Supporting information

Rapid and Sensitive detection of *Staphylococcus aureus* and *Klebsiella pneumonia* based on Bacitracin-modified Fe₃O₄@PDA magnetic beads combined with Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry

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Fig.S1 The effect of adding Fe_3O_4 @PDA@Bacitracin and Fe_3O_4 @PDA (negative control) on the enrichment *S. aureus*

The plate results of *S.aureus* (A) before enrichment, (B) enrichment by 60µL Fe₃O₄@PDA@Bacitracin, (C) enrichment by 60µL Fe₃O₄@PDA(negative control)



Fig.S2 Effect of the different volume of Fe₃O₄@PDA@Bacitracin on the enrichment of *Staphylococcus aureus*.

(A)The plate results of the count of $10^2 \text{ CFU} \cdot \text{mL}^{-1} S$. *aureus* (before enrichment); (B)-(F). The plate results of the count of $10^3 \text{ CFU} \cdot \text{mL}^{-1} S$. *aureus* after enriched by 20,40,60,80,100 µL Fe₃O₄@PDA@ Bacitracin in turn



Fig. S3 Effect of different bacterial protein extraction pre-treatment methods on MALDI-TOF MS assay.(A) 70% formic acid treatment before targeting, (B) 70% formic acid / acetonitrile (1:1) treatment before targeting, (C) target first and then treated with 70% formic acid



Fig.S4 Effects of different matrixes on the MALDI-TOF MS analysis (A) α -CHCA (B) DHB



Fig.S5 Imaging of *S. aureus* protein treated by different matrixes in visual field (A) α-CHCA (B) DHB



Fig.S6 MALDI-TOF-MS spectra of different concentrations of the *S. aureus* in aqueous solution after enrichment by Fe₃O₄@PDA@Bacitracin: (A) 10⁴CFU·mL⁻¹ (B) 10⁵CFU·mL⁻¹ (C) 10⁶CFU·mL⁻¹ (D) 10⁷CFU·mL⁻¹



Fig.S7 MALDI-TOF-MS spectra of different concentrations of the *K. pneumoniae* in aqueous solution after enrichment by Fe₃O₄@PDA@Bacitracin: (A) 10⁵CFU·mL⁻¹ (B) 10⁶CFU·mL⁻¹ (C) 10⁷CFU·mL⁻¹



Fig.S8 Reference MS spectra of (A) Staphylococcus aureus and (B) Klebsiella pneumonia

Name of bacteria	Plate counting results (CFU·mL ⁻¹)	Enrichment ratio
S.aureus	1.45×10 ³	45.2%
K.pneumoniae	4.3×10 ³	37.0%
P.aeruginosa	2.57×10 ³	20.9%
Listeria monocytogenes	1.47×10 ³	8.61%
Pyogeniccoccus	2×10 ³	6.62%
E. Coli	1.26×10 ³	6.42%
shigella flexneri	2.75×10 ³	1.92%
beta hemolytic streptococcus	1.42×10 ³	1.21%
<i>E. coli</i> O157:H7	8.85×10 ³	0%
Enterobacter sakazakii	4.95×10 ³	0%

Table S1 Enrichment of Fe₃O₄@PDA@Bacitracin to different bacteria