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Supplementary Information

Synthesis and characterization of mono S-lipidated peptide hydrogels: a

platform for the preparation of reactive oxygen species responsive materials

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1. Peptide Synthesis

1.1 Synthesis of 3



Peptide **3** was synthesised using Fmoc SPPS according to **general method A** and **B**, then purified according to **general method C**. Purified peptide **3** was obtained as a white solid after lyophilization (70.1% yield); $R_t = 9.1 \text{ min}; m/z$ (ESI-MS) 1028.4 ([M+H]⁺ requires 1028.4), Fig.S1.



Figure S1. RP-HPLC and ESI-MS traces of **3** (*ca.* 96% purity as judged by peak area of RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm × 150 mm; 5 μ m), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

1.2 Synthesis of 5a



Peptide **3** was synthesised using Fmoc SPPS following **general method A** and **B**. The crude product was purified following **general method C**. Purified **3** (10 mg, 10 µmol) was dissolved in 1 mL NMP solution, containing vinyl butyrate (23 µL, 0.2 mmol, 20 eq.), TIPS, TFA and *t*-nonyl mercaptan as per **general method D** for 15 min. The crude product of **5a** was purified following **general method C**. Purified lipidated peptide **5a** was obtained as a white solid after lyophilization (55.0 % yield); $R_t = 12.1$ min; m/z (ESI-MS) 1143.0 ([M+H]⁺ requires 1142.5), **Fig. S2**.



Figure S2. RP-HPLC and ESI-MS traces of **5a** (*ca.* 98% purity as judged by peak area of RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm × 150 mm; 5 μ m), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

1.3 Synthesis of 5b



Peptide **3** was synthesised using Fmoc SPPS following **general method A** and **B**. The crude product was purified following **general method C**. Purified **3** (10 mg, 10 µmol) was dissolved in 1 mL NMP solution, containing vinyl hexanoate (32 µL, 0.2 mmol, 20 eq.), TIPS, TFA and *t*-nonyl mercaptan as per **general method D** for 15 min. The crude product of **5b** was purified following **general method C**. Lyophilisation yielded purified **5b** as a white solid 51.3% yield); $R_t = 14.1 \text{ min}$; m/z (ESI-MS) 1170.5 ([M+H]⁺ requires 1170.6), **Fig. S3**.



Figure S3. RP-HPLC and ESI-MS traces of **5b** (*ca.* 99% purity as judged by peak area of RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm × 150 mm; 5 μ m), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

1.4 Synthesis of 5c



Peptide **3** was synthesised using Fmoc SPPS following **general method A** and **B**. The crude product was purified following **general method C**. Purified **3** (10 mg, 10 µmol) was dissolved in 1 mL NMP solution, containing vinyl octanoate (38 µL, 0.2 mmol, 20 eq.), TIPS, TFA and *t*-nonyl mercaptan as per **general method D** for 15 min. The crude product of **5c** was purified following **general method C**. The purified lipidated peptide **5c** was obtained as a white solid after lyophilization (50.2 % yield); $R_t = 15.9$ min; m/z (ESI-MS) 1198.5 ([M+H]⁺ requires 1198.6), **Fig. S4**.



Figure S4. RP-HPLC and ESI-MS traces of **5c** (*ca.* 99% purity as judged by peak area of RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm × 150 mm; 5 μ m), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.



Peptide **3** was synthesised using Fmoc SPPS following **general method A** and **B**. The crude product was purified following **general method C**. Purified **3** (10 mg, 10 µmol) was dissolved in 1 mL NMP solution, containing vinyl decanoate (45 µL, 0.2 mmol, 20 eq.), TIPS, TFA and *t*-nonyl mercaptan as per **general method D** for 15 min. The crude product of **5d** was purified following **general method C**. Purified lipidated peptide **5d** was obtained as a white solid after lyophilization (52.8 % yield); $R_t = 17.5$ min; m/z (ESI-MS) 1227.1 ([M+H]⁺ requires 1226.6), **Fig. S5**.



Figure S5. RP-HPLC and ESI-MS traces of **5d** (*ca.* 99% purity as judged by peak area of RP-HPLC at 210 nm); Agilent ZORBAX 300SB-C3, $(3 \text{ mm} \times 150 \text{ mm}; 3.5 \text{ }\mu\text{m})$, linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

1.6 Synthesis of 5e



Peptidyl **3** was synthesised using Fmoc SPPS following **general method A** and **B**. The crude product was purified following **general method C**. Purified **3** (10 mg, 10 µmol) was dissolved in 1 mL NMP solution, containing vinyl palmitate (65 µL, 0.2 mmol, 20 eq.), TIPS, TFA and *t*-nonyl mercaptan as per **general method D** for 15 min. The crude product of **5e** was purified following **general method C**. Purified lipidated peptide **5e** was obtained as a white solid after lyophilization (51.8 % yield); $R_t = 23.4$ min; m/z (ESI-MS) 1310.8 ([M+H]⁺ requires 1310.7), **Fig. S6**.



Figure S6. RP-HPLC and ESI-MS traces of **5e** (*ca.* 99% purity as judged by peak area of RP-HPLC at 210 nm); Agilent ZORBAX 300SB-C3, (3 mm × 150 mm; 3.5 µm), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

1.7 Synthesis of 5f



Peptidyl resin **2** was synthesised using Fmoc SPPS following **general method A** and **E**. The crude product of **5f** was purified following **general method C**. Purified lipidated peptide **5f** was obtained as a white solid after lyophilization (50.0 % yield); $R_t = 17.9$ min; m/z (ESI-MS) 1136.7 ([M+H]⁺ requires 1136.7), **Fig. S7**.



Figure S7. RP-HPLC and ESI-MS traces of **5f** (*ca.* 98% purity as judged by peak area of RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm × 150 mm; 5 μ m), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

1.8 Oxidation of peptide 5b in the presence of H_2O_2



Purified peptide **5b** was dissolved in the 100 mM H_2O_2 at a 2 wt% concentration at 37 °C. Reaction progress was monitored by RP-HPLC analysis (**Fig. S8**). Upon reaction completion, the desired peptide was purified using **General method C**. Purified **5g** was obtained as a white solid after lyophilization in a 70 % yield; R_t = 12.5 min; m/z (ESI-MS) 1187.0 ([M+H]⁺ requires 1185.6), **Fig. S9**.



Figure S8. RP-HPLC of an aliquot of 2wt% hydrogel **5b** containing 100 mM (left) and 20 mM (right) H_2O_2 at pH 7.0 after a) 1 min, b) 3 h and c) 24 h and 36 h (RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm × 150 mm; 5 µm), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.



Figure S9. RP-HPLC and ESI-MS traces of **5g** (*ca.* 98% purity as judged by peak area of RP-HPLC at 210 nm); XTerra® MS C18, (4.6 mm \times 150 mm; 5 μ m), linear gradient of 5% B to 95% B over 30 min, *ca.* 3% B per minute at r.t. 1.0 mL/min.

-100

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5.0 %

2. Other Figures



Figure S10. Storage (*G'*) and viscous (*G''*) moduli of a 2 wt% hydrogel of **5b** in water pH 7 at 37 °C as a function of time ($\omega = 6 \text{ rad.s}^{-1}$ and $\gamma = 0.2\%$ strain).



Figure S11. Storage (*G'*) and loss moduli (*G''*) of a 2 wt% hydrogel of peptide **5b** in H₂O at pH 7 as a function of frequency ($\gamma = 0.2\%$ strain). After first dynamic time sweep measurement (**Fig. 1**, region I) (left) and after temperature ramp sweep experiments (**Fig. 1**, region IV) (right).



Figure S12. Storage (*G'*) and loss moduli (*G''*) of a 2 wt% hydrogel of peptide **5b** in H₂O at pH 7 as a function of strain (frequency = 6 rad.s⁻¹). After first dynamic time sweep measurement (**Fig. 1**, region I) (left) after temperature ramp sweep experiments (**Fig. 1**, region IV) (right).



Figure S13. Thioflavin (ThT) fluorescence assay plots of 5a, 5c, 5d, 5e, and 5f. The fluorescence intensity was normalized relative to the one of the ThT fluorescence assay of 5b at the highest peptide concentration.



Figure S14. Circular dichroism spectrum of a 0.05 wt% solution of peptides **5a-5f** in H₂O at pH 7. The solution of **5b**was prepared by dilution of a 2 wt% hydrogel sample. The solutions of peptides **5a** and **5c-5f** were prepared from a 5 wt% solution aged for 12 h and then diluted to 0.05 wt%. All the spectra were acquired right after sample preparation.



Figure S15. AFM phase image of a 2 wt% sample of peptide 5b showing a fibrillar nanostructure.



Figure S16. SAXS profiles of samples of 5b prepared at 2 wt % (red open squares) and 1 wt % (green open triangle).



Figure S17. TEM image of 5b sample at pH 7 prepared from 2 wt% hydrogel for 12 h and then diluted to 1 wt%.



Figure S18. TEM image of **5b** at 1 wt% concentration aqueous solution at a) pH 2, b) pH 4, c) pH 6, pH 8 d) pH 10. The pH was adjusted by the addition of 1 M aqueous HCl or NaOH.



Figure S19. TEM image of samples of **5b** in water at 0.1 wt% (left) and 0.05 wt% (right). The yellow arrows point to examples of some fibres from which the width was measured.



Figure S20. TEM images of 5a, 5c, 5d, 5e and 5f samples prepared from a 5 wt% solution for 12 h and then diluted to 1 wt%.



Anti-parallel β-sheet

Parallel β-sheet



Figure S21. Anti-parallel and parallel β -sheet assembled structures of peptide 5b.



Figure S22. Predicted log D values for the side chain groups and S-lipidated fragment of 5b.



Figure S23. Storage (*G'*) and loss (*G''*) modulus of a 3 wt% hydrogel of **5b** in water and after addition of H₂O₂ 100 mM (indicated by the solid grey line) to give a 2 wt% final concentration as a function of time ($\omega = 6 \text{ rad.s}^{-1}$ and $\gamma = 0.2\%$ strain).



Figure S24. TEM image of a sample of the reaction mixture of 5b with H_2O_2 after oxidation completion. The concentration of the peptide in the sample deposited in the grid was 0.4 wt%



Figure S25. ATR-FTIR of a 2 wt% hydrogel sample of **5b**, 1 min after (blue), 3 h (orange), 24 h (grey) after addition of H_2O_2 ; and of a 2 wt% solution of purified peptide **5g** (yellow).



Figure S26. Circular dichroism spectrum of a solution of peptide 5g at a 0.05 wt% concentration.



Figure S27. Thioflavin (ThT) fluorescence assay plot of **5g**. The fluorescence intensity was normalized relative to the one of the ThT fluorescence assay of **5b** at the highest peptide concentration.

3. References

1 C. A. Schneider, W. S. Rasband and K. W. Eliceiri, *Nat. Methods*, 2012, 9, 671-675.