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## Bio-Inspired Casein-Derived Antioxidant Peptides Exhibiting a Dual Direct/Indirect Mode of Action

3 Gizella Csire, François Dupire, Laetitia Canabady-Rochelle, Katalin Selmeczi,\* and Loic Stefan\*



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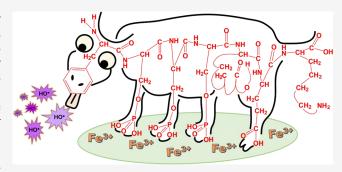
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4 ABSTRACT: Antioxidant compounds are chemicals of primary 5 importance, especially for their applications in nutrition and 6 healthcare, thanks to their abilities to prevent oxidation processes 7 and to limit and/or rebalance the oxidative stress, well-known for 8 its impact on a wide variety of diseases. While several biomolecules 9 are well-known for their antioxidant properties (e.g., ascorbic acid, 10 carotenoids, phenolic derivatives), bio-sourced antioxidants have 11 drawn considerable attention in the last decades, especially 12 bioactive peptides, mainly obtained by the hydrolysis process. 13 Antioxidant peptide sequences are mainly identified a posteriori, 14 thanks to fastidious and time-consuming approaches and 15 techniques, limiting the discovery of new efficient peptides. In



16 this context and taking inspiration from nature, we report herein on a new series of three bio-inspired antioxidant peptides derived 17 from the milk protein casein. These phosphopeptides, designed to chelate the redox-active iron(III) and forming highly soluble 18 complexes up to pH 9, act both as *indirect* (*i.e.*, inhibition of the metal redox activity) and *direct* (*i.e.*, radical scavenging) antioxidants.

#### 19 INTRODUCTION

20 Reactive oxygen species (or ROS) are essential highly reactive 21 components for the proper functioning of the body, acting as 22 signaling molecules to regulate biological and physiological 23 processes. 1,2 In healthy organisms, the control of the ROS 24 concentration is ensured by a complex endogenous antioxidant 25 defense system involving enzymes, proteins, or other 26 chemicals. However, the presence of endogenous or exogeneous stresses can trigger a dysregulation of the ROS 28 homeostasis, inducing the so-called oxidative stress reported to 29 cause a wide variety of diseases. 4,5 In order to prevent oxidative 30 stress and its detrimental effects, the intake of exogeneous 31 antioxidants is an efficient strategy to rebalance the 32 concentration of ROS. Depending on their inherent structure, 33 antioxidants act according to two main modes of action: (i) a 34 direct mode of action whereby the antioxidants directly react 35 with the ROS; 3,6,7 (ii) an indirect mode of action whereby the 36 antioxidants hinder the biometal-induced redox aerobic 37 process via the chelation of specific transition metals. Indeed, 38 in vivo, copper and iron are involved in the production of ROS 39 through the catalytic reduction of molecular oxygen (O2) to 40 the superoxide anion radical (O2 -), and subsequently to 41 hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) and a hydroxyl radical (HO') 42 (Figure 1a). Thus, their chelation induces a decrease of the 43 ROS production and positive effects on ROS-mediated 44 pathologies. Numerous metal-chelating antioxidants 45 have been reported<sup>3,8-11</sup> and, recently, peptides have emerged
46 as new bio-sourced antioxidants.<sup>12-14</sup> Peptides have been 47 already considered as therapeutics offering high selectivity and

effectiveness, safety, and tolerability, 15,16 and biologically 48 relevant peptides can be synthesized using proven, fast, and 49 efficient methods. 17 Up until now, the majority of antioxidant 50 peptides have been identified through tandem LC/MS/MS 51 analyses from protein hydrolysates of natural resour- 52 ces 12-14,18,19 known for their antioxidant properties. However, 53 this serendipitous approach is hazardous, fastidious, and time- 54 consuming, severely limiting the discovery of new effective 55 antioxidant peptides. Thus, we believe that a more rational 56 approach based on a finely tailored bio-inspired molecular 57 design could achieve the same goal in a more judicious and 58 efficient way. While a large body of work concerning copper- 59 chelating peptides has been reported in the recent years, 20,21 60 the development and studies of iron-chelating peptides are still 61 in their infancy and remain challenging. However, iron is 62 present in the body in iron labile pools in which iron is not or 63 slightly chelated and therefore accessible to exogeneous 64 antioxidants,<sup>22,23</sup> including in the lysosomes where iron is in 65 its free Fe(III) form.<sup>22,24</sup> For these reasons and because 66 Fe(III) is in a biologically unusual +III oxidation state, we 67 focus our attention on the development of antioxidant Fe(III)- 68

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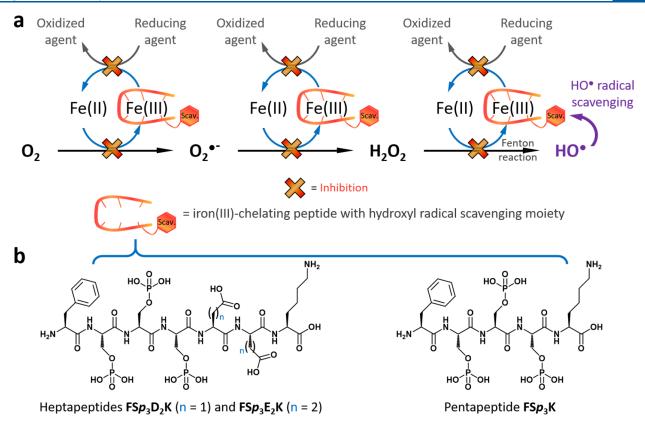


Figure 1. (a) Schematic representation of the *indirect* (i.e., inhibition of the metal redox activity) and *direct* (i.e., radical scavenging) antioxidants mode of actions and (b) chemical structures of the three phosphopeptides studied.

69 chelating peptides with both indirect and direct modes of 70 action (Figure 1a). While a first series of efficient antioxidant 71 pentapeptide has been previously reported by our group, 25 we 72 would like to report herein on new bioinspired phosphopep-73 tides offering both high antioxidant properties and high 74 Fe(III)-chelating capabilities.

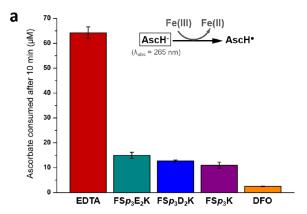
#### ■ RESULTS AND DISCUSSION

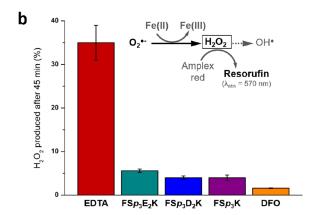
Bioinspired Design and Synthesis of Antioxidant 76 77 **Phosphopeptides.** To design such compounds, we took 78 inspiration from dietary phosphoproteins well-known for their 79 iron-chelating abilities, such as the egg yolk phosvitin<sup>26</sup> or the 80 milk caseins.<sup>27</sup> In particular, caseins are a family of four 81 proteins termed  $\alpha_{S1}$ -,  $\alpha_{S2}$ -,  $\beta$ -, and  $\kappa$ -caseins in which 82 phosphorylation occurs at the serine residues and reported 83 for their antioxidant properties. 28,29 Interestingly, a specific 84 recurring sequence of five amino acids, termed cluster, 85 comprising three phosphoserines (noted Sp) followed by two 86 glutamic acids (E), i.e., an SpSpSpEE cluster, appears once in 87 the  $\beta$ -casein and  $\alpha_{\rm S1}$ -casein, and twice in the  $\alpha_{\rm S2}$ -casein. $^{30-32}$ 88 Based on these observations and on our previous work,<sup>25</sup> we 89 design a first heptapeptide FSp<sub>3</sub>E<sub>2</sub>K (Figure 1b) comprised of 90 the above-mentioned casein-derived cluster on which we grafted (i) a lysine (K) at the C-term bringing an additional 92 charge to improve the peptide solubility at neutral pH, and (ii) 93 a phenylalanine (F) at the N-term to provide a further 94 antioxidant property to the peptide, i.e., a direct mode of action 95 based on the ability of phenylalanine to entrap hydroxyl 96 radicals (vide infra). To evaluate the impact of the two acidic 97 side chains on the chelation and antioxidant properties, two 98 other compounds have been considered: the heptapeptide 99 FSp<sub>3</sub>D<sub>2</sub>K in which the glutamic acids have been replaced by

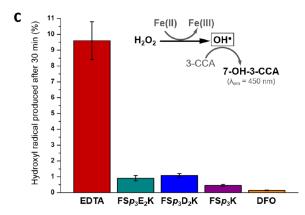
aspartic acids, and the acidic side chain-lacking pentapeptide 100  $FSp_3K$  in which the two E have been removed. The three 101 phosphopeptides were synthesized, purified, and fully characterized (Supporting Information (SI), part 2).

**Evaluation of the Direct and Indirect Antioxidant** 104 **Properties.** To evaluate the antioxidant performance of the 105 three Fe(III)-chelating peptides, we carried out a series of 106 three complementary assays, namely, ascorbate,  $^{33}$  Amplex 107 Red,  $^{34}$  and 3-CCA (for coumarin-3-carboxylic acid)  $^{35}$  tests to 108 measure the iron redox activity (*i.e.*, the reducibility of Fe(III) 109 to Fe(II)) and the formation of  $\mathrm{H_2O_2}$  and  $\mathrm{HO}$ , respectively. 110 To assess the antioxidant power of the three Fe(III)-chelating 111 phosphopeptides, two compounds were selected as references: 112 the positive control is the deferoxamine (DFO), an approved 113 siderophore drug used to treat iron overload and exhibiting 114 antioxidant properties both *in vitro*  $^{36-38}$  and *in vivo*,  $^{39}$  whereas 115 the negative control is the ethylenediaminetetraacetic acid 116 (EDTA), well-known for its good chelation abilities but weak 117 antioxidant properties.  $^{36,37,40}$ 

Thus, employing the ascorbate test, a first series of 119 experiments were carried out with 1 equiv Fe(III) for 1 120 equiv ligand at physiological pH 7.4. As depicted in Figure 2a, 121 f2 the maximum AscH<sup>-</sup> consumption is observed for EDTA 122 (negative control) with 64.2  $\mu$ M consumed after 10 min. 123 Interestingly, when phosphopeptides were used, this value 124 drastically drops to 15.0  $\mu$ M for the casein cluster-derived 125 heptapeptide FS $p_3$ E<sub>2</sub>K, to 12.7  $\mu$ M for the other heptapeptide 126 FS $p_3$ D<sub>2</sub>K, and to 11.0  $\mu$ M for the shortest FS $p_3$ K. These data 127 highlight the efficiency of the three peptides to reduce the 128 redox process of the metal ion from 77% (FS $p_3$ E<sub>2</sub>K) to 83% 129 (FS $p_3$ K) compared to EDTA, even if the positive control DFO 130 appears slightly more efficient (-96%, 2.5  $\mu$ M AscH<sup>-</sup> 131







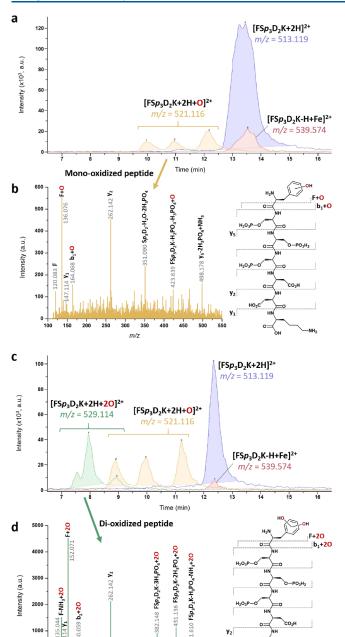
**Figure 2.** Evaluation of the antioxidant properties of all the phosphopeptides and of the negative (EDTA) and the positive (DFO) reference compounds. The corresponding reactions involved for each assay are inserted in each graph, with their corresponding absorbance or emission wavelengths. All experiments were carried out at pH 7.4 with (a) [AscH $^-$ ] = [Ligand] = [Fe(III)] = 100  $\mu$ M, in 50 mM HEPES, (b) [AscH $^-$ ] = 200  $\mu$ M, [Ligand] = [Fe(III)] = 50  $\mu$ M (except [EDTA] = [Fe(III)] = 10  $\mu$ M), in 50 mM HEPES, and (c) [AscH $^-$ ] = 125  $\mu$ M, [Ligand] = [Fe(III)] = 50  $\mu$ M (except [EDTA] = [Fe(III)] = 5.0  $\mu$ M and [AscH $^-$ ] = 12.5  $\mu$ M) in water.

132 consumed, Table S1). To gain further insights into the 133 antioxidant properties of the three phosphopeptides, we then 134 measured their abilities to limit the formation of two distinct 135 ROS, namely,  $H_2O_2$  and HO. Considering the negative 136 control EDTA in the presence of 1 equiv Fe(III), the amount 137 of  $H_2O_2$  formed after 45 min is equal to 35.0% relative to the 138 concentration of the complex (Figure 2b, Table S2). By

contrast, this value drops to 1.6% when the positive control 139 DFO is used. Phosphopeptides also exhibit significant 140 antioxidant properties, with an amount of H2O2 produced 141 from 5.6% for  $FSp_3E_2K$  to 4.0% for both  $FSp_3D_2K$  and  $FSp_3K$ . 142 Interestingly, the glutamic acid-incorporating peptide  $FSp_3E_2K$  143 with the closest casein cluster-like sequence has the weaker 144 antioxidant efficiency compared to the two others, as 145 previously observed in the AscH test (Figure 2a). 146 Subsequently, 3-CCA experiments were carried out, and 147 results (Figure 2c) highlight once more that the highest 148 quantity of HO (9.6%, corresponding to the formed HO / 149 Fe(III)-complex ratio) is formed in the presence of EDTA. 150 Oppositely, DFO and phosphopeptides show better antiox- 151 idant activities, with a ratio of HO produced of 0.14% for 152 DFO, while this value spans from 1.08 (for  $FSp_3D_2K$ ) to 0.46% 153 (FSp<sub>3</sub>K) for our compounds (Table S3). These experiments 154 confirm that all the three phosphopeptides act as effective 155 antioxidants, slightly less efficient than DFO but highly better 156 than EDTA.

However, these peptides have also been designed to behave 158 as direct antioxidants through the entrapment of the most 159 reactive ROS, i.e., the hydroxyl radical HO. Indeed, the 160 oxidation of phenylalanine to tyrosine by HO has already been 161 reported as a marker of oxidative stress. 41,42 To follow the 162 oxidation of the Phe moiety, a first series of four experiments 163 per peptide was performed by HRMS/MS.<sup>25</sup> Thus, the MS 164 spectrum (direct injection) of a stoichiometric peptide/ 165 Fe(III)/AscH<sup>-</sup>/H<sub>2</sub>O<sub>2</sub> solution for the heptapeptide FSp<sub>3</sub>D<sub>2</sub>K 166 (Figure S8, SI for FSp<sub>3</sub>E<sub>2</sub>K (Figure S14), and FSp<sub>3</sub>K (Figure 167 S2)) shows the presence of the native heptapeptide (m/z) = 168513.124 [M + 2H]<sup>2+</sup>), the FSp<sub>3</sub>D<sub>2</sub>K-iron(III) complex (m/z = 169539.579 [M-H + Fe]<sup>2+</sup>), which confirms the Fe(III)-chelating 170 properties of the peptide (discussed hereinafter), and of 171 another peak at m/z = 521.123 corresponding to the monooxidized peptide  $[M + 2H + O]^{2+}$ . To separate and precisely 173 analyze each of these species, all the samples were injected in 174 RP-HPLC/HRMS/MS, and the chromatogram for FSp<sub>3</sub>D<sub>2</sub>K 175 (Figure 3a, SI part 4 for FSp<sub>3</sub>E<sub>2</sub>K and FSp<sub>3</sub>K) reveals the 176 f3 presence of the native peptide (retention time  $t_{\rm R}$  = 13.4 min), 177 preceded by a set of three peaks with shorter retention times 178 ( $t_{\rm R}$  = 10, 11, and 12.2 min). The latter ones correspond to the mono-oxidized  $[M + 2H + O]^{2+}$  species attributable to the 180 oxidation of the phenyl ring of the N-term phenylalanine to its 181 monohydroxy-equivalent as determined by HRMS/MS 182 fragmentation (Figure 3b). The use of HPLC allows us to 183 separate the three phenolic isomers, 43,44 *i.e.*, the peptides 184 containing the ortho-, meta-, and para-hydroxyphenylalanine, 185 indicative of the reaction of the HO with all positions of the 186 phenyl ring. Interestingly, working with harsher conditions (i.e., 187  $FSp_3D_2K/Fe(III)/AscH^-/H_2O_2 1/1/10/10)$  in order to shift 188 the reaction equilibrium, we observed the formation of 189 additional broad peaks (in green, Figure 3c) with shorter  $t_{\rm R}$  190 = 7.5 to 8.9 min characterized by m/z = 529.116 191 corresponding to the di-hydroxylated peptide [M + 2H + 192 20]<sup>2+</sup>, i.e., the peptide in which the phenylalanine has been 193 oxidized twice (Figure 3d). These experiments confirm the 194 ability of our phosphopeptides to selectively react with HO 195 (control experiments were carried out to discard the possibility 196 of a H<sub>2</sub>O<sub>2</sub>-induced oxidation, SI part 4) via the N-term 197 phenylalanine, validating the relevance of our approach and the 198 direct antioxidant mode of action of our designed peptides. 199 Lastly, the HPLC/HRMS chromatograms also highlight the 200

presence of the mononuclear peptide-Fe(III) complex ( $t_R = 201$ 



**Figure 3.** HPLC-HRMS chromatograms highlighting (a) the mono-oxidation of FS $p_3$ D $_2$ K ([Fe(III)] = [FS $p_3$ D $_2$ K] = [AscH $^-$ ] = [H $_2$ O $_2$ ] = 1 mM), and (c) the di-oxidation of FS $p_3$ D $_2$ K ([Fe(III)] = [FS $p_3$ D $_2$ K] = 1 mM, [AscH $^-$ ] = [H $_2$ O $_2$ ] = 10 mM). Corresponding HRMS/MS spectra of the (b) mono-oxidized peptide at m/z = 521.116 and (d) di-oxidized peptide at m/z = 529.114. Experiments were carried out at pH 7.4 in water.

400

250 300

202 13.5 min, m/z = 539.574 [M-H + Fe]<sup>2+</sup> for FS $p_3$ D<sub>2</sub>K (in red, 203 Figure 3a); see SI part 4 for FS $p_3$ E<sub>2</sub>K and FS $p_3$ K), which have 204 been further characterized using potentiometric measurements, 205 circular dichroism (CD), and <sup>31</sup>P NMR spectroscopies (SI part 206 5A,B).

Studies on the Phosphopeptides Chelation Proper-208 ties. First, potentiometric studies were carried out on the 209 synthesized compounds, in the presence and absence of 210 Fe(III), to both access the species distributions as a function of pH and to obtain characteristic thermodynamic constants 211 (protonation, complexation) of the peptides and iron(III) 212 complexes. The deprotonation constants of each ligand alone 213 (i.e., in the absence of Fe(III)) are reported in Table 1. 214 t1

Table 1. Overall Protonation Constants ( $\log \beta$  H<sub>i</sub>L) and Deprotonation Constants (pK) of the Studied Peptides; T = 298 K, I = 0.1 M NaNO<sub>3</sub> (Standard Deviations Are in Parentheses)

$\log\!eta$	$FSp_3K$	$FSp_3E_2K$	$FSp_3D_2K$
HL	10.90 (4)	11.36 (1)	11.43 (1)
$H_2L$	19.16 (6)	19.90 (2)	20.01 (2)
$H_3L$	26.31 (6)	27.28 (2)	27.43 (2)
$H_4L$	32.49 (7)	33.86 (2)	34.02 (2)
$H_5L$	38.18 (8)	39.66 (3)	39.82 (3)
$H_6L$	42.27 (12)	44.74 (3)	44.72 (3)
$H_7L$		49.12 (3)	48.85 (3)
$H_8L$		52.78 (5)	52.37 (5)
pK			
$COOH_{term}$	4.09 (12)	3.66 (5)	3.52 (5)
COOH (1)		4.38 (3)	4.13 (3)
COOH (2)		5.08 (3)	4.90 (3)
$PO_3(OH)^-(1)$	5.69 (8)	5.80 (3)	5.80 (3)
$PO_3(OH)^-(2)$	6.18 (7)	6.58 (2)	6.59 (2)
$PO_3(OH)^-(3)$	7.15 (6)	7.38 (2)	7.42 (2)
$\mathrm{NH_{3}^{+}}\left(\mathrm{N_{term}}\right)$	8.26 (6)	8.54 (2)	8.58 (2)
$\varepsilon \mathrm{NH_3}^+$ (Lys)	10.90 (4)	11.36 (1)	11.43 (1)

Depending on the peptide sequence, these ligands have 9 215  $(FSp_3K)$  or 11  $(FSp_3D_2K, FSp_3E_2K)$  protonable groups, 216 notably one or three carboxylate (COOH/COO<sup>-</sup>), three 217 phosphates  $(PO_2(OH)_2/PO_4^{2-})$ , and two amino  $(-NH_3^+/-218)$  $NH_2$ ) groups. The first pK of phosphate groups  $(PO_2(OH)_2/219)$ PO<sub>3</sub>(OH)<sup>-</sup>) is too low to be easily determined in titration 220 experiments. Thus, the first measured dissociation step 221 corresponds to the deprotonation of terminal COOH whose 222 value is considerably higher (p $K_{COOHterm} = 3.52-4.09$ ) than 223 the pK value of free amino acids (2.0-2.5). This increase in 224 the pK values is probably due to the large network of hydrogen 225 bonds formed between the negatively charged phosphate 226 groups (as H acceptors) and the protonated amine and 227 COOH groups (as H donors). This H-bond network has also 228 an impact on the deprotonation process of terminal NH2 229  $(pK_{Nterm} = 8.26 - 8.58)$  and lysine  $NH_2$   $(pK_{Lys} = 10.90 - 11.43)$  230 groups, whose pK values are therefore higher than those 231 observed for the first series of pentapeptide FD<sub>3</sub>K (p $K_{Nterm}$  = 232 7.64,  $pK_{Lys} = 10.98$ ) and  $FE_3K$  ( $pK_{Nterm} = 7.50$ ,  $pK_{Lys} = 233$ 10.82). In the case of FSp<sub>3</sub>D<sub>2</sub>K, FSp<sub>3</sub>E<sub>2</sub>K peptides, two 234 additional deprotonation steps were measured corresponding 235 to the dissociation of aspartic and glutamic acid side chains 236  $(pK_{COOH(1)} = 4.13 \text{ and } 4.38, pK_{COOH(2)} = 4.90 \text{ and } 5.08 \text{ for } 237)$  $FSp_3D_2K$  and  $FSp_3E_2K$ , respectively). These pK values are 238 similar to those measured for the previously studied peptides, 25 239 showing that these groups are not involved in inter- or 240 intramolecular interactions. Regarding the PO<sub>3</sub>(OH)<sup>-</sup>/PO<sub>4</sub><sup>2-</sup> 241 dissociation processes of three phosphate groups in the peptide 242 sequence, the consecutive pK values are well separated ( $\Delta$ pK 243  $\approx 0.6-0.9$ ) and the first p $K_{PO3(OH)-(1)}$  value (5.69-5.80) <sup>244</sup> corresponds well to those of phosphoserine alone 245  $(pK_{PO3(OH)-serine} = 5.61).$ 

Then, stability constants  $(\log \beta)$  and stepwise deprotonation <sup>247</sup> constants (pK) of iron(III) complexes were assessed and are <sup>248</sup> f4

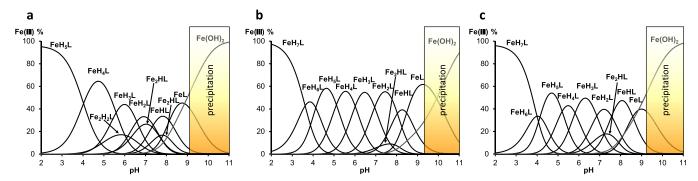


Figure 4. Speciation diagrams of Fe(III)-phosphopeptides 1:1 systems: (a) Fe(III)-FS $p_3$ K, (b) Fe(III)-FS $p_3$ D<sub>2</sub>K and (c) Fe(III)-FS $p_3$ E<sub>2</sub>K. For all experiments, [peptide]<sub>tot</sub> = 2.0 mM, T = 298 K, and I = 0.1 M NaNO<sub>3</sub>.

249 summarized in Table 2 and Figure 4. The results indicate that 250 mono- and binuclear complexes were formed, while bis(ligand) 251 complexes were not observed. For all the three phosphopeptides and at pH 2, the second deprotonation of one phosphate group occurs, meaning that the complexation of iron(III) is already performed by mono- and di-deprotonated phosphate groups. Accordingly, potentiometric titrations start with the 256 species FeH<sub>5</sub>L for FSp<sub>3</sub>K and FeH<sub>7</sub>L for both FSp<sub>3</sub>D<sub>2</sub>K and 257 FSp<sub>3</sub>E<sub>2</sub>K. The following steps up to the formation of FeH<sub>4</sub>L 258 species correspond to the deprotonation of COOH<sub>term</sub> for the 259 three phosphopeptides and also to the deprotonations of COOH(1) and COOH(2) in the case of  $FSp_3D_2K$  and FSp<sub>3</sub>E<sub>2</sub>K. The next two dissociation steps belong to the 262 formation of the FeH<sub>3</sub>L and FeH<sub>2</sub>L complexes where the other 263 two phosphate groups are deprotonated. All these pK values 264 from p $K(FeH_7L/FeH_6L)$  to p $K(FeH_3L/FeH_2L)$  are close to 265 the corresponding pK values in the ligands alone (Table 1), 266 demonstrating that the metal ion-promoted dissociation 267 process does not occur. The H-bond network, operating in

Table 2. Stability Constants ( $\log \beta$ ) and Derived Data (pK) for Fe(III) Complexes of the Studied Peptides; T = 298 K,  $I = 0.1 \text{ M NaNO}_3$  (Standard Deviations Are in Parentheses)

$\log\!eta$	$FSp_3K$	$FSp_3E_2K$	$FSp_3D_2K$
FeH <sub>7</sub> L		59.24 (5)	60.32 (4)
FeH <sub>6</sub> L		55.24 (7)	56.70 (4)
FeH <sub>5</sub> L	48.19 (6)	51.10 (5)	52.57 (4)
$FeH_4L$	44.16 (5)	45.90 (6)	47.45 (3)
FeH <sub>3</sub> L	38.57 (6)	40.08 (5)	41.43 (3)
FeH <sub>2</sub> L	31.99 (7)	33.22 (6)	34.48 (3)
FeHL	24.62 (6)	25.65 (3)	26.44 (2)
FeL	16.48(5)	17.07 (5)	17.96 (3)
Fe <sub>2</sub> H <sub>6</sub> L		62.05 (7)	62.83 (6)
Fe <sub>2</sub> H <sub>5</sub> L		58.78 (7)	59.99 (4)
Fe <sub>2</sub> H <sub>4</sub> L	50.71 (8)	54.81 (6)	56.20 (4)
Fe <sub>2</sub> H <sub>3</sub> L	47.70 (6)	49.87 (6)	51.54 (4)
$Fe_2H_2L$	43.07 (6)	44.33 (5)	45.86 (4)
$Fe_2HL$	36.78 (6)	37.73 (5)	39.09 (4)
$Fe_2L$	29.13 (10)		
p $K$ (FeH <sub>7</sub> L/FeH <sub>6</sub> L)		4.00	3.62
pK (FeH <sub>6</sub> L/FeH <sub>5</sub> L)		4.14	4.13
pK (FeH <sub>5</sub> L/FeH <sub>4</sub> L)	4.03	5.20	5.12
pK (FeH <sub>4</sub> L/FeH <sub>3</sub> L)	5.59	5.82	6.02
pK (FeH <sub>3</sub> L/FeH <sub>2</sub> L)	6.58	6.86	6.95
pK (FeH <sub>2</sub> L/FeHL)	7.37	7.57	8.04
pK (FeHL/FeL)	8.14	8.58	8.48
-			

the solution of phosphopeptide alone, cannot be formed with 268 terminal NH<sub>3</sub><sup>+</sup> because the phosphate groups are coordinated <sup>269</sup> to the metal ion; thus, the  $pK_{Nterm}$  (FeHL formation) value is 270 decreased to 7.37-8.04, a value close to that observed with the 271 previous FD<sub>3</sub>K series.<sup>25</sup> The last dissociation step corresponds <sup>272</sup> to water deprotonation coordinated to the metal ion with a pK 273 value of 8.14, 8.48, and 8.58 for FSp<sub>3</sub>K, FSp<sub>3</sub>D<sub>2</sub>K, and 274  $FSp_3E_2K$ , respectively. At a concentration of 100  $\mu$ M and at pH 275 7.4, while small precipitation occurred in the presence of 276 FD<sub>3</sub>K, homogeneous solutions were detected with the 277 phosphorylated FSp<sub>3</sub>K, FSp<sub>3</sub>D<sub>2</sub>K, and FSp<sub>3</sub>D<sub>2</sub>K peptides. At 278 this pH, the pFe value is, for instance, 1.5 logarithmic units 279 higher for  $FSp_3D_2K$  (pFe = 14.3) than for  $FD_3K$  (pFe = 280 12.8),<sup>25</sup> highlighting that the presence of phosphoserines in the 281 sequences significantly increases the stability of the formed 282 iron(III) complexes and are able to prevent metal ion 283 hydrolysis up to pH 9.2. Based on all these data 284 (potentiometric and spectroscopic, SI part 5), it can be 285 reasonably assumed that the mononuclear 1:1 Fe(III):peptide 286 complexes are formed through the chelation of the metal ion 287 mainly by the phosphate groups, even if other functions (e.g. 288 carboxylate) and/or water can be involved in the chelation as 289 previously reported.<sup>25</sup> However, to decipher the precise 290 complex structures of these phosphopeptides, further inves- 291 tigations are required. Interestingly, such data both confirm 292 and bring new insights into the role of the casein cluster Sp<sub>3</sub>E<sub>2</sub> 293 on the chelation and antioxidant properties of milk casein 294 proteins, as already reported.<sup>27–32</sup>

Finally, since our phosphopeptides are derived from a 296 specific sequence of casein, a milk protein well-known for its 297 ability to bind calcium mainly through its phosphoserines, 45,46 298 we performed competition experiments in the presence of 299 calcium ions. Using the AscH test, no loss of antioxidant 300 activity is observed upon addition of 1 and 10 equiv of the 301 non-redox active Ca(II) to a 1:1 mixture of Fe:peptide (Table 302 S1); conversely, a sensitive decrease of the ascorbate 303 consumption, *i.e.*, a better antioxidant activity, appears. 304 Additional CD, <sup>31</sup>P NMR, and ESI-MS studies of the 305 Fe(III):Ca(II):phosphopeptide ternary systems (Figures 306 S23-S27) show that iron remains chelated even with excess 307 calcium and suggest a possible reorganization of the peptides 308 around Fe(III) in the presence of Ca(II) (SI part 5B), 309 confirming the efficiency and selectivity of our bioinspired 310 antioxidant phosphopeptides. 311

#### CONCLUSIONS

Thus, we reported herein on the rational design of an original 313 series of bioinspired casein-derived phosphopeptides exhibiting 314

315 significant antioxidant properties thanks to a dual mode of 316 action. Indeed, these compounds show high Fe(III)-chelating 317 abilities with stability of the complexes formed up to pH 9.2, 318 leading to indirect antioxidant activities. Concomitantly, the 319 direct antioxidant behavior is provided by the phenylalanine 320 moiety and its inherent abilities to act as radical scavenger, 321 especially for the highly reactive hydroxyl radical, as observed 322 by HRMS experiments. Altogether, these two combined 323 antioxidant effects brought by the phosphopeptides reduce, 324 *inter alia*, the metal redox process to 83% and the formation of 325 HO to -95%.

Also, our study highlights that to design effective antioxidant peptides, the incorporation of aromatic amino acid(s) (herein, phenylalanine) is an easy way to induce a direct antioxidant mode of action (*i.e.*, radical scavenging), while the presence of highly chelating amino acids (herein, phosphoserines) are required to assess both an efficient Fe(III) chelation and effective indirect antioxidant properties.

#### **MATERIAL AND METHODS**

Peptide Synthesis (General Procedure). The three peptides 334 335 were synthesized at a 200  $\mu$ mol scale using an automated ResPep XL 336 synthesizer (Intavis AG) using an Fmoc/tBu strategy and double 337 couplings for each amino acid. Preloaded Wang-resin (0.70 mmol·g<sup>-1</sup>, 338 200-400 mesh) was used. The standard conditions for each coupling were Fmoc-amino acid (5 equivalents), 2-(1H-benzotriazol-1-yl)-340 1,1,3,3-tetramethyl-uronium tetrafluoroborate (HBTU, 5 equiva-341 lents), and 4-methylmorpholine (NMM, 10 equivalents) in 342 dimethylformamide (DMF) and N-methyl-2-pyrrolidone (NMP), 343 with a coupling time of 60 min. Fmoc-Ser(PO(OBzl)OH)-OH was 344 introduced at 4.0 equiv using HBTU (4.0 equiv), NMM (8.0 equiv) 345 in DMF, and NMP. Fmoc-deprotection steps were carried out using a 346 20% piperidine solution in DMF (3 × 15 min), and final cleavages 347 were achieved using a trifluoroacetic acid/triisopropylsilane/water 348 (92.5/5/2.5) mixture (3 h, 500 rpm). The crude peptides were 349 precipitated from cold diethylether (-20 °C), centrifuged ( $3 \times 5000$ 350 rpm, 5 min) washed with cold diethylether, dried under reduced 351 pressure, resolubilized in water, and finally, lyophilized. Then, 352 peptides were dissolved in a solvent A, sonicated, and purified by 353 semi-preparative HPLC equipped with a Nucleosil (Macherey-354 Nagel) 100-5  $C_{18}$  250 × 21 mm column using solvent A (95% water, 355 5% acetonitrile, 0.1% trifluoracetic acid) and solvent B (100% 356 acetonitrile, 0.1% trifluoroacetic acid) with a UV detection at 214 nm. 357 For FSp<sub>3</sub>K, the same procedure was applied except solvent A, which is 358 97% water, 3% acetonitrile, and 0.1% trifluoracetic acid. The resulting 359 solutions were evaporated under reduced pressure and were double-360 lyophilized. The purity of each peptide was evaluated by analytical 361 reversed-phase HPLC equipped with a Nucleosil (Macherey-Nagel) 362 100-5 C18 250  $\times$  4.6 mm column using solvent A' (97% water, 3% 363 acetonitrile, 0.1% formic acid) and solvent B' (100% acetonitrile, 0.1% 364 formic acid) with a UV detection at 214 nm. Peptides were obtained 365 with 31% (FSp<sub>3</sub>E<sub>2</sub>K), 37% (FSp<sub>3</sub>D<sub>2</sub>K), and 38% (FSp<sub>3</sub>K) yield. All 366 chemical characterizations (NMR, HRMS, and HPLC) are provided 367 in the SI.

368 Antioxidant Activity Assays (General Procedure). HEPES 369 (50 mM, pH 7.4) buffer was treated on a Chelex100 to remove trace 370 metal contaminants from the solution and prevent unwanted reactions 371 before the different tests. The stock solution at 2 mM of iron(III)-372 nitrate was prepared from an analytical grade reagent in an acidic 373 media (0.1 M HNO<sub>3</sub>) to keep iron(III) ion in solution; its exact 374 concentration was checked spectrophotometrically using complexo-375 metric titration. The stock solution at 2 mM of calcium(II)-chloride 376 was prepared in  $H_2O$ , and the concentration was checked as the same 377 method as in the case of iron(III). Peptides, EDTA, and DFO ligand 378 stock solutions at 2 mM were prepared using HEPES buffer. To 379 perform the tests, a semi-micro quartz cell with a 1 cm optical 380 pathway was used; the sample holder in the spectrophotometer was

thermostated at 25  $^{\circ}$ C by a Peltier temperature controller. For all the 381 experiments, the iron(III)-complexes were prepared 20 min before 382 each antioxidant analysis to assure a complete chelation of the metal 383 ion.

**Ascorbate Tests.** For a standard assay, a 5 mM ascorbate stock 385 solution was freshly prepared using HEPES buffer at pH 7.4. The 386 absorbance of ascorbate was followed over 10 min at 265 nm ( $\varepsilon$  = 387 14,500 M<sup>-1</sup> cm<sup>-1</sup>), and at least three independent measurements were 388 carried out. In a total volume of 1 mL, the final concentration of the 389 components was 100 μM with 1:1:1 AscH<sup>-</sup>:ligand:Fe(III) ratio. In 390 the ascorbate test and in the competition study with Ca(II), the final 391 concentration of the components was also 100 μM with a 1:1:1 (or 2 392 or 10) AscH<sup>-</sup>:ligand:Ca(II) ratio or with a 1:1:1:1 (or 10) 393 AscH<sup>-</sup>:ligand:Fe(III):Ca(II) ratio. The pH was checked at the end 394 of each assay. The reaction was triggered by the addition of ascorbate. 395 As a reference reaction, AscH<sup>-</sup> oxidation by ligands alone was also 396 studied. The initial ascorbate oxidation rate was calculated from the 397 slope of [AscH<sup>-</sup>] = f(t) (Figure S1).

Amplex Red Assays. The H<sub>2</sub>O<sub>2</sub> production was detected by an 399 Amplex Red reagent in the presence of horseradish peroxidase (HRP) 400 forming the compound Resorufin with  $\lambda_{abs}$  = 570 nm. For this assay, 401 all components except the iron(III) ion and Amplex Red reagent were 402 dissolved in 50 mM HEPES buffers. The Amplex Red reagent was 403 dissolved in a DMSO:buffer mixture at a 1:4 ratio. A working solution 404 was first prepared containing 0.8 U/mL HRP and 200  $\mu$ M Amplex 405 Red reagent in a light-protected tube. A total of 50  $\mu$ L of this solution 406 was placed in the total volume of 1 mL of reaction mixture. The final 407 concentration of the AscH $^-$  was 200  $\mu$ M, and the ligand (peptides or 408 DFO):Fe(III):AscH<sup>-</sup> ratio was 1:1:4. In the case of EDTA, a ratio of 409 1:1:20 of EDTA:Fe(III):AscH was applied. The AscH was the last 410 reagent added, thus initiating the reaction between O2 and Fe(III); 411 the pH was checked at the end of each reaction. A negative control 412 containing only the working solution in buffer was necessary in order 413 to monitor the background absorbance of the Amplex Red itself. The 414 Resorufin formation was monitored by visible absorption spectrosco- 415 py at 570 nm at RT over 45 min. To determine the H<sub>2</sub>O<sub>2</sub> 416 concentration, a standard calibration curve was used. To obtain this 417 calibration curve, a 20 mM hydrogen peroxide solution was previously 418 titrated by permanganometry and then diluted successively. The 419 results are summarized in Table S2.

Coumarin-3-carboxylic Acid (3-CCA) Assay. The HEPES 421 buffers disturbed the fluorescence spectra and also had efficient 422 HO radical scavenging activity, thus pure water treated on Chelex100 423 was used as solvent and the pH was adjusted by adding 0.1 M NaOH. 424 Thus, all the reagents were dissolved in Milli-Q water except the 3- 425 CCA, since it was only soluble in phosphate buffer (50 mM, pH 7.4). 426 The concentration of the 3-CCA stock solution was 5 mM. In the 427 total volume of 2 mL, the final concentrations of the ascorbate and 3-428 CCA were both 125  $\mu$ M, and the ratio of the reaction mixture was 429 ligand (peptides, EDTA or DFO):Fe(III):AscH<sup>-</sup> = 1:1:2.5. The pH 430 was checked at the end of the measurement. In the case of EDTA, the 431 amount of all components was 10 times less than that of the other 432 ligand systems (in order to avoid the saturation of the signal due to 433 the high activity of EDTA). For the greater reproducibility of the 434 measured data, each measurement was started by monitoring a blank 435 sample (without any ascorbate) for 5 min, and then the ascorbate was 436 added to the sample. The reaction between 3-CCA and hydroxyl 437 radical generating the fluorescent compound 7-OH-3-CCA was 438 followed over 30 min at 25 °C at  $\lambda_{\rm em}$  = 450 nm ( $\lambda_{\rm ex}$  = 395 nm) by 439 spectrofluorometry. The HO concentration was determined by a 440 standard calibration curve using commercially available 7-OH-3-CCA. 441

**High Resolution Mass Spectroscopy Studies.** For direct 442 injections (HRMS, HRMS/MS, and LC-HRMS experiments), the 443 reaction mixtures were prepared at room temperature in a total 444 volume of 1 mL at pH 7.4 in water (pH was adjusted by 0.1 M 445 NaOH) and analyzed after 24 h. Five different mixtures were studied, 446 including peptide alone, peptide + oxidant ( $H_2O_2$ ), peptide + Fe(III) 447 + oxidant ( $H_2O_2$ ), peptide + Fe(III) + oxidant ( $H_2O_2$ ) + ascorbate, 448 and also, as another control without oxidant, peptide + Fe(III) + 449 ascorbate. Similarly to antioxidant assays, the iron(III)-complexes 450

451 were prepared 20 min before the addition of the redox agent(s)  $452~(H_2O_2~and/or~ascorbate),$  to assure a complete chelation of the metal. 453~LC-HRMS experiments were carried out using solvent A (95% water, 454~5% acetonitrile, 0.1% trifluoracetic acid) and solvent B (100% 455 acetonitrile, 0.1% trifluoracetic acid) with a 15 min linear gradient 456~(0.5%~to~8%~solvent~B) followed by another 10 min linear gradient 457~(up~to~50%~B).

Potentiometric Measurements. The pH-potentiometric titra-459 tions were investigated in the pH range of 2.0-11.0 at I=0.1 M 460 NaNO<sub>3</sub> ionic force and at  $T = 298.0 \pm 0.1$  K. A Dosimat 715 461 (Metrohm) automatic burette and pH-meter equipped with a semi-462 micro combined electrode (Metrohm) was used for the titration. The 463 initial concentration of peptides was  $1.5 \times 10^{-3}$  M, using 1:1, 1:2, and 464 2:1 metal-to-ligand ratios. The titrations were performed with a 465 carbonate free stock solution of sodium hydroxide at known 466 concentration. During the titration, argon was bubbled through the 467 samples to ensure the absence of oxygen and carbon dioxide. The 468 recorded pH readings were converted to hydrogen ion concentration 469 as described by Irving et al. 47 Protonation constants of the ligands and 470 the overall stability constants ( $\log\!eta_{
m pqr}$ ) of the complexes were 471 calculated from at least three independent titrations (ca. 70 data 472 points per titration) by means of the general computational programs, 473 PSEQUAD<sup>48</sup> and SUPERQUAD using eqs 1 and 2.

$$pM + qH + rL \rightleftharpoons M_p H_q L_r \tag{1}$$

$$\beta_{pqr} = \frac{[M_p L_q H_r]}{[M]^p [L]^q [H]^r}$$
(2)

#### **476 ASSOCIATED CONTENT**

#### 7 Supporting Information

478 The Supporting Information is available free of charge at 479 https://pubs.acs.org/doi/10.1021/acs.inorgchem.1c03085.

Chemical analysis of synthetized peptides, results of antioxidant activity assays, high-resolution mass spectroscopy studies, and complexation studies (PDF)

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Notes

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