



Review

Metabolic Syndrome-Preventive Peptides Derived from Milk Proteins and Their Presence in Cheeses: A Review

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Abstract: The metabolic syndrome (MetS) is defined as the occurrence of diet-related diseases such as abdominal obesity, atherogenic dyslipidemia, hyperglycemia (insulin resistance) and hypertension. Milk-derived peptides are well-known agents acting against high blood pressure, blood glucose level, and lipoprotein disproportion. The aim of this review are metabolic syndrome-preventive peptides derived from milk proteins which were identified in cheeses. Special attention was paid to the sequences acting as angiotensin converting enzyme (ACE), dipeptidyl peptidase IV (DDP4), and α -glucosidase inhibitors, as well as antioxidative, hypocholesterolemic, antiobesity, and anti-inflammatory agents. Some results of meta-analyses concerning the consumption of cheese and the risk of MetS diseases were also presented.

Keywords: biopeptides; BIOPEP-UWM database; cheese; metabolic syndrome; milk proteins

1. Introduction

The metabolic syndrome (MetS) is defined as a clustering of four major cardiovascular risk factors, namely: abdominal obesity, atherogenic dyslipidemia, hyperglycemia (insulin resistance), and hypertension (Figure 1) [1]. Moreover, the pathogenesis of MetS involves both genetic and acquired factors contributing to the inflammation [2]. MetS also represents a significant risk factor for the development of cardiovascular disease (CVD), type 2 diabetes mellitus (T2D), and all-cause mortality [1]. Moore et al. [3] analyzed data derived from the National Health and Nutrition Examination Survey (NHANES). Based on the analysis including the period from 1988 to 2012, MetS was defined as at least three conditions from these listed further that have to occur in a patient: elevated waist circumference, elevated level of triglycerides, reduced high-density lipoprotein cholesterol level, high blood pressure, and elevated fasting blood glucose level. According to the above-mentioned definition of MetS, in 2012 there were more than 66 million adults in the US with recognized MetS symptoms. They were observed in every sociodemographic group [3].

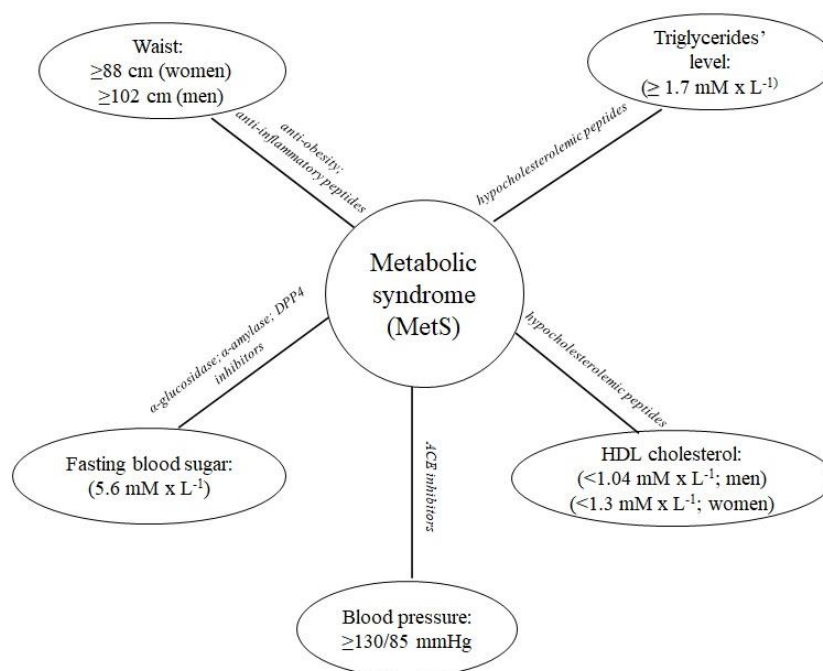


Figure 1. Metabolic syndrome criteria and related preventive agents according to Huang [4].

The following enzymes are involved in MetS: dipeptidyl peptidase IV (EC 3.4.14.5; DPP4), α -glucosidase (EC 3.2.1.20), α -amylase (EC 3.2.1.1), and angiotensin-converting enzyme (EC 3.4.15.1; ACE). The first three affect the postprandial blood glucose level, whereas ACE is the key enzyme involved in the regulation of the blood pressure. Thus, peptides being the inhibitors of the above-mentioned enzymes may be considered anti-MetS agents [5]. Additionally, peptides also reduce cholesterol level, antioxidative stress [6], act as anti-obesity [7], and anti-inflammatory agents [8].

Milk is the first functional food for every mammalian species. The contents of its nutrients, such as proteins (and their proportions), fat, sugar, vitamins, and minerals, vary between species as well as breeds. Moreover, milk composition is affected by feeding methods, age of an animal, and many other [9]. Milk and dairy products were also studied as anti-metabolic syndrome foods [10].

A cohort study made in 2018 on 136,384 individuals from 21 countries on five continents demonstrated that dairy product consumption (especially of milk and yogurt) can be associated with a lower risk, of not only CVD, but also mortality. According to this study, even one serving of milk or yogurt significantly lowered the risks of the above-mentioned events. Additionally, the higher consumption of dairy products can be associated with significantly lower risk of stroke [10]. According to another study, every 200 mL increase of daily milk consumption reduced the risk of CVD by 6%. Based on the data from 2015 it was shown that three main milk constituents were responsible for the MetS-preventive activity, namely calcium, fat, and proteins [11]. One of the MetS-preventive potentials of milk and dairy products resulting from the presence of proteins might be related to the action of biologically active peptides [5]. It also reflects vast information found in. e.g., Scopus database [www.scopus.com]. Providing the following words as the input query "ALL (metabolic AND syndrome AND milk AND peptides)" revealed 7459 hits (accessed in September 2019).

The first mentions about cheese manufacturing appeared around 5500 BC. Early adult humans were unable to digest lactose. Thus, making cheese was a good way to consume health beneficial nutrients derived from milk, without getting ill [11]. The casein, main milk protein, during cheese maturation is hydrolyzed into a variety of peptides. Their composition varies and depends on the activity of endogenous proteases and peptidases of milk, rennet, starter culture, and microbial flora [12]. Bovine casein has been shown to possess a variety of bioactivities, including MetS-preventive ones. In silico analyses and reviews conducted on bovine casein confirmed the casein-derived peptides to

exhibit these activities [13,14]. Hence, considering the increasing role of biopeptides in the prevention of MetS, the aim of this paper was to describe MetS-preventive peptides derived mainly from bovine milk proteins which were also identified in the ripened cheeses. The main attention was paid to the peptides exerting antidiabetic, antihypertensive, antioxidative, cholesterol-reducing, antiobesity, and anti-inflammatory effects.

2. Antidiabetic Agents

One of the body enzymes involved in the regulation of the glucose level is DPP4 [15]. This enzyme is involved in the incretin effect after meal consumption. Incretins, like glucagon-like peptide 1 (GLP-1) and glucose-dependent insulinotropic polypeptide (GIP), are hormones that decrease blood glucose levels by stimulating insulin secretion and inhibiting glucagon release. GIP serves several additional functions including promotion of growth and survival of the pancreatic beta-cell and stimulation of adipogenesis [15]. This effect is hindered by DPP4, therefore inhibition of DPP4 prolongs the incretin effect and lowers the postprandial blood glucose level (Figure 2) [16].

Uenishi et al. [17] identified DPP4 inhibitory peptides in water-soluble Gouda cheese extracts. Peptides derived from β -casein (LPQNIPPL) and α_{S1} -, α_{S2} -, β -caseins (LPQ) evoked the highest DPP4-inhibiting effect in vitro. Their IC_{50} values were 46.0 and 56.7 μ M, respectively. The bioactivity of the LPQNIPPL sequence was also tested on rats (Female Sprague Dawley, 8-week-old). The administration of this peptide at $300 \text{ mg} \times \text{kg}^{-1}$ body weight reduced significantly the postprandial glucose level in those rats [16]. Other peptides with DPP4 inhibitory bioactivity that were identified in Gouda cheese water-soluble-extracts (WSE) were characterized with the following sequences: VPITPTL (110 μ M), VPITPT (130 μ M), FPGPIPN (260 μ M), YPFPGPIPN (670 μ M), PGPIHNS (1000 μ M), IPPLTQTPV (1300 μ M), PQNIPPL (1500 μ M), and VPPFIQPE (2500 μ M). Their activity expressed in IC_{50} values was provided in brackets [17].

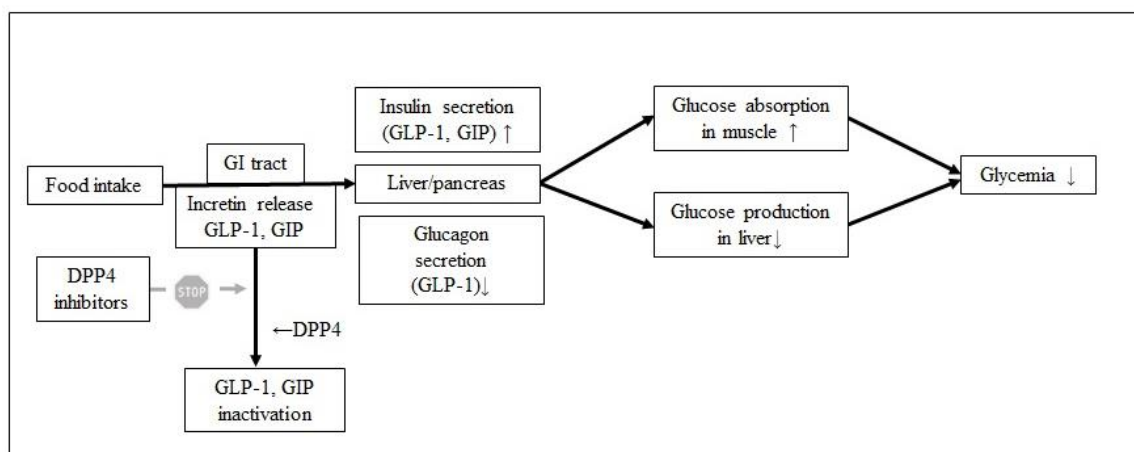


Figure 2. The involvement of the dipeptidyl peptidase IV inhibitors in prolonging the incretin effect according to Mkele [16].

Antidiabetic properties of peptides may also result from their ability to inhibit α -glucosidase (EC 3.2.1.20). α -Glucosidase is an enzyme located in the brush border of enterocytes of the jejunum in the small intestine and is a key enzyme in carbohydrate breakdown. It cleaves carbohydrate to easy absorbable glucose. Thus, one of the T2D preventive strategies is to decrease the postprandial hyperglycemia by delaying glucose absorption through the inhibition of α -glucosidase (Figure 3). α -Glucosidase inhibitors also play a crucial role in the secretion of GLP-1 in healthy and diabetic patients, due to the prolonged incretin effect [18].

α -Amylase is another enzyme involved in the regulation of the blood sugar level [5]. This enzyme is responsible for the initial hydrolysis of the α -D-(1-4) glycosidic bonds of polysaccharides like glycogen

and starch to oligosaccharides. Then, oligosaccharides are further hydrolyzed into monosaccharides by α -glucosidase (see above), which are next transferred into bloodstream [5].

Some milk proteins were reported as the sources of α -glucosidase inhibitors. Whey proteins: α -lactalbumin, β -lactoglobulin, serum albumin, and lactoferrin hydrolysates obtained by peptic digestion were confirmed to exert such an inhibiting effect. β -Lactoglobulin and whey protein isolate hydrolysates were found to inhibit α -glucosidase (IC_{50} value of 3.5 and 4.5 $mg \times mL^{-1}$, respectively) [18]. Dipeptide, like arginine-proline (RP), derived from whey proteins and being the component of commercial Pep2Dia[®] product, affected the inhibition of α -glucosidase with an IC_{50} value of 0.0025 $mg \times mL^{-1}$ [19]. To the best of our knowledge, there is no research papers about α -amylase inhibitors identified milk proteins and dairy products. Only one article concerned cheese as a source of α -glucosidase inhibitors. According to Grom et al. [20], Prato cheese inhibited α -glucosidase activity in vitro by 52.6%. Although, it was concluded that the consumption of probiotic Prato cheese can contribute to the reduction of postprandial glycemia in healthy individuals, but no peptide was identified [20].

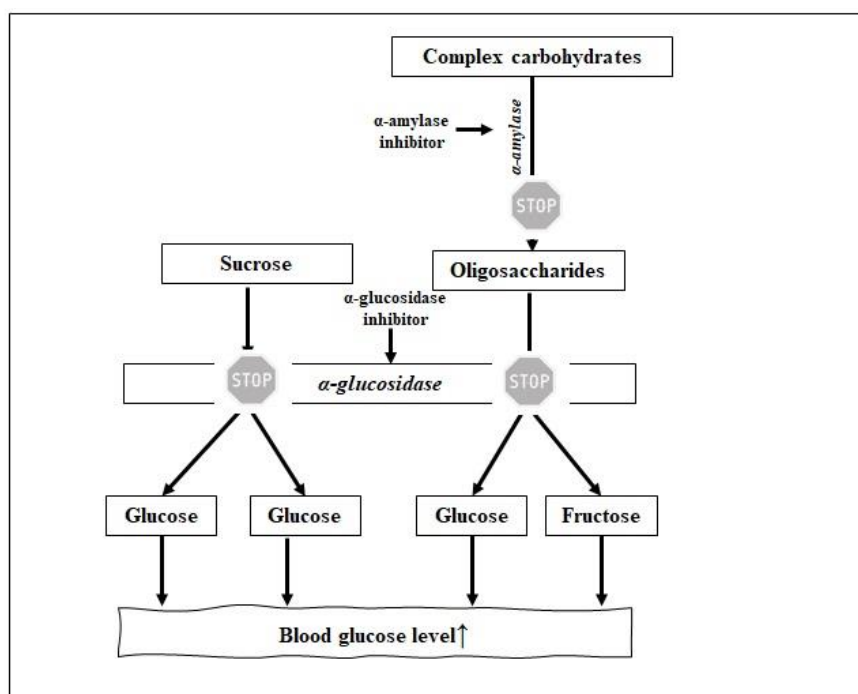


Figure 3. The delay of carbohydrate breakdown by α -glucosidase and α -amylase inhibitors according to Iwaniak et al. [5] and Patil et al. [18].

It was evidenced on mice that peptides from cheese effectively decreased glucose level. Some epidemiological surveys showed a slightly higher risk of T2D development with increasing cheese consumption, but most of the meta-analyses concluded that increasing cheese consumption was correlated with a significant decreasing in T2D development risk [21]. Another epidemiological and cohort studies showed that dairy products can be beneficial in type 2 diabetes prevention. The latest research carried out on 34,224 men in the Health Professionals Follow-Up Study (1986–2012), 76,531 women in the Nurses' Health Study (1986–2012), and 81,597 women in the Nurses' Health Study II (1991–2013) suggested that the decrease of total dairy intake by > 1.0 serving per day over a period of 4 years was associated with an 11% increase of T2D development risk [22]. However, there were some opposite remarks to these above. For example, it was stated that increasing cheese consumption by > 0.5 serving per day, caused a 9% increase of T2D development risk, when compared to the maintained, stable cheese intake [22]. In turn, these findings were not consistent with other studies. For example, the meta-analysis made by Gao et al. [23], concerning seven studies with a total of

178,429 subjects and 14,810 cases, led to opposite conclusions. They found an inverse linear correlation between the consumption of cheese and the risk of T2D development. The consumption of 30 g per day cheese decreased the relative risk (RR) of T2D development ($RR\ 0.80 \pm 0.12$). The obtained results suggested that a modest increase in cheese consumption might prevent from T2D but further research is needed [23]. Another eight cohort studies concerning the meta-analysis of dairy food consumption and its effect on the T2D risk revealed that the consumption of 50 g cheese per day decreased the risk of T2D development ($RR\ 0.92 \pm 0.07$) and a non-linear negative correlation was observed between cheese consumption and T2D risk [24]. The cohort study conducted on 2809 participants for 12 years showed a negative dose-response relation between cheese consumption and risk of T2D development. Out of the 1867 participants with no T2D symptoms, 902 were diagnosed as prediabetic. Consumption of ≥ 4 servings of cheese per week lowered the risk of T2D incident by 70%. Less than one cheese serving per week was associated with a 63% reduction of T2D incidents. None of the abovementioned meta-analyses, however, considered what type of cheese caused the risk-reducing effect [25].

3. Peptides with Antihypertensive Activity

The regulation of blood pressure that takes place in the organism involves several modes of ACE action. They include renin-angiotensin, kinin-nitric-oxide, and neutral endopeptidase systems. The first one (RAS; renin-angiotensin system), also called renin-angiotensin-aldosterone system (RAAS), is considered as the major in blood pressure regulation. Briefly, renin (EC 3.4.22.15) activates angiotensinogen and releases angiotensin I. Then, ACE (Figure 4) transforms angiotensin I into angiotensin II (known as potent vasoconstrictor) and induces the release of aldosterone. Moreover, the increase of the blood pressure caused by angiotensin II causes some dysfunctions like damages of body organs leading to e.g., heart and kidney failure. Thus, the action of ACE receptor blockers or ACE-inhibiting peptides contributes to the blood pressure reduction [26].

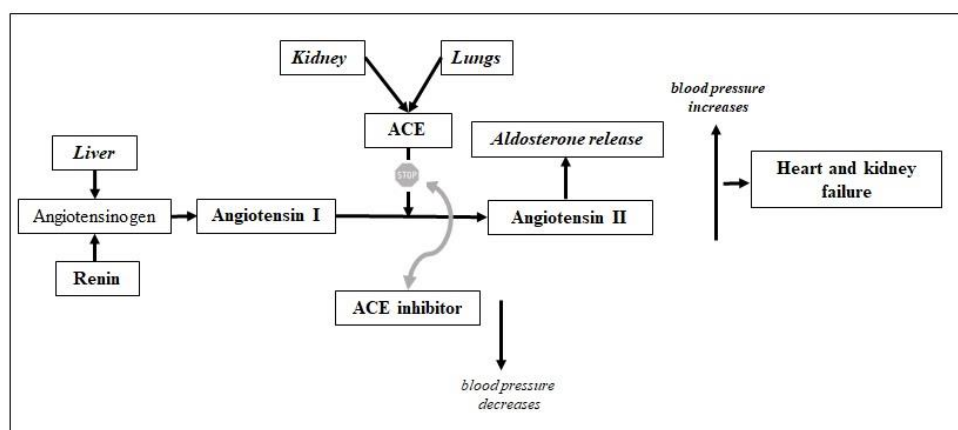


Figure 4. Brief description of the mechanism of action of ACE and its inhibitors according to Iwaniak et al. [26].

Milk proteins have been reported as the reservoir of ACE inhibitors for many years [27]. The most known sequences are VPP and IPP identified in β - and κ -casein, respectively. Due to their antihypertensive potential tested in animals and humans, they became the components of commercial milk drink nutraceuticals such as Calpis, Evolus, and Valio [28]. VPP and IPP sequences were also identified in 36 cheeses of Swiss origin (Table 1). Depending on the type of Swiss cheese, the content of these two sequences varied. For VPP, it ranged from 0 to 224 mg per kg of cheese, whereas for IPP from 0 to 95.4 mg. The highest content of both peptides, VPP and IPP, was identified in the hard cheese “Hobelkäse from the Bernese Oberland” (224.1 and $95.4\text{ mg} \times \text{kg}^{-1}$, respectively). That cheese sample had the highest ACE inhibitory activity of all analyzed cheeses, with an IC_{50} value of $2.6\text{ mg} \times \text{mL}^{-1}$ [29]. Different concentrations of these two peptides in cheeses were probably related

to milk pretreatment, cultures, scalding conditions, and ripening time. Thus, it is recommended to develop a reproducible cheese-making process to produce higher concentrations of these peptides that could be used for clinical trials [30].

The potential of some cheeses containing the high concentrations of VPP and IPP to reduce blood pressure was tested in spontaneously hypertensive rats (SHR). Blood pressure was monitored after 6 h since oral administration. It was observed that the decrease of systolic blood pressure (SBP) differed depending on the cheese variety. The level of the SBP reduction (mmHg) in SHR was as follows: -24.7 and -17.2 (Gouda, aged 8 and 24 months, respectively), -17.2 (Gouda, aged 24 months), -20.7 (Edam), -20.3 (Blue), -20.0 (Havarti), and -13.0 (Emmental) [31].

Table 1. Quantification of VPP and IPP in cheeses of Swiss origin with their corresponding ACE inhibitory activity expressed with IC_{50} values. Adapted with permission from [29], Copyright Elsevier, 2007.

Cheese	VPP	IPP	IC ₅₀
	(mg × kg ⁻¹)	(mg × kg ⁻¹)	(mg cheese × mL ⁻¹)
Hard cheeses			
Emmental:			
cave-aged, aged 12 months	100.1	23.5	7.1 ± 0.2
reserve, aged 8 months	68.6	13.2	9.9 ± 0.2
organic, aged 4 months	153.7	35.8	7.5 ± 1.8
classic, aged 4 months	42.6	4.6	10.5 ± 0.6
Gouda old	97.7	90.1	2.0 ± 0.3
Gruyere AOC:			
vieux, aged 12 months	13.7	7.9	29.4 ± 16.8
salé, aged 10 months	68.7	20.4	14.2 ± 0.9
mi-salé, aged 8 months	22.7	14.9	18.7 ± 6.1
doux, aged 5 months	53.0	9.3	14.9 ± 0.7
Hobelkäse from the Bernese Oberland	*224.1	95.4	2.6 ± 0.2
Parmino	2.0	0.8	28.6 ± 4.2
Sbrinz	28.0	9.4	16.4 ± 5.5
Semi-hard cheeses			
Appenzeller:			
aged 7–10 months (1/4 fat)	94.4	64.9	4.2 ± 1.6
extra, aged 6–7 months	29.3	14.5	16.8 ± 1.1
surchoix, aged 4 months	26.7	17.8	8.0 ± 1.0
aged 3 months (1/4 fat)	56.0	10.5	7.1 ± 1.9
classic, aged 3 months	22.0	8.5	17.6 ± 7.6
Edam	1.0	0.1	13.3 ± 0.4
Küsnachter	25.7	18.0	19.6 ± 2.4
Mountain cheese Graubünden:			
$\frac{1}{4}$ fat	32.0	22.7	4.8 ± 0.0
$\frac{1}{2}$ fat	10.4	7.1	8.9 ± 4.1
full fat	25.1	4.3	9.2 ± 1.4

Table 1. Cont.

Cheese	VPP	IPP	IC ₅₀
	(mg × kg ⁻¹)	(mg × kg ⁻¹)	(mg cheese × mL ⁻¹)
Raclette:			
from raw milk	12.6	2.6	8.8 ± 0.0
from pasteurized milk	1.0	0.1	9.0 ± 0.3
St. Paulin	5.2	1.0	23.5 ± 6.3
Tête de Moine	98.0	15.5	7.1 ± 1.9
Tilsit:			
from raw milk	120.2	30.0	4.4 ± 0.2
from pasteurized milk	1.2	0.1	18.0 ± 6.3
Vacherin fribourgeois	62.1	46.7	5.0 ± 0.3
Wangener Geissmutschli (goat cheese)	33.2	0.0	7.9 ± 0.5
Winzerkäse	97.7	27.1	9.9 ± 1.8
Soft cheeses			
Brie	0.4	0.3	25.9 ± 3.0
Camembert	0.2	0.2	Very weak
Mozarella:			
from buffalo milk	0.1	0.0	-
from cow milk	0.0	0.0	-
Tomme vaudoise	0.0	0.0	14.9 ± 3.7
Vacherin Mont d'Or	0.2	0.1	21.4 ± 9.6

***Bold** - the highest concentration of peptide in the analyzed category (hard, semi-hard, and soft cheese).

According to the scientific reports, sequential motif located between 60 and 70 residue of β -casein is considered as strategic zone due to its ACE-inhibiting, immunostimulating, and opioid potential [32]. The example of peptide derived from β -casein showing dual bioactivity is β -casomorphin-7 (YPPFGPI; fragment 60–66) acting as opioid and ACE inhibitory agent [26]. De Noni and Cattaneo [33] identified this peptide in different dairy products including commercial cheeses like e.g., Brie, Cheddar, Gouda, Fontina, and Gorgonzola (see Table 2).

Table 2. The Concentration of YPPFGPI in commercial cheeses before and after in vitro simulated gastro-intestinal digestion. Adapted with permission from [33], Copyright Elsevier, 2010.

Cheese	Concentration of YPPFGPI (mg × kg ⁻¹)	
	Before Simulated Digestion	After Simulated Digestion
Brie	0.15	2.07 ± 0.09
Caprino	0	1.51 ± 0.09
Cheddar	0.11	15.22 ± 0.13
Fontina	0.04	10.43 ± 0.12
Gorgonzola	0.01	5.41 ± 0.08
Gouda	0.10	21.77 ± 0.12
Grana Padano (aged 25 months)	0	11.06 ± 0.08
Grana Padano (aged 17 months)	0	12.55 ± 0.09
Grana Padano (aged 10 months)	0	8.79 ± 0.04
Taleggio	0	7.15 ± 0.07

Some ACE inhibitors which matched the sequences of caseins were identified in Spanish cheeses. Those peptides which sequences fulfilled the structural requirements for being a potent ACE inhibitors were synthesized [34]. Their structural characteristics included the presence amino acids like lysine, arginine or residues possessing hydrophobic side chains as well as C-terminal proline [35]. Finally, it was reported that eight peptides (out of 11) inhibited ACE. Among them DKIHP peptide was found as the most potent ($IC_{50} = 113.1 \mu M$) but the addition C-terminal phenylalanine (DKIHPPF) caused the dramatic reduction of its ACE-inhibiting potential ($IC_{50} = 2419.4 \mu M$) [36]. The ACE-inhibiting activity of synthesized peptide DKIHP (source Manchego cheese; β -casein) was also measured using different protecting groups, namely *tert*-butyloxycarbonyl (Boc) and fluorenyl-methyloxycarbonyl (Fmoc). The IC_{50} values for DKIHP were 113.18 and 577.92 μM , respectively. It was concluded that the differences above were attributed to the presence of less active conformers in the structure of the peptide [37]. The other sequences of ACE inhibitors identified in some Spanish cheeses are given in Table 3.

Table 3. ACE inhibitors identified in Spanish cheeses. Adapted by permission from Springer, European Food Research and Technology [36], 2006.

Cheese	Peptide	IC_{50} (μM)	Precursor	Reference
Manchego, Roncal, Idiazábal, Cabrales, goat	DKIHP	113.1	β -casein	[36]
	DKIHPPF	2419.4		
Goat	VRGP	120.9	various fragments	[36]
	PFP	144.4		
Manchego	FP	1215.7	α_{S1} -casein	[30]
	PP	2284.7		
	KKYNVPQL	77.2		
	VRYL	24.1		
Manchego, Roncal, Idiazabal, Goat	QP	598.1	various fragments	[36]
Roncal	PKHP	709.1	α_{S1} - casein	

Sequences of caseins were also the sources of other ACE-inhibiting peptides that were identified in cheeses: RPKHPIKHQ (source: α_{S1} -casein; Cheddar and Gouda; $IC_{50} = n.d.$ and $13.4 \mu M$, respectively), RPKHPIKHQGLPQ (source: α_{S1} -casein; Gouda; $IC_{50} = n.d.$), YPFPGPIP (source: β -casein; Gouda; $IC_{50} = 14.8 \mu M$), MPFPKYVPVQPF (source: β -casein; Gouda; $IC_{50} = n.d.$), RPKHPIK, RPKHPI, FVAPFPEVFGK (source: α_{S1} -casein, Cheddar; $IC_{50} = n.d.$), YQEPVLGPVVRGPFPIIV (source: β -casein; Cheddar; $IC_{50} = n.d.$), RPKHPIKHQGLPQEV, RPKHPIKHQGLPQEVLNEN, LLR, EVLNENLLRF (source: α_{S1} -casein, Fresco; $IC_{50} = n.d.$), FVAPFPEVFGK, YQEPVLGPVVRGPF, YQEPVLGPVVRGPFPI, YQEPVLGPVVRGPFPIIV (source: β -casein; Fresco; $IC_{50} = n.d.$) [30]. Tonouchi et al. [38] isolated two peptides (LQP and MAP) from Danish skimmed milk cheese. LQP was originally known as ACE inhibitor identified in α -zein hydrolyzed with thermolysin. MAP sequence was identified in β -casein and exhibited ACE inhibition ($IC_{50} = 0.8 \mu M$) and antihypertensive effect in SHR. Significant hypotensive effect of MAP tripeptide was observed at 8 h since oral administration (-17.0 mm Hg; dose $3 \text{ mg} \times \text{kg}^{-1}$ body weight) [38].

The potential of cheeses as diet components that may affect the reduction of blood pressure is also in the focus of interest of the scientists aiming at meta-analyses [39]. Nilsen et al. [39] carried out a cross-sectional study to analyze the impact of the consumption of Gamalost cheese (a traditional Norwegian cheese) on the reduction of the risk of hypertension development. Based on data obtained from the questionnaires of 168 participants, the average intake of Gamalost servings per week was 2 and the average value of systolic and diastolic blood pressures was 128 and 78 mmHg, respectively.

The increase of Gamalost servings reduced the systolic blood pressure by 0.72 mm Hg. Moreover, it was concluded that blood pressure reduction related to Gamalost consumption was associated with the presence of ACE inhibitory peptides [39]. Crippa et al. [40] investigated the antihypertensive effect resulting from the consumption of Grana Padano cheese and observed that supplementation of Grana Padano cheese at a dose 30 g per day significantly lowered the systolic and diastolic blood pressure in mild-hypertensive subjects (−3.5 and −2.4 mm Hg, respectively) [40].

One of the meta-analyses concerned studies evaluating the association of CVD, coronary heart disease (CHD), and stroke with high vs. low cheese consumption. This analysis revealed that high cheese consumption lowered the relative risk for: total CVD (RR 0.90; 7; 8076), CHD (RR 0.86; 8; 7631), and stroke (RR 0.90; 7; 10,449). Respective numbers in the brackets represent: relative risk, number of studies, and number of documented events provided. Finally, the results indicated a nonlinear relationship between cheese consumption and decreasing risks of CVD, CHD, and stroke. Consumption of approximately 40 g of cheese per day ensured the best results in reducing the risks of dysfunctions related to hypertension [41].

To conclude, epidemiological studies suggested that the antihypertensive potential of cheeses can be linked to the presence of biopeptides. According to above-mentioned analyses, the optimal daily cheese dose to reduce the risk of hypertensive dysfunctions was 30–40 g. These results suggest that cheese-as part of a diet-can prevent from the hypertension-related diseases.

4. Antioxidative Peptides

Free radicals are involved in many signaling processes, i.e., as defending the body from infections. Thus, the accumulation of free radicals in the organism can lead to unwanted cell damage. Oxidized cells can then initiate several body dysfunctions like e.g., rapid ageing, atherosclerosis, arthritis, diabetes, and cancer (Figure 5). It is the so-called oxidative stress defined as the process of formation of highly reactive molecules (that is, reactive oxygen species-ROS). In order to act against the destructive activity of ROS, it is important to look for diet-derived compounds acting as antioxidants [42].

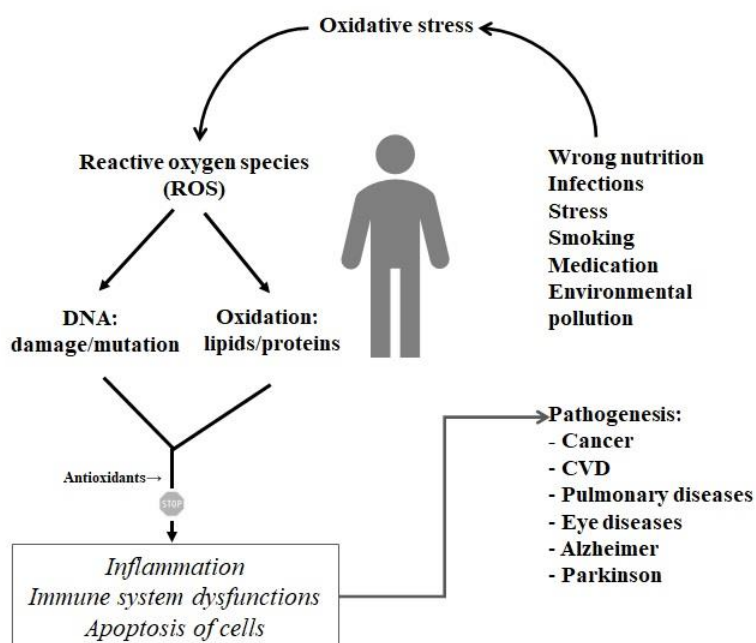


Figure 5. The preventive activity of antioxidants on the formation of diseases according to Sarmadi and Ismail [42].

Gupta et al. [43] produced Cheddar cheese using different *Lactobacillus casei* cultures and screened for antioxidant activity. Two antioxidative peptides matching the fragments of bovine β -casein

(VKEAMAPK) and α_{S1} -casein (HIQKEDVPSER) were identified in Cheddar. Both peptides exhibited 2,2-diphenyl-1-picryl-hydrazyl (DPPH) radical scavenging activity comparable to the commercial antioxidants such as butylated hydroxyanisole (BHA), *tert*-butyl hydroquinone (t-BHQ) or ferulic acid [44]. The results indicated that the antioxidant activity of the water-soluble cheese extracts (WSE) was dependent on the ripening period of cheese. The highest bioactivity was observed after 4 months of ripening. After that, the activity started to decrease, up until the 7th month of ripening and remained similar after 9 months [44]. Another research group evaluated the inhibition of DPPH radicals by the WSE of Australian Cheddar cheeses. They found a moderate antioxidant activity of the fractions containing peptides with a molecular weight > 10 kDa, although peptide identification was not carried out [45]. A study conducted to analyze the antioxidative potential of Mexican Cotija cheese revealed that the maximal antioxidant activity of this cheese was achieved after 24 weeks of ripening. Although no specific peptides were identified, it was concluded that the antioxidative potential of cheese was related to a significantly high content of bioactive sequences [46].

Timón et al. [47] produced hard cow cheese and analyzed the antioxidative potential of its WSE. After the fractionation of WSE, it was discovered that some of the extracts exhibited DPPH-radical scavenging and metal chelating bioactivity. Moreover, two peptides, namely EIVPN and DKIHFP which were known from their ACE-inhibiting function, were identified in WSE of the hard cheese. Another sequence i.e., VAPFPQ identified in cheese-derived extract was discovered as a new metal chelating peptide. To conclude, it was postulated that antioxidative potential of cheese water-soluble extracts may be the result of the presence of the above-mentioned peptides. However, the bioactivity of these sequences should be confirmed on synthetic peptides [47].

Some peptides with the antioxidative potential were identified in WSE of the Burgos-type cheese. It was shown that 22 fractions derived from Burgos-type cheese extracts exhibited radical scavenging activity (oxygen radical absorbance capacity with fluorescein (ORAC-FL) assay). Liquid chromatography and tandem mass spectrometry (LC-MS/MS) identification revealed the presence of the aromatic amino acids like tyrosine, phenylalanine or tryptophan in some fractions of cheeses extracts. These amino acids itself possess the antioxidative potential [48]. It was also reported that some of the Burgos-type cheese-originating peptide fractions contained the sequences, namely: SDIPNPIGSENSEKTTMPLW (source: α_{S1} -casein), YQQPVLGPVVRGPFPIIV (source: β -casein), and LLYQQPVLGPVVRGPFPIIV (source: β -casein) [49]. According to Timón et al. [49], peptides matching the sequences of β -casein have already been known as antioxidative agents. The sequence SDIPNPIGSENSEKTTMPLW is probably found as the new peptidic antioxidant, however further tests to measure this activity need to be performed [49].

5. Hypocholesterolemic Peptides

Plasma lipoproteins are composed of the following elements: cholesterol, triglycerides, phospholipids, and apoproteins. Their varying composition affects the density, size, and electrophoretic mobility of each lipoprotein particle which enables for the classification of lipoprotein disorders [50]. According to Cox et al. [50], cholesterol is an essential precursor of some substances as adrenal and gonadal steroid hormones and bile acids. Moreover, it is an element of the animal cell membranes. Triglycerides are the major component of dietary fat as well as animal fat depots. Abnormal concentrations of cholesterol and triglycerides in blood indicate some dysfunctions in synthesis, degradation, and transport of lipoproteins and their increase levels lead to the different variants of hypercholesterolemia and hypertriglyceridemia [50]. In turn, phospholipids are the constituents of the natural membranes. They are polar lipids which are e.g., the components of the milk fat globule membrane involved in the circulation of the lipid droplets secreted by the mammary gland cells [51].

It was reported that some peptides/protein-derived hydrolysates exhibited activity against metabolic lipids' dysfunctions and their effect was tested in cellular systems and animal models [52]. To the best of our knowledge, no peptides acting as hypolipidemic were identified in cheeses. When thinking about milk, most of the studies concerning their "anticholesterolemic effect" conducted so far,

have concerned the whey proteins. The ingestion of whey protein was observed to be correlated with a significant reduction in total cholesterol levels in rats fed cholesterol-free and cholesterol-enriched diets [53]. The cholesterol reducing activity of β -lactoglobulin was due to the presence of two major biopeptides, namely lactostatin (IIAEK) and β -lactotensin (HIRL) [54]. According to the literature, consumption of 300 mg IIAEK \times kg⁻¹ body weight per day reduced LDL level as well as increased HDL cholesterol level, and fecal steroid excretion. The lactostatin activity was linked to the degradation of cholesterol into bile acids in hepatic cells [55,56]. The HIRL sequence was shown to be involved in the reduction of total (22.7%), and LDL plus VLDL (32.0%) cholesterol levels in mice after 2 days of administration at the dose of 30 mg \times kg⁻¹ body weight [57]. The activity of both peptides was tested on mice fed a cholesterol-enriched diet. Despite the optimistic results, it should be noted that the exact mechanisms of the hypocholesterolemic activity of peptides are still unclear. It is speculated that it might be linked to the amino acid composition, but further research is needed in this respect [54].

6. Antiobesity and Anti-Inflammatory Peptides

The studies concerning the effectiveness of different types of diets and proportions of macronutrients on the weight loss is in the focus of the interest of the many scientists [58]. Some studies have shown that the high-protein content diet is the most effective when thinking about the reduction of body weight. It results from the fact that consumption of the protein contributes to the increased satiety and the reduction of the energy intake in addition to a greater dietary thermogenesis [59].

Although proteins accelerate caloric metabolism, some of them are more efficient on weight loss than others. It was observed that soybean-originating proteins more efficiently enhance weight loss than bovine milk proteins [60]. One of the explanations of this phenomenon results from the fact that peptides produced from proteins during gastrointestinal digestion initiate satiety signals from the gut and hence suppress food intake. Thus, scientists try to explain reliably the mechanisms of a protein-rich diet on the body's health [5]. Some of them include e.g., the impact of dietary protein on cholecystokinin (CKK) release, the activation of GLP-1 receptor to suppress appetite, and/or modulating the adipose genes affecting metabolic and vascular functions [60].

It was reported that opioid peptides may also contribute to the weight loss due to the reduction of food intake [61]. For example, casomorphins interact with gastric opioid receptors and slow down the motility of the gastrointestinal tract, which inhibits food intake [60]. The example of the opioid peptide identified both in bovine casein and several types of cheeses is β -casomorphin-7 (YPPFGPI) which was described above (see Section 2). It was indicated that dose of 179 mg \times kg⁻¹ body weight of modified β -casomorphins effectively impaired gastric emptying in pup rats. It was probably related to direct interaction of β -casomorphins with opioid receptors in the gut [62].

Some studies confirmed the causal direction of the relationship between adiposity and inflammation (the greater adiposity, the higher C-reactive protein level) [63]. Citing the words by Zielińska et al. [64], inflammation is defined as a “complex biological process, occurring through a variety of mechanisms and leading to changes in local blood flow and the release of several molecular mediators”. Uncontrolled inflammation may lead to chronic dysfunctions such as rheumatoid arthritis, atherosclerosis, and it is linked to cancer [65]. It was reported that VPP and IPP milk-derived ACE-inhibiting tripeptides exhibited anti-inflammatory activity. First reduced leukocyte-endothelial cell interaction in vitro and this effect was related to that of inhibition of MAPK (mitogen-activated protein kinase) signaling pathway. The anti-inflammatory effect of both peptides was also tested in a model of intestinal enterocolitis. It was found that VPP and IPP were protective against atherosclerosis in mice with defect of endogenous apoprotein E (ApoE). These peptides suppressed the mRNA for inflammatory cytokines, oxidized low density lipoprotein receptor, and transcription regulators [65]. These two peptides were also identified in the different types of cheeses (see Section 2).

To the best of our knowledge, there is no more specific anti-inflammatory peptides that could be identified in cheeses. However, it was found that water-soluble extracts of Cheddar

cheese possessed such an activity which was determined based on nitric oxide (NO) production in lipopolysaccharide-stimulated macrophage (RAW-264.7) cells [66].

7. Multi-Activity of Peptides in MetS Prevention: A Screening Using BIOPEP-UWM Database

Loads of publication data concern the analysis of the proteins as the sources of biopeptides exhibiting only one biological effect while the information about dual or multifunctional properties of peptides is rather poor [5]. Such statement may also refer to the MetS-preventive peptides. To the best of our knowledge, there is no a “universal” peptide sequence that would show all bioactivities described in this paper. Thus, databases of biological information like BIOPEP-UWM, available at: <http://www.uwm.edu.pl/biochemia>, may be a suitable option to acquire the knowledge about peptides exhibiting more than one biological function [67]. BIOPEP-UWM is a curated database providing comprehensive information on 3899 bioactive peptide sequences (accessed: March 2020).

We applied BIOPEP-UWM database to retrieve the profiles of potential biological activities of the four major casein fractions (see Table A1). Profile of the potential biological activity of the protein is defined as the type and the location of bioactive fragment in a protein chain [67]. Briefly, such analysis enables to answer the question: what bioactive fragments match the protein sequence? Then, each peptide sequence was pasted again to BIOPEP-UWM database of bioactive peptide sequences using the search option “sequence” and ticking “exact” [67]. Such searching was helpful to show what additional activities are exhibited by this peptide (the pasted one). This approach led us find that 66 peptides had dual i.e., ACE and DPP4 inhibitory function, and 17 were acting as ACE inhibitory and antioxidant agents. Some peptides exerted a triple bioactivity, i.e., eight dipeptides inhibited ACE and DPP4 as well as exerted an antioxidant activity. The example of the sequence with triple inhibitory bioactivity is YP peptide present in α_{S1} -, β -, and κ -casein sequence. It has proven ACE, α -glucosidase and DPP4 inhibitory function. The brief summary of peptides encrypted in the sequences of caseins and exerting e.g., more than one biological effect is shown in Table 4. For checking what particular sequences were found as dual or multiactive, see Table A1.

Table 4. The summary of casein-derived peptides showing more than one biological activity (based on BIOPEP-UWM database screening; <http://www.uwm.edu.pl/biochemia>; accessed on March 2020).

Dual and Triple Activity of Casein-Derived Peptides	Number of Peptides Encrypted in the All Casein Sequences
ACE and DPP4 inhibitor	66
ACE inhibitor and antioxidative	17
DPP4 and α -glucosidase inhibitor	1
DPP4 inhibitor and opioid	1
ACE inhibitor and opioid	5
DPP4 inhibitor and antioxidative	5
ACE, DPP4 inhibitor, and antioxidative	8
ACE, DPP4, and α -glucosidase inhibitor	1
Total	104
Peptides Showing one Bioactivity	
ACE inhibitor	171
α -glucosidase inhibitor	0
antioxidative	52
DPP4 inhibitor	127
opioid agonist	3
opioid antagonist	2
opioid	4
Total	359
Total Number of Bioactive Sequences	463

Although, our bioinformatic approach is useful for searching multifunctions of peptides, it has some limitations. The important one is based on the idea of the positive selection assuming that sequence of interest matches the peptide sequences present in databases. Thus, it may not include peptides that are “new” and they are, at the time of screening, not displayed in the database. Thus, some authors postulate for the regular update of the databases with newly-reported bioactive peptides [68]. Nevertheless, such databases, if curated and reliable, may contribute to acquiring of some information about the peptides which multi-activity can be described as “MetS-preventive”.

8. Conclusions

Some milk-derived peptides were identified in different types of cheeses. These biopeptides exhibited bioactivities playing an important role in the prevention from the dysfunctions related to metabolic syndrome. It needs to be noted that majority of studies concerned the identification of peptides in cheeses and did not included studies on animal and humans. Several epidemiological and cohort studies confirmed the positive effect of cheese consumption on the reduction of the risk of MetS diseases. A majority of studies concerning milk-derived peptides and their identification in cheeses focus on the determination of only one biological effect. Despite its limitations, bioinformatic-assisted approach can be helpful in the screening of multifunctional effect of peptides being the promising agents in the prevention of MetS.

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Abbreviations

ACE	Angiotensin-converting enzyme (EC 3.4.15.1)
ApoE	endogenous apoprotein E
BHA	butylated hydroxyanisole
BIOPEP-UWM	database of bioactive peptide sequences [68]
Boc	tert-butyloxycarbonyl
CHD	coronary heart disease
CVD	cardiovascular disease
DPP4	dipeptidyl peptidase IV (EC 3.4.14.5)
DPPH	2,2-diphenyl-1-picryl-hydrazyl
Fmoc	fluorenylmethyloxycarbonyl
GIP	glucose-dependent insulinotropic polypeptide
GLP-1	glucagon-like peptide 1
IP3	inositol 1,4,5-triphosphate
LC-MS/MS	Liquid Chromatography and Tandem Mass Spectrometry
LDL cholesterol	low density lipoprotein cholesterol
MAPK	mitogen-activated protein kinase
MetS	metabolic syndrome
NHANES	Health and Nutrition Examination Survey
NO	nitric oxide
NTR2	neurotensin receptor 2
ORAC-FL	Oxygen Radical Absorbance Capacity with Fluorescein
RAAS	renin-angiotensin-aldosterone system
RAS	renin-angiotensin system
ROS	reactive oxygen species
RR	relative risk
SBP	systolic blood pressure
SHR	spontaneously hypertensive rats
T2D	type 2 diabetes mellitus
t-BHQ	tert-butyl hydroquinone
VLDL cholesterol	very low density lipoprotein cholesterol
WSE	water-soluble cheese extracts

Appendix A

Table A1. Biopeptides (MetS-preventive sequences) Present in the Sequences of Bovine Caseins - Data Acquired from the BIOPEP-UWM Database (accessed on March 2020).

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
7590/8637	1*	AA	κ -casein	(86–87)
8758	12	AE	α_{S1} -casein	(77–78), (131–132)
			α_{S2} -casein	(77–78)
7583/8759	1	AF	β -casein	(189–190)
8193	9	AI	α_{S2} -casein	(42–43)
			κ -casein	(128–129)
8559	12	AL	α_{S1} -casein	(13–14)
			α_{S2} -casein	(13–14), (96–97), (190–191)
			κ -casein	(13–14), (70–71)
3177/7584	1	AP	α_{S1} -casein	(41–42), (191–192)
			β -casein	(103–104)
7742	9	AR	α_{S1} -casein	(15–16)
			κ -casein	(117–118)
8762	12	AS	κ -casein	(147–148)
8763	12	AT	α_{S2} -casein	(80–81)
			κ -casein	(165–166)
8764/8951	1	AV	α_{S1} -casein	(11–12)
			α_{S2} -casein	(11–12), (131–132)
			β -casein	(177–178)
			κ -casein	(87–88), (159–160), (189–190)
7543/8460/8695	7	AW	α_{S1} -casein	(178–179)
3563/7866/8765	7	AY	α_{S1} -casein	(158–159), (173–174)
7606	9	DA	α_{S1} -casein	(172–173), (190–191)
9075	9	DM	α_{S2} -casein	(155–156)
			β -casein	(184–185)
8768	12	DQ	α_{S1} -casein	(66–67)
			α_{S2} -casein	(125–126)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
7623	9	EA	α_{S1} -casein	(76-77)
			β -casein	(100-101)
			κ -casein	(158-159)
7622/8770	1	EG	α_{S1} -casein	(140-141)
8771	12	EH	α_{S2} -casein	(20-21)
7826/8772	1	EI	α_{S1} -casein	(85-86), (125-126)
			α_{S2} -casein	(99-100)
			β -casein	(11-12)
			κ -casein	(139-140), (179-180)
7840/8558	1	EK	α_{S1} -casein	(50-51), (207-208)
			α_{S2} -casein	(38-39), (172-173)
			β -casein	(31-32)
			κ -casein	(33-34)
7888	11	EL	α_{S1} -casein	(54-55), (156-157), (163-164)
			β -casein	(2-3), (5-6), (44-45)
8529	12	EP	α_{S1} -casein	(148-149)
			β -casein	(117-118), (195-196)
			κ -casein	(150-151)
8773	12	ES	α_{S1} -casein	(62-63), (78-79)
			α_{S2} -casein	(27-28), (75-76), (157-158)
			β -casein	(14-15), (21-22)
			κ -casein	(161-162), (175-176)
8774	12	ET	α_{S2} -casein	(33-34)
7828/8775	1	EV	α_{S1} -casein	(29-30), (45-46)
			α_{S2} -casein	(57-58), (78-79), (83-84), (160-161)
			β -casein	(91-92)
			κ -casein	(172-173)
7752/8777	1	EY	α_{S2} -casein	(66-67)
3176	12	FA	α_{S2} -casein	(189-190)
			β -casein	(52-53)
9342	11	FC	α_{S2} -casein	(54-55)
7605	9	FG	α_{S1} -casein	(47-48)
8555	12	FL	α_{S2} -casein	(178-179)
			β -casein	(87-88), (190-191)
			κ -casein	(6-7), (18-19), (76-77)
3502/8506	1	FP	α_{S1} -casein	(43-44)
			α_{S2} -casein	(107-108)
			β -casein	(62-63), (111-112), (157-158), (205-206)
8779/9076	1	FQ	β -casein	(33-34)
7592/8780	1	FR	α_{S1} -casein	(165-166)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
3556	9	FY	α_{S1} -casein	(160–161), (168–169)
			α_{S2} -casein	(103–104)
7598/8524	1	GA	α_{S1} -casein	(177–178)
			κ -casein	(20–21)
7615/8781	1	GE	β -casein	(10–11)
			κ -casein	(149–150)
7596/8785	1	GI	α_{S1} -casein	(141–142)
7611	9	GK	α_{S1} -casein	(48–49)
7599/8561	1	GL	α_{S1} -casein	(25–26)
			κ -casein	(60–61)
3169/7512	1	GP	α_{S2} -casein	(117–118)
			β -casein	(64–65), (199–200), (203–204)
7607	9	GS	α_{S1} -casein	(60–61), (202–203)
			α_{S2} -casein	(70–71)
7612	9	GT	α_{S1} -casein	(185–186)
7608/8786	1	GV	α_{S1} -casein	(152–153)
			β -casein	(94–95)
3532/8788	1	GY	α_{S1} -casein	(108–109)
3184	12	HA	α_{S1} -casein	(143–144)
8793	12	HI	α_{S1} -casein	(95–96)
7844	9	HK	β -casein	(106–107)
3317/7602/8557	7	HL	β -casein	(134–135)
			κ -casein	(123–124)
7842/8520	1	HP	α_{S1} -casein	(19–20)
			κ -casein	(119–120), (121–122)
			β -casein	(50–51)
8795	12	HS	α_{S1} -casein	(136–137)
8797	12	HV	α_{S2} -casein	(21–22)
3494/8799	1	HY	α_{S2} -casein	(92–93)
7562/8525	1	IA	κ -casein	(43–44), (146–147)
			β -casein	(30–31)
7827	9	IE	κ -casein	(174–175)
			β -casein	(30–31)
7593	9	IF	α_{S2} -casein	(5–6)
7595	9	IG	α_{S1} -casein	(59–60), (151–152), (201–202)
			α_{S2} -casein	(69–70)
8800	12	IH	α_{S1} -casein	(142–143)
			β -casein	(49–50)
8801	12	II	α_{S2} -casein	(29–30)
			β -casein	(207–208)
8802/9079	1	IL	α_{S1} -casein	(5–6)
			κ -casein	(11–12), (94–95)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8804	12	IN	α_{S2} -casein	(43–44), (100–101)
			β -casein	(26–27)
			κ -casein	(72–73), (143–144), (180–181)
7581/8501	1	IP	α_{S1} -casein	(197–198)
			α_{S2} -casein	(216–217)
			β -casein	(66–67), (74–75)
8805	12	IQ	κ -casein	(47–48), (129–130), (140–141)
			α_{S1} -casein	(96–97)
			α_{S2} -casein	(209–210)
3258/8215/8806	7	IR	β -casein	(187–188)
3174/7743	1	KA	κ -casein	(49–50)
			α_{S2} -casein	(30–31)
8134	11	KD	α_{S2} -casein	(95–96), (203–204)
			β -casein	(176–177)
7841/8808	1	KE	α_{S1} -casein	(57–58)
			κ -casein	(34–35)
			α_{S1} -casein	(49–50), (98–99), (139–140), (147–148)
7692/8809	1	KF	α_{S2} -casein	(47–48), (56–57)
			β -casein	(99–100), (107–108)
			α_{S2} -casein	(2–3), (106–107), (188–189)
8811	12	KH	β -casein	(32–33)
			α_{S1} -casein	(18–19), (22–23), (94–95)
			α_{S2} -casein	(91–92)
8812	12	KI	β -casein	(105–106)
			α_{S2} -casein	(85–86), (181–182)
			β -casein	(29–30), (48–49)
8813	12	KK	κ -casein	(42–43)
			α_{S1} -casein	(117–118)
			α_{S2} -casein	(151–152), (164–165), (180–81)
7693	9	KL	β -casein	(28–29)
			κ -casein	(132–133)
7810/8218/8519	7	KP	α_{S1} -casein	(2–3)
			α_{S2} -casein	(167–168)
7586/8814	1	KR	α_{S2} -casein	(206–207)
			κ -casein	(67–68), (84–85)
8815	12	KS	α_{S2} -casein	(128–129)
8816	12	KT	κ -casein	(3–4), (107–108)
			α_{S1} -casein	(208–209)
			α_{S2} -casein	(152–153), (165–166), (196–197), (212–213)
			κ -casein	(137–138)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8817	12	KV	α_{S1} -casein	(51–52), (120–121)
			α_{S2} -casein	(214–215)
			β -casein	(97–98), (169–170)
7691/8819	1	KY	α_{S1} -casein	(118–119)
			β -casein	(113–114)
			κ -casein	(45–46)
3175/7585	1	LA	α_{S1} -casein	(14–15), (157–158)
			α_{S2} -casein	(10–11), (14–15)
			κ -casein	(12–13)
3551	9	LF	α_{S1} -casein	(164–165)
7619	9	LG	α_{S1} -casein	(107–108), (184–185)
			β -casein	(198–199)
			κ -casein	(19–20)
3305/8820	6	LH	α_{S1} -casein	(135–136)
			β -casein	(133–134)
8821	12	LI	α_{S1} -casein	(4–5)
			κ -casein	(71–72)
8217	11	LK	α_{S1} -casein	(116–117)
			α_{S2} -casein	(179–180), (195–196)
3182	12	LL	α_{S1} -casein	(3–4), (35–36), (113–114)
			α_{S2} -casein	(9–10)
			β -casein	(139–140), (191–192)
7832/8823	1	LN	α_{S1} -casein	(31–32)
			α_{S2} -casein	(97–98), (121–122), (138–139), (176–177)
			β -casein	(6–7)
3180	12	LP	κ -casein	(61–62)
			α_{S1} -casein	(26–27)
			α_{S2} -casein	(191–192)
7831	9	LQ	β -casein	(70–71), (135–136), (137–138), (151–152), (171–172)
			κ -casein	(16–17), (77–78)
			α_{S2} -casein	(111–112)
9213	9	LR	β -casein	(45–46), (88–89), (140–141)
			κ -casein	(95–96)
			α_{S1} -casein	(36–37), (114–115)
8824	12	LT	α_{S1} -casein	(6–7)
			α_{S2} -casein	(168–169)
			β -casein	(77–78), (125–126), (127–128)
			κ -casein	(14–15)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8825	12	LV	α_{S1} -casein	(9–10)
			β -casein	(58–59)
			κ -casein	(7–8)
3389/8462/8688	7	LW	α_{S1} -casein	(213–214)
3381/7872	2	LY	α_{S2} -casein	(114–115)
			β -casein	(192–193)
3173	12	MA	α_{S2} -casein	(41–42)
			β -casein	(102–103)
			κ -casein	(116–117), (127–128)
7839/8826	1	ME	α_{S1} -casein	(69–70), (75–76)
			α_{S2} -casein	(19–20), (156–157)
3385/8827	1	MF	β -casein	(156–157)
7609/8828	1	MG	β -casein	(93–94)
8829	12	MH	β -casein	(144–145)
8830	12	MI	α_{S1} -casein	(150–151)
8831	12	MK	α_{S1} -casein	(1–2), (138–139)
			α_{S2} -casein	(1–2), (205–206)
			κ -casein	(2–3)
8833/9085/9086	7	MM	κ -casein	(1–2)
3171	12	MP	α_{S1} -casein	(211–212)
			β -casein	(109–110), (185–186)
8839	12	NA	α_{S2} -casein	(61–62), (130–131)
8841	12	NE	α_{S1} -casein	(32–33), (53–54)
			α_{S2} -casein	(63–64), (98–99)
7683/8842	1	NF	α_{S2} -casein	(177–178)
7698	9	NK	β -casein	(27–28)
8845	12	NL	α_{S1} -casein	(34–35)
			α_{S2} -casein	(49–50)
			β -casein	(132–133)
8846	12	NM	α_{S2} -casein	(40–41)
8847	12	NN	κ -casein	(73–74)
8530	12	NP	α_{S1} -casein	(199–200)
			α_{S2} -casein	(44–45), (122–123)
8848	12	NQ	α_{S1} -casein	(154–155)
			α_{S2} -casein	(101–102)
			κ -casein	(25–26), (74–75), (134–135)
8849	12	NR	α_{S2} -casein	(139–140), (174–175)
8850	12	NT	α_{S2} -casein	(17–18)
			κ -casein	(102–103), (144–145), (181–182)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8851	12	NV	β -casein	(7–8)
7682/8853	1	NY	κ -casein	(62–63)
3179	12	PA	κ -casein	(85–86), (91–92), (105–106)
			α_{S1} -casein	(42–43)
8854	12	PF	β -casein	(51–52), (61–62), (86–87), (110–111), (118–119), (204–205)
			κ -casein	(17–18)
7625/8855	1	PG	β -casein	(9–10), (63–64)
7843/8856	1	PH	β -casein	(147–148)
			κ -casein	(120–121), (122–123)
			α_{S1} -casein	(20–21), (200–201)
8857	12	PI	α_{S2} -casein	(118–119), (133–134)
			β -casein	(65–66), (186–187), (206–207)
			κ -casein	(29–30), (48–49)
			α_{S1} -casein	(17–18)
8858	12	PK	α_{S2} -casein	(211–212)
			β -casein	(104–105), (112–113)
			κ -casein	(131–132)
7513/8638	1	PL	α_{S1} -casein	(183–184), (212–213)
			β -casein	(76–77), (136–137), (138–139), (150–151)
8859	12	PM	α_{S1} -casein	(149–150)
8860	12	PN	α_{S1} -casein	(88–89), (128–129), (198–199)
			β -casein	(67–68)
3170/7836	1	PP	β -casein	(75–76), (85–86), (152–153), (158–159)
			κ -casein	(130–131), (177–178)
			α_{S1} -casein	(27–28), (122–123)
7837/8861	1	PQ	α_{S2} -casein	(108–109), (192–193)
			β -casein	(71–72), (159–160), (174–175), (181–182)
8862	12	PS	α_{S1} -casein	(102–103), (175–176), (192–193)
			α_{S2} -casein	(45–46)
			κ -casein	(57–58)
			α_{S2} -casein	(136–137)
7833/8863	1	PT	β -casein	(153–154)
			κ -casein	(113–114), (141–142), (151–152), (155–156)
8864	12	PV	β -casein	(81–82), (115–116), (172–173), (196–197), (200–201)
			κ -casein	(68–69)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8190/8865	6	PW	α_{S2} -casein	(123–124), (207–208)
8866	12	PY	α_{S2} -casein	(217–218)
			β -casein	(179–180)
			κ -casein	(78–79), (80–81)
8867	12	QA	α_{S1} -casein	(67–68)
			β -casein	(188–189)
			κ -casein	(110–111)
8868	12	QD	β -casein	(46–47)
			κ -casein	(135–136)
8869	12	QE	α_{S1} -casein	(28–29), (155–156)
			α_{S2} -casein	(32–33), (37–38)
			β -casein	(194–195)
			κ -casein	(22–23), (26–27)
8870	12	QF	α_{S1} -casein	(167–168)
			α_{S2} -casein	(102–103)
			κ -casein	(75–76)
7617/8871	1	QG	α_{S1} -casein	(24–25)
			α_{S2} -casein	(116–117)
8872	12	QH	α_{S2} -casein	(200–201)
8873	12	QI	κ -casein	(93–94)
			α_{S1} -casein	(93–94), (97–98), (146–147)
7680	9	QK	α_{S2} -casein	(94–95), (105–106), (187–188), (202–203)
			β -casein	(175–176)
			κ -casein	(66–67)
8874	12	QL	α_{S1} -casein	(112–113), (123–124), (170–171)
			α_{S2} -casein	(142–143)
8875	12	QN	β -casein	(72–73)
			κ -casein	(24–25)
8532	12	QP	α_{S2} -casein	(210–211)
			β -casein	(89–90), (146–147), (149–150)
			κ -casein	(28–29), (112–113)
8876	12	QQ	α_{S1} -casein	(145–146)
			β -casein	(38–39), (39–40)
			κ -casein	(65–66)
8877	12	QS	β -casein	(34–35), (56–57), (123–124), (141–142), (160–161), (167–168)
8878	12	QT	β -casein	(40–41), (54–55), (79–80)
8879	12	QV	α_{S2} -casein	(126–127)
			κ -casein	(98–99), (184–185)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8880	12	QW	κ -casein	(96–97)
8881	12	QY	α_{S1} -casein	(187–188)
			α_{S2} -casein	(109–110), (112–113), (193–194)
			κ -casein	(50–51)
3489	9	RF	α_{S1} -casein	(37–38)
			κ -casein	(37–38)
8882/9173	1	RG	β -casein	(202–203)
8883	12	RH	κ -casein	(118–119)
8884	12	RI	β -casein	(25–26)
3257/8886	1	RL	α_{S1} -casein	(115–116), (134–135)
			α_{S2} -casein	(175–176)
8888	12	RN	α_{S2} -casein	(60–61), (129–130)
7582/8518	1	RP	α_{S1} -casein	(16–17)
3380	9	RY	α_{S1} -casein	(105–106)
			α_{S2} -casein	(185–186), (220–221)
			κ -casein	(55–56)
7685/8891	1	SF	α_{S1} -casein	(193–194)
			κ -casein	(4–5), (125–126)
7618	9	SG	α_{S1} -casein	(176–177)
			κ -casein	(148–149)
8893	12	SI	α_{S1} -casein	(79–80)
			α_{S2} -casein	(28–29), (68–69)
			β -casein	(22–23)
8894	12	SK	α_{S1} -casein	(56–57)
			α_{S2} -casein	(46–47), (150–151)
			β -casein	(96–97), (168–169)
8560	12	SL	β -casein	(15–16), (57–58), (69–70), (124–125), (164–165)
8505	12	SP	κ -casein	(90–91), (170–171), (176–177)
9184	9	ST	α_{S1} -casein	(63–64)
			α_{S2} -casein	(52–53), (144–145), (158–159)
			κ -casein	(153–154), (162–163), (187–188)
8895	12	SV	α_{S1} -casein	(90–91)
			β -casein	(161–162)
8896	12	SW	β -casein	(142–143)
7684/8897	1	SY	κ -casein	(58–59)
8531	12	TA	κ -casein	(188–189)
8898	12	TD	α_{S1} -casein	(189–190)
			β -casein	(128–129)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
7830/8899	1	TE	α_{S1} -casein	(64–65)
			α_{S2} -casein	(81–82), (159–160), (169–170)
			β -casein	(41–42), (120–121)
			κ -casein	(138–139), (157–158)
8185/8900	1	TF	α_{S2} -casein	(53–54)
8903	12	TI	κ -casein	(10–11), (142–143), (145–146)
8904	12	TK	α_{S2} -casein	(163–164), (166–167), (213–214)
8905	12	TL	α_{S2} -casein	(137–138)
			β -casein	(126–127)
			κ -casein	(15–16), (166–167)
8906	12	TM	α_{S1} -casein	(210–211)
			α_{S2} -casein	(18–19)
			κ -casein	(115–116)
8503/9073	1	TP	α_{S2} -casein	(135–136)
			β -casein	(80–81)
			κ -casein	(154–155)
7834/8908	1	TQ	α_{S1} -casein	(186–187)
			β -casein	(55–56), (78–79)
8909	12	TR	β -casein	(24–25)
8910	12	TS	α_{S2} -casein	(145–146)
			κ -casein	(152–153), (186–187)
8911	12	TT	α_{S1} -casein	(209–210)
			κ -casein	(114–115), (156–157)
8912	12	TV	α_{S2} -casein	(87–88), (153–154), (197–198)
			β -casein	(154–155)
			κ -casein	(103–104), (163–164), (182–183)
8219/8914	6	TY	α_{S2} -casein	(34–35)
3172	12	VA	α_{S1} -casein	(10–11), (12–13), (40–41)
			α_{S2} -casein	(12–13), (79–80)
			κ -casein	(69–70), (164–165)
8915	12	VD	α_{S2} -casein	(88–89), (154–155)
7829/8916	1	VE	α_{S1} -casein	(91–92)
			β -casein	(13–14), (116–117), (130–131)
			κ -casein	(160–161)
3384/8917	1	VF	α_{S1} -casein	(46–47)
			α_{S2} -casein	(161–162)
8920	12	VI	α_{S2} -casein	(215–216)
			κ -casein	(173–174)
7558/8921	1	VK	α_{S2} -casein	(84–85), (127–128)
			β -casein	(98–99)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8922	12	VL	α_{S1} -casein	(30–31)
			α_{S2} -casein	(120–121)
			β -casein	(162–163), (170–171), (197–198)
			κ -casein	(52–53), (99–100)
8923	12	VM	β -casein	(92–93), (155–156)
8924	12	VN	α_{S1} -casein	(52–53), (153–154)
3181/7587	1	VP	α_{S1} -casein	(87–88), (101–102), (121–122), (127–128), (182–183)
			α_{S2} -casein	(132–133)
			β -casein	(8–9), (84–85), (173–174), (178–179)
			κ -casein	(104–105)
8925	12	VQ	κ -casein	(183–184)
7628/8594	1	VR	α_{S2} -casein	(59–60), (219–220)
			β -casein	(201–202)
			κ -casein	(88–89)
8926	12	VS	α_{S2} -casein	(22–23)
			β -casein	(95–96)
8927	12	VT	κ -casein	(9–10), (185–186)
3183	12	VV	α_{S2} -casein	(58–59)
			β -casein	(82–83), (83–84)
			κ -casein	(8–9)
3492/8224/8929	7	VY	α_{S2} -casein	(198–199)
			β -casein	(59–60)
8930	12	WD	α_{S2} -casein	(124–125)
8679	12	WI	α_{S2} -casein	(208–209)
8682/9090	1	WM	β -casein	(143–144)
8678	12	WQ	κ -casein	(97–98)
7898/8683	6	WY	α_{S1} -casein	(179–180)
7589/8932	1	YA	κ -casein	(82–83)
8935	12	YF	α_{S1} -casein	(159–160)
3553/8936	1	YG	κ -casein	(59–60)
8938	12	YI	κ -casein	(46–47)
7697/8939	1	YK	α_{S1} -casein	(119–120)
			α_{S2} -casein	(35–36)
8940	12	YL	α_{S1} -casein	(106–107), (109–110)
			α_{S2} -casein	(110–111), (113–114), (194–195), (221–222)
3666/8521/9548	8	YP	α_{S1} -casein	(161–162), (174–175)
			β -casein	(60–61), (114–115), (180–181)
			κ -casein	(56–57), (79–80)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8943	12	YQ	α_{S1} -casein	(169–170)
			α_{S2} -casein	(93–94), (104–105), (115–116), (186–187), (199–200)
			β -casein	(193–194)
			κ -casein	(64–65)
8945	12	YS	α_{S2} -casein	(67–68)
8696	12	YT	α_{S1} -casein	(188–189)
8946/9077	1	YV	α_{S1} -casein	(181–182)
			α_{S2} -casein	(218–219)
			κ -casein	(51–52)
8948	12	YY	α_{S1} -casein	(180–181)
			κ -casein	(63–64), (81–82)
9042	9	AFL	β -casein	(189–191)
3597	9	AIP	κ -casein	(128–130)
9029	9	ALP	α_{S2} -casein	(190–192)
3370	9	AVP	α_{S2} -casein	(131–133)
			β -casein	(177–179)
9359	11	CLV	α_{S1} -casein	(8–10)
7823	9	FAL	α_{S2} -casein	(189–191)
7560	9	FFL	κ -casein	(5–7)
3377	9	FGK	α_{S1} -casein	(47–49)
7554	9	GEP	κ -casein	(149–151)
7509	9	GLP	α_{S1} -casein	(25–27)
7545/9116	1	GPV	β -casein	(199–201)
3311	11	HPH	κ -casein	(119–121), (121–123)
9027	9	HQG	α_{S1} -casein	(23–25)
7626	9	IAK	κ -casein	(43–45)
3167	12	IPI	κ -casein	(47–49)
3522	9	IPP	β -casein	(74–76)
			κ -casein	(129–131)
7803	9	IPY	α_{S2} -casein	(216–218)
8184/8693	1	IQP	α_{S2} -casein	(209–211)
8149/8180	2	IQY	κ -casein	(49–51)
9365	11	KKY	α_{S1} -casein	(117–119)
8133	11	KVI	α_{S2} -casein	(214–216)
3558	9	LAY	α_{S1} -casein	(157–159)
9031	9	LEE	β -casein	(3–5)
9206	9	LFR	α_{S1} -casein	(164–166)
7508	9	LGP	β -casein	(198–200)
7995	11	LHL	β -casein	(133–135)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8000	11	LHS	α_{S1} -casein	(135–137)
8484	11	LLR	α_{S1} -casein	(35–37), (113–115)
8509	9	LNF	α_{S2} -casein	(176–178)
3544	9	LNP	α_{S2} -casein	(121–123)
9026	9	LNK	κ -casein	(61–63)
9040	9	LPF	κ -casein	(16–18)
8616	12	LPL	β -casein	(135–137), (137–139)
3391	9	LPP	β -casein	(151–153)
9339	12	LPQ	α_{S1} -casein	(26–28)
			α_{S2} -casein	(191–193)
			β -casein	(70–72)
3542/8689	1	LQP	β -casein	(88–90)
9163	11	LTC	α_{S1} -casein	(6–8)
8402	9	LVY	β -casein	(58–60)
9241	9	MAP	β -casein	(102–104)
8958	9	MKP	α_{S2} -casein	(205–207)
9363	11	NEN	α_{S1} -casein	(32–34)
7965	11	NYK	κ -casein	(62–64)
8139	11	PEL	α_{S1} -casein	(162–164)
8029	11	PHL	κ -casein	(122–124)
8032	11	PHQ	β -casein	(147–149)
3094/7510	5	PLG	α_{S1} -casein	(183–185)
2664	9	PLP	β -casein	(136–138), (150–152)
3531	9	PLW	α_{S1} -casein	(212–214)
7645	9	PPK	κ -casein	(130–132)
8652	12	PPL	β -casein	(75–77)
3373	9	PQR	β -casein	(181–183)
7559	9	PSY	κ -casein	(57–59)
8040	11	PWD	α_{S2} -casein	(123–125)
8044	11	PWI	α_{S2} -casein	(207–209)
3372	9	PYP	β -casein	(179–181)
			κ -casein	(78–80)
3307	11	PYY	κ -casein	(80–82)
9146	9	QGP	α_{S2} -casein	(116–118)
9543	9	RYL	α_{S1} -casein	(105–107)
			α_{S2} -casein	(220–222)
8382/8474	2	RYQ	α_{S2} -casein	(185–187)
7498	9	TVY	α_{S2} -casein	(197–199)
3521	9	VAP	α_{S1} -casein	(40–42)
7635	9	VAV	α_{S1} -casein	(10–12)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
2653	9	VLP	β -casein	(170–172)
8347	12	VPL	α_{S1} -casein	(182–184)
3524	9	VPP	β -casein	(84–86)
9046	9	VQV	κ -casein	(183–185)
8384	9	VRV	α_{S2} -casein	(219–221)
3505	9	VYP	β -casein	(59–61)
7962	11	WYY	α_{S1} -casein	(179–181)
7963	11	YFY	α_{S1} -casein	(159–161)
3973	9	YGL	κ -casein	(59–61)
7959	11	YLY	α_{S2} -casein	(113–115)
2870	15	YPF	β -casein	(60–62)
8617/9550	3	YPY	κ -casein	(79–81)
8487	11	YQL	α_{S1} -casein	(169–171)
8150	11	YVL	κ -casein	(51–53)
7634	9	YVP	α_{S1} -casein	(181–183)
7939	11	YYA	κ -casein	(81–83)
7948	11	YYQ	κ -casein	(63–65)
7942	11	YYV	α_{S1} -casein	(180–182)
8369	9	AFLI	β -casein	(189–192)
7668	9	AIPP	κ -casein	(128–131)
8380/8472	2	AYPS	α_{S1} -casein	(173–176)
8591	12	FLQP	β -casein	(87–90)
7499	9	FVAP	α_{S1} -casein	(39–42)
8489	11	FYQL	α_{S1} -casein	(168–171)
8371/8467	2	GTQY	α_{S1} -casein	(185–188)
3306	11	HPHL	κ -casein	(121–124)
7661	9	LFRQ	α_{S1} -casein	(164–167)
2668	9	LHLP	β -casein	(133–136)
2665	9	LPLP	β -casein	(135–138)
3397	9	LQSW	β -casein	(140–143)
9236	9	LRFF	α_{S1} -casein	(36–39)
3331	9	LVYP	β -casein	(58–61)
3374	9	PFPE	α_{S1} -casein	(42–45)
7896	11	PYPQ	β -casein	(179–182)
8357	9	RYLG	α_{S1} -casein	(105–108)
8378/8470	2	RYPS	κ -casein	(55–58)
8221	11	TSTA	κ -casein	(186–189)
8483	11	TVYQ	α_{S2} -casein	(197–200)
8593	12	VLGP	β -casein	(197–200)
8309	9	VRSP	κ -casein	(88–91)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
7801	9	VRYL	α_{S2} -casein	(219–222)
8308	9	VVPP	β -casein	(83–86)
8592	12	WIQP	α_{S2} -casein	(208–211)
8381/8473	2	YAKP	κ -casein	(82–85)
8485	11	YGLN	κ -casein	(59–62)
9420/9421	2	YLGY	α_{S1} -casein	(106–109)
8482	11	YLKT	α_{S2} -casein	(194–197)
8140	11	YPEL	α_{S1} -casein	(161–164)
2868	15	YFPF	β -casein	(60–63)
3218/8615	4	YPYY	κ -casein	(79–82)
7878	11	YQEP	β -casein	(193–196)
8477	11	YQLD	α_{S1} -casein	(169–172)
3399/7660	9	AMKPW	α_{S2} -casein	(204–208)
9229	9	ARHPH	κ -casein	(117–121)
3371	9	AVPYP	β -casein	(177–181)
3519	9	AYFYP	α_{S1} -casein	(158–162)
3497	9	FFVAP	α_{S1} -casein	(38–42)
8368	9	FPIIV	β -casein	(205–209)
9378	9	FPPQS	β -casein	(157–161)
8141	11	FYPEL	α_{S1} -casein	(160–164)
9375	9	GLPQE	α_{S1} -casein	(25–29)
8478	11	GYLEQ	α_{S1} -casein	(108–112)
2666	9	HLPLP	β -casein	(134–138)
8613	12	IPIQY	κ -casein	(47–51)
7627	9	KDERF	κ -casein	(34–38)
9552	9	KFPQY	α_{S2} -casein	(106–110)
8618	12	LPLPL	β -casein	(135–139)
7631/8614	1	LPYPY	κ -casein	(77–81)
2671	9	NLHLP	β -casein	(132–136)
8157	9	PGPIP	β -casein	(63–67)
9242	9	PLPLL	β -casein	(136–140)
9243	9	PPEIN	κ -casein	(177–181)
8373/8469	2	RHPHP	κ -casein	(118–122)
9244	9	RINKK	β -casein	(25–29)
8481	11	RLKKY	α_{S1} -casein	(115–119)
8358	9	RYLGY	α_{S1} -casein	(105–109)
8375	9	RYQKF	α_{S2} -casein	(185–189)
9246	9	SLPQN	β -casein	(69–73)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
3332	9	SLVYP	β -casein	(57–61)
8379/8471	2	SRYP	κ -casein	(54–58)
7485	9	TKVIP	α_{S2} -casein	(213–217)
8278/9556	6	VPYPQ	β -casein	(178–182)
8370/8466	2	YAKPA	κ -casein	(82–86)
8480	11	YPELF	α_{S1} -casein	(161–165)
3263	13	YPFPG	β -casein	(60–64)
8486	11	YQKFP	α_{S2} -casein	(104–108)
8372/8468	2	ARHPHP	κ -casein	(117–122)
3501	9	AYFYPE	α_{S1} -casein	(158–163)
7493	9	DKIHPF	β -casein	(47–52)
7629	9	EKDERF	κ -casein	(33–38)
3511	9	EMPFPK	β -casein	(108–113)
7480	9	FALPQY	α_{S2} -casein	(189–194)
7487	9	KVLPVP	β -casein	(169–174)
3513	9	LAYFYP	α_{S1} -casein	(157–162)
2667	9	LHLPLP	β -casein	(133–138)
8179	9	LKKISQ	α_{S2} -casein	(179–184)
9238	9	LVYFPF	β -casein	(58–63)
8152/8178	2	PYVRYL	α_{S2} -casein	(217–222)
3426	9	QSLVYP	β -casein	(56–61)
8479	11	RDMPIQ	β -casein	(183–188)
3128	14	RYLGYL	α_{S1} -casein	(105–110)
3217	14	SRYP	κ -casein	(54–59)
3127/3530	5	TTMPLW	α_{S1} -casein	(209–214)
9248	9	VLSRYP	κ -casein	(52–57)
8658	12	VPITPT	α_{S2} -casein	(132–137)
7481	9	VTSTAV	κ -casein	(185–190)
9251	9	VVVPPF	β -casein	(82–87)
3504	9	VYFPFG	β -casein	(59–64)
7887/8376	2	YFYPEL	α_{S1} -casein	(159–164)
7630	9	YIPIQY	κ -casein	(46–51)
3500	9	YKVPQL	α_{S1} -casein	(119–124)
3265	15	YLGYLE	α_{S1} -casein	(106–111)
9252	12	YPVEPF	β -casein	(114–119)
7662/9253	9	YQEPVL	β -casein	(193–198)
8383	9	YYQQKP	κ -casein	(63–68)
3480/7875	2	AVPYPQR	β -casein	(177–183)
8359	9	AYFYPEL	α_{S1} -casein	(158–164)
7496	9	FPEVFGK	α_{S1} -casein	(43–49)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
8659	12	FPGPIP	β -casein	(62–68)
7659	9	FPQYLQY	α_{S2} -casein	(107–113)
8181	9	FSDKIAK	κ -casein	(39–45)
8159	9	GPFPIIV	β -casein	(203–209)
8285	11	IPIQYVL	κ -casein	(47–53)
9373	9	KEDVPSE	α_{S1} -casein	(98–104)
9234	9	KKYKVPQ	α_{S1} -casein	(117–123)
3498	9	KVLPVPQ	β -casein	(169–175)
7565	9	LHLPLPL	β -casein	(133–139)
8657	12	LPQNIPP	β -casein	(70–76)
8155	9	LTLTDE	β -casein	(125–131)
2669	9	NLHLPLP	β -casein	(132–138)
9058	12	PQNIPPL	β -casein	(71–77)
3473	13	RYLGYLE	α_{S1} -casein	(105–111)
8476	11	TIASGEP	κ -casein	(145–151)
3333	9	TQSLVYP	β -casein	(55–61)
7877	11	VLPVPQK	β -casein	(170–176)
8475	11	VLSRYPS	κ -casein	(52–58)
9249	12	VPITPTL	α_{S2} -casein	(132–138)
9250	9	VPSELYL	α_{S1} -casein	(101–107)
3262/7665	5	YPPFGPI	β -casein	(60–66)
3214	15	YPSYGLN	κ -casein	(56–62)
9254	9	YQKFPQY	α_{S2} -casein	(104–110)
8377	9	AYFYPELF	α_{S1} -casein	(158–165)
3512	9	DAYPSGAW	α_{S1} -casein	(172–179)
9539	9	EKVNELSK	α_{S1} -casein	(50–57)
2672	9	ENLHLPLP	β -casein	(131–138)
7658/8151	2	FALPQYLK	α_{S2} -casein	(189–196)
7876	11	KVLPVPQK	β -casein	(169–176)
9235	12	LPQNIPPL	β -casein	(70–77)
3398	9	MKPWIQPK	α_{S2} -casein	(205–212)
7802	9	NMAINPSK	α_{S2} -casein	(40–47)
9168/9273	9	PFPEVFGK	α_{S1} -casein	(42–49)
3334	9	QTQSLVYP	β -casein	(54–61)
8160	9	RGPFPIIV	β -casein	(202–209)
7796	11	VKEAMAPK	β -casein	(98–105)
7492	9	VYPPFGPI	β -casein	(59–66)
2869/7501	5	YPPFGPIP	β -casein	(60–67)
7879	11	YQEPVLGP	β -casein	(193–200)
7804	9	ALNEINQFY	α_{S2} -casein	(96–104)

Table A1. Cont.

BIOPEP-UWM ID	Activity	Sequence	Protein	Location
3395/7497	9	AMKPWIQPK	α_{S2} -casein	(204–212)
3335	9	AQTQSLVYP	β -casein	(53–61)
8660	12	IPPLTQTPV	β -casein	(74–82)
9380	9	LGPVRGPFPP	β -casein	(198–206)
7494	9	LNVPGEIVE	β -casein	(6–14)
8396	9	QSWMHQPHQ	β -casein	(141–149)
7483	9	RPKHPIKHQ	α_{S1} -casein	(16–24)
9376	9	VEQKHIQKE	α_{S1} -casein	(91–99)
7566	9	VRGPFPIIV	β -casein	(201–209)
7486	9	YFPFGPIPNN	β -casein	(60–68)
8156	9	YPQRDMPIQ	β -casein	(180–188)
8397	9	YYAKPAAVR	κ -casein	(81–89)
3336	9	FAQTQSLVYP	β -casein	(52–61)
8169/8286	2	GPVRGPFPII	β -casein	(199–208)
9231	12	INNQFLPYPY	κ -casein	(72–81)
7495	9	NIPPLTQTPV	β -casein	(73–82)
3216/3495	5	YIPIQYVLSR	κ -casein	(46–55)
7805	9	ALNEINQFYQK	α_{S2} -casein	(96–106)
7797	11	ARHPHPHLSFM	κ -casein	(117–127)
9237	9	LTQTPVVVPPF	β -casein	(77–87)
3503	9	TPVVVPPFLQP	β -casein	(80–90)
3251	13	YFPFGPIPNSL	β -casein	(60–70)
9377	9	EIVPNSAEERLH	α_{S1} -casein	(125–136)
7800	9	FFVAPFPEVFGK	α_{S1} -casein	(38–49)
3337	9	HPFAQTQSLVYP	β -casein	(50–61)
3338	9	IHPFAQTQSLVYP	β -casein	(49–61)
9379	9	QEPVLGPVRGPFPP	β -casein	(194–206)
3339	9	KIHPFAQTQSLVYP	β -casein	(48–61)
9240	9	LVYFPFGPIPNSLPQN	β -casein	(58–73)
7564	9	LVYFPFGPIPNSLPQNIPP	β -casein	(58–76)

*1-ACE and DPP4 inhibitor; 2-ACE inhibitory and antioxidative activity; 3-DPP4 and α -glucosidase inhibitor; 4-DPP4 inhibitory and opioid activity; 5- ACE inhibitory and opioid activity; 6- DPP4 inhibitory and antioxidative activity; 7- ACE, DPP4 inhibitory, and antioxidative activity; 8- ACE, DPP4, and α -glucosidase inhibitor; 9-ACE inhibitor; 10- α -glucosidase inhibitor; 11- antioxidative activity; 12- DPP4 inhibitor; 13-opioid agonist activity; 14 - opioid antagonist activity; 15-opioid activity.

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