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#### Abstract

The worldwide consumption of high-protein food has notoriously increased in recent years. Meat industry generates substantial quantities of protein-rich raw material, which are often discarded as low-value by-products. However, several bioactive compounds can be isolated from these products giving an added value to them. In addition to conventional extraction methods, emerging technologies, including high pressure processing (HPP), ultrasound (US), pulsed electric fields (PEF) can be used for peptide isolation from meat by-products allowing to maintain the functional properties of these compounds. Antioxidant and antimicrobial activities are between the properties associated with peptides, what would enable their introduction in foods as ingredient and preservative. This review is focused to gather accurate information about the entire extraction process, from the source used until the final peptides obtained. In this sequence are included the pretreatment of the by-product, the extraction procedure, the fractionation and purification, as well as the assay used for the determination of their antioxidant and antimicrobial activities. 

 Keywords: Bioactive; Co-products; Enzymatic hydrolysis; Meat industry; Valorization

# Meat industry and by-products

Meat industry is one of the most important industrial sectors in the world, producing around of 330 million of tons. Europe provides approximately one fifth of the world's production, being Germany, Spain, France and Italy the countries with the largest population of primary livestock (FAOSTAT 2016). This great production gives an idea of the huge amounts of disposal generated each day in processing meats and slaughterhouses (Toldrá et al. 2012), since not all the parts of animals are suitable and safe to human consumption. It is estimated that more than half of live animal weight is not appropriate for human consumption (Irshad and Sharma 2015). Europe together with Asia are the two largest consumers of meat by-products, especially those that have as source beef and lamb (Liu and Ockerman 2001). Furthermore, the discard of these animal by-products may influence unfavorably the environment because of the increase in chemical (COD) and biochemical oxygen demand (BOD) (O'Sullivan et al. 2017; Bah et al. 2015).

Several studies have aimed to study meat by-products since a great number of bioactive compounds can be obtained from them (Ryder et al. 2016). These by-products have a high nutritional content, with proteins between the nutrients that could have functional properties (Matak et al. 2015; Toldrá et al. 2016; Pateiro et al. 2019). Bioactive peptides usually contain sequences of 2 to 20 amino acids, among them are included essential amino acids with high bioavailability that humans cannot synthesize, like hydroxymethyllysine and methylhistidine, which are not usually present in plant protein (Arihara et al. 2006; Wang et al. 2008; Vongsawasdi et al. 2014; Siti et al. 2016). Their amino acid content is going to determine the biological activity of the peptides (Aristoy and Toldrá 2011; Mullen et al. 2017). Due to their amino acid sequences and structural properties, peptides present several biological activities such as antioxidant, antimicrobial, anti-hypertensive (ACE-inhibitory), anticancer, antiviral, antithrombotic, opiate, hypocholesteolemic, immune-modulatory, among the most

outstanding (Helkar et al. 2016; Lafarga and Hayes 2014), which have influence on the human health (Lemes et al. 2016; Liu et al. 2017; Mullen et al. 2017). Moreover, they can be employed as natural preservatives in foodstuffs and as beneficial constituents in functional foods (Di Bernardini et al. 2011a). In fact, despite the lower activity of the peptides from meat by-products compared to those obtained from synthetic products, some scientific researches discuss the possibility of use them as potential substitute of synthetic products, avoiding the potential toxicity and carcinogenesis of artificial products (Lee et al. 2012a).

# Peptides from animal sources

A wide range of foods and by-products have been employed as source of biopeptides production (Lafarga and Hayes 2017). Among these by-products are included blood (Hu et al. 2016; Adje et al. 2011), bones (Chiang et al. 2019; Gallego et al., 2019; Salazar-Posada et al. 2012), collagen (Ryder et al. 2016; Fu et al. 2015; Saiga et al. 2008), gelatin (Lee et al. 2012a, Herregods. 2010), liver (Verma et al. 2017; Di Bernardini et al. 2011b), lungs (O'Sullivan et al. 2017; Lafarga and Hayes 2017), placenta (Teng et al. 2011), skin (Onuh et al. 2014; Lee et al. 2012b), and visceral mass (Bhaskar et al. 2007).

Collagen is the most abundant protein in multiple by-products obtained from meat industry, and the main component in bones, cartilages, hides and skin (Toldrá et al. 2016). Its partial hydrolysis results in gelatin, a soluble protein compound. Many factors have influence on the properties of the gelatins, such as the species and the age of the animal used (Gómez-Guillén et al. 2011). Elastin and keratin are other important fibrous structural proteins, which are part of connective tissues, feathers, hair, ligaments and skin (Ferraro et al. 2016). The peptides obtained can be used as food additives since they have a protective effect against lipid oxidation (Di Bernardini et al. 2011a).

Blood is an excellent reservoir of proteins where the most abundant complex is hemoglobin, an iron rich protein (Toldrá et al. 2016). Plasma is obtained by the addition of

anticoagulants, such as heparin or EDTA, to prevent blood clotting (Böttger et al. 2017). This portion of blood has a great interest due to its functional properties and its absence of color (Jayathilakan et al. 2012).

Most of the peptides have been extracted from bovine and porcine sources, nevertheless the appearance of some diseases and the restrictions established by the regulations or by the religions and culture traditions has made that new alternatives are sought (Sarbon et al. 2013). In this way, several authors support the use of poultry by-products (bones, feet and skin) as source of bioactive peptide (Lee et al. 2010; Jayathilakan et al. 2012; Onuh et al. 2014).

# Methods of peptide extraction from meat by-products

The amino acids that are part of the bioactive peptides are usually inactive in their parent protein (Najafian et al. 2014; Sarmadi et al. 2010). To have functional effects they can be released from the protein sequence through enzymatic hydrolysis, microbial fermentation (proteolytic action of microorganisms), or by acid and alkaline chemical hydrolysis (solvent extraction) (Korhonen et al. 2003; Lafarga and Hayes 2014; Mora et al. 2014; Lemes et al. 2016) (Fig. 1). Enzymatic hydrolysis is considered as the most relevant approach to acquire antioxidant peptides from meat protein (Najafian et al. 2014). Moreover, it is important to note that food processing and gastrointestinal digestion can also release these peptides (Arihara and Ohata, 2010; Escudero et al., 2012).

The hydrolysis conditions, type of enzyme utilized, sequence and configuration of the peptides and the amino acid composition are closely related to the activities of the protein hydrolysates, such as antimicrobial and antioxidant capacities that confer the main impact on the bioactivity (Bah et al. 2015; Wang et al. 2018; Vieira et al. 2017).

# Conventional extraction methods

Enzymatic hydrolysis

Enzymatic treatment is an attractive method to extract proteins since it employs soft procedure conditions, the control of reaction is not difficult, and the generation of by-products is minimal (Liu et al. 2010a). It is the most explored and common technique in literature due to its high grade of hydrolysis. In the bioactive peptide isolation, the success and effectiveness of the breakdown of the proteins depend on the conditions of the experiment, the raw material and the type of the enzyme (Table 1). The pH, temperature and time of the process must be fixed and controlled to ensure the final result. Researchers usually work with meat from farm origin due to their accessibility to raw materials and as well as with poultry, porcine and bovine due to their easy market introduction. Although, other kinds of products like venison (Bah et al. 2015; Kim et al. 2016) or buffalo (Liu et al. 2010b) have been studied as result of the new worldwide requirements. The enzymes used have three possible origins: animal, vegetable and microbial (Toldrá et al., 2018). The main peptidases used are papain (cysteine protease from papaya fruit), bromelain (sulfhydryl protease from pineapple fruit), thermolysin (metalloprotease from the bacteria Bacillus thermoproteolyticus), alcalase (serine proteinase from bacteria Bacillus licheniformis) (Mora et al. 2014), flavourzyme (mixture of proteases from the fungus Aspergillus oryzae) (Feng et al. 2014), pepsin and trypsin (enzymes secreted by the stomach and the pancreas of animals, respectively) (Wen et al. 2015). 

# Acid and alkaline hydrolysis

130 Currently, most research lines are focused on the enzymatic hydrolysis due to their 131 effectiveness of breaking down the structure of proteins and the amount of enzymes available. 132 However, the complexity and the cost of them are higher than other methods (like acid and 133 alkaline hydrolysis). The most common acid hydrolysis treatment for proteins is carried out 134 with HCl and with other sulfonic acids, like methanesulfonic and mercaptoethanesulfonic 135 acids (Fountoulakis et al. 1998), that are taken as a reference to be compared with other 136 treatments like sub-critical water hydrolysis (Rogalinski et al. 2005). On the other hand, the most frequent alkaline hydrolysis is the NaOH treatment, although the KOH is very usual as well. Peptide bonds are cleaved during the alkaline hydrolysis and, depending on the base used, low molecular weight peptides and sodium and potassium salts of free amino acids are formed (Kalambura et al. 2016). These methods have been used in some animal by-products as porcine blood (Álvarez et al. 2013) and chicken feathers, where the use of microwave treatment supposes a reduction in the hydrolysis time (Lee et al. 2016).

#### Microbial fermentation

Another possible source of bioactive peptides is that results from the natural metabolism of bacteria, where their own enzymes release peptides, because they need these proteins as a source of nitrogen, which is essential in their lives. Most of the bioactive peptides isolated by microbial fermentation have been obtained from milk (Aguilar-Toalá et al. 2017), eggs (Nimalaratne et al. 2015) and its derivatives like casein in the case of milk. Lactobacillus are the most common bacteria used due to its greater proteolytic activity (Raveschot et al. 2018). However, this activity is not enough to apply in meat fermentations. As a result, bioactive peptides have barely been isolated from muscle proteins and by-products by this method, perhaps because of the poor proteolytic activity of the Lactobacillus used (Arihara, 2006; Ryan et al. 2011). The lack of literature shows the need to focus the future research in this regard. Even though, *Lactobacillus* are usually used in the microbial proteolysis, other bacteria have also been studied over the years (Da Silva 2017) in Chryseobacterium with chicken feathers (Fontoura et al. 2014) and Monascus purpureus in porcine liver (Yu et al. 2017). The results obtained with the last one strain showed that despite achieving a lower extraction yields compared to those obtained with an enzymatic hydrolysis, the antioxidant activity of the extracted compounds was higher. 

#### Non-conventional extraction methods

Sub-critical water hydrolysis

Other technique used for the extraction of proteins is sub-critical water hydrolysis (SWH). This extraction uses water as solvent in sub-critical state, from 100 °C to 374 °C at 0.10 MPa and 22 MPa, respectively. In the sub-critical state, water has the ability to extract ionic and polar compounds as well as non-polar compounds, leading to the hydrolysis and the breakdown of the protein structure, thus releasing the peptides. Moreover, if the extraction time is increased, it allows the extraction of amino acids. Some researchers have been used this technique for the extraction of peptides from meat by-products (Esteban et al. 2010; Zhu et al. 2010; Álvarez et al. 2012). Álvarez et al. (2012) managed to improve the antioxidant and functional characteristics of the peptides extracted from porcine hemoglobin. The peptides obtained were more soluble and had higher reducing antioxidant power, while a double-step enzymatic hydrolysis would be needed to achieve the same results (Chang et al. 2007). Moreover, the extraction yields obtained were high, obtaining values up to 80% with conditions of time and temperature of 360 minutes and 180 °C. Despite being cheap and clean (not use solvents, so it can be considered as green technology), this extraction can lead to modifications in the structure of the proteins that suppose the loss of their techno functional properties (Lynch et al. 2018; Drummond et al. 2019). 

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# Isoelectric solubilization/precipitation

Other emerging technology is the isoelectric solubilization/precipitation (ISP) that allow to recover the protein in several steps through changes of pH. The process involves a solubilization of the protein in an alkaline medium, a purification and concentration using a membrane technology with a pH adjustment, to finally centrifuge and thus get an isoelectric precipitation (Tahergorabi et al. 2011). The yield of this extraction will depend on both the material from which the extraction is made and the extraction conditions (pH, temperature, extraction time) (Selmane et al. 2008).

# Emerging technologies as pretreatment of meat by-products

There is not much information about the extraction methods of hydrolyzed peptides from meat by-products, being the aforementioned methods those commonly used. However, others can also be applied as alternative techniques to support the extraction procedures of peptides. This is the case of microwave processing that is also used as pretreatment in the chemical alkali and acid hydrolysis of chicken feathers (Lee et al. 2016) and sheep skin (Gousterova et al. 2005).

Recently, in order to avoid the alteration of the functional properties of these proteins, alternative methods are being carried out (Lynch et al. 2018). It is about emerging technologies which avoid using high temperatures and reduce the use of solvents. Between these processing techniques are high hydrostatic pressures, pulsed electric fields, and ultrasounds (Mullen et al. 2017). Pulsed electric fields (PEF) and ultrasounds (US) are considered emerging non-thermal technologies, since in this case heat comes from internal energy generation (Tahergorabi and Hosseini 2017), thus reducing the temperature effects on peptides (cross bonds, peptide breakdown, protein denaturation) (Korhonen et al. 1998). Despite not having much information about the applicability of PEF on the extraction of peptides from meat by-products, some studies suggested its potential use (Gudmundsson and Hafsteinsson 2001). In this way, Ghosh et al. (2019) used this non-thermal technology for the extraction of proteins from waste chicken meat. The application of high voltage, short pulses followed by low voltage, long pulses result on chemical-free extracts with potential antioxidant properties (Gómez et al. 2019). 

High pressure processing technology (HPP) can be considered as an alternative to thermal processes since it employs pressures in the range 400-700 MPa at refrigerated to moderate temperatures (under 50 °C) (Escobeda-Avellaneda et al. 2011). Some researchers have evaluated it as a pretreatment to the extraction of peptides from meat by-products

(Toldrá et al. 2011). The results obtained allow through the changes in the structure of protein to improve the enzymatic hydrolysis and decrease the incubation period.

Ultrasound technique allows through cavitation, wherein microbubbles are formed, grow in size and eventually collapse due to propagation of ultrasounds in the biological matrix (Kadam et al. 2015). This method is faster, and its results show a smaller molecular weight and higher peptide concentrations in porcine cerebral hydrolysate (Zou et al. 2016). Moreover, its combination with enzymatic hydrolysis favors the hydrolysis yields and the antioxidant activity of the peptides obtained (Vidal et al. 2018a,b). 

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# **Purification of protein hydrolysates**

Once the protein hydrolysate is obtained, it is advisable to carry out a fractionation and purification of the mixture of peptides (Agyei and Danquah 2011; Lafarga and Hayes 2014) (Fig. 1). This is a critical step that suppose percentages up to 70% of the peptide production costs (Agyei and Danquah 2011). It is possible isolate the hydrolysates obtained depending on their molecular weight using ultrafiltration with molecular-weight cut-off (MWCO) membranes (Korhonen et al. 2003; Sohaib et al. 2017; Power et al. 2014). The smaller molecular weights of the peptides, the greater the antioxidant activity (Lee et al. 2016). Therefore, a purification process is necessary to introduce them as bioactive substances in food or pharmaceutical products. Size exclusion chromatography (SEC), ionic chromatography (IC) and high-performance liquid chromatography (HPLC) are the three most common techniques used in this stage (Korhonen et al. 2003; Mullen et al. 2017). The last mentioned could be considered the most important due to its high ability to separate biopeptides (Sohaib et al. 2017).

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# **Bioactive peptides as natural antioxidants**

Antioxidants can be natural or synthetic substances with the ability to inhibit or delay oxidative cell damage on oxidizable substrates at relatively low concentrations (Apak et al.

2016). The action of antioxidants positively affects the shelf life and the quality of meat products, since they delay the lipid oxidation and reduce rancidity without modifying negatively their nutritional or sensory properties (Kumar et al. 2015). Many synthetic chemical antioxidants have been recognized, such as tert-Butylhydroquinone (TBHQ), butylated hydroxytoluene (BHT), butylated hydroxyanisole (BHA) and propyl gallate (PG) that have a strong antioxidant activity as food additives. They prevent the deterioration and were extensively applied to extend the shelf life of food (Guo et al. 2015). Nevertheless, they also have unfavorable consequences on DNA and human enzyme system (Liu et al. 2016), so their use should be under stringent limitation (Yang et al. 2018). Thus, natural antioxidants are becoming importance in the meat industry because of the non-acceptance of the customers over the synthetics antioxidants (Kumar et al. 2015). 

The antioxidant importance of peptides was the focus of several researches (Di Bernardini et al. 2011a; Lorenzo et al. 2018). They allow to reduce the oxidative degradation, since they react quickly with the reactive oxygen species (Chakka et al. 2015; Liu et al. 2016). These free radicals, highly chemically reactive, can react spontaneously with cellular components and harm proteins, lipids and DNA (Gupta et al. 2019). They play a critical role causing serious disorders like cancer and cardiovascular diseases (Teng et al. 2011). Also, they can cause damage tissues or cells and lead to aging, coronary heart affections, inflammation, stroke and diabetes mellitus (Ye et al. 2018). Radicals can be deactivated employing antioxidants that act through two major mechanisms, hydrogen atom transfer (HAT) and single electron transfer (ET) (Lorenzo et al. 2018). It must be emphasized that the antioxidant action is a multiple chemical reaction where sometimes many complex mechanisms take part (Li et al. 2018). 

There are many antioxidant peptides reported in meat products like dry-cured hams (Escudero, Aristoy, Nishimura, Arihara, & Toldrá, 2012; Mora, Escudero, & Toldrá, 2016).

In Spanish dry-cured ham, peptides SAGNPN and GLAGA exerted very high antioxidant activity and reducing power, respectively (Escudero, Mora, Fraser, Aristoy, & Toldrá, 2013), peptide SNAAC had an IC50 of 75.2 μM for DPPH assay and 205 μM in reducing power (Mora, Escudero, Fraser, Aristoy & Toldrá, 2014) and was stable against heating and gastrointestinal digestion (Gallego, Mora & Toldrá, 2018a). On the contrary, peptide AEEEYPDL that was also a powerful antioxidant peptide but was cleaved by pepsin during the gastrointestinal digestion (Gallego, Mora & Toldrá, 2018b).

Total antioxidant capacity (TAC) measurement methods are commonly noncompetitive electron transfer (ET) and mixed-mode (ET/HAT) assays (usually involving a redox reaction with the oxidant). The ferric reducing antioxidant power (FRAP) mechanism is completely electron transfer rather than mixed ET/HAT, so in combination with other methods can be very useful in discerning dominant mechanisms with distinct antioxidants. Copper reduction (CUPRAC) assay is a variant of the FRAP assay, it uses copper instead of iron, so it is also included in ET methods (Prior et al. 2005). Despite the 2,2'-azino-bis (3-ethylbenzothiazoline-6-sulphonic acid) (ABTS), trolox equivalent antioxidant capacity and 2,2-diphenyl-1-picrylhydrazyl (DPPH) assessments are generally categorizes as ET reactions. These two radicals may be deactivated either by direct reduction through ET mechanism or by radical quenching via HAT. In contrast, antioxidant activity (AOA) assays, oxygen radical absorbance capacity (ORAC), total radical-trapping antioxidant parameter (TRAP), and total oxyradical scavenging capacity (TOSC) are normally competitive and work on HAT mechanism (Apak et al. 2016). 

The use of various antioxidant methods helps to understand which type or types of mechanisms are involved in the activity of peptides, although the lack of standardization complicates the comparison between assays (Lorenzo et al. 2018; Huang et al. 2005). Among the diverse methods employed to determine the antioxidant capacity (Fig. 2), the most widely

used are 2,2-diphenyl-1-picrylhydrazyl (DPPH), 2,2'-azino-bis (3-ethylbenzothiazoline-6sulphonic acid) (ABTS), metal chelating activity (MCA), reducing antioxidant power (FRAP), oxygen radical absorbance capacity (ORAC) and hydroxyl and superoxide anion radical scavenging activity (Lorenzo et al. 2018).

## Noncellular antioxidant activity

The hydrolysates obtained from the enzymatic hydrolysis of skin by-products displayed antioxidant properties (Table 1). The gelatin hydrolysates obtained stand out for their emulsifying, foaming and gelling properties with important applications in cosmetic, food and pharmaceutical fields (Sarbon et al. 2013). As mentioned before, the enzymes used for their extraction is going to condition their activity, being pepsin and its combination with collagenase which showed the hydrolysate with the highest free radical scavenging activity (Lee et al. 2012a,b). Li et al. (2007) also found that the combination of proteases from bovine pancreas and from *Streptomyces* and *Bacillus polymyxa* resulted in higher antioxidant values.

In contrast, Onuh et al. (2014) found that alcalase showed higher antioxidant capacity than those found by the mixture of pepsin and pancreatin (35% vs 33%, 95% vs 80%, and 3800  $\mu$ M Trolox equivalent/g vs 3200  $\mu$ M Trolox equivalent/g for DPPH, metal chelating effect and ORAC, respectively). These results could be related with the fact that different sources of chicken skin (breast and thigh hydrolyzed with pepsin and pancreatin or alcalase, respectively) were used for these extractions.

Other source of bioactive peptides is bovine lung (O'Sullivan et al. 2017a). It hydrolysis with Alcalase, papain and pepsin results in hydrolysates with antioxidant activity. FRAP assay showed low ferric reducing activity power in the lung hydrolysates. The values were approximately 6  $\mu$ M Fe<sup>+2</sup>/mg/mL, 8  $\mu$ M Fe<sup>+2</sup>/mg/mL and 10  $\mu$ M Fe<sup>+2</sup>/mg/mL for Alcalase, papain and pepsin hydrolysates, respectively. Within ORAC determination, papain and pepsin

hydrolysates showed similar values (400 µmol Trolox equivalents), while Alcalase displayed the highest values (around 450 µmol Trolox equivalents). 

Liver is an important source of peptides. Di Bernardini et al. (2011b) demonstrated the antioxidant activity of the peptidic fractions obtained from bovine liver. Thermolysin was the enzyme used in the hydrolysis of sarcoplasmic proteins. DPPH, FRAP and Fe<sup>2+</sup> chelating ability assays were utilized to evaluate their in vitro antioxidant activity. FRAP showed similar activity in the three types of hydrolysates studied with values of 8.78, 8.37 and 9.37 µg Trolox equivalents/mg for non-ultrafiltered, ultrafiltered 10-kDa MWCO and ultrafiltered 3-kDa MWCO hydrolysates, respectively. Within DPPH, non-ultrafiltered samples displayed the highest scavenging activity (84.09% vs 83.89% and 82.24% for non-ultrafiltered, ultrafiltered 10-kDa and ultrafiltered 3-kDa samples, respectively). On the contrary, ultrafiltered samples displayed the highest metal chelating ability (67.36% and 60.59% vs 55.57% for ultrafiltered 10-kDa and 3-kDa vs non-ultrafiltered samples, respectively).

Alcalase, papain and trypsin were also used for the extraction of hydrolysates from porcine liver (Verma et al. 2017). The samples hydrolyzed with trypsin displayed the highest antioxidant activity in the three assays studied. ABTS radical scavenging activity showed values of 86.8%, while the samples obtained with Alcalase and papain displayed values lower than 75%. The highest values of trypsin could be related with the degree of hydrolysis achieved with trypsin (26.82% vs 23.56% and 19.12% for trypsin vs Alcalase and papain, respectively), since it could be considered as one of the factors that has influence on the ability of hydrolysates to eliminate ABTS<sup>+</sup> radicals (Phanturat et al. 2010). In all cases FRAP values were above 12 mM equivalent to FeSO<sub>4</sub>.7H<sub>2</sub>O. The different values obtained for this method could be related with the type of proteinase used, since the higher reducing power **332** observed in trypsin hydrolysates could be associated with the smallest size peptides obtained from the hydrolysis (Ajibola et al. 2011). In DPPH radical scavenging activity, trypsin 

hydrolysates presented values of 57.5%, whereas in Alcalase and papain hydrolysates the values were below 50%. Yu et al. (2017) observed a similar DPPH radical scavenging activity in porcine liver proteins hydrolyzed with Alcalase and papain (42% and 37%, respectively), while the hydrolysates obtained with pepsin showed the highest values, similar than those obtained for the aforementioned authors (55% vs 57.5%). Chou et al. (2014) evaluated the antioxidant ability of chicken liver hydrolysates obtained by hydrolysis with pepsin. The results obtained showed a higher DPPH ability than those obtained with the same enzyme in porcine liver. However, similar values were obtained for the chelating ability of ferrous ion (about 90%). 

In addition to the known antioxidant activity of the liver, Damgaard et al. (2014) demonstrate that porcine by-products as appendix, colon, heart, liver, lung, pancreas and rectum can also generate antioxidant hydrolysates. Appendix (77.0%, 84.4% and 17.1% for FRAP, ABTS and DPPH, respectively) and colon (79.3%, 86.4% and 17.6% for FRAP, ABTS and DPPH, respectively) showed the highest antioxidant capacity, only surpassed by liver in FRAP, lung in ABTS and heart in DPPH. Similar results were also found by Damgaard et al. (2015), who confirmed that hydrolysates obtained from porcine (colon, heart and neck) and bovine (lung, kidney and pancreas) tissues displayed antioxidant activity. In this case, bovine kidney and lung showed the higher ion chelation (63.4% and 55.0%, respectively), while porcine colon and heart presented the higher ABTS values (49.6% and 44.6%, respectively). 

Protein hydrolysates were also obtained from poultry viscera (Jamdar et al. 2012). The extraction of the hydrolysates was carried out though an autolytic degradation of protein of connective tissues, gall bladder, intestine and spleen. Hydrolysates showed higher values than those obtained in peptide fractions (<3 kDa, 3-10 kDa, >10 kDa) in all of the radical

scavenging activity assays (721.6 μM TEAC/mg, 244.2 μM TEAC/mg, 47.0% and 43.9% for
ABTS, DPPH, Hydroxyl and Superoxide radical scavenging activity, respectively).

Besides to viscera, there are other duck processing by-products such as feet, bones and heads that have been evaluated as a source of valuable bioactive compounds (Lee et al. 2010). In this study, eight proteases (Alcalase,  $\alpha$ -chymotrypsin, flavourzyme, neutrase, papain, pepsin, protamex and trypsin) were used to isolate antioxidant peptides. The hydroxyl radical scavenging activity of the obtained enzymatic hydrolysates displayed IC<sub>50</sub> values of 571 mg/mL. Lee et al. (2012b) obtained similar values when these enzymes were used in duck skin by-products (554 µg/mL).

Recent results also showed the potential of dry-cured ham bones as a source of antioxidant peptides that retain their bioactivity after household cooking preparations and gastrointestinal digestion. In fact, cooking using conventional household methods increased the antioxidant activity of ham by-products whereas simulated gastrointestinal digestion showed no significant effect on the antioxidant activity of ham by-products and antioxidant activity decreased when assessed using the ORAC and  $\beta$ -carotene bleaching assays. In this study, collagen peptides were responsible for the differences in antioxidant activities observed in cooked samples after simulated gastrointestinal digestion. This fact confirmed the traditional believe of bioactive properties of bone broths and stews (Gallego et al. 2017).

Blood is other edible by-product considered as a source of bioactive compounds due to its important protein contents (Bah et al. 2013). It is composed by two fractions: cellular (hemoglobin) and plasma (immunoglobulins, fibrinogen and serum albumin). Hydrolysates with significant antioxidant activity were found in bovine blood (O'Sullivan et al. 2017b). Papain was used for the hydrolysis of the blood protein fractions ( $\alpha/\beta$ -globulins, serum albumin,  $\gamma$ -globulins and fibrinogen).  $\gamma$ -globulin and fibrinogen hydrolysates were those that showed the highest activity. The values of fibrinogen were around 25  $\mu$ M Fe<sup>+2</sup>/mg/mL in

FRAP assay and 1000 μmol Trolox equivalents in ORAC determination. γ-globulin displayed an intermediate activity with values around 17 µM Fe<sup>+2</sup>/mg/mL and 800 µmol Trolox equivalents for FRAP and ORAC assays, respectively. Finally,  $\alpha/\beta$ -globulins showed the lowest values both in FRAP and ORAC assays (less than 5  $\mu$ M Fe<sup>+2</sup>/mg/mL and less than 250 umol Trolox equivalents, respectively).

The hydrolysates isolated from plasma has several properties, highlighting its antioxidant activity (Liu et al. 2010a,b; Adje et al. 2011; Nyberg et al. 2013). Bah et al. (2015) also claimed that plasma of deer, sheep and pig allow to obtain bioactive peptides and peptide-rich protein hydrolysates. Their antioxidant activity was tested through DPPH, FRAP and ORAC assays. The hydrolysates obtained with the use of proteases from fungal displayed higher DPPH values than those found with other enzymes from plants (papain and bromelain). The hydrolysates obtained from pig plasma with fungal enzymes also displayed higher DPPH (55.2%) activities than those found in other porcine plasma hydrolysates obtained using pepsin (48.4%) and trypsin (11.5%) (Wei and Chiang 2009; Xu et al. 2009), but lower than those obtained with Alcalase (76.5%; Liu et al. 2010a). A similar behavior was observed with FRAP and ORAC essays. Higher capacity was obtained with fungal proteases and in deer and sheep plasma hydrolysates. This could be due to these fungal enzymes are capable of producing small peptides associated with highest antioxidant properties (Di Pierro et al. 2014).

# Cellular antioxidant activity

In vivo determinations are necessary to evaluate bioavailability and functionality of antioxidant peptides. These assays could be due through cell or animal models. Both models allow to know more information about the metabolism of these antioxidant compounds (Liu et al. 2016). The cellular antioxidant activity of the hydrolysates could be tested though the comet assay. This method, based on the method described by Phelan et al. (2009), allows to

evaluate the potential ability of hydrolysates to preserve the cells from DNA damage. This harm could be due to the reaction between free radicals and cellular components (Gupta et al., 2019). In this procedure, U937 lymphocytes cells were incubated with hydrolysates for 24 hours to then start the cell damage with H<sub>2</sub>O<sub>2</sub> for 30 minutes at 37 °C. The results can be expressed as tail DNA damage or as a relative percentage to  $H_2O_2$  control values. The last ones were conducted in order to know the concentration needed to produce a significant DNA damage. Some researchers have been used this assay in hydrolysates obtained from meat byproducts. O'Sullivan et al. (2017a,b) evaluated the cellular antioxidant activity of bovine lung and blood hydrolysates. These hydrolysates did not display ability to prevent DNA damage. In contrast, peptides obtained from buffalo horn showed protection against the damage of rat cerebral cells (Liu et al. 2010b). Other studies were carried out in rat liver cells (Ac2F) to evaluate the effect that the presence or the absence of peptides has on cell viability (Kim et al. 2001). The MTT method (role of mitochondrial succinate dehydrogenase) will allow to ascertain the possible mechanisms of non-enzymatic cellular defence induced by these compounds.

Animal models assess the bioavailability of antioxidant peptides through different methods such as enzymatic defence catalase (CAT), superoxide dismutase (SOD) and glutathione peroxidase (GPx) (Liu et al. 2016). The toxicity studies are usually carried out using Wistar rats and the bioactive compounds are determined in the gastrointestinal tract, liver and kidney (Lee et al. 2012b).

# Antimicrobial activity of bioactive peptides

Conventional antibiotics have been used for a long time due to their active properties; however, the emergence of resistance to them has led to seek new sources (Lafarga and Hayes 2014). Peptides could meet this demand since there are studies that assert that they are able to inhibit the action of strains of pathogenic and altering microorganisms (Table 2) such as

Gram-positive (Listeria monocytogenes, Bacillus cereus, *Staphylococcus* aureus, Micrococcus luteus) and Gram-negative (Escherichia coli, Yersinia enterocolitica) (Adje et al. 2011; Di Bernardini et al. 2011a). The highest antibacterial activity was observed when the molecular weight of these peptide hydrolysates was between 400 and 1400 Da (Ryan et al. 2011). Moreover, some authors confirmed that a  $\alpha$ -helical conformation favors their action against microorganisms (Daoud et al. 2005). In fact, an antibacterial peptide was obtained from a fragment of the  $\alpha$ -chain of bovine hemoglobin. This peptide had activity against Escherichia coli, Listeria innocua, Micrococcus luteus and Salmonella enteritidis. Similar results were also obtained by Nedjar-Arroume et al. (2006). Another peptide RHGYM, isolated from dry-cured ham, was able to inhibit the growth of L. monocytogenes, showing a MIC value of 6.25 mM (Castellano et al. 2016). 

The commonly method used to evaluate this activity is the disk diffusion method (Kirby-Bauer method). It is a technique based on the method described by Bauer et al. (1966), which consists on the relationship between the concentration of the substance necessary to inhibit a bacterial strain and the inhibition halo of growth on the surface of an agar plate with a suitable culture medium and seeded homogeneously with the bacteria to be tested (Fig. 2). As a positive control for gram-positive bacteria could be used Nisin and Gentamicin as a positive control against gram-negative bacteria.

452 Many researchers have tested the antibacterial activity of protein hydrolysates. Chakka 453 et al. (2015) found that the hydrolysates obtained from the enzymatic hydrolysis of chicken 454 liver only displayed a moderate inhibition against *Micrococcus luteus* (12 mm). Whereas, 455 hydrolysates obtained from porcine liver with Alcalase, papain and trypsin showed 456 antimicrobial activity against *Bacillus cereus, Escherichia coli, Listeria monocytogenes and* 457 *Staphylococcus aureus*. As it happened with the antioxidant activity, trypsin hydrolysates 458 displayed the highest antimicrobial activity in all of the strains evaluated. These results could be due to the influence that the degree of hydrolysis had in the antimicrobial activity and the hydrophobic and cationic properties of the peptides obtained. These abilities allow to modify the permeability of microbial membranes leading to their breakdown. Moreover, the smaller size peptides obtained from the hydrolysis with trypsin would favor the interaction with the microbial membrane (Verma et al. 2017).

No antimicrobial activity was found by Bah et al. (2015) in the hydrolysates obtained from deer, pig, and sheep plasma with plant and fungal proteases against strains usually found in meat (E. coli, Pseudomonas aeruginosa and Staphylococcus aureus). A possible explanation for this behavior could be related to the fact that the enzymes used are not able to extract peptides from the plasma proteins with this ability. In this way, alternative hydrolysis methods were developed in order to obtain peptides with antibacterial properties. These methods consist in limiting the hydrolysis using an alcohol, which results in a modification of the protein conformation. Adje et al. (2011) studied this controlled hydrolysis using several alcohols (butanol, ethanol, methanol or propanol) and pepsin. The results obtained were satisfactory since it was possible to isolate antimicrobial peptides. These compounds were effective against Escherichia coli, Listeria innocua and Micrococcus luteus. In his case, the activity of the peptides was calculated through the minimum inhibitory concentration (MIC), that allow to know the minimum concentration enough to completely inhibited the microbial growth on agar plates after 24 hours at 37 °C (Froidevaux et al. 2001). 

Conclusions

Slaughtered by-products are an important source of novel bioactive compounds, what makes them become into products with a great added-value with the consequent positive economic impact for meat sector and environmental benefits. The hydrolysates extracted from these waste materials have the potential to be a protein rich ingredient for use in formulated food products and possible help in reduction of oxidative and microbial deterioration, since

the antimicrobial and antioxidant properties associated with these peptides would allow to extend the shelf life of food, even improving the results obtained with synthetic products. The results of noncellular antioxidant activity are validated by in vivo models, which allow to evaluate their toxicity, bioaccessibility and bioavailability, and above all their safety for their use. The pretreatment with emerging technologies would increase the quality of these compounds since they hardly affect the composition and the structure of these hydrolysate compounds, increasing the extraction yields as well as their antioxidant properties. In this way, the combination of HPP, PEF or US with enzymatic hydrolysis is effective as it improves the hydrolysis yields and the antioxidant activity of the peptides obtained. Finally, besides bovine and porcine sources, this review confirms the possibility of use poultry byproducts as valuable bioactive peptides resource. 

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# **Compliance with Ethical Standards**

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# **Conflict of Interest**

Paula Borrajo declares that she has no conflict of interest. Mirian Pateiro declares that she has no conflict of interest. Francisco J. Barba declares that he has no conflict of interest. Leticia Mora declares that she has no conflict of interest. Daniel Franco declares that he has

1	508	no conflict of interest. Fidel Toldrá declares that he has no conflict of interest. José M.
1 2 3	509	Lorenzo declares that he has no conflict of interest.
4 5	510	Ethical Approval
6 7 8	511	This article does not contain any studies with human or animal subjects.
9 10	512	Informed consent
11 12 13	513	Not applicable.
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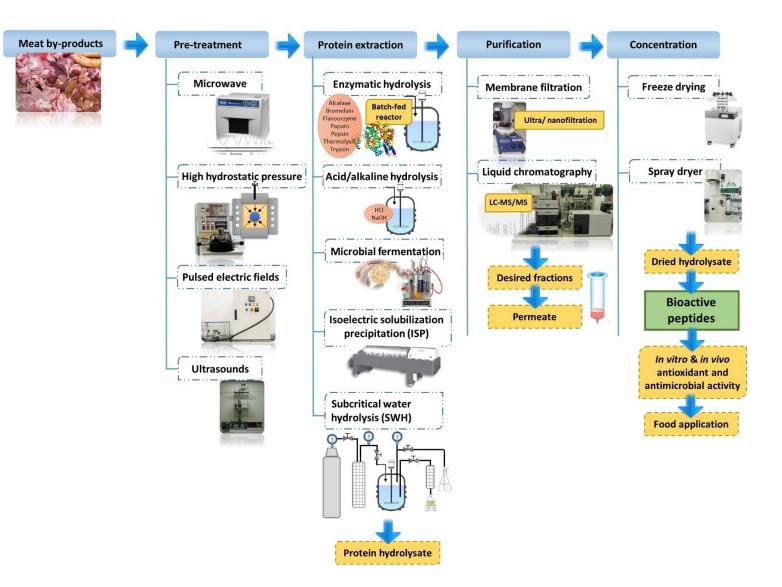
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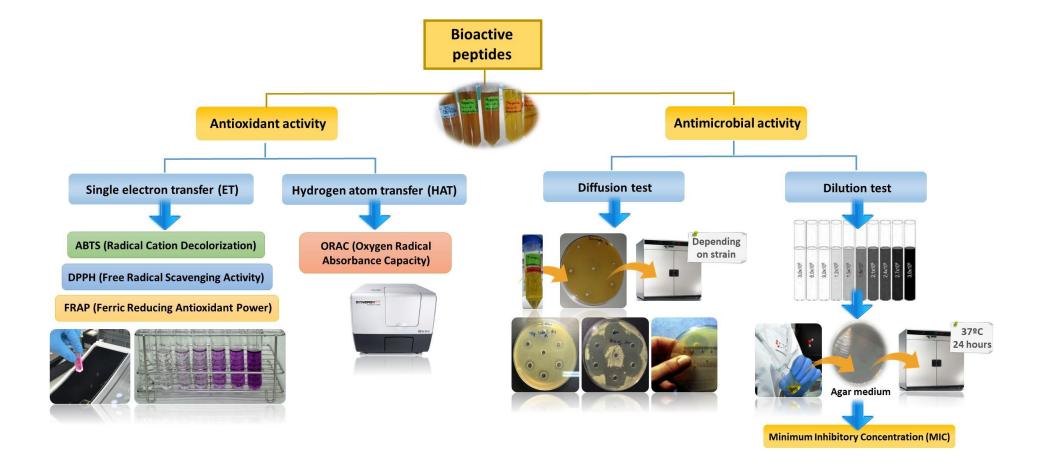
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	831	FIGURE CAPTIONS:
1 2 3	832	Figure 1. Extraction process of bioactive peptides from meat by-products
4 5	833	Figure 1. Antioxidant and antimicrobial determinations generally used to assess the activity of
6 7	834	peptides extracted from meat by-products
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Figure



		Incubation		Antioxidant assays					
Source	Enzyme	conditions	Deactivation	ABTS	DPPH radical scavenging activity	FRAP	Fe <sup>2+</sup> chelating ability	ORAC (Trolox equivalents)	Reference
					Poultry				
	Pepsin	37°C, 5h	95°C, 15 min	n/a	90.7%	n/a	94.2%	n/a	Chou et al. (2014)
Liver	Alcalase	45°C, 150min, 120 rpm	90°C, 10 min	19.3%	92.8%	n/a	n/a	n/a	Chakka et al. (2015)
	Alcalase	pH 8.0, 55°C, 4h		n/a	≅ 35%	n/a	≅ 95%	$\cong 3800 \ \mu M/g$	
Skin	Pancreatin, pepsin	pH 2.0, 37°C, 2h	95°C, 15 min	n/a	≅ 33%	n/a	≅ 80%	$\cong 3200 \ \mu M/g$	Onuh et al. (2014)
Viscera	Autolytic	pH 2.8, 55°C, 6h		721.6 µM/mg	244.2 µM/mg	n/a	n/a	n/a	Jamdar et al. (2012)
					Bovine				
Blood (fibrinogen)	Papain	pH 6.5, 65°C, 24h	95 °C, 10 min	n/a	n/a	$25 \ \mu M \ Fe^{+2}/mg \ mL^{-1}$	n/a	1000 µmol	O' Sullivan et al. (2017b)
-	Papain	pH 6.5, 65°C, 24h		n/a	n/a	$8 \ \mu M \ Fe^{+2}/mg \ mL^{-1}$	n/a	400 µmol	
Lung	Pepsin	pH 2.0, 37°C, 24h	95 °C, 10 min	n/a	n/a	$10 \ \mu M \ Fe^{+2}/mg \ mL^{-1}$	n/a	400 µmol	O' Sullivan et al. (2017a)
	Alcalase	pH 9.5, 60°C, 24h		n/a	n/a	$6 \ \mu M \ Fe^{+2}/mg \ mL^{-1}$	n/a	450 µmol	(2017a)
Liver	Thermolysin	pH 9.5, 37°C, 2h, 150 rpm	99°C, 10 min	n/a	*84.1%, 82.2% and 83.9%	*8.8, 9.4 and 8.4, μg Trolox equivalents/mg	*55.6%, 60.6% and 67.4%	n/a	Di Bernardini et al. (2011)
	Alcalase	pH 8.0, 50°C, 6h		n/a	6 mg/mL (IC <sub>50</sub> )	n/a	n/a	n/a	
	Flavourzyme	pH 7.0, 50°C, 6h		n/a	4 mg/mL (IC <sub>50</sub> )	n/a	n/a	n/a	
Yak skin	Pepsin, trypsin	pH 2.0, 37°C, 6h pH 7.0, 37°C, 6h	100 °C, 10 min	n/a	2.6 mg/mL (IC <sub>50</sub> )	n/a	n/a	n/a	Tian et al. (2017)
	Proteinase K	рН 7.5, 37°С, 6h		n/a	5 mg/mL (IC <sub>50</sub> )	n/a	n/a	n/a	
	Protamex	pH 6.5, 50°C, 6h		n/a	4.5 mg/mL (IC <sub>50</sub> )	n/a	n/a	n/a	
					Porcine				

**Table 1.** Antioxidant activity of bioactive peptides isolated from meat by-products after enzymatic hydrolysis.

	Alcalase, Flavourzyme	pH 8.5, 50°C, 4h; pH 7.5, 50°C, 6h	90°C, 20 min	n/a	41.9%	0.23%	63.5%	n/a	Chang et al. (2007)
	Alcalase	pH 7.5, 55°C, 24h		n/a	30.8%	n/a	n/a	n/a	
Hemoglobin	Chymotrypsin, thermolysin and trypsin	pH 7.5, 35°C, 24h	100°C, 15 min	n/a	24.4%	n/a	n/a	n/a	Wei et al. (2009)
	Trypsin	pH 7.0, 38°C, 24h		n/a	11.5%	n/a	n/a	n/a	
	Papain	pH 6.5, 50°C, 6h		70.6%	40.3%	12.7 mM equivalent FeSO <sub>4</sub> .7H <sub>2</sub> O	n/a	n/a	
	Alcalase	pH 8.0, 50°C, 6h	85°C, 15 min	74.6%	42.3%	13.7 mM equivalent FeSO <sub>4</sub> .7H <sub>2</sub> O	n/a	n/a	Verma et al. (2017)
T	Trypsin	pH 8.0, 37°C, 6h		86.8%	57.5%	14.9 mM equivalent FeSO <sub>4</sub> .7H <sub>2</sub> O	n/a	n/a	
Liver	Alcalase	55°C, 2h	95°C, 10 min	79.2%	9.9%	n/a	92%	n/a	Daamgard et al. (2014)
	Papain	pH 6.5, 37°C, 12h		n/a	37%	n/a	n/a	n/a	Yu et al. (2017)
	Pepsin	pH 3.0, 37°C, 12h	95-100°C, 10 min	n/a	55%	n/a	n/a	n/a	
	Alcalase	pH 8.0, 50°C, 12h	11111	n/a	42%	n/a	n/a	n/a	
	Bromelain	pH 6.5, 50°C, 24h		n/a	≅ 30%	≅ 400 mM equivalent FeSO₄/g	n/a	$\cong 10 \text{ mM}$	
	Papain	pH 6.5, 55°C, 24h	90°C, 20 min	n/a	≅ 30%	≅ 600 mM equivalent FeSO₄/g	n/a	$\cong 10 \text{ mM}$	Bah et al. (2015
Plasma	Fungal proteases	pH 6.5, 50°C, 24h		n/a	≅ 50-60%	≅ 750-1000 mM equivalent FeSO₄/g	n/a	> 10 mM	
	Alcalase	pH 8.0, 55℃, 5h	95°С, 5 min, pH 7.0	n/a	76.5%	1407.9 μM	12.0%	n/a	Liu et al. (2010)
	Pepsin	pH 2.0, 37°C, 5h	10090 2	n/a	48.4%	n/a	≅ 25 %	n/a	X
	Papain	pH 8.0, 37°C, 16h	100°C, 3 min	n/a	43.1%	n/a	$\cong 8\%$	n/a	Xu et al. (2009)
Skin collagen	Cocktail proteases	pH 7.5, 37°C, 24h	100°C, 3 min	n/a	87.2%	n/a	37.4%	n/a	Li et al. (2007)
					Venison				

	Bromelain	pH 6.5, 50°C, 24h		n/a	≅ 25%	$\cong$ 500 mM equivalent FeSO <sub>4</sub> /g	n/a	$\cong 14 \text{ mM}$	_
Deer Plasma	Papain	pH 6.5, 55°C, 24h	90°C, 20 min	n/a	≅ 30%	≅ 650 mM equivalent FeSO₄/g	n/a	≅ 17.5 mM	Bah et al. (2015)
	Fungal proteases	pH 6.5, 50°C, 24h		n/a	≅ 55-65%	$\cong$ 700-1100 mM equivalent FeSO <sub>4</sub> /g	n/a	≅ 18-22 mM	
	Celluclast	pH 4.5, 50°C, 24h		n/a	3-5 mg/mL	n/a	n/a	n/a	
Malaset anti-	Flavourzyme	pH 7.0, 50°C, 24h	100% 10	n/a	1.3-2.7 mg/mL	n/a	n/a	n/a	$V_{int} = t = 1$ (2016)
Velvet antler	Pepsin	pH 7.0, 37°C, 24h	100°C, 10 min	n/a	n/a	n/a	n/a	n/a	Kim et al. (2016)
	Termamyl	pH 6.0, 60°C, 24h		n/a	n/a	n/a	n/a	n/a	
					Caprine				
Goat Placenta	Pepsin	рН 5.0, 36.5°С	100°C, 10 min	n/a	96.7%	n/a	n/a	n/a	Teng et al. (2011)
	Bromelain	pH 6.5, 50°C, 24h		n/a	≅ 28%	≅ 400 mM equivalent FeSO₄/g	n/a	$\cong 16 \text{ mM}$	
Sheep Plasma	Papain	pH 6.5, 55°C, 24h	90°C, 20 min	n/a	≅ 30%	≅ 700 mM equivalent FeSO₄/g	n/a	$\cong 15 \text{ mM}$	Bah et al. (2015)
	Fungal proteases	pH 6.5, 50°C, 24h		n/a	≅ 55-63%	≅ 800-1200 mM equivalent FeSO₄/g	n/a	≅ 15-18 mM	

\*for non-ultrafiltered, ultrafiltered with 3 kDa and 10 kDa; n/a: data not available.

<b>G</b>			Antin	-			
Source	Extraction method	MW (Da)	Strains	MIC (µM)	Inhibition (mm)	Reference	
				Bovine			
		2236.9	Micrococcus luteus	671	n/d	Froidevaux et al. (2001)	
	-		Listeria innocua	38	25		
		2150	Escherichia coli	76	17		
		3150	Micrococcus luteus	76	15	Daoud et al. (2005)	
			Salmonella enteritidis	76	16		
	Hydrolysis		Salmonella enteritidis	5-87			
	(Pepsin)		Escherichia coli	8-87			
Hemoglobin		655-3152	Listeria innocua	1-71	n/d	Nedjar-Arroume et al. (200	
			Micrococcus luteus	9-87			
	-		Micrococcus luteus				
		653.7- 5437.3	Listeria innocua	1-187.1	n/d	Adje et al. (2011)	
			Escherichia coli				
	<b>.</b>		Candida albicans		7.6		
	Lysis (Ammonium	1992.4	Escherichia coli	n/d	10.3	Hu et al. (2011)	
	chloride)		Staphylococcus aureus		9.5	···· 、/	
				Porcine			
			Bacillus cereus		16.8		
<b>T</b> ·	Hydrolysis	(1	Listeria monocytogenes	<i>.</i> •	23.4		
Liver	(Trypsin)	n/d	Escherichia coli	n/d	18.2	Verma et al. (2017)	
			Staphylococcus aureus		18.4		
	Hydrolysis		Escherichia coli				
Plasma	(Bromelain, Papain,	n/d	P aeruginosa	n/d	n/d	Bah et al. (2015)	
	Fungal proteases)		Staphylococcus aureus				
				Poultry			
			Bacillus cereus		28		
<b>T</b>			Listeria monocytogenes		30	(1,1)	
Liver	Fermentation	n/d	Micrococcus luteus	n/d	18	Chakka et al. (2015)	
			Yersinia enterocolitica		18		

<b>Table 2.</b> Antimicrobial activity of bioactive	peptides isolated from meat by-	products.
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Hydrolysis (Alcalase) Micrococcus luteus	12	
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n/a: data not available; n/d: not determinated.