

1 **Sensitive and simultaneous detection of different pathogens by**
2 **surface-enhanced Raman scattering based on aptamer and Raman**
3 **reporter co-mediated gold tags**

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14 **ABSTRACT**

15 A biosensor based on novel SERS tags, consisting of gold nanorods (GNRs) complexed with
16 oligonucleotide aptamers and the Raman reporters, was developed for the sensitive and
17 simultaneous detection of different food pathogens. The aptamers not only act as bio-recognition
18 molecules, but along with the Raman reporters, induce the GNRs to grow to specific shapes,
19 which in turn enhance the Raman signal and facilitate sensitive detection. Signal interference
20 during the simultaneous detection of pathogens is avoided, due to the stable anchored aptamers
21 and embedded Raman reporters. We combined the novel SERS tags with antibody-modified
22 magnetic nanoparticles to create a biosensor capable of simultaneous detection of *E. coli*
23 O157:H7 and *S. typhimurium* with good linear response (10^1 to 10^6 cfu/mL), high detection
24 sensitivity (< 8 cfu/mL) and recovery rate (95.26%-107.88%) in spiked food samples. This
25 strategy achieves the goal of sensitive and simultaneous quantitative detection of pathogens.

26 **Keywords:** surface enhanced Raman scattering (SERS); pathogens; simultaneous; aptamer;
27 biosensor.

28 **1 Introduction**

29 Foodborne pathogens are a global public health issue. *Escherichia coli* O157:H7 and
30 *Salmonella typhimurium* are the most commonly reported foodborne pathogens. They cause
31 serious issues such as bloody diarrhea [1,2], intestinal infectious diseases [3], food poisoning [4],
32 and even death. Simultaneous detection of multiple foodborne pathogens is of the utmost
33 importance to offer more useful information for illness control, in comparison to single species
34 pathogen, due to the multiple contaminations. Currently, the screening and detection pathogen is
35 mainly done by culture plate [5], polymerase chain reaction [6], enzyme-linked immunosorbent
36 assay [7], or by biosensors based on electrochemistry [8], quartz crystal microbalance [9], or
37 surface plasmon resonance methods [10]. However, the sensitive and simultaneous detection of
38 multiple foodborne pathogens is still restricted by signal interference, lack of sufficient
39 sensitivity, or difficulty achieving quantitative detection, which is important information for
40 early screening and disease control.

41 Surface-enhanced Raman scattering (SERS) is an ultrasensitive, vibrational spectroscopic
42 technique that enables the detection of molecules on or near the surface of a noble metal
43 nanostructure, which, due to local surface plasmon resonance, greatly enhances the strength of
44 the Raman signal [5]. SERS-based biosensors, possessing fingerprint precision, high sensitivity
45 and quantitative detection capability [11], have been widely used for pathogen detection. Label
46 –based SERS method has higher sensitivity than label-free method attributed to the specific
47 binding of the aptamers, leading to sharp characteristic peaks in the spectra of the Raman
48 reporters, and to the strong and stable enhancement of the noble metal substrates [12-14].
49 However, existing SERS tags used for the simultaneous detection of pathogens are generally
50 prepared by adsorbing Raman reporters onto the surface of nanomaterials, which are easily

51 eliminated and destroyed during subsequent washing treatments [15]. The limited surface area of
52 nanomaterials also affects the adsorption of Raman reporters onto their surfaces [16].

53 To solve these problems, SERS tags containing core-shell structures have been developed to
54 improve the stability, as well as the signal reproducibility, of the tags [17]. A variety of coating
55 materials have been applied to the surface of SERS tags to create these core-shell structures
56 [18-20]. Once the basic core-shell structure is established, chemical or biological groups
57 modified molecules may be subsequently added [21-23]. The core-shell SERS tags-based
58 method can provide a relatively stable and reproducible SERS signal, but the complicated
59 syntheses and high cost limit its application. There is a need for simple fabrication methods for
60 SERS tags, possessing stable and strong Raman signals and specific bio-recognition to realize
61 the simultaneous detection of foodborne pathogens [24-27].

62 DNA binds to gold via surface-binding moieties such as carbonyls and amides, with an
63 adsorption ability order of G > A > C > T [28]. Meanwhile, the morphology of nanomaterials can
64 be regulated by DNA to form nanoparticles and nanorods with specific sharp shapes [29,30].
65 According to the electromagnetic field enhancement principle, a sharp morphology produces a
66 “hotspot” effect to facilitate Raman signal enhancement [31]. Furthermore, during the regrowth
67 of the nanostructures, DNA molecules not only provide a stable anchor in the outer layer of the
68 nanomaterial, but also retain their bio-recognition properties [32], providing an opportunity for
69 aptamer-sequence design and multipurpose usage in SERS tag fabrication. However, producing
70 embedded reporter and DNA-based SERS tags directly and controllably with stable and strong
71 Raman signals is difficult. Our team has recently developed nanobones shaped SERS tags for the
72 detection of *E. coli* O157:H7 [33], while more challenging work is developing a multiplex
73 biosensor capable of recognizing multiple pathogens simultaneously, which could overcome the

74 limitations of signal interference and sensitivity.

75 To realize the sensitive and simultaneous detection of *E. coli* O157:H7 and *S. typhimurium*,
76 we have developed a novel biosensor based on different morphologically controllable SERS tags
77 via Raman reporter and aptamer co-mediated gold nanorods (GNRs). The tags, grown using
78 one-pot synthesis, provide stable and strong characteristic signals, as well as bio-recognition
79 specificity, thus avoiding signal interference during the simultaneous detection of pathogens.
80 SERS tags with cracked octahedral shapes and small protrusion morphologies were fabricated
81 using two different Raman reporters and specifically designed aptamers during mediated
82 regrowth of the GNRs. Assisted by antibodies conjugated onto modified magnetic nanoparticles,
83 *E. coli* and *S. typhimurium* were simultaneously and quantitatively detected within 10^1 – 10^6
84 cfu/mL with LODs of 5 and 8 cfu/mL, respectively. This biosensor, benefiting from the novel
85 SERS tags, realizes the goal of sensitive and simultaneous quantitative detection of bacteria, and
86 may possibly become a universal detection tool for the daily simultaneous detection of
87 foodborne pathogens.

88 **2 Materials and methods**

89 *2.1 Chemical and biochemical materials*

90 Chloroauric acid trihydrate (HAuCl₄), cetyltrimethylammonium bromide (CTAB), sodium
91 borohydride, ascorbic acid, silver nitrate, DTNB, MBA, bovine serum albumin (BSA),
92 hydroxylamine hydrochloride (HA), and sodium hydroxide were purchased from Sigma-Aldrich
93 (St. Louis, MO, USA). Streptavidin-modified magnetic nanoparticles (MNPs) of 150 nm
94 diameter were obtained from Ocean nanotech (Dunedin, FL, USA). The Luria–Bertani medium
95 (LB), alkaline peptone water medium, and agar were purchased from Aoboxing Biotech (Beijing,

96 China). Phosphate-buffered saline (PBS), *E. coli* O157:H7 (ATCC 43888), *S. typhimurium*
97 (ATCC14028), *E. coli* (ATCC 25922), *Vibrio parahemolyticus* (ATCC 17802), and *S. aureus*
98 (ATCC 25923) were purchased from Solarbio Life Sciences (Beijing, China). Rabbit antibody
99 against *E. coli* O157:H7, and mouse antibody against *S. typhimurium*, were purchased from
100 Meridian Life Science (Memphis, TN, USA). A Long-Arm Biotin Labeling Kit from
101 Elabscience Biotechnology (Wuhan, China) was used for the modification of antibodies. The
102 designed aptamers for *E. coli* O157:H7 (TTTTT TTTTT TTTTT TTTTT CCGGA CGCTT
103 ATGCC TTGCC ATCTA CAGAG CAGGT GTGAC GG) [34] and *S. typhimurium* (CCCCC
104 CCCCC CCCCC TATGG CGGCG TCACC CGACG GGGAC TTGAC ATTAT
105 GACAG) [35] were synthesized by Sangon Biotech (Shanghai, China). Ultrapure water was
106 prepared using a Milli-Q system (Bedford, MA, USA).

107 2.2 Preparation of novel SERS tags

108 The novel SERS tags consisted of regrown GNRs co-mediated by aptamers and Raman
109 reporters, prepared using one-pot synthesis as illustrated in **Fig. 1A**. The GNRs were prepared
110 via CTAB-mediated seed growth, as previously described [31]. Meanwhile, the aptamers were
111 heated to 90°C to open the secondary structure and then immediately put on ice to maintain the
112 open chain structure. Then, 90 μL the prepared GNRs were incubated with 2.97 μL 100 μM
113 prepared aptamers for 1 h, followed by addition of 1 μL 1 mM Raman reporters. The tag-1 and
114 tag-2 were synthesized separately. Aptamers of *E. coli* O157:H7 and DTNB were used for tag-1
115 syntheses, while aptamers of *S. typhimurium* and MBA for tag-2 syntheses. After another 1 h
116 incubation, 2.5 μL 40 mM HA solution (pH 5.0) as mild reducing agent, and 1 μL 1% (w/w)
117 HAuCl₄ solution as the gold precursor, were added to the solution, which then was shaken
118 vigorously for 1.5 h to obtain the novel SERS tags. The SERS tags were purified by

119 centrifugation at $5000 \times g$ for 3 min to remove the supernatant and then re-suspended in
120 ultrapure water.

121 *2.3 Characterization of SERS tags*

122 UV-vis spectra were recorded on a Shimadzu UV-1780 spectrometer. Transmission electron
123 microscopy (TEM) (HITACHI HT7700, Japan) was used to characterize the morphology of the
124 nanostructures. Energy-dispersive spectrometry (EDS) and high-resolution transmission electron
125 microscopy (HRTEM) were conducted using a JEOL JEM-2100F microscope at an accelerating
126 voltage of 200 kV to obtain images representing the microscopic surface structures and element
127 distributions. SERS detection was performed on a DXR Raman microscope (JY H-800,
128 HORIBA), equipped with a 633 nm excitation laser and a $10 \times$ objective confocal microscope
129 ($2 \mu\text{m}$ spot diameter and 5 cm^{-1} spectral resolution). Samples were excited using 15-mW laser
130 power and a slit width of $50 \mu\text{m}$, with a total integration time of 10 s for each SERS spectrum.
131 Five spots were chosen randomly for each sample measurement.

132 *2.4 Separation and detection of bacteria*

133 Fe_3O_4 -antibody capture probes were prepared by conjugating monoclonal antibodies onto
134 MNPs. Briefly, $30 \mu\text{L}$ 1 mg/mL streptavidin-modified MNPs with 150 nm diameters was
135 incubated with $0.6 \mu\text{L}$ 1 mg/mL biotinylated antibody for 45 min at 15 rpm in a 1% BSA-soaked
136 centrifuge tube, which conducted in a disc rotary mixer at room temperature ($25 \text{ }^\circ\text{C}$). The surplus
137 antibodies were removed by magnetic separation and PBS washing twice, and the monoclonal
138 antibody-modified MNPs were suspended in PBS.

139 *E. coli* O157:H7 and *S. typhimurium* were cultured in LB medium at 37°C for 12 h and then
140 serially diluted 10-fold with sterile PBS buffer to final concentrations from 10⁸ to 10¹ cfu/mL.
141 Then two kinds of antibody-modified MNPs (30 μL) were added. The mixture was incubated for
142 45 min at 15 rpm to form MNPs-bacteria complexes. After capture, the MNPs-bacteria
143 complexes were separated and washed twice with PBS under a magnetic field. Then, 180 μL
144 prepared SERS tags (0.4 nM each of tag-1 and tag-2) were added, followed by another 45-min
145 incubation at 15 rpm. The capture probe–pathogen–SERS tag sandwich complexes were
146 collected and washed twice with PBS under a magnetic field and then used to measure the SERS
147 spectra.

148 *2.5 Optimization and evaluation of the established biosensor*

149 The Raman signal intensity of the tags is a key performance factor of the biosensor and
150 depends on the aptamers and Raman reporters. The concentrations of aptamer-1 and aptamer-2
151 (1.5, 2.0, 2.5, 3.0, 3.5, and 4.0 μM) and of DTNB and MBA (2.5, 5, 10, 20, 30, 40, and 50 μM)
152 were tested to optimize the conditions for fabrication of tag-1 and tag-2. The various
153 concentrations were evaluated according to the Raman signal intensity of the fabricated SERS
154 tags to maximize the detection sensitivity. Then, we determined the optimum concentration of
155 tags by adding various concentrations (0.1, 0.2, 0.3, 0.4, 0.5, and 0.6 nM) of tags into 10⁶ cfu/mL
156 pathogens.

157 Suspensions of *E. coli* O157:H7 and *S. typhimurium* at concentrations of 10⁸–10¹ cfu/mL were
158 used to determine the range of pathogen concentrations over which the biosensor response was
159 linear. Partial least-squares (PLS) analyses were used to evaluate the linear relationship; the
160 relationship was considered linear if the correlation coefficient, R, was close to 1, and the

161 root-mean-square error of calibration (RMSEC) was close to 0. The LOD was calculated
162 according to the formula $LOD = 3N/S$, where N is the standard deviation of measurements on a
163 blank sample and S is the slope of the standard curve. To evaluate the specificity of the biosensor,
164 *E. coli*, *V. Parahemolyticus*, and *S. aureus* were used as interference bacteria. In addition, 20
165 random samples from a mixture of *E. coli* O157:H7 and *S. typhimurium* at 10^4 cfu/mL were
166 selected to evaluate the reproducibility of the biosensor.

167 *2.6 Analyses of food samples*

168 Samples of tap water, cucumber, and chicken were used for realistic tests of the detection of
169 *E. coli* O157:H7 and *S. typhimurium*. The samples were prepared according to previous reports
170 with small modifications [14]. The food samples of cucumber and chicken were washed with
171 ultrapure water and homogenized, filtered to remove solid precipitates, and then mixed with PBS
172 at a ratio of 1: 9 (w/v). Then, *E. coli* O157:H7 and *S. typhimurium* at concentrations of 3.5×10^2
173 and 2.0×10^3 cfu/mL were added to each sample, respectively. After that, antibody-modified
174 MNPs and SERS tags were added into spiked sample subsequently for separation and labelling
175 according to Section 2.4. Finally, the Raman detection was conducted to calculate the
176 concentration of bacteria. Traditional plate counting was also used to assess the recovery and
177 accuracy of the Raman detection method. The recovery rate was calculated as the ratio of
178 concentrations measured using SERS to the concentrations measured using the plate counting
179 method. The accuracy was represented by relative standard deviation (RSD).

180 *2.7 Data analyses*

181 SERS spectra were recorded and analyzed using LabSpec application software and TQ analyst
182 software (v8.0, Thermo Fisher Scientific). The raw SERS spectra were pre-processed by

183 smoothing the data and adjusting the baseline for later analyses. All the experiments were
184 repeated three times. For each sample analyzed by SERS, five spots were selected randomly and
185 scanned in the range of 400-2000 cm^{-1} . The results are presented as means and standard deviation,
186 calculated using Origin 8.0.

187 **3 Results and discussion**

188 *3.1 Design principles of the novel SERS biosensor for pathogens*

189 The sensitive and simultaneous detection of *E. coli* O157:H7 and *S. typhimurium* were carried
190 out based on two kinds of morphologically controllable, novel SERS tags, which exhibited
191 strong Raman signals and bio-recognition specificities (**Fig. 1**). The SERS tag-1, which was
192 synthesized using DTNB and an aptamer complementary to *E. coli* O157:H7, acquired a cracked
193 octahedral shape, while the SERS tag-2, which was synthesized using MBA and an aptamer
194 complementary to *S. typhimurium*, acquired small protrusions (**Fig. 1A**). Both aptamers and
195 Raman reporters, when incubated with GNRs in the one-pot synthesis, successfully adsorbed
196 onto the surfaces of the GNRs, which then grew into morphologically controllable tags. The
197 cracked octahedral shape of tag-1 and the small protrusion morphology of tag-2 created more
198 hotspots (localized regions of intense electromagnetic field enhancement), which greatly
199 enhanced the Raman signal [31]. Meanwhile, the embedded Raman reporters in the outer layers
200 of the SERS tags resulted in stable and strong Raman signal enhancements [17], which
201 contributed to the sensitivity and overcame any signal interference arising from the simultaneous
202 detection of multiple pathogens. During the regrowth process, the mononucleotide repeats at the
203 5' ends of the aptamers stably anchored the SERS tags into the GNRs [29], comparable to
204 thiol-modified aptamers. The aptamers not only tuned the regrowth process of the GNRs to a

205 programmed morphology, but also retained bio-recognition specificity for *E. coli* O157:H7 and
206 *S. typhimurium*. The SERS tags specifically labeled the target pathogens to form sandwich
207 complexes with antibody-modified MNPs, thereby realizing simultaneous SERS detection of the
208 pathogens (**Fig. 1B**).

209 3.2 Characterization of the SERS tags

210 The SERS tags, being key factors for the performance of the biosensor, were characterized by
211 UV-vis absorption spectroscopy, TEM, HRTEM, EDS, and SERS. The prepared GNRs
212 displayed a blue-green color with transverse and longitudinal localized surface plasmon
213 resonance (T-LSPR and L-LSPR) peaks at 516 and 650 nm, respectively. The colors for Tag-1
214 and Tag-2 changed to reddish purple and blueish purple, and the T-LSPR and L-LSPR peaks
215 were red-shifted to 527 and 670 nm, and 538 and 677 nm, respectively (**Fig. 2A**). The red shift of
216 the LSPR peaks and their separations suggest that the re-grown GNRs in the one-pot synthesis of
217 the SERS tags were larger and more uniform in shape than the original GNRs. To investigate the
218 shape of synthesized SERS tags, TEM was used to characterize the morphology of the SERS
219 tags. The original GNRs had a well-defined rod shape and an average aspect ratio of 2.5 (width
220 ~22 nm, length ~55 nm). The GNRs grown with DTNB and aptamer-1 exhibited a cracked
221 octahedral shape with a smooth surface (tag-1), while the GNRs grown with MBA and
222 aptamer-2 exhibited small protrusions and a rough surface (tag-2). The average aspect ratios of
223 tag-1 and tag-2 were both about 2.0 (tag-1: width ~32 nm, length ~65 nm; tag-2: width ~40 nm,
224 length ~80 nm). The TEM results confirmed the larger and more uniform morphologies of the
225 GNRs in tag-1 and tag-2, in agreement with the red-shifted LSPR peaks. Differences in the
226 UV-vis absorption spectra of tag-1 and tag-2 may reflect the distinct shapes and gap distances
227 between pointed ends. Notably, the different morphologies of tag-1 and tag-2 may be attributed

228 to the sequence of nucleotides at the 5' ends of the aptamers. As previously reported, GNRs
229 grown with homopolymeric thymine (T20) grew into cracked octahedral with smooth facets [31],
230 as did our own tag-1. However, the GNRs in tag-2, which were grown with MBA and Aptamer-2,
231 which contained homopolymeric cytosine (C20) at the 5' end, grew small protrusions and had a
232 rough surface. Therefore, we speculated that the overgrowth time and concentrations of Raman
233 reporters both have effect on the morphology of tags.

234 The successful fabrication of the SERS tags was further confirmed by HRTEM imaging and
235 EDS. The magnified HRTEM images of the edges of tag-1 (**Fig. S1A** and **S1B**) and tag-2 (**Fig.**
236 **S1C** and **S1D**) show that the aptamers were uniformly distributed around the SERS tags in a
237 layer about 3.2 nm and 2.7 nm thick, respectively. The EDS image shows the presence of
238 aptamer-1 (element P) and DTNB (element S) in cracked octahedral-shaped tag-1 (element Au,
239 **Fig. 2B**) and the presence of aptamer-2 and MBA in small protrusion-shaped tag-2 (**Fig. 2C**),
240 demonstrating the successful embedding of aptamers and Raman reporters in the gold shell of the
241 SERS tags.

242 The SERS tags also exhibited greater enhancement effects, compared to the GNRs,
243 presumably due to the adhered and embedded Raman reporters (**Fig. 2D** and **2E**). Raman
244 enhancement factors (EFs) were 1.71×10^5 and 2.03×10^5 for tag-1 and tag-2, respectively,
245 which were four times greater than the EFs for the Raman reporters adsorbed on the surface of
246 GNRs without the aptamers. The Raman signal intensity depends largely on electromagnetic
247 enhancement of the geometrically defined surface plasmon resonances of the nanomaterials.
248 Here, the Raman signal enhancement was facilitated not just by the adhered and embedded
249 Raman reporters, but also by the tuned, irregular shapes, which produced more Raman hotspots.
250 These stronger EFs could facilitate the detection sensitivity of the biosensor.

251 3.3 Optimization of the SERS-based biosensor

252 During the one-pot synthesis of the SERS tags, the aptamers and Raman reporters compete for
253 limited binding space on the growing GNRs [32]. Thus, the concentrations of aptamers and
254 Raman reporters needed to be optimized to obtain the strongest possible Raman signal
255 enhancement. With increasing concentrations of aptamer and Raman reporter, the Raman signal
256 intensity of the SERS tags increased, reached an optimum, and then decreased (**Fig. 3A-3D**). To
257 facilitate the sensitive and simultaneous detection of *E. coli* O157:H7 and *S. typhimurium*, the
258 optimal concentrations of aptamer-1, aptamer-2, DTNB, and MBA for the one-pot syntheses of
259 the SERS tags were 3.0, 2.5, 10.0, and 10.0 μM , respectively. To determine the optimal
260 concentration of tags in the biosensor system, various concentrations of tag-1 and tag-2 were
261 added separately into a 10^6 cfu/mL suspension of *E. coli* O157:H7 and *S. typhimurium*. The
262 strongest characteristic Raman signal for sandwich complexes was obtained when the
263 concentration of the added tag was 0.4 nM, for both tag-1 and tag-2 (**Fig. 3E** and **3F**,
264 respectively). This result also demonstrated specific binding between the pathogens and the
265 SERS tags.

266 The formation of capture probe–pathogen–SERS tag sandwich complexes were further
267 indicated by TEM imaging. The cracked octahedral-shaped tag-1 was observed binding to *E. coli*
268 O157:H7 (**Fig. 4A**), while the small protrusion-shaped tag-2 was observed binding to
269 *S. typhimurium* (**Fig. 4B**). When both pathogens were present, tag-1 and tag-2 were each
270 specifically bound to their respective pathogens: tag-1 to *E. coli* O157:H7 and tag-2 to
271 *S. typhimurium* (**Fig. 4C**). The capture capability of the antibody-modified MNPs was confirmed
272 by traditional plate counting. The capture rates for *E. coli* O157:H7 and *S. typhimurium* were
273 above 90% in bacterial suspensions up to 10^6 cfu/mL (**Fig. 4D** and **4E**, respectively). The

274 stability and specificity of the capture probes contributed to the sensitive and simultaneous
275 detection of the pathogens by the biosensor.

276 3.4 Evaluation of the analytical performance

277 The individual and simultaneous detection of *E. coli* O157:H7 and/or *S. typhimurium* were
278 carried out under optimized conditions. For the individual detection, the favorable linear range of
279 concentrations was first explored by detecting gradient concentrations (10^1 to 10^8 cfu/mL) of *E.*
280 *coli* O157:H7 and *S. typhimurium*. Raman peaks of tag-1 at 1331 cm^{-1} and tag-2 at 1074 cm^{-1} ,
281 attributed to DTNB and MBA, respectively, were chosen for the quantitative analyses of *E. coli*
282 O157:H7 and *S. typhimurium*. The Raman signal intensity grew rapidly with increasing
283 concentrations of pathogens from 10^1 to 10^6 cfu/mL, and PLS analyses showed a good linear
284 relationship (**Fig. S2**). There were strong linear correlations between the intensity of the Raman
285 signal and the logarithms of the concentrations of *E. coli* O157:H7 ($y = 105.4x + 254.7$, $R^2 =$
286 0.9942) and *S. typhimurium* ($y = 87.6x + 59.5$, $R^2 = 0.9955$) with LOD values of 3 and 5 cfu/mL,
287 respectively (**Fig. S3**). These results demonstrate that the biosensor was sensitive and
288 quantitative for single-pathogen detection. For the simultaneous detection of *E. coli* O157:H7
289 and *S. typhimurium*, the characteristic peaks were readily separated, and both peak intensities
290 gradually increased with increasing concentration of pathogens (**Fig. 5A**). There were still good
291 linear correlations between 10^1 cfu/mL and 10^6 cfu/mL for *E. coli* O157:H7 ($y = 91.0x + 185.7$,
292 $R^2 = 0.9914$) and *S. typhimurium* ($y = 64.2x + 99.7$, $R^2 = 0.9800$) with LOD values of 5 and 8
293 cfu/mL, respectively (**Fig. 5B**). The LOD values for *E. coli* O157:H7 and *S. typhimurium* were of
294 the same order, indicating strong specificity for the tags and capture probes, with little signal
295 interference between them. Compared to other SERS methods for the simultaneous detection of
296 pathogens (**Table S1**) [36-38], the lower LOD values and wider linear range observed with our

297 biosensor confirmed its outstanding quantitative performance, which we attribute to the stable
298 and strong Raman signals of the novel SERS tags. With regard to localized surface plasmon
299 resonance (LSPR) method, bacteria could be monitored by measuring the wavelength shift in the
300 LSPR resonance peak, however, LOD values are relatively high [39, 40]. As for electrochemical
301 biosensors, the non-specific adsorption on the electrode affects the sensitivity and specificity of
302 detection [34]. In this study, the specific affinity of antibodies and aptamers enable high
303 detection specificity and sensitivity. In combination with specific shaped SERS tags, which
304 could provide strong characteristic Raman signals, the established biosensor could achieve more
305 sensitive detection for pathogens compared with the label-free method.

306 Good reproducibility was verified by taking 20 random samples from a mixed suspension of
307 *E. coli* O157:H7 and *S. typhimurium* at 10^4 cfu/mL. Their RSDs were 8.33% and 9.91%,
308 respectively (**Fig. 5C**). To verify the specificity of the biosensor, *E. coli*, *V. Parahemolyticus* and
309 *S. aureus* were used as interference bacteria. The signal intensities of *E. coli* O157:H7 and *S.*
310 *typhimurium* were almost 100-fold stronger than those of the non-target pathogens (**Fig. 5D**). A
311 very weak signal intensity in non-target samples (with MNPs but no tags) indicated that Raman
312 signals enhancement was contributed from SERS tags rather than MNPs. Furthermore, *E. coli*
313 O157:H7 and *S. typhimurium* were precisely detected in mixed samples, which consisted of *E.*
314 *coli* O157:H7, *S. typhimurium*, *E. coli*, *S. aureus*, and *Vibrio parahemolyticus*, attributed to the
315 high specific affinity of the tags and capture probes for their target pathogens. Good
316 reproducibility and specificity further demonstrate the reliability of our detection method.

317 To further investigate the applicability of the SERS biosensor for *E. coli* O157:H7 and
318 *S. typhimurium* in practical use, tap water, cucumber, and chicken were examined as models, and
319 our detection results were compared to the classic plate counting method as a standard [36]. As

320 shown in **Table 1**, the recovery rate for *E. coli* O157:H7 and *S. typhimurium* ranged from
321 95.92% to 105.32%, and 95.26% to 107.88%, respectively, indicating applicability and good
322 accuracy of the proposed biosensor for the quantification of *E. coli* O157:H7 and *S. typhimurium*
323 counts in food samples. "Plate counting" method needs at least 12 hours for culture, the SERS
324 biosensor method established in this study was much faster (about 1 hour). Besides, the detection
325 specificity can be guaranteed by the affinity of antibodies and aptamers, as well as the
326 characteristic Raman signals produced by different SERS tags, while plate counting method
327 needs further morphological identification. These results further demonstrate the bio-recognition
328 specificity of the two novel SERS tags and the absence of signal interference during
329 simultaneous SERS detection of pathogens.

330 **4 Conclusions**

331 The sensitive and simultaneous SERS detection of *E. coli* O157:H7 and *S. typhimurium* was
332 achieved without signal interference by using a biosensor containing two kinds of aptamers and
333 Raman reporters as novel SERS tags, which exhibit strong, stable Raman signals and
334 bio-recognition capabilities. The cracked octahedral-shaped tag-1 and small protrusion-shaped
335 tag-2 provided strong EFs of 1.71×10^5 and 2.03×10^5 , respectively, which boosted detection
336 sensitivity. The embedded Raman reporters and anchored aptamers with specific bio-recognition
337 capacity made possible the avoidance of signal interference during the simultaneous detection of
338 multiple pathogens. With monoclonal antibody-modified MNPs as capture probes, we were able
339 to specifically separate and enrich the target pathogens, *E. coli* O157:H7 and *S. typhimurium*,
340 demonstrating effective, sensitive and specific SERS detection of multiple pathogens. This
341 biosensor had good quantitative capability between the Raman signal intensity and the

342 logarithms of the concentrations of *E. coli* O157:H7 and *S. typhimurium* from 10^1 to 10^6 cfu/mL
343 in a suspension of multiple pathogens, with LOD values of 5 and 8 cfu/mL, respectively. The
344 recovery rates from three spiked food samples ranged from 95.26% to 107.88%, demonstrating
345 promising potential of the biosensor method to detect various pathogens, facilitating early
346 screening of contaminated food. This biosensor, benefiting from the flexible combination of
347 aptamers and Raman reporters in novel SERS tags, may possibly become a universal detection
348 tool for the simultaneous detection of other foodborne pathogens.

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355 **Declaration of Competing Interest**

356 The authors declare no competing financial interest.

357 **Appendix A. Supplementary data**

358 Supplementary data associated with this article can be found in the online version.
359 High-resolution TEM images of SERS tags-1 and tags-2 (**Fig. S1**); Partial least-squares (PLS)
360 analyses of *E. coli* O157: H7 and *S. typhimurium* concentration prediction (**Fig. S2**); Linear
361 correlation between the logarithm of pathogen concentrations and SERS intensity for *E. coli*.

362 O157: H7 and *S. typhimurium* in single detection system (**Fig. S3**); Comparison of various
363 methods for the simultaneous detection of pathogens by SERS (**Table S1**).

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491 **Captions**

492 **Fig. 1** Schematic illustration of the SERS detection biosensor. Principles of the one-pot synthesis
493 of SERS tags co-mediated with aptamers and Raman reporters (A), and the separation
494 and SERS detection of *E. coli* O157:H7 and *S. typhimurium* (B).

495 **Fig. 2** Characterization of the SERS tags. UV absorption spectra, appearance, and transmission
496 electron microscopy (TEM) images of GNRs and SERS tags (A); high-resolution TEM
497 (HR TEM) images and energy dispersive spectrometer images of tag-1 (B) and tag-2 (C):
498 P = aptamer, S = Raman reporter, Au = gold nanorod; Raman enhancement of tag-1 (D)
499 and tag-2 (E).

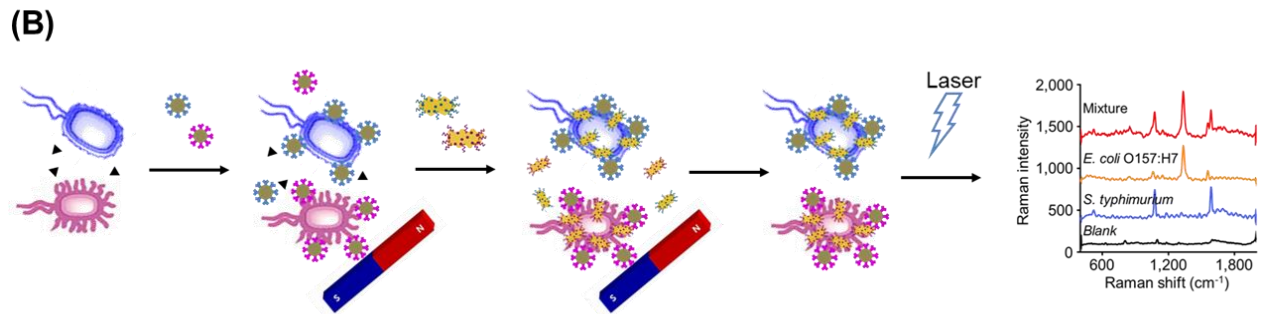
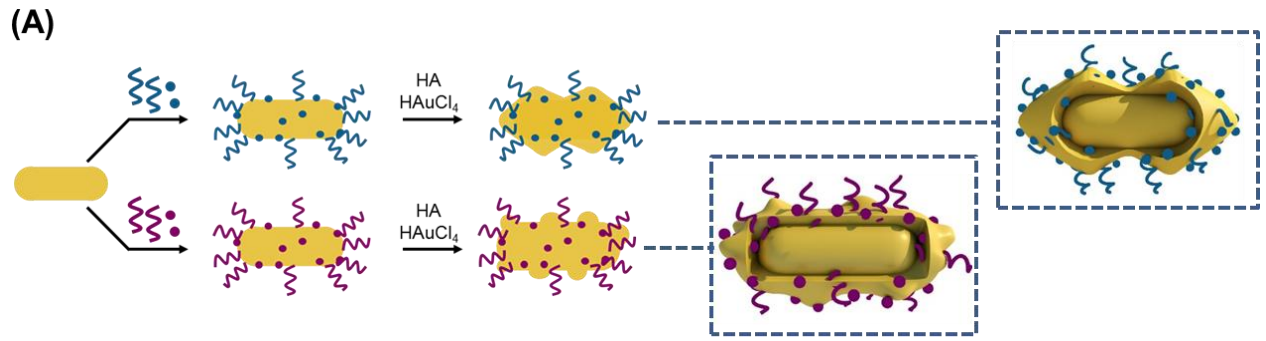
500 **Fig. 3** Optimization of the SERS biosensor. Effects of aptamer-1 concentration on tag-1 (A),
501 aptamer-2 concentration on tag-2 (B), concentration of 5,5'-dithiobis-(2-nitrobenzoic
502 acid) on tag-1 (C), concentration of 4-mercaptobenzoic acid on tag-2 (D), tag-1
503 concentration on *E. coli* O157:H7 detection (E), and tag-2 concentration on *S.*
504 *typhimurium* detection (F).

505 **Fig. 4** Bio-recognition specificity of the capture probes and SERS tags. TEM image of the
506 capture probe–pathogen–SERS tag sandwich complexes and corresponding magnified
507 image, obtained from the square area, of *E. coli* O157:H7 (A), *S. typhimurium* (B), and
508 the mixture of *E. coli* O157:H7 and *S. typhimurium* (C); capture rates of the capture
509 probes at different concentrations of *E. coli*. O157: H7 (D) and *S. typhimurium* (E).

510 **Fig. 5** Evaluation of quantitative detection, signal stability and specificity properties of the SERS
511 biosensor. (A) SERS spectrum for *E. coli*. O157: H7 and *S. typhimurium* in a mixed
512 system; (B) Linear correlation between the logarithms of pathogen concentrations and
513 SERS intensities of the peaks at 1330 cm⁻¹ for *E. coli*. O157: H7 and 1074 cm⁻¹ for *S.*

514 *typhimurium*; (C) Raman intensities of 20 random samples separately measured from a
515 suspension of *Escherichia coli*. O157: H7 and *Salmonella typhimurium* at 10^4 cfu/mL; (D)
516 Raman intensities for *E. coli* O157:H7, *S. typhimurium* and three interfering bacteria. The
517 mixture sample in Fig. 5D consisted of 5 different bacteria, including *E. coli* O157:H7, *S.*
518 *typhimurium*, *E. coli*, *S. aureus*, and *Vibrio parahemolyticus*.

519 **Table 1** Comparison of the simultaneous detection of *Escherichia coli* O157:H7 and *Salmonella*
520 *typhimurium* in three real samples by the surface-enhanced Raman scattering biosensor,
521 versus classic plate counting.

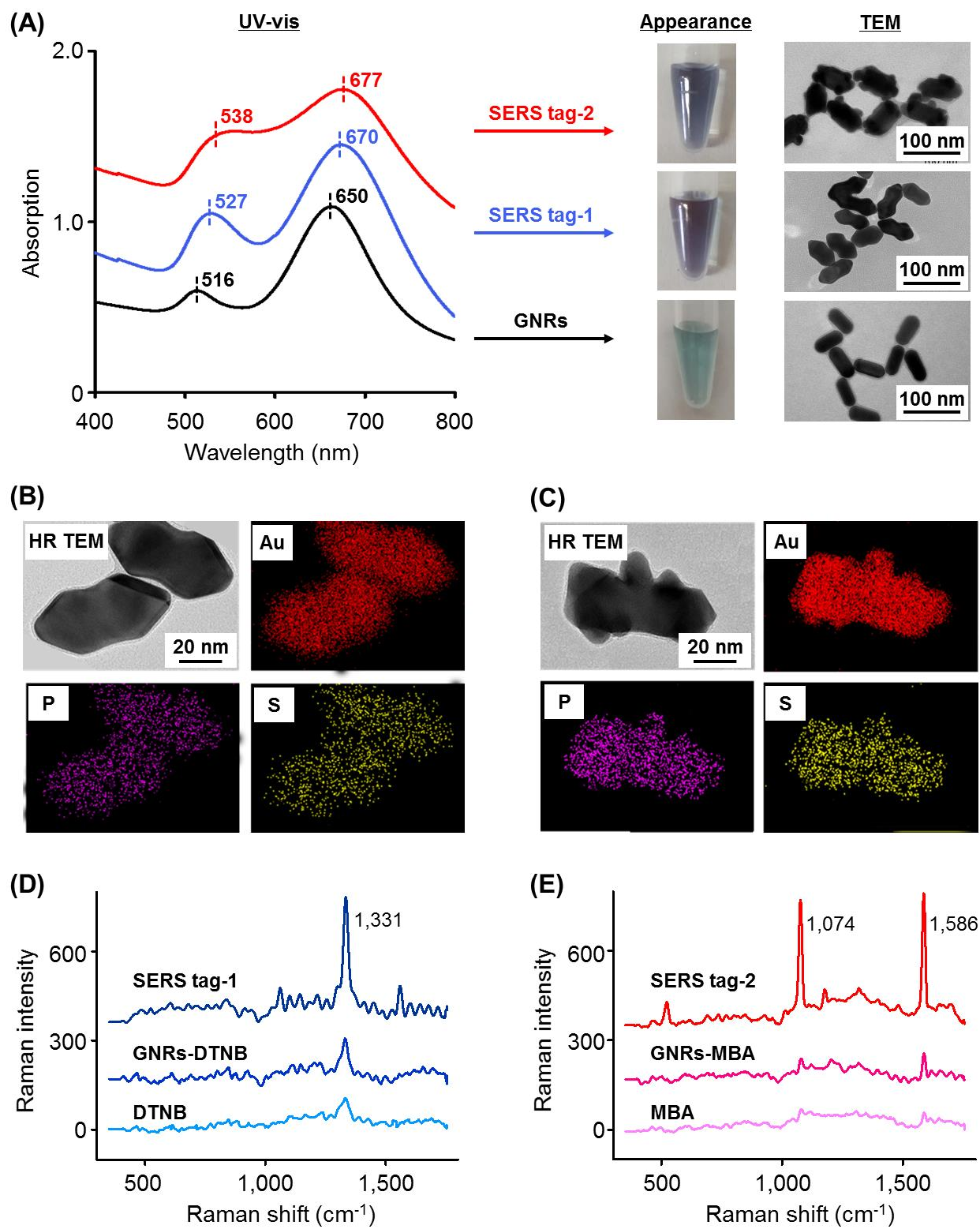


▬ GNRs
 ~ Aptamer-1
 ● Reporter-1
 ⬢ SERS tag-1 (cracked octahedral shape)
 ◉ *E. coli* O157:H7
 ⬢ Fe₃O₄-antibody-1
▲ Matrix
 ~ Aptamer-2
 ● Reporter-2
 ⬢ SERS tag-2 (small protrusion shape)
 ◉ *S. typhimurium*
 ⬢ Fe₃O₄-antibody-2

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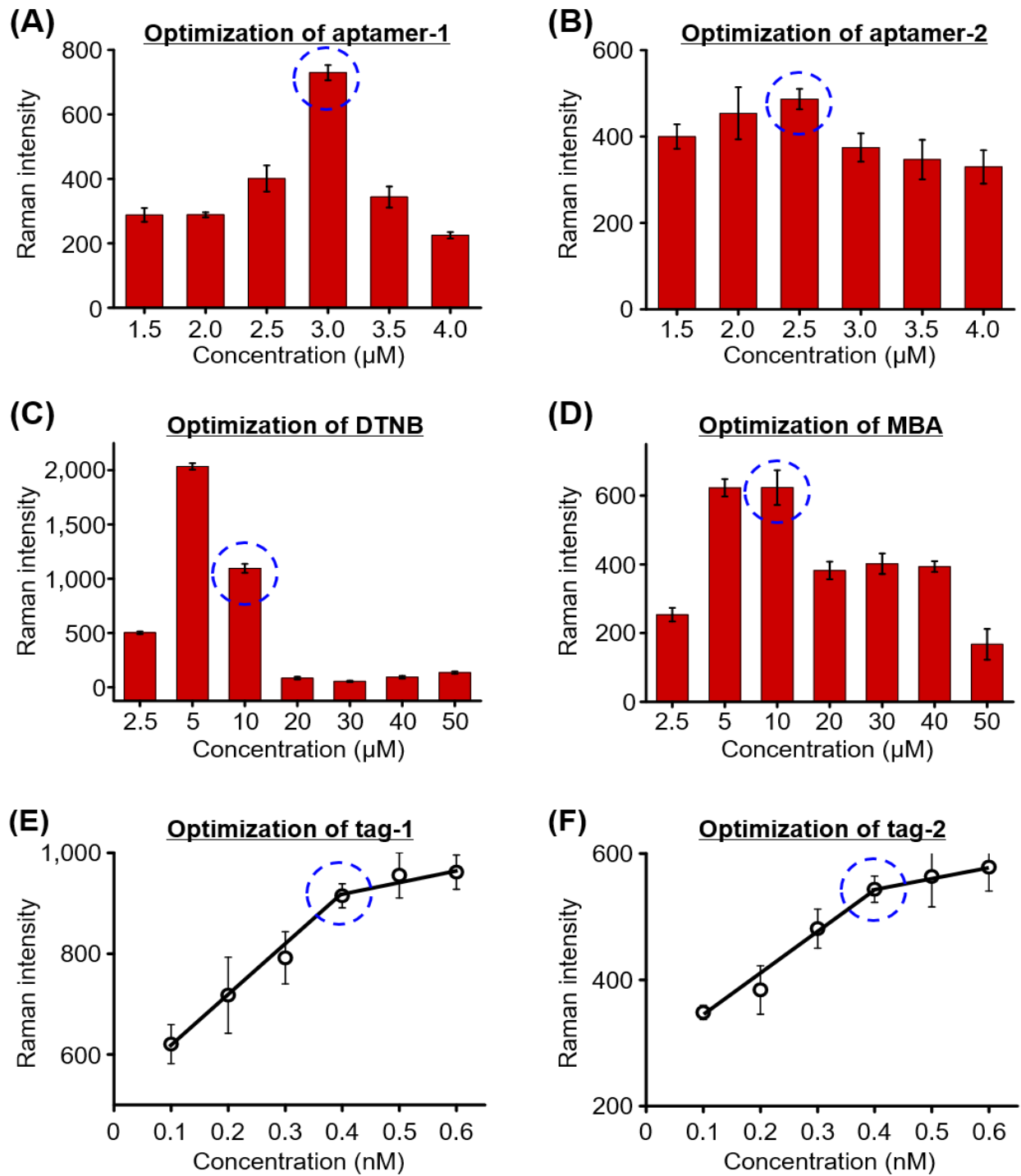
Fig. 1



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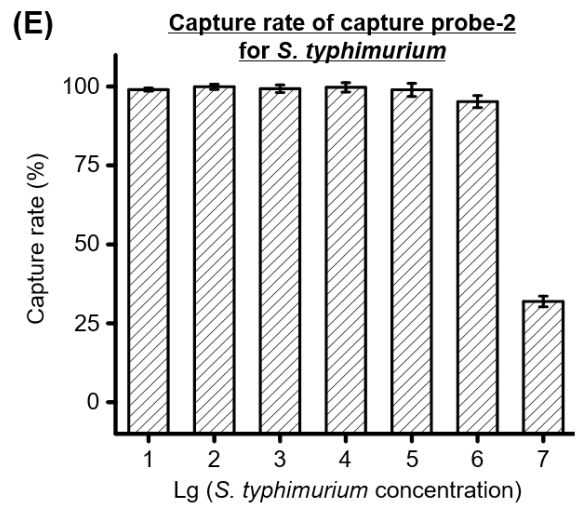
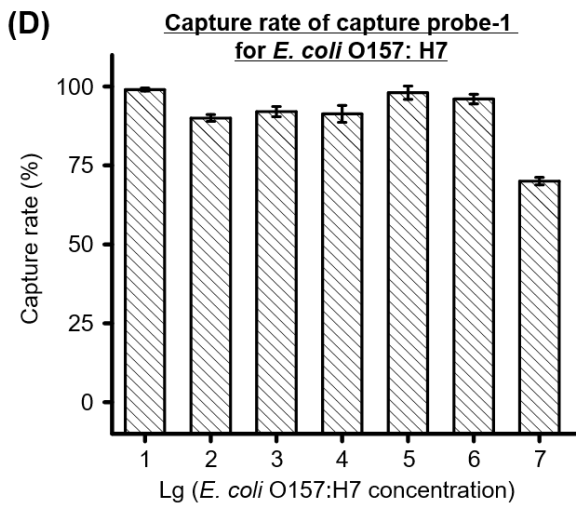
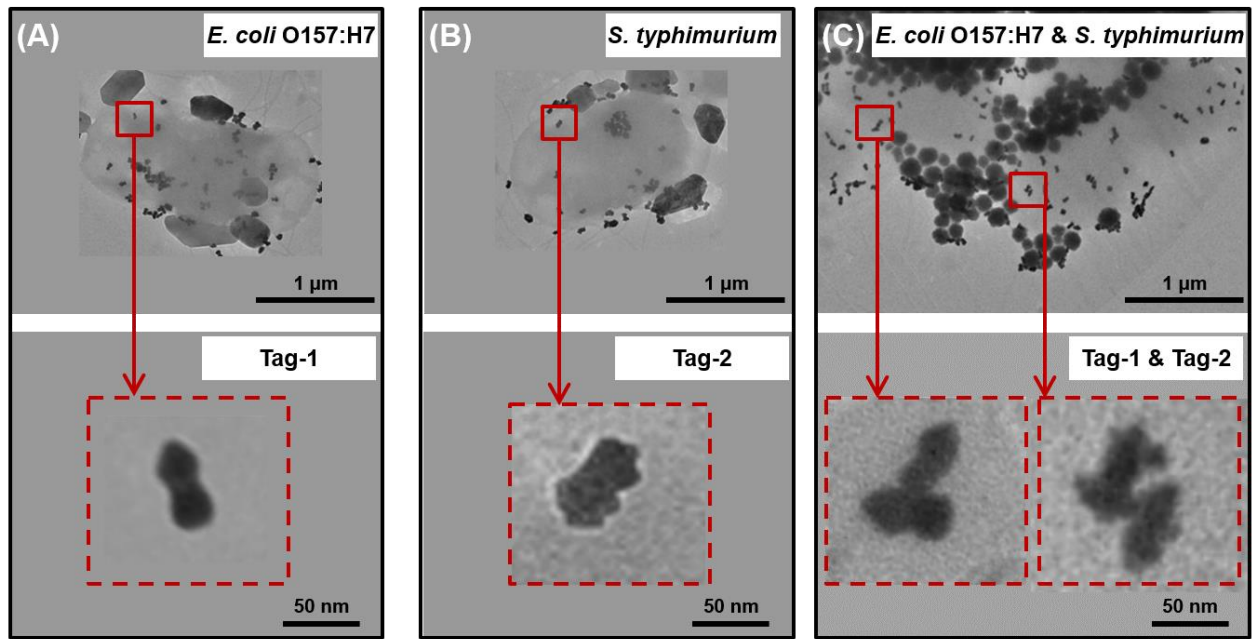
Fig. 2



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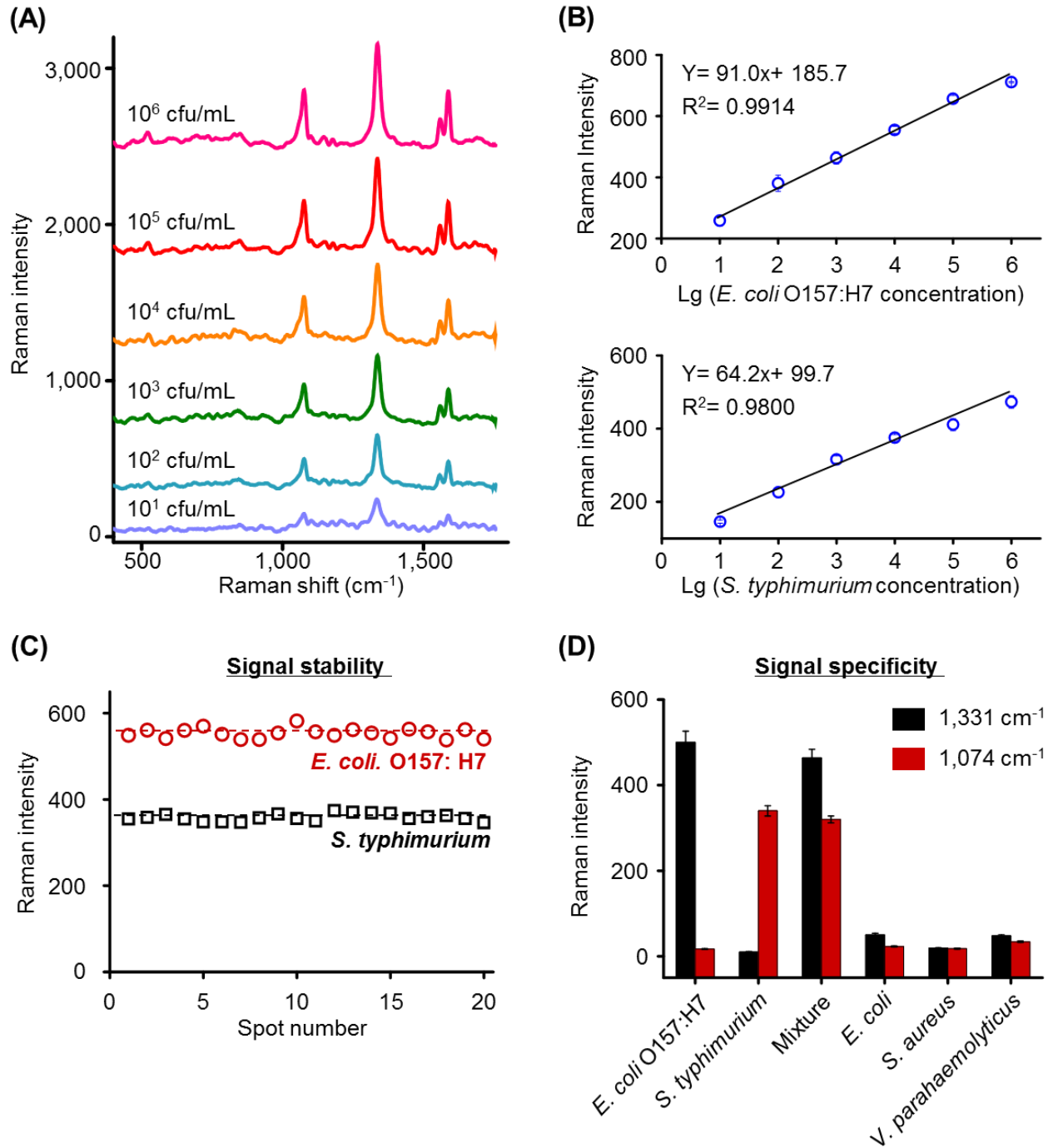
Fig. 3



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Fig. 4



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531

Fig. 5.

Table 1

	Our method		Plate counting method		Recovery rate (%)	
	<i>E. coli</i> O157:H7 (cfu/mL)	<i>S. typhimurium</i> (cfu/mL)	<i>E. coli</i> O157:H7 (cfu/mL)	<i>S. typhimurium</i> (cfu/mL)	<i>E. coli</i> O157:H7	<i>S. typhimurium</i>
Tap water	329 ± 26	194 ± 13	343 ± 12	193 ± 8	95.92	100.52
Cucumber	357 ± 21	181 ± 15	367 ± 5	190 ± 8	97.28	95.26
Chicken	376 ± 32	219 ± 17	357 ± 16	203 ± 10	105.32	107.88