

Supporting Information

A biomineralized bi-functional hybrid nanoflower to effectively combat bacteria *via* a glucose-powered cascade catalytic reaction

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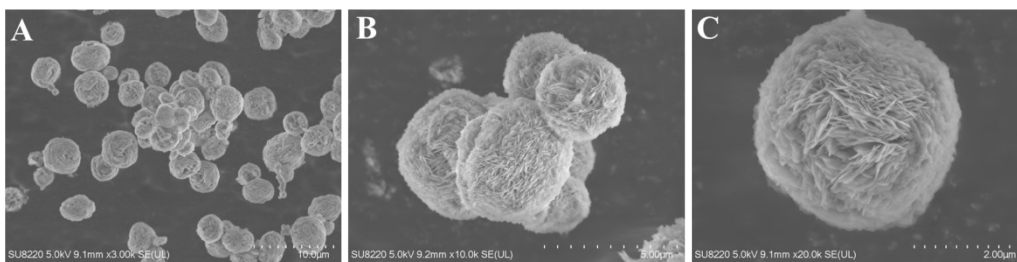


Figure S1. SEM images of Cu-GMPNF (A-C).

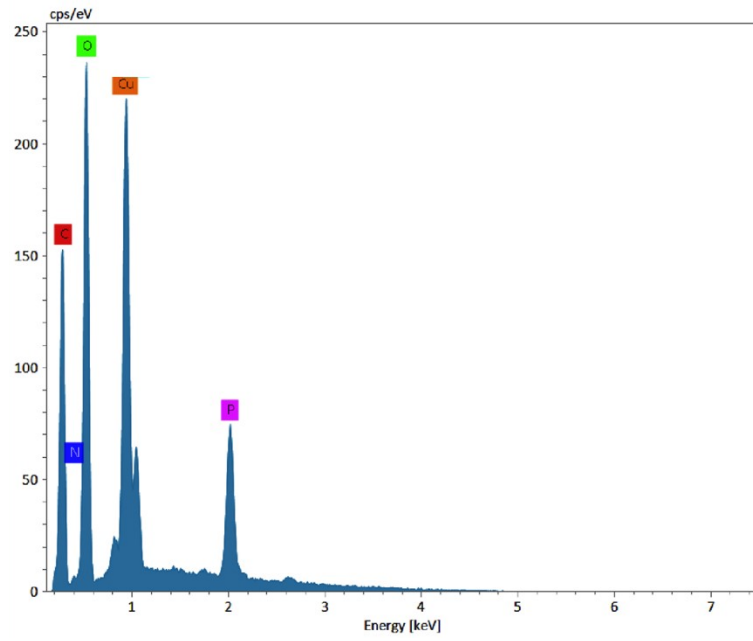


Figure S2. EDS of Cu-GMP/GODNF

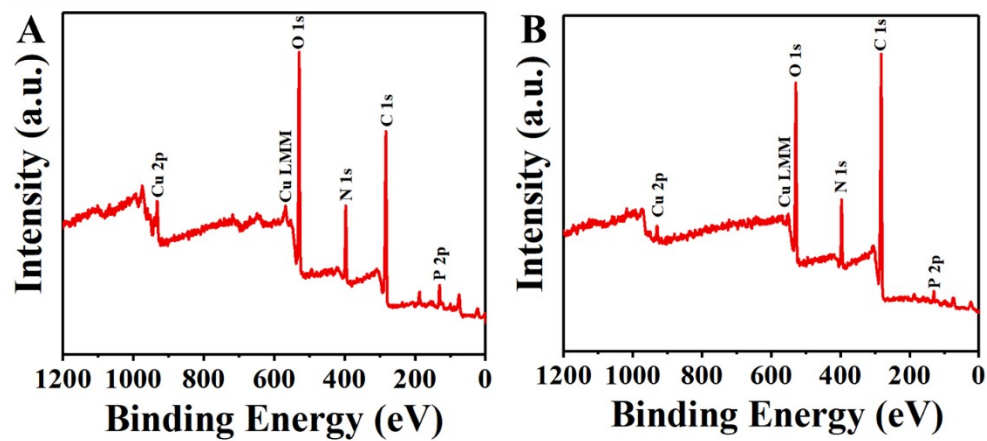


Figure S3. XPS spectra of Cu-GMPNF (A) and Cu-GMP/GODNF (B)

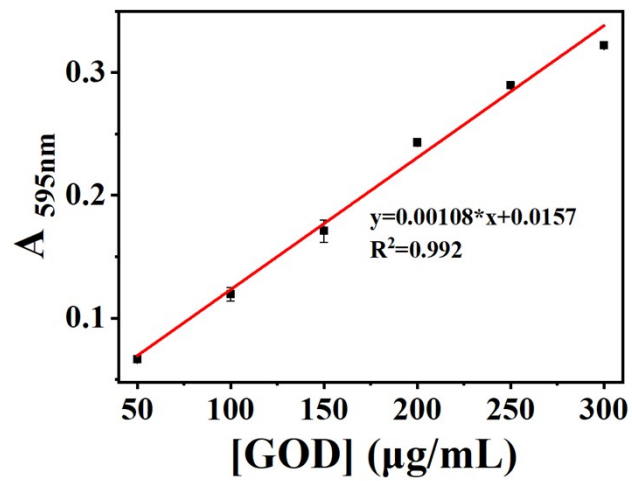


Figure S4. Calibration curves for different concentrations of GOD solutions (50 ~ 300 µg · mL⁻¹) according to Modified Bradford Protein Assay Kit.

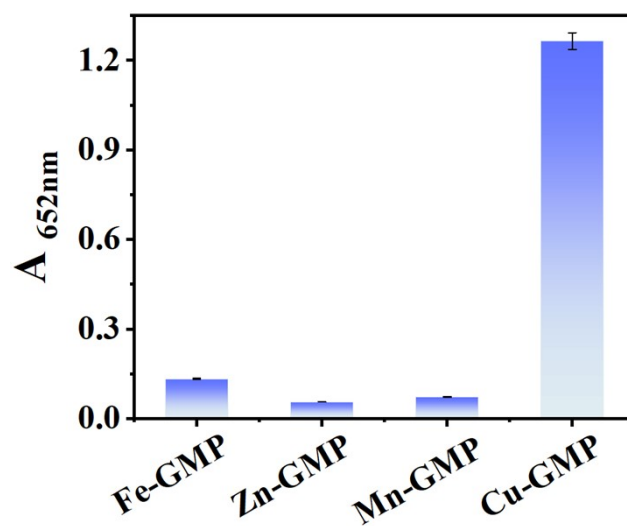


Figure S5. The POD-like activity of different metal/GMP (0.1 mg/mL) by measuring the absorbance of oxTMB at 652 nm under 2.0 mM H₂O₂ (NaAc-HAc buffer, pH 5.0).

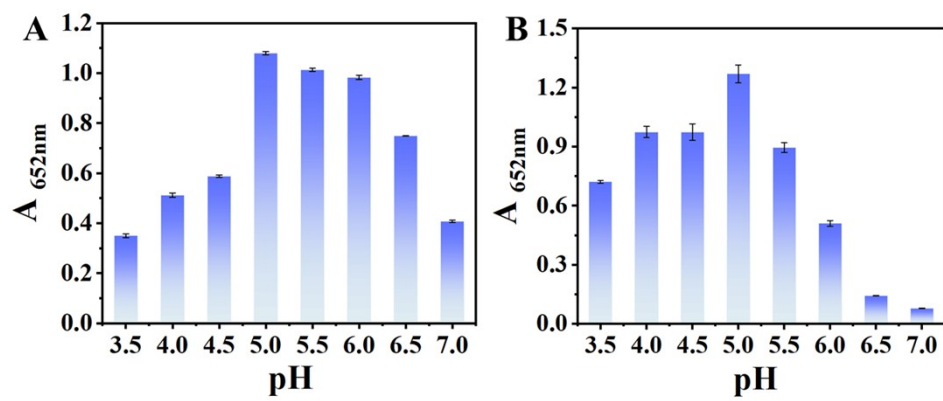


Figure S6. pH-dependence POD-like activity of Cu-GMPNF and cascade catalytic activity of Cu-GMP/GODNF.

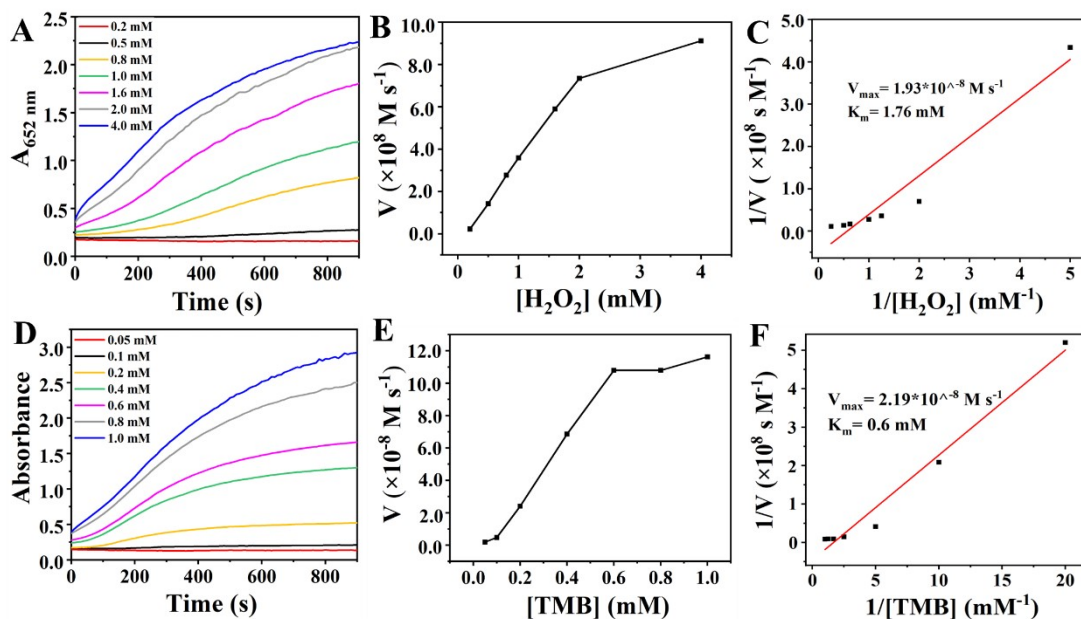


Figure S7. Typical Michaelis-Menten curves and kinetic parameters of Cu-GMPNF. Absorbance at 652 nm vs time of reaction solutions catalyzed by the Cu-GMPNF under different concentrations of H₂O₂ (A) or TMB (D). The velocity (*v*) of the reaction changes in the presence of different concentrations of H₂O₂ (B) or TMB (E). The double-reciprocal plot in the presence of different concentrations of H₂O₂ (C) or TMB (F). All experiments were conducted in 1.0 mg/mL of Cu-GMPNF. When the H₂O₂ concentration was varied, the TMB concentration was fixed at 1.0 mM. When the TMB concentration was varied, the H₂O₂ concentration was fixed at 5.0 mM.

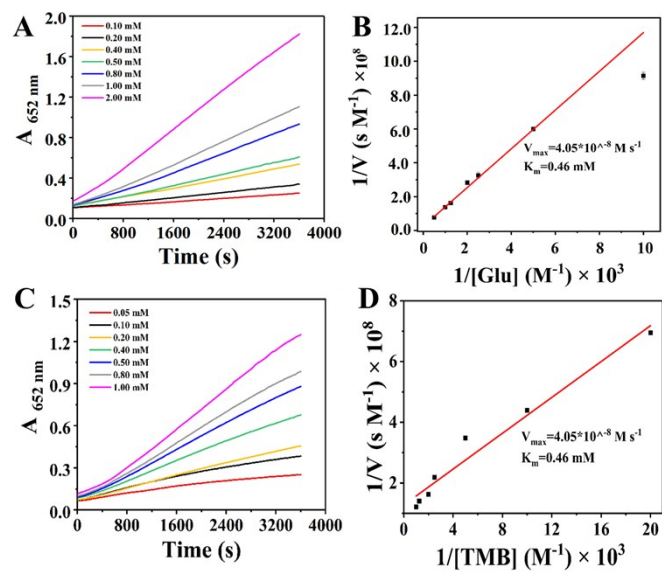


Figure S8. Steady-state kinetic data of the Cu-GMP/GODNF. Absorbance at 652 nm vs time of reaction solutions catalyzed by the Cu-GMP/GODNF under different concentrations of glucose (A) or TMB (D). Velocity (v) of the reaction changes in the presence of different concentrations of glucose (B) or TMB (E). Double-reciprocal plot in the presence of different concentrations of glucose (C) or TMB (F).

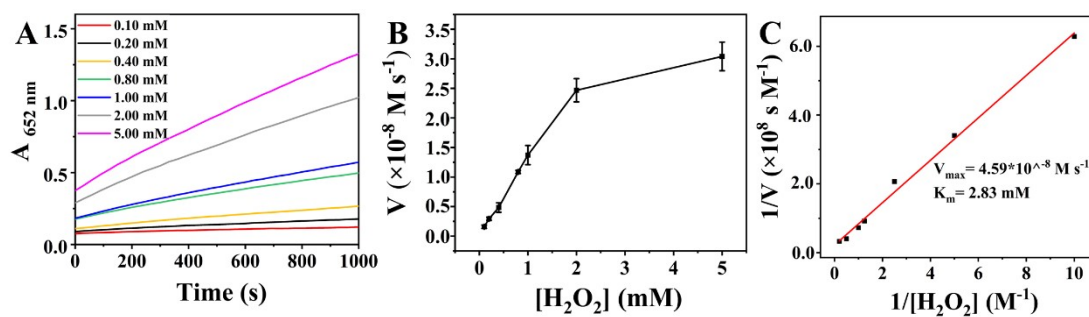


Figure S9. Typical Michaelis-Menten curves and POD-like kinetic parameters of Cu-GMP/GODNF. Absorbance at 652 nm vs time of reaction solutions catalyzed by the Cu-GMP/GODNF under different concentrations of H_2O_2 (A). The velocity (v) of the reaction changes in the presence of different concentrations of H_2O_2 (B). The double-reciprocal plot in the presence of different concentrations of H_2O_2 (C). All experiments were conducted in 1.0 mg/mL of Cu-GMP/GODNF. When the H_2O_2 concentration was varied, the TMB concentration was fixed at 1.0 mM.

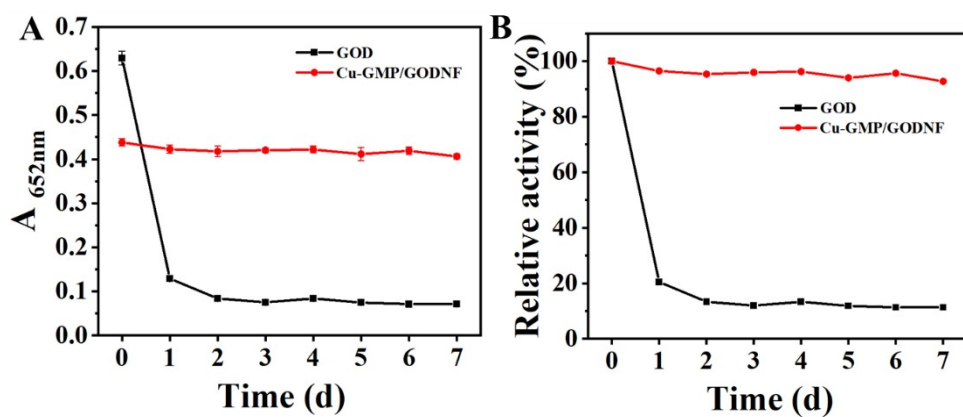


Figure S10. Absorbance of free GOD + Cu-GMPNF or Cu-GMP/GODNF catalyzed glucose/TMB at 652 nm (A), catalytic activity of free GOD and Cu-GMP/GODNF (B) under different storage times.

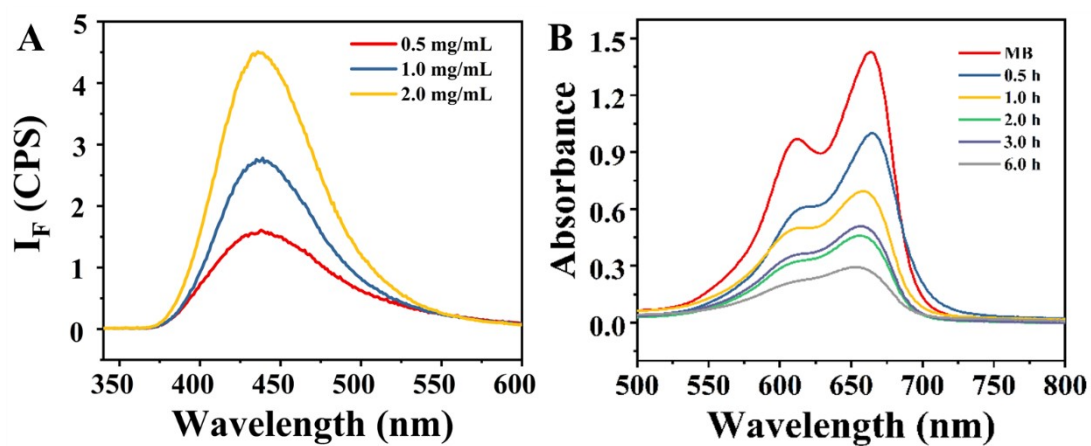


Figure S11. TA fluorescent signal generated with increasing Cu-GMP/GODNF concentration at 435 nm (A). The degradation of MB at different time points (B).

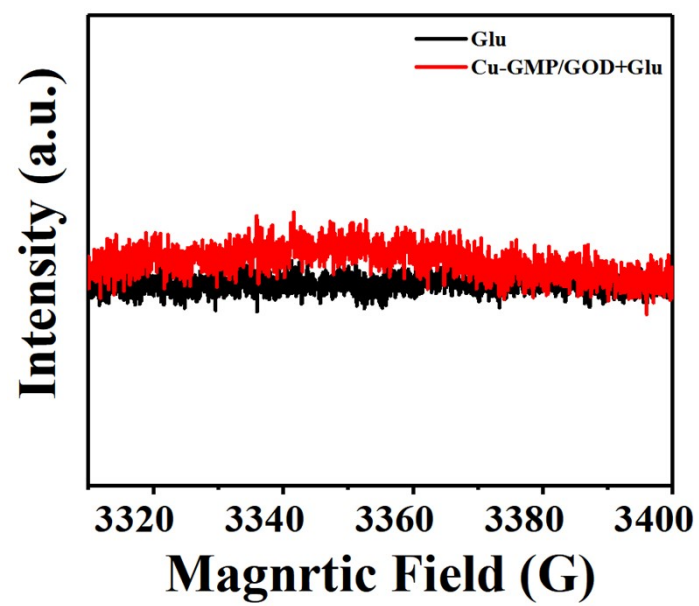


Figure S12. ESR spectra of different reaction systems with and $^1\text{O}_2$.

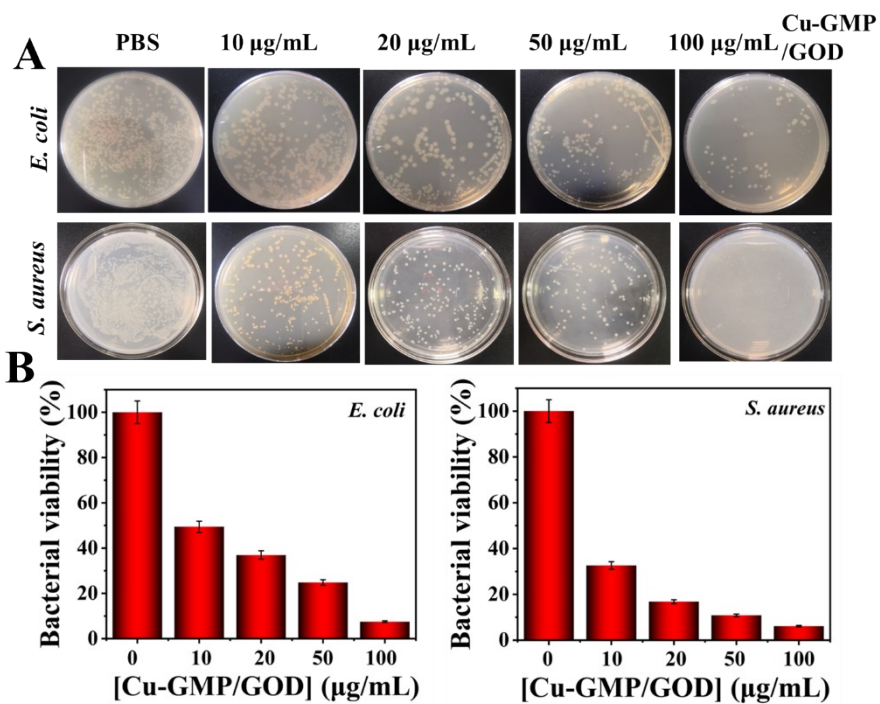


Figure S13. Agar plate results (A) and the corresponding bacterial viabilities (B) of *E. coli* and *S. aureus* in the presence of Cu-GMP/GODNF with different concentrations.

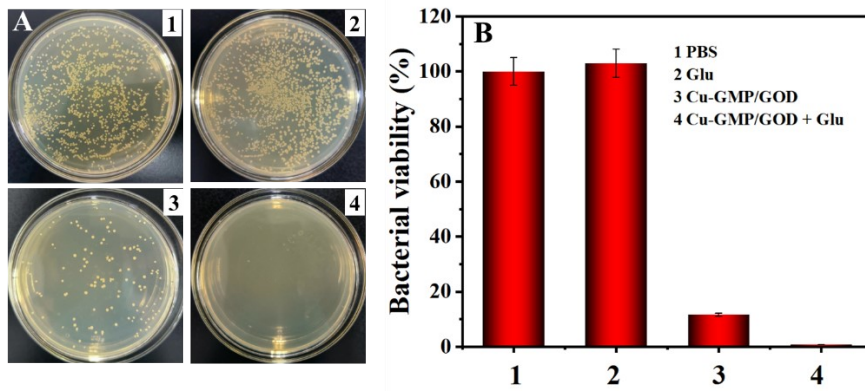


Figure S14. In vitro antibacterial efficacy of Cu-GMP/GODNF. Photographs of bacterial colonies formed (A) and the corresponding bacterial viabilities (B) of MRSA after different treatments.

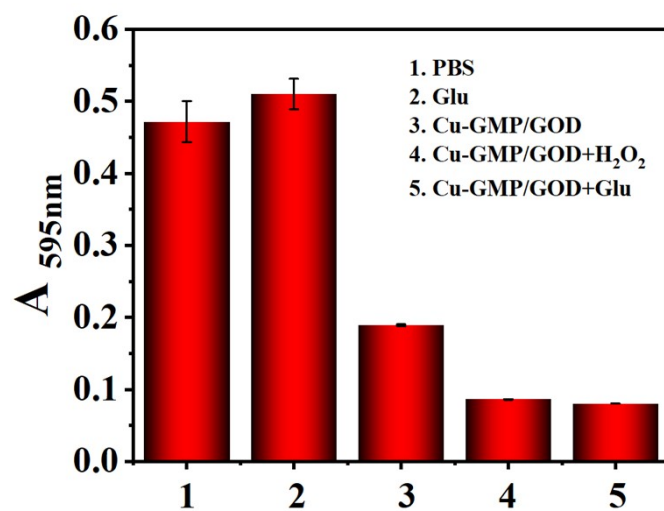


Figure S15. Biomass of MRSA biofilms under different treatments by measuring OD values at 595

nm.

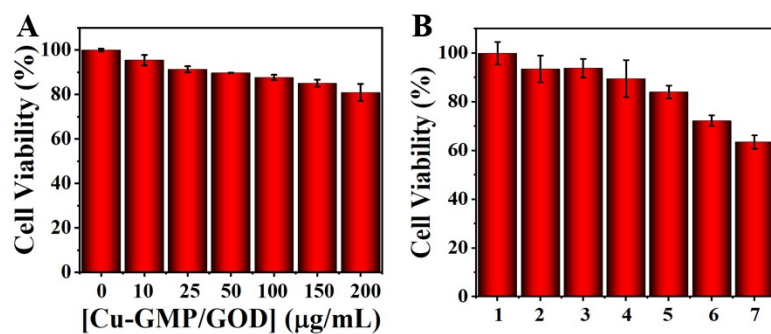


Figure S16. In vitro cell viability of L929 cells incubated with Cu-GMP/GODNF under different concentrations (A), and different treatments (1 PBS, 2 Cu-GMP, 3 Cu-GMP+Glu, 4 Cu-GMP/GODNF, 5 Cu-GMP/GODNF+1.0 mM Glu, 6 Cu-GMP/GODNF+2.0 mM Glu, 7 Cu-GMP/GODNF+5.0 mM Glu), the concentration of Cu-GMP and Cu-GMP/GODNF were 0.1 mg/mL, respectively (B).

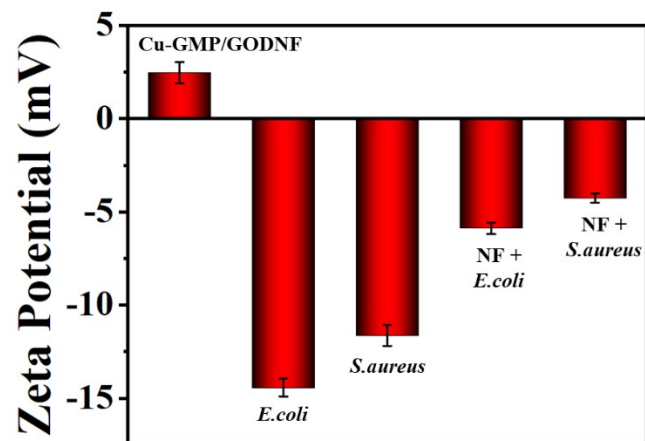


Figure S17. Zeta potential of bacteria before and after incubation with Cu-GMP/GODNF (1 mg mL⁻¹).

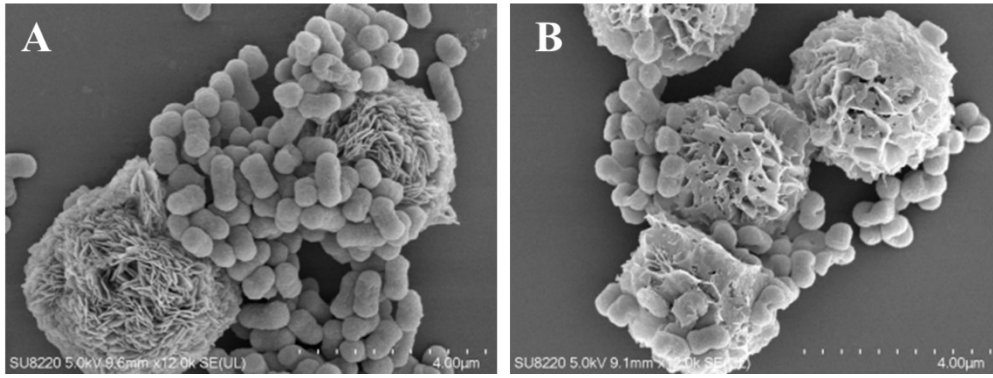


Figure S18. SEM images for verifying adhesion of Cu-GMP/GODNF to *E. coli* (A) and *S. aureus* (B).

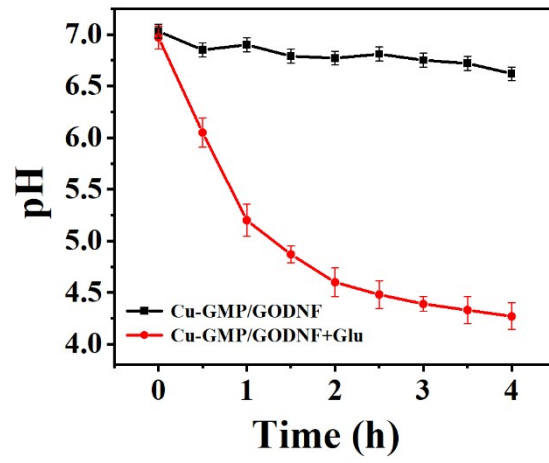


Figure S19. Changes in pH of bacteria after different treatments within 4 h.

Table S1. Kinetic parameters of Cu-GMPNF and NPs Cu-GMP/GODNF

Nanozymes	TMB		H ₂ O ₂		Glu	
	K_m	V_{max}	K_m	V_{max}	K_m	V_{max}
	(mM)	(10 ⁻⁸ M/s)	(mM)	(10 ⁻⁸ M/s)	(mM)	(10 ⁻⁸ M/s)
Cu-GMPNF	0.60	2.19	1.76	1.93		
Cu-GMP/GODNF	0.23	0.783	2.83	4.59	1.78	1.04