Food-inspired peptides from spinach Rubisco endowed with antioxidant, antinociceptive and anti-inflammatory properties

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22	ABSTRACT: Rubiscolin-6 (amino acid sequence: YPLDLF) is a selective δ -opioid receptor peptide
23	isolated from spinach Rubisco. Its synthetic analogue, peptide YPMDIV is the most potent described
24	so far for its increased opioid activity, thus in this work it was considered as lead compound for the
25	design of twelve new analogues e.g. LMAS1-12. Firstly all the novel compounds have been tested
26	for their antinociceptive and anti-inflammatory capacity in vitro and in vivo in order to evaluate their
27	ability to maintain or loss the original activity. Among them peptides LMAS5-8 gave the best results,
28	thus their antioxidant properties have been investigated along with their enzymatic inhibitory ability.
29	Peptide LMAS6 shows a strong antioxidant (154.25 mg TE/g CUPRAC) and inhibitor activity on
30	tyrosinase (84.49 mg KAE/g), indicating a potential role in food industry as anti-browning agent,
31	while peptides LMAS5 and LMAS7 possess a modest cholinesterase inhibitory activity suggesting

32 a conceivable use for nutraceuticals production.

33 **KEYWORDS:** rubiscolin-6, Rubisco, antioxidants, anti-browning agent, nutraceuticals

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44 **1. INTRODUCTION**

Plant and animal food proteins are considered rich sources of bioactive peptides, which are classified 45 on the basis of their biological effects, e.g., antithrombotic, antimicrobial, antihypertensive, opioid, 46 immunomodulatory and antioxidant (Sánchez & Vázquez, 2017; Udenigwe & Aluko, 2012). Given 47 their small size, sample preparation's process, high potency and selectivity, they are an excellent 48 example of tools to inhibit protein-protein interactions. Food derived bioactive peptides may play an 49 important role in the regulation of various human functions acting directly through food's 50 consumption or after in vivo or in vitro hydrolysis of the proteins in which they are encrypted 51 (Hartmann & Meisel, 2007). In fact they are generally a mixture of small size-peptides because of the 52 high cost of purification and low yield of extraction; after purification, some of them may lose their 53 additive or synergistic activity with polyphenols or other food components, resulting in inactive 54 compounds that are useless as nutraceuticals. However, the literature reports examples of isolated 55 bioactive compounds with proven physiological effects that can be added to food products (Day, 56 Seymour, Pitts, Konczak, & Lundin, 2009). For instance, antioxidant peptides derived from the 57 digestion process of different proteins may be added to food products to provide them antioxidant 58 benefits (Lafarga & Hayes, 2014). Indeed, bioactive peptides with antioxidant properties have been 59 used as free radical scavengers in meat products (Sohaib et al., 2017; Tkaczewska, 2020). The 60 antioxidant effect of protein hydrolysates is related to their amino acids content (Sánchez et al., 2017). 61

Rubiscolin-6 (amino acid sequence: YPLDLF) is an interesting natural peptide recently evaluated for 62 its antioxidant activity; it was identified in the spinach Rubisco large subunit (residues 103-108) 63 (Stefanucci et al., 2020; Yang et al., 2001). Rubisco (D-ribulose-1,5-bisphosphate 64 carboxylase/oxygenase; EC 4.1.1.39) represents the 10-30% of the whole amount of leaves proteins. 65 It is composed of two subunits A and B, combined in a type-structure A₈B₈ (Kobbi et al., 2017; Taylor 66 & Andersson, 1997; Yoshikawa, 2015). This enzyme plays an important role in carbon dioxide 67 fixation and photorespiration (Yoshikawa, 2015), and it is involved in food's production for the whole 68 69 biosphere. A recent study showed that Rubisco hydrolysate from alfa-lfa green juice has significant antioxidant properties preventing linoleic acid oxidation, decreasing ferric ion and inducing stable 70 ABTS^{*+} compared to the unhydrolysate protein (Kobbi et al., 2017). 71

Rubiscolin-6 is a selective δ-opioid receptor peptide agonist highly conserved in a huge variety of plants (Yang et al., 2001). It exhibits several effects, *e.g.*, antinociceptive activity, anxiolytic-like

effect, orexigenic action, enhancement of glucose uptake in skeletal cell lines, memory consolidation 74 and reduction of skin inflammation (Chajra, Amstutz, Schweikert, Auriol, Redziniak, & Lefevre, 75 2015; Hirata, Sonoda, Agui, Yoshida, Ohinata, & Yoshikawa, 2007; Kairupan et al., 2019; Kaneko 76 et al., 2012; Miyazaki et al., 2014; Yang, Kawamura, & Yoshikawa, 2003; Yang et al., 2001). 77 Numerous rubiscolin-6 analogues have been already synthesized to evaluate its structure-activity 78 79 relationships in order to improve its δ -opioid activity (Yang, Sonoda, Chen, & Yoshikawa, 2003). The structural features of rubiscolin-6 analogues required to guarantee a high δ -opioid activity have 80 been also determined through 3D quantitative structure-relationship (QSAR) analysis (Caballero, 81 Saavedra, Fernández, & González-Nilo, 2007). Thus researchers have considered the hypothesis of 82 using rubiscolin-6 and its derivatives for the production of functional foods and as lead compounds 83 for the development of new analogues (Stefanucci et al., 2020). 84

Rubiscolin-6 and its C-terminal amide analogue have been recently tested in vitro for their antioxidant 85 activity, however rubiscolin-6 C-amide showed low DPPH radical scavenging activity (2.72 mg 86 TE/g), while rubiscolin-6 gave the best result in ABTS assay (8.86 mg TE/g) (Stefanucci et al., 2020). 87 Among the novel analogues, YPMDIV showed an increased δ -opioid activity on mouse vas deferens 88 (MVD) assay in comparison with rubiscolin-6 (Yang, Sonoda, et al., 2003). In the current study, 89 YPMDIV has been assumed as *lead compound* for the design of twelve new analogues named 90 LMAS1-12 (Figure 1), with the aim to find bioactive peptides endowed with antioxidant and enzyme 91 inhibitory activities potentially useful as food ingredients or in the development of new nutraceuticals. 92

93 **2. MATERIALS AND METHODS**

94 2.1 Design

A rational structure-based design approach has been applied to the development of novel peptide 95 analogues of rubiscolin-6. Considering the multi-residue replacement described by a SAR study in 96 97 literature (Yang, Sonoda, et al., 2003), we selected the synthetic peptide YPMDIV as the lead compound for further structural modifications. The following findings have been taken in 98 consideration in order to design the novel bioactive peptides: *i*) the protonated Tyr at the *N*-terminus 99 fits the essential requirement of message domain for opioid receptor binding; *ii*) the Pro² residue 100 contributes to the restriction of peptide conformation and stability; *iii*) the presence of Met³ enhances 101 the opioid activity in MVD about four times, while Ala³ and Val³, Phe³ and Trp³ decreased the activity 102 in MVD assay; iv) Asp⁴ is the ideal residue for high activity and selectivity on δ -opioid receptor 103 (DOR); v) in the fifth position of Rubiscolin-6, hydrophobic residues increase the potency at DOR, 104 the preferential order is Leu > Ile > Met > Val > Ala; vi) the Phe⁶ in rubiscolin-6 can be replaced by 105 aliphatic residue without losing δ -opioid activity. In a first attempt we explored the substitution of 106 Met³ in the *lead compound* with an unnatural residue of (2S,4S)-4-(methylthio)pyrrolidine-2-107 carboxylic acid (Proline-Methionine chimeras) prepared in laboratory following the procedure 108 previously described by Mollica et al. (Mollica, Paradisi, Varani, Spisani, & Lucente, 2006) linked 109 to both D/L-Pro² (compounds LMAS1-4), with the aim to improve the metabolic stability of the 110 former peptides as C-terminal acids and amides. Since methionine is one of the most documented 111 amino acid responsible of antioxidant activity in several peptides (Lorenzo et al., 2018), the 112 replacement with Cys and the shortening of the Met³ side chain have been also investigated in 113 peptides of the series LMAS5-8 and LMAS9-12 respectively. According to the SAR studies (Yang, 114 Sonoda, et al., 2003), all the other amino acids in position first, fourth, fifth and sixth have been 115 conserved. These modifications are intended to understand the role exerted by this amino acid in 116 varying the biological activity of such modified analogues of rubiscolin-6. The lead compound has 117 been prepared as reference for the biological assays (Figure 1). 118



Tyr-Pro-Met-Glu-Ile-Val-OH



LMAS1 sequence: Tyr-Pro-(2*S*,4*S*)-4-(methylthio)pyrrolidine-2-carboxylic acid-Glu-Ile-Val-NH₂ LMAS2 sequence: Tyr-pro-(2*S*,4*S*)-4-(methylthio)pyrrolidine-2-carboxylic acid-Glu-Ile-Val-NH₂ LMAS3 sequence: Tyr-Pro-(2*S*,4*S*)-4-(methylthio)pyrrolidine-2-carboxylic acid-Glu-Ile-Val-OH LMAS4 sequence: Tyr-pro-(2*S*,4*S*)-4-(methylthio)pyrrolidine-2-carboxylic acid-Glu-Ile-Val-OH



LMAS5 sequence: Tyr-Pro-Cys-Glu-Ile-Val-NH₂ LMAS6 sequence: Tyr-pro-Cys-Glu-Ile-Val-NH₂ LMAS7 sequence: Tyr-pro-Cys-Glu-Ile-Val-OH LMAS8 sequence: Tyr-Pro-Cys-Glu-Ile-Val-OH



LMAS9 sequence: Tyr-Pro-(S)-2-amino-3-(methylthio)propanoic acid-Glu-Ile-Val-NH₂ LMAS10 sequence: Tyr-pro-(S)-2-amino-3-(methylthio)propanoic acid-Glu-Ile-Val-NH₂ LMAS11 sequence: Tyr-Pro-(S)-2-amino-3-(methylthio)propanoic acid-Glu-Ile-Val-OH LMAS12 sequence: Tyr-pro-(S)-2-amino-3-(methylthio)propanoic acid-Glu-Ile-Val-OH

120 Figure 1. Structure of the *lead compound* (YPMDIV) and peptides LMAS1-12.

121 *2.2 Chemistry*

119

Coupling reagents and solvents have been purchased by VWR (Radnor, PN, USA). Fmoc-Tyr(t-Bu)-122 OH, Fmoc-Pro-OH, Fmoc-D-Pro-OH, Fmoc-Met-OH, (S)-2-amino-3-(methylthio)propanoic acid, 123 Fmoc-Cys(Trt)-OH, Fmoc-Asp(t-Bu)-OH, Fmoc-Ile-OH and Fmoc-Val-OH by Chem-Impex (Wood 124 Dale, IL, USA). The synthetic amino acid residue (2S,4S)-4-(methylthio)pyrrolidine-2-carboxylic 125 acid has been synthesized as previously reported in literature (Mollica, Paradisi, Varani, Spisani, & 126 Lucente, 2006). Commercial amino acid (S)-2-amino-3-(methylthio)propanoic acid has been 127 protected with fluorenylmethoxycarbonyl protecting group (Fmoc) before its use (Carpino and Han, 128 1979). Fmoc-Rink amide and 2-chlorotrityl chloride resins (Loading coefficient: 0.74 mmol/g and 129 130 1.60 mmol/g respectively) by IRIS Biotech GmbH (Marktredwitz, DH, Germany). Peptides have been prepared by manual solid-phase synthesis (SPPS) using TBTU/HOBt/DIPEA and TBTU/2,4,6-131 trimethylpyridine for Fmoc-Cys(Trt)-OH reaction, as coupling mixture; a solution of piperidine 132 20%/DMF for Fmoc-deprotection. A strong cleavage has been applied to the peptide-resin system 133

using TFA/TIPS/H₂O for LMAS1-4,9-12 or TFA/TES/H₂O for LMAS5-8 as cocktails (95:2.5:2.5) 134 (Scheme S1, see SI). The final peptides as TFA salts have been purified by RP-HPLC system 135 equipped with Waters XBridge Prep BEH130 C18, 5.0 µM, 250-10 mm, 7.0 mL/min, Waters Binary 136 1525 pump, H₂O + 0.1% TFA/ACN + 0.1% TFA gradient from 5% to 95% of ACN in 35 minutes 137 (wavelengths: 213 nm, 254 nm and 275 nm). Peptides purity as TFA salts was assessed using ¹H-138 NMR (Varian Inova 300MHz) in DMSOd₆ and RP-HPLC analytic column C-18, 4.6-150 mm, 1 139 mL/min, H₂O + 0.1% TFA/ACN + 0.1% TFA gradient from 5% to 95% of ACN in 30 minutes. Mass 140 spectrometer LCQ (Thermo Finnigan, San Jose, CA, USA) with capillary temperature 300°C, 141 electrospray ion Source (ESI), 4.00 Kv, auxiliary and preservative gas N₂ has been used to confirm 142 the compound's identity. The purity of all the final peptides resulted to be \geq 95% after purification 143 (see SI). 144

145 *2.3 Opioid binding assay and GTP stimulation*

146 The opioid receptor binding affinity has been calculated performing displacement assay on μ -opioid 147 receptor (MOR), δ -opioid receptor (DOR) and κ -opioid receptor (KOR) on rat and guinea pig brain 148 membrane homogenates. For the detailed procedures see the supporting information.

149 *2.4 In vitro Antioxidant assay*

Methodologies applied in our previous literature have been used (Uysal et al., 2017), considering the following parameters: mg Trolox equivalents (TE)/g extract in 2,2-diphenyl-1-picrylhydrazyl (DPPH) and 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulfonic acid) (ABTS) radical scavenging, cupric reducing antioxidant capacity (CUPRAC) and ferric reducing antioxidant power (FRAP) tests, and mmol TE/g extract in phosphomolybdenum assay.

155 2.5 In vitro Enzymatic inhibitory assay

The enzyme inhibitory assays were performed following the procedure reported in literature (Zengin, 2016). The following parameters have been choosen: acetylcholinesterase (AChE) and butyrylcholinesterase (BChE) inhibition in mg galanthamine equivalents (GALAE)/g extract; tyrosinase inhibition in mg kojic acid equivalents KAE/g extract; amylase and glucosidase inhibition in mmol acarbose equivalents (ACAE)/g extract.

161 *2.6 In vivo antinociceptive assays*

Tail flick and formalin tests, and zymosan-induced edema formation have been carried out following the procedures previously decribed (Pieretti, Saviano, Mollica, Stefanucci, Aloisi, & Nicoletti, 2022; Della Valle et al., 2021) using groups of 7-8 animals. The experimental protocols performed in the present study were in accordance with Italian Legislative Decree 27/92 and approved by the local ethics committee (approval number: 198/2013-B). Detailed procedures are described in the Supporting Information.

168 2.7 Data analysis and statistics

169 Data obtained from *in vivo* experiments are reported as mean \pm s.e.m. Data analysis of tail flick and 170 formalin test was carried out using one-way ANOVA followed by Dunnett's multiple comparisons 171 test. Data from zymosan-induced paw edema experiments were analysed using two-way ANOVA 172 followed by Dunnett's multiple comparisons test. GraphPad Prism 9.0 software (San Diego, CA, 173 USA) for data elaboration. Means are considered statistically significant at P \leq 0.05.

174 **3. RESULTS AND DISCUSSION**

175 *3.1 Chemical synthesis*

The novel peptides as TFA salts were prepared following a straightforward and highly efficient 176 solution phase peptide synthesis protocol, both as C-terminal amides and acids (Table S1, Scheme S1 177 see SI). Their amino acid sequences and overall yields are shown in Table S1 (see SI). All peptides 178 contain a L or D-proline in position 2 in combination with (2S,4S)-4-(methylthio)pyrrolidine-2-179 carboxylic acid (LMAS1-4), cysteine (LMAS5-8) or (S)-2-amino-3-(methylthio)propanoic acid 180 (LMAS9-12) in third position. The synthetic procedure applied in this work allowed us to optimize 181 the reactions time, reagent/solvents costs and the purification steps since the crude final products 182 present less detectable by-products in RP-HPLC chromatographic traces, they are completely soluble 183 in water/methanol medium and they are easier to purify than those obtained by solution phase peptide 184 185 synthesis.

186 *3.2 C*

3.2 Opioid binding assay

All the final peptides and the *lead compound* have been tested for their ability to bind the MOR, DOR and KOR in presence of the reference ligands [³H]DAMGO, [³H]Ile^{5,6}-delthorphin II and [³H]HS665 respectively (Figure S1, Table S2, see SI). Peptide **LMAS4** seems to be selective for MOR with a moderate binding affinity (Ki: 137.4 \pm 0.15 nM). All the other compounds are not able to bind the three opioid receptors at 10 μ M concentration. It's worth to note that none of them exhibits a significant binding affinity for δ -opioid receptor, for which the *lead compound* shows a IC₅₀ value of 0.12 μ M in [³H]DPDPE binding assay (Yang, S., Sonoda, S., Chen, L., & Yoshikawa, M., 2003).

187 This result is not surprising since the sequence D-Pro-(2S,4S)-4-(methylthio)pyrrolidine-2-carboxylic 188 acid (proline-methionine chimera) contained in LMAS4 has been already reported to confer an 189 increased metabolic stability to plasma and enzyme's degradation (Mollica et al., 2012), as well as to 190 be responsible of an improved μ -opioid receptor selectivity, thanks to the ability of such sequence to 191 assume a well-defined 3D structure (Stefanucci et al., 2011).

3.3 GTP stimulation assay

In the GTP stimulation assay, all the peptides were tested to evaluate their capacity to stimulate the 193 G-protein coupled receptor (Figure S2, Table S2 see SI). Their efficacy almost stayed around the 194 basal activity, with the only exception of compound LMAS4 (E_{max} : 139.8 \pm 2.8%). In agreement with 195 the binding data, this analogue of rubiscolin-6 is more efficacious and potent in GTP stimulation 196 assay than the *lead compound*, which is in turn about 20 times more potent than rubiscolin-6 in MVD 197 assay (EC₅₀ value of 5.65 µM) (Yang, S., Sonoda, S., Chen, L., & Yoshikawa, M., 2003). The novel 198 analogue shows a diverse affinity/selectivity profile in vitro and a more potent antinociceptive activity 199 than rubiscolin-6 and its analogue previously described by Yang et al. 200

- 201 *3.4 In vivo experiments*
- 202 *3.4.1 Tail flick test*

In the first series of experiments, the effects of the *lead compound* and LMAS1-12 peptides were investigated in an animal model of acute nociception induced by thermal stimuli as the tail flick tests. Peptides were injected via intracerebroventricular route (i.c.v.) at the dose of 10 μ g/mouse (Figure 2). Peptides LMAS1-12 induce a greater antinociceptive effect than vehicle-treated animals, which reaches statistical significance for LMAS1,5,7,8,12. All of them also increased antinociceptive response times in comparison with *lead compound*-treated animals. Statistical analysis reveals that this effect is significant in LMAS5,7,8,12 treatment groups. All of them exert an antinociceptive

- effect in this assay higher than that of rubiscolin-6 and its *C*-terminal amide derivative previously
- described by us (Stefanucci et al., 2020), which could be due to an improved enzymatic stability at a
- 212 central level.



Tail Flick test

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Figure 2. Antinociceptive effects induced by the *lead compound* (LC) and LMAS1-12 peptides in the tail flick test. Peptides were administered i.c.v. at the dose of 10 μ g/mouse. Tail flick data were reported as area under %MPE curve (AUC). * is for P<0.05, ** is for P<0.01 and *** is for P<0.001 vs vehicle-treated animals (V, DMSO 0.1% in saline); ° is for P<0.05, °° is for P<0.01 and °°°° is for P<0.001 vs LC-treated animals. N=7.

218 *3.4.2 Formalin test*

In the formalin test, the *lead compound* and LMAS1-12 peptides were administered subcutaneously 219 (s.c.) into the mice hind paw at a dose of 100 µg/mouse, 15 min before formalin (Figure 3). In the 220 early phase of the formalin test, LMAS3,4,5,6,7,9,11 reduced the licking behaviour induced by 221 formalin, although only the antinociceptive effects of LMAS5-7 peptides reach statistical 222 significance in comparison with vehicle treated animals (Figure 3, left panel). In the late phase of the 223 formalin test, LMAS1,2,4-9 reduced formalin-induced nociceptive effect, even if only the LMAS4 224 peptide is able to significantly reduce the licking activity induced by the aldehyde (Figure 3, right 225 panel). These results partially support our findings described in paragraph 3.2, where peptide LMAS4 226 exerted the best binding affinity for u-opioid receptor probably due to an improved plasma or 227 metabolic stability. On the contrary, the antinociceptive effect of this peptide and analogues LMAS5-228 7 is lower than that of rubiscolin-6 C-terminal amide, previously described by us (Stefanucci et al., 229 2020) in the early phase of the formalin test, while it appears to be stronger than the parent compound 230 in the late phase. 231



Figure 3. Antinociceptive effects induced by the *lead compound* (LC) and LMAS1-12 peptides in the formalin test. In the left panel, the result obtained in the early phase of the test are reported. The right panel reports the licking activity recorded in the late phase of the formalin test. Peptides were administered s.c. into the hind paw at the dose of 100 μ g/mouse. * is for P<0.05, ** is for P<0.01 and **** is for P<0.001 *vs* vehicle-treated animals (V, DMSO:saline, ratio 1:3 *v/v*). N=7.

238 *3.4.3 Zymosan-induced edema formation*

Since some of the tested peptides show the best in vivo antinociceptive profile, we decided to study 239 the possible anti-inflammatory effects of LMAS6,7 and lead compound (Figure 4). All of them 240 induced an anti-inflammatory effect, although a significant reduction of edema volume for the lead 241 compound was observed after 2 hours and for the LMAS7 peptide, 1 and 2 hours after zymosan 242 administration. The LMAS6 peptide appears to be the most active, as it was able to significantly 243 reduce the formation of edema for the entire duration of the observation period. This result pairs with 244 the data obtained by the formalin test in the early phase after subcutaneous administration, leading us 245 to suppose a possible analgesic activity at the periphery, however further investigation is required to 246 support such hypothesis raising by an embryonic stage of the work. 247

Zymosan-induced edema



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Figure 4. Anti-inflammatory effects induced by the *lead compound* (LC) and LMAS6,7 peptides. Inflammatory paw edema was induced by zymosan, and peptides were administered s.c. at the dose of 100 μg/mouse 30 min before. The increase in paw volume was evaluated as the percentage difference between the paw volume at each time point and the

basal paw volume. * is for P<0.05 and **** is for P<0.0001 vs vehicle-treated animals (V, DMSO:saline, ratio 1:3 v/v). N=8.

Overall, the cluster of peptides LMAS5-8 exhibits the best in vivo antinociceptive results. 254 Surprisingly they are not able to bind opioid receptors neither to stimulate G protein coupled to them, 255 thus their activity in vivo should be related to the activation of other systems involved in nociceptive 256 stimuli control. It is well known that several non-opioid peptides, some of them recently marketed, 257 are powerful antinociceptive agents as evidenced in pre-clinical studies. Non-opioid peptides acting 258 directly or indirectly at different ion channels or non-opioid G-protein coupled receptors (GPCRs) 259 localized in the nociceptive pathways include peptides targeting Ca²⁺, Na⁺ and K⁺ voltage-gated ion 260 channels, the neuronal nicotinic receptors (nAChR), transient receptor potential channels (TRP), the 261 calcitonin gen-related peptide (CGRP), cannabinoid, bradykinin and neurotensin receptors. (Pérez de 262 Vega MJ, Ferrer-Montiel A, González-Muñiz R. 2018; Dimmito et al., 2021; Mollica et al., 2017; 263 Mollica et al. 2015). These are all targets involved in the mechanism of pain through interaction with 264 peptidic endogenous ligands, also showing a certain overlapping positioning with opioid system in 265 the central nervous system (CNS) (Dvoracsko et al. 2015). Thus we cannot exclude the possibility of 266 a single or multitarget interaction with some of them located at central level or periphery. These 267 peptides are characterized by the presence of a cysteine residue in position 3 in place of methionine 268 contained in the *lead compound*. In light of these data, we decided to test the most active compounds, 269 e.g. LMAS5-8 in a battery of antioxidant and enzyme inhibition assays. 270

271 *3.5 Antioxidant activity*

Peptides LMAS5-8 have been evaluated for their antioxidant properties using radical scavenging assays (DPPH and ABTS), reducing power assays (CUPRAC and FRAP) and Phosphomolybdenum assay (Table 1).

275

276	Table 1. Antioxidant ac	ctivities of	f lead	compound	and	LMAS5-8	in	DPPH,	ABTS,	FRAP,	CUPRAC	and
277	Phosphomolybdenum assay	ys*.										

Compounds	DPPH mg TE/g	ABTS mg TE/g	FRAP mg TE/g	CUPRAC mg TE/g	Phosphomolyb denum mg TE/g
Lead compound	na	8.69 ± 0.64	14.21 ± 0.64	19.35 ± 0.64	0.02 ± 0.00
LMAS5	88.84 ± 2.25	136.03 ± 0.61	24.71 ± 0.37	96.79 ± 0.98	1.67 ± 0.03
LMAS6	91.02 ± 0.74	154.25 ± 0.21	27.01 ± 0.11	156.91 ± 2.16	2.18 ± 0.01
LMAS7	62.43 ± 0.46	139.45 ± 1.58	18.53 ± 0.86	99.67 ± 0.97	1.72 ± 0.15
LMAS8	75.75 ± 2.65	131.28 ± 1.78	19.71 ± 0.17	99.64 ± 1.02	1.61 ± 0.03

*Values are reported as mean \pm SD of three parallel experiments. TE: Trolox Equivalent; na: not active.

Peptide LMAS6 was the most remarkable antioxidant compound showing the best activity in DPPH 279 and ABTS assays, among the other analogues. Furthermore, it shows a good activity in FRAP, 280 CUPRAC and phosphomolybdenum assays. Interestingly its antioxidant potential is higher than that 281 of the *lead compound*, which seems to be only slightly active. The improved antioxidant activity of 282 LMAS6 could be due to its amino acid sequence, in particular the presence of cysteine in position 3. 283 Indeed, sulfur-containing amino acids already showed a paramount effect in the reduction of Fe³⁺-284 ferricyanide complex (Nwachukwu & Aluko, 2019; Udenigwe & Aluko, 2011). Sulphur group 285 contained in cysteine may neutralize free radicals forming cysteine sulfoxide, a stable oxidation 286 compound (He, Ju, Yuan, Wang, Girgih, & Aluko, 2012). Furthermore, proline and tyrosine could be 287 involved in direct electrons transfer causing the enhancement of free radical scavenging activity 288 (Ketnawa, Wickramathilaka, & Liceaga, 2018; Nwachukwu et al., 2019). 289

3.6 Enzyme inhibitory activity

Oxidative stress is the principle cause of a huge number of medical diseases like neurological 291 disorders, inflammatory processes, ischemic diseases, hypertension etc (Lobo, Patil, Phatak, & 292 Chandra, 2010). Bioactive agents able to reduce the oxidative damages simultaneously inhibiting the 293 main enzymes involved in this kind of diseases could be useful for the production of functional foods 294 295 or nutraceuticals in combination with commercial drugs. For example, J. acutus, J. maritimus and J. inflexus leaves and roots extracts have been tested as acetylcholinesterase and butyrylcholinesterase 296 inhibitors and for their antioxidant activity; the results suggest their potential role as sources of 297 bioactive compounds useful for the production of nutraceuticals with cognitive improvement 298 properties or food additives (Rodrigues et al., 2017). 299

For this reason, peptides LMAS5-8 have been studied to investigate their in vitro inhibitory activity 300 against acetylcholinesterase, butyrylcholinesterase, amylase, glucosidase and tyrosinase (Table 2). 301 302 Acetylcholinesterase (AChe) and butyrylcholinesterase (BChE) inhibition causes the increase of acetylcholine levels enhancing cognitive functions (Greig et al., 2005; Rodrigues et al., 2017), 303 representing a promising approach for the management of Alzheimer disease. Peptides LMAS5 and 304 LMAS7 exhibit an increased inhibitory activity against acetylcholinesterase compared to the lead 305 compound. Peptide LMAS5 is more effective against butirylcholinesterase than the reference 306 compound which is not active. Hyperglycaemia-induced reactive oxygen species has widely 307 described in literature (Brownlee, 2001; Vanessa Fiorentino, Prioletta, Zuo, & Folli, 2013). The 308 inhibition of carbohydrate hydrolysing enzymes a-amylase and a-glucosidase, is important in the 309 management of hyperglycaemia (Cardullo et al., 2020; Hakamata, Kurihara, Okuda, Nishio, & Oku, 310 2009). For instance, C-glucosidic ellagitannins and some galloylated glucopyranosis have been 311 recently evaluated for their potential use as food ingredients with anti-diabetic effect due to their 312 inhibitor action on amylase and glucosidase (Cardullo et al., 2020). We tested the lead compound and 313 LMAS5-8 for their amylase and glucosidase inhibitory activities (Table 2). Surprisingly the lead 314 compound resulted to be the only active glycosidase inhibitor, all the other analogues show low 315 inhibitory activity against amylase. Tyrosinase inhibitory activity has been checked for LMAS5-8 316 (Table 2). Tyrosinase is involved in browning reactions in food causing the variation of aspect and 317 organoleptic properties of food products reducing their shelf-life and market value (Chazarra, 318 Escribano, & Cabanes, 2001). The use of tyrosinase inhibitors is a promising strategy to prevent 319 browning phenomenon in food industry (Zheng, Cheng, To, Li, & Wang, 2008). In contrast to the 320 321 lead compound, peptides LMAS5-8 are active against tyrosinase; among them LMAS6 gives the best result, suggesting a possible use as anti-browning agent in food industry. 322

323 Table 2. Cholinesterase, tyrosinase, amylase and glucosidase inhibitory activity of *lead compound* and LMAS5-8*.

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Compounds	AChE mg GALAE/g	BChE mg GALAE/g	Tyrosinase mg KAE/g	Glucosidase mmol ACAE/g	Amylase mmol ACAE/g			
Lead compound	3.17 ± 0.09	na	na	1.57 ± 0.03	0.08 ± 0.00			
LMAS5	6.24 ± 0.03	9.57 ± 0.08	59.84 ± 2.28	na	0.09 ± 0.01			
LMAS6	2.73 ± 0.05	na	84.49 ± 1.27	na	0.09 ± 0.02			
LMAS7	5.87 ± 0.04	na	39.49 ± 2.22	na	0.09 ± 0.00			
LMAS8	2.94 ± 0.03	0.48 ± 0.02	49.94 ± 2.28	na	0.11 ± 0.01			

*Values are reported as mean±SD of three parallel experiments. AChE: Acetylcholinesterase; BChE:
butyrylcholinesterase; GALAE: galantamine equivalents; KAE: kojic acid equivents; ACAE: acarbose equivalents; na:
not active.

Overall the combined in vitro and in vivo data confirm the group of synthetic peptides LMAS5-8 as 327 the most active antinociceptive and anti-inflammatory agents, antioxidants and enzyme inhibitors. 328 These peptides possess a molecular weight very close each other's and a chemical structure 329 characterized by common amino acids in several positions of the primary sequence. The C-terminal 330 acid and amide functions don't seem to be responsible of any changes in biological activity, while 331 the presence of a D-Pro² is crucial to guarantee efficient antinociceptive and anti-inflammatory 332 activity in vivo, as well as the best antioxidant power and anti-tyrosinase activity (e.g. LMAS6). This 333 could be due to a stronger metabolic or enzymatic stability in vivo or in vitro, as also recently reported 334 by us for their parent compounds (Stefanucci et al., 2020). Furthermore Cys³ is responsible of a strong 335 effect encountered by CUPRAC, DPPH, and ABTS assays, being also involved in several 336 neuroprotective roles at the CNS (Su et al., 2020). It's interesting to note that all these described 337 features are not present in the lead compound, which resulted to be less active than LMAS6 in all the 338 performed assays. 339

4. CONCLUSION

Isolation and identification of bioactive peptides require high-cost equipment to elucidate their secondary and tertiary structures. This aspect is fundamental to correlate the specific 3D-structure with their biological effects. Regarding the antioxidant peptides, their application as prophylactic agents should be emphasized in order to improve the quality of life. In this work, three series of rubiscolin-6 analogues have been designed following modifications in position 2 and 3 of a previously described rubiscolin-6 analogue YPLDLF. The synthetic procedure is easy and straightforward allowing to obtain high purity compounds through a simple isolation technique. The peptides LMAS5-8 give the best results *in vivo* for their antinociceptive and anti-inflammatory effect. Among them, peptide LMAS6 shows the best antioxidant and tyrosinase inhibitory activities suggesting a possible use in food industry as preservatives or/and anti-browning agents. In addition, data obtained for the anti-cholinesterase peptides LMAS5 and LMAS7 indicate their potential development in new nutraceuticals with cognitive-enhancing properties.

CONFLICT OF INTEREST: Declared none.

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SUPPORTING INFORMATION DESCRIPTION: Peptides sequences and overall yields, RP-HPLC traces, LRMS spectra, opioid binding and GTP stimulation assays, *in vivo* procedure.

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Food-inspired peptides from spinach Rubisco endowed with antioxidant, antinociceptive and anti-inflammatory properties

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HIGHLIGHTS

- Rubiscolin-6 is a linear peptide isolated by the spinach Rubisco.
- New analogues have been prepared via SPPS with high purity and good overall yields.
- Their activity on opioid receptors has been detected in vitro and in vivo.
- Some of them possess a strong antioxidant and tyrosinase inhibitor activity.
- Peptide LMAS6 exerts also a significant anti-inflammatory *in vivo*.