

1 **Antihypertensive peptides: production, bioavailability and incorporation into foods**

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15 **Abstract**

16 Bioactive food peptides are encrypted within the source protein but can exert
17 physiological properties once released by enzymatic hydrolysis during gastrointestinal transit,
18 fermentation or maturation during food processing, or proteolysis by food-grade enzymes derived
19 from microorganisms or plants. Among the bioactive food peptides, those with antihypertensive
20 activity are receiving special attention due to the high prevalence of hypertension in the Western
21 countries and its role in cardiovascular diseases. This paper reviews the current literature on
22 antihypertensive food peptides, focusing on the existing methodologies for their production, such
23 as enzymatic hydrolysis, fermentation and genetic recombination in bacteria. This paper also
24 evaluates the structure/activity relationship of angiotensin-converting enzyme (ACE) inhibitory
25 peptides, as well as their bioavailability, physiological effects demonstrated by both *in vitro* and
26 *in vivo* assays, and the existence of mechanisms of action other than ACE inhibition. Finally,
27 current reported strategies for incorporation of antihypertensive peptides in foods and their
28 effects on both availability and activity of these peptides are revised in this manuscript.

29
30 **Keywords:** Antihypertensive peptides, angiotensin-converting enzyme inhibitory activity,
31 enzymatic hydrolysis, fermentation, genetic recombination, bioavailability.

32 **1. Introduction**

33 In recent years, increasing epidemiological evidence is linking the prevalence of diseases,
34 such as cardiovascular disease, obesity, hypertension, diabetes, and even cancer to dietary factors.
35 Manufacture of new foods termed functional foods is emerging in response to the increased
36 perception about the relation of food and health. A functional food is generally any food which
37 can provide a health benefit to one or more bodily functions beyond that of basic nutrition [1].
38 Recently, it has been recognized that apart from their basic nutritional role, many dietary proteins
39 contain, encrypted within their primary structure, different peptide sequences that exert beneficial
40 effects upon human health once released by digestive enzymes during gastrointestinal transit or
41 by fermentation or ripening during food processing. Bioactive peptides range in size from 2 to 50
42 amino acid residues and exhibit different activities, such as antimicrobial, antioxidant,
43 antithrombotic, antihypertensive, immunomodulatory, opioid, and antiproliferative activities,
44 among others [2-4], affecting the major body systems – namely, the cardiovascular, digestive,
45 endocrine, immune and nervous systems. The potential of these bioactive peptides to reduce the
46 risk of chronic diseases and to promote human health has aroused increasing scientific and
47 commercial interest over the past decade [5].

48 High blood pressure or hypertension, which is estimated to affect one third of the Western
49 population [6], is a risk factor for cardiovascular diseases including coronary heart disease,
50 peripheral artery disease and stroke. In view of its high prevalence and importance, changes in
51 life-style, dietary approaches and pharmacological treatments are broadly applied to treat
52 hypertension. It has been recognized that nutritional factors play a significant role in the
53 prevention and/or treatment of hypertension, and therefore, efforts are being put into the
54 production of foods with antihypertensive activity. Angiotensin-converting enzyme (ACE, EC
55 3.4.15.1) is one of the main regulators of blood pressure through its action on two body systems.

56 Firstly, ACE forms part of the rennin-angiotensin system (RAS), converting angiotensin I to a
57 potent vasoconstrictor, angiotensin II, which also induces the release of aldosterone and therefore,
58 increases the sodium concentration and blood pressure. ACE also takes part of the kinin-kallicrein
59 system as it hydrolyzes bradykinin, which has a vasodilator action. By inhibiting this enzyme,
60 bioactive peptides have been shown to lower blood pressure in animal and clinical studies. First
61 ACE inhibitors were discovered in snake venom. Currently, different ACE inhibitors, such as
62 Captopril and Enalapril, are being extensively used to treat essential hypertension. However, their
63 undesirable effects, such as hypotension, cough, increased potassium levels, reduced renal
64 function, angioedema, etc. [7], have promoted the search of ACE inhibitory peptides derived
65 from food natural sources. To date, milk from different species is the main source of ACE
66 inhibitory peptides [3, 8, 9]. Other animal protein sources of these peptides are muscle [10],
67 ovalbumin [11], blood [12], and fish proteins [13, 14]. Plant protein sources include, among
68 others, pea [15], garlic [16], rice [17], soybean [18, 19], wheat [20], and Amaranth proteins [21].

69 This article reviews current literature on the subject of ACE inhibitory and
70 antihypertensive peptides, their structure-activity relationship, mechanism of action and
71 bioavailability. Evaluation of their activity in humans as well as their possible incorporation into
72 food products will be also covered.

73

74 **2. Release and identification of antihypertensive peptides**

75 Biologically active peptides can be released from their parent protein by enzymatic
76 hydrolysis during gastrointestinal digestion, fermentation or maturation during food processing or
77 proteolysis by food-grade enzymes derived from microorganisms, animals or plants [22]. If the
78 peptidic sequence is known, it is also possible to synthesize the peptide by chemical or enzymatic
79 synthesis or by recombinant DNA technology [23].

80 2.1. *Gastrointestinal digestion*

81 It has been recognized that dietary proteins and peptides are susceptible to hydrolysis
82 during the different stages of gastrointestinal digestion, namely ingestion, digestion and
83 absorption [24]. Once ingested, these proteins and peptides are subjected to hydrolysis by
84 different enzymes present in the gastrointestinal tract such as pepsin, trypsin, chymotrypsin and
85 peptidases at the surface of epithelial cells to release peptides of various lengths. Some of these
86 peptides may exert a direct function at the gastrointestinal tract. However, other peptides can be
87 absorbed to reach target organs and tissues through systemic circulation [25].

88 In order to examine the effect of gastrointestinal proteases on the release of and
89 breakdown of ACE inhibitory peptides from food proteins, simulated gastrointestinal digestion
90 processes have been carried out on various protein sources, such as milk proteins (Table 1) [26-
91 30], egg proteins [11], meat proteins [31, 32], fish proteins [33, 34], as well as vegetal proteins
92 [21, 35-37]. As an example, Hernandez-Ledesma et al., [28] identified peptides with ACE
93 inhibitory and antioxidant activity in hydrolyzates of several samples of human milk and infant
94 formulas after digestion with pepsin and pancreatin simulating infant gastrointestinal conditions.
95 Recently, Majumder and Wu [38] have studied the effect of simulated gastrointestinal digestion
96 of cooked eggs on the release of ACE inhibitory peptides. These authors found that fried egg
97 digests showed more potent ACE inhibitory activity than boiled egg digests, and postulated that
98 the lower protein denaturation in boiled eggs may results in a lower protein digestibility.

99

100 2.2. *Fermentation and maturation process*

101 During fermentation process, lactic acid bacteria (LAB) hydrolyze milk proteins, mainly
102 caseins, into peptides and amino acids which are used as nitrogen sources necessary for their
103 growth [39]. Hence, bioactive peptides can be generated by starter and non-starter bacteria used

104 in the manufacture of fermented dairy products (Table 1) [40-43]. Many of these peptides have
105 been reported to exert ACE inhibitory and antihypertensive properties. Proteolytic system of
106 *Lactobacillus helveticus*, *Lactobacillus delbrueckii* ssp. *bulgaricus*, *Lactococcus lactis* ssp.
107 *diacetylactis*, *Lactococcus lactis* ssp. *cremoris*, and *Streptococcus salivarius* ssp. *thermophilus*
108 strains have been demonstrated to hydrolyze milk proteins releasing ACE inhibitory peptides
109 [reviews 3, 43, 44]. Some of the peptides identified also have been shown to lower blood
110 pressure in hypertensive rats [45-47], and humans [48, 49]. The best characterized ACE
111 inhibitory peptides are VPP and IPP found in milk fermented with *Lactobacillus helveticus* and
112 commercialized in Japan (Calpis, Calpis Co. Ltd., Tokyo, Japan) and Finland (Valio Evolus
113 Double Effect, Valio Ltd., Finland). This fermented milk has shown beneficial effects on blood
114 pressure in several rat models and human studies [47, 48, 50-55].

115 Ashar and Chand [42] identified an ACE-inhibitory peptide from milk fermented with
116 *Lactobacillus delbrueckii* ssp. *bulgaricus*, and Pihlanto et al. [56] reported two peptides
117 responsible for the ACE inhibitory activity of milk fermented with *Lactobacillus jensenii*. In
118 combination with *Streptococcus salivarius* ssp. *thermophilus* and *Lactococcus lactis* biovar.
119 *diacetylactis*, a hypotensive structure with a sequence of SKVYP was obtained from β -casein.
120 Quirós and co-workers [57] identified two peptides in fermented milk with *Enterococcus faecalis*
121 that corresponded to β -CN fragments LHLPLP and LVYFPFGPIPNLQNIIP, with potent
122 ACE-inhibitory activity and proven antihypertensive effect when orally administered to
123 spontaneously hypertensive rats after acute and long-term administration.

124 During the maturation of cheese, the major milk proteins are degraded into a large number
125 of peptides due to the action of endogenous milk enzymes, added coagulants and microbial
126 enzymes. A number of studies have shown that ACE inhibitory peptides can be produced during

127 cheese making, in particular during ripening process. These peptides have been characterized in
128 different commercial cheeses, such as Edam, Gouda, Camembert, Havarti and Blue cheese [58],
129 Italian and Spanish cheeses [29, 59, 60], and Asiago cheeses [30]. Tri-peptides VPP and IPP have
130 also been identified and quantified in different cheese varieties by Butikofer and co-workers [61,
131 62]. A low-fat cheese containing ACE-inhibitory peptides derived from α_{s1} -casein and named as
132 “Festivo” is commercialized in Eastern countries.

133 Fermented soy products, traditionally consumed in Eastern countries, have been also
134 found to be an important source of ACE-inhibitory and antihypertensive peptides. A potent
135 antihypertensive peptide has been identified and characterized in a Korean soy product
136 denominated “chunggugjang” and obtained by soy fermentation with *Bacillus subtilis* CH-1023
137 [63]. Other ACE-inhibitory and antihypertensive peptides have been identified in soy paste [64],
138 soy sauce [65, 66], natto and tempeh [67], and other fermented soy products [19, 68, 69].

139

140 2.3. *Enzymatic hydrolysis*

141 The most common way to produce bioactive peptides is through enzymatic hydrolysis of
142 whole protein molecules (Table 1). A large number of studies have demonstrated the release of
143 ACE inhibitory and/or antihypertensive peptides from food proteins, by hydrolysis with
144 gastrointestinal enzymes, such as pepsin, trypsin, and chymotrypsin [24, 70-72]. Manso and
145 Lopez-Fandino [73] described occurrence of ACE-inhibitory peptides in hydrolyzates of bovine,
146 ovine and caprine κ -casein upon hydrolysis with various digestive enzymes. Pepsin was used by
147 Contreras and co-workers [74] to hydrolyze total isoelectric casein and three peptide sequences
148 derived from α_{s1} -casein and α_{s2} -casein were characterized as ACE inhibitors and antihypertensive
149 peptides.

150 In addition to live microorganisms, proteolytic enzymes from bacterial and fungal sources
151 have been used to generate bioactive peptides from various proteins. The use of commercially
152 available microbial-derived food grade proteinases to hydrolyze food proteins is advantageous as
153 these enzymes are low-cost and safe, and the product yields are very high [75]. Ueno et al. [76]
154 purified and characterized an endopeptidase from *Lb. helveticus* CM4 and demonstrated that this
155 peptidase can generate antihypertensive peptides using synthetic pro-peptides as substrates.
156 Mizuno et al. [77] measured the ACE-inhibitory activity of casein hydrolyzates upon treatment
157 with nine different commercially available proteolytic enzymes. Among these enzymes, a
158 protease extracted from *Aspergillus oryzae* acted specifically on casein to release VPP and IPP,
159 and the obtained casein hydrolyzate demonstrated a significant dose-dependent antihypertensive
160 effect in a rat model with spontaneously hypertensive rats.

161 Recently, the interest of food technologists has turned to the use of different techniques,
162 such as high-pressure and heat denaturing and power ultrasound to modify protein structure and
163 increase enzymatic hydrolysis. As compared to the proteolysis at atmospheric pressure,
164 qualitative and quantitative differences were detected in the hydrolysis pattern when proteolysis
165 with trypsin was carried out under high pressure treatments [78-79]. Hernandez-Ledesma et al.
166 [80] reported that heating of β -Lg during enzyme treatments with thermolysin enhances the
167 formation of peptides with ACE-inhibitory activity, and one of the peptides released under these
168 heat-denaturing conditions was LQKW that had previously been described as a potent ACE
169 inhibitor [81]. Prolonged exposure to high-intensity ultrasound has been shown to inhibit the
170 catalytic activity of a number of food enzymes [82]. However, in some cases, solutions
171 containing enzymes have been found to have increased activity following short exposures to
172 ultrasound [83]. Jia et al. [84] found that the use of ultrasonic treatment during proteolysis could

173 facilitate the enzymatic hydrolysis of peptide DWGP, whereas ultrasonic pre-treatment could
174 promote the release of ACE inhibitory peptides from this peptide.

175

176 *2.4. Genetic recombination in bacteria*

177 Industrial preparation of ACE inhibitory and/or antihypertensive peptides by enzymatic
178 hydrolysis and microbial fermentation showed to be a low efficient process because of low yield
179 and high cost of separation and purification processes [85-88]. To solve these issues, during last
180 years a new technique based on genetic engineering is being developed. One of the challenges of
181 this technique is the susceptibility of short antihypertensive peptides to degradation by protease
182 or peptidase. Moreover, the expression products may be harmful to the host, impacting the high-
183 level expression of the gene. This shortcoming has been conquered by expression of
184 antihypertensive peptides in the forms of a fusion protein or a tandem gene. Antihypertensive
185 peptides with sequences HHL, HVLPVP, FFVAPFPEVFGK, and GHIATFQER have been
186 expressed successfully in *Escherichia coli* [89-92], although special proteases are needed to
187 release the target active protein, thus increasing the cost of separation and purification after
188 enzymatic hydrolysis. Recently, Rao et al. [93] expressed an antihypertensive peptide multimer, a
189 common precursor of 11 kinds of antihypertensive peptides, and the release was confirmed by
190 simulated gastrointestinal digestion. Because of the fact that currently genetic modified
191 microorganisms are difficult to be used in food products, further studies should be needed.

192

193 **3. *In vitro* and *in vivo* assays**

194 The search for ACE inhibitory activity is the most common strategy followed in the
195 selection of antihypertensive peptides derived from food proteins. In order to study ACE
196 inhibition, simple, rapid, sensitive and reliable analytical methods are desirable. *In vitro*

197 inhibitory activity is generally measured by monitoring the conversion of an appropriate substrate
198 by ACE in the presence and absence of the potential inhibitors. There are several methods, but
199 those based on spectrophotometric and high-performance liquid chromatography (HPLC) assays
200 are most commonly utilized. The spectrophotometric method of Cushman and Cheung [94] is
201 based on the hydrolysis of Hippuryl-His-Leu (HHL) by ACE to hippuric acid and His-Leu, and
202 the extent of hippuric acid released is measured after its extraction with ethyl acetate. The
203 inhibitory potency is expressed as the IC_{50} value, or concentration needed to inhibit 50% of the
204 enzyme activity. Extraction of reaction product is tedious and may overestimate ACE activity if
205 unhydrolyzed HHL is also extracted. Another broadly used spectrophotometric method is based
206 in the hydrolysis of a furanocryloyl tripeptide (FAPGG) to FAP and the di-peptides GG [95].
207 Using HPLC methods, the peak of hippuric acid may be interfered with by the added ACE
208 inhibitors in the reaction mixture, so the mobile phase needs to be adjusted to different tested
209 compounds. Moreover, the HPLC method show lower detection sensitivity and longer analysis
210 time to obtain good results. Doig and Smiley [96] and Mehanna and Dowling [97] have improved
211 HPLC methods by applying ultraviolet detection, and Van Elswijk et al. [98] have developed an
212 alternative strategy for the screening of complex food samples applying an HPLC method with
213 biochemical detection. In that approach, separation and activity detection are combined within
214 one step. However, as spectroscopic detection is used to monitor the enzymatic conversion, these
215 methods are restricted to artificial substrates as well. Direct, extraction-free methods have been
216 published recently [99, 100]. Siemerink et al. [101] have optimized a new robust HPLC coupled
217 with electrospray ionization mass spectrometry (HPLC/ESI-MS)-based screening method for
218 ACE-inhibiting substances in crude samples. Similarly, an ultra-performance-liquid
219 chromatography (UPLC) coupled with MS (UPLC-MS) for determination of the ACE activity
220 has been recently developed by Geng et al. [102]. This new method is more sensitive, accurate

221 and reproducible. The small total reaction volume, the short analysis time, high selectivity and
222 lower expense are the advantages of this method in comparison with the conventional methods.

223 The antihypertensive effects can only be reliably assessed by *in vivo* experiments using
224 spontaneously hypertensive rats that constitute an accepted model for human essential
225 hypertension [7]. A great number of studies have addressed the effects of both short-term and
226 long-term administration of potential antihypertensive peptides using this animal model [88, 103-
227 106]. Moreover, many *in vivo* studies include the evaluation of the effect of antihypertensive
228 peptides on arterial blood pressure of normotensive Wistar-Kyoto rats. Recently, Nakahara et al.
229 [66] have used the Dahl salt-sensitive rats as a model of salt-sensitive hypertension to evaluate
230 the antihypertensive effect of a peptide-enriched soy-sauce like seasoning. The results of these
231 tests have highlighted an important lack of correlation between the *in vitro* ACE inhibitory
232 activity and the *in vivo* action. This fact has provided doubts on the use of the *in vitro* ACE
233 inhibitory activity as the exclusive criteria for potential antihypertensive substances because other
234 mechanisms of action than ACE inhibition might be responsible for the antihypertensive effect.
235 Also, it should be needed to take into consideration the physiological transformations
236 determining the bioavailability of the peptides.

237 The antihypertensive effect of some food proteins-derived peptides has been conducted in
238 human studies to determine whether these peptides possess an antihypertensive effect on human
239 subjects with high-normal blood pressure and mild hypertension [50, 107, 108]. The most
240 substantiated antihypertensive activity in humans has been obtained for the commercial
241 fermented milk products and hydrolyzates containing the ACE-inhibitory peptides IPP and VPP.
242 The antihypertensive effect of the sour milk product Calpis, commercialized in Japan, was tested
243 in a clinical study with mildly hypertensive patients [50]. Recently, a study has been conducted
244 among patients with high-normal blood pressure and mild hypertension, evaluating the effect of

245 different doses of a casein hydrolyzate produced by *Aspergillus oryzae* containing IPP and VPP
246 and commercialized as AmealPeptide[®] by Calpis [77]. Similarly, a milk product Evolus[®]
247 fermented with *Lactobacillus helveticus* LBK-16H and produced by Valio Ltd. (Finland) has
248 been tested in hypertensive humans [48]. This product, containing peptides IPP and VPP, showed
249 to exert a long-term blood pressure-lowering effect after normal daily ingestion during a 21-
250 weeks intervention period.

251

252 **4. Structure-activity relationship**

253 Although the structure-activity relationship of ACE-inhibitory peptides derived from
254 foods has not yet been fully elucidated, several structural features influencing potency of these
255 peptides have been identified [7, 109]. Recently, it has been reported that artificial neural
256 networks (ANN) and quantitative structure-activity relationship (QSAR) modelling may be used
257 to develop statistical computer models potentially capable of identifying ACE inhibitory peptides
258 based on structure-activity data [110]. Several descriptor variables such as molecular mass and
259 shape, hydrophobicity, charge and electronic properties have been recognized as critical in this
260 QSAR modelling. The majority of ACE inhibitory peptides are relatively short sequences
261 containing from 2 to 12 amino acids. This in agreement with the results of Natesh and coworkers
262 [111], which demonstrated from crystallography studies, that the active site of ACE cannot
263 accommodate large peptide molecules. However, some studies have identified ACE inhibitory
264 peptides with up to 27 amino acids [58, 112, 113]. Of many ACE-inhibitory peptides identified
265 from different food sources, structure-activity correlation indicated that C-terminal tri-peptide
266 residues play a predominant role in competitive binding to the active site of ACE. It has been
267 reported that this enzyme prefers substrates or inhibitors containing hydrophobic (aromatic or
268 branched side chains) amino acid residues at each of the three C-terminal positions. The most

269 effective ACE inhibitory peptides identified contain Tyr, Phe, Trp, and/or Pro at the C-terminal.
270 Gomez-Ruiz and coworkers [29] have suggested that amino acid Leu may contribute
271 significantly to increase ACE inhibitory potential. Furthermore, other branched chain aliphatic
272 amino acids such as Ile and Val are predominant in highly peptide inhibitors. In addition,
273 structure-activity data suggest that the positive charge of Lys (ϵ -amino group) and Arg (guanidine
274 group) as the C-terminal residue may contribute to the inhibitory potency [114-117]. Other
275 characteristics have also been found to play important roles for ACE inhibition. It has been
276 recognized that ACE inhibitory peptides possess a characteristic pattern (i.e. a similar positive
277 potential located at the C-terminal end) different from that of inactive peptide molecules [112,
278 118]. For long chain peptides, it is expected that peptide conformation, i.e. the structure adopted
279 in the specific environment of the binding site, will influence binding to ACE [117, 118]. It has
280 also been demonstrated that ACE has a requirement for the L-configuration of the amino acid at
281 position three from the C-terminal. Moreover, changes in cis-trans conformations of Pro at the C-
282 terminal position of an ACE inhibitory peptide may cause significant changes in its interaction
283 with the enzyme [119].

284

285 **5. Mechanism of action**

286 Blood pressure is determined by cardiac output and vascular peripheral resistance, and is
287 regulated by a complex system involving the RAS, the sympathetic nervous system (SNS), and
288 the kidney and fluid balance mechanism [118]. Most food-derived peptides usually display
289 higher *in vivo* activities than the efficacy levels extrapolated from the *in vitro* ACE inhibitory
290 activity. This may be an indication of the existence of an additional mode of action [120]. In fact,
291 increasing evidence is being provided that different mechanisms, others than ACE inhibition, are

292 involved in the blood-pressure-modulating effect exerted by many of these peptides. *In vitro*, tri-
293 peptides VPP and IPP have been shown to inhibit ACE at micromolar concentrations [40, 41]. *In*
294 *vivo*, long-term treatment of spontaneously hypertensive rats with fermented milk containing
295 these peptides has been found to decrease serum ACE activity [121-123]. However, according to
296 Jauhiainen et al. [54], the mechanistic theory of ACE inhibition of IPP and VPP remains to be
297 confirmed and other effects have to be taken into consideration. Some of these effects have been
298 evaluated in animal models and clinical studies. Plasma rennin activity and levels have been
299 found to be raised in spontaneously hypertensive rats receiving IPP and VPP for 14 weeks. This
300 raising can be due to the lack of negative feedback by angiotensin II, which supports that ACE
301 was inhibited [47]. Other authors have reported the protective effects exerted by these peptides on
302 endothelial function of isolated mesenteric arteries of rats after 24 h incubation with them [124].
303 In humans with mild hypertension, administration of a casein hydrolyzate containing VPP and
304 IPP increases maximum blood flow forearm during reactive hyperemia, thus demonstrating an
305 improvement in the vascular endothelial dysfunction. Yamaguchi et al. [125] studied effect of a
306 5-day repeated administration of VPP and IPP on gene expression of spontaneously hypertensive
307 rats abdominal aorta using DNA microarray microanalysis, reporting a significant increase for the
308 endothelial nitric oxide synthase (eNOS) gene and the connexin 40 gene, which are involved in
309 blood pressure regulation. Expression of these genes was restored in the aortic tissue after
310 treatment with these tri-peptides [126, 127], suggesting that VPP and IPP might act *in vivo* as
311 ACE inhibitors in the aorta and also have preventive potential in cardiovascular function.

312 Fuglsang et al. [46] reported that ingestion of two milks fermented with *Lactobacillus*
313 *helveticus* provokes a decrease of the response to an intravenous injection of angiotensin I in
314 unconscious normotensive rats, whereas response to bradikinin was increased, confirming the
315 inactivation of ACE. Dried bonito-ACE inhibitory peptides slightly inhibit angiotensin I-induced

316 contractions in rat-isolated aorta as compared with Captopril, but unlike this drug, peptides exert
317 a direct action on vascular smooth muscles [128]. Similarly, peptide lactokinin (ALPMHIR)
318 inhibits the release of ET-1, an endothelial factor that evokes contractions in smooth muscle cells
319 through mechanisms both dependent and independent of ACE-inhibition [129]. It is also likely
320 that opioid receptors are involved in the antihypertensive effect of some peptides, such as α -
321 lactorphin, β -lactorphin and human casein-derived fragments, as this was abolished by the opioid
322 receptor antagonist naloxone. As an example, it has been demonstrated that α -lactorphin, a tetra-
323 peptide (YGLF) formed by *in vitro* proteolysis of α -lactalbumin with pepsin and trypsin, lowers
324 blood pressure in spontaneously hypertensive rats and produces an endothelium-dependent
325 relaxation of their mesenteric arteries that is inhibited by an eNOS inhibitor [47]. Therefore, a
326 mechanism of action driven by the stimulation of peripheral opioid receptors and subsequent
327 nitric oxide (NO) release causing vasodilation has been proposed for this peptide. Although α -
328 lactorphin interacting with opioid receptors does not elicit effects typical of centrally active
329 opioids such as antinociception and sedation [130]. It has been suggested that these opioid
330 peptides might lower blood pressure through receptors expressed in the gastrointestinal tract,
331 which implies that no absorption is required [131].

332 Strong epidemiological evidence indicates that oxidative stress and associated oxidative
333 damage are mediators in cardiovascular diseases. In experimental and human hypertension
334 studies, it has been demonstrated an increased production of superoxide anion and hydrogen
335 peroxide, reduced NO synthesis, and decreased bioavailability of antioxidants [132]. Therefore,
336 food-derived peptides with antioxidant properties might also have effect on blood pressure
337 modulation. Many of these peptides have been identified and characterized from casein and whey
338 proteins hydrolyzed with different enzymes [133, 134].

339 Taking together the results of all these studies, more thorough mechanistic research
340 should be probably needed to detect the changes in the factors affecting blood pressure and
341 vascular tone to show the exact mechanisms also *in vivo* of antihypertensive peptides.

342

343 **6. Bioavailability and clinical studies**

344 The physiological effects of bioactive peptides depend on the ability to reach in an active
345 form their target organs. This implies resistance to gastrointestinal enzymes and brush border
346 membrane peptidases and absorption through the intestinal epithelium. The resistance of peptides
347 to these processes is usually performed by sequential hydrolysis with pepsin and pancreatic
348 extracts mimicking the gastrointestinal conditions and with *in vitro* studies with epithelial
349 intestinal cells. Although peptides were thought to be rapidly metabolized to constituent amino
350 acids, these studies have demonstrated that several peptides are resistant to these physiological
351 processes and can reach the circulation. This is the case of the short tri-peptides IPP and VPP
352 [135, 136], but also has been demonstrated for longer proline-rich peptides, such as, LHLPLP.
353 This latter peptide resists simulated gastrointestinal digestion but it is hydrolysed to a shorter
354 active form, HLPLP, by cellular peptidases prior to transport across the intestinal epithelium [137,
355 138]. Figure 1 shows the formation of the shorter peptide during incubation of the peptide
356 LHLPLP in the apical chamber of the Caco-2 cell culture. The penta-peptide appeared in
357 approximately 3 min and its concentration increased with the incubation time up to 60 min. This
358 shorter form has also been detected in human plasma after oral administration which
359 demonstrates intestinal absorption of the pentapeptide in humans [139]. In some cases, the active
360 form is released during gastrointestinal processes. For instance, the active form of peptide
361 KVLVPVQ is generated by hydrolysis of the glutamine residue at the C-terminal end during
362 pancreatic digestion [140]. This is also the case of the egg-derived antihypertensive peptides

363 YAEERYPIL and RADHPFL that were hydrolysed to other active forms after simulated
364 gastrointestinal digestion [11]. The pharmacokinetic behaviour of the tri-peptides IPP and VPP
365 has also been studied and an absolute bioavailability of 0.1% respect the administered dose has
366 been calculated in pigs [141]. In humans, these two tri-peptides were detected in plasma after oral
367 administration at picomolar concentrations and their absorption was enhanced when ingested in
368 the form of an enriched yogurt beverage. In addition, a further increase (1.2-fold) in the plasmatic
369 concentration of IPP was found when the enriched yogurt was administered after a meal [142].

370 Several clinical studies have evaluated the antihypertensive effect of the tri-peptides IPP
371 and VPP after long-term administration in humans. Most of them are included in two meta-
372 analyses recently published [143, 144]. The meta-analysis by Xu et al. [143] includes 12 trials
373 with a total of 623 participants and found significant decreases in systolic and diastolic blood
374 pressure (4.8 mmHg and 2.2 mmHg, respectively). Similar results arose from the meta-analysis
375 published by Pripp et al. [144] with a total of 15 clinical trials included. Although two long-term
376 studies have not found statistical differences with these tri-peptides [145, 146], most recent
377 reviews on the subject identify several factors such as component of the final product, dose,
378 method for blood pressure measurement that can influence the results in different trials [147, 148].

379

380 **7. Incorporation into food products**

381 For an industrial application of protein hydrolyzates containing antihypertensive peptides,
382 main considerations would be the organoleptic characteristics of these ingredients and the
383 evaluation of the resistance of the active peptides to processing conditions. The practical use of
384 protein hydrolyzates in food systems is hindered due to the presence of low molecular weight
385 peptides composed mainly of hydrophobic amino acids that results in a bitter taste [149]. In fact,
386 this problem has limited the use of some of the developed hydrolyzates with proved

387 antihypertensive effect. In addition to the bitterness, the pH of the hydrolysis reaction needs to be
388 regulated because the substrate susceptibility and the enzyme activity are strongly influenced by
389 the pH. In order to achieve the desired hydrolysis degree to obtain biologically active peptides,
390 the addition of some alkali or acid is required to neutralize the hydrolysis products. This leads to
391 undesirable high ash build up in the hydrolyzates and the development of salty off-flavors.
392 Different strategies have been applied for debittering protein hydrolyzates. These include
393 absorption of bitter peptides on activated carbon, chromatographic removal using different
394 matrices and selective extraction with alcohols [149]. The most extended approaches include
395 hydrolysis of bitter peptides with enzymes such as aminopeptidase, alkaline/neutral protease and
396 carboxypeptidase, condensation reactions of bitter peptides using protease and use of
397 *Lactobacillus* as debittering starter adjunct [150]. However, the application of all these methods
398 in biologically active hydrolyzates is limited because the enzymatic activity used for debittering
399 can hydrolyze the previously generated bioactive peptides. Although it has not been found a
400 significant correlation between bitterness and the ACE-inhibitory activity of di- and tri-peptides
401 [151], it is recognized the importance of hydrophobic amino acid residues for both bitterness and
402 ACE-inhibitory peptides. Therefore, those methods based in the removal of bitter peptides (either
403 absorption, extraction or hydrolysis) have to be carefully applied to hydrolyzates containing, for
404 instance, ACE-inhibitory peptides. For bioactive hydrolyzates, the application of masking
405 methods by using monosodium glutamate or glutamylglutamic acid [152], the addition of
406 cyclodextrins [153], encapsulation [154], or the addition of phospholipids and lysophospholipids
407 [155] could be preferably used. For instance, for the encapsulation of casein hydrolyzates,
408 different materials such as soy proteins isolates alone or in mixtures with gelatin [154, 156],
409 maltodextrins [157], or lipospheres [158] have been successfully applied. The presence of
410 proteins and reducing carbohydrates in a food ingredient will lead to the formation of Maillard

411 compounds that can have a positive effect on flavor but this reaction has to be controlled to avoid
412 the generation of undesirable compounds [159, 160]. Therefore, the improvement of the flavor of
413 biologically active protein hydrolyzates including possible protein-flavor interactions and the
414 effect of these processes on biological activity are of interest in relation to the development of
415 novel protein foods.

416 Processing provides an additional value to foods in improving their safety, shelf-life,
417 palatability, nutritive and functional value, but the conditions of processing and storage may be
418 detrimental to peptides. At this regard, changes in the molecular structure of an amino acid may
419 lead to changes in the bioactivity and in the absorption of the peptide of interest. The most
420 relevant degradation pathways of amino acids during processing were recently reviewed by
421 López-Fandiño & co-workers [3]. For example, thermal processing favors racemization, amino
422 acids decomposition (e.g. ornithine from arginine), glycation (non-enzymatic browning or
423 Maillard reaction), and cross-linking. Furthermore, amino acids can be also oxidized during food
424 processing [161], and even *D*-amino acids can be synthesized out from *L*-amino acids by LAB.
425 Dehydration by spray-drying may produce some negatively effects on food protein
426 hydrolyzates, such as changes in peptide composition, reduction of amino acid content and non
427 enzymatic browning [162-164]. There are little data about the effects of other processes on
428 bioactive peptides. Recently, the stability of two α_{s1} -casein-derived antihypertensive peptides to
429 spray-drying, homogenization and pasteurization when they were incorporated into fermented
430 milk has been demonstrated [165]. Incorporation of active hydrolyzates to fermented milks
431 implies that these peptides have to survive in the presence of LAB because of their cell-
432 associated proteinases/peptidases systems that could further hydrolyze the bioactive sequences.
433 For instance, it has been reported that an 12 amino acid long antihypertensive peptide can be

434 digested by exposure to yogurt starter culture strains and therefore, the incorporation of this
435 bioactive peptide should preferably done at the end of the yogurt-making process [166].

436

437 **8. Future prospects**

438 Among the different groups of bioactive peptides defined, antihypertensive peptides have
439 received special attention, their activity has been tested *in vitro*, animal models and humans, and
440 they have been incorporated into different food products. Controversial results on clinical studies
441 and the different health claim legislations will contribute to increase research in this area. In this
442 sense, different studies have been performed to demonstrate stability of the peptide, absorption
443 and identification of the active form in the organism. It has been postulated that physiologically
444 relevant concentrations and elimination kinetics will be mandatory for food derived bioactive
445 components as it is now for pharmaceutical compounds. At the same time, the recent advances on
446 specific analytical techniques able to follow small amounts of the peptides or derivatives from
447 them in complex matrices and biological fluids will allow performing these kinetic studies in
448 model animals and humans. Similarly, advances in new disciplines such as nutrigenomic and
449 nutrigenetic will open new ways to follow bioactivity in the organism by identifying novel and
450 more complex biomarkers of exposure and/or of activity. All these advances will be done
451 simultaneously with the knowledge food technologists since the final formulation of the food
452 product is crucial to ensure activity and bioavailability of bioactive peptides.

453

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459 **Figure captions**

460 Figure 1: Changes of the synthetic peptide LHLPLP when added to apical chamber of a Caco-2
461 cell culture at different periods of time. Extracted ion chromatograms obtained by HPLC-MS
462 analysis of the apical chamber after a) 5 min, b) 10 min, c) 30 min, and d) 60 min of incubation.
463 The extracted ion chromatogram was obtained by extraction of ions with m/z 689.4, 711.4 and
464 727.4, which correspond to molecular ion of peptide LHLPLP and its sodium and potassium
465 adducts, and ions with m/z 598.3, 614.3 and 620.3, corresponding to molecular ion of peptide
466 HLPLP and its sodium and potassium adducts. Reproduced from [137] with permission.

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Table 1. ACE inhibitory and antihypertensive activity in spontaneously hypertensive rats of peptide derived from caseins and whey proteins by fermentation and enzymatic hydrolysis

Peptide	Sequence	IC ₅₀ (μM) ^a	Decrease of SBP (mmHg) ^b	Origin	Reference
α _{s1} -CN f(1-9)	RPKHPIKHQ	13.4	-9.3	Gouda cheese	[58]
α _{s1} -CN f(146-147)	YP	720	-32.1	Fermentation with <i>Lb. helveticus</i> CPN4	[140]
β-CN f(58-76)	LVYFPFGPIPNSLPQ NIPP	5.2	-15.0	Fermentation with <i>En. faecalis</i>	[11, 57]
β-CN f(60-68)	YFPFGPIP	14.8	-7.0	Gouda cheese	[58]
β-CN f(74-76)	IPP	5.0	-28.3 (-10.1) ^d	Fermentation with <i>Lb. helveticus</i> y <i>Sc. cerevisiae</i>	[40, 41]
β-CN f(84-86)	VPP	9.0	-32.1 (-10.1) ^d	Fermentation with <i>Lb. helveticus</i> y <i>Sc. cerevisiae</i>	[40, 41]
β-CN f(133-138)	LHLPLP	5.5	-21.9	Fermentation with <i>En. faecalis</i>	[11, 57]
β-CN f(133-139)	LHLPLPL	425	-7.7	Fermentation with <i>En. faecalis</i>	[11, 57]
β-CN f(197-206)	VLGPVRGPF	137	-16.2	Fermentation with <i>En. faecalis</i>	[11, 57]
β-CN f(201-209)	VRGPFPIIV	599	-16.1	Fermentation with <i>En. faecalis</i>	[11, 57]
α _{s1} -CN f(23-34)	FFVAPFPGVFGK	77	-34.0	Hydrolysis with trypsin	[167]
α _{s1} -CN f(104-109)	YKVPQL	22	-13.0	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]
α _{s1} -CN f(194-199)	TTMPLW	16	-13.6	Hydrolysis with trypsin	[167]
α _{s2} -CN f(189-192)	AMPKPW	580	-5.0	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]
α _{s2} -CN f(190-197)	MKPWIQPK	300	-3.0	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]
α _{s2} -CN f(198-202)	TKVIP	400	-9.0	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]
α _{s2} -CN f(203-208) ^d	PYVRYL	1.9	23.4	Hydrolysis with pepsin	[168]
β-CN f(59-61)	VYP	288	-21.0	Hydrolysis with proteinase K	[169]
β-CN f(59-64)	VYFPFG	221	-22.0	Hydrolysis with proteinase K	[169]
β-CN f(80-90)	TPVVVPPFLQP	749	-8.0	Hydrolysis with proteinase K	[169]
β-CN f(140-143)	LQSW	500	-2.0	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]
β-CN f(169-174)	KVLPVP	5	-32.2	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]
β-CN f(169-175)	KVLPVPQ	1000	-31.5	Hydrolysis with a proteinase from <i>Lb. helveticus</i> CP790	[140]

925

926 **Table 1. (Continuation).** ACE inhibitory and antihypertensive activity in spontaneously hypertensive rats of peptide derived from caseins and whey proteins
 927 by fermentation and enzymatic hydrolysis
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 929

Peptide	Sequence	IC ₅₀ (μM) ^a	Decrease of SBP (mmHg) ^b	Origin	Referente
β-CN f(177-183)	AVPYPQR	15	-10.0	Hydrolysis with trypsin	[167]
α-La f(50-53)	YGLF	733	-23.0	Hydrolysis with gastric and pancreatic enzymes	[170]
β-Lg f(58-61) ^c	LQKW	34.7	-18.1	Hydrolysis with thermolysin	[103]
β-Lg f(78-80)	IPA	141	-31.0	Hydrolysis with proteinase K	[169]
β-Lg f(103-105) ^e	LLF	79.8	-29.0	Hydrolysis with thermolysin	[104]
BSA f(221-222)	FP	315	-27.0	Hydrolysis with proteinase K	[169]

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 931
 932 ^a: Peptide concentration needed to inhibit 50% ACE activity

933 ^b: Systolic blood pressure

934 ^c: Antihypertensive effects in humans

935 ^d: Ovine protein

936 ^e: Caprine protein
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