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**- Food Science of Animal Resources -**  
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5

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## 34 **Introduction**

35

36           Campylobacteriosis is an infectious disease that can cause gastrointestinal  
37 symptoms including diarrhea, abdominal pain, and vomiting. Guillaine-Barre syndrome  
38 is one of the complications of campylobacteriosis that can damage nerve system of  
39 humans (Yang et al., 2013). The European Center for Disease Prevention and Control  
40 and the European Food Safety Authority ranked campylobacteriosis as the most  
41 common zoonosis in Europe (Euro surveillance Editorial Team, 2012). The most  
42 common species that can cause campylobacteriosis in humans is *C. jejuni*, followed by  
43 *C. coli* (Tam et al., 2003). Contact with live animals and consuming of raw poultry have  
44 been defined as the main sources of *Campylobacter* infection (Tam et al., 2003).

45           To confirm species origin for pathogens, analytical methods are mostly based  
46 on protein or DNA analysis (Ayaz et al., 2006). However, protein-based isolation  
47 method has a weakness. When the sample is exposed to high temperatures and pressure,  
48 the protein has a general tendency to degenerate (Murugaiah et al., 2009). Compared  
49 with protein-based methods, DNA based methods are more dependable because of their  
50 unique variability and high stability. Among DNA based methods, PCR methods,  
51 including conventional PCR, real-time PCR (Camm`a et al., 2012; Kesmen et al., 2012)  
52 and PCR-RFLP (Chen et al., 2010; Haider et al., 2012), have high specificity and  
53 accuracy. Recently, specific nanoparticles, including silver (AgNPs) and gold (AuNPs)  
54 nanoparticles, have been widely used as components of new technologies to detect  
55 pathogens (Du et al., 2013), hazardous materials (Li et al., 2009), DNA (Benedetto et al.,  
56 2011), small molecules (Lv et al., 2013), and aptamers (Ping et al., 2012). Aptamer-  
57 based detection has been carried out in our previous research (Kim et al., 2018). When  
58 various solutions change from dispersion to aggregation state, specific nanoparticles

59 exhibit significant color change because of their unique optical properties, that can be  
60 observed using a UV-visible wave-based spectrophotometer or the naked eye (Du et al.,  
61 2020; Shams et al., 2019). Thus, this research was designed to use AuNPs. A few types  
62 of AuNP-based colorimetric assays have been previously reported. In particular, the  
63 colorimetric assay based on thiol-labeled PCR primer and gold nanoparticles has been  
64 previously reported for the detection or diagnosis of disease (Htoo et al., 2019; Osmani  
65 Bojd et al., 2017.). The nanoparticle method is favored in the surveillance of pathogens  
66 over other PCR methods, due to its advantages over other PCR methods such as  
67 conveniency and time saving. To our knowledge, however, few studies have focused on  
68 the selective detection of fastidious foodborne pathogens such as *Campylobacter*. Since  
69 conventional ways to detect *Campylobacter* based on culture method require more than  
70 approximately 4 days (Masdor et al., 2016), various rapid detection methods have been  
71 developed for decades (Yang et al., 2013).

72 In this study, we aimed to develop a colorimetric assay integrating PCR and  
73 AuNP conjugation technology. The current study also outlined a colorimetric assay for  
74 the PCR-amplified *Campylobacter* genes, that could be directly identified by the naked  
75 eye.

76

## 77 **Materials and Methods**

78

### 79 **Bacterial strains and cultivation**

80 All oligonucleotides (Denis et al., 1999) were synthesized and purified by  
81 Bionics (Seoul, Korea) (Table 1). For the inclusivity test, some *C. jejuni* (A total of four  
82 *C. jejuni* strains: ATCC 33560 and 3 wild type strains from chicken carcasses) and *C.*  
83 *coli* strains (A total of four *C. coli* strains: ATCC 33559 and 3 3 wild type strains from

84 chicken carcasses) were used. *Campylobacter* strains were isolated from chicken  
85 carcasses from a local slaughterhouse and a poultry farm. Non-*Campylobacter* strains  
86 for the exclusivity test including *Salmonella* Enteritidis (*S. Enteritidis*) and *Escherichia*  
87 *coli* (*E. coli*), were also isolated from the collected chicken carcasses.

88 All *Campylobacter* strains were incubated in Bolton broth (Oxoid, Hampshire,  
89 UK) at 42°C for 42 h under microaerobic condition (5% O<sub>2</sub>, 10% CO<sub>2</sub>, and 85% N<sub>2</sub>),  
90 while non-*Campylobacter* strains were incubated in tryptic soy broth (Oxoid) at 37°C  
91 for 24 h. Additionally, phosphate-buffered saline was applied to dilute samples in the  
92 sensitivity test of gold nanoparticle (GNP)-PCR and 4 log CFU of *Campylobacter* and  
93 non-*Campylobacter* strains were incubated in 30 g of chicken meat samples with 30 ml  
94 of Bolton broth (42°C for 42 h) and tryptic soy broth (37°C for 24 h). Thereafter, 10 µl  
95 of cultured bacteria was used for GNP-PCR direct detection test.

96

### 97 **Genomic DNA isolation**

98 Approximately 1 ml cultures were used for DNA extraction. DNA was  
99 extracted by using QIAamp minikit (Qiagen, Germany). The concentration of extracted  
100 DNA was measured by using Nanodrop 2000 (Thermo Scientific, Wilmington, DE,  
101 USA) and genomic DNA was stored at 4°C before PCR amplification.

102

### 103 **PCR amplification**

104 PCR was conducted under the following conditions for the amplification of the  
105 *mapA* (*C. jejuni*) and *ceuE* (*C. coli*) genes. The reaction was carried out in a 50 µl  
106 mixture with 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, 10 mM Tris-HCl, 0.5 mM of  
107 deoxynucleoside triphosphate, 200 nM of forward/backward primer, 1 unit of Taq DNA  
108 polymerase (TaKaRa, Japan), and 2 µl (200 ng) of template DNA. The details of

109 primers used in this study is presented in Table 1. The amplification initiated with 10  
110 min at 95°C, followed by 30 cycles of 30 s at 94°C, 90 s at 59°C, and 60 s at 72°C, and  
111 then 10 min at 72°C. PCR was conducted in a Veriti 96 well thermal cycler (Applied  
112 Biosystems, Waltham, Massachusetts, USA). Exactly 2 µl of PCR products were then  
113 used for a gel electrophoresis with 1.0% agarose gel. The samples were run at 100 V for  
114 30 min, followed by some amount of PCR products being visualized with image  
115 analysis software (image lab software V3, Bio-Rad, CA, USA). The length of expected  
116 PCR product was 589 bp and 462 bp for *mapA* and *ceuE* gene, respectively.

117

### 118 **AuNPs synthesis**

119 Sodium citrate dehydrate and HAuCl<sub>4</sub> were from Sigma Aldrich (St. Louis,  
120 USA). GNPs (13 nm in diameter) were prepared via HAuCl<sub>4</sub> citrate reduction. All  
121 glassware was cleaned in KOH-IPA (Potassium hydroxide + isopropyl alcohol) solution  
122 and washed with pure H<sub>2</sub>O. A 50 ml of a 38.8 mM trisodium citrate solution was  
123 rapidly mixed with HAuCl<sub>4</sub> solution (1 mM, 500 ml), resulting in a change of color  
124 from yellow to dark red. The solution was then refluxed for another 15 min to cool it  
125 down to 25°C. The characteristics of the GNPs were verified using a surface plasmon  
126 band centered at 520 nm (Grabar et al., 1995).

127

### 128 **Colorimetric assay-based GNPs**

129 The center of the surface plasmon band of GNPs was 520 nm. A 6 µl of PCR  
130 product was mixed with 40 µl of the gold colloid for the colorimetric assay. They were  
131 mixed for 1 min of mixing, and another 10 µl of 1 M NaCl was added. The color was  
132 identified by the naked eye, and they were quantified using Nanodrop 2000 (Thermo  
133 Scientific, Wilmington, DE, USA). Since the aggregations between GNPs at 520 and at

134 650 nm are different, the ratio of absorbance 520/650 (Abs520/Abs650) was chosen.

135

### 136 **Transmission electron microscopy (TEM)**

137 The colloidal gold solution was dropped onto 1000 mesh of copper grids that  
138 was coated by carbon, followed by being dried in ambient conditions. Images were  
139 acquired using TEM JEM-2100 (JEOL Ltd, Japan) from Seoul National University. At  
140 least ten locations on the TEM copper grid were examined.

141

### 142 **Data analysis**

143 Analysis of different Abs520/Abs650 data for *Campylobacter* spp. for  
144 specificity was compared to that for non-*Campylobacter* spp. and the data for various  
145 concentrations of PCR products in the sensitivity analysis were analyzed using one-way  
146 ANOVA, which was conducted using GraphPad Prism software (GraphPad Software,  
147 San Diego, CA).

148

### 149 **Results and Discussion**

150

#### 151 **The design of colorimetric assay for the detection of *Campylobacter***

152 Fig. 1 represents the main principle of the suggested method. The main idea  
153 behind the GNP colorimetric assay was the dispersion and aggregation of colloidal  
154 GNPs. It was effectively demonstrated that the interparticle movement and pulsive  
155 forces are the main reason for GNP stabilization or aggregation (Napper, 1983; Zhao et  
156 al., 2008). GNP electrostatic stabilization relies on the surface charge of the repulsive  
157 electric double layer. The layer can stabilize the colloids against Van der Waals forces  
158 (Evans and Wennerstrom, 1999). When the electric double layer of GNP is suppressed,

159 the electrostatic repulsion force dramatically decreases at high concentrations of salt.  
160 That is why citrate-combined GNPs are stabilized in water but aggregated in salty  
161 conditions (Zhao et al., 2008). Thus, the covalently combined complex of thiolated  
162 double-strand DNA (dsDNA) and monodispersed GNP can resist the aggregation  
163 regardless of salt addition.

164 AuNPs maintain a red color in stable state. When they aggregated, the state of  
165 AuNPs becomes unstable, causing change of color from red to blue or purple. The main  
166 focus of the current research was that AuNPs can be modified by thiolated PCR  
167 products, that prevented AuNPs from salt-induced aggregation (Chan et al., 2014; Fu et  
168 al., 2013; Jyoti et al., 2010). Because of this attribute, we modified unique  
169 *Campylobacter* target primers with an unlabeled backward primer and a thiol-labeled  
170 forward primer. Therefore, thiol-labeled target gene sequence was obtained. When  
171 AuNPs were mixed with thiolated PCR products, they were surrounded by a thick layer  
172 that consisted of dsDNA. Therefore, this structure tended to make a change of color.

173 For a successful result, GNPs need a thick obstacle with a lot of negative  
174 charges to inhibit their aggregation. We modified the primer for *mapA* of *C. jejuni* (589  
175 bp) and *ceuE* of *C. coli* (462 bp) for PCR amplification. *mapA*, which is usually called  
176 membrane-associated protein A gene, does not cross-react with *C. coli* proteins  
177 (Gonzalez et al., 1997; Stucki et al., 1995). *ceuE* is a lipoprotein-encoding gene which  
178 has been used for *C. coli* identification (Gonzalez et al., 1997). To confirm PCR  
179 efficiency, thiol-labeled PCR primers used for *C. jejuni* and *C. coli* were the same as  
180 unlabeled primers used for the conventional PCR.

181 PCR and gel electrophoresis were conducted to analyze the PCR products. Fig. 2  
182 shows that PCR products from thiol-labeled- and unlabeled primers are identical in size.  
183 It shows that the result obtained using the thiolated primers did not show any size

184 difference to conventional PCR products.

185

### 186 **Thiolated PCR products**

187 For the detection of *Campylobacter* spp., thiol-labeled PCR products prevented  
188 GNP aggregation based on salt-inducing. Both unlabeled and thiol-labeled PCR  
189 products were added into the GNP solution and the color change was compared. After  
190 adding NaCl solution, GNP solution with unlabeled PCR products turned from red to  
191 blue and that with thiol-labeled products remained red. Spectroscopic analysis verified a  
192 sharp peak of surface plasmon resonance (SPR) at approximately 520 nm. This narrow  
193 SPR peak appeared due to the red colored solution containing thiol-labeled PCR  
194 products after salt adjusting. A broad SPR band (530-650 nm) seemed in the blue  
195 colored solution with unlabeled PCR products, indicating GNP aggregation after salt  
196 addition (Fig. 3). To confirm the characteristic features of GNPs in the colorimetric  
197 assay, different behaviors of the salt-induced aggregation of GNPs were observed using  
198 TEM JEM-2100 (JEOL Ltd, Japan) (Fig. 4).

199

### 200 **The specificity of the assay**

201 A color change was observed in the GNP solution containing non-labeled PCR  
202 products, whereas the GNP solution with thiol-labeled PCR products are still red (Fig.  
203 5). This can be identified with the naked eye.

204 Abs<sub>520</sub>/Abs<sub>650</sub> of *C. jejuni* and *C. coli* isolates were significantly different (*C.*  
205 *jejuni*: 4.86±0.27; *C. coli*: 4.81±0.28) compared to that of *S. Enteritidis* and *E. coli* (*C.*  
206 *jejuni* *P*-value: 0.0085; *C. coli* *P*-value: 0.0239) in five replicates. Thus, 200 ng/μl of  
207 *mapA* and *ceuE* were detected in the sample with thiol-labeled products after salt  
208 adjusting.

209 **The sensitivity of the assay**

210 For the determination of the sensitivity, different concentrations of *mapA* from  
211 *C. jejuni* were tested. Abs520/Abs650 of the detected samples are shown in Fig. 6. Fig.  
212 6 (A) shows that at least 15 ng/ $\mu$ l of *C. jejuni mapA* PCR product was needed to detect  
213 *C. jejuni* using the optical method. At this concentration, Abs520/Abs650 was  
214  $3.38\pm 0.17$  (mean $\pm$ standard deviation (S.D.)). Fig. 6 (B) shows that at least  $1 \times 10^3$   
215 CFU/ml of *C. jejuni* was needed to use the colorimetric method. At this concentration,  
216 Abs520/Abs650 was  $3.26\pm 0.35$  (mean $\pm$ S.D.). These results represented that the  
217 Abs520/Abs650 increased with the increasing number of PCR products and cells. To  
218 verify the precision of the assay, one-way ANOVA was used to determine the *P*-value  
219 after measuring the absorption at different concentrations in five replicates.

220 More PCR products can be obtained by applying more amplification cycles.  
221 However, too many amplification cycles lead to low efficiency because of the formation  
222 of non-specific bands due to DNA degeneration (Cha and Thilly, 1993). Therefore, 30  
223 cycles of PCR protocol was selected in this study. The colorimetric system was  
224 controled by NaCl. The concentration of NaCl affects the dynamics of the method as  
225 well as the sensitivity of the detection. The limit of detection (LOD) of various types of  
226 detection methods are compared in Table 2. qPCR is slightly more sensitive than GNP-  
227 PCR (Table 2). However, the detection limit of the GNP-PCR was similar or superior to  
228 other detection methods such as ELISA, GNP-adjusted aptasensor, and Lateral flow  
229 (Table 2).

230 As we stated above, the thiol-labeled PCR primer and gold nanoparticles have  
231 been applied in many research fields for the detection or diagnosis of disease. Htoo et al.  
232 (2019) used thiol-labeled PCR primer and unmodified gold nanoparticles for the  
233 detection of target RNA for the diagnosis of prostate cancer and found that the assay

234 was specific and sensitive for the RNA detection in prostate cancer cell lines with a  
235 visual detection limit of 31.25 ng/reaction. Osmani Bojd et al. (2017) reported  
236 application of thiolated AuNP probes and multiplex PCR for the detection of  
237 *Staphylococcus epidermidis*. They found that the minimum quantity of target DNA for  
238 multiplex PCR was 1 ng/mL and for color and absorption alteration of solution in  
239 colorimetric assay was 20 ng/mL. Since *Campylobacter* is fastidious bacteria that  
240 requires long incubation time and labors, development of this types of rapid and  
241 convenient isolation method is highly useful for the screening of the pathogen. In the  
242 present study, the newly developed method showed slightly lower sensitivity compared  
243 to qPCR according to Table 2. However, this method could be favored over qPCR  
244 especially in field tests as the assay could be conducted without further steps that  
245 require cost, labor, and detection device. With the assay, the positives could be easily  
246 distinguished from the negatives with the naked eye. Thus, the developed assay could  
247 be finished only with thermal cycler and optical detection, while conventional or qPCR  
248 still need further steps after PCR cycles. Even though qPCR is superior to this assay in  
249 terms of sensitivity, the developed method could be widely used in many research fields  
250 considering its convenience for the detection of pathogens.

251

### 252 **Direct detection of *Campylobacter* in foods**

253 To estimate the detection ability of GNP-PCR in food pathogenic bacteria, the  
254 method was applied to a food sample. Three groups of chicken meat samples were  
255 purchased from local retails. A group was artificially spiked with an overnight culture of  
256 *C. jejuni*, while the others were contaminated with *S. Enteritidis* and *E. coli*. Artificially  
257 contaminated samples were cultured at 42°C overnight and 10 µl of each sample was  
258 directly adjusted to the GNP-PCR assay. The results of these assays are shown in Fig. 7.

259 When thiol-labeled primers were used in the rinsate of fresh chicken meat, the color of  
260 the GNP solution changed from red to blue after salt adjusting. Abs520/Abs650 was  
261  $1.06 \pm 0.22$  (mean  $\pm$  S.D.). However, if the *C. jejuni* template was present, the color of  
262 GNP solution is still red after salt adjusting. This result suggested that the GNPs  
263 remained dispersed, while the Abs520/Abs650 of *mapA* was  $4.30 \pm 0.3$ . With the  
264 spectroscopic results, the sample groups can be clearly distinguished.

265 The detection ability of the GNP-PCR assay for *C. jejuni* and *C. coli* was well  
266 defined according to sensitivity assays results. The length of PCR amplicons and  
267 amplification efficiency were important factors for the detection. Previous research  
268 found that GNPs highly tend to aggregate when the PCR amplicons were shorter than  
269 400 bp. When the amplicons were longer than 400 bp, it became a limiting factor for  
270 efficient detection (Fu et al., 2013). As we stated above, GNPs require a thick barrier  
271 with a lot of negative charges to inhibit the aggregation. We selected the 589 bp *mapA*  
272 for *C. jejuni* detection and the 462 bp *ceuE* for *C. coli* detection.

273

## 274 **Conclusion**

275 The suggested assay enables the generation of a PCR product with high  
276 amplification efficiency and the convenience of a colorimetric assay. Moreover, leading  
277 to a limit of optical detection of density around 15 ng/ $\mu$ l of *Campylobacter* verified high  
278 specificity. The suggested assay has a couple of advantages. Above all, the PCR  
279 colorimetric method has high sensitivity, specificity, and amplification efficiency.  
280 Furthermore, this method avoids bacterial culture steps, reduces the time needed, and  
281 requires no complicated apparatus. Moreover, the proposed method allows the detection  
282 of pathogens even by the naked eye. Therefore, it appears that this assay has high  
283 potential for a various bio-related detections. Lastly, as we stated above, this method

284 does not require complex devices, while detecting conveniently with the naked eye.

285 The current research reported a novel and unique colorimetric assay for the  
286 detection of *Campylobacter* spp. in raw poultry. Considering high specificity of the  
287 assay, the accurate identification of *Campylobacter* spp. from other environmental  
288 samples is also available with this method.

289

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399 **Table 1. Primer pairs for *Campylobacter* spp.**

Target gene	Sequence (5'- 3')	Amplicon size	Reference
<i>mapA</i> ( <i>C. jejuni</i> ) forward	SH-C12-CTA TTT TAT TTT TGA GTG CTT GTG	589 bp	(Denis et al., 1999)
<i>mapA</i> ( <i>C. jejuni</i> ) reverse	GCT TTA TTT GCC ATT TGT TTT ATT A		
<i>ceuE</i> ( <i>C. coli</i> ) forward	SH-C12-AAT TGA AAA TTG CTC CAA CTAT TG	462 bp	(Denis et al., 1999)
<i>ceuE</i> ( <i>C. coli</i> ) reverse	TGA TTT TAT TAT TTG TAG CAG CG		

400 \*Thiol (-SH) formations were applied to 5' end of the primer sequence.

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413 **Table 2. Comparison of the analytical performance of the developed GNP-PCR assay with several**  
 414 **reported assays.**

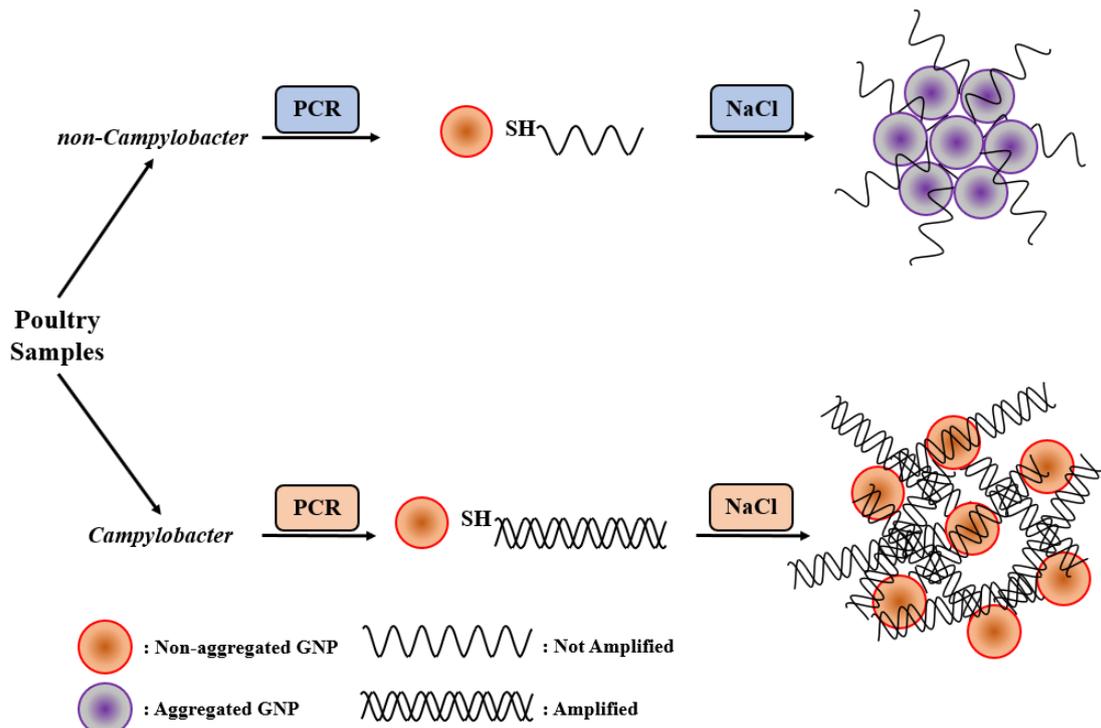
<b>Methods</b>	<b>Target bacteria</b>	<b>LOD (cfu/mL)</b>	<b>Reference</b>
ELISA	<i>C. jejuni</i>	$4.5 \times 10^4$ cfu/mL	(Che, Li, & Slavik, 2001)
GNP-adjusted aptasensor	<i>C. jejuni</i>	$7.2 \times 10^5$ cfu/mL	(Kim et al., 2018)
QD adjusted Fluorescent biosensor	<i>C. jejuni</i>	$1 \times 10^3$ cfu/mL	(He et al., 2018)
Lateral flow	<i>C. jejuni</i>	$1 \times 10^7$ cfu/mL	(Wadl et al., 2009)
qPCR	<i>C. jejuni</i>	$1.25 \times 10^2$ cfu/ml	(Suh, Dwivedi & Jaykus 2014)
GNP-PCR assay	<i>C. jejuni</i>	$1 \times 10^3$ cfu/mL	The present study

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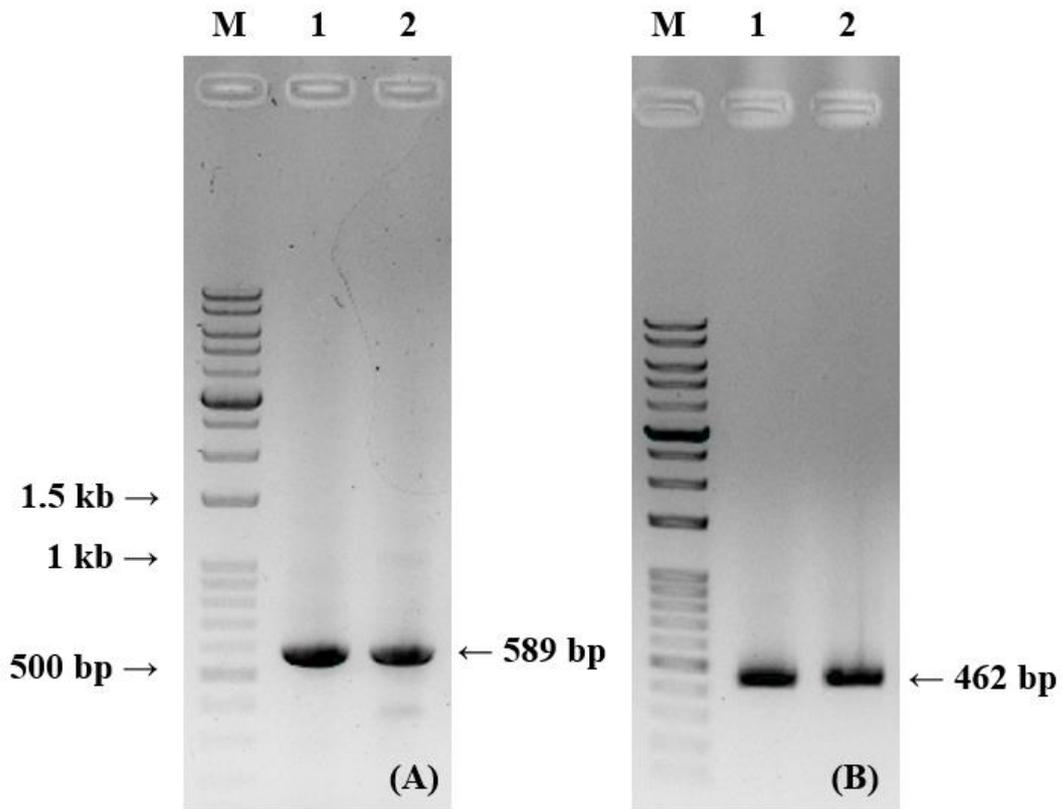
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**Fig. 1. The schematic diagram for the PCR-based gold nanoparticle colorimetric assay for detection of *Campylobacter*. *mapA* and *ceuE* were used for *C. jejuni* and *C. coli*, respectively.**



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427 **Fig. 2. Verification of PCR.** PCR products were electrophoresed in a 1% agarose gel.

428 The DNA ladder is indicated in lane M. Lane 1 indicates PCR products with no thiol-

429 labeled primers and lane 2 represents PCR products with thiol-labeled primers. Thiol-

430 labeled and non-labeled PCR products had similar sizes. Thiolated primer did not cause

431 size differentiation. (A) Agarose gel electrophoresis of *mapA* from *C. jejuni* ATCC

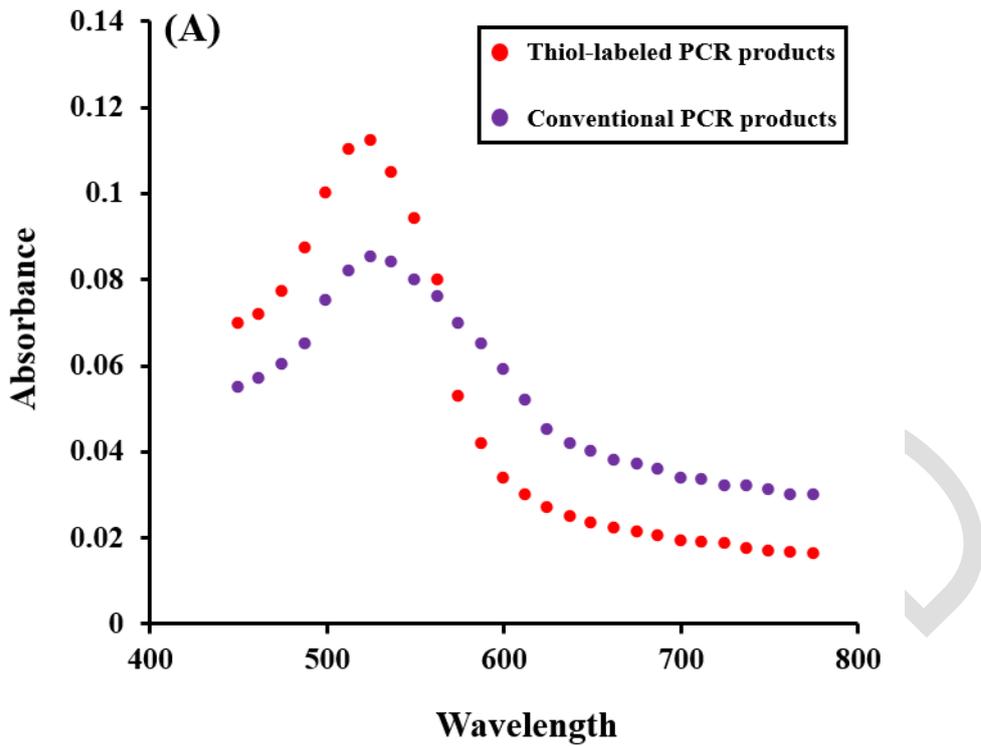
432 33560 (589 bp). (B) Agarose gel electrophoresis of *ceuE* gene from *C. coli* ATCC

433 33559 (462 bp).

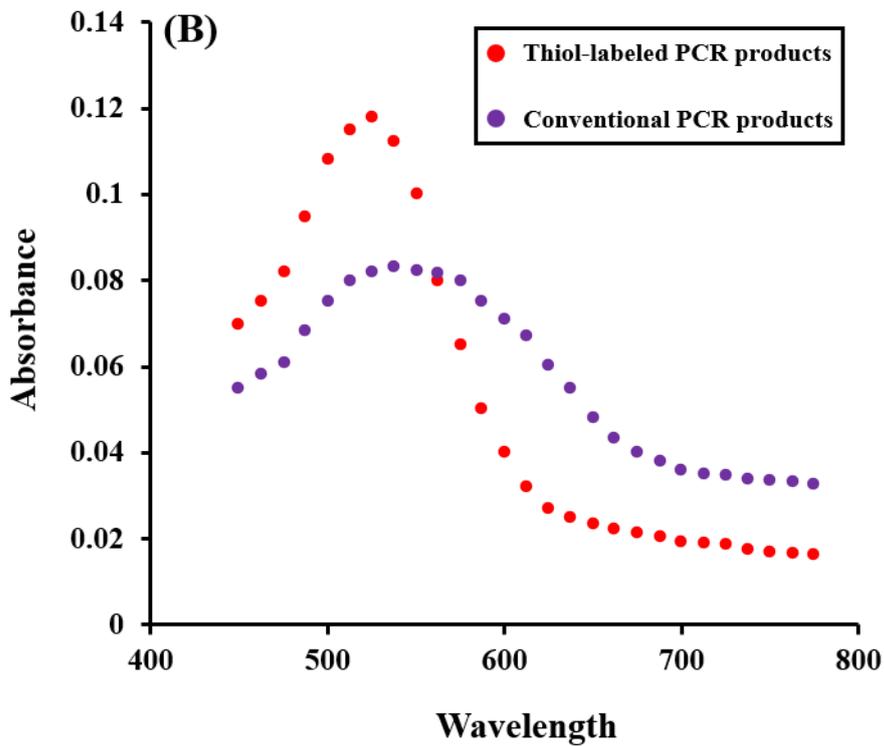
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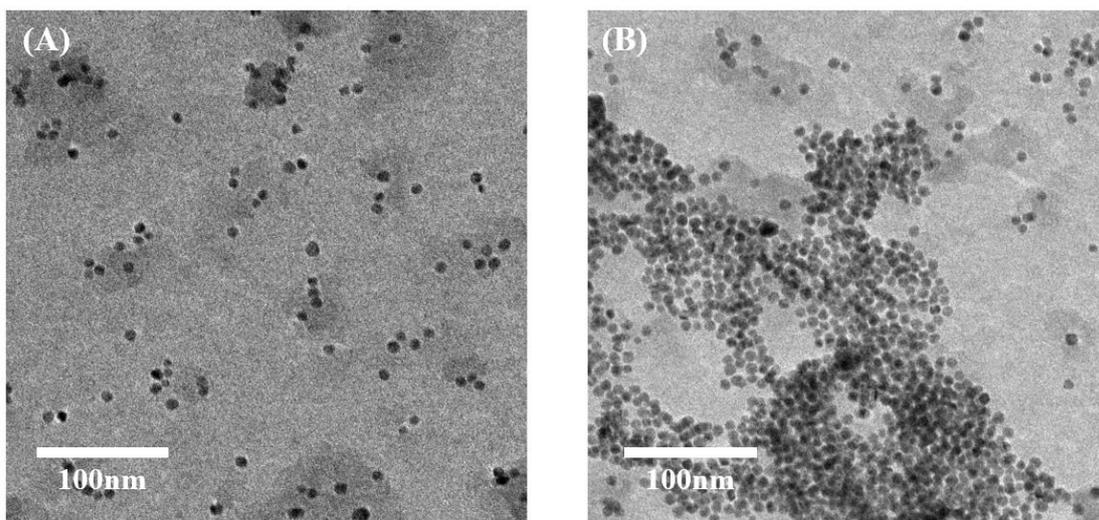


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439 **Fig. 3.** Colorimetric assays. The effect of thiol-labeled PCR products is verified based  
440 on *mapA* of *C. jejuni* (A) and *ceuE* of *C. coli* (B) isolated from chicken carcass samples.



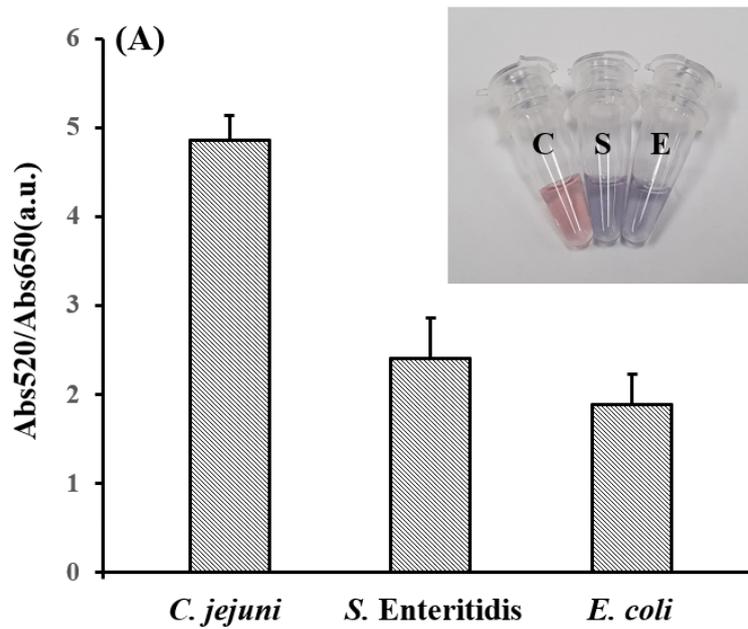
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442 **Fig. 4.** TEM images of AuNPs after the addition of thiol-labeled PCR product with salt

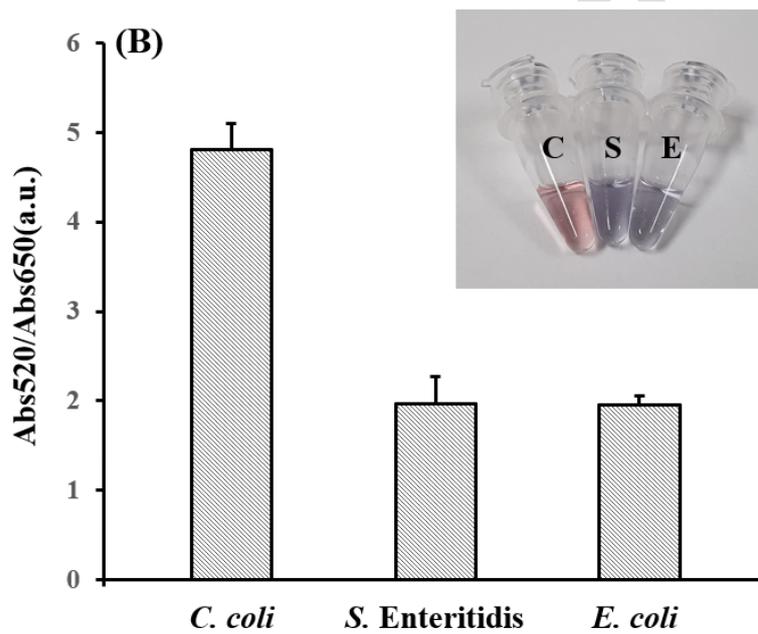
443 (A) and unlabeled PCR product with salt (B). TEM, transmission electron microscopy.

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449 **Fig. 5. Specificity analysis.** Specificity evaluation of absorption spectra analysis of  
 450 GNPs mixed with the PCR products of *C. jejuni* and *C. coli*. (A) Specificity evaluation  
 451 of absorption spectra analysis of GNPs mixed with the PCR products of *C. jejuni*. Tube  
 452 C shows the result of the colorimetric assay for *C. jejuni*, tube S shows that for *S.*  
 453 *Enteritidis*, and tube E shows that for *E. coli*. (B) Specificity evaluation of absorption  
 454 spectra analysis of GNPs mixed with the PCR products of *C. coli*. Tube C shows the

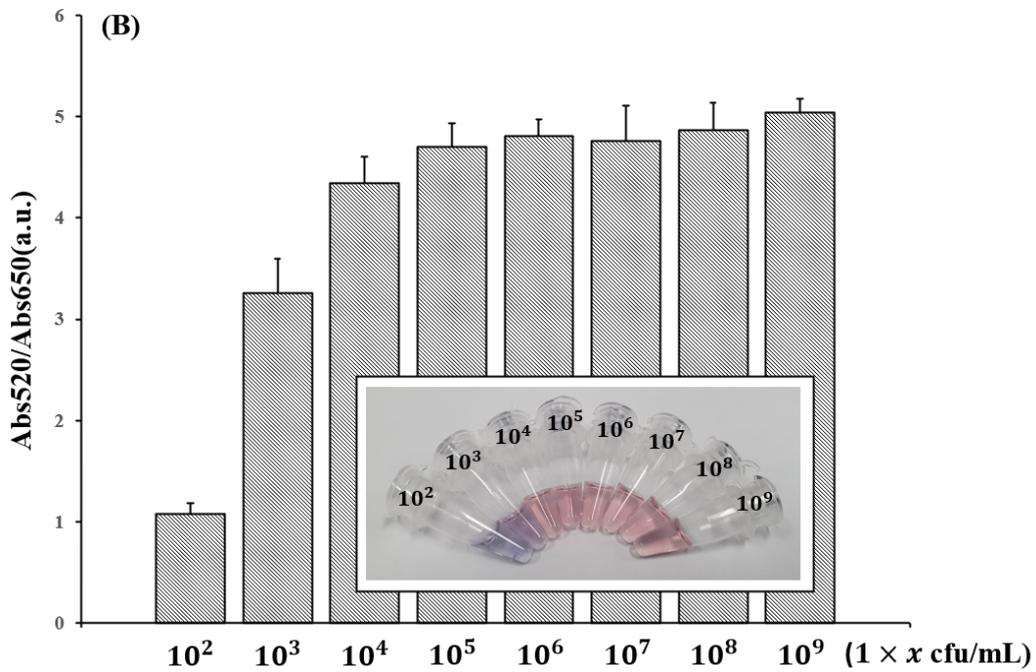
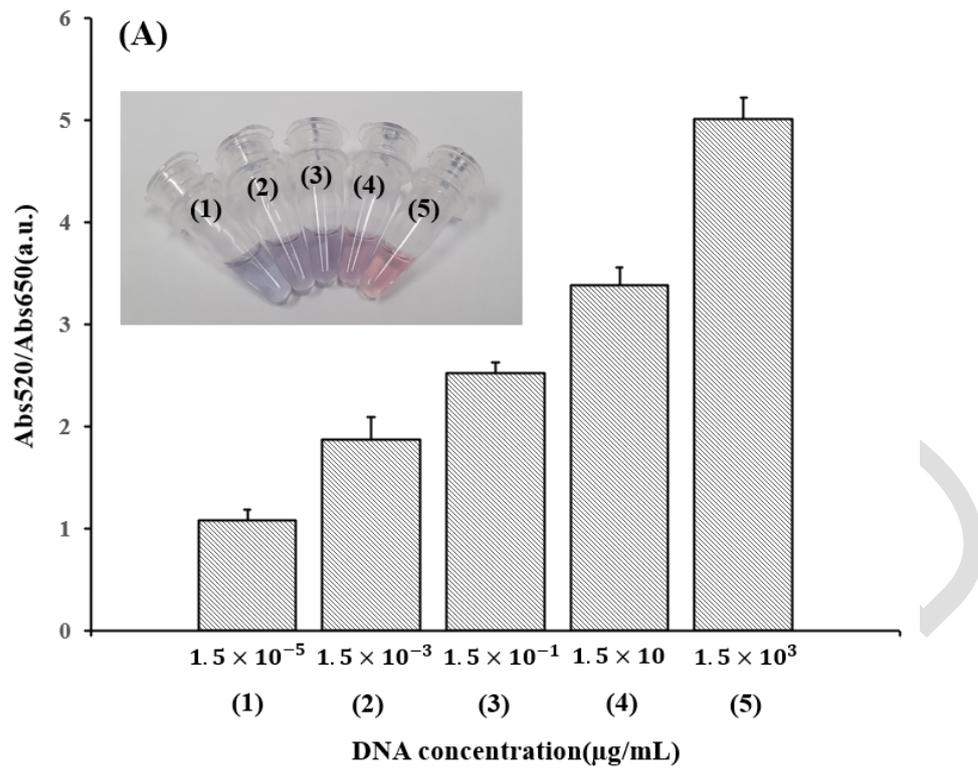
455 result of the colorimetric assay for *C. coli*, tube S shows that for *S. Enteritidis*, and tube

456 E shows that for *E. coli*. GNP, gold nanoparticle.

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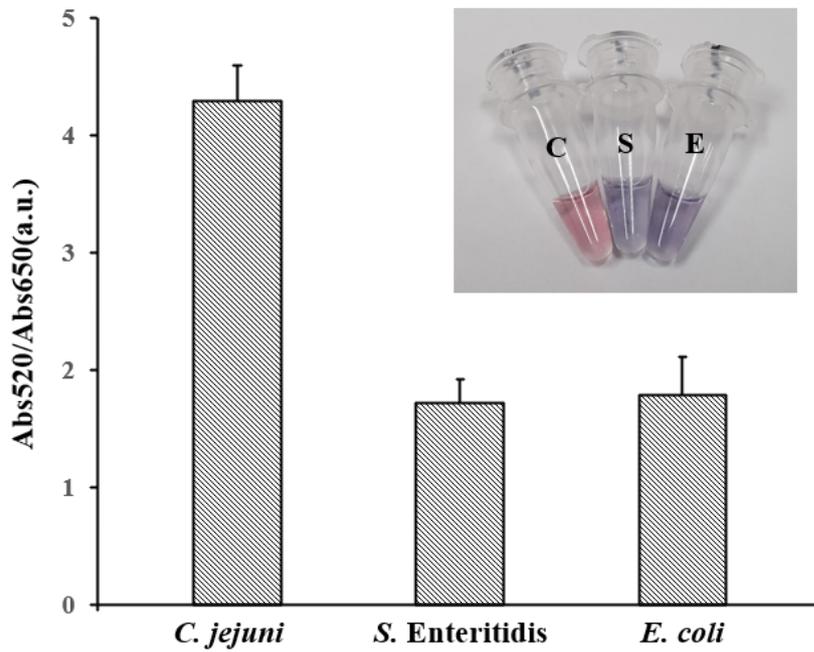
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462 **Fig. 6. Sensitivity analysis.** The sensitivity of different concentrations of GNP-PCR

463 colorimetric assay for DNA (A) and cells (B) based on *mapA* of *C. jejuni*.



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468 **Fig. 7. *C. jejuni* detection in artificially contaminated chicken meat.** Tube C shows  
469 the result of the colorimetric assay for *C. jejuni*, tube S shows that for *S. Enteritidis*, and  
470 tube E shows that for *E. coli*. All three samples were obtained from artificially  
471 contaminated chicken meat samples.

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