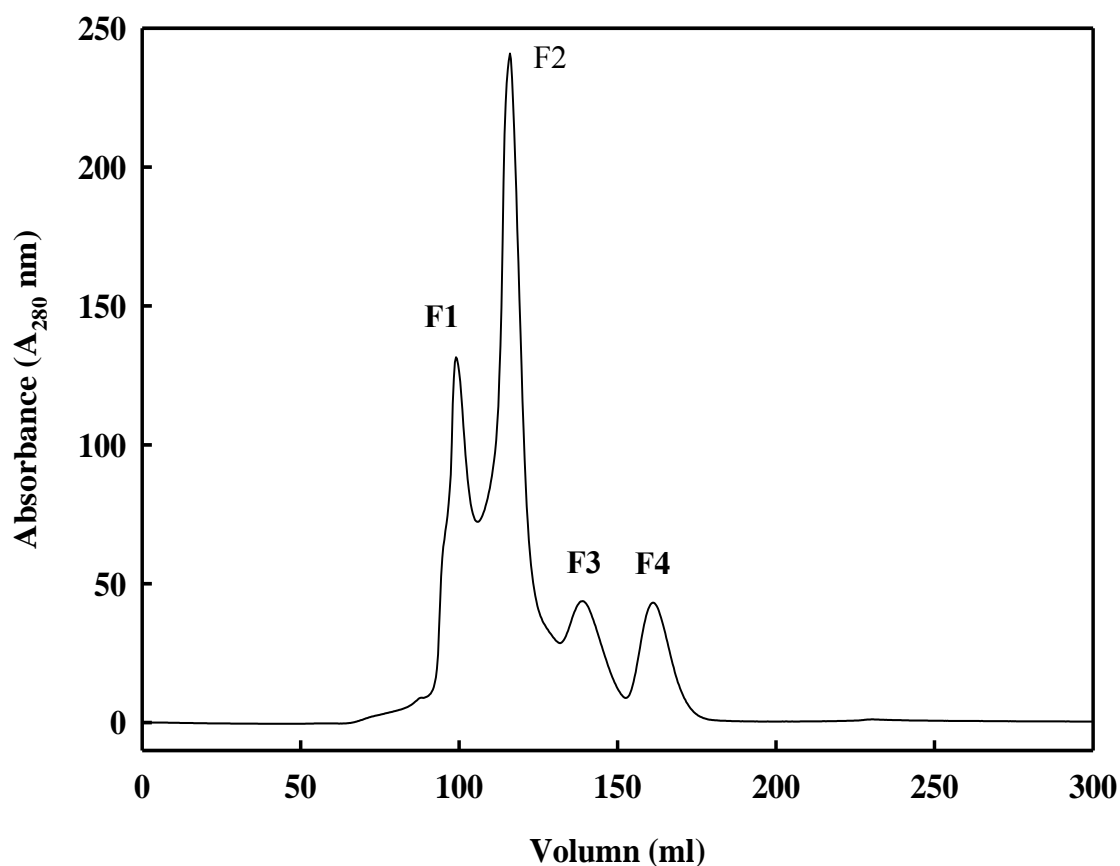


Figure 4. 1 Gel filtration chromatogram of 1% flavourzyme hydrolysate from chicken feather meal (MW < 0.65 kDa) from Sephacryl S-200 gel filtration chromatography

4.5 MTT assay for measuring of cells proliferation

The fraction F2 from gel filtration chromatography was measured cells proliferation of Macrophage RAW 264.7 cells using MTT assay. Mostly, MTT assay is mainly use as determined cells viability, cells proliferation and cytotoxicity of drug, herbal extract and bioactivity compound (Han *et al.*, 2010). The results are given in Figure 4.2. The F2 fraction did not exhibit any cytotoxic effect on macrophage RAW 264.7 cells within the range of concentration up to 40 $\mu\text{g/ml}$. These studies confirm that 1% flavourzyme hydrolysate was not cytotoxic to macrophage RAW 264.7 cells.

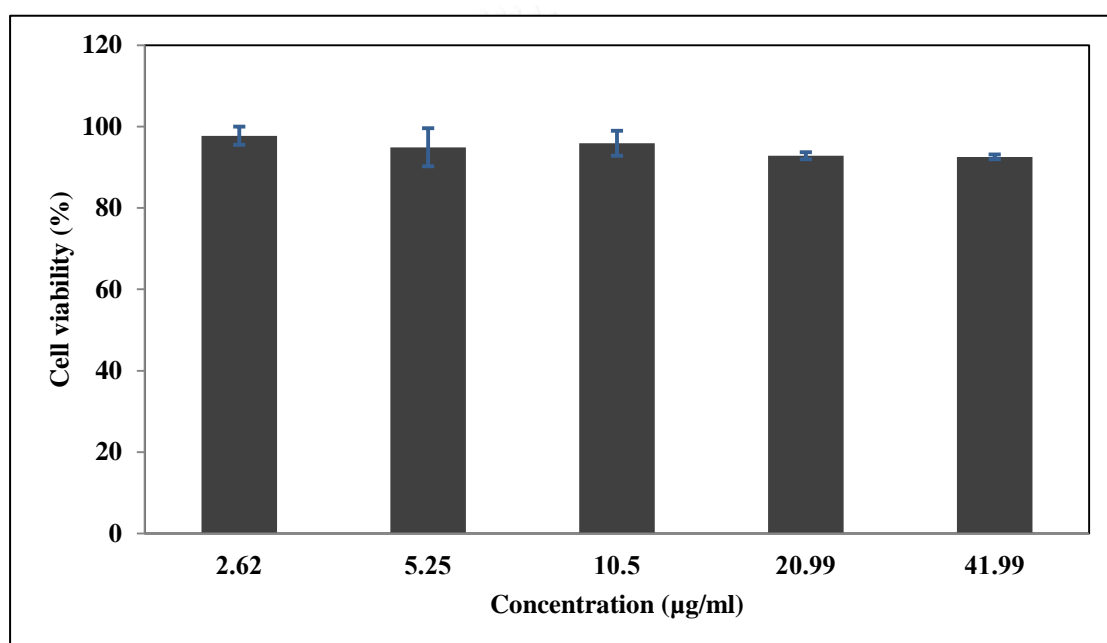


Figure 4. 2 Cell viability of macrophage RAW 264.7 cells in the presence of the F2 fraction, which show no toxicity against the growth of the macrophage.

4.6 Determination of nitric oxide (NO) production from Macrophage RAW 264.7

Nitric oxide (NO) plays an important role in the inflammatory mechanism, immune response and is mainly released from macrophage cells. The treatment of macrophage RAW 264.7 cells with LPS activation resulted in increased NO production in the culture medium. Then the F2 fraction was added into the culture medium to LPS activation in normalized NO production. As can be seen from the results in Figure 4.3, the F2 fraction from gel filtration chromatography was observed to exhibit LPS-induced NO production from macrophage RAW 264.7.

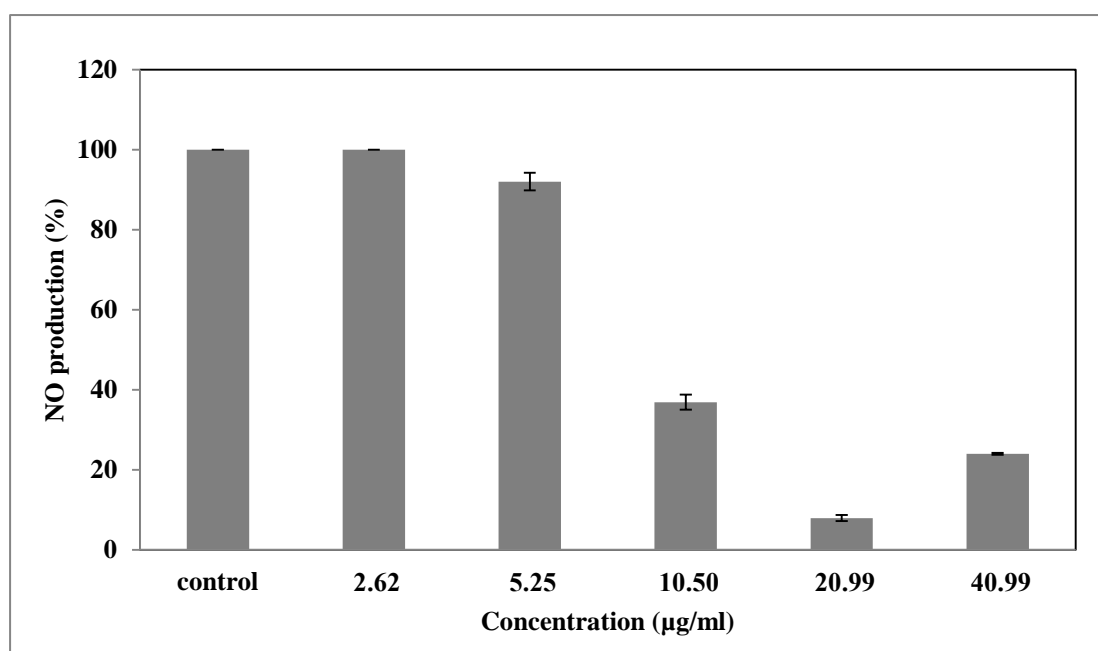


Figure 4. 3 Inhibitory effect of 1% Flavourzyme protein hydrolysate from chicken feather meal on LPS-induced Macrophage RAW 264.7 cell.

4.7 Purification of 1% flavourzyme hydrolysate by RP-HPLC

Throughout the purification process, the potent fraction F2 obtained using gel filtration chromatography showed the highest nitric oxide radical scavenging activity. These fractions were further separated by reverse-phase HPLC on a Shimpack C-18 column (250x46 mm) using a gradient of trifluoroacetic acid (TFA) 0.1% and acetonitrile containing a trifluoroacetic acid 0.05% solvent system at a flow rate of 0.7 ml/min. Elution was detected at 280 nm. In this purification step, peptides were isolated into five fractions: F2-1, F2-2, F2-3, F2-4 and F2-5. The RP-HPLC chromatogram is shown in Figure 4.4. From the result, the nitric oxide radical scavenging activities of F2-1, F2-2, F2-3 and F2-5 reached 16.79 ± 1.47 , 15.34 ± 0.66 , 17.43 ± 0.39 and 33.99 ± 1.93 $\mu\text{g/ml}$, but F2-4 had a low concentration of protein so the IC_{50} value of nitric oxide radical scavenging activity could not be calculated. The fractions which showed nitric oxide radical scavenging activities were subjected to further identification of the amino acid sequence by LC-MS/MS Q-TOF mass spectrometry.

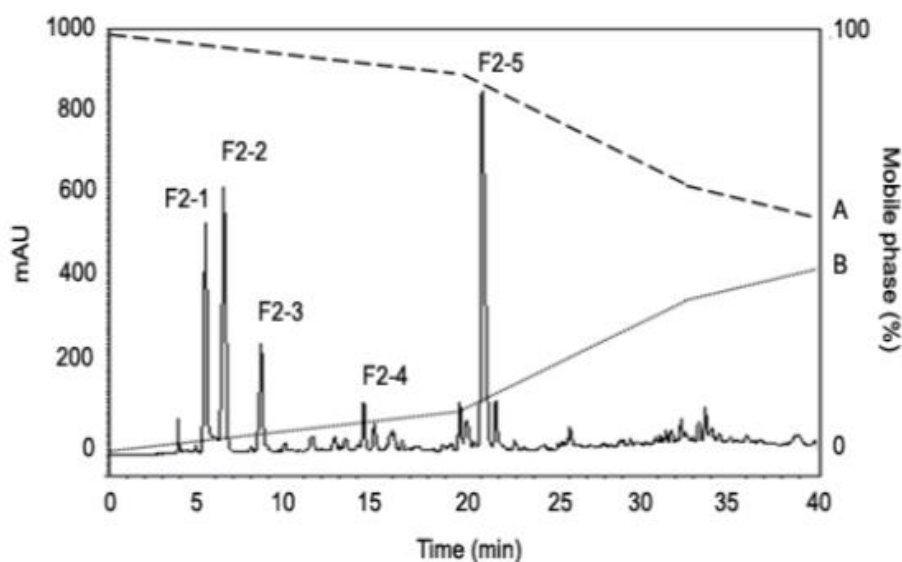


Figure 4. 4 RP-HPLC Chromatogram of 1% flavourzyme hydrolysate from chicken feather meal on a Luna C₁₈ (4.6 mm x 250 mm) column.

4.8 Identification of anti-inflammatory peptide

The mass spectrometry technique has widely emerged as a technology of choice with high throughput and sensitivity to the analysis of amino acid sequence and peptide identification. The peptide fractions from RP-HPLC which had nitric oxide radical scavenging activity (F2-1, F2-2, F2-3 and F2-5) were further identified by LC-MS/MS Q-TOF mass spectrometers for the characterization of peptides. Normally, there are two methods to analyze peptide interpretations: the database search and de novo sequencing (Frank *et al.*, 2007; Li and Deinzer, 2007). The results in this experiment were identified by de novo sequencing in fractions F2-1, F2-3 and F2-5 and F2-2 and were analyzed by the MASCOT database program. From purification by RP-HPLC, 1% flavourzyme hydrolysate was separated into five fractions which were analyzed by LC/MS/MS, and only four fractions inhibited nitric oxide production. The amino acid sequences and molecular masses of fractions F2-1, F2-2, F2-3 and F2-5 are shown in Table 4.5, and identified in four peptide chains. Each sequence contained around 9-13 amino acid residues. Fraction 1 (F2-1) was identified as SNPSVAGVR, Fraction 2 (F2-2) as SLFLHTHSIVADK, Fraction 3 (F2-3) as AVLKKKVTSTFGR and Fraction 4 (F2-5) as LSPWPVKGV. Regarding the relationship between the properties and anti-inflammatory activity of amino acids, all of the peptide sequences contained mainly amino acids as the anti-inflammatory of Gly (G) and His (H). Thus, there are some hydrophilic aminos: Ala (A) Phe (F), Leu (L), Pro (P) and Trp (W). These have also been reported to have high antioxidant activity (Bougatef *et al.*, 2010; Mendis *et al.*, 2005; Sacchetti *et al.*, 2008). The anti-inflammatory activity of glycine and histidine has been reported by Hasegawa *et al* (Hasegawa *et al.*, 2012) in their study on glycine, histidine and cysteine. It has been suggested that glycine, histidine and cysteine also acts as an anti-inflammatory by reducing NF-KB activation and inhibiting the expression of IL-6 in human coronary arterial endothelial cells.

Table 4. 5 Peptide sequence of 1% flavourzyme hydrolysate from chicken feather meal identified by Q-TOF LC/MS/MS

| Fractions | Amino acid sequence | Protein name | Organism | Accession number |
|------------------|----------------------------|--------------------------------------------------------|------------------------|-------------------------|
| F2-1 | SNPSVAG VR | putative E3 ubiquitin-protein ligase SH3RF2 | <i>Gallus gallus</i> | XP_01515 7384.1 |
| | | putative E3 ubiquitin-protein ligase SH3RF2 isoform X2 | <i>Gallus gallus</i> | XP_01515 7384.1 |
| | | putative E3 ubiquitin-protein ligase SH3RF2 isoform X1 | <i>Gallus gallus</i> | XP_41466 2.3 |
| | | PRELI domain containing protein 3B [Gallus gallus] | <i>Gallus gallus</i> | NP_00102 6037.1 |
| | | hypothetical protein RCJMB04_17b4, partial | <i>Gallus gallus</i> | CAG3206 5.1 |
| F2-2 | SLFLHTHS IVADK | Lactate dehydrogenase | <i>Phodilus badius</i> | GI629677 270 |
| F-2-3 | AVLKKKV TSTFGR | cystine/glutamate transporter | <i>Gallus gallus</i> | XP_42628 9.3 |
| | | dynein heavy chain 1, axonemal | <i>Gallus gallus</i> | XP_01514 8334.1 |
| | | unconventional myosin-XVI isoform X2 | <i>Gallus gallus</i> | XP_00493 8593.1 |
| | | unconventional myosin-XVI isoform X1 | <i>Gallus gallus</i> | XP_41695 0.3 |
| | | transcription initiation factor TFIID subunit 9B | <i>Gallus gallus</i> | NP_00126 4725.1 |

Table 4.5 Peptide sequence of 1% flavourzyme hydrolysate from chicken feather meal identified by Q-TOF LC/MS/MS

| Fractions | Amino acid sequence | Protein name | Organism | Accession number |
|------------------|----------------------------|---------------------------------------------------|----------------------|-------------------------|
| F2-5 | LSPWPVKGV | zinc finger matrin-type protein 1-like isoform X1 | <i>Gallus gallus</i> | XP_004936323.1 |
| | | neogenin isoform X6 | <i>Gallus gallus</i> | XP_015134538.1 |
| | | neogenin isoform X5 | <i>Gallus gallus</i> | XP_015134537.1 |
| | | neogenin isoform X4 | <i>Gallus gallus</i> | XP_015134536.1 |
| | | neogenin isoform X3 | <i>Gallus gallus</i> | XP_015134535.1 |

4.9 Anti-inflammatory effect of the F2 Fraction in LPS induced Macrophage RAW 264.7 cells

NO and pro-inflammatory cytokine plays a critical role in the physiology and pathology of diverse tissues including the immune system of the body. The F2 fraction was tested for nitric oxide radical scavenging activity and inhibitory effect against NO production on LPS-induced macrophage RAW 264.7 cells and was shown to have the highest activity against NO production with an IC₅₀ value of 34.60±3.45 µg/ml. In order to investigate the mechanism of purified F2 fraction on LPS-induced NO production and inflammatory cytokines such as IL-6, TNF-α and COX-2 expression which are known to be secreted by LPS-induced macrophage RAW 264.7 cells by RT-PCR, it was compared with budesonide (positive control). The results showed that unstimulated macrophage RAW 264.7 cells iNOS, Cox-2, TNF-α and IL-6 were undetectable. However, iNOS, Cox-2, TNF-α and IL-6 were expressed at the mRNA level by LPS and the F2 fraction significantly inhibited these expressions. When the F2 fraction was compared with budesonide as a positive control, it was found that both the F2 fraction and budesonide strongly inhibited the LPS mediating increased incellular iNOS, IL-6 and TNF-α mRNA level (Figure 4.5). Nevertheless, the fraction F2 did not affect the expression of COX-1, for maintaining normal physiological functions, and β-actin, the house keeping gene. Inducible nitric oxide synthase (iNOS) is involved in the response to various pro-inflammatory cytokines including interferon-γ (INF-γ), tumor necrosis factor-α (TNF-α), and interleukin-6 (IL-6), and mediates several inflammatory responses (Jung *et al.*, 2007). Therefore, the results also agreed with the results of the inhibitory effect of nitric oxide production. Moreover, it has been reported that the inducible isoforms of NOS and COX-2 are involved in the production of large amounts of NO and prostaglandin E₂ (PGE₂), respectively (Lappas *et al.*, 2002; Wang S. Y. *et al.*, 2008). The activation of nuclear factor kappaB (NF-KB) is responsible for the expression of iNOS and COX-2 (Brown *et al.*, 2008; Park *et al.*, 2006).

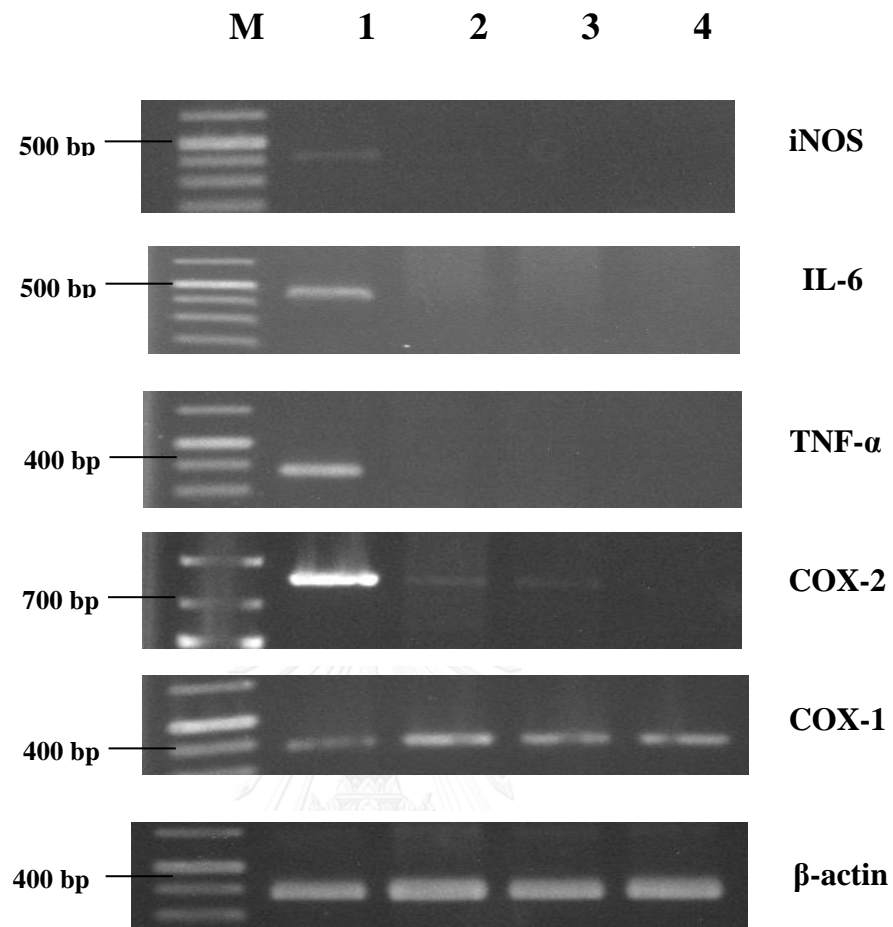


Figure 4.5 The effect of the F2 fraction on LPS induced iNOS (423 bp), IL-6 (417 bp), COX-1 (382 bp), COX-2 (861 bp) and TNF- α (375 bp) expression in macrophage RAW 264.7 cells. For all panels: Lane 1: LPS, Lane 2: LPS + F2 fraction (40 μ g/ml), Lane 3: LPS+budesonide (positive control), Lane 4: no addition and Lane M: 1000 bp

CHAPTER V

CONCLUSION

In conclusion, the present study demonstrated chicken feather meal peptide produced by microbial enzyme which had anti-inflammatory activity was evaluated by nitric oxide radical scavenging assay. The results showed that 1% flavourzyme hydrolysate had the highest nitric oxide radical scavenging activity. Then, the peptide was fractionated by ultrafiltration indicating that the lowest molecular weight peptides ($MW \leq 0.65$ kDa) were effective in their nitric oxide radical scavenging activity. The F2 fraction from Sephacryl S-200 gel filtration chromatography was not toxic and helped prevent NO production in macrophage RAW 264.7 cell-induced by LPS. The anti-inflammatory peptide was further purified by RP-HPLC and the amino acid sequence identified in 4 chains: SNPSVAGVR, SLFLHTHSIVADK, AVLKKKVTSTFGR and LSPWPVKGV. The peptide sequence contained amino acids with anti-inflammatory activity especially glycine and histidine. Moreover, the F2 fraction is a potent inhibitor of LPS-induced NO, PGE₂, TNF- α and IL-6 production at mRNA level which is a pro-inflammatory cytokine. These results suggest that chicken feather meal hydrolysate could be used as a natural anti-inflammatory in functional food or pharmaceutical products.

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APPENDIX



จุฬาลงกรณ์มหาวิทยาลัย
CHULALONGKORN UNIVERSITY

APPENDIX A

Preparation of 20 mM PBS with 0.15 M NaCl 1 L

| | |
|--------------------------|--------|
| KH_2PO_4 | 2.72 g |
| K_2HPO_4 | 3.48 g |
| NaCl | 8.77 g |

Adjusted pH to 7.2 with 1M KOH and adjusted volume to 1 L with deionized water.

Preparation of Bradford solution

Bradford Stock Solution 300 ml

| | |
|---------------------|--------|
| 95% ethanol | 100 ml |
| 88% phosphoric acid | 200 ml |
| Serva Blue G | 350 mg |

Bradford Working buffer 500 ml

| | |
|-------------------------|--------|
| 95% ethanol | 15 ml |
| 88% phosphoric acid | 30 ml |
| Bradford stock solution | 30 ml |
| Deionized water | 425 ml |

Preparation of solution for nitric oxide radical scavenging assay

10 mM sodium nitroprusside (SNP) in PBS pH 7.2

0.29 g of sodium nitroprusside was dissolve in 100 ml PBS pH 7.2

0.33% (w/v) sulfanilamide solution

0.33 g of sulfanilamide was dissolved in 5% phosphoric acid, and adjusted total volume to 100 ml by deionized water.

0.1% (w/v) N- (1-Naphthyl) ethylenediamine dihydrochloride (NED)

0.1 g of NED was dissolved in 100 ml

APPENDIX B

Media: Complete medium

| | |
|--------------------------|------|
| DMEM | 100% |
| Fetal Bovine Serum (FBS) | 10% |

MTT solution

| | |
|----------------------|------|
| 5 mg/ml MTT solution | |
| MTT | 5 mg |
| Deionized water | 1 ml |

LPS 1 µg/ml

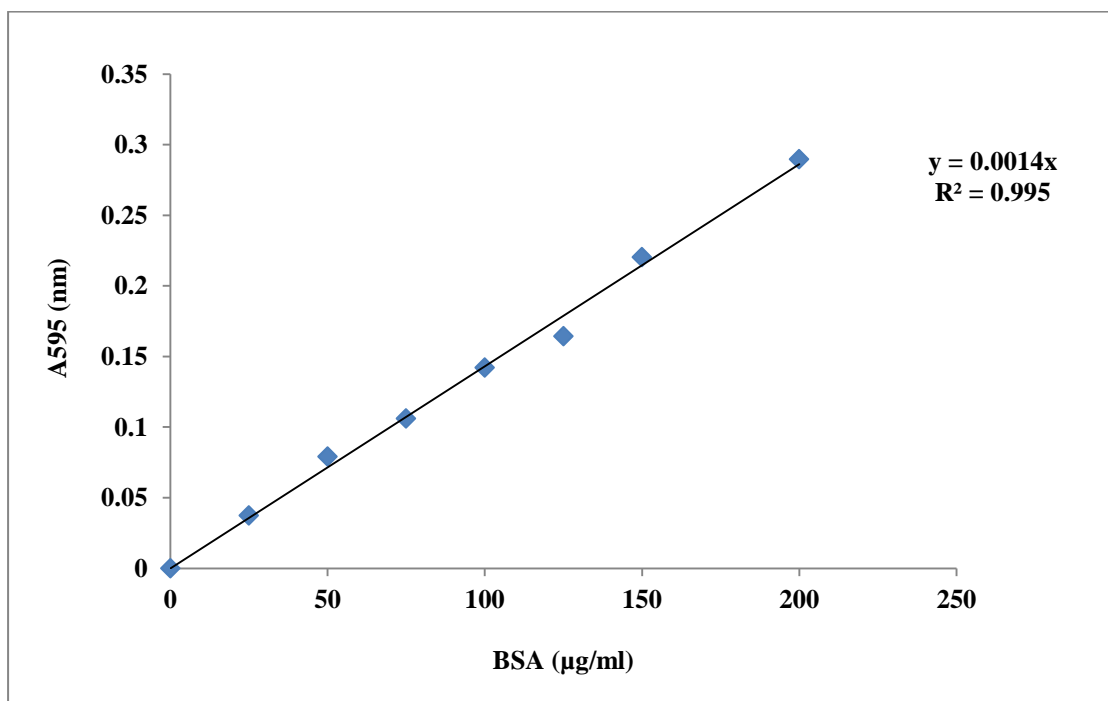
| | |
|---------------|------|
| LPS 500 µg/ml | 4 µl |
| DMEM | 2 µl |

TE buffer

50X Tris-acetic acid EDTA (TAE) buffer solution

APPENDIX C

Standard curve of protein determination by Bradford method



APPENDIX D

Amino acid abbreviations

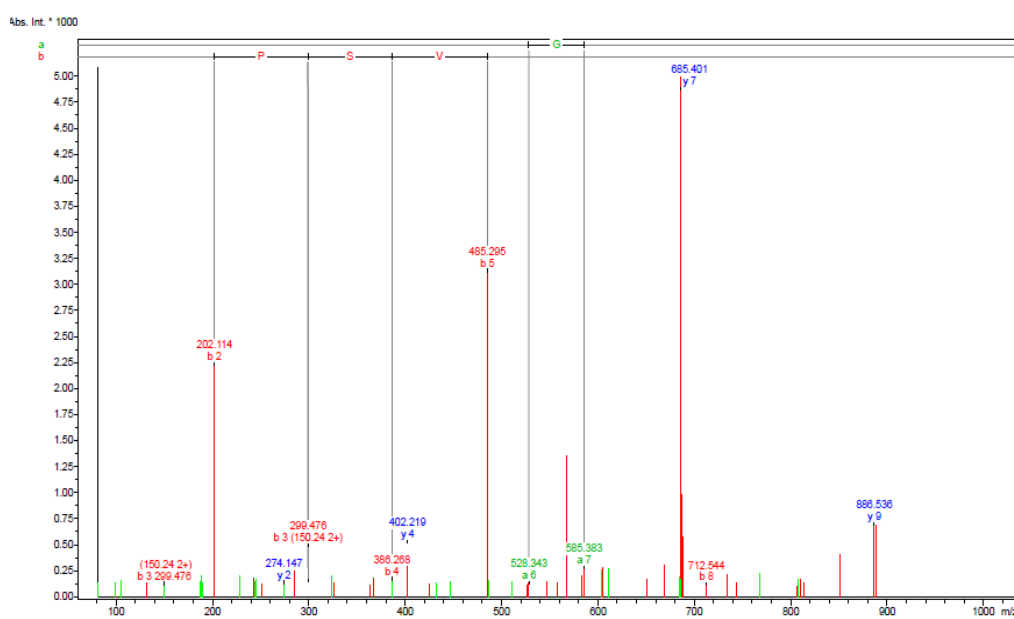
| Amino acid | Three-letter | One-letter |
|---------------|--------------|------------|
| Alanine | Ala | A |
| Arginine | Arg | R |
| Asparagine | Asn | N |
| Aspartic-acid | Asp | D |
| Cysteine | Cys | C |
| Glutamine | Gln | Q |
| Glutamic acid | Glu | E |
| Glycine | Gly | G |
| Histidine | His | H |
| Isoleucine | Ile | I |
| Leucine | Leu | L |
| Lysine | Lys | K |
| Methionine | Met | M |
| Phenylalanine | Phe | F |
| Proline | Pro | P |
| Serine | Ser | S |
| Threonine | Thr | T |
| Tryptophan | Trp | W |
| Tyrosine | Tyr | Y |
| Valine | Val | V |

APPENDIX E

Sequence F2-1 SNPSVAGVR

Mass error 0.061 MH+ (mono): 886.475
Threshold (a.i): 0.000 Tolerance (Da): 0.500
Above Threshold: 60 Assigned Peaks: 11

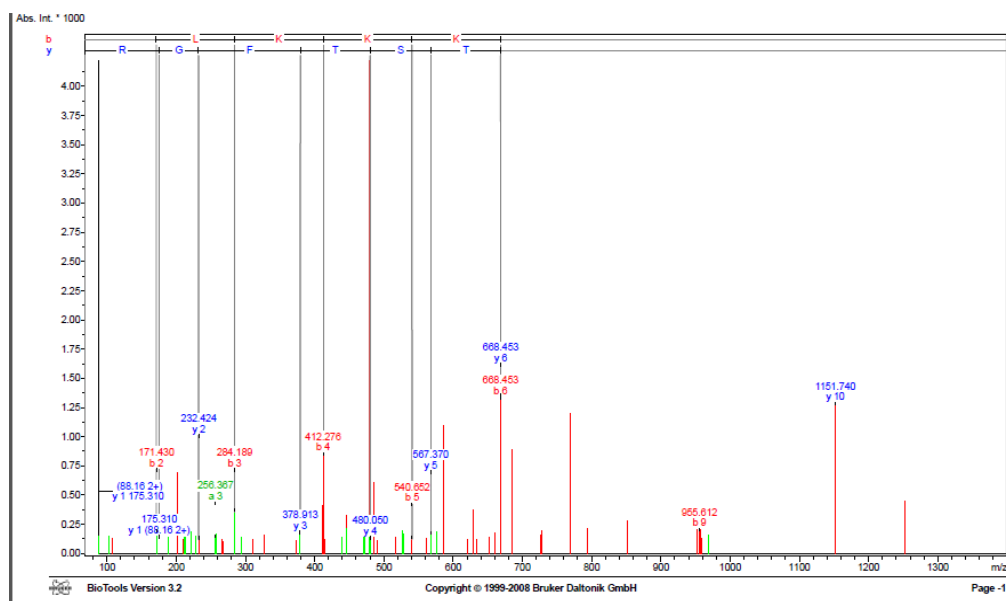
Parentmass 886.536
MH+ (avq): 886.975
Number of peak: 60
Not assigned Peaks: 49



Sequence F2-3 AVLKKKVTSTFGR

Mass error 0.037 MH+ (mono): 1434.880
Threshold (a.i): 0.000 Tolerance (Da): 0.500
Above Threshold: 72 Assigned Peaks: 13

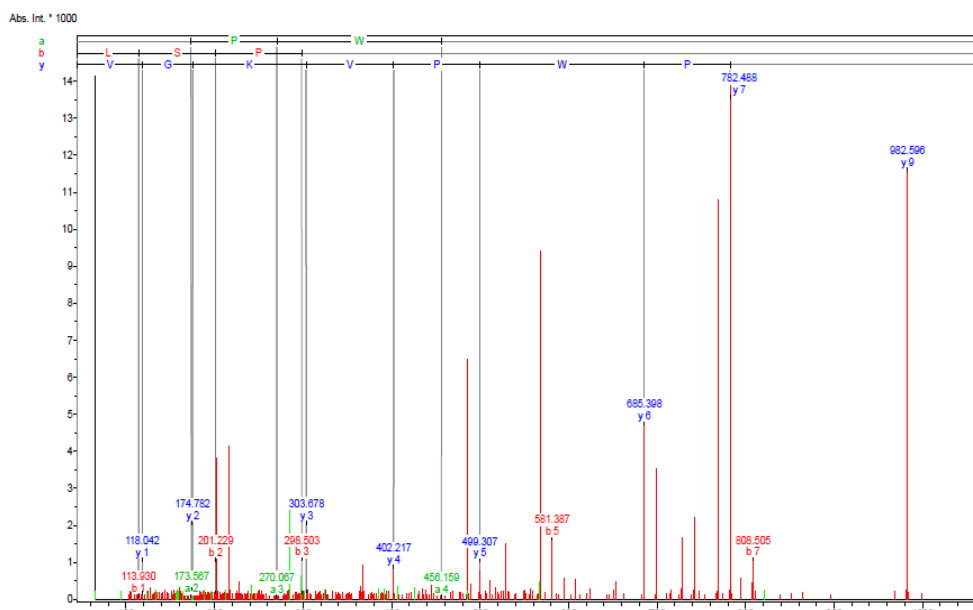
Parentmass 1434.917
MH+ (avq): 1436.737
Number of peak: 72
Not assigned Peaks: 59



Sequence F2-5 LSPWPVKGV

Mass error 0.061 MH+ (mono): 886.475
Threshold (a.i): 0.000 Tolerance (Da): 0.500
Above Threshold: 60 Assigned Peaks: 11

Parentmass 886.536
MH+ (avq): 886.975
Number of peak: 60
Not assigned Peaks: 49



VITA

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Academic presentation;

1) Sukaboon, R., Sangtanoo, P., Saisavoey, T., and Karnchanatat, A. Inhibitory activity against nitric oxide production in macrophage RAW 264.7 by peptide hydrolysate derived from chicken feather meal. In "The 5th International Biochemistry and Molecular Biology Conference" 26-27 May 2016, B.P. Samila Beach Hotel, Bangkok Thailand (Proceeding book)

