

Supporting Information

Developing Chemically Modified Redox-responsive Protein As Smart Therapeutics

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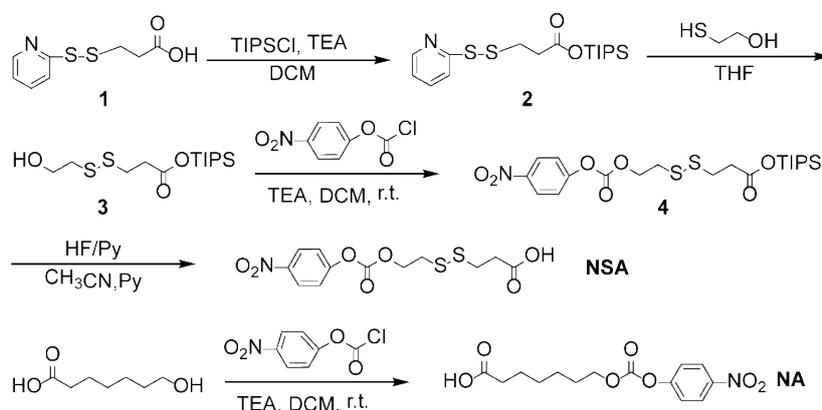
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General

All chemical used for lipid synthesis and protein modification were purchased from Aladdin, TCI or Sigma-Aldrich and used as received. RNase A was purchased from Sigma, the ribonuclease activity was assayed according to previous report,¹ the intracellular GSH level was determined using DTNB assay according to previous report.²⁻⁴ The lipids were synthesized and characterized according to our previous reports.⁵

Synthesis of NSA and NA



Scheme S1. Synthetic route of NSA and NA for protein modification.

Synthesis of **2**: 3-(pyridine-2-yl)propanoic acid **1** was synthesized following an early report⁶: ¹H NMR (400 MHz, CDCl₃) δ 8.43 (d, *J* = 4.7 Hz, 1H), 7.67 (d, *J* = 8.1 Hz, 1H), 7.64 – 7.56 (m, 1H), 7.09 – 7.02 (m, 1H), 3.01 (t, *J* = 7.0 Hz, 2H), 2.77 (t, *J* = 7.1 Hz, 2H), 1.27 (dt, *J* = 14.8, 7.5 Hz, 3H), 1.04 (d, *J* = 7.5 Hz, 21H). To a solution of **1** (6.8g, 31.6 mmol) and triethylamine (TEA, 4.6 mL, 38.00 mmol) in CH₂Cl₂ (20 mL) was added chlorotriisopropylsilane (TIPSCl, 6.09g, 31.6 mmol) dropwise at 0 °C. After stirring at room temperature for 4 h, the reaction mixture was dried by rotary evaporator, then diluted with ethyl acetate (100 mL). The organic layer was separated, washed by saturated aqueous NaHCO₃ and brine, dried over MgSO₄, and concentrated under reduced pressure. Flash chromatography of the residue on silica gel using Petroleum ether /EtOAc (10/1) as the eluent gave **2** (11 g, 98 % yield) as colorless oil: ¹H NMR (400 MHz, CDCl₃) δ 8.43 (d, *J* = 4.7 Hz, 1H), 7.67 (d, *J* = 8.1 Hz, 1H), 7.64 – 7.56 (m, 1H), 7.09 – 7.02 (m, 1H), 3.01 (t, *J* = 7.0 Hz, 2H), 2.77 (t, *J* = 7.1 Hz, 2H), 1.27 (dt, *J* = 14.8, 7.5 Hz, 3H), 1.04 (d, *J* = 7.5 Hz, 21H). ESI-MS Calcd for C₁₇H₃₀NO₂S₂Si [M+H]⁺: 372.1, Found: 372.1.

Synthesis of **3**: To a solution of **2** (5 g, 13.461 mmol) in THF (15 mL) was added 2-mercaptoethanol (1.26 g, 16.15 mmol) dropwise, the mixture stirring at room temperature overnight. The reaction mixture was concentrated in vacuum. The residue was purified on a silica gel column using petroleum/EtOAc (4/1) as the eluent to give **3** (3.87 g, 85%) as a colorless oil: ¹H NMR (300 MHz, CDCl₃) δ 3.89 (s, 2H), 2.94 (t, *J* = 6.9 Hz, 2H), 2.87 (t, *J* = 5.6 Hz,

2H), 2.80 (t, $J = 6.8$ Hz, 2H), 2.12 (s, 1H), 1.29 (dd, $J = 14.6, 7.5$ Hz, 3H), 1.08 (d, $J = 7.3$ Hz, 19H). ^{13}C NMR (75 MHz, DMSO) δ 171.16, 59.45, 41.03, 34.78, 33.39, 17.58, 11.32. ESI-MS Calcd for $\text{C}_{14}\text{H}_{30}\text{O}_3\text{S}_2\text{SiNa}$ $[\text{M}+\text{Na}]^+$: 361.1, Found: 361.0.

Synthesis of **4**: To a solution of **3** (1g, 2.96 mmol) in DCM (10 mL) was added triethylamine (0.5 mL, 0.36 g) dropwise with stirring. After stirring for 30 minutes, was added 4-Nitrophenyl chloridocarbonate (0.76 mg, 3.55 mmol) dropwise at 0 °C. The mixture stirred for 4 h, and extracted with CH_2Cl_2 . The combined organic layer was dried over MgSO_4 and concentrated under reduced pressure. Flash chromatography of the residue on silica gel using Petroleum ether /EtOAc (10/1) as the eluent gave **4** (1.2 g, 80%). ^1H NMR (300 MHz, CDCl_3) δ 8.29 (d, $J = 8.8$ Hz, 2H), 7.40 (d, $J = 8.7$ Hz, 2H), 4.54 (t, $J = 6.5$ Hz, 2H), 3.03 (t, $J = 7.1$ Hz, 2H), 2.96 (d, $J = 6.9$ Hz, 2H), 2.80 (t, $J = 6.8$ Hz, 2H), 1.31 – 1.24 (m, 3H), 1.07 (d, $J = 7.3$ Hz, 18H). ^{13}C NMR (75 MHz, CDCl_3) δ 171.62, 155.55, 152.44, 145.60, 125.47, 121.93, 67.00, 36.66, 35.63, 33.93, 17.90, 12.02. MALDI-FTICR-MS Calcd for $\text{C}_{21}\text{H}_{33}\text{NO}_7\text{S}_2\text{SiNa}$ $[\text{M}+\text{Na}]^+$: 526.13654, Found: 526.13627.

Synthesis of **NSA**: To a solution of **4** (1 g, 1.98 mmol) in pyridine (3 mL) and acetonitrile (3 mL) was added HF/pyridine (70% weight, 5 mL) at 0 °C with stirring. After stirring for 30 minutes, the reaction was quenched by adding EtOAc (50 mL). The organic layer was separated, washed by water, saturated aqueous CuSO_4 and brine, dried over MgSO_4 , and concentrated. The crude product was purified on a silica gel column (eluent: hexanes/EtOAc = 1/1) to give **5** (412.2 mg, 60% yield) as a yellow oil: ^1H NMR (300 MHz, CDCl_3) δ 8.28 (d, $J = 8.8$ Hz, 1H), 7.39 (d, $J = 8.8$ Hz, 1H), 4.54 (t, $J = 6.4$ Hz, 1H), 3.03 (t, $J = 6.4$ Hz, 1H), 2.97 (t, $J = 6.9$ Hz, 1H), 2.82 (t, $J = 6.9$ Hz, 1H). ^{13}C NMR (75 MHz, CDCl_3) δ 177.74, 155.51, 152.44, 145.60, 125.47, 121.92, 66.94, 36.72, 33.90, 32.96. ESI-MS Calcd for $\text{C}_{12}\text{H}_{13}\text{NO}_7\text{S}_2\text{Na}$ $[\text{M}+\text{Na}]^+$: 370.0, Found: 370.0.

NA is synthesized in a similar procedure to that of **NSA**. ^1H NMR (300 MHz, CDCl_3) δ 8.46 – 8.08 (m, 2H), 7.46 – 7.28 (m, 1H), 4.60 – 3.97 (m, 3H), 2.77 – 2.36 (m, 2H), 1.97 – 1.17 (m, 8H). ^{13}C NMR (75 MHz, CDCl_3) δ 175.33, 158.36, 152.85, 145.95, 125.46, 121.95, 68.25, 33.68, 29.32, 28.64, 26.31, 22.17. ESI-MS Calcd for $\text{C}_{14}\text{H}_{17}\text{NO}_7\text{Na}$ $[\text{M}+\text{Na}]^+$: 334.1, Found: 334.1.

Chemical conjugation of RNase A with NSA and NA

RNase A-NSA and RNase A-NA were prepared by reacting RNase A with an excess amount of NSA or NA. Briefly, 2 mg RNase A was dissolved in 450 μL 0.1 M NaHCO_3 buffer solution (pH = 8.5). To above solution was added 20 μL DMSO solution containing 12 mg NSA or NA dropwise. The reaction mixtures were then stirred at room temperature for additional 8 h, followed by ultrafiltration purification using Amicon® Ultra Centrifugal Filters (MWCO = 10000, Millipore, MA). The fluorescent labeled FITC-RNase A-NSA was prepared by reacting RNase A-NSA with

twice weight amount of NHS-FITC for 2 h in 0.1 M NaHCO₃ buffer solution (pH = 8.5), the mixtures were diluted by 50 mM Tris (pH = 8.0) using Desalt column and protein were concentrated and stored at -20 °C.

Activation of RNase A-NSA by GSH

RNase A-NSA or RNase A-NA were incubated with GSH at 37 °C for 2h, 4h, 6 h, respectively, followed by dialysis against DEPC water to remove excessive GSH and then concentrated by ultrafiltration purification using Amicon® Ultra Centrifugal Filters (MWCO = 10000, Millipore, MA). The proteins were then subjected to MALDI-TOF characterization and ribonuclease activity assay. To verify the selective RNase A-NSA activation by GSH, RNase A-NSA (14 mM) was incubated with several main reductive substances of cellular concentration for 6 h at 37 °C, followed by RNase A activity assay as described above. The concentrations of the reductive species were listed below: 0.1 mM Cys; 10 μM Hcy; 4 mM GSH; 1 mM AA; 250 μM Na₂S (for generation of H₂S); 200 μM NADH.

Cell culture

HeLa, SiHa, 3T3 and HEK-293T cells were purchased from National Infrastructure of Cell Line Resource (China) and cultured in high-glucose dulbecco's modified eagle medium (DMEM) containing 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin (P/S) at 37 °C in the presence of 5% CO₂. For the intracellular delivery, the cells were sub-cultured and seeded in 48-well or 96-well plate 24 h prior to experiment.

Intracellular delivery of RNase A-NSA

Cells were seeded in 96-well plate at a density of 1×10^4 per well and treated with different lipid/protein nanocomplexes for 6 h before changing fresh culture medium, the cells were incubated for additional 42 h before MTT assay to measure cell viability. For CLSM imaging study, HeLa cells were treated with PBPA-O16B (5.0 μg/mL)/FITC-RNase A-NSA (250 nM) nanocomplexes for 6 h, then the cells were washed with cold DPBS twice before fluorescence imaging. The endosome was stained using 100 nM LysoTracker@Red according to the manufacturer's instruction.

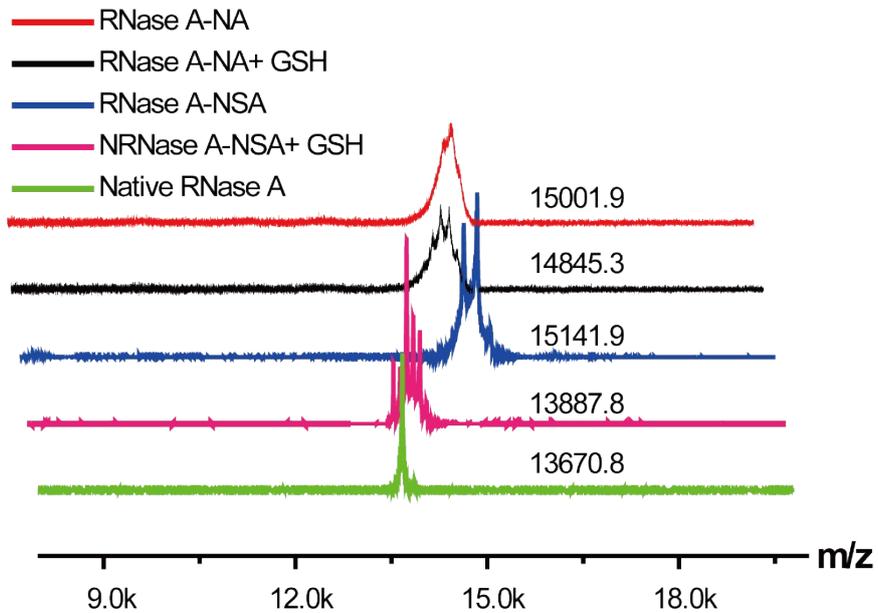


Fig. S1. Mass characterization of native RNase A and RNase A-NSA and RNase A with and without 4 mM GSH treatment.

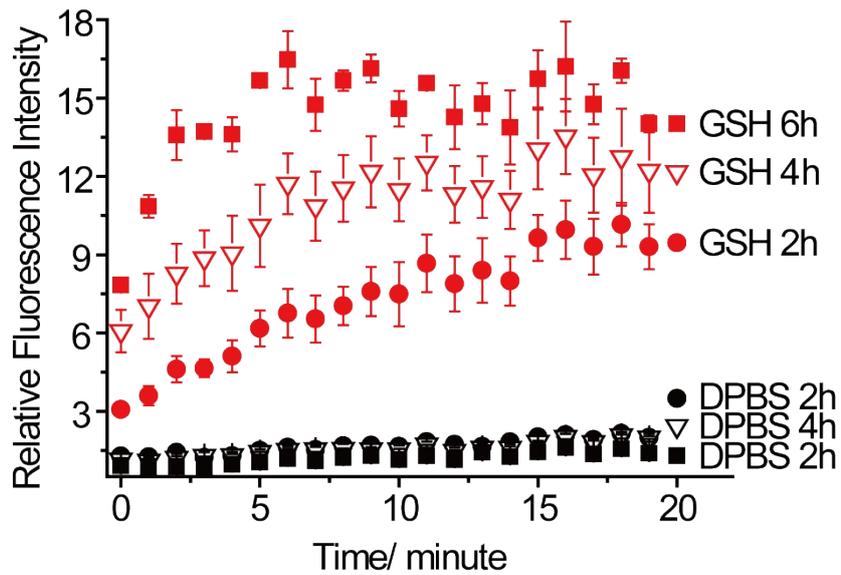


Fig. S2. Ribonuclease activity RNase A-NSA treated with 4 mM GSH at different time points. 14 mM RNase A-NSA was treated with GSH before enzyme activity assay.

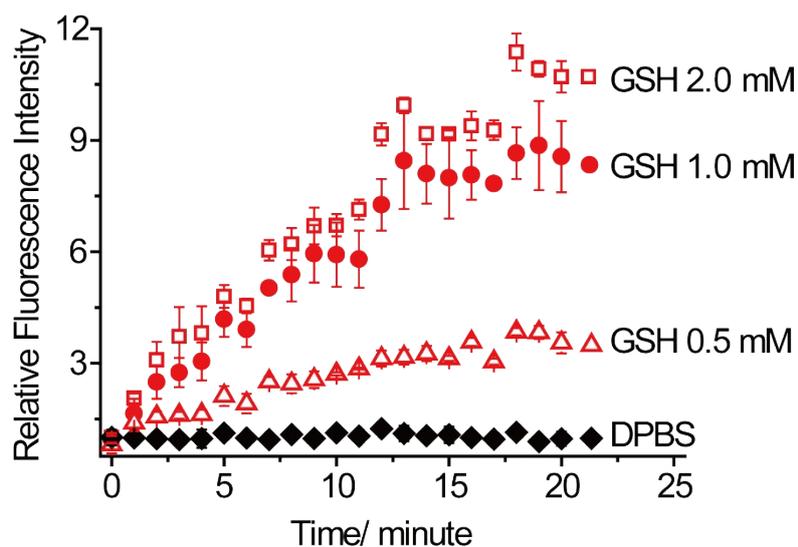
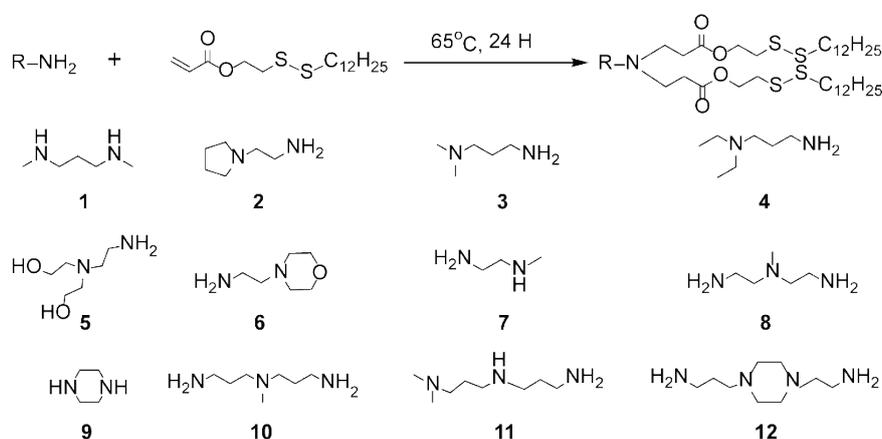


Fig. S3. Ribonuclease activity assay of RNase A-NSA treated with GSH at indicated concentration.



Scheme S2. The synthesis route and the chemical structure of amines used for RNase A delivery.

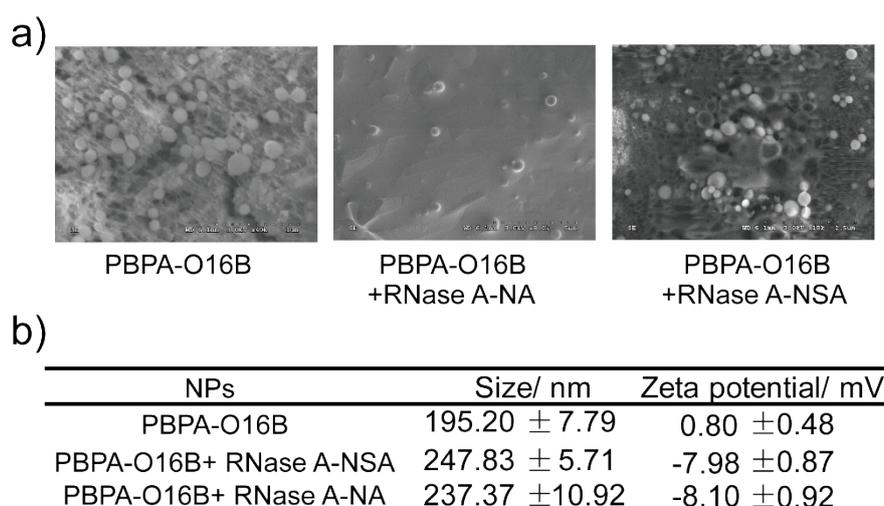


Fig. S4. (a) Diameter and zeta potential characterization of PBPA-O16B, PBPA-O16B/RNase A-NA and PBPA-O16B/RNase A-NSA nanoparticles; (b) Cryo-SEM image of PBPA-O16B, PBPA-O16B/RNase A-NA and PBPA-O16B/RNase A-NSA nanoparticles.

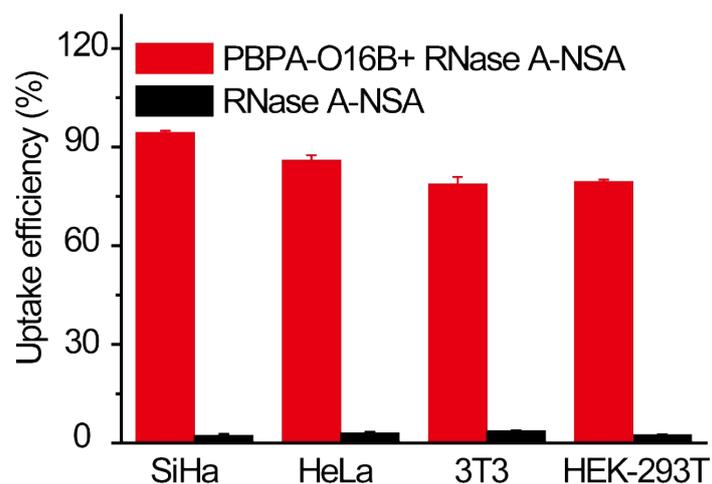


Fig. S5. Uptake efficiency of PBPA-O16B/RNase A-NSA nanoparticles on normal cell lines and cancer cell lines after treatment of 250 nM protein dose for 8 hours.

References

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