ANTIOXIDANT ACTIVITY OF FOOD PROTEINS AND FOOD PROTEIN HYDROLYSATES

By

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Suggested Short Title:

ANTIOXIDANT ACTIVITY OF FOOD PROTEINS

This thesis is dedicated to my beloved grandparents, Hira Singh and Jaswant Kaur; my mother, Surinder Kaur and my sister, Sukhwinder Kaur.

ABSTRACT

The objective of this research was to study the antioxidant activity of soybean protein hydrolysates (SPH) and chickpea protein hydrolysates (CPH) at different concentrations, and to measure the antioxidant activity of fractions collected from the RP-HPLC analysis of SPH and CPH. Protein hydrolysates were prepared by the proteolytic enzyme trypsin. The hydrolysates obtained were subjected to DPPH (1, 1-diphenyl-2 picrylhydrazyl) radical scavenging assay. The SPH and CPH at concentration of 2.5-10 mg/ml showed antioxidant activity of 16.5-32 % and 3.4-26.8 %. SPH and CPH were fractionated by using RP-HPLC on C18 column. The antioxidant activity of four SPH and CPH fractions (F I, F II, F III, and F IV) was measured by using DPPH radical scavenging assay. For SPH, antioxidant activity of F III (47.7 %) was higher than other fractions at protein concentration of 1 mg/mL and for CPH; F II showed maximum antioxidant activity 27.9 % at protein concentration 1 mg/mL. The results from the SDS-PAGE confirmed the hydrolysis of protein samples.

The second part of the study was to measure the impact of high pressure processing (HPP) on the degree of hydrolysis and antioxidant activity of proteins. High pressure processing (HPP) of isolated soybean protein (ISP) and isolated chickpea protein (ICP) was done at 400 MPa and 600 MPa for 5 min and 10 min. The degree of hydrolysis of isolated soybean protein and isolated chickpea protein treated with high pressure processing and with trypsin hydrolysis showed continuous increase from 12.4 to 24.9 % for SPH and 13.6 to 26.2 % for CPH. The DPPH radical scavenging assay showed a more than two fold increase in antioxidant activity of SPH and CPH: 67 % as compared to the 32 % of SPH without HPP and 56.6 % as compared to the 26.8 % of CPH without HPP at concentration 10 mg/mL. These results show that HPP increased the degree of hydrolysis and antioxidant activity of protein hydrolysates.

RÉSUMÉ

Le but principal de cette recherche constituait l'analyse du potentiel antioxydant, à diverses concentrations, d'hydrolysats de protéine de soya (HPS) et d'hydrolysats de protéine de pois chiche (HPP). Les hydrolysats de protéine ont été isolés à l'aide de l'enzyme protéolytique trypsine. Les HPS et HPP démontraient respectivement un potentiel antioxydant de 16.5 à 32% et 3.4 à 26.8 % lorsque présents à des concentrations de 2.5 à 10 mg/mL. L'utilisation d'une colonne C18 a permis de séparer, par CLHP-PI, les HPS et HPP en quatre fractions (F I, F II, F III, et F IV) qui furent dosées avec du DPPH (1,1-diphényl-2-picrylhydrazyle) afin de comparer leur pouvoir de scavenging sur les radicaux. Pour les HPS, le potentiel antioxydant de F III (47.7 %) était supérieur à celui des autres échantillons alors que pour les HPP, 27.9 % (F II) était le seuil maximal. Dans les deux cas, les hydrolysats étaient concentrés à 1mg/mL. L'hydrolyse des échantillons de protéine a été confirmée par SDS-page.

La deuxième partie de l'étude visait à mesurer l'impact de la pascalisation sur le degré d'hydrolyse et le potentiel antioxydant des protéines. Des isolats de protéine de soya (IPS) et de protéine de pois chiche (IPP) ont été traités à haute pression (400 MPa et 600 MPa) pendant 5 et 10 min. Le degré d'hydrolyse des IPS et IPP soumis à la pascalisation et à la trypsin ont démontré une augmentation constante allant de 12.4 à 24.9 % pour les isolats de protéine de soya et de 13.6 à 26.2 % pour les isolats de protéine de pois chiche. L'analyse au DPPH du pouvoir d'épuration des radicaux a montré que le potentiel antioxydant des hydrolysats a plus que doublé, passant de 32 à 67 % pour les HPS et de 26.8 à 56.6 % pour les HPP, lorsqu'ils étaient traités par hautes pressions. Cela démontre que la pascalisation améliore le degré d'hydrolyse et le potentiel antioxydant des hydrolysats de protéines.

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ABBREVIATIONS

| AA | Antioxidant Activity |
|---------|---|
| ABTS | 2, 2'-azinobis (3-ethylbenzothiazaline-6-sulfonic acid |
| BHA | Butylated Hydroxyanisole |
| BHT | Butylated Hydroxytoluene |
| СРН | Chickpea Protein Hydrolysates |
| DH | Degree of Hydrolysis |
| DPPH | 1, 1-diphenyl-2-picrylhydrazyl |
| EPR | Electron Paramagnetic Resonance |
| ESR | Electron Spin Resonance |
| FTC | Ferric Thiocyanate |
| ICP | Isolated Chickpea Protein |
| ISP | Isolated Soybean Protein |
| LDL | Low-density Lipoprotein |
| MetO | Methionine Sulfoxide |
| MW | Molecular Weight |
| NHREP | Neutrase Hydrolysed Rice Endosperm Protein Hydrolysates |
| OPA | o-Phthaldialdehyde |
| ORAC | Oxygen Radical Absorbance Capacity |
| PAGE | Polyacrylamide Gel Electrophoresis |
| RP-HPLC | Reverse Phase- High Performance Liquid Chromatography |
| SDS | Sodium Dodecyl Suplhate |
| SOD | Superoxide Dismutase |
| SPH | Soybean Protein Hydrolysates |
| TBARS | Thiobarbituric Acid Reactive Substances |
| TBHQ | Tert-butyl Hydroquinone |

- TBHQ Tert-butyl Hydroquinone
- TCA Trichloroacetic Acid
- WGPH Wheat Germ Protein Hydrolysates

CHAPTER 1

INTRODUCTION

1.1 General Introduction

Proteins are important components of foods and ingredients of food products. In addition to their nutritional role, proteins can act as functional food ingredients, gelling agents, and emulsifiers; have water and oil binding capacity and foaming characteristics. Proteins can also act as free radical scavengers, chelating agents for transition metals, quenchers of singlet oxygen molecule and decreases the radical damage in biological systems (Zielinski and Kozlowska, 2000). Recently due to concern over the safety of synthetic compounds, interest has been given to naturally occurring antioxidants for the replacement of potentially toxic synthetic antioxidants. As the demand for all natural products is increasing, the new technologies with improved natural antioxidants are also needed. This has led to the new investigations into measuring the antioxidant potential of biologically active peptides from protein hydrolysates (Li et al., 2007). Proteins also have ability to inhibit lipid oxidation. Therefore the oxidative stability of the foods can be increased by protecting the endogenous antioxidant enzymes, enhancing the activity of the proteins by altering the structure, and by using proteins and peptides with antioxidant activity as food additives.

Various food protein hydrolysates, for example pea protein hydrolysates (Aluko et al., 2010), soy protein hydrolysates (Chen et al., 1996), milk protein hydrolysates (Kunio et al., 2000) and wheat germ protein hydrolysates (Zhu et al., 2006) have ability to inhibit lipid oxidation in foods. The antioxidant activity of proteins and protein hydrolysates is mainly due to, inactivation and sequestration of pro-oxidative metals, their ability to scavenge free radicals,

non-radical reduction of lipid hydroperoxides; inhibition of lipid oxidation by altering the physical state of food (Roberto, 2008).

Soybean (*Glycine max*) and chickpea (*Cicer arietinum* L.) are important food crops in Canada and many other countries. Because of their relatively high protein contents (30-46 % for soybean, 14.9-24.6 % for chickpea), both seeds can be good sources of peptides with antioxidant potential. There is only little information on the antioxidant and free radical scavenging of soybean and chickpea protein. Furthermore, there is no information on the effect of high pressure processing on the solubility and functional properties of these proteins. Therefore it is of interest to study the antioxidant activity and effect of high pressure processing on soybean and chickpea proteins.

1.2 Objectives of Research

The overall objective of this research was to study the antioxidant activity of proteins and protein hydrolysates. The specific objectives of this work were:

- 1. To investigate the tryptic hydrolysis of soybean and chickpea proteins and to determine the antioxidant activity of soybean and chickpea protein hydrolysates.
- 2. To separate soybean and chickpea protein hydrolysates by using RP-HPLC and to collect fractions at different retention time from RP-HPLC analysis and to investigate the protein fraction with high antioxidant activity from RP-HPLC analysis.
- 3. To characterize soybean and chickpea protein hydrolysates using SDS-PAGE.
- 4. To evaluate the impact of high pressure processing on the degree of hydrolysis and antioxidant activity of soybean and chickpea proteins.

CHAPTER 2

LITERATURE REVIEW

2.1 Proteins and Peptides as Antioxidants

There are several studies that demonstrate the ability of proteins and peptides to inhibit lipid oxidation in food products. Proteins from various sources like milk protein (Diaz et al., 2003), soy protein (Park et al., 2005), wheat protein (Zhu et al., 2005), rice protein (Zhang et al., 2009), potato protein (Wang and Xiong, 2005), and corn protein (Kong and Xiong, 2006) exhibit antioxidant activity. Procine blood plasma (2.5 %) retards the formation of thiobarbituric acid reactive substances (TBARS) in both salted ground pork (Faraji et al., 1991) and cooked ground beef (Shantha and Decker, 1995) due to the presence of antioxidant proteins such as albumin and transferrin. Whey protein concentrate act as antioxidant in cooked beef (Shantha and Decker, 1995), and soy and whey proteins inhibit lipid oxidation in cooked pork patties containing 2 % protein (Pena-Ramos and Xiong, 2003). Whey proteins have also been found to inhibit lipid oxidation in oil-in-water emulsions (Taylor and Richardson, 1980; Allen and Wrieden, 1981a and b; Donnelly et al., 1998; Tong et al., 2000; Elias et al., 2005).

Peptides resulting from hydrolysis of food proteins also act as antioxidants and inhibit lipid oxidation in foods. Hydrolysates of whey, casein, soy, and egg yolk proteins inhibit lipid oxidation in various muscle foods, such as beef, pork, and tuna (Sakanaka and Tachibana, 2006; Diaz and Decker, 2005; Sakanaka et al., 2005, Pena-Ramos and Xiong, 2003). According to Chan and Decker (1994), carnosine and anserine are histidine-containing dipeptides found in skeletal muscle that exhibit antioxidant properties. Lipid oxidation and myoglobin discolouration in muscle foods can be inhibited by addition of carnosine to muscle foods (Calvert and Decker,

1992; Decker and Crum, 1991). Wu et al., (2011) reported that free aromatic amino acids, tryptophan and tyrosine from egg yolk have ability to inhibit lipid oxidation in foods. Aluko et al., (2009) showed that tryptophan released from mother's milk exhibit antioxidant properties.

2.2 Antioxidant Mechanisms of Proteins

2.2.1 Lipid Oxidation

Autooxidation is a process in which oxygen attacks at the susceptible lipids leading to complex chemical changes, resulting in the rancidity and production of off flavours in the food. The mechanism of oxidative rancidity or autooxidation of lipids has been well established (Bateman et al., 1953; Labuza, 1971; Eriksson, 1987). Atmospheric oxygen reacts spontaneously with organic compounds and degrades the structure of organic compounds, which is mainly responsible for the loss in quality of the chemical products of industrial importance. The spontaneous oxidative reaction in food systems can result in the deterioration of lipids (Jadhav et al., 1996) because fats, oils and vitamins contains different amounts of unsaturated linkages in their hydrocarbon chains and these unsaturated sites are susceptible to oxidation (Coulter, 1988). Free-radical chain reaction is the direct reaction of a lipid molecule with a molecule of oxygen (Jadhav et al., 1996) and the mechanism of autooxidation can be described in three steps: chain initiation, chain propagation and chain termination (Bateman et al., 1953; Labuza, 1971; Eriksson, 1987).

The chain initiation occurs with the formation of free radicals.



 $R^{\circ} + O_2 \longrightarrow ROO^{\circ}$

The formation of lipid free radical R° mainly occurs due to light or heat, trace metals, irradiation and alkaline conditions. After oxidation reaction, lipid hydroperoxides break down to yield radicals.

ROOH
$$\longrightarrow$$
 RO° + HO°, and
2ROOH \longrightarrow RO° + ROO° + H₂O

In chain propagation reaction, the lipid radicals are converted into peroxy radicals by reacting with molecular oxygen (Jadhav et al., 1996).

$$R^{\circ} + O_2 \longrightarrow ROO^{\circ}$$
$$ROO^{\circ} + RH \longrightarrow ROOH + R^{\circ}$$

Chain initiation reaction occurs due to the presence of lipid peroxy radicals (ROO°) and it results into the formation of lipid hydroperoxides and lipid free radicals. The chain propagation reaction continuous as long as unsaturated lipid molecules are available. Chain termination occurs when there is a reduction in the amount of unsaturated lipid molecules present, radicals bond to one another, forming a stable non-radical compound. Chain propagation stage can be prevented by the presence of natural antioxidants, which break the chain reaction (Coulter, 1988). Vitamin E is a major lipid soluble chain breaking antioxidant and prevents lipid oxidation by scavenging or converting free radicals into less reactive forms.

$$TH + X^{\circ} \longrightarrow T^{\circ} + XH$$

 $X^{\circ} =$ Free radical, TH = Tocopherol

 $T^{\circ} = Tocopheroxyl radical$

Tocopheroxyl radical is further converted into tocopherol by using water soluble antioxidants such as ascorbate (Chan et al., 1991; Mukai et al., 1990).

2.2.2 Sources of Antioxidants

2.2.2.1 Synthetic Antioxidants

Synthetic antioxidants (Figure 2.1) are mainly phenolic antioxidants and include butylated hydroxyanisole (BHA), tert-butyl hydroquinone (TBHQ), butylated hydroxytoluene (BHT), and propyl gallate. The use of synthetic antioxidants is limited to 100-200 ppm of BHA, BHT, or TBHQ or 200-500 ppm of the propyl gallate for the stabilization of fats and oils.



BHA butylated hydroxyanisole



OН

- СН₂— СН₃

BHT butylated hydroxytoluene





Propyl gallate

Figure 2.1 Structures of BHA, BHT, TBHQ, and Propyl Gallate (Madhavi et al., 1996).

2.2.2.2 Natural Antioxidants

In recent years, the demand for "all natural" products is increasing because of the increasing limitations on the use of synthetic antioxidants and enhanced public awareness of health issues. Natural antioxidants are "generally recognized as safe" (GRAS) to be use in food

components. Many natural food components like oils and oilseeds, proteins and protein hydrolysates, fruits and vegetables, oat and rice bran, spices, herbs and tea have antioxidant properties. Natural antioxidants from these food components provide oxidative stability to the food product.

As antioxidative properties of proteins contribute to the endogenous antioxidant capacity of foods, therefore proteins can also be used as potential antioxidant additives. Lipid oxidation in foods can be inhibited by using proteins by their biologically designed mechanisms (e.g. antioxidant enzymes and iron-binding proteins) or by nonspecific mechanisms (e.g. scavenging of free radicals and reactive oxygen species). Overall, the antioxidant activity of proteins is mainly due to interactions between their ability to inactivate reactive oxygen species, chelate prooxidative transition metals, scavenge free radicals, and reduction of hydroperoxides.

2.2.3 Antioxidant Enzymes

2.2.3.1 Peroxidases

Hydrogen and lipid peroxides are commonly found in foods where they can decompose to form free radicals. For example, reduced state of transition metals (e.g., Fe and Cu) decomposes hydrogen peroxide to the hydroxyl radical, an extremely reactive radical that can oxidize lipids, and most organic matter at diffusion-limited rates. Catalase is a heme-containing enzyme found in various biological systems that catalyzes the conversion of hydrogen peroxide to water by the following pathway:

$$2H_2O_2 \longrightarrow 2H_2O + O_2$$

Ascorbate peroxidase also removes hydrogen peroxide in plants by following mechanism:

Glutathione peroxidase (GSH-Px) is present in various biological tissues. Glutathione also inactivates both lipid and hydrogen peroxides. Glutathione peroxidase reacts with hydrogen peroxide and it forms water and oxidized glutathione (GSSG).

$$H_2O_2 + 2GSH \rightarrow 2H2O + GSSG$$

2.2.3.2 Superoxide Dismutase

The production of superoxide anion occurs by the addition of an electron to the molecular oxygen. Superoxide anion is prooxidative due to its ability to reduce transition metals, and form perhydroxyl radical which can catalyze lipid oxidation under acidic conditions (pH< 4.8) (Decker, 2002). Isoforms of superoxide dismutase (SOD) contain manganese or copper plus zinc in the active site, and both catalyze the conversion of superoxide anion to hydrogen peroxide by the following reaction:

 $2O_2^- + 2H + \longrightarrow O_2 + H_2O_2$

2.2.4 Inactivation and Sequestration of Prooxidative Metals by Proteins

Transition metals such as iron and copper are important prooxidants in food lipids. These are capable of catalyzing the reduction of hydroperoxides to reactive radical species as shown in the following reaction:

$$Metal^{n+} + Lipid-OOH \longrightarrow Metal^{(n+1)+} + OH^{-} + Lipid - O$$

Proteins can inhibit the oxidative reaction by chelating the metals and by changing the physical location of transition metals (e.g. partitioning metals away from oxidatively labile lipids

or hydroperoxides), forming insoluble metal complexes, reducing the chemical reactivity of the transition metals, and by sterically hindering the interaction of metals and dispersed lipids (Diaz et al., 2003). Various food proteins function as binding prooxidant metals. Proteins whose biological function is to chelate and store or transport catalytically inactive metals are found in most biological systems. Table 2.1 shows the metal binding protein antioxidants.

 Table 2.1 Metal binding protein antioxidants (Symons and Gutteridge, 1998)

| Protein | Function |
|--------------|--|
| Ferritin | Binds and stores ferric ions within helical fold |
| Transferrin | Binds ferric ions |
| Lactoferrin | Binds ferric ions at low Ph |
| Haptoglobins | Binds hemoglobulin |
| Hemopexin | Binds heme |

Ferritin is mainly known as the iron storage protein, it can bind up to 4,500 ferric ions (Fennema, 1996; Weinberg, 1990). Transferrin, lactoferrin, ovotrasferrin and ferritin controls iron reactivity by binding iron in its less active ferric state. Ferritin mainly reduces iron reactivity by interfering with its redox cycling capacity. In the case of ferritin, ferrous (Fe⁺²) ions are oxidized to ferric (Fe⁺³) oxidation state after binding to a specific site located inside the subunit helical fold (i.e. the ferroxidase centre) (Arosio and Levi, 2002). After oxidation, iron migrates to the protein's cavity where it aggregates to form a ferric hydroxide core (Arosio et al., 2002; Carrondo, 2003). Thus, iron is maintained in its ferric state after sequesteration, and is unavailable to participate in redox reaction (e.g. lipid hydroperoxide decomposition). Lactoferrin is also an effective antioxidant in foods. (Satue-Gracia et al., 2000) found that lactoferrin inhibits

lipid oxidation in infant formula. Lactoferrin also inhibits lipid oxidation in milk and mayonnaise (Nielsen et al., 2004). Haptoglobins and hemopexins are extracellular antioxidants that are capable of binding hemoglobulin and heme, respectively (Symons and Gutteridge, 1998).

Copper is also a prooxidative transition metal mainly present in food at lower concentration than iron. Copper is a more effective catalyst than iron in decomposition of hydroperoxide (Halliwell and Gutteridge, 1990). Proteins also form complexes with copper and thus can affect lipid oxidation. In various biological tissues, copper is bound by proteins such as serum albumin (one cupric ion per protein) and ceruloplasmin (six cupric ions per protein). These proteins inhibit lipid oxidation as in copper-catalyzed oxidation of low-density lipoprotein (LDL), where oxidation rates are inversely proportional to the concentration of bovine serum albumin (BSA) (Bourdon et al., 1999; Schnitzer et al., 1997). Various amino acid residues such as histidine, glutamic acid, aspartic acid, threonine, and phosphorylated serine are known to bind metals. The ability of a protein to chelate metals is mainly dependent on pH. For example, if the pH value is above the isoelectric point, a net anionic charge will be established on a protein. This results in electrostatic attraction between the protein and cationic transition metals, which ultimately inhibits lipid oxidation reactions.

Various proteins have been reported to bind prooxidative transition metals. These include casein (Diaz et al., 2005; Diaz et al., 2003), whey protein (Faraji et al., 2004; Tong et al., 2000), soy proteins (Faraji et al., 2004), bovine serum albumin (Villiere et al., 2005), zein (Kong and Xiong, 2006), and potato protein (Wang and Xiong, 2005).

The metal-catalyzed decomposition of lipid hydroperoxides is thought to be dominant oxidative pathway in processed foods, and in particular oil-in-water emulsions (McClements and

Decker, 2000). This reaction mainly occurs at the surface of the emulsion droplet because lipid hydroperoxides are surface active and migrate to the water-oil interface. One way in which proteins can inhibit lipid oxidation reaction in oil-in-water emulsions is by changing the physical location of aqueous prooxidants, in this way the metals are moved away from the surface of emulsion droplet, reducing its ability to decompose lipid hydroperoxides.

2.2.5 Inhibition of Lipid Oxidation by Proteins by Altering the Physical State of Foods

In various food products, lipid oxidation reactions do not occur randomly but instead occur at specific locations based on the physicochemical properties of the food product. For example, in the case of metal-catalyzed oxidation in lipid dispersions, surface active lipid hydroperoxides concentrate at oil-water interfaces. Aqueous phase transition metals can catalyze the decomposition of these hydroperoxides and produces radical species. Proteins inhibit lipid oxidation in such lipid dispersions by hindering access of metals to the oil-water interface through electrostatic repulsion. In protein-stabilized oil-in-water emulsions where lipid oxidation rates are slower at pH values below the isoelectric point of the proteins. A net cationic charge is established at the water-oil interface, repelling iron and physically hindering the ability of cations from binding to the droplet interface (Donnelly et al., 1998; Hu et al., 2003a; Hu et al., 2003b; Kellerby et al., 2006).

Lipid oxidation rate is also affected by thickness or existence of a thick emulsion droplet interface that physically inhibits the ability of iron to access lipid hydroperoxides at the droplet surface. In protein-stabilized oil-in-water emulsions, protein could form a thick interfacial layer that inhibits metal-lipid interactions. For example, casein forms an interfacial layer around dispersed oil droplets up to 10 nm compared to 1-2 nm for whey proteins (Dalgleish et al., 1995).

2.2.6 Scavenging Free Radicals and Reactive Oxygen Species by Proteins

Proteins also have the ability to scavenge free radicals and reactive oxygen species. Many studies have been conducted to check the free radical scavenging activity of proteins. Figure 2.2 shows the reaction of the DPPH with the antioxidant.









Figure 2.2 DPPH chemical structure and its reaction with a scavenger, indicated by A-H (Roberto, 2008).

These studies have typically employed the use of free radical generators, and a determination is made if a given protein can scavenge the radical species produced. For example, a radical scavenging assay using 1, 1-diphenyl-2-picrylhydrazyl (DPPH) has shown that rice endosperm protein (Zhang et al., 2009), wheat germ protein (Zhu et al., 2005) and egg yolk proteins (Sakanaka and Tachibana, 2006) can scavenge free radicals.

A second radical scavenging assay using oxygen radical absorbance capacity (ORAC) is a common method that measures the ability of a free radical scavenger to quench peroxyl radicals generated by 2, 2-azo-bis (2-methylpropionamidine) dihydrochloride (AAPH) and protect the fluorescent marker, fluorescein. Using ORAC, β -lactoglobulin (Elias et al., 2006) and casein (Diaz et al., 2005) were shown to scavenge peroxyl radicals. Radical scavenging activity has also been measured by using free radicals generated by 2, 2'-azinobis (3ethylbenzothiazoline-6sulfonic acid) (ABTS) to show that zein (Kong and Xiong, 2006) and potato (Wang and Xiong, 2005) proteins quench free radicals. Proteins and free radicals interactions can also be measured by electron paramagnetic resonance (EPR) (Davies and Hawkins, 2004) or electron spin resonance (ESR). Using electron paramagnetic resonance, bovine serum albumin, β -lactoglobulin, and lactoferrin were shown to scavenge free radicals generated from cumene hydroperoxide and iron (Pazos et al., 2006).

2.2.7 Non-Radical Reduction of Lipid Hydroperoxides

Proteins also have ability to reduce lipid hydroperoxides to relatively non-reactive lipid hydroperoxides by non-radical reactions (Garner et al., 1998a; Garner et al., 1998b; Pryor et al., 1994; Pryor and Squadrito, 1995). The mechanism for the reduction of lipid hydroperoxides to lipid hydroxides is as following:



Figure 2.3 Proposed two-electron reduction of lipid hydroperoxide (LOOH) by thioethercontaining side chain of methionine (Garner et al., 1998a; Garner et al., 1998b; Panzenbock and Stocker, 2005).

The mechanism for the reduction of lipid hydroperoxides to lipid hydroxides involves a direct two-electron transfer from the sulfide of the methionine's thioether group, resulting in the oxidation of methionine to methionine sulfoxide (MetO) (Garner et al., 1998a; Garner et al, 1998b; Panzenbock and Stocker, 2005). In effect, methionine residues reduce lipid hydroperoxide to non-reactive species.

2.3 Methods for Measuring Antioxidant Activity of Proteins

There are various methods used to measure the antioxidant activity of proteins and protein hydrolysates. These mainly includes the ability of proteins to scavenge free radicals like 1,1-diphenyl-2-picrylhydrazyl (DPPH), superoxide radical, to test the reducing power of proteins, to test the inhibition of linoleic acid autooxidation.

2.3.1 DPPH Radical-Scavenging Assay

The scavenging effect of proteins and protein hydrolysates fractions on 1, 1-diphenyl-2picrylhydrazyl (DPPH) free radical is mainly measured by according to the method of shimada et al., (1992) with little modification. Firstly, a protein solution with different concentrations is prepared and after centrifugation at 8000 RPM for 20 min, 2 ml of the upper layer is added to the 2 ml solution of 0.1 mM DPPH dissolved in 95 % ethanol. The mixture is then shaken and left for 30 min in dark at room temperature, and the absorbance of resulting solution is read at 517 nm. A lower absorbance represents a higher DPPH scavenging activity. The scavenging effect is mainly expressed as shown in the following equation:

DPPH scavenging activity (%) = [(Control absorbance-Sample absorbance)/Control absorbance] x 100

Control consists of 2 ml of ethanol and 2 ml of 0.1 mM DPPH.

2.3.2 Superoxide Radical Scavenging Assay

The superoxide radical scavenging activity of proteins and protein hydrolysates is measured at 25°C using the spectrophotometric monitoring of the inhibition of pyrogallol autoxidation as described by Marklund and Marklund (1974) with some modifications.

Superoxide radical scavenging assay is dependent on the reducing activity of a test compound by an O_2^- dependent reaction, which releases chromophoric products.

Firstly, protein solution with different concentrations is prepared, and 0.1 ml of the protein solution is added into 2.8 ml Tris-Hcl-EDTA buffer (0.1 M, pH 8.0), and the mixture is shaken and heated at 25° C for 10 min. After 10 min of heating, the reaction is initiated by adding 0.1 ml of pyrogallol solution (3 mM) and then optical density is measured at 325 nm using a spectrophotometer. The scavenging activity is given as following:

Scavenging activity (%) = $[(CA-SA)/CA] \times 100$

CA = Control absorbance, SA = Sample absorbance

2.3.3 Test of Inhibition of Linoleic Acid Autoxidation by Proteins

The antioxidant activity of proteins and protein hydrolysates with different periods of incubation is mainly measured in a linoleic acid model system according to the method of Osawa and Namiki (1985) with little modification. 10 mg of protein or protein hydrolysate is dissolved in 10 ml of 50 mM phosphate buffer (pH 7.0) and then the mixture is added to a solution of 0.15 ml linoleic acid and 10 ml 99.5 % ethanol. Total volume is then adjusted to 25 ml with double distilled water. The mixture is than incubated in a conical flask at 40°C in a dark room for 6 or 8 days, and degree of oxidation is mainly evaluated by measuring the ferric thiocyanate (FTC) values.

FTC value is mainly measured according to the method of Mitsuta et al., (1996). According to this method, 100 μ l of the above reaction is mixed with 4.7 ml of 75 % ethanol, 0.1 ml 30 % ammonium thiocyanate and 0.1 ml of 0.02 M ferrous chloride solution in 3.5 % HCL.

After the reaction time of 3 min, the FTC value is measured by reading the absorbance at 500 nm. This absorbance is measured daily up to 6-8 days to know the inhibition of linoleic acid autooxidation.

2.3.4 Test for Reducing Power

The reducing power of proteins is mainly measured according to Oyaizu (1988) with little modification. Different concentrations of proteins are added to 2.5 ml of 0.2 M phosphate buffer (pH 6.6) and 2.5 ml of 1 % potassium ferricyanide. The mixture is incubated at 50°C for 20 min, and then 2.5 ml of 10 % trichloroacetic acid (TCA) is added to the reaction. After the reaction, centrifugation is done at 6500 g for 10 min. After centrifugation, 2.5 ml of the upper layer obtained is mixed with 2.5 ml of double distilled water and 0.5 ml of 0.1 % ferric chloride in 10 ml test tube. After 10 min of reaction time, the absorbance of the resulting solution is measured at 700 nm. Increased absorbance of the reaction mix indicates increased reducing power.

2.3.5 Antioxidant Activity of Food Protein Hydrolysates

In recent years, interest in utilizing natural antioxidants has increased substantially (Shahidi et al., 2006). Due to this the interest has led to new investigations to know the antioxidant potential of biologically active peptides from protein hydrolysates such as wheat protein (Zhu et al., 2005), rice protein (Zhang et al., 2009), fish protein (Kim et al., 2001; Wu et al., 2003), soy protein (Chen et al., 1996; Moure et al., 2006), and milk casein (Kunio et al., 2000). Enzymatic hydrolysis is widely applied to improve the functional and nutritional properties of food proteins. Protein hydrolysates derived from enzymatic hydrolysis have various

physiological activities such as antioxidant, antimicrobial, antihypertensive, and mineral binding (Korhonen and Pihlanto, 2003).

2.3.5.1 Antioxidant Activity of Wheat Germ Protein Hydrolysates

Antioxidant and free radical-scavenging activities of wheat germ protein hydrolysates was studied by Zhu et al., (2005). The main by-product of the oil extraction process is a defatted wheat germ meal, which has protein content up to 30 % (Ge et al., 2000). The authors used alcalase 2.4L for the enzymatic hydrolysis of wheat germ protein. Protein hydrolysis of 10 % (w/v) protein solution with alcalase was carried out at 50°C and pH 8.0 for 6 hours. The enzyme-substrate ration (E/S) was 0.4 AU/g of protein. The authors measured the antioxidant activity of the wheat germ protein hydrolysates by measuring the inhibition of linoleic acid autoxidation, scavenging effect of 1, 1-diphenyl-2-picrylhydrazyl (DPPH), determination of superoxide radical-scavenging activity, and determination of reducing power.

Figure 2.4 shows the antioxidative activity of wheat germ protein hydrolysate measured in linoleic acid emulsion system and compared with those of α -tocopherol and BHT by Zhu et al., (2005). From this the authors conclude that wheat germ protein hydrolysates have antioxidant activity as comparable to α -tocopherol.



Figure 2.4 The antioxidant activity of WGPH. WGPH was incubated in a linoleic acid emulsion system for 6 days. The degree of linoleic acid oxidation was measured by the ferric thiocyanate method at every 24 h interval. Butylated hydroxytoluene and α -tocopherol were used as positive controls (Zhu et al., 2005).

The scavenging effect on 1, 1-diphenyl-2-picrylhydrazyl (DPPH) was also measured to check the antioxidant activity of wheat germ protein hydrolysates. The DPPH free radicals are stable in ethanol and shows maximum absorbance at 517 nm. When DPPH radicals react with antioxidant, the radicals would be scavenged and the absorbance is reduced. Thus, the DPPH radicals were widely used to investigate the scavenging activity of protein hydrolysates. Figure 2.5 shows the results of scavenging DPPH radical ability of wheat germ protein hydrolysates (WGPH) and they concluded that with the increase in the concentration of wheat protein hydrolysates the antioxidant activity increase:



Figure 2.5 Scavenging effect on DPPH free radical of different concentrations of WGPH. Butylated hydroxytoluene (BHT) and ascorbic acid were used as positive controls to compare the scavenging effect of WGPH. Regression equation was obtained from linear regression of the concentrations of WGPH and DPPH radical-scavenging effects. Each value is expressed as mean S.D. (n = 3) (Zhu et al., 2005).

For the radical scavenging assay, reducing power assay and superoxide radicalscavenging assay, Zhu et al., (2005) found that with the increase in the concentration of wheat germ protein hydrolysate the antioxidant activity increases. The authors concluded that wheat germ protein hydrolysates can be used as a functional food ingredient in pharmaceutical and food industries.

2.3.5.2 Antioxidant Activity of Rice Endosperm Protein Hydrolysates

Zhang et al., (2009) studied the antioxidant activity of the rice endosperm protein hydrolysate. For the hydrolysis of the rice endosperm protein, they used five different enzymes
Alcalase, Chymotrypsin, Neutrase, Papain, and Flavorase based on their optimum hydrolysis conditions.

After 4 hrs of hydrolysis, the reaction was heated to 85° C for 10 min to inactivate the enzyme. The results of the hydrolysis showed that degree of hydrolysis (DH) of rice endosperm protein by Neutrase was 20 %, by Chymotrypsin (22.23 %), Alcalase (16.24 %), Papain (11.72 %), Flavorase (8.23 %) respectively. The authors measured the antioxidant activity of the rice endosperm protein hydrolysates by measuring the DPPH radical scavenging assay, superoxide radical scavenging assay, hydroxyl radical scavenging assay, reducing power, ferrous ion-chelating activity and test of inhibition of linoleic acid autoxidation.

The antioxidant activities of different hydrolysates were evaluated by the DPPH radical scavenging assay, and compared with that of α -tocopherol, a commercial antioxidant commonly used in the food industry. Zhang et al., (2009) found that the protein hydrolysates prepared from the enzyme Neutrase showed maximum antioxidant activity as shown in Figure 2.6.



Figure 2.6 DPPH radical scavenging activities of the various hydrolysates from the defatted rice endosperm protein (REP) at different concentrations. α -tocopherol was used as a positive control. Each value is expressed as mean \pm SD (Zhang et al., 2009).

Figure 2.7 shows the results of Zhan et al., (2009), they found that the percentage inhibition of autooxidation in linoleic acid system by Neutrase hydrolysed rice endosperm protein (NHREP) was 82.09 %, similar to that of α -tocopherol (86.59 %) but lower than that of BHT (99.69 %), respectively, on day 5.



Figure 2.7 Inhibition of linoleic acid autoxidation by the Neutrase hydrolysate from rice endosperm protein (NHREP). The degree of linoleic acid oxidation was measured by the ferric thiocyanate (FTC) method at every 24 h interval. BHT and a-tocopherol were used as positive controls. Vertical bars indicate mean values \pm SD. (Zhang et al., 2009).

Zhang et al., (2009) in their study concluded that Neutrase hydrolysed rice endosperm protein (NHREP) can be used as suitable natural antioxidant to prevent oxidation reactions in food processing and become ingredient of functional foods.

2.4 High Pressure Processing of Proteins

High pressure processing causes unfolding of proteins depending upon protein type, processing conditions and applied pressure. Solubility of proteins can be increased by using high pressure processing (Thakur and Nelson, 1998). High pressure processing ruptures covalent interactions within protein molecules and further cause's reformation of inter molecular bonds. Different types of interactions are responsible for secondary, tertiary and quaternary structure of proteins. The quaternary structure is mainly held by hydrophobic interactions that are sensitive to pressure. Even tertiary structure show considerable changes beyond 200 MPa. High pressure processing can increase the degree of hydrolysis and antioxidant activity of proteins.

2.5 Increasing the Antioxidant Activity of Proteins

Antioxidant activity of proteins can be increased by changing the protein concentration, reactivity, and physical structure. Many protein antioxidant mechanisms like free radical scavenging, metal chelation, hydroperoxide reduction are dependent on the amino acids composition. However, the antioxidant activity of these amino acids residues is limited by the tertiary structure of the protein, as many amino acids with antioxidant potential can be buried within the protein core where they are not accessible to prooxidants.

Antioxidant activity of proteins can be increased through the disruption of tertiary structure (i.e. partial denaturation), in this way it increases the accessibility of the amino acids with antioxidant potential to the prooxidants. Taylor and Richardson (1980) determined the antioxidant activity of the heated skim milk in a methyl linolate emulsion with hemoglobin used as a lipid oxidation catalyst. Taylor and Richardson (1980) showed that heat treatment (70 to 130° C for up to 30 min) increased the antioxidant activity of skim milk. Heat treatment resulted in an increase in reactive sulfhydryls (starting reactive sulfhydryl concentration of 48 μ M was increased to 71 μ M after heating at 130° C for 30 min).

Enzymatic hydrolysis of proteins also increases the exposure of antioxidant amino acids. Increased antioxidant activity in hydrolyzed proteins has been reported for zein (Kong and Xiong, 2006), dairy (Ostdal et al., 1999; Rival et al., 2001; Pena-Ramos and Xiong, 2003; Hernandez-Ledesma et al., 2005; Sakanaka et al., 2005; Diaz et al., 2005; Elias et al., 2005), potato (Wang and Xiong, 2005), gelatin (Park et al., 2005), egg yolk (Sakanaka and Tachibana, 2006), wheat (Zhu et al., 2005) and rice (Zhang et al., 2009) proteins. The observed increase in antioxidant activity due to hydrolysis may result directly from increased solvent exposure of amino acids. The higher solubility of proteins increased the free radical scavenging activity (Ostdal et al., 1999; Rival et al., 2001; Wang and Xiong, 2005; Park et al., 2005; Hernandez-Ledesma et al., 2005; Sakanaka et al., 2005; Diaz and Decker, 2005; Elias et al., 2006; Kong and Xiong, 2006; Sakanaka and Tachibana, 2006). Enzymatic hydrolysis of proteins is an important process to improve the physical, chemical, functional, and nutritional properties of original proteins. Enzymatic hydrolysis is an effective method to prepare active peptides, which possess antioxidative properties. Enzymatic hydrolysis of proteins is mainly done by using various enzyme depending on the substrate. Various enzymes, for e.g. Alcalase, Typsin, Chymotrypsin, Pepsin, Neutrase etc are used for the hydrolysis of proteins. The hydrolysis is carried out at specific temperature conditions of enzyme, and hydrolysis is stopped by heating the reaction mixture at high temperature to inactivate the enzyme.

CHAPTER 3

MATERIALS AND METHODS

3.1 Materials

A commercial soybean isolate was obtained by ADM Protein Technologies Inc. (Decatur, Illinois). Chickpea protein isolate prepared by Chang (2010). Trypsin type IX-S from procine pancreas, E.C. 3.4.21.4; activity 13100 units/mg protein, DPPH (1, 1-diphenyl-2-picrylhydrazyl) and OPA (*o*-phthaldialdehyde) were purchased from Sigma-Aldrich, Canada. All other chemicals used were of analytical and HPLC grade.

3.2 Preparation of Soybean and Chickpea Protein Hydrolysates

3.2.1 Enzymatic Hydrolysis of Protein Isolates and HPP treated Proteins

The enzymatic hydrolysis of isolated soybean and chickpea protein was carried out by using the method described by Chanput et al., (2009) and Adebiyi et al., (2008). Dispersions of soybean and chickpea protein (1 %, w/v) were brought to pH 8.0 with 2N NaOH under mixing and were incubated with trypsin at 1:20 enzyme to protein ration, in 50 mM sodium phosphate buffer at 37° C for 2 h. At time intervals between 0 to 120 min, tubes of the digest were removed from the water bath at 0, 15, 30, 45, 60 and 120 min. The enzyme action was stopped by heating at 95° C for 10 min. The protein hydrolysates were then centrifuged at 8,000x*g* for 30 min, and the supernatant was lyophilized. Figure 3.1 shows the steps involved in the enzymatic hydrolysis, antioxidant activity assay and characterization of soybean and chickpea protein.



Figure 3.1 Steps involved for enzymatic hydrolysis, antioxidant activity assay and characterization of soybean and chickpea protein.

3.2.2 Determination of Degree of Hydrolysis

The degree of hydrolysis (DH) was measured by using the o-phthaldialdehyde (OPA) method described by Church et al., 1983. The OPA reagent was prepared by combining the 25 mL 100 mM sodium tetraborate, 2.5 ml 20 % SDS (w/w) and 40 mg OPA (dissolved in 1 mL methanol) and 100 μ L β -mercaptoethanol and diluting to a final volume of 50 mL with distilled water. 10-50 μ L of hydrolysates was added to 2 mL OPA reagent, the solution was mixed and incubated for 2 min at ambient temperature and the absorbance was measured at 340 nm. The degree of hydrolysis was calculated by using the equation:

DH (%) = (MW
$$\Delta_{340nm}$$
)/ (d.e.p) x 100

Where MW = Average molecular weight of amino acids (120)

 Δ_{340nm} = Absorbance at 340 nm

d = Dilution factor

e = Average molar absorption of amino acids (6000 M⁻¹ cm⁻¹)

P = protein concentration

3.2.3 High Pressure Processing (HPP) of Proteins

Soybean and chickpea protein dispersions at different concentration were packed in lowdensity polyethylene bags (Whirl-Pak^R, USA) and the bags were heat sealed. The air in the bags was squeezed out as much as possible before sealing. Samples were then transferred to the 5L pressure treatment chamber (ACIP 6500/5/12VB; ACB Pressure Systems, Nantes, France) equipped with temperature and pressure regulators. Water was used as the pressure transmitting medium. Soybean and chickpea protein samples were pressure treated at specified pressure levels (400, and 600 MPa) for 5 and 10 min. The pressurization rate was about 4.4 MPa/s and released at 26 MPa/s. The initial temperature of the medium was 18° C which quickly increased to 25 and 28.5° C during adiabatic effect during pressurization of 400 and 600 MPa and equilibrated to 22 and 26°C during holding period at those pressure levels. All experiments were carried out in duplicate.

Soybean and chickpea protein dispersions were subjected HPP treatment at 400 MPa and 600 MPa for 5 min and 10 min. High pressure treated protein dispersions were subjected to trypsin hydrolysis. To measure the impact of high pressure processing, degree of hydrolysis (%DH) was measure according to the method described in section 3.2.2. Antioxidant activity assay was performed according to the procedure described in section 3.3 on the soybean and chickpea protein hydrolysates (SPH/CPH) to measure the effect of high pressure processing. Figure 3.2 shows the steps involved in effect of High Pressure Processing (HPP) on soybean and chickpea protein.



Figure 3.2 Steps involved in effect of High Pressure Processing (HPP) on soybean and chickpea protein.

3.3 Antioxidant Activity Assay

The antioxidant activity of soybean and chickpea protein isolates/hydrolysates was measured by using DPPH (1, 1-diphenyl-2-picrylhydrazyl) radical scavenging assay method described by Shimada et al., (1992). 2 mL each protein sample at different concentrations (0-10 mg/ml) was added to 2 mL 0.1 mM DPPH dissolved in 95 % ethanol. The solution was mixed well and incubated for 30 min in dark at room temperature. After incubation the absorbance of the mixture was measured at 517 nm. Scavenging activity was calculated by using equation:

Scavenging activity (%) = Blank absorbance – Sample absorbance/ Blank absorbance x 100

3.4 Characterization of Proteins and Protein Hydrolysates

3.4.1 Polyacrylamide Gel Electrophoresis (PAGE)

The soybean and chickpea protein/protein hydrolysates were subjected to sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE). Samples of lyophilized protein and protein hydrolysates were subjected to SDS-PAGE. Electrophoresis was carried out on a Mini-Protein III Electrophoresis Cell unit (Bio-Rad, Hercules, CA). SDS-PAGE was carried out according to the method described by Laemmli, (1970).

3.4.1.1 Sodium Dodecyl Sulfate Polyacrylamide Gel Electrophoresis (SDS-PAGE)

3.4.1.1.1 Sample Preparation

Freeze dried protein samples were dissolved in sample buffer consisting of Tris-HCL (1.5M, pH 8.8), glycerol, SDS (2 %), 2-mercaptoethanol, bromophenol blue (0.1 %) and distilled water at a final concentration of 5-10 mg/ml. Samples were heated at 95° C for 5 min before loading into each sample well.

3.4.1.1.2 Gel Concentration and Preparation

Gels of 12 % and 4 % (w/v) acrylamide as separation and stacking gels were used. A 12 % acrylamide separation gel was prepared with the following reagents: acrylamide, Tris-HCL (1M, pH 8.8), glycerol, SDS (10 %), distilled water, ammonium persulfate (APS) (10 %) and tetramethylethylene-diamine (TEMED). The glass plates were fixed and the liquid was pour between the glass plates and allowed to polymerize for 35-40 min. Similarly, the 4 % acrylamide gel was prepared with acrylamide, Tris-HCL (0.5 M, pH 6.8), SDS (10 %), distilled water, APS (10 %) and TEMED. The mixture for stacking gel was poured over the separation gel and a comb was inserted to form wells for sample loading and allowed to polymerize for 30min. A Hamilton syringe was used to load 10-15 μ L protein samples at a concentration of 5-10 mg/mL into wells. Protein bands were compared in relation to the mobility of the following marker proteins and a standard curve of log MW verses RF (relative mobility) values was established for determination of molecular weights of unknown proteins: myosin (200 kDa), β -galactosidase (116 kDa), phosphorylase b (97 kDa), serum albumin (66 kDa), ovalbumin (45 kDa), Carbonic anhydrase (31 kDa), trypsin inhibitor (22 kDa), lysozyme (14 kDa) and aprotinin (7 kDa).

3.4.1.1.3 Electrophoresis Running Conditions

The running buffer was consisting of tris-base glycine buffer with 1 % SDS. Electrophoresis was performed at a constant voltage of 100 V/gel. Electrophoresis run time varied between 1.5-2 h and terminated when the tracking dye front reached the bottom of the gel.

3.4.1.1.4 Protein Staining and Destaining

Gels were removed from the glass plates and immersed in a staining solution containing Coomassie Blue R-250 (0.1 % w/v) for approximately 18 h. The gels were destained with the destaining solution containing 20 % methanol and 10 % acetic acid until the background colour was removed.

3.4.2 RP-HPLC Separation of Protein and Protein Hydrolysates

Separation of protein and peptides in protein hydrolysates was performed by using reversedphase high performance liquid chromatography (RP-HPLC) according to the method described by El-Ramahi, (2003) with some modifications. The HPLC was equipped with a dual pump system for high pressure solvent delivery, a programmable solvent module (model 126, Beckman, CA) and a UV detector (model 166). The samples of protein and protein hydrolysates at concentration 5-10 mg/mL were filtered by using 0.45 μ m MilliporeTM membrane filters, subjected to RP-HPLC analysis. An Octadecyl (C18) reverse phase column (5 μ m, 250 x 4.6 mm column, Mallinckrodt Baker, Inc., New Jersey, USA) was used. 100 μ L sample was injected into a 200 μ L sample loop and elution was done at a flow rate of 1 mL/min using two solvent gradient system. Solvent A, 0.1 % trifluoracetic acid (TFA) in water; solvent B, 0.1 % TFA in 70/30 acetonitrile/water. Elution was carried 60 min and detection was carried out at 215 nm. Chromatographic data were analyzed by Beckman Gold System (version V810, USA) then translated from print (PRN) format to Microsoft Excel[®] worksheet.

CHAPTER 4

RESULTS AND DISCUSSION

4.1 Enzymatic Hydrolysis

4.1.1 Trypsin Hydrolysis of Isolated Soybean Protein (ISP) and Isolated Chickpea Protein (ICP)

Trypsin is an alkaline protease that has been used not only for production of protein hydrolysates with better functional and nutritional characteristics than the original proteins, but also for the generation of bioactive peptides. Figures 4.1 and 4.2 shows the hydrolysis curve of isolated soybean protein (ISP) and isolated chickpea protein (ICP). Table 4.1 shows the values of % DH. The results reveal that degree of hydrolysis (% DH) increased with increase in time up to 105 min and then decreased. The maximum DH for soybean protein hydrolysates (SPH) and chickpea protein hydrolysates (CPH) were 12.4 % and 13.6 % respectively after 105 min. The DH for controls (Isolated soybean protein (ISP) and isolated chickpea protein (ICP) without enzyme) were 1.1 % and 1.2 % respectively. These findings are in agreement with the results of enzymatic hydrolysis of other proteins reported by other workers. Ayad (2010) reported that when flaxseed protein isolate was hydrolysed for 2 h using trypsin, the DH was 9.4 %-24.5 % after 0.5-2 h of hydrolysis; the % DH increased with increase in enzyme/substrate ratio and time of hydrolysis. The DH for rice endosperm protein (Zhang et al., 2009) and wheat germ protein hydrolysates (Zhu et al., 2005) was reported to be 11.7 % and 25 % after 6 h of hydrolysis respectively; the % DH increased with increase in time of hydrolysis.



Figure 4.1 Hydrolysis curve of soybean protein hydrolysates (SPH) treated with trypsin. Reaction conditions: pH, 8.0; 37° C, enzyme:protein ratio- 1:20 (w/w), and curve of isolated soybean protein (ISP) as control (without enzyme).



Figure 4.2 Hydrolysis curve of chickpea protein hydrolysates (CPH) treated with trypsin. Reaction conditions: pH, 8.0; 37° C, enzyme:protein ratio- 1:20 (w/w), and curve of isolated chickpea protein (ICP) as control (without enzyme).

| Time (min) | % DH of ISP(control) | % DH of SPH | % DH of ICP(control) | % DH of CPH |
|------------|-------------------------|-------------|-------------------------|-------------|
| 0 | 0.7 | 2.7 | 0.6 | 4.5 |
| 15 | 1.1 | 3.8 | 0.9 | 6.3 |
| 30 | 1.1 | 4.7 | 1.1 | 8.2 |
| 45 | 0.9 | 6.4 | 1.1 | 9.4 |
| 60 | 1.1 | 8.4 | 1.2 | 10.5 |
| 75 | 1.1 | 9.6 | 1.2 | 11.7 |
| 90 | 0.9 | 10.7 | 1.1 | 12.4 |
| 105 | 0.9 | 12.4 | 1.2 | 13.6 |
| 120 | 0.9 | 11.5 | 1.2 | 12.3 |

Table 4.1 % DH of isolated soybean protein (ISP), isolated chickpea protein (ICP), soybean protein hydrolysates (SPH) and chickpea protein hydrolysates (CPH).

Results are average of duplicate determination

DH = Degree of Hydrolysis

4.2 Antioxidant Activity of Protein and Protein Hydrolysates

4.2.1 Antioxidant Activity of Isolated Soybean Protein (ISP) and Soybean Protein Hydrolysates (SPH); Isolated Chickpea Protein (ICP) and Chickpea Protein Hydrolysates (CPH)

Figure 4.3 shows the antioxidant activity of isolated soybean protein (ISP) and soybean protein hydrolysates (SPH). The results showed a continuous increase in antioxidant activity of ISP and SPH with increase in protein concentration from 2.5-10 mg/mL with antioxidant activity from 6.0 % to 10.6 % for ISP and from 16.5 % to 32 % for SPH. Soybean protein hydrolysates showed higher antioxidant activity as compared to the isolated soybean protein at same protein concentration; this is because for protein hydrolysates and peptides an increase in hydrophobicity will increase their solubility in lipid and therefore enhances their antioxidant activity (Zhu et al., 2005). Zhu et al., (2005) also reported the increase in the antioxidant activity of wheat germ protein hydrolysates (WGPH) with increase in protein concentration and suggested that released antioxidant peptides are the reason for increased antioxidant activity of wheat germ protein hydrolysates as compared to the isolated wheat germ protein.

Figure 4.4 shows the results for the antioxidant activity of isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH). The results showed similarity with the results for isolated soybean protein and soybean protein hydrolysates with an increase in antioxidant activity of ICP and CPH with increase in protein concentration from 2.5-10 mg/mL. Antioxidant activity was 1.8 % to 8.3 % for ICP and 3.4 % to 26.8 % for CPH. Chickpea protein hydrolysates also showed higher antioxidant activity as compared to the isolated chickpea protein at same protein concentration; the reason is the same as discussed above for SPH. ICP and CPH showed

lower antioxidant activity as compared to the ISP and SPH, the reason for this can be lower content of hydrophobic amino acids in chickpea as compared to the soybean protein. Zhang et al., (2009) also reported an increase in antioxidant activity of rice endosperm protein with an increase in protein concentration, and that rice endosperm protein hydrolysates have higher antioxidant activity as compared to the isolated rice endosperm protein. Table 4.2 shows the % antioxidant activity of soybean/chickpea proteins and hydrolysates at different protein concentration.



Figure 4.3 Antioxidant activity curve for isolated soybean protein (ISP) and soybean protein hydrolysates (SPH).



Figure 4.4 Antioxidant activity curve for isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH).

Table 4.2 Antioxidant activity of isolated soybean protein (ISP) and soybean protein hydrolysates (SPH); isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH) at different protein concentrations.

| Conc. (mg/mL) | % AA (ISP) | % AA (SPH) | % AA (ICP) | % AA (CPH) |
|------------------|------------|------------|------------|------------|
| 2.5 | 6 | 16.5 | 1.8 | 3.4 |
| 5.0 | 7.5 | 25.0 | 4.8 | 10.8 |
| 7.5 | 8.7 | 27.8 | 5.3 | 17.9 |
| 10 | 10.6 | 32.0 | 8.3 | 26.8 |

Results are average of duplicate determinations

AA = Antioxidant Activity

4.3 Characterization of Proteins and Protein hydrolysates

4.3.1 SDS-PAGE of Isolated Soybean Protein (ISP), Soybean Protein Hydrolysates (SPH), Isolated Chickpea Protein (ICP) and Chickpea Protein Hydrolysates (CPH)

Figure 4.5 shows the SDS-PAGE patterns of the isolated soybean protein (ISP) and soybean protein hydrolysates (SPH). Figure 4.6 shows the standard curve used for the estimation of molecular weight (MW) of protein fractions. Isolated soybean protein showed four protein bands. The MW of one protein band was 80 KDa reported for β -conglycinin subunits, a major soy protein; the estimated MW of two other bands was 58 and 26 KDa were for glycinin subunits, a second major soy protein (Chang, 2010). The estimated MW of one last band was 16 KDa reported could be a trypsin inhibitors (Burks et al., 1991). No major protein bands were identified in soybean protein hydrolysates, confirming the proteolytic action of the trypsin enzyme. These findings are in agreement with the results reported by other workers. The estimated MW of glycinin subunits (26 and 58 KDa) are similar to those of glycinin subunits (26 and 61 KDa) reported by Helm et al., (2000). The estimated MW of 80 KDa is in the range of that reported for β -conglycinin subunits (52 to 82 KDa) reported by Abdolgader (2000). The molecular weight of 16 KDa is similar to those of trypsin inhibitors (8-21.5 KDa) reported by Burks et al., (1991).



Figure 4.5 SDS-PAGE of isolated soybean protein (ISP) and soybean protein hydrolysates (SPH). (M) Standard protein markers; (A) Isolated soybean protein (conc. 5 mg/mL); (B) Soybean protein hydrolysates (conc. 5 mg/mL).



Figure 4.6 Standard curve generated by plotting the log of the molecular weight of protein standards vs. the relative mobility.

Figure 4.7 shows the SDS-PAGE patterns of isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH). Isolated chickpea protein showed four protein bands. The estimated MW of one protein band was 62 KDa reported for lipoxygenase (Chang, 2010). The estimated molecular weight of two other bands was 34 KDa and 16 KDa reported for legumin αsubunits and vicilin subunits. The MW of one protein band was 133 KDa reported as unknown. In comparison to ICP, no major protein bands were characterized for CPH confirming the hydrolysis of chickpea protein; confirming the proteolysis of the protein. These findings are in agreement with the results reported by other workers. The estimated MW of lipoxygenase (62 KDa), legumin α-subunits (34 KDa) and vicilin subunits (16 KDa) are similar to those of lipoxygenase (62 KDa), legumin α-subunits (32-43 KDa), vicilin subunits (16 KDa) reported by Chang (2010), Lasztity (1996) and Gueguen (1991). Chang (2010) also reported an unknown protein band with MW 130-200 KDa for isolated chickpea protein. The SDS-PAGE results indicate that the isolated soybean protein and isolated chickpea protein substantially hydrolyzed although the DH data indicated of limited hydrolysis ranging from 2.7 % -11.5 % and 4.5 %-13.6 %.



Figure 4.7 SDS-PAGE of isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH). (M) Standard protein markers; (A) Isolated chickpea protein (conc. 5 mg/mL); (B) Chickpea protein hydrolysates (conc. 5 mg/mL).

4.3.2 Reversed Phase HPLC (RP-HPLC)

Figure 4.8 shows the profile of isolated soybean protein (ISP) and soybean protein hydrolysates (SPH) chromatogram by RP-HPLC. Soybean protein hydrolysates showed at least 30-35 peaks confirming the hydrolysis of the soybean protein, by comparison the isolated soybean protein without hydrolysis showed four minor peaks. From the RP-HPLC profile of SPH, peptides with antioxidant activity were expected to be present within the retention time of 15 min to 35 min. Four different RP-HPLC fractions of SPH (F I, F II, F III and F IV) were collected from 15 to 35 min to evaluate their antioxidant activity according to the method described in section 3.3 and to find fraction with high antioxidant activity. The results reported for the RP-HPLC analysis of ISP and SPH are in agreement with the results reported by Aludatt (2006) for ISP and SPH; the RP-HPLC analysis showed approximately similar number of peaks.

Figure 4.9 shows the profile of isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH) chromatogram by RP-HPLC. RP-HPLC analysis of the ICP and CPH showed agreement with the profile of ISP and SPH. Few minor peaks were reported for isolated chickpea protein, while chickpea protein hydrolysates showed at least 30-35 peaks confirming the hydrolysis of chickpea protein. Peptides with high antioxidant activity were expected to be with in the retention time of 15-35 min. Four different fractions of CPH (F I, F II, F III, and F IV) were collected to evaluate their antioxidant activity and to find fraction of CPH with high antioxidant activity. The results for the RP-HPLC analysis of ICP and CPH are in agreement with the results reported by Liu et al., (2007) and Chang (2010); approximately similar number of peaks were obtained for both ICP and CPH.



Figure 4.8 RP-HPLC chromatogram of isolated soybean protein (ISP) and soybean protein hydrolysates (SPH).



Figure 4.9 RP-HPLC chromatogram of isolated chickpea protein (ICP) and chickpea protein hydrolysates (CPH).

4.4 Antioxidant Activity Assay of Fractions collected from RP-HPLC

Antioxidant activity of fractions collected was measured according to the method described in section 3.3. Figure 4.10 shows the standard curve for the bovine serum albumin (BSA), used to calculate the protein concentration of fractions collected. Table 4.3 shows the antioxidant activity of fractions collected from RP-HPLC analysis of SPH and CPH at protein concentration 1 mg/mL for each fraction. The results revealed that F III of SPH showed highest antioxidant activity of 47.71 % and F II of CPH showed highest antioxidant activity of 27.96 % at protein concentration of 1 mg/mL suggesting that the F III of SPH and F II of CPH are rich in peptides or amino acids with antioxidant potential. The results reported for CPH and SPH are in agreement with the results reported by other workers. Fractions of CPH showed lower antioxidant activity as compared to the fractions of SPH; could be related to the lower hydrophobic amino acids in CPH (Zhu et al., 2005). Aluko et al., (2011) and Liu et al., (2007) also reported that antioxidant activity of pea protein hydrolysates and chickpea protein hydrolysates is due to the antioxidant potential of hydrophobic amino acids. Aluko et al., (2009) reported that tryptophan, a hydrophobic amino acid released from mother's milk has antioxidant properties. Garner et al., (1998a) reported methionine, a hydrophobic amino acid has high antioxidant potential. In comparison to CPH, amount of hydrophobic amino acids could be the reason for high antioxidant activity of SPH. Soybean has high percentage of methionine and this could be the reason for high antioxidant activity of SPH.



Figure 4.10 Standard curve for the bovine serum albumin used to calculate the protein concentration of fractions collected from RP-HPLC.

| Table 4.3 | Antioxidant | activity of | of soybean | protein | hydrolysates | (SPH) | and | chickpea | protein |
|------------|---------------|-------------|--------------|---------|--------------|-------|-----|----------|---------|
| hydrolysat | es (CPH) frac | tions coll | ected from l | RP-HPL | C. | | | | |

| Fractions | % Antioxidant Activity (SPH) | % Antioxidant Activity (CPH) |
|----------------------|---------------------------------|---------------------------------|
| F I (RT 15-20 min) | 28.0 | 25.3 |
| F II (RT 20-25 min) | 29.8 | 27.9 |
| F III (RT 25-30 min) | 47.71 | 26.2 |
| F IV (RT 30-35 min) | 22.9 | 20.8 |

Protein concentration 1 mg/mL

Results are average of duplicate determination

4.5 Effect of High Pressure Processing (HPP) on Degree of Hydrolysis (% DH) of Soybean and Chickpea protein

Figure 4.11 and Figure 4.12 shows graphs for the degree of hydrolysis (% DH) of soybean and chickpea proteins treated with High Pressure Processing (HPP) and trypsin hydrolysis. After high pressure processing treatment followed by trypsin hydrolysis of soybean and chickpea proteins, the degree of hydrolysis (% DH) showed an increase with pressure and time when compared to the protein samples with no HPP treatment.

Table 4.4 and Table 4.5 shows the effect of high pressure processing on the degree of hydrolysis of soybean and chickpea proteins. The highest rate of hydrolysis was achieved for the protein samples treated at 600 MPa for 10 min with an almost two fold increase in degree of hydrolysis for both soybean and chickpea protein samples as compared to the soybean and chickpea protein hydrolysates with no high pressure processing. For soybean protein samples treated at 400 MPa for 5 min, % DH showed an increase from 0-60 min then decreased; for protein samples treated at 400 MPa for 10 min, % DH showed an increase from 15-75 min and then decreased; for protein samples treated at 600 MPa for 5 min and 10 min, % DH showed an increase from 0-60 min then decreased. The results showed a noticeable relation between % DH and applied pressure. For SPH treated with HPP, the % DH reached 17.9 %-24.9 % as compared to the 2.7 %-12.4 % of the soybean protein hydrolysates without high pressure processing.

The results for % DH of CPH with high pressure processing are similar to the results of SPH treated with high pressure processing. For chickpea protein samples treated at 400 MPa for 5 min and 10 min, % DH showed an increase from 0-90 min then decreased; for chickpea protein samples treated at 600 MPa for 5 min and 10 min, % DH showed an increase from 0-90 min then

decreased. For chickpea protein hydrolysates treated with high pressure processing, the % DH reached 13.0 %-25.5 % as compared to the 4.5 %-13.6 % of the chickpea protein hydrolysates without high pressure processing.



Figure 4.11 Trypsin hydrolysis curve of soybean protein hydrolysates (SPH) treated with high pressure processing at 400 MPa and 600 MPa for 5 and 10 min with trypsin and curve for isolated soybean protein (ISP) without trypsin and without high pressure processing, and soybean protein hydrolysates (SPH) without high pressure treatment and with trypsin. Reaction conditions: pH, 8.0; 37°C, enzyme-protein ratio- 1:20 (w/w).



Figure 4.12 Trypsin hydrolysis curve of chickpea protein hydrolysates (CPH) treated with high pressure processing at 400 MPa and 600 MPa for 5 and 10 min, and curve for isolated chickpea protein (ICP) without trypsin and without high pressure processing, and chickpea protein hydrolysates (CPH) without high pressure treatment and with trypsin. Reaction conditions: pH, 8.0; 37°C, enzyme-protein ratio- 1:20 (w/w).

| Time (min) | % DH (ISP) | % DH (SPH) | % DH (SPH-400 MPa-5 min) | % DH (SPH-400 MPa-10 min) | % DH (SPH-600 MPa-5 min) | % DH (SPH-600 MPa-10 min) |
|---------------|---------------|---------------|--------------------------------|------------------------------------|-----------------------------------|------------------------------------|
| 0 | 0.7 | 2.7 | 10.0 | 11.6 | 17.2 | 17.9 |
| 15 | 1.1 | 3.8 | 11.8 | 14.6 | 19.1 | 20.1 |
| 30 | 1.1 | 4.7 | 13.9 | 14.9 | 21.2 | 22.5 |
| 45 | 0.9 | 6.4 | 14.2 | 15.7 | 21.9 | 23.5 |
| 60 | 1.2 | 8.4 | 16.3 | 17.6 | 23.9 | 24.9 |
| 75 | 1.1 | 9.6 | 16.2 | 17.8 | 21.3 | 22.2 |
| 90 | 0.9 | 10.7 | 15.5 | 16.3 | 20.2 | 21.8 |
| 105 | 0.9 | 12.4 | 15.9 | 16.0 | 22.6 | 23.3 |
| 120 | 0.9 | 11.5 | 15.7 | 17.1 | 21.2 | 23.4 |

Table 4.4 Degree of Hydrolysis (% DH) of Soybean Protein after High Pressure Processing (HPP) and Trypsin Hydrolysis.

Results are average of duplicate determinations.

DH = Degree of Hydrolysis

| Time (min) | % DH (ICP) | % DH (CPH) | % DH (CPH-400 MPa-5 min) | % DH (CPH-400 MPa-10 min) | % DH (CPH-600 MPa-5 min) | % DH (CPH-600 MPa-10 min) |
|---------------|---------------|---------------|--------------------------------|------------------------------------|-----------------------------------|------------------------------------|
| 0 | 0.6 | 4.5 | 10.3 | 12.9 | 13.5 | 13.0 |
| 15 | 0.9 | 6.3 | 11.8 | 13.9 | 14.5 | 15.3 |
| 30 | 1.1 | 8.2 | 14.6 | 15.7 | 16.4 | 17.4 |
| 45 | 1.1 | 9.4 | 17.4 | 18.9 | 19.3 | 21.6 |
| 60 | 1.2 | 10.5 | 19.4 | 20.8 | 22.0 | 24.0 |
| 75 | 1.2 | 11.7 | 21.6 | 23.5 | 24.0 | 25.5 |
| 90 | 1.1 | 12.4 | 22.3 | 24.1 | 25.7 | 26.2 |
| 105 | 1.2 | 13.6 | 21.8 | 23.2 | 23.8 | 24.1 |
| 120 | 1.2 | 12.3 | 20.6 | 23.4 | 23.4 | 23.2 |

Table 4.5 Degree of Hydrolysis (%DH) of Chickpea Protein after High Pressure Processing (HPP) and Trypsin Hydrolysis.

Results are average of duplicate determinations.

DH = Degree of Hydrolysis

Proteins are flexible molecules and become compress when subjected to high pressure processing; high pressure affects the non-covalent bonds, especially hydrophobic and electrostatic interactions of proteins (Ledward, 2006). High pressure processing leads to the modification of the protein structure (Messens et al., 1997). Chapleau et al., (2003) reported that

high pressure treatments of proteins induces the formation of a structure closely resembling protein folding intermediate; this molecular state is characterized by the presence of a native-like degree of secondary structure but a loss of tight packing in the hydrophobic core of protein, leading to increased exposure of peptides or amino acids for hydrolysis. Stress applied by the high pressure processing partially unfolds the protein structure resulting into ease in protein hydrolysis and peptides with antioxidant activity were expected to release leading to the increase in antioxidant activity (Volk, 2009). This could explain why the observed degree of hydrolysis increased after trypsin hydrolysis for the soybean and chickpea protein samples treated with high pressure processing (HPP). The results reported for the effect of high pressure processing on soybean and chickpea protein supported earlier reports on high pressure effect by Ahmed et al., (2006).

4.6 Effect of High Pressure Processing on the Antioxidant activity

4.6.1 Effect of High Pressure Processing (HPP) on the Antioxidant Activity of Soybean Protein and Chickpea Protein

Figure 4.13 shows the results for the effect of HPP on the antioxidant activity of soybean protein hydrolysates. Table 4.6 shows the antioxidant activity of soybean protein hydrolysates after high pressure processing followed by trypsin hydrolysis, the results reveal that antioxidant activity showed an almost two fold increase with increase in pressure and time. The % antioxidant activity reached 29.9 %-66.7 % for the soybean protein sample treated with HPP at 600 MPa for 10 min followed by trypsin hydrolysis as compared to the 6.0 %-10.6 % of the isolated soybean protein; 16.5 %-32.0 % of the soybean protein hydrolysates without HPP at protein concentration 2.5-10 mg/mL.



Figure 4.13 Histograms showing the effect of high pressure processing on the antioxidant activity of soybean protein hydrolysates (SPH) at different concentrations with different pressure and time interval.
| Conc. (mg/mL) | % AA*- ISP | % AA- SPH | % AA- SPH-400 MPa-5min | % AA-SPH- 400 MPa- 10min | % AA- SPH-600 MPa- 5min | % AA- SPH-600 MPa- 10min |
|------------------|---------------|--------------|------------------------------|--------------------------------|----------------------------------|-----------------------------------|
| 2.5 | 6.0 | 16.5 | 25.4 | 29.3 | 27.3 | 29.9 |
| 5.0 | 7.5 | 25.0 | 33.3 | 36.3 | 36.7 | 42.8 |
| 7.5 | 8.7 | 27.0 | 43.3 | 45.1 | 47.4 | 55.9 |
| 10 | 10.6 | 32.0 | 49.1 | 54.1 | 53.2 | 66.7 |

Table 4.6 Effect of high pressure processing (HPP) on antioxidant activity of soybean protein hydrolysates (SPH).

Results are the average of duplicate determinations.

AA = Antioxidant Activity

Figure 4.14 shows the effect of high pressure processing on the antioxidant activity of chickpea protein hydrolysates (CPH). Table 4.7 shows the antioxidant activity of chickpea protein hydrolysates after high pressure processing followed by trypsin hydrolysis, the results are similar to the results of soybean protein treated with HPP; the antioxidant activity showed an almost two fold increase with increase in pressure and time. The % antioxidant activity reached 28.9 %-56.6 % for chickpea protein sample treated with HPP and trypsin hydrolysis as compared to the 1.8 %-8.3 % of isolated chickpea protein; 3.4 %-26.8 % of chickpea protein hydrolysates without high pressure processing at protein concentration 2.5-10 mg/mL. The results reported are in agreement with the results reported by other workers. Volk (2009) also reported an almost two fold increase in antioxidant activity of egg proteins after high pressure processing.



Figure 4.14 Histograms showing the effect of high pressure processing on the antioxidant activity of chickpea protein hydrolysates (CPH) at different concentrations with different pressure and time interval.

| Conc. (mg/mL) | % AA- ICP | % AA- CPH | % AA- CPH- 400MPa- 5min | % AA-CPH- 400MPa- 10min | % AA- CPH- 600MPa- 5min | % AA- CPH- 600MPa- 10min |
|------------------|--------------|--------------|----------------------------------|-------------------------------|----------------------------------|-----------------------------------|
| 2.5 | 1.8 | 3.4 | 22.2 | 25.9 | 26.1 | 28.9 |
| 5.0 | 4.8 | 10.8 | 27.6 | 34.4 | 38.9 | 37.3 |
| 7.5 | 5.3 | 17.9 | 36.7 | 41.5 | 45.6 | 48.9 |
| 10 | 8.3 | 26.8 | 45.4 | 47.4 | 48.6 | 56.6 |

Table 4.7 Effect of high pressure processing (HPP) on the antioxidant activity of chickpea protein hydrolysates (CPH).

Results are the average of duplicate determinations.

AA = Antioxidant Activity.

The findings are in agreement with the results reported by Cano et al., (2000) for effect of high pressure processing on radical scavenging activity of persimmon fruit purees; the results showed an almost two fold increase in radical scavenging activity of persimmon fruit purees. The results reported by Tauscher et al., (2006) showed an agreement with our results; an increase in antioxidant activity was reported for tomato puree after high pressure processing.

GENERAL CONCLUSION

Trypsin hydrolysis of the soybean and chickpea protein as measured by degree of hydrolysis (% DH) showed a continuous increase with time of hydrolysis. Maximum degree of hydrolysis for soybean and chickpea protein hydrolysates was reported to be 12.42 % and 13.6 %, respectively. Soybean and chickpea protein hydrolysates showed higher antioxidant activity as compared to the isolated soybean and chickpea protein. Maximum antioxidant activity for soybean and chickpea protein hydrolysates were 32 % and 26.8 % respectively at protein concentration 10 mg/mL. SDS-PAGE confirmed the hydrolysis of soybean and chickpea protein, four protein bands with different molecular weight were observed for isolated soybean and chickpea protein hydrolysates. Fractions with high antioxidant activity were identified from the RP-HPLC analysis of soybean and chickpea protein hydrolysates. The fraction resulting from 25 to 30 min of soybean protein hydrolysates showed maximum antioxidant activity of 47.71 % and the fraction resulting from 15 to 20 min of chickpea protein hydrolysates showed maximum antioxidant activity of 27.96 % at protein concentration 1 mg/mL.

High Pressure Processing showed increase in degree of hydrolysis (% DH) and antioxidant activity of soybean and chickpea protein with increase in pressure and time. Maximum degree of hydrolysis was reported to be 24.91 % and 26.21 % for the soybean and chickpea protein samples treated at 600 MPa for 10 min. Antioxidant activity increased by approximately two folds after high pressure processing. Maximum antioxidant activity was 66.67 % and 56.59 % for the soybean and chickpea protein hydrolysates treated with 600 MPa for 10 min and with trypsin hydrolysis.

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