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<b>Article Type</b>	Research article
<b>Article Title</b>	Angiotensin-I-Converting Enzyme Inhibitory Peptides in Goat Milk Fermented by Lactic Acid Bacteria Isolated from Fermented Food and Breast Milk
<b>Running Title (within 10 words)</b>	ACEI Peptides in Goat Milk Fermented by Lactic Acid Bacteria
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**Abstract** In this study, the inhibitory activity of Angiotensin-I-Converting Enzyme (ACEI) was evaluated in fermented goat milk fermented by lactic acid bacteria (LAB) from fermented foods and breast milk. Furthermore, the potential for ACEI peptides was identified of fermented goat milk with the highest ACEI activity. The proteolytic specificity of LAB was also evaluated. The 2% isolate was inoculated into reconstituted goat milk (11% w/v), then incubated at 37 °C until pH 4.6 was reached. The supernatant produced by centrifugation was analyzed for ACEI activity and total peptide. Viable counts of LAB and titratable acidity were also evaluated after fermentation. Peptide identification was carried out using Nano LC/MS/MS, and potential as an ACEI peptide was carried out based on a literature review. The result revealed that ACEI activity was produced in all samples (20.44-60.33%). Fermented goat milk by *Lc. lactis* ssp *lactis* BD17 produced the highest ACEI activity (60.33%;  $IC_{50}$   $0.297 \pm 0.10$  mg/mL) after 48 h incubation, viable counts  $>8$  Log CFU/mL, and peptide content  $4.037 \pm 0.27$ /mL. A total of 261 peptides were released, predominantly casein (93%). The proteolytic specificity of *Lc. lactis* ssp *lactis* BD17 through cleavage on the amino acid tyrosine, leucine, glutamic acid, and proline. A total of 21 peptides were identified as ACEI peptides. This study showed that one of the isolates from fermented food, namely *Lc. lactis* ssp *lactis* BD17, has the potential as a starter culture for the production of fermented goat milk which has functional properties as a source of antihypertensive peptides.

**Keywords** ACE inhibitory activity, antihypertensive peptides, goat milk fermented, proteolytic specificity, *Lc. lactis* ssp *lactis* BD17

## Introduction

Hypertension is a primary risk factor for cardiovascular diseases (CVDs), including stroke, heart attack, heart failure, and other complications related to structural damage to the cardiovascular system. In 2021, the World Health Organization (WHO) reported that CVDs are the leading cause of death globally. People who died from CVDs in 2019 were estimated at 17.9 million, representing 31% of all deaths worldwide. Heart attacks and strokes are the main causes of these deaths (85%) (WHO, 2021). Human blood pressure is regulated by a system called the "Renin Angiotensin Aldosterone system", in which Angiotensin-I-Converting Enzyme inhibitory (ACE) plays an important role. ACE could catalyze the conversion of the decapeptide angiotensin I to the potent vasoconstrictor angiotensin II. Furthermore, this enzyme hydrolyzes bradykinin and stimulates the release of aldosterone which causes vasoconstriction and fluid retention which increases blood pressure (Rai et al., 2017). Therefore, the treatment of clinical hypertension could be done by controlling ACE activity. ACE inhibitors such as captopril, enalapril, alacepril, lisinopril and ramipril are widely used in the clinical treatment of hypertension. However, the use of these synthetic drugs in some cases causes side effects such as coughing, increased blood calcium levels, decreased kidney function, angioedema, and skin rashes (Zeng et al., 2013). Several researchers through in vivo studies on rats with spontaneous hypertension (SHR) and humans with hypertension showed that ACE inhibitors without side effects could be obtained from food protein (Bravo et al., 2019; Chen et al., 2014; Seppo et al., 2003)

Food protein from milk and dairy products such as fermented milk is a source of ACE inhibitory (ACEI) peptides (Begunova et al., 2021; Wu et al., 2019). Among them have been reported from fermented goat milk. The presence of ACE inhibitors in fermented milk is associated with the

55 presence of lactic acid bacteria (LAB). Lactic acid bacteria are the dominant group of bacteria  
56 involved in fermenting milk such as yogurt and kefir. Kefir is a fermented goat milk that has been  
57 consumed for hundreds of years and is believed not only as a source of antihypertensive peptides  
58 but also as a source of antioxidants and immunological agents (Ibrahim et al., 2017; Parmar et al.,  
59 2020)

60 During fermentation, milk protein could be hydrolyzed by LAB into peptides and amino acids.  
61 The abundance and characteristics of peptides released from milk proteins by LAB are strain-  
62 dependent (Wang et al., 2015). Among these peptides, the presence of bioactive peptides could be  
63 identified (Li et al., 2017). Bioactive peptides differ in size and sequence. Bioactive peptides that  
64 have functional properties as ACEI peptides have the characteristics that their molecular weight is  
65 generally >3 kDa and the presence of the amino acids proline and phenylalanine in the sequence  
66 (Gonzalez-Gonzalez et al., 2013; Wu et al., 2006)

67 The ability of LAB to release bioactive peptides in fermented milk (Ayyash et al., 2020; Kim et  
68 al., 2017), and the status of LAB as “generally recognized as safe” (GRAS) for application in food,  
69 have increased the utilization of certain strains of LAB for production of fermented milk with  
70 certain functional properties. The purpose of this study was to investigate ACEI activity in goat  
71 milk fermented using LAB from fermented foods and breast milk. The potential of ACEI peptides  
72 was identified in the <3 kDa fraction of fermented goat milk with the highest ACEI activity. The  
73 proteolytic specificity of the LAB used was also evaluated. The ten strains used were selected  
74 because they effectively released ACEI peptides in fermented cow milk in our previous study  
75 (Rubak et al., 2020).

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## **Materials and methods**

### **Lactic acid bacteria**

The ten LAB isolates from fermented foods and breast milk used in this study were culture collections from the Laboratory of Food Microbiology, Southeast Asian Food and Agricultural Science and Technology (SEAFAST) Center, IPB University (Bogor Agricultural University). The isolates were refreshed in de Man Rogosa and Sharpe broth and incubated at 37 °C for 24 h, then adapted in fresh skimmed milk for 2 rounds (24 h, 37 °C) before being used as a starter culture in the experiment.

### **Fermentation of goat milk**

Goat skimmed milk 11% (w/v) was pasteurized at 95 °C for 10 min. After cooling (45 °C), LAB starter culture (2%) was inoculated followed by incubation at 37 °C until pH 4.6 (700 Eutech) was reached. The fermentation process was stopped by heating (75 °C for 1 min) followed by centrifugation (Hettich, Zentrifugen, Mikro 22R) at 6000 x g for 10 min, 4°C. The supernatant was collected for analysis of peptide content and ACE inhibitory activity (Chusman, 1971). Viable counts of LAB and titratable acidity were also analyzed from unheated samples.

### **Determination of ACE Inhibitory Activity**

Hippuryl-L-Histidyl-L-Leucine (HHL, Sigma, USA) was used as an enzyme-substrate. A total of 50  $\mu$ L of the substrate (50 mM HHL in 0.1 M sodium borate buffer containing 0.3 M NaCl at pH 8.3) was added into a 50  $\mu$ L sample and incubated at 37 °C for 5 min. To initiate the reaction, 50  $\mu$ L of 0.1 U/mL ACE (Rabbit lung, Sigma, USA) solution was added, and the mixture was incubated at 37 °C for 5 min. The reaction was stopped by adding 250  $\mu$ L 1 M HCl. The resulted hippuric acid (HA) was extracted with 1.5 mL ethyl acetate and centrifuged at 2000 x g for 5 min. An aliquot (0.8 mL) of the ethyl acetate layer was transferred to a clean tube and evaporated at 85 °C for 60 min. Distilled water (4 mL) was then added to dissolve the HA in the tube, and the amount of HA formed was measured by measuring the optical density at 228 nm (UV-2800, Hitachi, JPN). The extent of inhibition was calculated as 100% [(B-A)/B] where A is the optical density in the presence of ACE and ACEI components, and B is the optical density without the ACEI component.

#### **IC<sub>50</sub> and inhibitory efficiency ratio (IER) value**

The IC<sub>50</sub> of the sample having the highest ACEI activity was calculated from the linear regression equation by plotting the ACE inhibition (%) versus the inhibitory concentration for each sample dilution. The percentage of ACEI activity was divided by the peptide concentration to obtain the IER value.

#### **Ultrafiltration**

The supernatant of fermented goat milk (4 mL) was pipetted into ultrafiltration centrifuge tubes [molecular weight (MW) cut-off of 3 kDa; Merck, 4 mL, IRL], then centrifuged at 4000 x g for 30 min, 4 °C. The fractions (<3 kDa and >3 kDa) were collected and the volume was adjusted to 4 mL by addition of water. Fractions were analyzed for ACEI activity.

### **Identification of peptides by mass spectrometry**

Peptides in <3 kDa fraction were analyzed by using LC Ultimate 3000 series system Tandem Q Exactive Plus Orbitrap HRMS (Thermo scientific, GER). The samples (5 µL) were injected into the LC Nano MS/MS system. The samples were trapped on a trap column (164649, 30 µm x 5 mm; Thermo scientific) and washed for 6 min with a gradient of 98% solvent A [water/acetonitrile (98:2, v/v), 0.1% formic acid] and 2% solvent B [Water/acetonitrile (2:98, v/v), 0.1% formic acid] at a flow rate of 5 µL/min. The eluted peptides were loaded and separated on a capillary column (PepMap RSLC-C18, 75-µm ×150 mm, 3.5 µm particle size, 100 pore size, Thermo Scientific ES800) at a flow rate of 300 nL/min with a gradient at 2% to 35% solvent B over 30 min, then from 35% to 90% over ten min, followed by 90% solvent B for 5 min, and finally 5% solvent B for 15 min. Electrospray was performed at an ion spray voltage of 3500 eV. Automatically, the peptides were analyzed using Proteomic Discoverer 2.2 software. The range of m/z values was 200–2000.

Identification of ACEI peptides was carried out through a literature search. The investigated is a peptide that provides 100% similarity to the ACEI peptide that has been previously reported by the researchers.



## Statistical analysis

The data were analyzed using Analysis of Variance (ANOVA) performed using SPSS version 22.  $P \leq 0.05$  was considered significant. Each experiments were repeated three times, and the data were presented as mean  $\pm$  SD.

## Result and Discussion

### Characteristics of goat milk fermented by LAB from fermented food and breast milk

Milk has been known as a suitable growth medium for LAB. The population of the ten LAB in fermented goat milk reached 9 log CFU/mL. Viable counts of the LAB ranged from  $9.18 \pm 0.46$  to  $9.79 \pm 0.39$  log CFU/mL. When the pH reached 4.6, there was no difference in population (Table 1), which is in accordance with previous results obtained by Elkhtab et al. (2017) and from fermented cow milk using similar cultures (Rubak et al., 2020). However, the fermentation time to reach pH 4.6 was different between isolates that ranged from 18 to 48 h with titratable acidity ranging from  $0.77 \pm 0.06$  to  $0.94 \pm 0.04\%$ . A short fermentation time (18 h) was observed in the fermentation by *Lb. rhamnosus* R2, while the longest fermentation time (48 h) occurred in the fermentation by *Lb. fermentum* S206, *Lb. delbrueckii* BD7, and *Lc. lactis* ssp *lactis* BD17. In our research the fermentation was ended at pH 4.6 to obtain high production of peptides. The release of a bioactive peptide from the protein matrix by culture could decrease when the pH value falls below 4.5 (Gonzalez-Gonzalez et al., 2013). Increase in coagulation could inhibit bacterial cell

diffusion to protein tissue, thus inhibiting access of CEP to milk protein for hydrolysis. Further acidification could be avoided by stopping fermentation when it reaches pH 4.6, or the pH must be controlled by adding alkaline solutions such as sodium hydroxide (Chen et al., 2015).

Insert Table 1

### **ACE inhibitory activity**

ACE inhibitory activity was detected in all supernatants of goat milk fermented in the range of  $20.44 \pm 2.33$  to  $60.79 \pm 8.78\%$  (Table 2). The highest percentage of ACEI activity (>50%) was obtained in goat milk fermented by *Lb. delbrueckii* BD7 and *Lc. lactis ssp lactis* BD17, but it was not significantly different (>0.05) from that of *Lb. kefir* YK4 and *Lb. kefir* JK17. It has been reported that goat milk could be used as a potential precursor for the production of ACE inhibitors through the fermentation process (Izquierdo-González et al., 2019). Starter cultures of LAB, growth conditions, and substrate are factors that influence ACEI production in fermented milk (Wang et al., 2015; Shu et al., 2015; Li et al., 2017). *Lactobacillus* species are known to produce high ACEI activity (>50%) in fermented milk (Hati et al., 2018; Wu et al., 2019). The variation of ACEI activity between LAB in milk fermented is related to its proteolytic activity. The proteolytic activity of LAB is determined by the specificity of its proteolytic components (Chen et al., 2015).

The proteolytic activity of LAB in goat milk was measured by the OPA method, with results ranging from  $3.55 \pm 0.26$  to  $5.69 \pm 0.21$  mg/mL (Table 2). Goat milk fermented by *P. pentasaceus* 1 W2SR04 and *Lb. kefir* YK4 (>5 mg/mL) showed high peptide content. In a study by Toe et al. (2019), *P. pentasaceus* species also showed high proteolytic activity. However, high peptide

content was not always associated with high ACEI activity in samples. This is also seen in the results of our study. ACE inhibitory activity is more related to the abundance of ACEI peptides that could be released during fermentation.

The IER values evaluated in ten samples showed that the highest IER values were obtained in fermented goat milk of *Lc. lactis* ssp *lactis* BD17. The IC<sub>50</sub> value was also determined in this sample and the result was  $0.297 \pm 0.10$  mg/mL. The IC<sub>50</sub> value reflects the peptide concentration required to inhibit 50% ACE. The IC<sub>50</sub> value in fermented milk <1 mg/mL (Gútiérrez et al., 2013). Our results show that the obtained IC<sub>50</sub> value is lower than that of fermented milk of other *Lactobacillus* species, as reported by Qian et al. (2011) in fermented milk by *Lb. delbrueckii* (IC<sub>50</sub>:  $67.71 \pm 7.62$  mg/mL), by Moslehishad et al. (2013) in fermented milk by *Lb. rhamnosus* (IC<sub>50</sub>:  $3.947 \pm 0.029$  mg/mL) and by Chen et al. (2007) in fermented milk using several isolates (IC<sub>50</sub>: 0.65 mg/mL) and in koumiss ( $52.47 \pm 2.87$  mg/mL) (Chen et al., 2010). Barla et al. (2016) have also reported from fermented milk by *Lb. brevis*, *Lb. buchnery* and *W. hellenica* (IC<sub>50</sub>: 0.28-0.83 mg/mL).

Insert Table 2

#### **ACE inhibitory activity of > 3 kDa and <3 kDa fraction**

Filtration using a 3 kDa MW cut-off showed that the ACEI activity was concentrated in the MW fraction <3 kDa (Table 3). There was no significant difference (>0.05) in the ACEI activity of the <3 kDa fraction compared to the supernatant (without filtration). Bioactive peptides with

ACEI activity have been reported as peptides with MW of < 3 kDa (Gonzalez-Gonzalez et al., 2013).

Insert Table 3

#### **Characteristics of peptides in the <3 kDa fraction of fermented goat milk of *Lc. lactis* ssp *lactis* BD17**

A total of 261 peptides were released in fermented goat milk by *Lc. lactis* ssp *lactis* BD17 (Table 4 and Supplementary Data). Most of the peptides were hydrolyzed from casein (97%) and whey (3%). The main fraction of goat milk protein is casein, which is 80% of the total milk protein (Jandal, 1996). This explains the abundance of peptide hydrolyzed from casein in our results. Another thing that casein has a very flexible and open structure so it is very sensitive to proteolysis. While whey protein is more resistant which is explained by the presence of a globular structure (Swaisgood, 1993). According to the results,  $\beta$ -casein (54.02%) was the most accessible to the proteolytic system of *Lc. lactis* ssp *lactis* BD17 to release a number of peptides. The cleavage site's dominance on the  $\beta$ -casein was also shown by *Lb. rhamnosus* CGMCC11055 (Guo et al., 2016) and *L. delbrueckii* subsp. *lactis* ACA-DC 178 (Hebert et al., 2008).

Insert Table 4

The spectrum of MS analysis revealed a large number of peaks with retention times of 8 to 65 min (Figure 1), representing an abundance of released peptides with peptide mass/molecular

charge (m/z) ranging from 310.1 to 1146.0. The diversity of detected ions indicates that the nine peaks with retention time (RT) of 11.29 to 30.97 corresponded to thirteen peptides identified in goat milk fermented by *Lc. lactis* ssp *lactis* BD17 (Table 5). ARHPPHLSFM ( $\kappa$ -casein; RT 11.29, 11.89; m/z 665.33, 673.33) was a peptide that exhibited a prominent peak according to its abundance in the sample. This peptide was also identified as being present in goat milk kefir (Izquierdo-Gonzales et al., 2019) and goat milk (Ibrahim et al., 2017). Another signal was associated with the abundance of peptides in goat milk fermented by *Lc. lactis* ssp *lactis* BD17 from the parent protein  $\beta$ -casein, namely MPFPKYPVEPF (RT 20.54; m/z 676.34), QEPVLGPVRGPFPI (RT 29.61; m/z 753.43) and RDMPIQAFL (RT 23.59; m/z 602.33). This peptide has also been identified in goat milk kefir (Izquierdo-Gonzales et al., 2019) and bovine kefir (Ebner et al., 2015).

Insert Fig 1

Insert Table 5

The release of peptides from the protein matrix is initiated by *Cell Envelope Proteinase* (CEP), one of the essential enzymes in the LAB proteolytic system (Griffiths & Tellez, 2013) which cleaves the proteins resulting in peptides with 4 to 30 amino acid. The CEP of LAB is classified into three types based on the hydrolysis of casein (Kunji, 1996): (1) CEP type PI which specifically hydrolyzes  $\beta$ -casein, (2) CEP type PIII which hydrolyzes  $\alpha$ S1-casein and  $\kappa$ -casein, (3) CEP intermediate type PI/PIII which hydrolyzes  $\beta$ -casein and  $\alpha$ S1-casein. Based on this classification, our results indicate that the CEP of *Lc. lactis* ssp *lactis* BD17 represents CEP types I and III. These

types of CEP were also reported from *Lb. casei* PRA205 and *Lb. rhamnosus* PRA331(Solieri et al., 2018), and *Lb. paracasei* ssp. *Paracasei* (Nikolić et al., 2009).

Types and domains in the CEP region of each LAB provide variations in the specificity of the hydrolyzed substrate which have implications for the diversity of molecular weights and amino acid sequences of the released peptides (Raveschot et al., 2020). The molecular weight of the peptide released by *Lc. lactis* ssp *lactis* BD17 ranged from 659 to 2201.1 Dalton with amino acid residues ranging from 6 to 20. An investigation was carried out to determine the specificity of cleavage of CEP *Lc. lactis* ssp *lactis* BD17 in goat milk parent protein during fermentation to release a number of peptides. The results are presented in Figure 2. It appears that sites of *Lc. lactis* ssp *lactis* BD17 dispersed throughout the parent protein, although certain amino acids are favorite for cleavage by CEP of *Lc. lactis* ssp *lactis* BD17. In the  $\alpha$ S1-casein region, the cleavage sites were frequently at serine, aspartate, and phenylalanine amino acid (f183-f184, f186-f187, and f184-f185), in  $\alpha$ S2-casein region the sites were at amino acids tyrosine, leucine, and glutamine (f116-f117, f114-f115, f112-f113), and in the  $\beta$ -casein region the sites were at amino acids tyrosine, leucine, glutamic acid and proline (f208-f209, f207-f208, f123-f124, f125-f126). Moreover, in a  $\kappa$ -casein region the cleavage sites were at amino acids leucine, tyrosine, and alanine (f53-f54, f51-f52, f44-f45). These results indicate that the cleavage sites of casein by *Lc. lactis* ssp *lactis* BD17 occur mostly in hydrophobic and aromatic amino acids. It seems that hydrophobic and aromatic amino acids are more easily accessed and released from parent protein. Similar results have been reported by Lozo et al. (2011) on *Lb. subsp. paracasei* BGHN14 (prtp), *Lb. rhamnosus* BGT10 (prtR), and *Lb. helveticus* BGRA43 (prtP), and Hebert et al. (2008) reported on *Lb. delbrueckii* subsp. *lactis* CRL 581.

Insert Fig 2

### ACE inhibitory peptides

Twenty-one of the 261 peptides released by *Lc. lactis ssp lactis* BD17 were identified as ACEI peptides (Table 5), most of which were released from  $\beta$ -casein. One of peptides namely ARHPHPHLSFM from parent protein  $\kappa$ -casein; 116-f117 was reported as an ACEI peptide (Ibrahim et al., 2017), and VLNENLR ( $\alpha$ S1-casein; f39-f40) (Swaisgood, 1993). ACE inhibitory peptides could be identified based on their amino acid sequence (Lunow et al., 2015).

Insert Table 6

The three amino acids located at the C-terminus could determine whether a peptide could act as an ACEI peptide (Wu et al., 2006). Amino acids from aliphatic groups (proline, isoleucine, valine) and aromatic amino acids (phenylalanine) are the dominant amino acids found in the ACEI peptide. Our investigation of other peptide released by *Lc. lactis ssp lactis* BD17 which has not been identified as an ACEI peptide according to a literature search, demonstrated its potential as an ACEI peptide. A total of 36% of these peptides had a proline amino acid residue and 21% a phenylalanine amino acid residue at the C-terminus.

The characteristics of the ACEI peptide were not only observed in the presence of amino acids at the C-terminus. By other researchers, the presence of amino acids at the N-terminal was also evaluated. Aslam et al. (2018) showed that three identified ACEI peptides were released in goat milk fermented by *Lb. helveticus* cicc22171 has hydrophobic/aliphatic amino acids not only at the

C-terminus but also at the N-terminus (valine and proline). Daliri et al. (2018) also presented their research results that four peptides identified as ACEI peptides were associated with the presence of negative amino acids (glutamate) and uncharged amino acids (glutamine) at their N-terminus. The presence of this amino acid in the *Lc. lactis* ssp *lactis* BD17 peptide was also identified, namely in the peptide released from the parent protein  $\beta$ -casein and  $\alpha$ -S1-casein (Table 4).

In addition to the presence of certain amino acids in the ACEI peptide sequence. Another characteristic of ACEI peptides is their molecular weight. ACE inhibitory peptides are generally short peptides with molecular weight < 3 kDa, may consist of 6 to 16 amino acids (Ibrahim et al., 2017). However, ACEI peptides with 20 amino acid residues have also been reported (Elkhtab et al., 2017). The ACEI peptide identified in our study has a molecular weight of <2 kDa (657-1994 Da), consisting of 6 to 18 amino acids. Short peptides are known to easily bind to the active site of ACE (Aslam et al., 2018). ACE inhibitory peptide binding to the active of ACE is facilitated by hydrogen bonding, hydrophobic interactions, and disrupting the stability of the  $Zn^{+2}$  ion. The presence of ACEI peptide in fermented milk is highly dependent on the type of LAB used for the fermentation process. it is therefore very important to use isolates that have been shown to have the ability to release ACEI peptides. *Lc. lactis* ssp *lactis* BD17 used in this study was an isolate isolated from kefir. Although the ability of this strain has only been explored in this study, as a comparison, the results of studies using kefir grain could be presented. Ebner et al. (2015) stated that kefir microbes were able to release 12 ACEI peptides in fermented milk. The same thing was also conveyed by Dallas et al. (2016) that kefir microbes release 29 bioactive peptides, including ACEI peptides in fermented cow milk. A recent study by Izquierdo-González et al. (2019) showed that in goat milk using kefir grains, five ACEI peptides were identified.



## Conclusion

ACE inhibitory activity was detected in all fermented goat milk using isolates from fermented food and breast milk. Goat milk fermented using *Lc. lactis* ssp *lactis* BD17 produced the highest ACE inhibitory activity ( $IC_{50}$ :  $0.297 \pm 0.10$  mg/mL) after 48 h of incubation. A total of 261 peptides were hydrolyzed by *Lc. lactis* ssp *lactis* BD17 during fermentation, most of which were released from casein ( $\beta$ -casein). The peptide has a molecular weight of 659 to 2201.1 Dalton, consisting of 6-20 amino acid residues. The CEP specificity of *Lc. lactis* ssp *lactis* BD17 in goat milk parent protein, dominant with cleavage on the amino acids tyrosine, leucine, glutamic acid, and proline. A total of 21 peptides were identified as ACEI peptides, having 100% homology to the reported ACEI peptides. Several characteristics of ACEI peptides are present in peptides hydrolyzed by *Lc. lactis* ssp *lactis* BD17. These peptides mostly have hydrophobic and aromatic amino acids at the C-terminus. The results of this study add to the information that *Lc. lactis* ssp *lactis* BD17 is a candidate that could be considered as a starter culture to obtain fermented milk that has functional properties as a source of ACE inhibitory peptides.

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## Conflicts of interest

The authors declare no conflicts of interest

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## Tables and Figures

**Table 1.** Profile of goat milk fermented by LAB from kefir grains and breast milk

Culture	Fermentation time to reach pH 4.6 (h)	Titrateable acidity (%)	Viable count (Log CFU/mL)
<i>Lactobacillus rhamnosus</i> R2	18	0.90 ± 0.01	9.69 ± 0.11
<i>Pediococcus pentasaceus</i> 1 W2SR04	24	0.94 ± 0.04	9.18 ± 0.46
<i>Lactobacillus kefir</i> YK4	24	0.80 ± 0.03	9.79 ± 0.39
<i>Lactobacillus kefir</i> JK17	24	0.79 ± 0.02	9.67 ± 0.07
<i>Lactobacillus fermentum</i> R6	24	0.84 ± 0.01	9.58 ± 0.03
<i>Lactobacillus plantarum</i> 1W22408	32	0.77 ± 0.06	9.25 ± 0.39
<i>Lactobacillus</i> R7F	36	0.81 ± 0.02	9.46 ± 0.52
<i>Lactobacillus fermentum</i> S206	48	0.70 ± 0.06	9.53 ± 0.17
<i>Lactobacillus delbrueckii</i> BD7	48	0.75 ± 0.05	9.73 ± 0.36
<i>Lactococcus lactis</i> ssp <i>lactis</i> BD17	48	0.75 ± 0.05	9.55 ± 0.38

**Table 2.** ACE inhibitory activity, peptide content, and IER of goat milk fermented by LAB after reaching pH 4.6

Culture	Inhibition of ACE (%)	Peptide content (mg/mL)	IER (% per mg/mL)
<i>Pediococcus pentasaceus</i> 1 W2SR04	20.44 ± 2.33 <sup>c</sup>	5.696 ± 0.21 <sup>a</sup>	3.60 ± 0.49 <sup>c</sup>
<i>Lactobacillus kefir</i> YK4	58.65 ± 8.87 <sup>a</sup>	5.691 ± 0.25 <sup>a</sup>	10.31 ± 1.55 <sup>b</sup>
<i>Lactobacillus fermentum</i> S206	26.64 ± 3.90 <sup>c</sup>	4.809 ± 0.23 <sup>b</sup>	5.59 ± 1.04 <sup>c</sup>
<i>Lactobacillus delbrueckii</i> BD7	60.79 ± 8.78 <sup>a</sup>	4.767 ± 0.27 <sup>b</sup>	12.81 ± 2.12 <sup>ab</sup>
<i>Lactobacillus kefir</i> JK17	56.94 ± 2.81 <sup>a</sup>	4.750 ± 0.30 <sup>b</sup>	12.06 ± 1.23 <sup>b</sup>
<i>Lactobacillus fermentum</i> R6	48.50 ± 6.92 <sup>ab</sup>	4.089 ± 0.10 <sup>c</sup>	11.83 ± 1.42 <sup>b</sup>
<i>Lactobacillus</i> R7F	42.85 ± 5.10 <sup>b</sup>	4.007 ± 0.23 <sup>c</sup>	10.66 ± 0.73 <sup>b</sup>
<i>Lactococcus lactis</i> ssp <i>lactis</i> BD17	60.33 ± 4.73 <sup>a</sup>	4.037 ± 0.27 <sup>c</sup>	14.99 ± 1.26 <sup>a</sup>
<i>Lactobacillus rhamnosus</i> R2	40.51 ± 6.15 <sup>b</sup>	3.658 ± 0.19 <sup>c</sup>	11.03 ± 1.10 <sup>b</sup>
<i>Lactobacillus plantarum</i> I W22408	39.69 ± 4.70 <sup>b</sup>	3.548 ± 0.26 <sup>c</sup>	11.15 ± 0.53 <sup>b</sup>

<sup>a-c</sup> Different superscript in the same column indicated significant ( $p < 0.05$ )

**Table 3.** ACE inhibitor activity of supernatant (without ultrafiltration), fractions of >3 kDa and <3 kDa of goat milk fermented by *Lc. lactis* ssp *lactis* BD17

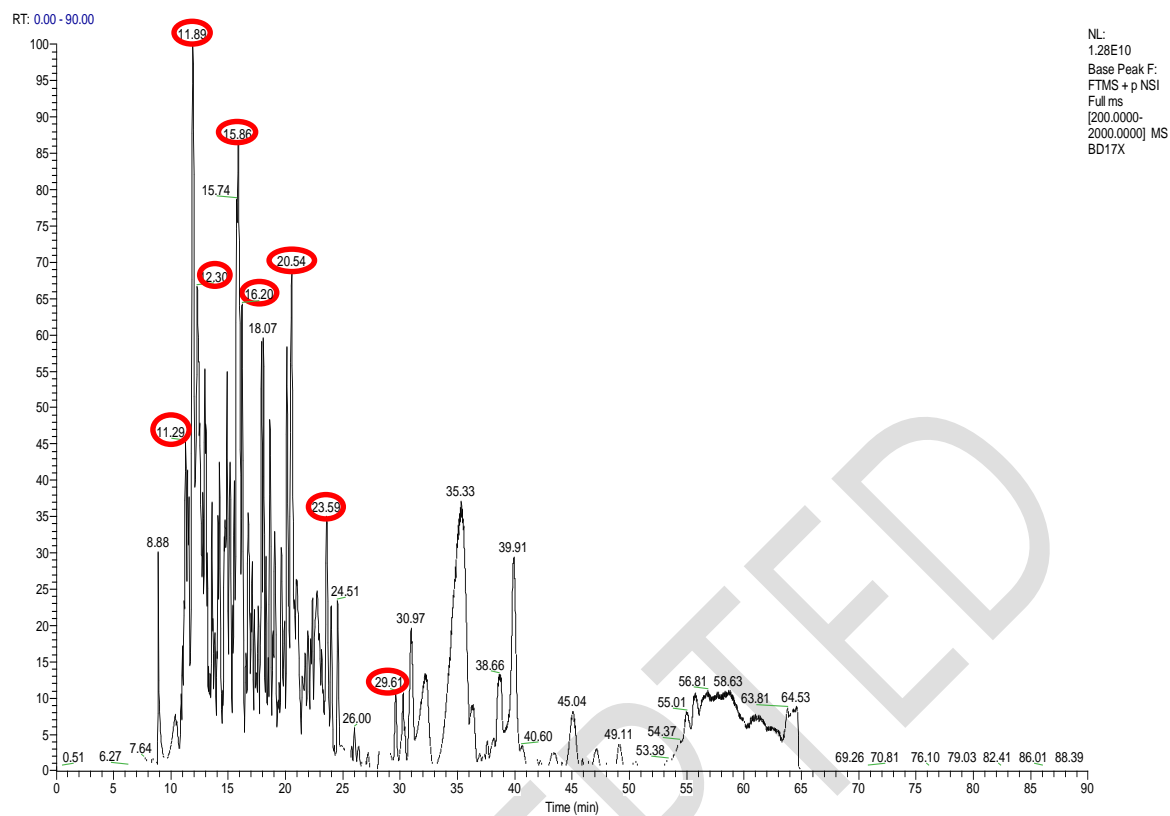
ACE inhibitory activity (%)		
Supernatant (Without ultrafiltration)	>3 kDa	<3 kDa
60.33 ± 4.73 <sup>a</sup>	24.57 ± 2.36 <sup>b</sup>	57.31 ± 2.41 <sup>a</sup>

**Table 4.** Characteristics of peptides (<3 kDa) of fermented goat milk of *Lc. lactis* ssp *lactis*

BD17

Parent protein*	Total Peptides	Range of precursor m/z	Range of precursor MW	Amino acid residue	Dominant peptide
$\alpha$ S1-casein	41	330.1-1146.0	659.3-2291.1	6 - 20	FSDIPNPIGSE, FSDIPNPIGSENSGKTTMP, NSGKTTMPLW
$\alpha$ S2-casein	37	310.1- 906.9	772.4-1812.9	6 - 15	QGPIVLNPWDQVQR
$\beta$ -casein	141	326.7-1072.5	652.40-2201.1	6 - 18	QEPVLGPVRGPFPII, QEPVLGPVRGPFPI, QEPVLGPVRGPFPIIV, VLGPVRGPFPIIV, TQTPVVVPPFLQPE
$\kappa$ -casein	36	376.5-750.8	761.45-1500.7	6 - 10	ARHPHPLSFM
$\alpha$ -lactalbumin	3	497.7-563.7	994.4-1126.4	8 - 10	AFHTSGYDTQ
$\beta$ -lactoglobulin	3	420.5-482.2	903.57-1259.7	8 - 11	

\*Protein access code at <https://www.uniprot.org/>. A: Alanine, D: Aspartic Acid, E: Glutamic Acid, F: Phenylalanine, G: Glysin, I: Isoleucine, K: Lysine, L: Leucine, M: Methionine, N: Asparagine, P: Proline, Q: Glutamine, R: Arginine, S: Serine, T: Threonin, V: Valine, W: Tryptophan, Y: Tyrosine, MW: Molecular Weight.



**Fig. 1.** Nano LC-MS/MS chromatogram of goat milk fermented by *Lc. lactis* ssp *lactis* BD17

577 **Table 5.** Peptides identified of goat milk fermented by *Lc. lactis* ssp *lactis* BD17 with Retention  
578 Time (RT) detected on Nano LC-MS/MS. Peptides in bold letters indicate a match with RT.  
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Retention Time (min)	Retention time (min)	m/z [Da]	MH <sup>+</sup> [Da]	Peptide identified	Parent Protein
11.29	11.29	665.33	1329.66	<b>ARHPPHLSFM</b>	$\kappa$ -casein (A0A140T8A9 <sup>*</sup> )
	11.22	495.75	990.49	ITVDDKHY	$\alpha$ 2-casein (P02663 <sup>*</sup> )
11.89	11.81	563.74	1126.48	AFHTSGYDTQ	$\alpha$ -lactalbumin (B6V3I5 <sup>*</sup> )
	11.87	581.34	1161.67	EKNRLNFLK	$\alpha$ 2-casein (P02663 <sup>*</sup> )
	11.89	673.33	1345.66	<b>ARHPPHLSFM</b>	$\kappa$ -casein (A0A140T8A9 <sup>*</sup> )
12.30	12.30	559.75	1118.49	<b>QQQTEDELQ</b>	$\beta$ -casein (P02666 <sup>*</sup> )
	12.30	546.79	1092.57	<b>YQKALNEIN</b>	$\alpha$ 2-casein (P02663 <sup>*</sup> )
15.86	15.86	470.74	940.47	<b>IPNPIGSEN</b>	$\alpha$ 1-casein (P02662 <sup>*</sup> )
	15.86	521.27	1041.54	<b>RYLGYLEQ</b>	$\alpha$ 1-casein (P02662 <sup>*</sup> )
16.20	16.20	413.72	826.43	<b>IPNPIGSE</b>	$\alpha$ 1-casein (P02662 <sup>*</sup> )
	16.20	560.78	1120.55	<b>FYQKFPQY</b>	$\alpha$ 2-casein (P02663 <sup>*</sup> )
	16.24	621.31	1241.61	DELQDKIHPF	$\beta$ -casein (P02666 <sup>*</sup> )
	16.26	457.24	913.48	RYLGYLE	$\alpha$ 1-casein (P02662 <sup>*</sup> )
	16.26	610.29	1219.57	SRYPYGLNY	$\kappa$ -casein (A0A140T8A9 <sup>*</sup> )
20.54	20.54	676.34	1351.68	<b>MPFPKYPVEPF</b>	$\beta$ -casein (P02666 <sup>*</sup> )
	20.59	329.20	657.40	VVVPPF	$\beta$ -casein (P02666 <sup>*</sup> )
23.59	23.50	683.86	1366.72	RDMPIQAFLLY	$\beta$ -casein (P02666 <sup>*</sup> )

Retention Time (min)	Retention time (min)	m/z [Da]	MH <sup>+</sup> [Da]	Peptide identified	Parent Protein
	23.51	559.80	1118.59	MFPPQSVLSL	β-casein (P02666*)
	23.53	385.74	770.48	VVVPPFL	β-casein (P02666*)
	23.54	948.04	1895.07	LYQEPVLGPVRGPFPII	β-casein (P02666*)
	23.56	795.47	1589.94	EPVLGPVRGPFPIIV	β-casein (P02666*)
	23.59	602.33	1203.66	<b>RDMPIQAFL</b>	β-casein (P02666*)
24.51	24.50	859.50	1718.00	QEPVLGPVRGPFPIIV	β-casein (P02666*)
	24.55	484.79	968.58	TPVVVPPFL	β-casein (P02666*)
26.00	26.07	548.30	1095.59	MPIQAFLLY	β-casein (P02666*)
	26.08	809.97	1618.93	QEPVLGPVRGPFPII	β-casein (P02666*)
29.61	29.61	753.42	1505.84	<b>QEPVLGPVRGPFPI</b>	β-casein (P02666*)
30.97	30.98	776.42	1551.83	TQTPVVVPPFLQPE	β-casein (P02666*)

580

581 \*Protein access code at <https://www.uniprot.org/>: Alanine, D: Aspartic Acid, E: Glutamic Acid, F:  
582 Phenylalanine, G: Glysin, I: Isoleucine, K: Lysine, L: Leucine, M: Methionine, N: Asparagine, P: Proline,  
583 Q: Glutamine, R: Arginine, S: Serine, T: Threonin, V: Valine, W: Tryptophan, Y: Tyrosine

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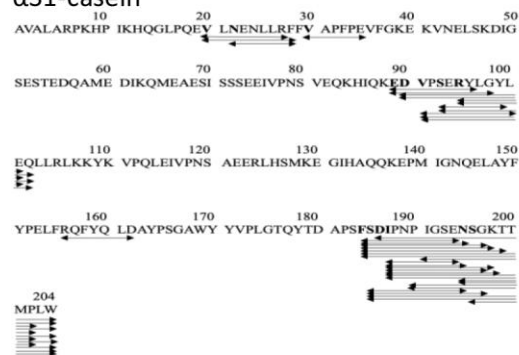
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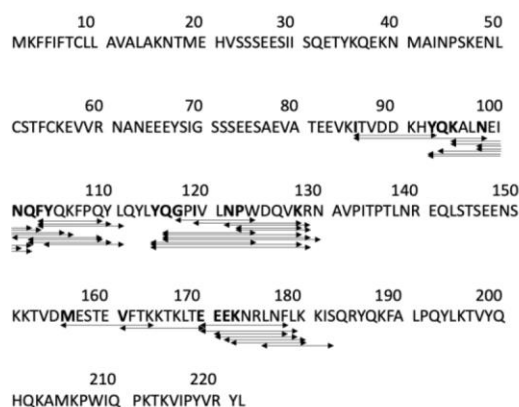
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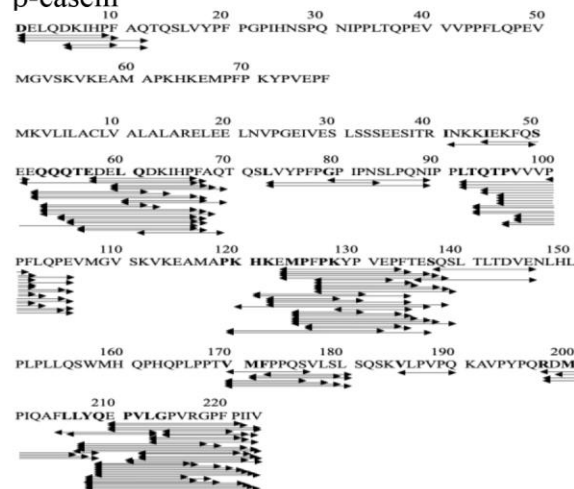
# $\alpha$ S1-casein



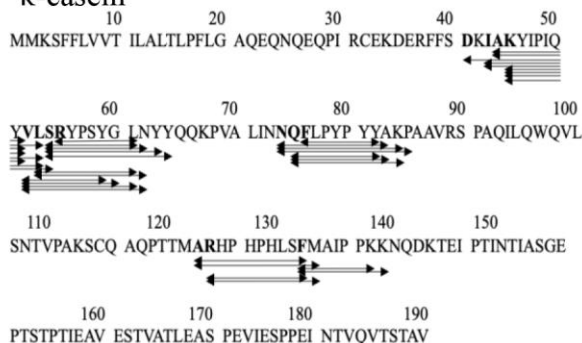
# $\alpha$ S2-casein



# $\beta$ -casein



# $\kappa$ -casein



**Fig. 2.** Cleavage specifics of the peptide of the parent protein  $\alpha$ S1-casein,  $\alpha$ S2- casein,  $\beta$ - casein, and  $\kappa$ - casein

596 **Table 6.** ACE inhibitory peptides in fermented goat milk of *Lc. lactis* ssp *lactis* BD17 as  
597 reported in the literature

Parent Protein*	Sequence	m/z [Da]	MH+ [Da]	Charge	References
κ-cas.	ARHPHPLSFM	665.33	1329.66	3	Ibrahim et al. (2017)
β-cas.	DELQDKIHPF	621.30	1241.61	2	Rodríguez-Figueroa et al. (2012)
β-cas.	DKIHPF	378.71	756.41	2	Fan et al. (2018)
β-cas.	DKIHPFAQ	478.25	955.50	2	Gobbetti et al. (2000)
β-cas.	EMPFPKYPVEPF	740.86	1480.71	2	Papadimitriou et al. (2007)
β-cas.	ELQDKIHPF	563.80	1126.59	2	Fan et al. (2018)
β-cas.	GPVRGPFPI	470.27	939.54	2	Amorim et al. (2019)
β-cas.	LGPVRGPFPI	470.27	939.54	2	Hernandes-Ledesma et al. (2004)
β-cas.	LTQTPVVVPPF	599.34	1197.68	2	Villegas et al. (2014)
β-cas.	LVYPFPGPIHNSLPQN	896.96	1792.93	2	Quirós et al. (2009)
β-cas.	LYQEPVLGPVRGPFPIIV	997.58	1994.14	2	Pihlanto et al. (2010)
β-cas.	MPFPKYPVEP	602.80	1204.60	2	Contreras et al. (2009)
β-cas.	MPFPKYPVEPF	676.34	1351.67	2	Hayes et al. (2007)
β-cas.	QEPVLGPVRGPFPI	696.88	1392.76	2	Hernandes-Ledesma et al. (2004)
β-cas.	QEPVLGPVRGPFPIIV	859.50	1718.00	2	Perpetuo et al. (2003)

Parent Protein*	Sequence	m/z [Da]	MH <sup>+</sup> [Da]	Charge	References
$\alpha$ s1-cas.	VLNENLR	485.78	970.56	2	Zhao et al. (2019)
$\beta$ -cas.	VLGPVRGPFP	519.81	1038.61	2	Gútiéz et al. (2013)
$\beta$ -cas.	VVVPPF	329.20	657.39	2	Torres-Llanez et al. (2011)
$\beta$ -cas.	YQEPVLGPVRGPFPI	834.95	1668.90	2	Zhao et al. (2019)
$\beta$ -cas.	YQEPVLGPVRGPFPIIV	627.69	1881.06	3	Zhao et al. (2019)
$\beta$ -cas.	YQEPVLGPVR	579.31	1157.63	2	Kalyankar et al. (2013)

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599 \*Protein access code at <https://www.uniprot.org/>. A: Alanine, D: Aspartic Acid, E: Glutamic Acid, F:

600 Phenylalanine, G: Glycine, I: Isoleucine, K: Lysine, L: Leucine, M: Methionine, N: Asparagine, P: Proline,

601 Q: Glutamine, R: Arginine, S: Serine, T: Threonine, V: Valine, W: Tryptophan, Y: Tyrosine, cas: casein

602

**Supplementari Dataset 1.** Peptide profile (<3 kDa) released from goat milk proteins during fermentation using *Lc. lactis* ssp *lactis* BD17

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
$\alpha$ s1-	DIPNPIGSENSGKTTMPLW	2057.00	1029.00	2057.00	1029.00	2	21.97
casein	DVPSERYLGYLE	1440.70	720.85	1440.70	720.85	2	19.95
	DVPSERYLGYLEQ	1568.76	784.88	1568.76	784.88	2	19.81
	EDVPSERY	994.45	497.73	994.45	497.73	2	12.21
	EDVPSERYLG	1164.55	582.78	1164.55	582.78	2	14.95
	FSDIPNPIGSE	1175.56	588.28	1175.56	588.28	2	19.45
	FSDIPNPIGSEN	1289.60	645.30	1289.60	645.30	2	19.16
	FSDIPNPIGSENSG	1433.65	717.33	1433.65	717.33	2	19.04
	FSDIPNPIGSENSGK	1561.75	781.38	1561.75	781.38	2	16.64
	FSDIPNPIGSENSGKT	1662.79	831.90	1662.80	831.90	2	16.62
	FSDIPNPIGSENSGKTTMP	1991.93	996.47	1991.94	996.47	2	17.96
	FSDIPNPIGSENSGKTTMPLW	2291.10	1146.05	2291.10	1146.05	2	21.96
	IGSENSGKTTMPLW	1520.74	760.87	1520.74	760.87	2	18.41
	IPNPIGSE	826.43	413.72	826.43	413.72	2	16.20
	IPNPIGSEN	940.47	470.74	940.47	470.74	2	15.86
	IPNPIGSENSEKT	1385.69	693.35	1385.69	693.35	2	14.46
	IPNPIGSENSG	1084.52	542.77	1084.53	542.77	2	15.65
	IPNPIGSENSGK	1212.62	606.81	1212.62	606.81	2	13.16
	IPNPIGSENSGKTTMP	1642.81	821.91	1642.81	821.91	2	15.69
	IPNPIGSENSGKTTMPLW	1941.97	971.49	1941.97	971.49	2	20.38
	NENLLRF	905.48	453.24	905.48	453.25	2	17.8

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	NSGKTTMPL	948.48	474.74	948.48	474.74	2	12.03
	NSGKTTMPLW	1134.56	567.78	1134.56	567.78	2	17.5
	PIGSENSGKTTMP	1318.63	659.82	1318.63	659.82	2	12.1
	PIGSENSGKTTMPLW	1617.79	809.40	1617.79	809.40	2	18.7
	RQFYQLD	969.48	485.24	969.48	485.24	2	16.04
	RYLGYLE	913.48	457.24	913.48	457.24	2	16.26
	RYLGYLEQ	1041.54	521.27	1041.54	521.27	2	15.86
	SDIPNPIGSEN	1142.53	571.77	1142.53	571.77	2	17.19
	SDIPNPIGSENSG	1286.58	643.79	1286.59	643.80	2	17.07
	SDIPNPIGSENSGKTTMP	1844.87	922.94	1844.87	922.94	2	16.72
	SDIPNPIGSENSGKTTMPLW	2144.03	1072.52	2144.03	1072.52	2	21.19
	SERYLG Y	887.42	444.22	887.43	444.22	2	14.12
	SERYLGYLE	1129.55	565.28	1129.55	565.28	2	17.28
	SGKTTMPLW	1020.52	510.76	1020.52	510.76	2	17.59
	VAPFPE	659.34	330.17	659.34	330.17	2	17.23
	VLNENLLR	970.57	485.79	970.57	485.79	2	15.29
	VLNENLLRF	1117.64	559.32	1117.64	559.32	2	19.78
	VPSEYR LGYL	1196.63	598.82	1196.63	598.82	2	18.44
	VPSEYR LGYLE	1325.67	663.34	1325.67	663.34	2	18.13
	VPSEYR LGYLEQ	1453.73	727.37	1453.73	727.37	2	17.95
$\alpha$ S2-	EEKNRLNF	1178.58	589.79	1178.58	589.79	2	12.64
casein	EEKNRLNFL	1291.66	646.33	1291.66	646.34	2	15.88
	EEKNRLNF	1049.54	525.27	1049.54	525.27	2	11.90

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	EEKNRLNFL	1162.62	581.81	1162.62	581.81	2	15.54
	EKNRLNFLK	1161.67	581.34	1161.67	581.34	2	11.87
	FYQKFPQ	957.48	479.24	957.48	479.25	2	14.06
	FYQKFPQY	1120.55	560.78	1120.55	560.78	2	16.20
	FYQKFPQYL	1233.63	617.32	1233.63	617.32	2	18.63
	GPIVLNPW	895.50	448.25	895.50	448.26	2	22.37
	ITVDDKHY	990.49	495.75	990.49	495.75	2	11.22
	ITVDDKHYQKALN	1544.80	515.61	1544.81	515.61	3	12.22
	IVLNPWDQVK	1211.68	606.34	1211.68	606.34	2	18.66
	KALNEINQ	929.51	465.26	929.51	465.26	2	8.98
	KALNEINQF	1076.57	538.79	1076.57	538.79	2	15.44
	KNRLNFLK	1032.63	344.88	1032.63	344.88	3	10.93
	LNFLKKIS	962.60	481.80	962.60	481.81	2	15.05
	MESTEVFTK	1071.50	536.25	1071.50	536.25	2	13.62
	NEINQFYQ	1055.48	528.24	1055.48	528.24	2	17.52
	NEINQFYQK	1183.57	592.29	1183.57	592.29	2	14.66
	NPWDQVKR	1042.54	348.19	1042.54	348.19	3	12.96
	NQFYQKFPQ	1199.58	600.30	1199.58	600.30	2	15.36
	PWDQVK	772.40	386.70	772.40	386.70	2	12.47
	PWDQVKR	928.50	310.17	928.50	310.17	3	10.59
	QFYQKFPQ	1085.54	543.27	1085.54	543.27	2	14.89
	QFYQKFPQY	1248.60	624.80	1248.60	624.81	2	16.83
	QGPIVLNPW	1023.56	512.28	1023.56	512.28	2	22.27
	QGPIVLNPWDQVK	1493.81	747.41	1493.81	747.41	2	19.51

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	QGPIVLNPWDQVQR	1649.90	825.46	1649.91	825.46	2	17.90
	QGPIVLNPWDQVKRN	1763.95	882.48	1763.96	882.48	2	17.75
	QKALNEINQ	1057.56	529.28	1057.56	529.29	2	10.95
	VFTKKTKLTE	1194.71	597.86	1194.71	597.86	2	9.81
	YQGPIVLNPW	1186.62	593.82	1186.63	593.82	2	23.21
	YQGPIVLNPWDQVK	1656.87	828.94	1656.87	828.94	2	20.32
	YQGPIVLNPWDQVQR	1812.97	906.99	1812.98	906.99	2	18.51
	YQKALNEIN	1092.57	546.79	1092.57	546.79	2	12.30
	YQKALNEINQ	1220.62	610.82	1220.63	610.82	2	12.46
	YQKFPQYL	1086.56	543.78	1086.56	543.78	2	17.42
β-casein	DELQDKIHP	1094.55	547.78	1094.55	547.78	2	12.66
	DELQDKIHPF	1241.61	621.31	1241.62	621.31	2	16.24
	DELQDKIHPFAQ	1440.71	480.91	1440.71	480.91	3	16.44
	DKIHPF	756.40	378.70	756.40	378.71	2	20.37
	DKIHPFAQ	955.50	478.25	955.50	478.25	2	12.10
	EDELQDKIHP	1223.59	408.53	1223.59	408.53	3	13.45
	EDELQDKIHPF	1370.66	685.83	1370.66	685.83	2	16.69
	EEQQQTEDELQDKIHP	1966.90	656.30	1966.90	656.30	3	14.60
	ELQDKIHPF	1126.59	563.80	1126.59	563.80	2	14.92
	ELQDKIHPFA	1197.62	599.32	1197.63	599.32	2	15.04
	ELQDKIHPFAQ	1325.68	663.35	1325.68	663.35	2	14.89
	EMPFPKYPVEP	1333.65	667.33	1333.65	667.33	2	18.71
	EMPFPKYPVEPF	1480.72	740.86	1480.72	740.86	2	21.40
	EMPFPKYPVEPFTE	1726.80	863.90	1726.80	863.91	2	19.59

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	EMPFPKYPVEPFTES	1813.83	907.42	1813.84	907.42	2	20.89
	EMPFPKYPVEPFTESQ	1941.89	971.45	1941.89	971.45	2	20.65
	EPVLGPVRGPFPP	1264.70	632.86	1264.70	632.86	2	18.78
	EPVLGPVRGPFPI	1377.79	689.40	1377.79	689.40	2	20.98
	EPVLGPVRGPFPII	1490.87	745.94	1490.87	745.94	2	22.40
	EPVLGPVRGPFPIIV	1589.95	795.48	1589.94	795.47	2	23.56
	EQQQTEDELQ	1247.54	624.27	1247.54	624.27	2	12.66
	EQQQTEDELQDK	1490.66	745.83	1490.66	745.83	2	10.45
	EQQQTEDELQDKIHP	1837.85	613.29	1837.86	613.29	3	13.82
	EQQQTEDELQDKIHPF	1984.92	992.96	1984.92	992.97	2	16.78
	FLLYQEPVL	1121.62	561.32	1121.62	561.32	2	24.24
	FPKYPVE	879.46	440.23	879.46	440.23	2	14.87
	FPKYPVEPF	1123.58	562.29	1123.58	562.29	2	19.12
	FPKYPVEPFT	1224.63	612.82	1224.63	612.82	2	18.77
	FPKYPVEPFTES	1440.70	720.85	1440.70	720.86	2	18.66
	FPKYPVEPFTESQ	1568.76	784.88	1568.76	784.89	2	18.65
	FPKYPVEPFTESQS	1655.79	828.40	1655.80	828.40	2	18.47
	FPPQSVLSL	987.55	494.28	987.55	494.28	2	22.89
	GPIPNSLPQN	1036.54	518.77	1036.54	518.77	2	16.11
	GPVRGPFPP	826.46	413.73	826.46	413.73	2	14.30
	GPVRGPFPI	939.54	470.27	939.54	470.27	2	17.48
	GPVRGPFPII	1052.62	526.82	1052.63	526.82	2	20.18
	HKEMPFPKYP	1273.64	637.32	1273.64	637.32	2	13.54
	HKEMPFPKYPVEPF	1745.87	873.44	1745.87	873.44	2	17.82



Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	IEKFQSE	880.44	440.72	880.44	440.72	2	10.32
	INKKIEKFQ	1147.68	574.34	1147.68	574.35	2	9.73
	KEMPFPK	876.46	438.74	876.46	438.74	2	11.13
	KEMPFPKYP	1136.58	568.79	1136.58	568.79	2	14.79
	KEMPFPKYPVEPF	1624.81	542.27	1624.81	542.27	3	17.85
	KEMPFPKYPVEPFTESQ	2053.99	1027.50	2053.99	1027.50	2	18.37
	KHKEMPFPKYPVEPLTE	2070.09	518.28	2070.07	518.27	4	16.31
	KYPVEPF	879.46	440.23	879.46	440.23	2	16.30
	LGPVRGPF	939.54	470.27	939.54	470.27	2	16.86
	LGPVRGPFPI	1052.62	526.82	1052.63	526.82	2	19.27
	LGPVRGPFPII	1165.71	583.36	1165.71	583.36	2	21.23
	LLYQEPVL	974.55	487.78	974.56	487.78	2	20.67
	LQDKIHPF	997.55	499.28	997.55	499.28	2	13.46
	LQDKIHPFAQ	1196.64	598.82	1196.64	598.82	2	13.32
	LTQTPVVVPP	1050.62	525.81	1050.62	525.81	2	17.35
	LTQTPVVVPPF	1197.69	599.35	1197.69	599.35	2	22.14
	LTQTPVVVPPFLQPE	1664.93	832.97	1664.93	832.97	2	24.04
	LVYPFPGPIHN	1253.67	627.34	1253.67	627.34	2	19.39
	LVYPFPGPIHNSLPQN	1792.93	896.97	1792.94	896.97	2	19.59
	LYQEPVLGPVR	1270.72	635.86	1270.72	635.86	2	16.42
	LYQEPVLGPVRGP	1424.79	712.90	1424.79	712.90	2	17.40
	LYQEPVLGPVRGPFPI	1782.00	891.50	1781.99	891.50	2	21.94
	LYQEPVLGPVRGPFPII	1895.07	948.04	1895.08	948.04	2	23.54
	LYQEPVLGPVRGPFPIIV	1994.14	997.58	1994.15	997.58	2	24.37

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	MFPPQSVLS	1005.51	503.26	1005.51	503.26	2	19.98
	MFPPQSVLSL	1118.59	559.80	1118.59	559.80	2	23.51
	MPFPKYP	879.44	440.22	879.44	440.23	2	19.47
	MPFPKYPVEP	1204.61	602.81	1204.61	602.81	2	16.50
	MPFPKYPVEPF	1351.68	676.34	1351.68	676.34	2	20.54
	MPFPKYPVEPFT	1452.72	726.86	1452.72	726.87	2	20.24
	MPFPKYPVEPFTE	1597.76	799.38	1597.76	799.38	2	19.13
	MPFPKYPVEPFTES	1668.80	834.90	1668.80	834.90	2	20.16
	MPFPKYPVEPFTESQS	1883.89	942.45	1883.89	942.45	2	19.98
	MPIQAFLLY	1095.59	548.30	1095.59	548.30	2	26.07
	PFPKYPVEP	1073.57	537.29	1073.57	537.29	2	16.40
	PFPKYPVEPF	1220.63	610.82	1220.64	610.82	2	19.42
	PKHKEMPFKYPV	1597.85	533.29	1597.86	533.29	3	12.70
	PKHKEMPFKYPVEPFTE	2201.13	734.38	2201.11	734.37	3	16.73
	PKYPVEPFTE	1206.60	603.80	1206.60	603.81	2	16.75
	PKYPVEPFTES	1293.63	647.32	1293.64	647.32	2	16.52
	PVLGPVRGPFPI	1248.74	624.88	1248.75	624.88	2	20.30
	PVLGPVRGPFPIIV	1460.90	730.95	1460.90	730.95	2	23.12
	PVVVPPFLQP	1092.64	546.83	1092.65	546.83	2	23.69
	PVVVPPFLQPE	1221.69	611.35	1221.69	611.35	2	23.43
	QDKIHPFAQ	1083.56	542.28	1083.56	542.28	2	11.59
	QEPVLGPV	838.47	419.74	838.47	419.74	2	17.73
	QEPVLGPVR	994.57	497.79	994.57	497.79	2	13.75
	QEPVLGPVRGP	1148.64	574.82	1148.64	574.82	2	15.26

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	QEPVLGPVRGPF	1295.71	648.36	1295.71	648.36	2	18.59
	QEPVLGPVRGPF	1392.76	696.88	1392.76	696.89	2	18.86
	QEPVLGPVRGPFPI	1505.85	753.43	1505.85	753.43	2	26.61
	QEPVLGPVRGPFPII	1618.93	809.97	1618.93	809.97	2	26.08
	QEPVLGPVRGPFPIIV	1718.00	859.50	1718.00	859.50	2	24.50
	QQQTEDELQ	1118.49	559.75	1118.50	559.75	2	12.30
	QQQTEDELQD	1233.52	617.26	1233.52	617.27	2	12.81
	QQQTEDELQDKIHP	1708.81	854.91	1708.81	854.91	2	13.18
	QQQTEDELQDKIHPF	1855.88	619.30	1855.88	619.30	3	16.44
	QQQTEDELQDKIHPFA	1926.92	642.98	1926.92	642.98	3	16.39
	QQTEDELQDKIHP	1580.75	790.88	1580.76	790.88	2	13.22
	QQTEDELQDKIHPF	1727.82	864.41	1727.82	864.42	2	16.54
	QTEDELQDKIHP	1452.70	726.85	1452.70	726.85	2	13.75
	QTEDELQDKIHPF	1599.76	800.39	1599.76	800.39	2	16.45
	QTPVVVPPFL	1096.64	548.82	1096.64	548.82	2	24.27
	QTPVVVPPFLQPE	1450.79	725.90	1450.79	725.90	2	23.43
	RDMPIQAFL	1090.57	545.79	1090.57	545.79	2	20.65
	RDMPIQAFL	1203.65	602.33	1203.66	602.33	2	23.59
	RDMPIQAFLLY	1366.72	683.86	1366.72	683.86	2	23.50
	SEEQQQTEDELQDKIHP	2053.92	1027.47	2053.93	1027.47	2	15.05
	SQSLTLTDVE	1092.54	546.77	1092.54	546.77	2	18.13
	TEDELQDKIHP	1324.64	662.82	1324.64	662.82	2	13.33
	TEDELQDKIHPF	1471.70	736.36	1471.71	736.36	2	16.02
	TESQSLTLTDVE	1322.63	661.82	1322.63	661.82	2	18.26

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	TPVVVPPF	855.50	428.25	855.50	428.25	2	21.27
	TPVVVPPFL	968.58	484.79	968.58	484.79	2	24.55
	TPVVVPPFLQP	1193.69	597.35	1193.69	597.35	2	23.75
	TPVVVPPFLQPE	1322.73	661.87	1322.74	661.87	2	23.43
	TQTPVVVPPF	1084.60	542.80	1084.60	542.81	2	21.60
	TQTPVVVPPFL	1197.69	599.35	1197.69	599.35	2	24.43
	TQTPVVVPPFLQP	1422.80	711.90	1422.80	711.90	2	23.63
	TQTPVVVPPFLQPE	1551.84	776.43	1551.84	776.42	2	30.97
	VLGPVRGPF	1038.61	519.81	1038.61	519.81	2	18.19
	VLGPVRGPFPI	1151.69	576.35	1151.69	576.35	2	20.03
	VLGPVRGPFPII	1264.78	632.89	1264.78	632.89	2	22.78
	VLGPVRGPFPIIV	1363.84	682.43	1363.85	682.43	2	28.17
	VLPVPQ	652.40	326.70	652.40	326.71	2	15.32
	VMFPPQS	805.39	403.20	805.39	403.20	2	16.30
	VMFPPQSVL	1017.54	509.27	1017.54	509.28	2	22.36
	VMFPPQSVLS	1104.57	552.79	1104.58	552.79	2	21.62
	VMFPPQSVLS	1120.57	560.79	1120.57	560.79	2	19.49
	VMFPPQSVLSL	1233.65	617.33	1233.65	617.33	2	23.33
	VMFPPQSVLSL	1217.66	609.33	1217.66	609.33	2	24.26
	VVVPPF	657.40	329.20	657.40	329.20	2	20.59
	VVVPPFL	770.48	385.74	770.48	385.74	2	24.53
	VVVPPFLQPE	1124.63	562.82	1124.64	562.82	2	24.08
	YQEPVLGPV	1001.53	501.27	1001.53	501.27	2	18.92
	YQEPVLGPVR	1157.63	579.32	1157.63	579.32	2	15.35

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	YQEPVLGPVRG	1214.66	607.83	1214.65	607.83	2	15.39
	YQEPVLGPVRGP	1311.70	656.36	1311.71	656.36	2	16.47
	YQEPVLGPVRGPFPP	1555.82	778.42	1555.83	778.42	2	19.51
	YQEPVLGPVRGPFPI	1668.91	834.96	1668.91	834.96	2	64.61
	YQEPVLGPVRGPFPII	1781.99	594.67	1781.99	594.67	3	22.93
	YQEPVLGPVRGPFPIIV	1881.06	627.69	1881.06	627.69	3	23.72
κ-casein	AKYIPIQY	995.56	498.28	995.56	498.28	2	16.34
	AKYIPIQYVL	1207.70	604.36	1207.71	604.36	2	20.15
	ARHPHPLSF	1198.62	599.81	1198.62	599.82	2	10.28
	ARHPHPLSFM	1329.66	665.33	1329.66	665.34	2	11.29
	ARHPHPLSFM	1345.65	673.33	1345.66	673.33	2	11.89
	DKIAKYIPIQY	1351.76	676.38	1351.76	676.38	2	17.78
	FLPYPPY	962.47	481.74	962.47	481.74	2	21.87
	FMAIPPK	819.44	410.23	819.44	410.23	2	11.73
	FMAIPPK	803.45	402.23	803.45	402.23	2	16.49
	FMAIPPKK	931.54	466.27	931.54	466.28	2	11.40
	IAKYIPIQY	1108.64	554.82	1108.64	554.82	2	17.45
	IAKYIPIQYVL	1320.79	660.90	1320.79	660.90	2	21.29
	KYIPIQ	761.45	381.23	761.46	381.23	2	13.23
	KYIPIQY	924.52	462.76	924.52	462.76	2	16.01
	KYIPIQYVL	1136.67	568.84	1136.67	568.84	2	20.10
	KYIPIQYVLS	1223.70	612.35	1223.70	612.36	2	19.15
	LSRYPSYGL	1055.55	528.28	1055.55	528.28	2	15.88
	LSRYPSYGLN	1169.59	585.30	1169.59	585.30	2	15.07

Parent protein*	Sequence	Prec MW	Prec m/z	Theor MW	Ther m/z	Charge	RT
	NQFLPYPPYYA	1275.60	638.30	1275.60	638.31	2	22.27
	NQFLPYPPYYAK	1403.70	702.35	1403.70	702.35	2	18.61
	NQFLPYPPYYAKP	1500.75	750.88	1500.75	750.88	2	19.05
	QFLPYPPYY	1090.52	545.77	1090.52	545.77	2	22.46
	QFLPYPPYYA	1161.56	581.28	1161.56	581.28	2	22.24
	QFLPYPPYYAK	1289.65	645.33	1289.66	645.33	2	18.58
	RHPHPHLSF	1127.58	376.53	1127.59	376.53	3	10.16
	RHPHPHLSFM	1258.62	629.82	1258.63	629.82	2	12.78
	RHPHPHLSFm	1274.62	637.81	1274.62	637.81	2	10.50
	RYPSYGL	855.43	428.22	855.44	428.22	2	14.89
	SRYPYGL	942.47	471.74	942.47	471.74	2	15.20
	SRYPYGLN	1056.51	528.76	1056.51	528.76	2	14.19
	SRYPYGLNY	1219.57	610.29	1219.57	610.29	2	16.26
	SRYPYGLNYY	1382.64	691.82	1382.64	691.82	2	17.51
	VLSRYPY	984.51	492.76	984.51	492.76	2	13.81
	VLSRYPYSG	1041.53	521.27	1041.54	521.27	2	13.44
	VLSRYPYGL	1154.62	577.81	1154.62	577.81	2	16.60
	VLSRYPYGLN	1268.66	634.83	1268.66	634.84	2	15.66
$\alpha$ -Ib	AFHTSGYDTQ	1126.48	563.74	1126.48	563.74	2	11.81
	DKVGINYW	994.50	497.75	994.50	497.75	2	18.44
	FHTSGYDTQ	1055.44	528.22	1055.44	528.23	2	11.48
$\beta$ -Ig	KTKIPAVF	903.57	452.29	903.57	452.29	2	14.03
	KTKIPAVFKID	1259.77	420.59	1259.77	420.60	3	15.07
	LEKFDKAL	963.55	482.28	963.55	482.28	2	13.25

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607 \*Protein access code at <https://www.uniprot.org/>. A: Alanine, D: Aspartic Acid, E: Glutamic Acid,  
608 F: Phenylalanine, G: Glycine, I: Isoleucine, K: Lysine, L: Leucine, M: Methionine, N: Asparagine,  
609 P: Proline, Q: Glutamine, R: Arginine, S: Serine, T: Threonine, V: Valine, W: Tryptophan, Y:  
610 Tyrosine, lb: lactalbumin, lg: lactoglobulin, MW: Molecular Weight, RT: Retention Time.

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ACCEPTED