Supporting Information

A Glycoconjugate-Based Gold Nanoparticle Approach for the Targeted Eradication of *Pseudomonas aeruginosa* Biofilms

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S1. Experimental

General Information and Protocols. All purchased reagents were analytical grade and used as received. Hydrogen tetrachloroaurate (III) hydrate (HAuCl₄·XH₂O, 99.9%-Au) was purchased from J&K Chemical. Trisodium citrate was purchased from J&K Chemical. Ceftazidime was purchased from J&K Chemical. Proteins including UEA-1 (Ulex European Agglutinin 1), PNA (Peanut Agglutinin), LCA (*Lens culinaris* Lectin), SBA (Soybean Agglutinin), Con A (Conconvalin A), WGA (Wheat Germ Agglutinin), BSA (Bovine Serum Albumin) and PeP (Phospholipid Exchange Protein) were purchased from Sigma-Aldrich *P. aeruginosa* strain (ATCC 27853, ATCC BAA2110), *E. coli* (ATCC 25922) and *MRSA* (ATCC 43300) were obtained from the American Type Culture Collection (ATCC). UV-Vis absorption measurements were performed on a Varian Cary Eclipse fluorescence spectrophotometer. Dynamic Light Scattering (DLS) analysis was carried out using a Horiba LB-550 Dynamic Light Scattering nano-analyzer. Fluorescence bacteria imaging was achieved using a Nikon AIR Confocal Laser-Scanning Microscope equipped with a 100x oil-immersion objective lens.

Synthetic procedure of Fuc-AuNP/Gal-AuNP. Cyclooctyne disulfide Cyc-S, azido fucoside (Fuc-N), and azido galactoside (Gal-N) were obtained using a previously reported procedure.¹ AuNPs were prepared by adding trisodium citrate solution (38.8 mM, 5 mL) to an aqueous solution of HAuCl₄ (0.29 mM, 50 mL) that was stirring under reflux at 140 °C. The mixture was stirred under reflux for an additional 10 min and then cooled to room temperature, to obtain a red solution (AuNP). Subsequently, 10 mL of the freshly prepared AuNP solution was added to an aqueous solution of cyclooctyne disulfide Cyc-S (100 μ M, 100 μ L) and an aqueous solution of Fuc-N (100 μ M, 100 μ L) or Gal-N (100 μ M, 100 μ L). The mixture was removed to afford a residual solid product, which was re-dispersed in Tris-HCl buffer solution (20 mL, 0.01 M, pH 7.4). This process was repeated for three times to remove excessive reactants. The concentration of **Fuc-AuNP** or **Gal-AuNP** was determined as 0.8 nM via a previously established method (Anal. Chem. 2007, 79, 4215-4221). Finally, **Fuc-AuNP** and **Gal-AuNP** were used directly in the following experiments.

Lectin Binding and Dynamic Light Scattering analysis. Freshly prepared Fuc-AuNP (0.2 nM) solution was added to a cuvette for UV-Vis absorption and Dynamic Light Scattering (DLS) measurements. Then, different amounts of UEA-1 (Ulex European Agglutinin 1, (0.01, 0.03, 0.05, 0.1, 0.2 μ M)) dissolved in Tris-HCl buffer solution (0.01 M, pH 7.4) was added to the Fuc-AuNP solution and the UV-Vis absorption and DLS measurements were conducted. Similarly, the freshly prepared Gal-AuNP solution (0.2 nM) was measured UV-Vis absorption with the addition of PNA (Peanut Agglutinin). For confirming protein-selectivity, UEA-1, PNA, and several common proteins including LcA (*Lens culinaris* Lectin), SBA (Soybean Agglutinin), Con A (Conconvalin A), WGA (Wheat Germ Agglutinin), BSA (Bovine Serum Albumin), and PeP (Phospholipid Exchange Protein) (0.05 μ M), were dissolved in Tris-HCl buffer solution (0.01 M, pH 7.4) and added to 0.2 nM Fuc-AuNP/Gal-AuNP respectively, and UV-Vis changes before and after protein addition were obtained.

High-Resolution Transmission Electron Microscope. A drop of the sample was placed onto 200 mesh holey carbon copper grids. Then, the images were recorded with JEOL 2100 equipped with a Gatan Orius charged-coupled device camera and Tridiem energy filter operating at 200 kV. The three samples: AuNP (0.02 nM), **Fuc-AuNP** (0.02 nM), and **Fuc-AuNP** + UEA-1 (**Fuc-AuNP** = 0.02 nM, UEA-1 = 0.02 μ M). **Gal-AuNP** and **Gal-AuNP** + PNA were tested using the same procedure.

Procedures for CAZ loading onto Fuc-AuNP/Gal-AuNP and the evaluation of Reactive Oxygen Species (ROS) production. A solution of either Fuc-AuNP or Gal-AuNP (0.8 nM) was added to an aqueous solution of Ceftazidime (32 µg/mL). The combined mixture was stirred at room temperature for 4 h and centrifuged at 10000 rpm for 5 min. The supernatant was removed to afford residual solid products, which were washed with Tris-HCl buffer. After washing, the products were respectively redispersed in Tris-HCl buffer (0.01 M, pH 7.4) to obtain Fuc-AuNP@CAZ and Gal-AuNP@CAZ (Fuc-AuNP/Gal-AuNP = 0.8 nM, CAZ = 8 µg/mL). For ROS testing, dihydrorhodamine-123 (DHR123) was used as the ROS fluorescent probe. Fuc-AuNP (0.2 nM) was accumulated by UEA-1 (0.2 µM), and then DHR123 (6 µM) was added. The sample was subjected to light irradiation (600 nm, 1 W cm⁻²) for 90 min, and the fluorescence was measured every 15 minutes with an excitation of 485 nm.

Raman Spectroscopy. Raman spectra of each sample was obtained using a Renishaw In Via Reflex Raman system (Renishaw plc, Wotton-under-Edge, UK) equipped with a grating spectrometer, Peltier-cooled charge-coupled device detector coupled to a confocal microscope. The raw data was then processed with Renishaw WiRE 3.2 software. The Raman scattering was excited by an argon ion laser (I = 514.5 nm). Samples tested: AuNP (0.2 nM), **Fuc-AuNP** (0.2 nM), and **Fuc-AuNP@CAZ** (0.2 nM and 2 μ g/mL). **Gal-AuNP** and **Gal-AuNP@CAZ** were tested using the same procedure.

Bacteria Culture. A monocolony of *P. aeruginosa* strain (ATCC 27853, ATCC BAA2110) *E. coli* (ATCC 25922) and *MRSA* (ATCC 43300) on Luria-Bertani (LB) agar plate were transferred to LB culture medium (20 mL) and grown at 37°C for 12 h, followed by the dilution using LB medium to a density of 10⁶ cfu/mL. The bacteria sample was used in the following procedures.

Fluorophore-Loading of Fuc-AuNP/Gal-AuNP and Fluorescence imaging by Confocal Laser-Scanning Microscope (CLSM). Fuc-AuNP or Gal-AuNP (0.8 nM) was mixed with a solution of dicyanomethylene 4H-pyran (DCM)² (10 μ M) for 4 h, then centrifuged at 10000 rpm for 5 min, washed with Tris-HCl buffer, re-dispersed, to obtain Fuc-AuNP@DCM or Gal-AuNP@DCM (Fuc-AuNP/Gal-AuNP = 0.8 nM, DCM = 4 μ M). 10⁶ cfu/mL *P. aeruginosa* fluid was incubated with **Fuc-AuNP@DCM** (0.2 nM and 1μ M) for 30 min. For the competition assay, free L-fucose (10 mM) was pre-incubated with P. aeruginosa fluid at 37 °C for 30 min, before the incubation of Fuc-AuNP@DCM. After incubation, the suspensions of the two groups were centrifuged at 8000 rpm for 5 min, and the supernatants were removed. The bacteria of two groups were washed for three times by PBS buffer, suspended in pure water, placed (100 µL each) onto two microscope slides and dried at room temperature, respectively. CLSM images were obtained using a Nikon AIR Confocal Laser-Scanning Microscope equipped with a 100x oil-immersion objective lens. The excitation and emission wavelengths are at 460 nm and 600-640 nm, respectively. Then, Gal-AuNP@DCM was similarly used for imaging *P. aeruginosa* with or without pre-incubation of free D-galactose as discussed above.

Chemical structure of the DCM dye is shown below:



In vitro Bacteria Viability Test. In two 96-well plates, 10⁶ cfu/mL P. aeruginosa strain (ATCC 27853, ATCC BAA2110), E. coli (ATCC 25922) or MRSA (ATCC 43300) fluid was added to each well and treated with: Fuc-AuNP@CAZ (0.2 nM and 2 µg/mL), Fuc-AuNP (0.2 nM), Ceftazidime (2 μ g/mL), and sterile water (control). One 96-well plate was irradiated under 600 nm light (1 W cm⁻²) for 30 min, and another plate was placed in the dark for 30 min. Both plates were put in an incubator (37°C) in dark for 3 h. After incubation, the mixtures were further diluted to 10¹, 10², 10³, 10⁴, and 10⁵ times by sterile water. For viability testing, diluents of each group with different diluting times (10³, 10⁴, and 10⁵) were all dropped on one LB (Luria-Bertani) agar plate for three times (5 µL for each drop), and incubated at 37°C for 18 h. Then, the numbers of average colony forming units (CFUs) and the standard deviations (between three repetitions) of 10⁴ dilution groups were counted and calculated. Defining negative control group as 100% viability, average CFUs of all groups were transferred to relative bacterial viabilities, along with errors calculated from standard deviations. Other drops of 10³ or 10⁵ dilution groups were used to confirm result accuracy. For wholeplate culturing, the above diluent (10^4 dilution times, 100μ L) of each group was respectively spread onto one individual LB agar plate by spread plate method, then all the plates were incubated at 37°C for 18 h. Photographs of the plates were taken. Gal-AuNP@CAZ and Gal-AuNP were tested using the same procedure.

The Inhibition of Biofilms Formation.

PROCEDURE :

(1) Preparation of Precultures and Biofilms Growth.³

(2) Treatment of Biofilms. To the established *P. aeruginosa* biofilms on glass slides, sterile water (100 µL, control), Ceftazidime (32 µg/mL),¹ Fuc-AuNP@CAZ (0.2 nM and 32 µg/mL), and Gal-AuNP@CAZ (0.2 nM and 32 µg/mL) were added. Slides treated with Fuc-AuNP@CAZ or Gal-AuNP@CAZ were subjected to continuous light irradiation (600 nm, 1 W cm⁻²) for 30 min, the additional slides were placed into dark for 30 min along with control and Ceftazidime groups. After this procedure, each slice was further incubated for 24 h at 37 °C. To the established *P. aeruginosa* biofilms on steel surface, Fuc-AuNP@CAZ (0.2 nM and 32 µg/mL) and Gal-AuNP@CAZ (0.2 nM and 32 µg/mL) were added respectively. The two groups were then subjected to the same procedure as above.

(3) Staining of Biofilms. For each group, 20 μ L of staining solution (Syto9: 14 μ M and PI: 83 μ M) was added onto the glass/steel slice directly to achieve a final concentration

of 1.4 μ M of Syto9 and 8.3 μ M of PI. Each slice was placed inside the wells of a wellplate and the well-plates were covered with a fresh air-permeable foil, then the wellplates were placed into the incubator and incubated for 12 h at 37 °C before imaging. (4) Microscopy.²

(5) Data Analysis. The different proportions of blue (live bacteria) and yellow (dead bacteria) biovolumes were calculated from the analyzed imaging stacks.



S2. Additional Figures and Tables

Figure S1. Zeta Potential analysis (a,c) and Raman spectra (b,d) of AuNPs (0.2 nM), **Fuc-AuNPs** (0.2 nM), **Gal-AuNPs** (0.2 nM), **Fuc-AuNP@CAZ** (0.2 nM and 2 μ g/mL), and **Gal-AuNP@CAZ** (0.2 nM and 2 μ g/mL).



Figure S2. Plotting the UV-vis absorbance changes of 0.2 nM **Fuc-AuNPs** (a) and **Gal-AuNPs** (b) as a function of UEA-1 (0-0.05 μ M) and PNA (0-0.05 μ M) concentration for the determination of the limit of detection (3 σ /k).



Figure S3. High-Resolution Transmission Electron Microscope (HR-TEM) images: (a) AuNPs (0.02 nM) (b) **Gal-AuNPs** (0.02 nM) (c) **Gal-AuNPs** (0.02 nM) + PNA (0.02 μ M). Dynamic Light Scattering (DLS): (d) AuNPs (0.2 nM) (e) **Gal-AuNPs** (0.2 nM) (f) **Gal-AuNPs** (0.2 nM) + PNA (0.05 μ M).



Figure S4. Dynamic Light Scattering (DLS) analysis: (a) AuNPs (0.2 nM), **Fuc-AuNPs** (0.2 nM), and **Fuc-AuNPs** (0.2 nM) with the addition of different concentrations of UEA-1 (0.01-0.2 μ M). (b) AuNPs (0.2 nM), **Gal-AuNPs** (0.2 nM), and **Gal-AuNPs** (0.2 nM) with the addition of different concentrations of PNA (0.01-0.2 μ M).



Figure S5. Plotting the Zeta Potential (a) and Dynamic Light Scattering (DLS) (b) analysis of **Fuc-AuNP@CAZ** (0.2 nM and 2 µg/mL) or **Gal-AuNP@CAZ** (0.2 nM and 2 µg/mL) stored in dark during 8-day period.



Figure S6. The change in fluorescence emission intensity at 540 nm (dihydrorhodamine, 6 μ M) with increasing light irradiation time (600 nm, 0-90 min, 1 W cm⁻²). (a) **Fuc-AuNPs** (0.2 nM) with and without UEA-1 (0.2 μ M) (b) **Gal-AuNPs** (0.2 nM) with and without PNA (0.2 μ M). Change in UV-Vis absorbance at 650 nm (c) pH; (d) Temperature. Experiments were carried out using 0.2 nM **Fuc-AuNPs** or **Gal-AuNPs** and 0.5 mM TMB in Tris-HCl buffer solution (0.01 M, pH 4) and the resultant solution was irradiated with light (30 min, 600 nm, 1 W cm⁻²).



Figure S7. (a) IR thermal images of EP tube solutions of AuNPs (0.2 nM), **Fuc-AuNP@CAZ** (0.2 nM and 2 μ g/mL), and **Fuc-AuNP@CAZ** + UEA-I (**Fuc-AuNP@CAZ** = 0.2 nM and 2 μ g/mL, UEA-I = 0.1 μ M) under light irradiation (600 nm, 1 W cm⁻², 4 min). (b) Temperature changes of the EP tube solutions monitored by the IR thermal camera during irradiation.



Figure S8. Temperatures of Fuc-AuNP@UEA-1 and Gal-AuNP@PNA solutions as a function of laser irradiation time. The photothermal conversion e□ciency of Fuc-AuNP and Gal-AuNP at 600 nm were calculated to be 26.2% and 25.7% respectively.



Figure S9. (a) IR thermal images of EP tube solutions of UEA-1 (0.1 μ M) and **Fuc-AuNP@CAZ** (0.2 nM and 2 μ g/mL) bound to *P. aeruginosa* (ATCC 27853) (b) Overall temperature changes of the EP tube solutions that were monitored by the IR thermal camera during light irradiation.



Figure S10 - Time-dependent release of CAZ (measuring the absorption of CAZ at 257 nm) from (a) **Fuc-AuNP@CAZ** or (b) **Gal-AuNP@CAZ** in Tris-HCl solution (0.01 M, pH 7.4) with and without light irradiation (600 nm, 1 W cm⁻²) during 8-hour period.



Figure S11 - Time-dependent release of CAZ (measuring the absorption of CAZ at 257 nm) from (a) **Fuc-AuNP@CAZ** or (b) **Gal-AuNP@CAZ** in Tris-HCl solution (0.01 M, pH 7.4) with and without UEA-1 or PNA during 8-hour period.



Figure S12 - (a) Relative viabilities of *P. aeruginosa* (ATCC 27853) treated with AuNPs (0.2 nM), **Gal-AuNPs** (0.2 nM), and **Fuc-AuNPs** (0.2 nM) with or without 30 min irradiation by 600 nm light (1 W cm⁻²). (b) *P. aeruginosa* (ATCC 27853) colonies on Luria-Bertani (LB) agar plates treated with the respective systems with or without light irradiation (30 min, 600 nm, 1 W cm⁻²).



Figure S13 - Confocal Laser-Scanning Microscope (CLSM) images of *P. aeruginosa* (ATCC BAA2110) treated with sterile water (control), **Gal-AuNPs** (0.2 nM), **Fuc-AuNPs** (0.2 nM), Ceftazidime (CAZ, 2 μ g/mL), **Gal-AuNP@CAZ** (0.2 nM and 2 μ g/mL), **Fuc-AuNP@CAZ** (0.2 nM and 2 μ g/mL) with or without 30 min irradiation by 600 nm light (1 W cm⁻²). the performances were evaluated using the LIVE/DEAD Biofilm viability assay (PI, dead – red and Syto9[®], live – green).



Figure S14 - HR-TEM images of *P. aeruginosa*: (a) untreated, (b) **Fuc-AuNPs** (0.2 nM) incubated for 3 h, (c) **Fuc-AuNP@CAZ** (0.2 nM and 2 μ g/mL) with 30 min irradiation by 600 nm light and incubated for 3 h. Arrows indicate AuNPs.



Figure S15 - Relative viabilities of (a) *E. coli* (ATCC 25922) and (b) Methicillin-Resistant *S. aureus* (ATCC 43300) treated with sterile water (control), Ceftazidime (CAZ, 2 µg/mL), **Fuc-AuNPs** (0.2 nM), **Gal-AuNPs** (0.2 nM), **Fuc-AuNP@CAZ** (0.2 nM and 2 µg/mL), and **Gal-AuNP@CAZ** (0.2 nM and 2 µg/mL) with or without 30 min irradiation by 600 nm light (1 W cm⁻²). (c) *E. coli* (ATCC 25922) and (d) Methicillin-Resistant *S. aureus* (ATCC 43300) colonies on Luria-Bertani (LB) agar plates treated with the respective systems with or without light irradiation (30 min, 600 nm, 1 W cm⁻²).



Figure S16 - Viability of MDA-MB-231 and 293T cell lines in the absence and presence of (a) Fuc-AuNPs (0.2-1.0 nM) or (b) Gal-AuNPs (0.2-1.0 nM).

S3. Additional References

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