

1	IDENTIFICATION OF ANTIOXIDANT PEPTIDES OF HEN EGG WHITE
2	LYSOZYME AND EVALUATION OF INHIBITION LIPID PEROXIDATION AND
3	CYTOTOXICITY IN THE ZEBRAFISH MODEL
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6	Carrillo W ^{1, 2*} , Gómez-Ruiz J.A ¹ , Miralles B ¹ , Ramos M ¹ , Barrio D ³ , Recio I ¹ .
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8	¹ Instituto de Investigación en Ciencias de la Alimentación, CIAL (CSIC-UAM), Nicolás
9	Cabrera, 9, 28049 Madrid, Spain
10	² Facultad de Ciencia e Ingeniería en Alimentos, Universidad Técnica de Ambato. Av.
11	Los Chasquis y Rio Payamino. Campus Huachi, CP 1801334, Ambato-Ecuador.
12	³ Universidad Nacional de Rio Negro, Don Bosco y Leloir s/n (8500) Río Negro-
13	Argentina.
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27	[*] Corresponding author: Dr. Carrillo W
28	Phone: +593 (3) 2400987 Ext. 121
29	Fax: +593 980281086
30	E-mail address: wi.carrillo@uta.edu.ec

31 Abstract

Hen egg lysozyme was hydrolyzed with pepsin in situ on a cation-exchange column to 32 isolate antioxidant peptides. The most cationic fraction was eluted with 1 M NaCl. Five 33 positively charged peptides f(109-119) VAWRNRCKGTD, f(111-119) WRNRCKGTD, 34 f(122-129) AWIRGCRL, f(123-129) WIRGCRL and f(124-129) IRGCRL were identified 35 using tandem mass spectrometry. Using ORAC-FL, all five peptides presented 36 37 antioxidant activity with values of (1,970; 3,123; 2,743; 2,393 and 0.313 µmol Trolox/ 38 µmol peptide) respectively. Using method TBARS in zebrafish larvae, all five synthetic peptides were found to efficiently inhibit lipid peroxidation (36.8; 51.6;55.56; 63.2; 39 40 61.0 % inhibition of lipid peroxidation) respectively. None of the five peptides were toxic in zebrafish eggs and larvae at concentrations lower than 50 µg/ml. 41 42 Concentrations higher than 50μ g/ml were toxic for both zebrafish eggs and larvae.

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Keywords: lysozyme, antioxidant activity in zebrafish larvae, bioactive peptides,
hydrolysate, cation exchange column and toxicity in zebrafish egg.

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47 Introduction

Lysozyme is a basic protein consisting of 129 amino acids with a molecular weight of 48 14.3 kDa. These amino acid residues are cross-linked by four disulfide bridges, and 49 50 have an isoelectric point of 10.7. Hen egg is the richest source of lysozyme, accounting for 3.5% of total egg white proteins [1]. Lysozyme belongs to a type of enzymes that 51 52 lyses the cell wall of certain Gram-positive bacteria by splitting β (1-4) linkages 53 between N-acetylmuramic acid and N-acetylglucosamine of the peptidoglycan [2]. As a well-known antimicrobial protein, hen egg white lysozyme has been commercialized 54 for applications as a natural preservative to control lactic bacteria in meat products 55 56 such as sausages, salami, pork, beef or turkey. Lysozyme has also been used to prevent growth of *Clostridium tyrobutyricum* in cheese production or to control lactic bacteria 57 in wine and beer production [3, 4, 5, 6, 7]. Lysozyme can also be used in other 58 pharmaceutical and cosmetics applications [3, 7]. Moreover, lysozyme is an enzyme 59 60 widely used as food additive (E1105) due to its numerous properties [8]. Lysozyme has many other functions, including antiviral [9, 10], immune modulatory [11], anti-61

inflammatory [12] and antitumor [13] activities. At pH 7.0, lysozyme is positively
charged, whereas the rest of the proteins of the egg white are negatively charged.
Many cationic proteins such as lactoferrin, lactoperoxidase and lysozyme may be
purified using ion exchange, as this fact has already been demonstrated [14]. Lysozyme
has also been purified with cation-exchange membranes and resins [15, 16].

67 Bioactive peptides have between 3-20 amino acid residues; their bioactivity depends 68 on the sequence and amino acid compositions [17-19]. Recently, attention has mainly focused on the antioxidant peptides generated from food proteins, being these 69 70 peptides safer and healthier than synthetic drugs [20]. Antioxidant peptides contain 5-16 amino acid residues. Their antioxidant activities can be related to ion chelating, 71 radical scavenging and inhibition of lipid peroxidation. The importance of positively 72 73 charged amino acids in determining the strength of peptides as antihypertensive and 74 antioxidants has been indicated in different studies. Strong antimicrobial peptides are cationic charged. Those cationic charged peptides content amino acids as Lys, Arg and 75 76 His [21, 3]. Lysozyme has an isoelectric point of 10.7, with a high content of positively 77 charged amino acids. Lysozyme may be a great substrate for production of bioactive peptides with antioxidant activity. You et al., (2010)[3] have described two 78 79 antioxidants fractions of hydrolysate of lysozyme with pepsin containing positively charged amino acids such as f(13-20)KRHGLDNY, f(14-23)RHGLDNYRGY and f(13-80 81 23)RHGLDNYRGY. Moreover, many researchers have reported that peptides and 82 protein hydrolyzed from various food sources have significant antioxidant activity [22]. 83 Furthermore, hen egg white lysozyme suppresses reactive oxygen species (ROS) 84 generation and protects against acute and chronic oxidant injuries [20, 23]. Some 85 peptides have shown to have multifunctional activities [3, 24]. Different bioactive 86 peptides from lysozyme have been reported with antimicrobial, antioxidant and 87 antihypertensive activities [25-29].

The zebrafish (*Danio rerio*) has become a promising model organism for experimental studies in different biomedical areas. Zebrafish is an ideal animal model for laboratory research. These animals are inexpensive, low-maintenance, and abundantly produced all year round [30-33]. Zebrafish genes are highly conserved sharing a 70 – 80% homology to those of humans [34]. The transparent embryos rapidly develop externally. Organogenesis is completed within the first 48 hours of development. Since

94 zebrafish embryos develop externally, changes in development may be continuously monitored and observed, which greatly facilitates developmental time course studies. 95 96 Zebrafish development has been well characterized and therefore results from 97 zebrafish are comparable to mammalian developmental studies [35-37]. Moreover, zebrafish is a vertebrate model for modeling behavioral and functional parameters 98 99 related to human pathogenesis and for clinical treatment screening. More recently, 100 zebrafish has become also a valuable model to environmental and toxicological 101 studies. Therefore, zebrafish model can be an interesting model to evaluate toxicology of new ingredients of functional foods such as antioxidant peptides. 102

103 In this study, ion-exchange chromatography has been used to isolate the bioactive 104 peptides from hen egg white lysozyme. Lysozyme was hydrolyzed in situ with pepsin to 105 generate positive charged peptides. Those peptides were separated on a cation-106 exchange column by selective elution. The objective was to identify new antioxidant 107 peptides and evaluate their toxicity in the Zebrafish model (*Danio rerio*).

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109 Materials and methods

110 Chemicals

Hen egg white lysozyme 58,000 U/ml, pepsin crystalline 3,440 U/mg obtained from
porcine stomach mucus, 2,20-azobis (2-methylpropionamide)-dihydrochloride (AAPH),
6-hydroxy-2,5,7,8-tetramethylchroman- 2-carboxylic acid (Trolox), fluorescein
disodium (FL), and dithiothreitol (DTT) were obtained from Sigma Chemical (Saint
Louis, MO, USA). The rest of chemicals used were of HPLC grade.

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Pepsin hydrolysis of lysozyme in situ on an ion-exchange column and isolation ofpeptides

Denaturation of lysozyme was performed as previously described by Carrillo et al., 2014 [38]. Lysozyme was denatured using heat treatment. Lysozyme at 5mg/ml was suspended in buffer phosphate pH 6.0 and heat at 95°C during 20 minutes. Then, lysozyme was lyophilized and stored at -20°C. To carry out the hydrolysis of columnbound native and denatured lysozyme, 1000 ml of a 0.1 mg/ml solution of protein in 10 mM NaCl, adjusted to pH 2.0 with HCl, were pumped through a column cation125 exchange column. The binding of native and denatured lysozyme was carried out at 126 room temperature (25°C) with at a flow rate of 20 ml/min, which was generated by a peristaltic pump (Verder-Vleuten, Vleuten, The Netherlands). The process was 127 128 monitored by a UV detector with a 2 mm light path flow cuvette (Model EM-1 Econo UV Monitor, Bio-Rad) at 280 nm. Prior to be used, the ion-exchange column was pre-129 130 equilibrated with water acidified with HCl (pH 2.0). The native and denatured lysozyme bound to the column were hydrolyzed at 37°C by recycling with 100 ml of an aqueous 131 132 solution (pH 2.0) of porcine pepsin (25 mg/ml) at 20 ml/min during 6 hours. The 133 column was washed sequentially with acidified water at pH 2.0. Solvent A was 10 mM ammonium hydrogen carbonate acidified to pH 7 with formic acid, and solvent B was 134 3M and 5M of ammonia solution and finally the column was treated with 1 M NaCl to 135 136 remove more cationic peptides. The effluent was monitored at 280 nm. All fractions were collected with Fast Protein Liquid Chromatography (FPLC) of GE-Pharmacia, 137 freeze-dried, and analyzed with a high-performance liquid chromatography-138 139 electrospray ionization-tandem mass spectrometry (HPLC-ESI-MS-MS).

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141 Identification of peptides by ESI-MS/MS

142 The selected fractions separated and collected from FPLC were analyzed by RP-HPLC-ESI-MS/MS, on an Agilent 1100 HPLC System (Agilent Technologies, Waldbron, 143 Germany) connected on-line to an Esquire 3000 ion trap (Bruker Daltonik GmbH, 144 145 Bremen, Germany) and equipped with an electrospray ionization source, as described 146 by López-Expósito et al. (2006) [39]. The variable-wavelength detector was set at 214 147 nm. A C18-guard column (Nova-Pak[®] 20 mm × 2826 x 3.9 × 4 µm of particle size; 148 Waters Corp., Milford, MA, USA) was used to protect the analytical column (HiPore® 149 RP318 C18 column 250×4.6 mm and 5 μm of particle size; Bio-Rad, Richmond, CA, 150 USA). The samples were eluted at 0.8 mll/min with a linear gradient from 0 to 45% of 151 solvent B (acetonitrile and TFA, 1,000:0.270, v/v) in solvent A (water and TFA, 152 1,000:0.370, v/v) in 60 minutes. The injection volume was 50 μ l and duplicate of 153 injection was made for each point of the standard curve and the samples. The flow 154 from HPLC was divided approximately 1:3 previous to ionization source, and the first 6 min of the eluent flow was directed to waste to reduce salt deposit on the transfer 155 156 capillary of the MS instrument and to reduce interferences. For HPLC-MS, spectra were

recorded over the mass-to-charge (m/z) range of 100 to 1,500. Helium was used as collision gas with an estimated pressure of 5×10–3 bar. About 15 spectra were averaged in the MS analyses and about five spectra in the tandem MS analyses. Using Data AnalysisTM (version 3.0; Bruker Daltoniks), the m/z spectral data were processed and transformed to spectra representing mass values. The acquired MS/MS spectra were interpreted using BioTools (version 2.1; Bruker Daltoniks).

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164 **Peptide synthesis**

165 The synthetic peptides from lysozyme VAWRNRCKGTD, f(109-119), WRNRCKGTD, 166 f(111-119) AWIRGCRL, f (122-129), WIRGCRL, f (123-129) and IRGCRL, f (124-129) 167 were prepared using a conventional FMOC solid-phase synthesis method with a 431A 168 peptide synthesizer (Applied Biosystems Inc. Überlingen, Germany).

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170 Oxygen radical absorbance capacity-fluorescein (ORAC-FL) assay

171 The ORAC-FL assay was based on the assay proposed by Ou, Hampsch- Woodill, & Prior 172 (2001) and Dávalos et al., 2004[40, 41]. The reaction was made at 40°C in 75 mM phosphate buffer (pH 7.4). The final assay mixture (200 mL) contained FL (70 nM), 173 174 AAPH (14 mM), and antioxidant [Trolox (0.2-1.6 nmol) or samples of the five synthetic peptides (at different concentrations)]. The fluorescence was recorded during 137 min 175 176 (104 cycles). A FLUOstar OPTIMA plate reader (BMG Labtech, Offenburg, Germany) 177 with 485 nm excitation and 520 nm emission filters was used. The equipment was 178 controlled by the FLUOstar Control software version (1.32 R2) for fluorescence 179 measurement. Black polystyrene 96-well microplates (Nunc, Denmark) were used. 180 AAPH and Trolox solutions were prepared daily and FL was diluted from a stock 181 solution (1.17 mM) in 75Mm phosphate buffer (pH 7.4). All reaction mixtures were 182 prepared in duplicate and at least three independent runs were performed for each 183 sample. Final ORAC-FL values were expressed as µmol of Trolox equivalent/ µmol of 184 peptide [42].

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186 Thiobarbituric acid reactive substances (TBARS)

The thiobarbituric acid reactive species method was used as described by Westerfield,
1995 [43]. The Zebrafish colony was established in the laboratory, in a glass aquarium,

189 containing an internal filter and an activated carbon aerator for water oxygenation. 190 The population of animals was fed three times a day with food chips for fish. Adult fish 191 were kept on 16 hours light and 10 hours dark cycles. Embryos were obtained by 192 photo-induced spawning over green plants and cultured at 28°C in a fish tank water. 5 193 days post fecundation (dpf) larvae were then incubated in 24-well plates, 30 larvae per 194 well, with 50 µg/ml of lysozyme peptide in each well. Lipid peroxidation was initiated by adding 1 ml 500 μ M H₂O₂ and incubated during 8 hours at 28°C. Groups from 30 195 196 larvae/well in aquarium water were used as controls. Then, H_2O_2 was removed with a micropipette and 500 μl of Tween 0.1% was added. All groups were mixed and 197 198 homogenized with a T25 Ultra turrax IKA. Then, absorbance of the solution of zebrafish 199 larvae and peptides was measured at 532 nm. The decrease of absorbance indicates an 200 increase of antioxidant activity. The values of antioxidant activity were expressed as 201 the percentage inhibition of lipid peroxidation in larvae homogenate as follows:

202 The total antioxidant activity % Inhibition of lipid peroxidation = $[(A_b - A_s)/A_b = X 100]$

where A_b is the absorbance of blank and A_s is the absorbance the sample.

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205 Test of toxicity in the Zebrafish model

Zebrafish of the AB strain (wild-type, wt) embryos were obtained from natural spawning. Embryos were raised and fish were maintained as described by Westerfield, (1995) [43]. After collection and disinfection, eggs were placed in 24-well microplates with 1 mL of water. To study the in vivo toxicity of all peptides coming from lysozyme with the zebrafish model, the FET test was employed.

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212 FET Test

The assay was based on the OECD draft guideline on Fish Embryo Toxicity (FET) Test [44] and is described in detail by (Domingues et al., 2010) [45]. The Test Guideline is based on chemical exposure of newly fertilized zebrafish eggs for up to 48 hours and is expected to reflect acute toxicity in fish in general. After 24 and 48 hours of exposure to the peptides, four apical endpoints were recorded as indicators of acute lethality in fish: coagulation of fertilized eggs, lack of somite formation, lack of detachment of the tail-bud from the yolk sac and lack of heart-beat. The eggs were considered dead whenthey exhibit at least one of the previous mentioned indicators.

In the control wells, there should be less than 10% of the eggs with one of the mentioned indicators after 48 hours, (29) [46]. Ten eggs per treatment (3 replicates) were selected and distributed in 24-well microplates. The test started with newly fertilized eggs exposed to the nominal concentrations of 50; 156; 312; 625; 1250; 2500 and 5000 μ g/ml of peptides and run during 2 days. Embryos were observed at 24 and 48 hours under a stereomicroscope (magnification used in the stereomicroscope for observations was 40X).

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229 Results and discussion

230 Hydrolysis in situ from lysozyme with pepsin in a cation-exchange column.

Hen egg white lysozyme was subject to hydrolysis in situ with pepsin in a cation-231 232 exchange column (Figure 1). The objective was to obtain, in one step, rich peptides 233 with positively charged amino acids derived from the hydrolysis of lysozyme and, to assess whether denatured lysozyme could generate peptides other than native 234 lysozyme. It is known that lysozyme has resistance to the hydrolysis with pepsin, but it 235 236 has been recently described that lysozyme at pH 1.2 has total susceptibility to the 237 hydrolysis with pepsin [39, 47-49]. Fu, Abbott, and Hatzos (2002) [50] have reported 238 that lysozyme resisted more than 60 minutes at pH 1.2, at an E: S of (13:1) wt:wt. 239 Thomas et al., (2004) [49] described that hen egg white lysozyme is resistant to 240 hydrolysis with pepsin at pH 2.0. Ibrahim et al. (2005)[51] found that 40% of the original lysozyme was hydrolyzed after 120 minutes of digestion at an E: S of 1:50 241 (wt:wt) and pH 4.0. There is then controversy about the hydrolysis of hen egg white 242 lysozyme and this can be due to the different methods used. In this study, lysozyme 243 was hydrolyzed at pH 2.0 with an excess of pepsin. 244

245 Many antimicrobial and antioxidant peptides contain positively charged amino acids 246 thus determining the strength of their activity [21, 52, 53]. Lysozyme has an isoelectric 247 point of 10.7 with a high content of positively charged amino acids. Hen egg white 248 lysozyme has 17 positively charged (6 Lys, 11 Arg) and nine negatively charged residues 249 (7 Asp, 2 Glu), thus leading to a net positive charge at pH below the isoelectric point (10.7). This positive charge makes hen egg white lysozyme even more attractive forinvestigation with the negative charged peptides [54].

Therefore, lysozyme may be a good substrate for production of antimicrobial and 252 253 antioxidant peptides. You et al., 2010 [53] have reported antioxidant hydrolysates from lysozyme obtained with alcalase. They found that the fractions were rich in 254 255 cationic peptides with high percentage of Arg and Lys (positively charged amino acids). Samples of native and denatured lysozyme were loaded in the cation-exchange 256 257 column. Then, those samples were treated over night at 37°C with recirculation of 258 pepsin solution. Immediately after, the hydrolysate was eluted with a gradient of 3 M 259 and 5 M of ammonia. Two different fractions were successively collected respectively, 260 and then a third fraction was eluted with sodium chloride (NaCl) 1 M. This fraction 261 contained the peptides with maximum net positive charge and, therefore, those with the highest affinity for the cation-exchange column (Figure 2). 262

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264 Identification of peptides sequences

265 After treatment with dithiothreitol all fractions from native and denatured lysozyme were analyzed with RP-HPLC-ESI-MS-MS to characterize their molecular mass and 266 267 amino acid sequences. Table 1 shows the identified peptides eluted with NaCl 1 M. The peptides from 3 M and 5 M ammonia fractions were discarded as those peptides are 268 269 less cationic. Both fractions were very complex with high content of peaks. As 270 expected, this fraction contained peptides with abundant positively charged amino 271 acids (Arg and Lys). For this reason, it was decided to work with this fraction. All 272 sequences are located in the C-terminus of lysozyme, in the α -dominium in the zone of 273 helix 90-129. It can be seen that there was no difference between the peptides 274 identified in both hydrolysates, indicating that the process of heat denaturation of 275 lysozyme results in a lack of production of new hydrolysis sites in the protein. Ibrahim 276 et al., (2001, 2005) [51, 55] have reported antimicrobial peptides with high activity 277 present in the α -dominium (1-40 and 90-129), specifically in the regions 1-38 and 87-278 114 from the lysozyme. However, we only found peptides located in the α -dominium 279 C-terminal 90-129. This could be caused by the ionic separation performed, where cationic peptides have predominantly been recovered. The peptides identified were 280 281 synthesized to be used in the antioxidant assay.

282 Antioxidant peptides sequences

283 Five peptides from the fraction NaCl were assayed for their antioxidant activity, against peroxyl radicals, by using ORAC-FL assay. Table 2 shows results of antioxidant activities 284 285 for the five peptides assayed. ORAC-FL values of peptides from hen egg white lysozyme were very high, indicating very high antioxidant activity. Four peptides f(109-119) 286 287 VAWRNRCKGTD, f(111-119) WRNRCKGTD, f(122-129) AWIRGCRL and f(123-129) WIRGCRL (1,970; 3,123; 2,743 and 2,393 µmol Trolox/ µmol peptide) respectively were 288 289 more active than vitamin C (1.65 μ mol Trolox equivalents/ μ mol vitamin C). Peptides 290 f(122-129) AWIRGCRL and f(123-129) WIRGCRL were more active than synthetic 291 antioxidant peptide named butylated hydroxyanisole (BHA) (2,430 µmol Trolox 292 equivalents/ μ mol BHA) used in food industry for its high antioxidant activity [56,42]. 293 Only the peptide f(124-129) IRGCRL presents low activity with 0.313 μ mol Trolox/ μ mol peptide, this might be due to the absence of Trp in its sequence. As shown in the ORAC 294 295 database prepared by Li and Li (2013) [57], the length of peptides derived from food 296 sources with peroxyl radical scavenging activity, ranges from 4 to 20 amino acids. 297 Peptides described in our study with peroxyl radical scavenging activity have between 1 to 11 amino acids. Peptides of our study are small peptides with high antioxidant 298 299 activity using ORAC-FL. On the other hand, Hernández-Ledesma et al., (2005) [58] have 300 described a peptide from soybean named lunasin with high antioxidant activity 3.44 \pm 301 0.07 μ mol Trolox equivalents/ μ mol lunasin. Potent activity of lunasin was attributed to 302 the presence of amino acids Trp, Cys, and Met in its sequence. All peptides in our study 303 contain in their sequence Trp, Cys or both. Possibly, the higher antioxidant activity of 304 peptides in this study is explained by the presence of Trp and Cys amino acids in a 305 particular site of their sequence. Peptide number 2 IRGCRL was compared to peptide 306 number 3 WIRGCRL. We observed that IRGCRL peptide has not Trp (W) in its sequence, 307 and a value of ORAC-FL of 0.311 µmol Trolox/ µmol peptide was detected. On the 308 other hand, peptide number 3 WIRGCRL presented antioxidant activity with a value of 309 2,393µmol Trolox/ µmol peptide. The difference in the antioxidant activity can be 310 related to the presence of Trp in peptide number 3 WIRGCRL sequence.

The molecular weight of the identified peptides are in the range of most of the antioxidant peptides derived from food sources isolated previously of 4 to 20 residues amino acids [59]. Moreover, antioxidant peptides often possess hydrophobic amino

- acid residues such as Pro, His, Tyr, Trp, Met, or Cys in their sequences and Val or Leu at
- the N-terminus [60]. One of our peptides showed Val at the N-terminus.
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317 Potential of synthetic peptides from lysozyme to inhibit lipid peroxidation

318 The antioxidant action is assessed by inhibiting the damage caused by free radicals and 319 the mechanisms involved in many human diseases such as hepatotoxicities, 320 hepatocarcinogenesis, diabetes, and skin cancer to include lipid peroxidation as a main 321 source of cellular damage. Lipid peroxidation in biological systems has been thought to 322 be a toxicological phenomenon leading to various pathological consequences. MDA formed from lipid peroxidation of unsaturated phospholipid reacts with TBA to 323 324 produce a pink MDA-TBA adducts. MDA is reactive and active in crosslinking with DNA 325 and proteins and damages liver cells [61]. Phospholipids are believed to be present in 326 high amounts in cell membranes [62]. Lipid peroxidation has been a major contributor 327 to the loss of cell function under oxidative stress [63, 64]. To determine oxidative 328 stress, inhibition of lipid peroxidation in zebrafish larvae model was used to determine 329 damage cellular in vivo. Figure 3A presents the inhibition of lipid peroxidation by synthetic peptides from lysozyme at a concentration of 50 µg/ml. This assay confirmed 330 331 that these synthetic peptides were not toxic for zebrafish larvae. Zebrafish larvae presented normal aspect after 24 hours of assay. When zebrafish larvae were 332 examined, no morphological abnormalities are shown such as crooked bodies, spinal 333 334 deformities or any significant effects in the growth of the body (Figure 3B). The values 335 of percentage inhibition of lipid peroxidation indicated that all synthetic peptides were 336 efficient to inhibit the lipid peroxidation in zebrafish larvae. For example peptide P4 337 (AWIRGCRL) had a result of 63.2 % TBARS inhibition, (Figure 3A). The antioxidant 338 results showed the higher activity of peptides 3, 4 and 5 in both assays. The presence 339 of tryptophan seems important for the ORAC activity. However, in the case of the 340 TBARS inhibition, the peptide size is probably contributing to the increased values of 341 activity in peptides P2 to P5.

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345 Test in Zebrafish embryo

Based on the in vitro studies antioxidant activity described above, we decided to evaluate the toxicity of the peptides in a model of zebrafish eggs. Zebrafish has become a widely used model organism for studies of developmental biology and drug discovery. This model helps drug development by combining the tools of medicinal chemistry and zebrafish biology.

351 Figure 4 shows a representative curve doses-response result of FET test for AWIRGCRL 352 peptide, this peptide presents the highest TBARS inhibition percentage. This sample 353 was only taken as an example as all samples presented identical results. The test was carried out for all peptides of this study, however, no significant differences were 354 355 observed in the rest of peptides. Again, as in the zebrafish larvae test, the AWIRGCRL 356 peptide does not present toxicity at a concentration of 50 μ g/ml for zebrafish eggs. 357 However, concentrations higher than 50 µg/ml of peptides were cytotoxic to zebrafish 358 egg after 24 hours of incubation. Mortality was identified with an absence of 359 embryonic development and coagulation of nuclear material of eggs. The control eggs 360 were totally normal in their development (Figure 5A). However, eggs treated with the 361 peptides of this study presented no embryonic development and coagulation total of nuclear material of eggs. Around the eggs, material of the chorion due to ruptures of 362 363 the eggs was observed (Figure 5B).

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365 As a conclusion, hen egg lysozyme was hydrolyzed with pepsin in situ using a cationexchange. Hen egg white lysozyme is a good source of antioxidant peptides using 366 pepsin for hydrolysis at low pH. The zebrafish model was efficient to measure the 367 inhibition of lipid peroxidation and cytotoxicity of synthetic peptides from lysozyme. 368 369 Development of zebrafish is sensitive to the exposure to all lysozyme synthetic 370 peptides used in this study at concentrations higher than 50µg/ml. However, further 371 investigations would need to be carried out to evaluate the death mechanisms of 372 these peptides on zebrafish embryos, for eventual pharmaceutical and medical 373 applications.

374

375 **Conflict of interest**

The authors declare that they have no conflict of interest.

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- 622 List of figures and tables
- **Table 1. Identification of the molecular mass and amino acid sequence of the fraction eluted**
- 624 with 1 M NaCl using LC-ESI-MS/MS.

N°	Fragment	Mass	Mass Calc ^a	m/z ^b	Sequence ^c	NLZ	LZ95°C
		0.055.	Calc.				
1	f(109-119)	1306,0	1306,0	653,5 (2)	VAWRNRCKGTD	+	+
2	f(124-129)	716,6	717,6	717,6 (1)	IRGCRL	+	+
3	f(123-129)	902,7	903,7	903,7 (1)	WIRGCRL	+	+
4	f(122-129)	973,7	974,7	974,7 (1)	AWIRGCRL	+	+
5	f(111-119)	1134,7	1135,7	1135,7 (1)	WRNRCKGTD	+	+

627 Table 2. In vitro antioxidant activity of peptides from lysozyme.

		628
N⁰	Peptides	ORAC (micromol Trolox
		equivs/micromol peptide) ± SD
1	VAWRNRCKGTD	1.970 ± 0.171 630
2	IRGCRL	0.313 ± 0.029 631
3	WIRGCRL	2,393 ± 0.280 632
4	AWIRGCRL	2,743 ± 0.193 634
5	WRNRCKGTD	3,123 ± 0.266
		635







649 **oven at 37°C over night.**

650







654 Figure 3. A) TBARS result of synthetic peptides from lysozyme. Data is expressed as % TBARS

655 inhibition compared to positive control (error bars expressed as ± SD). P1= VAWRNRCKGTD;

656 P2= IRGCRL; P3= WIRGCRL; P4= AWIRGCRL; P5= WRNRCKGTD. B) Photography of zebrafish

657 larvae with peptide and without peptide after assay. All peptides were assay at 50 μg7ml.

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660 Figure 4. Mortality percentage of Zebrafish embryo treated with AWIRGCRL peptide from

661 Iysozyme at different concentrations at 48 hours for three replicates.



Figure 5. Peptides from lysozyme induced inhibition cellular in embryo Zebrafish. A) Control
without peptides and B) embryo with AWIRGCRL peptide from lysozyme. Magnification was
of 40X. AWIRGCRL peptide was incubated with zebrafish eggs during 24 hours at 26°C.

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