Both metal-chelating and free radical-scavenging synthetic pentapeptides as efficient inhibitors of reactive oxygen species generation

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- Supporting information -

Table of contents:	
1. Chemical characterization	p.2
2. Solution equilibrium studies	p.8
3. Antioxidant activity assays	p.11
3.A. Ascorbate test	p.11
3.B. Nitro-blue tetrazolium (NBT) test	p.12
3.C. Amplex red assay	p.13
3.D. Coumarin-3-carboxyilic acid (3-CCA) assay	p.14

1. CHEMICAL ANALYSIS

NMR analysis:

1.A. Pentapeptide H₂N-Phe-Asp-Asp-Asp-Lys-OH, FD₃K

Compound H₂N-Phe-Asp-Asp-Asp-Lys-OH was obtained as a white solid (176 mg, 203 µmol, 51% yield).



¹**H** NMR (400 MHz, H₂O, pH 4.0) δ (ppm): 1.32 (m, 2H, H19), 1.60 (m, 2H, H20), 1.63-1.84 (m, 2H, H18), 2.52-2.82 (m, 6H, H12, H14, H16), 2.93 (m, 2H, H21), 3.15 (m, 2H, H11), 4.14 (m, 1H, H9), 4.21 (t, *J* = 7.4 Hz, 1H, H1), 4.53-4.60 (m, partially suppressed, H3, H5, H7), 7.21-7.3 (m, 5H, H aromatic), 7.80, 8.31, 8.66 (m, NH amide).



¹³C NMR (100 MHz, H₂O, pH 4.0) δ (ppm): 21.9 (C19), 26.3 (C20), 30.7 (C18), 36.6, 36.8, 37.1, 37.2 (C12, C14, C16, C11), 39.6 (C21), 51.0, 51.1, 51.2 (C3, C5, C7), 52.8 (C9), 54.1 (C1), 128.0, 129.2, 129.4, 133.8 (C aromatic), 169.1 (C2), 172.0, 172.1 (C4, C6), 172.6 (C8), 175.9, 176.0, 176.1 (C13, C15, C17), 177.8 (C10).

Reverse-phase analytical HPLC chromatogram:



HRMS (ESI): m/z calculated for C₂₇H₃₉N₆O₁₂ [M+H]⁺ (100%), 639.2620, found 639.2636.

1.B. Pentapeptide H₂N-Phe-Glu-Glu-Glu-Lys-OH, FE₃K

Compound H2N-Phe-Glu-Glu-Glu-Lys-OH was obtained as a white solid (257 mg, 283 µmol, 71% yield).

NMR analysis:



¹**H** NMR (400 MHz, H₂O, pH 6.0) δ (ppm): 1.31 (m, 2H, H22), 1.57 (m, 2H, H23), 1.65 (m, 2H, H21), 1.72-2.04 (m, 6H, H12, H15, H18), 2.14 (m, 2H, H19), 2.23 (m, 4H, H13, H16), 2.91 (t, 2H, H24), 3.13 (m, 2H, H11), 4.10 (m, 1H, H9), 4.21 (m, 1H, H1), 4.15-4.26 (m, 3H, H3, H5, H7), 7.17-7.37 (m, 5H, H aromatic), 7.92, 8.45 (m, NH amide).



¹³**C NMR** (100 MHz, H₂O, pH 6.0) δ (ppm): 22.0 (C22), 26.4 (C23), 27.6 (C18), 27.8 (C12, C15), 31.1 (C21), 33.4 (C19), 33.5, 33.6 (C13, C16), 36.9 (C11), 39.5 (C24), 53.5 (C7), 53.9, 54.0 (C3, C5), 54.6 (C1), 54.8 (C9), 128.0, 129.1, 129.4, 133.8 (C aromatic), 169.3 (C2), 172.6, 172.8 (C4, C6), 173.5 (C8), 178.5 (C10), 181.2 (C20), 181.3, 181.4 (C14, C17).

Reverse-phase analytical HPLC chromatogram:



<u>HMRS (ESI)</u>: m/z calculated for C₂₇H₃₉N₆O₁₂ [M+H]⁺ (100%), 639.2620, found 639.2636.

1.C. Pentapeptide H₂N-Phe-Asp-Asp-Ala-OH, FD₃A

Compound H₂N-Phe-Asp-Asp-Asp-Ala-OH was obtained as a white solid (162 mg, 233 µmol, 58% yield).





¹**H NMR** (400 MHz, H₂O, pH 2.3) δ (ppm): 1.35 (d, 3H, H18), 2.66-2.92 (m, 6H, H12, H14, H16), 3.14 (m, 2H, H11), 4.21 (t, 1H, H1), 4.29 (m, 1H, H9), 4.25-4.32 (m, 3H, H3, H5, H7), 7.19-7.40 (m, 5H, H aromatic), 8.05, 8.23, 8.44, 8.68 (m, N*H* amide).



¹³C NMR (100 MHz, H₂O, pH 2.3) δ (ppm): 16.3 (C18), 35.3-35.5 (C12, C14, C16), 36.9 (C11), 49.0 (C9), 50.1-50.4 (C3, C5, C7), 54.5 (C1), 128.1-133.7 (C aromatic), 169.0 (C2), 171.4-171.9 (C4, C6, C8), 174.1-174.2 (C13, C15, C17), 176.4 (C10).

Reverse-phase analytical HPLC chromatogram:



HMRS (ESI): *m/z* calculated for C₂₄H₃₂N₅O₁₂ [M+H]⁺ (100%), 582.2042, found 582.2071.

1.D. Pentapeptide H2N-Phe-Glu-Glu-Glu-Ala-OH, FE3A

NMR analysis:

Compound H₂N-Phe-Glu-Glu-Glu-Ala-OH was obtained as a white solid (140 mg, 190 µmol, 48% yield).

KSCG-LST053.3.fid FEEEA pH 1.8 HO 21 CH3 OH H_2N 1 HO HO ò Ò 8.5 8.0 7.5 6.5 5.5 5.0 4.5 4.0 3.5 3.0 7.0 6.0 2.5 2.0 1.5 f1 (ppm)

¹**H** NMR (400 MHz, H₂O, pH 1.8) δ (ppm): 1.36 (d, 3H, H21), 1.81-2.11 (m, 6H, H12, H15, H18), 2.34 (t, 2H, H19), 2.43 (t, 4H, H13, H16), 3.14 (d, 2H, H11), 4.22 (t, 1H, H1), 4.25-4.34 (m, 3H, H3, H5, H7), 4.30 (m, 1H, H9), 7.17-7.36 (m, 5H, H aromatic), 8.5-8.46 (m, NH amide).



¹³C NMR (100 MHz, H₂O, pH 1.8) δ (ppm): 16.2 (C21), 26.1-26.3 (C12, C15), 26.5 (C18), 29.7 (C19), 29.8 (C16), 29.8 (C13), 36.9 (C11), 48.9 (C9), 52.7 (C7), 52.9 (C5), 53.1 (C3), 54.4 (C1), 128.0.0-133.6 (C aromatic), 169.0 (C2), 172.0 (C6), 172.7 (C8), 172.9 (C4), 176.4 (C10), 177.0 (C20, C17), 177.1 (C14).

Reverse-phase analytical HPLC chromatogram:



HMRS (ESI): m/z calculated for C₂₇H₃₈N₅O₁₂ [M+H]⁺ (100%), 624.2511, found 624.2548.

2. SOLUTION EQUILIBRIUM STUDIES



Figure S1. pH dependent CD spectra of FD_3K ligand (left), and pH-dependence of CD signal at 223 nm (right, secondary axis) with the corresponding speciation diagram (right, primary axis). $[FD_3K]_{tot} = 1 \text{ mM}$



Figure S2. pH dependent CD spectra of FD₃K ligand in the presence of 1 equivalent iron(III) ion (left), and pHdependence of CD signal at 282 nm (right, secondary axis) with the corresponding speciation diagram (right, primary axis). $[FD_3K]_{tot} = [Fe^{3+}] = 1 \text{ mM}$









8.6 8.4 8.2 8.0 7.8 7.6 7.4 7.2 4.8 4.6 4.4 4.2 4.0 3.8 3.6 3.4 3.2 3.0 2.8 2.6 2.4 2.2 2.0 1.8 1.6 1.4 1.2 **Figure S5.** ¹H NMR spectra of FD₃K alone (red), complexes Fe(III):FD₃K 1:6 ratio (green) at pH 4.



Figure S6. ¹H NMR spectra of FE₃K alone (red), complexes Fe(III):FE₃K 1:5 ratio (green) at pH 6.

3. ANTIOXIDANT ACTIVITY ASSAYS

3.A. ASCORBATE TEST

pH 6.0 50 mM MES	$v_i \times 10^{-8} (Ms^{-1})$	[AscH ⁻] after 10 min (µM)	AscH ⁻ consumed (%)
AscH ⁻	0.14(2)	99.3	0.7
FD ₃ K	0.088(3)	99.6	0.4
Fe(III):FD ₃ K	4.4(4)	80.5	19.5
Fe(III):DFO	1.3(2)	93.4	6.6
Fe(III):EDTA	5.70(2)	68.0	32.0
FE ₃ K	0.10(2)	99.7	0.3
Fe(III):FE ₃ K	4.1(2)	81.7	18.3

Table S1. Initial reaction rate (standard deviations are in parentheses) and AscH⁻ concentration after 10 min reaction at pH 6.0 (50 mM MES buffer). [AscH⁻]₀ = [Ligand]₀ = 100 μ M and [Fe(III)]₀ = 100 μ M if present T = 298 K

Table S2. Initial reaction rate (standard deviations are in parentheses) and AscH⁻ concentration after 10 min reaction at pH 7.4 (50 mM HEPES buffer). [AscH⁻]₀ = [Ligand]₀ = 100 μ M and [Fe(III)]₀ = 100 μ M if present, *T* = 298 K.

pH 7.4 50 mM HEPES	$v_i {\times} 10^{-8}(Ms^{-1})$	[AscH ⁻] after 10 min (µM)	AscH ⁻ consumed (%)
AscH ⁻	0.1(1)	99.4	0.6
FD ₃ K	0.08(3)	99.5	0.5
Fe(III):FD ₃ K	1.2(2)	92.8	7.2
Fe(III):DFO	0.42(2)	97.5	2.5
Fe(III):EDTA	10.7(4)	35.8	64.2



Figure S7. Absorbance at 265 nm *versus* time graph for the ascorbate consumption at pH 7.4 (50 mM HEPES buffer). [AscH⁻]₀ = [Ligand]₀ = 100 μ M and [Fe(III)]₀ = 100 μ M if present, *T* = 298 K.

3.B. NITRO-BLUE TETRAZOLIUM (NBT) TEST

To measure the quantity of superoxide anion radical formed, the classical NBT test was chosen. The formation of the blue product NBT-diformazan ($\lambda_{max} = 560$ nm, $\varepsilon = 12300$ M⁻¹cm⁻¹, Scheme S1) was followed spectrophotometrically (see C. Beauchamp and I. Fridovich, *Anal. Biochem.*, 1971, **44**, 276-287). In a general test reaction the following conditions were used: [AscH⁻]₀ = 300 µM, [Ligand]₀ = 100 µM, [Fe(III)]₀ = 100 µM and [NBT] = 500 µM in 50 mM buffer solution. Two major problems have been identified during the control measurements: inadequacy of generally used buffers (HEPES and MES) and the reaction between AscH⁻ and NBT (see K. Reybier, S. Ayala, B. Alies, J. V. Rodriguez, S. B. Rodriguez, G. La Penna, F. Collin, C. M. Gomes, C. Hureau and P. Faller, Angew. Chem. Int. Ed., 2015, 55, 1085-1089). Other buffers, TRIS or phosphate have also been tested and discarded due to the radical scavenging capacity in the case of TRIS, and the strong iron(III) complexation ability in the case of phosphate buffer at a concentration level of 50 mM. Secondly, the reaction between NBT and AscH⁻ is significant and depends on the pH and the concentration of AscH⁻ as shown in Figure S5. Because of these interfering side effects, the NBT test was discarded.



Scheme S1. Reaction between NBT and superoxide anion radical $(O_2^{\bullet-})$.



Figure S8. The formation of NBT-diformazan followed at 560 nm as a function of time, showing the pH dependence of the reaction between NBT and AscH⁻. [AscH⁻]₀ = 300 μ M and [NBT] = 500 μ M in water (pH was adjusted by addition of 0.1 M NaOH).

3.C. AMPLEX RED ASSAY



Scheme S2. Reaction between Amplex Red and H₂O₂.

Table S3. Formed H₂O₂ concentrations (standard deviations are in parentheses) after 45 min reaction at pH 6.0 (50 mM MES buffer or water). [AscH⁻]₀ = 200 μ M, [Ligand]₀ = 50 μ M and [Fe(III)]₀ = 50 μ M, T = 298 K.

	pH 6.0 in MES		pH 6.0 in water	
	$[H_2O_2]$	Formed H ₂ O ₂ /	$[H_2O_2]$	Formed H ₂ O ₂ /
	(µM)	complex ratio (%)	(µM)	complex ratio (%)
Fe(III):FD ₃ K	0.9(3)	1.8	2.0(2)	4.0
Fe(III):FE ₃ K	1.1(2)	2.2	1.8(2)	3.6
Fe(III):EDTA	9.5(6)	19.0	16.7(4)	33.4
Fe(III):DFO	2.3(3)	4.6	1.9(2)	3.8

Table S4. Formed H2O2 concentrations (standard deviations are in parentheses) after 45 min reaction at pH 7.4(50 mM HEPES buffer or water). [AscH⁻]0 = 200 μ M, [Ligand]0 = 50 μ M and

	$[Fe(III)]_0 = 50 \ \mu M, T = 298 \ K.$			
	pH 7.4 in HEPES		pH 7.4 in water	
	[H ₂ O ₂] Formed H ₂ O ₂ /		$[H_2O_2]$	Formed H ₂ O ₂ /
	(µM)	complex ratio (%)	(µM)	complex ratio (%)
Fe(III):FD ₃ K	0.87(9)	1.8	1.2(1)	2.3
Fe(III):EDTA	7.0(8)	14.0	8.2(2)	16.4
Fe(III):DFO	0.81(4)	1.6	1.9(3)	3.8



Figure S9. Absorbance at 570 nm *versus* time graph for the H_2O_2 formation at pH 7.4 (50 mM HEPES buffer). [AscH⁻]₀ = 200 μ M, [Ligand]₀ = 50 μ M and [Fe(III)]₀ = 50 μ M, T = 298 K.

3.D. COUMARIN-3-CARBOXYILIC ACID (3-CCA) ASSAY



Scheme S3. Reaction between 3-CCA and hydroxyl radical (HO[•]).

Table S5. Formed HO[•] concentrations (standard deviations are in parentheses) after 30 min reaction at pH 6.0 and 7.4 in water. [AscH⁻]₀ = 125 μ M, [Ligand]₀ = 50 μ M and [Fe(III)]₀ = 50 μ M in the case of peptides (FD₃K, FE₃K) and DFO. [AscH⁻]₀ = 12.5 μ M, [Ligand]₀ = 5.0 μ M and [Fe(III)]₀ = 5.0 μ M in the case of EDTA, *T* = 298

К.				
	pH 6.0 in water		pH 7.4 in water	
	[HO•]	Formed HO•/	[HO•]	Formed HO•/
	(µM)	complex ratio (%)	(µM)	complex ratio (%)
Fe(III):FD ₃ K	0.033(2)	0.07	0.25(1)	0.50
Fe(III):FE ₃ K	0.051(2)	0.10	-	-
Fe(III):EDTA	0.31(4)	6.20	0.48(6)	9.60
Fe(III):DFO	0.062(5)	0.12	0.07(1)	0.14



Figure S10. Normalised emission at 450 nm ($\lambda_{ex} = 395$ nm) *versus* time graph for the HO[•] radical formation at pH 7.4 in water. [AscH⁻]₀ = 125 μ M, [Ligand]₀ = 50 μ M and [Fe(III)]₀ = 50 μ M in the case of FD₃K and DFO. [AscH⁻]₀ = 12.5 μ M, [Ligand]₀ = 5.0 μ M and [Fe(III)]₀ = 5.0 μ M in the case of EDTA, *T* = 298 K.