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# Peptide separation of commercial fermented milk during refrigerated storage

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#### Abstract

Milk is an important source of bioactive compounds. Many of these compounds are released during fermentation and refrigerated storage. The aim of this study was to determine the release of peptides by lactic acid bacteria in commercial fermented milk during refrigerated storage. The size and profile of peptides were analyzed by polyacrylamide gel electrophoresis and size-exclusion HPLC. During electrophoresis, it was observed that the peptides were released from caseins, whereas  $\beta$ -lactoglobulin was the whey protein with the highest degradation. HPLC analysis confirmed the pattern of peptide formation observed in electrophoresis. Two fractions lower than 2 kDa with aromatic amino acids in their structure were separated. These results were consistent with those reported for structures of peptides with antihypertensive activity. Therefore, the presence of aromatic amino acids in the peptide fractions obtained increases the likelihood of finding peptides with such activity in refrigerated commercial fermented milk. In conclusion, during cold storage, peptides with different molecular weights are released and accumulated. This could be due to the action of proteinases and peptidases of the proteolytic system in lactic acid bacteria.

Keywords: bioactive peptide; fermented milk; cold storage; antihypertensive activity; lactic acid bacteria.

#### 1 Introduction

Functional foods can be obtained by the addition of probiotics, prebiotics, synbiotic, or functional ingredients. Probiotics are live microorganisms that exert positive effects on health by improving the balance in the intestinal microflora of consumers (Yerlikaya, 2014; Stanton et al., 2005; Vinderola et al., 2002; Gorbach, 2000). The growth of microorganisms in the gastrointestinal tract may be stimulated by prebiotics. Prebiotics are non-digestible oligosaccharides fermentable by different species of lactic acid bacteria (LAB) (Roberfroid, 2000). Synbiotics (a combination of probiotics and prebiotics) improve survival of probiotics enhancing their effects in the large bowel (Roberfroid, 2000). Several studies have shown additional benefits of probiotics, such as the increase in immune system response and the release of bioactive compounds (Hempel et al., 2012; Bomba et al., 2012). The release of bioactive peptides from a protein sequence can be produced by enzymatic hydrolysis (Udenigwe & Aluko, 2012; Hartmann & Meisel, 2007), by fermentation processes (Hafeez et al., 2014; Udenigwe & Aluko, 2012; Quirós et al., 2005), or by a combination of both and during gastrointestinal digestion of food (Saito, 2008).

Bioactive peptides are included in the group of protein ingredients which produce functional foods (Nagpal et al., 2011). Within the group of bioactive peptides, the ones that have attracted higher scientific and industrial interest are those having antihypertensive activity. Interest in studying these compounds has arisen from the increase in mortality in industrialized countries due to hypertension and/or renal, heart or brain complications (Rojas-Ronquillo et al., 2012; Contreras et al., 2009; Nielsen et al., 2009; Saito, 2008; Hartmann

& Meisel, 2007; Quirós et al., 2005; Wu & Ding, 2001; Takano, 1998).

Over the past few decades, fermented milks have been considered beneficial to health. This has been ascribed to the microorganisms involved in fermentation. Recent studies have proven that these benefits are also due to the different compounds released during the fermentation process, such as metabolites and other biologically active molecules (Figueroa-González et al., 2010). Numerous peptides with diverse biological activities have been isolated from fermented milks. The first fermented milk with proven antihypertensive activity was produced using a combination of Lactobacillus helveticus and Saccharomyces cerevisiae (Nakamura et al., 1995); the milk contained two angiotensin-converting enzyme inhibiting tripeptides (ACE). Accordingly, it has been demonstrated that different lactic bacteria (Lactobacillus, Lactococcus, and Streptococcus) are capable of producing bioactive peptides during the fermentation process and during cold storage (Phelan et al., 2009). Due to the short fermentation period during the production of fermented milks (2 to 4 hours), proteolytic activity is low. This activity represents the hydrolysis of 1 to 2% of the existing proteins, of which 80% is carried out on caseins (Meisel & Bockelmann, 1999). However, at the end of fermentation and during refrigerated storage, BAL are capable of surviving (Vinderola et al., 2000) and their proteolytic activity is maintained (Nielsen et al., 2009). Despite the fact that the optimum temperature for the enzymes belonging to the proteolytic system of lactic bacteria ranges between 35 and 37°C, their activity is not lost during refrigeration at 4°C (González-Olivares et al., 2011). During refrigeration; peptides

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are generated by the action of the enzymes associated to live bacteria present in the medium or in enzymes released after cell lysis (Nighswonger et al., 1996). Many studies, focused on strain improvement and preservation techniques, have shown the viability of probiotic bacteria in commercial fermented milk under refrigerated storage (Adhikari et al., 2000; Donkor et al., 2006; Donkor et al., 2007b). However, studies on peptide production during cold storage have merely focused on comparing proteolytic profiles at the beginning and the end of the storage period (Nielsen et al., 2009). Other studies have primarily focused on the combination of indirect measurement of microbial activity of bacteria in yogurt through proteolytic profile comparison during storage at 4 °C (Donkor et al., 2006). It has also been proven that after 28 days of refrigeration at 4°C, the antihypertensive activity of yogurt increases from 70% to 90%, as calculated from half maximal inhibitory concentration  $(IC_{50})$  (Donkor et al., 2007a).

Nevertheless, there is no information on the molecular weight of peptides and the presence of aromatic amino acids in the polypeptide chains in commercial fermented milks stored under refrigeration. It has been reported that the presence of aromatic amino acids in the peptide chain contributes to the inhibitory effect of the ACE (Wu et al., 2006, Cheung et al., 1980). Therefore, the aim of this study was to determine the molecular weight and the presence of aromatic amino acids of peptides released by LAB in a commercial fermented milk stored at 4°C for 13 days.

#### 2 Materials and methods

# 2.1 Sample preparation

Two samples (in duplicate) of two different batches of a commercial fermented milk were purchased from a convenience store; their expiration date was 15 days after the date of purchase. The commercial fermented milk was produced with *Lactobacillus acidophilus* and *Streptococcus thermophilus*. The samples were stored at 4°C for 13 days. Sample collection was carried out every third day, under sterile conditions. A volume of 5 mL of sample was used to determine the protein concentration.

For the analysis of peptides, 5mL of each sample were mixed with 5 mL of 12% (w/v) trichloroacetic acid in order to precipitate high molecular weight proteins (Jiménez-Guzmán et al., 2002). Both samples were centrifuged at 18000 rpm for 15 minutes at 4°C (Beckman J2-MI, IL, USA). The supernatant was cooled at 4°C and used for further analyses.

# Polyacrylamide gel electrophoresis (SDS-PAGE)

The changes in the protein concentration in the fermented milk during cold storage were determined according to Laemmli (1970). Tris-Glycine gel with 15% (w/v) acrylamide was used. Broad range molecular weight standards (Bio-Rad, Berkeley, CA, USA) were used. The gels were stained with Sypro Flamingo (Bio-Rad, Berkeley, CA, USA). Peptides of low molecular weight were separated by denaturing electrophoresis (Tris-tricine), as described by González-Olivares et al. (2011). These gels were analyzed using a Bio-Rad's (Berkeley, CA, USA) Gel-Doc image

analyzer. A solution of  $\beta$ -lactoglobulin (2 mg/mL) was used to determine the concentration of peptides and proteins.

# 2.2 Separation of peptides by Size exclusion HPLC

Supernatants of fermented milks were injected on a Lab alliance HPLC system (NY, USA) using a size exclusion column (Biosep-SEC 2000 Phenomenex  $^{\circ}$ , Torrance, CA, USA, particle size 5 mm, pore size 145 Å, 300 x 7.8 mm). A phosphate buffer solution at pH 6.8 was used as a mobile phase. The volume of the sample injected was 20  $\mu L$ . The analysis was performed at room temperature using a flow of 0.25 mL/min for 60 min. Detection was performed at 210, 257, and 280 nm using a diode array detector (Spectra System UV 6000 LP, Providence, RI, USA).

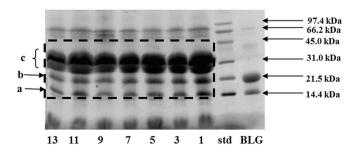
#### 2.3 Statistical analysis

All analyses were performed in triplicate. The results were analyzed by ANOVA and Tukey comparison test with a significance level of 0.05 using the SPSS Statistics 17.0 software (1993-2007 Polar Engineering and Consulting, Nikiski, AK, USA).

## 3 Results and discussion

# 3.1 Protein separation by electrophoresis

Casein concentration at the beginning of the experiment was 5.2 mg/mL, and it decreased to 2.9 mg/mL at the end of the study. The concentration of  $\beta$ -lactoglobulin (BLG) in the whey proteins decreased more than that of  $\alpha$ -lactoalbumin (ALA) (Figure 1). This can be explained because caseins are the main substrate of the proteolytic system of LAB, which enables the association of the accumulation of peptides in the medium with the decrease in the concentration of caseins. Furthermore, it is known that the nitrogen source of LAB in fermented milk comes from casein degradation (Gasson & de Vos, 1994; Savijoki et al., 2006). Moreover, even after the fermentation process is completed, LAB or at least the enzymes associated with their system maintain proteolytic activity during cold storage (Nielsen et al., 2009; Nighswonger et al., 1996). This may be due to the fact that the enzymes of the proteolytic system of LAB are proline-specific and there are four times more proline in the primary structure of BLG than in the structure of ALA (Savijoki et al., 2006).



**Figure 1**. Proteins for commercial fermented milk during refrigerated storage (4°C) in Tris-Glicine SDS-PAGE. BLG (β-lactoglobulin); std (standard of molecular weight); numbers (1-13) refer to the day of the experiment; a:  $\alpha$ -lactoalbumin, b:  $\beta$ -lactoglobulin, c: caseins.

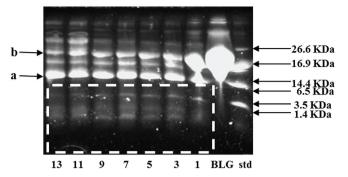
## 3.2 Peptide separation by electrophoresis

Seven peptides with a molecular weight lower than 14.4 kDa were separated and identified from day 1 of experiment. Similarly, bands of peptides of less than 1.4 kDa were identified (Figure 2). These results have raised the expectation of finding peptides with bioactive functions in refrigerated fermented milks since it has been reported that many of these compounds have a molecular weight of less than 10 kDa (Meisel, 2005). In addition, a decrease in the concentration of BLG and ALA with the same pattern of that found in the separation of proteins by electrophoresis was observed. Even after the preparation and refrigeration of fermented milk at 4°C, the proteolytic activity of LAB remained active (Nielsen et al., 2009). Moreover, several authors have asserted that the concentration of peptides derived from whey proteins is higher when they are released from BLG. This is due to the fact that the proline content in BLG is four times higher than that in ALA. This is related to the specificity of proteinases and peptidases in the proteolytic system of proline-specific lactic acid bacteria (Ortiz-Chao et al., 2009; Savijoki et al., 2006).

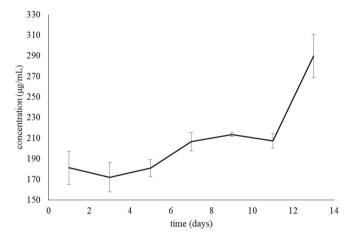
Initial concentration of peptides was 2.4 mg/mL; then there was a decrease (up to 1.8 mg/mL) in the first 48 hours of experiment. However, from day 5 to the end of the storage period, there was a tendency to peptide concentration increase, reaching a maximum value of 2.8 mg/mL. The decrease observed in the early days of storage could be due to the formation of smaller peptides from the peptides released in the fermentation by the endopeptidase action of LAB (Savijoki et al., 2006; Poolman et al., 1995). Concentration of BLG and ALA remained constant from day 7. This could be attributed to the fact that peptides are released from caseins during fermentation and frozen storage since they are the primary substrate of enzymes belonging to the proteolytic system of LAB (Gasson & de Vos, 1994). Donkor et al. (2007a) reported that an increase in peptide concentration in fermented milk was observed between weeks 1 and 3 of refrigerated storage, immediately after the end of the fermentation process, which is in agreement with the results obtained in this study.

# 3.3 Size exclusion HPLC

An increase in the concentration of peptides from day 5 was identified by HPLC analysis (Figure 3). Similarly, there was an increase in the number of peptides with great variability in molecular weight (mainly 6.21, 2.68, and 1.61 kDa) on each day of experiment. Statistical analysis showed a significant increase ( $P \ge 0.05$ ) in peptide concentration on day 7. However, it remained constant on days 9 and 11, reaching its maximum value on day 13 (exceeding 250 mg/mL). The two peptides with the highest concentration at the end of experiment were those of 2.68 kDa (59.55 mg/mL) and 0.87 kDa (101.84 mg/mL). It has been observed that during storage at 4°C, the production of peptides is higher in culture media with some strains of S. thermophilus and L. acidophilus compared with other probiotics in symbiotic fermentation or when they are alone in the fermentation (Vinderola et al., 2002). The commercial fermented milk under study had microorganisms of the same species. The viability of bacteria under cooling conditions is



**Figure 2.** Peptides in commercial fermented milk during refrigerated storage (4°C) in Tris-Tricine SDS-PAGE. BLG (β-lactoglobulin); std (standard of molecular weight); numbers (1-13) refer to the day of experiment. a:  $\alpha$ -lactoalbumin, b:  $\beta$ -lactoglobulin.



**Figure 3**. Peptide concentration of commercial fermented milk during 13 days of refrigerated storage (HPLC).

decisive in the formation of peptides (Donkor et al., 2007b), which was tested in this study (data not shown) through viable count.

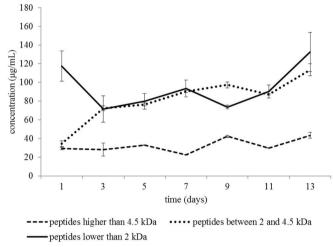
The calculations of concentration performed show both appearance and disappearance of peptides, especially those of more than 4.5 kDa. A peptide of 7.86 kDa appeared on the first and last days of refrigerated storage (Table 1). Similarly, a peptide of 5.62 kDa appeared only on day 3, and another one of 6.21 kDa reached maximum concentration (30.4 mg/mL) on day 13. Furthermore, a peptide of 4.5 kDa showed unstable behavior since its concentration increased and decreased frequently during the study. This peptide was observed throughout the entire refrigerated storage period. This behavior may have occurred due to the fact that these peptides are the source of formation of lower molecular weight peptides (Gasson & de Vos, 1994).

Peptide concentration was analyzed by groups of molecular weight: greater than 4.5 kDa, between 2 and 4.5 kDa, and smaller than 2 kDa (Figure 4). It was observed that the concentration of peptides in the first group was the lowest. This group also showed a varied behavior with a constant concentration on day 7 and increases and decreases

Table 1. Size of peptides found and their variation over refrigerated storage of commercial fermented milk at 4°C.

Molecular weight kDa	Presence of peptides over refrigerated storage						
	Day 1	Day 3	Day 5	Day 7	Day 9	Day 11	Day 13
7.86	$P^0$	A	A	A	A	A	P(+)*
6.21	A	$P^0$	P(+)	A	P(-)	P(=)	$P(+)^*$
5.62	P	A	A	A	A	A	A
4.55	P	$P(-)^0$	P+	$P(+)^*$	P(=)	P(-)	P(+)
3.86	P	P(=)	P(=)	$P(-)^0$	A	P(+)	P(+)*
3.10	A	A	A	$P^{0}$	P(+)	P(+)*	A
2.68	A	$P^0$	P(+)	P(=)	P(+)	A	$P(+)^*$
1.61	A	P	P(+)	$P(-)^0$	P(+)	P(+)*	P(-)
1.43	$\mathbf{P}^*$	$P(-)^0$	P(=)	P(+)	P(-)	A	P(-)
1.14	$\mathbf{P}^*$	P(-)	P(=)	P(=)	P(+)	P(-) <sup>0</sup>	P(+)
0.87	P	P(-)	P(+)	P(+)	$P(-)^0$	P(+)	$P(+)^*$

P:presence, A:absence, (-)Decreased concentration compared to the preceding measurement, (+)Increased concentration compared to the preceding measurement, (=)Same concentration compared to the preceding measurement, one concentration, \*Higher concentration.



**Figure 4**. Protein concentration of commercial fermented milk under refrigerated storage in molecular weight groups.

up to the end of the refrigerated storage period. With respect to the second group, the concentrations increased on days 3, 7, 9, and 13. The third group exhibited behavior similar to that of the second group; however, there was a significant decrease (P = 0.05) in the concentration from days 7 to 9. This group showed a significant increase in the concentration at the end of the experiment and had the highest concentration. Gasson & de Vos (1994) have reported that the formation of low molecular weight peptides during fermentation follows a cascade pattern. In this model, the source for the formation of high molecular weight peptides are proteins; the substrate for the intermediate molecular weight peptides are peptides of high molecular weight; and finally, they are the substrate for low molecular weight peptides. Similarly, it has been reported that the proteolytic system of LAB is associated with the accumulation of peptides of different sizes and types, depending on the nutritional needs of each LAB (Poolman et al., 1995). With regard to the proteolytic system of LAB, the optimum temperature of peptidases and proteinases ranges between 25 and 37°C (Juille et al., 2005). However, it was found that their activity continues even under refrigeration since these enzymes remain breaking proteins and releasing peptides (Nielsen et al., 2009; Nighswonger et al., 1996).

Peptide separation was performed by chromatography using three different wavelengths: 210 nm for determination of peptide bonds, 257 nm for the determination of residues with phenylalanine (Phe), and 280 nm for tyrosine (Tyr) and tryptophan (Trp) residues (Creighton, 1990). During separation, more peptides with Phe residues than peptides with Tyr and Trp residues were detected. It was also observed that the peptides showing absorption at 280 nm had molecular weights of less than 3kDa (2.9, 1.3 and 0.87 kDa). It is known that the concentration and amino acid characteristics of the peptides released depend on the specificity of the enzymes in the proteolytic system of LAB (Marshall & Tamime, 1997). Furthermore, it has been reported that the hydrophobic character of peptides promotes ACE inhibitory activity (Saito, 2008). Moreover, the importance of the presence of aromatic amino acids has been highlighted, especially phenylalanine with proline in peptides with high ACE inhibitory activity in ending C-terminal (Rojas-Ronquillo et al., 2012; Wu & Ding, 2001; Iroyukifujita et al., 2000; Matsumura et al., 1993; Miyoshi et al., 1991).

#### 4 Conclusions

It was determined that the accumulation of low molecular weight peptides in refrigerated commercial fermented milk follows a cascade pattern. This could be due to the effect of proteinases and peptidases of the proteolytic system on lactic acid bacteria, which promote the formation and accumulation of low molecular weight peptides.

The aromatic amino acids in the peptide fractions obtained increase the likelihood of finding peptides with antihypertensive activity in refrigerated commercial fermented milk since several authors have proven the importance of aromatic amino acids in peptides showing that function.

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