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Supporting Information

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1. Preparation of Zn-TPP-Ald

TPP-AId was solubilized in refluxing THF, and $Zn(OAc)_2$ (5 equivalents) was added. After 3 hours, the solvent was removed and the crude material was solubilized in dichloromethane and filtered on silica gel to obtain the desired **Zn-TPP-AId**. ¹H NMR (400 MHz, Acetone) δ 10.43 (s, 4H), 8.87 (s, 8H), 8.45 (d, *J* = 7.9 Hz, 8H), 8.37 (d, *J* = 8.3 Hz, 8H). MS (ESI) m/z calculated C₄₈H₂₈N₄O₄Zn [M]⁺: 788.14; observed: 788.10.



Figure S1: 1H NMR spectrum of Zn-TPP-Ald.



Figure S2: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of Zn-TPP-Ald.

2. Peptide syntheses

2.1. Synthetic route



Scheme S1: Synthetic route used for preparing the protected and deprotected peptide hydrazide building blocks.

2.2. Characterization data

Cys(Trt)-Hyd: Isolated by preparative reverse-phase HPLC (linear gradient from 95/5 A/B eluents to 5/95 A/B eluents in 45 min). ¹H NMR (400 MHz, MeOD) δ 7.44 - 7.40 (m, 6H), 7.34 (dd, *J* = 8.7, 6.7 Hz, 6H), 7.29 - 7.25 (m, 3H), 3.59 (dd, *J* = 6.0, 3.9 Hz, 1H, Ha), 2.69 (dd, *J* = 12.4, 7.3 Hz, 1H, Ha), 2.58 (dd, *J* = 12.4, 6.5 Hz, 1H, Ha). MS (ESI) m/z calculated C₂₂H₂₃N₃OS [M+H]⁺: 378.168; observed: 378.10.



Figure S3: ¹H NMR (top) and COSY NMR (bottom) spectra of Cys(Trt)-Hyd.



Figure S4: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of Cys(Trt)-Hyd.

Cys-Hyd: ¹H NMR (400 MHz, D₂O) δ 4.35 (t, *J* = 5.7 Hz) & 4.17 (t, *J* = 6.0 Hz) (1H), 3.18 - 2.98 (m, 2H). MS (ESI) m/z calculated C₃H₉N₃OS [M+H]⁺: 136.058; observed: 136.06.



Figure S5: ¹H NMR (top) and COSY NMR (bottom) spectra of Cys-Hyd.



Figure S6: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of Cys-Hyd.

Arg(Pbf)Cys(Trt)-Hyd: Isolated by preparative reverse-phase HPLC (linear gradient from 95/5 A/B eluents to 5/95 A/B eluents in 30 min, keeping this composition for 10 more min). ¹H NMR (400 MHz, CD₃OD) δ 7.35 (t, J = 1.8 Hz, 2H), 7.33 (dd, J = 1.5, 0.9 Hz, 4H), 7.28 (t, J = 1.5 Hz, 2H), 7.26 (t, J = 2.0 Hz, 2H), 7.25 – 7.23 (m, 2H), 7.21 (t, J = 1.4 Hz, 1H), 7.19 (t, J = 2.4 Hz, 1H), 7.18 – 7.17 (m, 1H), 4.22 (t, J = 7.5 Hz, 1H), 3.86 (t, J = 6.4 Hz, 1H), 3.20 – 3.09 (m, 2H), 2.95 (s, 2H), 2.65 – 2.55 (m, 2H), 2.53 (s, 3H), 2.46 (s, 3H), 2.04 (s, 3H), 1.87 – 1.75 (m, 2H), 1.65 – 1.50 (m, 2H), 1.41 (s, 6H). MS (ESI) m/z calculated C₄₁H₅₁N₇O₅S₂; [M+H]⁺: 786.348; observed: 786.40.



Figure S7: ¹H NMR (top) and COSY NMR (bottom) spectra of Arg(Pbf)Cys(Trt)-Hyd.



Figure S8: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of Arg(Pbf)Cys(Trt)-Hyd.

Arg-Cys-Hyd: ¹H NMR (400 MHz, D₂O) δ 4.63 (t, *J* = 5.1 Hz) & 4.50 (t, *J* = 6.7 Hz) (1H), 4.09 (t, *J* = 5.4 Hz, 1H), 3.22 (t, *J* = 5.9 Hz, 2H), 3.05 - 2.81 (m, 2H), 1.94 (m, 2H), 1.65 (m, 2H). MS (ESI) m/z calculated C₉H₂₁N₇O₂S; [M+H]⁺: 292.155; observed: 291.70.



Figure S9: ¹H NMR (top) and COSY NMR (bottom) spectra of ArgCys-Hyd.



Figure S10: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of ArgCys-Hyd.

BA-Arg(Pbf)Cys(Trt)-Hyd : Isolated by preparative reverse-phase HPLC (linear gradient from 80/20 A/B eluents to 5/95 A/B eluents in 30 min, keeping this composition for 10 more min). ¹H NMR (400 MHz, CD₃OD) δ 7.82 (d, *J* = 8.2 Hz, 2H), 7.75 (br, 2H), 7.36 – 7.33 (m, 6H), 7.27 – 7.17 (m, 9H), 4.51 (dd, *J* = 8.6, 5.9 Hz, 1H), 4.10 (t, *J* = 7.2 Hz, 1H), 3.27 – 3.13 (m, 2H), 2.95 (s, 2H), 2.71 – 2.61 (m, 2H), 2.55 (s, 3H), 2.49 (s, 3H), 1.95 – 1.75 (m, 2H), 1.71 – 1.55 (m, 2H), 1.42 (s, 6H). MS (ESI) m/z calculated C₄₈H₅₆BN₇O₈S₂; [M+H]⁺: 934.378; observed: 934.25.



Figure S11: ¹H NMR (top) and COSY NMR (bottom) spectra of BA-Arg(Pbf)Cys(Trt)-Hyd.



Figure S12: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of BA-Arg(Pbf)Cys(Trt)-Hyd.

BA-ArgCys-Hyd: ¹H NMR (400 MHz, D₂O) δ 7.85 (dt, *J* = 6.4, 2.0, 2H), 7.78 (dt, *J* = 8.4, 2.0, 2H), 4.55 (ddd, *J* = 8.7, 6.0, 3.6 Hz, 2H), 3.24 (t, *J* = 6.9 Hz, 2H), 2.95 (dq, *J* = 14.1, 6.7 Hz, 2H), 2.01 - 1.83 (m, 2H), 1.80 - 1.64 (m, 2H). MS (ESI) m/z calculated C₁₆H₂₆BN₇O₅S [M+H]⁺: 440.18; observed: 441.15.



Figure S13: ¹H NMR (top) and COSY NMR (bottom) spectra of BA-ArgCys-Hyd.



Figure S14: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of BA-ArgCys-Hyd.

PPh₃-Cys(Trt)-Hyd: Isolated by preparative reverse-phase HPLC (linear gradient from 95/5 A/B eluents to 5/95 A/B eluents in 40 min). ¹H NMR (400 MHz, CD₃OD) δ 7.89 – 7.83 (m, 3H), 7.80 – 7.76 (m, 4H), 7.76 – 7.69 (m, 8H), 7.37 – 7.33 (m, 6H), 7.25 (dt, *J* = 13.8, 4.8 Hz, 6H), 7.21 (dt, *J* = 9.5, 4.2 Hz, 3H), 4.16 (dd, *J* = 7.9, 6.7 Hz, 1H), 3.51 – 3.33 (m, 2H), 2.66 – 2.54 (m, 2H), 2.29 (q, *J* = 7.7 Hz, 2H), 1.84 (m, 2H), 1.70 (m, 2H). MS (ESI) m/z calculated C₄₅H₄₅N₃O₂PS [M+H]⁺: 723.308; observed: 722.15.



Figure S15: ¹H NMR (top) and COSY NMR (bottom) spectra of PPh₃-Cys(Trt)-Hyd.



Figure S16: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of PPh₃-Cys(Trt)-Hyd.

PPh₃-Cys-Hyd: ¹H NMR (400 MHz, D₂O) δ 7.70 (dd, *J* = 5.5, 1.7 Hz, 3H), 7.63 – 7.49 (m, 12H), 4.38 (dd, *J* = 8.1, 5.2 Hz, 1H), 3.17 (br, 2H), 2.72 (ddd, *J* = 22.3, 14.2, 6.7 Hz, 2H), 2.25 (m, 2H), 1.71 (br, 2H), 1.58 (br, 2H). MS (ESI) m/z calculated C₂₆H₃₁N₃O₂PS [M+H]⁺: 481.198; observed: 480.15.



Figure S17: ¹H NMR (top) and COSY NMR (bottom) spectra of PPh₃-Cys-Hyd.



Figure S18: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of PPh₃-Cys-Hyd.

Arg(Pbf)Ser('Bu)-Hyd: Isolated by preparative reverse-phase HPLC (linear gradient from 95/5 A/B eluents to 5/95 A/B eluents in 30 min, keeping this composition for 10 more min). ¹H NMR (400 MHz, CD₃OD) : δ 4.56 (t, J = 5.8 Hz, 1H), 4.01 (t, J = 6.4 Hz, 1H), 3.67 (qd, J = 9.2, 5.8 Hz, 2H), 3.21 (t, J = 6.6 Hz, 2H), 3.00 (s, 2H), 2.57 (s, 3H), 2.51 (s, 3H), 2.08 (s, 3H), 1.91 (m, 2H), 1.64 (m, 2H), 1.45 (s, 6H), 1.20 (s, 9H). MS (ESI) m/z calculated C₂₆H₄₅N₇O₆S; [M+H]⁺: 584.328; observed: 584.55.



Figure S19: ¹H NMR (top) and COSY NMR (bottom) spectra of Arg(Pbf)Ser(tBu)-Hyd.



Figure S20: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of Arg(Pbf)Ser(^tBu)-Hyd.

ArgSer-Hyd: ¹H NMR (400 MHz, D₂O): δ 4.65 (t, *J* = 5.5 Hz) & 4.54 (t, *J* = 5.4 Hz) (1H), 4.14 (d, *J* = 2.5 Hz, 1H), 3.98 - 3.86 (m, 2H), 3.26 (t, *J* = 6.6 Hz, 2H), 2.05 - 1.91 (m, 2H), 1.79 - 1.61 (m, 2H). MS (ESI) m/z calculated C₉H₂₁N₇O₃ [M+H]⁺: 276.178; observed: 276.00.



Figure S21: ¹H NMR (top) and COSY NMR (bottom) spectra of ArgSer-Hyd.



Figure S22: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of ArgSer-Hyd.

Arg₂(Pbf)₂Ser(^tBu)-Hyd: Isolated by preparative reverse-phase HPLC (linear gradient from 80/20 A/B eluents to 5/95 A/B eluents in 30 min, keeping this composition for 10 more min). ¹H NMR (400 MHz, CD₃OD) δ 4.24 (t, J = 5.5 Hz, 1H), 4.20 (dd, J = 8.4, 5.1 Hz, 1H), 3.74 (t, J = 6.2 Hz, 1H), 3.37 (ddd, J = 15.3, 9.2, 5.6 Hz, 2H), 2.93 (t, J = 6.5 Hz, 4H), 2.70 (s, 4H), 2.28 (s, 6H), 2.22 (s, 6H), 1.79 (s, 6H), 1.60 (m, 3H), 1.40 (m, 5H), 1.16 (s, 12H), 0.88 (s, 9H). MS (ESI) m/z calculated C₄₅H₇₄N₁₁O₁₀S₂ [M+H]⁺: 992.506; observed: 992.40.



Figure S23: ¹H NMR (top) and COSY NMR (bottom) spectra of Arg₂(Pbf)₂Ser(^tBu)-Hyd.



Figure S24: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of Arg₂(Pbf)₂Ser(^tBu)-Hyd.

Arg₂Ser-Hyd^{: 1}H NMR (400 MHz, D₂O) δ 4.49 (br, 1H), 4.39 (br, 1H), 4.04 (br, 1H), 3.86 (br, 2H), 3.19 (br, 4H), 1.87 (br, 4H), 1.64 (br, 4H). MS (ESI) m/z calculated $C_{15}H_{33}N_{11}O_4$ [M+H]⁺: 432.278; observed: 432.20.



Figure S25: ¹H NMR (top) and COSY NMR (bottom) spectra of Arg₂Ser-Hyd.



Figure S26: HPLC chromatogram (top) and mass spectrometry analysis (bottom) of $Arg_2Ser-Hyd$.

3. Characterization of TPP conjugates and cages



Figure S27: Representative ¹H NMR spectra of TPP Conjugates and TPP Cages in DMSO-d₆.



Figure S28: Representative HPLC chromatograms of TPP Conjugates and TPP Cages.

Zn-CAGE-PPh3



Figure S29: Representative HPLC chromatograms of Zn-TPP Conjugates and Zn-TPP Cages.



Figure S30: HPLC chromatogram (top) and MALDI-TOF mass spectrometry analysis (bottom) of CAGE-H.



Figure S31: HPLC chromatogram (top) and MALDI-TOF mass spectrometry analysis (bottom) of CAGE-BA.



Figure S32: HPLC chromatogram of CAGE-PPh3.



Figure S33: HPLC chromatogram of Zn-CAGE-H.



Figure S34: HPLC chromatogram analysis of Zn-CAGE-Arg.



Figure S35: HPLC chromatogram of Zn-CAGE-BA.



Figure S36: HPLC chromatogram of Zn-CAGE-PPh3.



Figure S37: HPLC chromatogram (top) and MALDI-TOF mass spectrometry analysis (bottom) of Zn-TPP-Arg4.





3. DOSY NMR

Entry	Compounds	log(D/m²s¹)	D (m²s¹)	R _{hyd} [Å]	V _{sph} [ų]
-		0.04	0.000.40-10	10	400 F
1.	IPP-Ald	-9.64	2.239×10 ¹⁰	4.9	492.5
2.	CAGE-H	-10.23	9.484×10 ⁻¹¹	11.5	6367.39
3.	TPP-Arg₄	-10.167	6.807×10 ⁻¹¹	16.1	17472.1
4.	TPP-Arg ₈	-10.336	4.613×10 ⁻¹¹	23.7	52279.61
5.	Zn-CAGE-H	-9.82	1.513×10 ⁻¹⁰	7.2	1560.17
6.	Zn-TPP-Arg₄	-9.897	1.267×10 ⁻¹⁰	8.6	2681.57
7.	Zn-TPP-Arg ₈	-10.07	8.511×10 ⁻¹¹	12.8	8766.09

Table S1: Hydrodynamic radii and spherical volumes calculated using the Stokes-Einstein equation.



Figure S39: DOSY NMR spectra.

4. UV and Fluorescence analyses



Figure S40: UV-Vis absorption (solid lines) and fluorescence (dashed lines) spectra in DMSO of representative free base (M=2H) (top) and metallated (M=Zn) (bottom) porphyrin derivatives.



Figure S41: UV-Vis absorption (solid lines) and fluorescence (dashed lines) spectra in H₂O of representative free base (M=2H) (top) and metallated (M=Zn) (bottom) porphyrin derivatives.

5. Singlet oxygen generation



Figure S42: Singlet oxygen generation probed by UV absorption spectroscopy using ABDA as a probe at varying light irradiation times (green light, λ = 525 nm).

6. PDT on cells



Figure S43: PDT study in human breast cancer (MCF-7) cells incubated with 1.31μ M of compounds for 24 h then exposed to green light for different irradiation times. Data are presented as mean ± SEM.

7. ROS production in cells



Figure S44: Fluorescence microscopyimaging of ROS using DCFDA assayin MCF-7 cells treated (or not) with 1.31 µM of compounds for 24 h, then, exposed (or not) to green light irradiation for 10 min.

8. Cellular uptake kinetics



Figure S45: Uptake kinetics study of compounds incubated with human breast cancer (MCF-7) cells at 1.31 μ M concentration for different time intervals. Data are presented as mean \pm SEM.