

Production of Iron-Binding Peptides from Colostral Whey by Enzymatic Hydrolysis

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Abstract

Colostrum whey prepared from colostrum (pooled from first six post-partum milkings) was heated for 10 min at 100°C. Heated colostrum whey was incubated with 1% enzymes (protein equivalent basis) for 15, 30, 60, 90, and 120 min at 50°C. Papain, pepsin, trypsin, and alcalase produced different degrees of hydrolysis (DH), 10.66%, 12.42%, 10.83%, and 25.31%, respectively, at an incubation time of 120 min. The SDS-PAGE reveals that significant amounts of bovine serum albumin (BSA), β -lactoglobulin (β -LG), and α -lactalbumin (α -LA) survived papain digestion. In contrast, pepsin completely removed BSA but not β -LG present in heated colostrum whey. Alcalase completely eliminated BSA, β -LG, and α -LA. This differential hydrolysis was confirmed by reversed-phase HPLC analysis. Using ion-exchange chromatography, fraction-1 (F-1) was obtained from alcalase hydrolysate at a NaCl gradient concentration of 0.25 M. Reversed-phase HPLC chromatograms of alcalase F-1 showed numerous small peaks, which probably indicate that a variety of new peptides were produced. Iron content of alcalase F-1 was 28.94 ppm, which was the highest among all enzyme fractions, whereas iron content of colostrum whey was 36.56 ppm. Main amino acids contained in alcalase F-1 were Thr (15.45%), Glu (14.12%), and Ser (10.39%). Therefore, alcalase can be used to generate good iron-binding peptides in heated colostrum whey, and the resulting iron-binding peptides could be suitable as a value-added food ingredient for food supplements.

Key words: Colostral whey, hydrolysis, alcalase, iron-binding peptide

Introduction

Bovine colostrum refers to a mixture of lacteal secretions, and can be harvested immediately preceding or following parturition. The colostrum has been known to contain serum immunoglobulins, antimicrobial peptides, and growth factors which accumulate in the mammary gland during the late pregnancy (Foley and Otterby, 1978; Playford *et al.*, 2000).

Iron is one of the essential trace elements for human nutrition, which is obtained only through dietary intake. Iron deficiency is prevalent (WHO, 2001), due to either insufficient intake or the presence of iron absorption inhibitors. Liang *et al.* (2008, 2009) have tried different ways to decrease contents of phytic acid, which is men-

tioned as an important inhibitor of iron absorption, in rice for improved bioavailability of minerals. Our previous studies (Kim *et al.*, 2007) showed that alcalase can be used for the production of iron-binding protein/peptides from normal milk whey protein. Compared to normal milk, colostrums contains higher levels of proteins (14%); 4.8% casein, 0.9% albumin, and 6.0% immunoglobulin (Foley and Otterby, 1978). If bovine colostrums can be used to produce iron-binding protein/peptides, more protein/peptides could be generated from colostrum whey fraction than from normal whey proteins. Further, these different compositions can affect peptide production from colostrum whey by enzymatic hydrolysis. This could change iron-binding capacity of protein/peptides generated from colostrum whey. Therefore, it is important to study hydrolysis of colostrum whey protein by various enzymes to prepare possible iron-binding protein/peptides without having antigenic protein fractions, β -lactoglobulin (β -LG) and α -lactalbumin (α -LA) (Pintado *et al.*, 1999). The objectives of this study were to evaluate the influence of

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4 kinds of enzymes on colostrum whey hydrolysis, with particular emphasis on the identification and quantification of peptides having iron-binding ability.

Materials and Methods

Colostrum and enzymes

Colostrum samples (n=10) from multiparous Holstein cows were taken for first six milkings in plastic bottles on April, shipped to laboratory, and analyzed immediately with a Lactoscope (MK2; Delta Instruments, Drachten, Netherlands) that was previously standardized for colostrum. To harvest colostrum whey, the frozen, pooled colostrum samples were allowed to thaw at room temperature, mixed with deionized water at a ratio of 1:4, and then defatted using ultracentrifugation at 12,000×g for 20 min (Supra 25K, Hanil Sci., Incheon, Korea). Dialysis tubing cellulose membrane of 33×21 mm size (Sigma-Aldrich, NY, USA) was used to de-mineralize the colostrum. 10% acetic acid was slowly added to de-fatted and de-mineralized colostrum samples to achieve pH 4.6. Acidified samples were then centrifuged at 12,000 g for 25 min for three times at 4°C to separate casein and colostrum whey. The pH of the obtained colostrum whey was soon raised to 7.0 with 1 N NaOH to protect the immunoglobulins from denaturation. The whey was stored at -80°C for later usage.

Alcalase EC 3.4.21.14, [*Bacillus globigii* (*Bacillus licheniformis*), activity 2.4 units/g protein], pepsin (EC 3.4.23.1, porcine gastric mucosa, activity 25,000-35,000 units/g protein), papain (EC 3.4.22.2, Carica Papaya L., activity 14,000 units/g protein), and trypsin (EC 3.4.21.4, porcine pancreas, activity 161,000 units/g protein) were purchased from Sigma-Aldrich. Analytical grade BSA, trinitrobenzenesulfonic acid, trifluoroacetic acid (TFA; HPLC grade), and all other reagents were purchased from Sigma-Aldrich.

Preparation of colostrum whey hydrolysates

The colostrum whey hydrolysates were prepared from heated colostrum whey according to the procedures described by Kim *et al.* (2007). Briefly, the frozen colostrum whey was allowed to thaw at room temperature. The pH of colostrum whey was adjusted to 8 using 0.5 M NaOH to avoid any coagulation during thermal treatment, and then the solution was heated for 10 min at 100°C. The pH of heated colostrum whey samples was adjusted using 0.5 M NaOH to 8 for alcalase and trypsin, and 7.5 for papain treatments. For peptic hydrolysis the pH of the heated colostrum whey was adjusted to 2 using 1 N HCl. The en-

zymes were dissolved in deionized water to prepare a 1% solution on a protein-equivalent basis. Enzyme solutions were added to heated colostrum whey samples at the ratio of 1:100 (enzyme:substrate, v/v, protein basis). The protein hydrolysis was done at 50°C. The pH of each reaction mixture was further maintained at a constant level using a pH-stat technique (Metrohm Ltd., Herisau, Switzerland).

Degree of hydrolysis

For the measurement of degree of hydrolysis (DH), samples were withdrawn 15, 30, 60, 90, and 120 min after the addition of enzymes to reaction mixture. The enzymes were inactivated by heating the reaction mixture for 10 min at 90°C, and then cooling to -4°C. The supernatants were taken as colostrum whey hydrolysates and the precipitates were discarded. The hydrolysates were stored at -20°C for subsequent estimation of DH, peptide identification, and iron-binding capacity. The DH of colostrum whey by various enzymes in the different incubation times was estimated according to Adler-Nissen (1979).

SDS-PAGE

SDS-PAGE was used to observe changes in the bands of α -LA, β -LG, and BSA that may be present in colostrum whey hydrolysates obtained at different reaction times.

Separation of iron-binding peptides

Separation of the iron-binding peptide from the colostrum whey hydrolysate was conducted by the method of Rose *et al.* (1969). Diethylaminoethylcellulose (DEAE; Whatman DE52, Whatman, Brentford, Middlesex, UK) was equilibrated in 500 mL of 20 mM Tris-HCl buffer at pH 7.8. A slurry of equilibrated DEAE was packed in a glass column (20×2.5 cm). Enzymatic hydrolysates were dissolved in the same buffer (pH 7.8) and loaded onto the column, then eluted by a step gradient with the same buffer containing 0.25, 0.5, and 0.75 M NaCl. The flow rate was 3 mL/min, the fraction volume was 30 mL per tube, and elutions were monitored at 280 nm. The 40 mL of injection volume contained approximately 200 mg of protein. Samples were filtered through 0.45 μ m membrane filters (Acrodisc Syringe Filters, Gelman Laboratory, MI, USA) prior to application to the column.

Reversed-phase HPLC

Four fractions 1 (F-1) from 4 enzymatic colostrum whey hydrolysates, which was eluted at the 0.25 M NaCl gradient, were analyzed by RP-HPLC Zorbax 300SB (Agilent Technologies Inc., Palo Alto, CA, USA) C₁₈ column (4.6

×250 mm) equilibrated with solvent A (0.1% TFA in H₂O) and eluted with a linear hygradient of solvent B (0.1% TFA in acetonitrile) for 40 min. Runs were conducted at room temperature using an HPLC system (Agilent 1200, Agilent Technologies Inc., Palo Alto, CA, USA) at a flow rate of 1 mL/min, and the absorbance was monitored at 214 nm. The injection volume was 10 mL and the concentration of protein material applied was approximately equivalent to 0.5 mg/mL of protein. All samples were filtered through 0.2 µm membrane filters (Acrodisc Syringe Filters, Gelman Laboratory, MI, USA) prior to application to the C₁₈ column.

Determination of the iron-binding ability

The heated colostrum whey, its hydrolysates (supernatants), and fractions from DEAE chromatography were used to prepare 1% protein (w/v) solutions in deionized water with pH adjusted to 5.0. 0.1% (w/v) FeSO₄ was added to these solutions, and these mixtures were incubated for 1 h at 37°C. The iron content of these mixture were determined using inductively coupled plasma (ICP) spectroscopy (ICP-OES, Varian, USA) at a wavelength of 259.94 nm. ICP spectroscopy was operated at a radio frequency power of 1.2 kW, coolant gas of 14 µL/min, plasma gas of 1.2 µL/min, and carrier gas of 0.7 µL/min. All samples were filtered through 0.2 µm membrane filters prior to their application to the ICP. The ICP standard iron (AnApex Co., Ltd., Korea) was used to make calibration curve. The iron solubility of these mixtures was calculated as an indicator of the iron-binding capacity of protein/peptides.

Amino acid analysis and protein determination

Amino acid analysis of the iron-binding protein/peptides fraction of the alcalase hydrolysate was performed by the method of Moore *et al.* (1958). Iron-binding protein/peptides (1 mg) were exhaustively dialyzed against distilled water and lyophilized, which was then hydrolyzed in 1 mL of 6 M HCl in evacuated tubes for 24 h at 110°C. After speed-vacuum concentration (Mivac concentrator Range, Genevac Limited, Suffolk, UK), the sample was dissolved in 0.2 M sodium citrate loading buffer (pH 2.2), and filtered through 0.2 mm membrane filters. Amino acid determination was carried out on an amino acid analyzer (Biochem 20, Pharmacia, Uppsala, Sweden). Protein concentrations in the enzyme preparations, hydrolysates, and fractions were determined by the dye-binding method of Bradford (1976). Bovine serum albumin (Sigma-Aldrich, NY, USA) was used as the stan-

dard. Data on DH of colostrum whey hydrolysates, and iron contents of the hydrolysate fractions are presented as mean±SD.

Statistical analysis

Data on DH of colostrum whey hydrolysates, iron contents of the hydrolysate fractions and amino acid composition of heated colostrum whey and fraction 1 (F-1) are analyzed by the SAS package (SAS, 2002).

Results and Discussion

Protein hydrolysis

Mean DH (±SD) of heated colostrum whey by 4 enzymes (Fig. 1) ranged from 5.67% to 25.31%. Extensive hydrolysis of heated colostrum whey was observed during the first 15 min of enzymatic treatment, followed by a slow increase in hydrolysis with increasing reaction time. Alcalase showed the highest DH of heated colostrum whey compared with other enzymes at all reaction times. The respective DHs of heated colostrum whey with these enzymes at 120 min were 25.31% (alcalase), 12.42% (pepsin), 10.66% (papain), and 10.83% (trypsin). These DHs were much higher than those for whey protein concentrate where DHs of whey protein concentrate were 13% (alcalase), 8% (pepsin), 7% (papain), and 9% (trypsin) (Kim *et al.*, 2007). This could be due to more substrates present in colostrum whey than in whey protein concentrate. More extensive hydrolysis of whey protein with alcalase than other enzymes used in our study has been demonstrated previously by Smyth and Fitz-Gerald (1998) and Kim *et al.* (2007). Alcalase has been widely used to prepare soluble hydrolysates of soy protein (Fox, 1989), fish protein (Rebeca *et al.*, 1991), and whey protein (Kim *et al.*, 2007). As alcalase is relatively crude bacterial

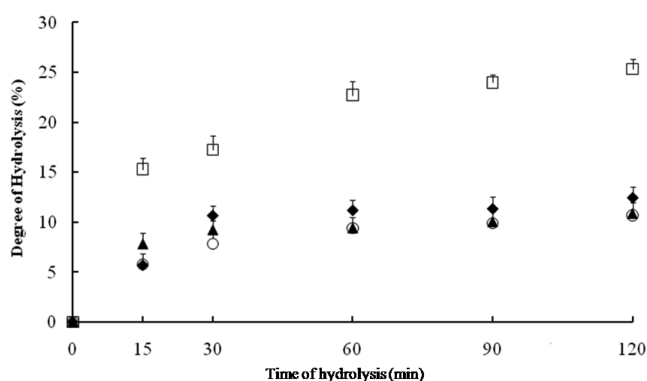


Fig. 1. Effects of alcalase (□), pepsin (◆), trypsin (△), and papain (○) on the degree of hydrolysis of heated colostrum whey at 50°C.

extract of *B. licheniformis* (Sukan and Andrews, 1982), it contains several different proteinases, each with different specificities. This property of alcalase could result in a significantly higher DH of heated colostrum whey compared with enzymes such as pepsin, trypsin, or papain.

The SDS-PAGE patterns of colostrum whey protein

hydrolysates obtained for 120 min's incubation by enzymes are shown in Fig. 2. Alcalase (Fig. 2A) successfully hydrolyzed all the major protein fractions of colostrum whey protein. Pepsin (Fig. 2B) did not completely digest α -LA and β -LG present in colostrum whey. Treatment of colostrum whey by papain (Fig. 2C) and trypsin

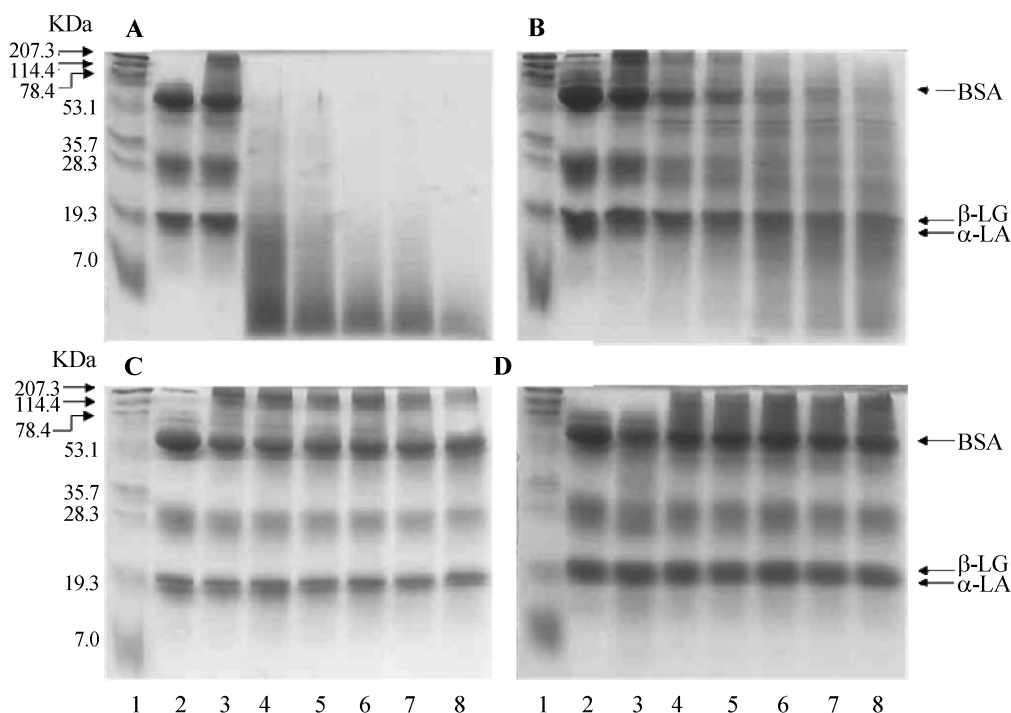


Fig. 2. SDS-PAGE patterns of heated colostrum whey hydrolyzed at 50°C with alcalase (A), pepsin (B), papain (C), and trypsin (D). 1: Standard board range marker (Bio-rad, USA), 2: Colostrum whey, and 3: Heated colostrum whey. Lanes 4-8 are different incubation times of 15, 30, 60, 90, and 120 min, respectively.

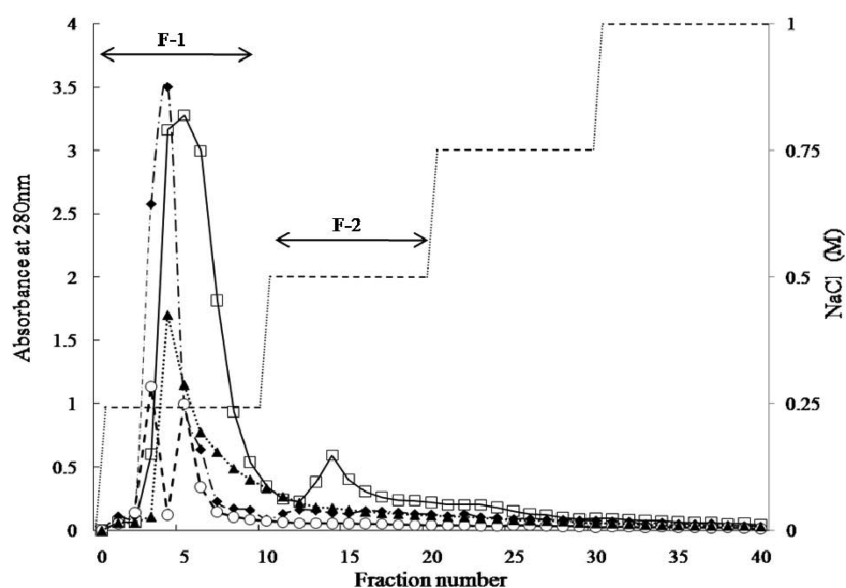


Fig. 3. Ion-exchange chromatograms of heated colostrum whey hydrolysate by alcalase (□), pepsin (◆), trypsin (△), and papain (○). Their hydrolysates dissolved in 20 mM Tris-HCl buffer (pH 7.8) were applied to the column. The column packed with DEAE-cellulose was washed with the same buffer, and then eluted with a step gradient of NaCl as indicated. The flow rate was 3 mL/min, fraction volume was 30 mL per tube, and elution was monitored at 280 nm.

(Fig. 2D) failed to hydrolyze bovine serum albumin, β -LG, and α -LA after 120 min of incubation.

Separation of iron-binding protein

Ion-exchange chromatograms of heated colostrals whey hydrolysate by alcalase, pepsin, trypsin, and papain are shown in Fig. 3. Fractions 1 (F-1) and 2 (F-2) were eluted from hydrolysates at the 0.25 and 0.5 M NaCl chromatographic step gradients, respectively. The RP-HPLC chromatogram patterns of each F-1 obtained from iron-exchange chromatography shown in Fig. 4. Many new peaks of shorter length of peptides appeared on the chromatograms of enzymatic hydrolysates of heated colostrals whey, which indicates degradation of whey protein into new shorter peptides. Heated colostrals whey hydrolysates derived with alcalase (Fig. 4A) and pepsin (Fig. 4B) produced greater numbers of shorter peaks than those derived with papain and trypsin. However, the chromatograms of hydrolysates by papain (Fig. 4C) and trypsin (Fig. 4D) showed that β -LG still survived these enzyme treatments.

Iron contents of fractions (F-1 and F-2) obtained from ion-exchange chromatography of heated colostrals whey hydrolysate are shown in Fig. 5. The highest iron contents were observed in F-1 that was derived with alcalase (28.94 mg/L). This value was much higher than that (0.21 mg/L) observed for alcalase F-1 prepared from whey protein concentrate (Kim *et al.*, 2007). Iron contents of trypsin F-1, pepsin F-1, and papain F-1 were 7.39, 7.04, and 3.27 mg/L, respectively.

Amino acid composition

The amino acid compositions of heated colostrals whey and the F-1 eluted by ion-exchange chromatography of the alcalase hydrolysate are shown in Table 1. The heated colostrals whey contained greater amount of Ala (16.09%) and Glu (14.85%) and lower amount of Ser (4.86%). However, Kim *et al.* (2007) reported that whey protein concentrate had higher amount of Glu and Leu, and lower amount of Ala. This difference can be due to higher proportion of immunoglobulins in heated colostrals whey.

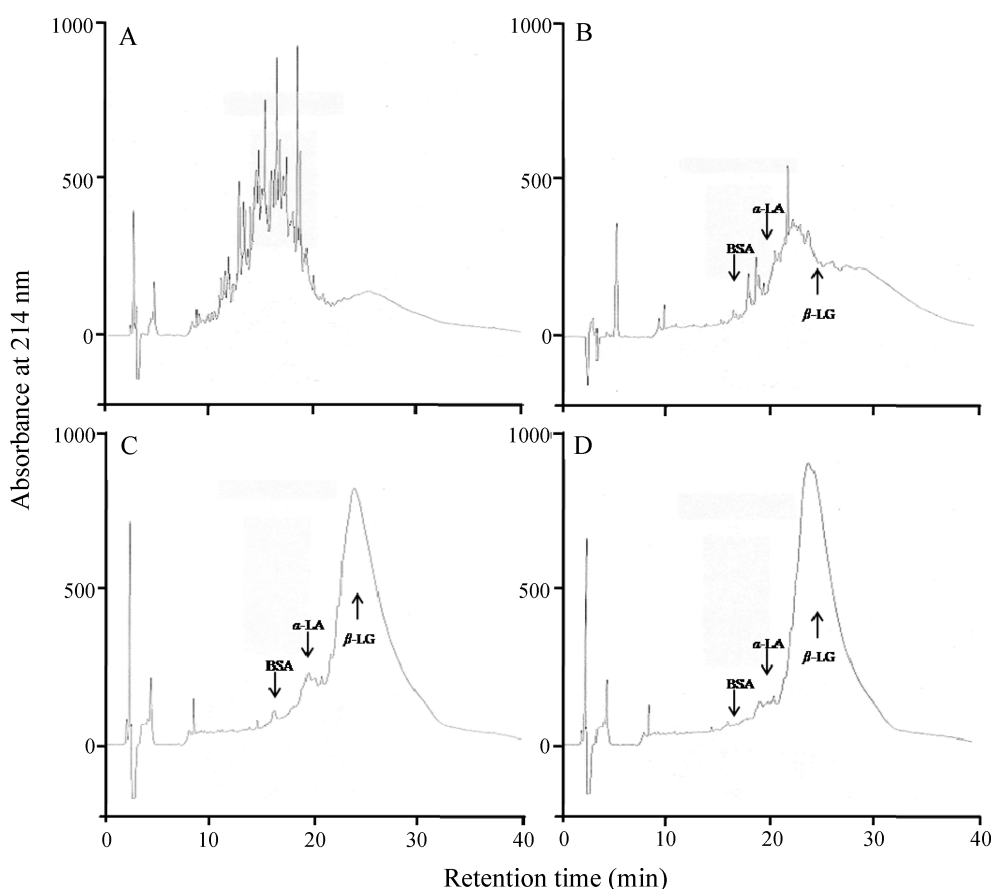


Fig. 4. RP-HPLC chromatogram of fraction 1 (F-1) isolated by ion-exchange chromatography from heated colostrals whey hydrolysates treated with alcalase (A), pepsin (B), papain (C), and trypsin (D). The column was equilibrated with solvent A (0.1% TFA in H_2O) and eluted with a linear gradient of solvent B (0.1% TFA in acetonitrile). The flow rate was 1 mL/min, injection volume was 10 μ L, and detection was at 214 nm.

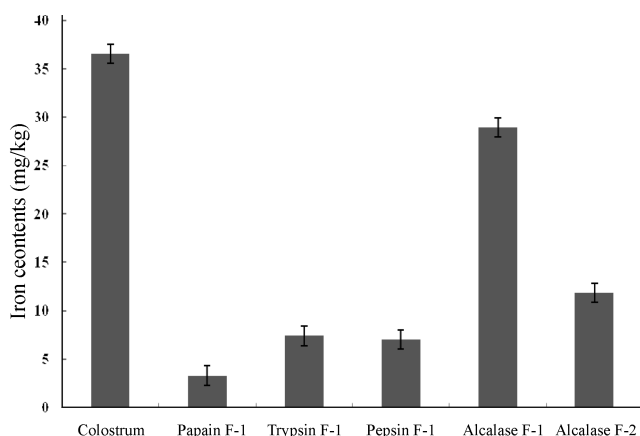


Fig. 5. Mean (\pm SD) iron contents (mg/kg) of heated colostrum whey and fractions from respective enzymatic treatments of heated colostrum whey. Each suspension was dissolved in de-ionized water (1% protein basis and pH 5.0). Iron-binding protein/peptides were precipitated by the addition of 0.1% (w/v) FeSO_4 , and incubation at 37°C for 1 h. The iron contents were determined using ICP (Inductively Coupled Plasma Spectrometer, ICP-OES, Varian, USA).

Table 1. Amino acid composition of heated colostrum whey and fraction 1 (F-1) eluted from ion-exchange chromatography of alcalase hydrolysate

Amino acids	Colostrum whey protein (%)	Alcalase F-1 (%)
Aspartic acid (Asp)	9.50 \pm 0.1 ¹⁾	8.02 \pm 0.08
Threonine (Thr)	6.07 \pm 0.08	15.45 \pm 0.09
Serine (Ser)	4.86 \pm 0.06	10.39 \pm 0.07
Glutamic acid (Glu)	14.85 \pm 0.1	14.12 \pm 0.15
Proline (Pro)	1.15 \pm 0.02	2.44 \pm 0.04
Glycine (Gly)	4.25 \pm 0.01	6.31 \pm 0.16
Alanine (Ala)	16.09 \pm 0.09	5.89 \pm 0.11
Valine (Val)	5.90 \pm 0.07	5.51 \pm 0.06
Cysteine (Sys)	2.31 \pm 0.06	1.46 \pm 0.08
Methionine (Met)	3.86 \pm 0.11	4.11 \pm 0.15
Isoleucine (Ile)	4.47 \pm 0.11	4.46 \pm 0.18
Leucine (Leu)	5.61 \pm 0.09	8.16 \pm 0.07
Tyrosine (Tyr)	2.64 \pm 0.04	2.16 \pm 0.04
Phenylalanine (Phe)	3.38 \pm 0.04	2.06 \pm 0.08
Lysine (Lys)	3.91 \pm 0.03	4.15 \pm 0.1
Histidine (His)	1.43 \pm 0.01	2.89 \pm 0.1
Arginine (Arg)	9.73 \pm 0.07	2.41 \pm 0.05
Total mol. ratio	100.00	100.00

¹⁾ Values are means \pm SE.

Immunoglobulins account for about 50% of the total amount of colostrum proteins, and about 90% of immunoglobulin is IgG₁ (Georgiev, 2008). Ser (13.57%) is highest amounts of amino acid of all amino acids in IgG₁, and Ala and Glu constitute 5.76 and 8.20%, respectively (Larson, 1992). Therefore, lower Ser content found in our

study could be due to lower content of Ser in α -LA and β -LG. Higher proportion of Glu in IgG₁, α -LA, and β -LG is responsible for higher content of Glu in our heated colostrum whey. Alcalase F-1 had high percentage of Thr (15.45%), Glu (14.12%), and Ser (10.39%), whereas Kim *et al.* (2007) reported whey protein concentrate had high percentage of Ala, Phe, and Lys.

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