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## Visual detection of *Listeria monocytogenes* using unmodified gold nanoparticles based on a novel marker

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### Abstract:

*Listeria monocytogenes* (*L. monocytogenes*) make people and animals have Listeria disease. After infection, the main symptoms of Listeria disease are sepsis, meningitis and mononucleosis. For this purpose, a visual detection of *L. monocytogenes* was developed with unmodified gold nanoparticles. The species-specific probes, in the presence of PCR target products of *L. monocytogenes*, cause the gold nanoparticles to aggregate irreversibly, producing a red to purple colorimetric change. As little as 1.304 fg/ $\mu$ l of DNA of *L. monocytogenes* were thus detected visually, by the naked eye, without the need for any sophisticated, expensive instrumentation and biochemical reagents, which was approximately seventy-fold higher sensitivity over conventional PCR gel electrophoresis method. The results indicate that this assay is highly species-specific, simple, low-cost, and visual for easy detection of *L. monocytogenes*.

**Keywords:** visual detection; *Listeria monocytogenes*; gold nanoparticles

### Introduction

*L. monocytogenes* has been recognized as a cause of disease in humans and animals and has been responsible for listeriosis. It can be found in dairy products, meat, poultry, vegetables, as well as in frozen food because of growing at refrigeration temperatures. The consumption of food products contaminated with this bacterium can cause listeriosis, a serious disease with a 30% mortality rate<sup>1</sup>. At home and abroad the damage of *L. monocytogenes* gets more and more valued, and WHO listed it as one of the four major food-borne pathogens in 1990s. Recent consumption of frozen food in China has increased, the potential risk of *L. monocytogenes* is becoming more and more outstanding, simultaneously, risk assessment data from the United States ranks several categories of dairy products as high-to moderate risk in terms of predicted listeriosis cases on a per serving basis<sup>2</sup>. The ubiquitous distribution of this pathogen in nature, its ability to grow at refrigeration temperatures and its tolerance to certain preservative agents make its elimination from food very difficult. A fast and sensitive detection of this pathogen would be extremely important for early prevention of disease spreading.

Nowadays, many different kinds of methods have been evaluated for detecting *L. monocytogenes* in food samples, including conventional culture-, biochemical-, and immunological-based assays<sup>3</sup>, polymerase chain reaction (PCR) and RT-PCR based methods<sup>4-6</sup>, loop-mediated isothermal amplification assays<sup>7-9</sup>, biosensors<sup>10,11</sup>, matrix-assisted laser desorption ionization time-of-flight (MALDI-TOF) mass spectroscopy<sup>12</sup>, Fourier transform infrared (FT-IR) spectroscopy<sup>13</sup>. Among these methods, calculating the number, biochemical characteristics and identification of colony are time-consuming, immunological assays can suffer from low specificity, resulting in false-positive results<sup>14</sup>, because most commercially developed antibodies are not

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3 species-specific. RT-PCR, biosensors, MALDI-TOF, and FT-IR assays are not routinely used  
4 because of the expensive requirements. LAMP can easily produce false-positive results even  
5 though a very small amount of contaminant is in reaction systems<sup>7</sup>.

7 The polymerase chain reaction (PCR) assay has the advantages in that it can be highly specific,  
8 sensitive, rapid and may permit direct detection of the pathogen in foods and other samples  
9 without the need for isolation of pure cultures. Various genes have been targeted for the detection  
10 of *L. monocytogenes*, including *hlyA*<sup>15</sup>, *iap*<sup>16</sup>, *inlA*<sup>17</sup>, *inlAB*<sup>15</sup>, *actA*<sup>18</sup> and *lmo0733*<sup>19</sup>. Although  
11 *lmo0202hly* gene fragment is a potential molecular detection marker for *L. monocytogenes* in  
12 bioinformatics analysis in our previous research<sup>20</sup>, it still needs to be identified by biological  
13 experiments in this study.

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17 Gold nanoparticles (AuNPs) have been used as sensing material for colorimetric detection due  
18 to their unique optical properties<sup>21</sup>. It presents a color change of AuNPs colloids based on  
19 electrostatic interaction between AuNPs with nucleotide sequences. Therefore, AuNPs can  
20 selectively aggregate due to the different characteristics of the single and double strand DNA, and  
21 provide a simple, inexpensive and colorimetric detection of various DNA sequences. However,  
22 many color assays are time consuming due to about a two-day's preparation for modified AuNPs<sup>22</sup>.  
23 In order to overcome limitations, a reliable, simple, cost-effective, and visual detection of *L.*  
24 *monocytogenes* based on unmodified gold nanoparticles and PCR has been developed in the study.

## 25 26 27 28 **Experimental**

### 29 30 **Materials and reagents**

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32 *L.monocytogenes*, *L.ivanuui*, *L.innocua*, *L.seeligeri*, *L.grayi*, *Listeria welshimeri*, *Salmonella*  
33 *choleraesuis*, *Salmonella typhimurium*, *Serratia marcescens* were supplied by Shanghai  
34 Entry-Exit Inspection and Quarantine Bureau, Shanghai, China. *Staphylococcus aureus* and  
35 *Carnation ringspot virus* (CRSV) were kept in our laboratory. HAuCl<sub>4</sub> (99.999%) and sodium  
36 citrate dehydrate were purchased from Sinopharm Group Chemical Reagent Co., Ltd., Shanghai,  
37 China. All glass was cleaned in aqua regia (3 parts HCl, 1 part HNO<sub>3</sub>) rinsed with pure H<sub>2</sub>O. The  
38 species-specific fragment from *Staphylococcus aureus* (289bp) and the species-specific fragment  
39 from *Carnation ringspot virus* (476bp) were supplied by our laboratory and used as controls. All  
40 oligonucleotides used in the study were synthesized and purified by Sangon Biotech.

### 41 42 43 44 **Extraction of Genomic DNA or RNA**

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47 For DNA isolation, 1 mL of overnight pure culture BHI broth was centrifuged and washed once  
48 with physiologic saline. Genomic DNA was isolated from the pellet using the TIANamp Bacteria  
49 DNA Kit (TianGenBiotech, Beijing, China) according to the manufacturer's instructions and  
50 dissolved in ddH<sub>2</sub>O. Viral genomic RNA was extracted using TIANamp Virus RNA Kit  
51 (TianGenBiotech, Beijing, China) according to the manufacturer's instructions. The RNA was  
52 eluted with 60 µL of elution buffer and stored at -80°C pending analysis.

### 53 54 55 **Preparation of negative control**

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57 289-bp fragment from *Staphylococcus aureus* and 475-pb fragment from *Maize chlorotic mottle*  
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3 *virus* (MCMV) were used to be controls and prepared as followed, respectively. The 289 bp of  
4 PCR products were gained according to our previous research<sup>23</sup>. 475-pb fragments from CMMCV  
5 were amplified with the primers (Forward primer: 5'-TCAGGTTTCATGCCCTCT-3', reverse  
6 primer: 5'- ATGCTTGCTCCATCCACT-3'). RT-PCR assays were performed on Eppendorf  
7 gradient thermocycler using One-Step PrimeScript™ RT-PCR Kit (Takara, China). All sets of  
8 reactions were carried out in a final volume of 25µl each containing: 0.5 µl of RNA (MCMV),  
9 12.5 µl of 2×OneStep RT-PCR Buffer, 1µl of PrimeScript enzyme (5U/µl), 0.5µl of MCMV PF(20  
10 µM), 0.5 µl of MCMV PR(20 µM), and 10 µl of RNase-free ddH<sub>2</sub>O. Amplification reactions were  
11 consisted of a reverse transcript at 42 °C for 25 min, an initial denaturing step at 94 °C for 2 min  
12 followed by 35 cycles at 94 °C for 30 s, 55 °C for 30 s and 72 °C for 30s followed by a final  
13 extension at 72 °C for 10 min.  
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### 17 **PCR procedure**

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19 In order to amplify the species-specific fragment for *L.monocytogenes*, the primes and probe  
20 were designed to target *lmo0202hly* gene fragment and were synthesized as follows: F:  
21 5'-ATCAACCAgATgTTCTCCCTg TA-3', R: 5'-GATTCACTgTAAgCCATTTTCgTC-3'. PCR  
22 was carried out in a 50 µl reaction mixture composed of 5.0 µl of 10×Taq DNA polymerase buffer,  
23 3.0 µl of 2.5 mmol/L dNTPs, 1.0 µl of Taq DNA polymerase, 1.0 µl of prime and reverse primer,  
24 38 µl of ddH<sub>2</sub>O and 1.0 µl of DNA. PCR reaction was performed in an Eppendorf gradient  
25 thermocycler. The cycling procedures for amplifying target sequence were set as follows: an  
26 initial step of denaturation at 95 °C for 2 min; 30 cycles of denaturation at 95 °C for 45 s, anneal at  
27 53 °C for 30 s and elongation at 72 °C for 55 s; final extension at 72 °C for 10 min. 5.0 µl of PCR  
28 product was analyzed by gel electrophoresis using a 1.5% agarose gel containing 0.5 µg of EB dye.  
29 The running conditions were stable voltage at 120 V for 30 min. After electrophoresis, the relative  
30 amount of PCR products was analyzed by image analysis software (Quantity One™, Bio-Rad,  
31 CA, USA). 600-bp DNA markers, which contain 600, 500, 400, 300, 200, and 100 bp DNA  
32 fragments, were used as standards for the evaluation of the gel electrophoresis results. The  
33 expected PCR product is the length of 226 bp.  
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### 39 **Preparation of gold nanoparticles (AuNPs)**

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41 AuNPs with the average diameter of 13nm were synthesized by a citrate reduction method.  
42 Briefly, AuNPs were prepared by boiling an aqueous solution of 0.01% HAuCl<sub>4</sub> (100 ml) under  
43 rapid stirring and adding 3.5 ml of 1% sodium citrate. After boiling for 15 min and further rapid  
44 stirring for 30 min, the solution was allowed to be cooled to room temperature and filtered through  
45 a 0.8 µm membrane. Concentration of the as-prepared AuNPs was estimated to be 3.5 nM, which  
46 was calculated from the quantity of starting material (HAuCl<sub>4</sub>) and the size of the nanoparticles.  
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### 49 **Effect of NaCl concentration on the complex of AuNPs with DNA**

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51 In order to achieve the optimal NaCl concentration to the complex of AuNPs with DNA, six  
52 different concentration of NaCl were induced in to the complex of AuNPs with double strand  
53 DNA, single strand DNA (probe) respectively.  
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### 55 **Specificity of visual detection of *L. monocytogenes***

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57 1 µl of 10-fold diluted PCR product and 1 µl of 10 mM probe (5'-  
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3 AAAAAATGCCACTAAATCAAA -3') were mixed completely in a sterile PCR tube. The  
4 mixture was denatured at 95°C for 2min, annealed at 53°C for 6min, and then cooled to room  
5 temperature for 10 min. After addition of 30 µl of colloidal AuNPs to the reaction mixture, 2.89 µl  
6 of 0.11M PBS buffer and ddH<sub>2</sub>O, the total volume reached 35µl and kept for 10 min at room  
7 temperature, the color changes in the solutions were observed visually and recorded by cameras.  
8 There existed 65 mM NaCl in the final mixture. To validate the specificity of this method, 39  
9 strains were tested using the above procedures.  
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### 12 **Sensitivity of visual detection of *L. monocytogenes***

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14 The sensitivity of the assay was evaluated by performing the above procedures at a series of  
15 diluted DNA of *L.monocytogenes*.  
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### 17 **Results**

#### 18 **The design of visual detection of *L.monocytogenes* based on AuNPs colorimetric assay**

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20 The detection strategy is shown in Scheme 1. Single strand DNA (ssDNA) adsorbs on  
21 citrate-coated AuNPs, and this adsorption increases the negative charge on the AuNPs, which give  
22 rise to increasing repulsion between the particles, thus preventing aggregation. The adsorption of  
23 ssDNA on AuNPs occurs due to the fact that ssDNA can uncoil and expose its nitrogenous bases,  
24 and the attractive electrostatic forces between the bases and the AuNPs allow adsorption of the  
25 ssDNA. Inversely, double strand DNA (dsDNA) does not adsorb on AuNPs due to the repulsion  
26 between its negatively-charged phosphate backbone and the negatively-charged coating of citrate  
27 ions on the surfaces of the AuNPs<sup>24, 25</sup>. Therefore, when AuNPs are added to a saline solution  
28 containing the target DNA and its complementary, specific probes, AuNPs aggregate (since the  
29 probes are not free to stabilize the AuNPs) and the solution color changes to grey blue. However,  
30 in the absence of the target or the presence of a non-complementary target, the probes remain free  
31 to stabilize the AuNPs thus preventing their aggregation, and the solution color remains red  
32 (Scheme.1). The denaturation and annealing help species-specific probes sufficiently absorb the  
33 targets. In appearance, the color difference of AuNPs can indicate whether the RCR product can  
34 combine complementally with species-specific probe or not, and identified the samples.  
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#### 41 **Optimal concentration of NaCl to color change of the complex of AuNPs colloid**

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43 Single strand DNA probes can absorb into the surface of AuNPs through electric interaction to  
44 increase the stability of colloidal of AuNPs, and the electric double layers of colloidal of AuNPs  
45 can also retain good stability. However, there exist a critical salt concentration, which can induce  
46 the complex of AuNPs with double strand DNA to deposit, but the complex of AuNPs with probes  
47 is still steadily stable. As shown in Fig.1, the color of four samples was clearly red at the  
48 concentration of NaCl below 60 mM (Fig.1A, B,C), however, the color of three samples turned to  
49 grey blue at the concentration of NaCl over 65 mM (Fig.1F), the color of one sample begin  
50 turning to grey blue when the concentration of NaCl was 62 mM (Fig.1D), only one sample was  
51 grey blue and others appeared red color (Fig.1E), this results demonstrated that the critical  
52 concentration of NaCl was 65 mM, which was sufficient for aggregation of AuNPs and  
53 guaranteed the color change of visual detection.  
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#### 57 **Specificity of visual detection of *L. monocytogenes***

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3 Based on our previous result<sup>20</sup>, *lmo0202hly* gene fragment was used to be a novel molecular  
4 marker for detection of *L. monocytogenes*, and PCR product was expected as the 226 bp fragment.  
5 As we supposed that all of *L. monocytogenes* (39 strains) could gain about 226 bp fragments in  
6 lane 1-39 in Fig.2A according to our designing specific primers and simultaneously, other non- *L.*  
7 *monocytogenes* (8 strains) cannot be successfully amplified in lane 40-47 in Fig. 2A with the same  
8 primers, the results showed that the novel molecular marker and the PCR assay are very high  
9 species-specific to *L. monocytogenes*.  
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12 The color change is the prominent indicator which determines the accuracy of the visual  
13 detection of *L. monocytogenes* assay. In the experiment, there were many non-specific  
14 amplification based on negative controls. Simultaneously there existed some specific  
15 amplification because of *L. monocytogenes*. The PCR of negative controls could not combine with  
16 the specific probe because that there were lack of target templates in lane 3-10 in Fig. 2B, which  
17 lead to free probes to absorb on the surface of AuNPs, increased NaCl resistance, and prevented  
18 the aggregation of AuNPs induced by NaCl, and the color in mixtures is always red in lane 1 and  
19 3-10 in Fig. 2B. In the other hand, the specific probes and the specific amplifications hybridized  
20 polymeric substance in lane 2 in Fig.2B, which lead to absence of probes, thus AuNPs could not  
21 be protected from salt induction to cause aggregation, and the color changed grey blue in lane 2 in  
22 Fig.2B. Compared with PCR assay, visual detection assay was highly species-specific to *L.*  
23 *monocytogenes*; In addition, only 0.1 $\mu$ l of PCR product in visual detection assay could make the  
24 same detection conclusion as the 7  $\mu$ l of PCR assay, which indicated that visual detection assay  
25 was seventy-fold (7/0.1) higher sensitivity than conventional PCR assay.  
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### 31 **High sensitivity of the visual detection assay**

32 In order to know the lowest detection limit of this new colorimetric assay system, PCR was  
33 performed on decreasing amounts of genome DNA. Mixtures of extensively amplified target,  
34 probes and AuNPs (lanes 1-7 in Fig.3B), changed gray blue after salt addition. Simultaneously, in  
35 cases where low DNA amplification yields could not appear in electrophoresis (lane 8-9) in  
36 Fig.3A, the solutions retained the red color after salt addition. the color difference among the  
37 various concentrations of DNA templates were clearly distinguishable owing to the strong  
38 aggregation of AuNPs occurring in the solution not containing enough templates. The  
39 observations described that as few as 1.304 fg/ $\mu$ l of the DNA of *L. monocytogenes* could be  
40 detected by observing the color change arising during the extended incubation time of 10 min  
41 following salt addition, it suggested that over the 1.304 fg/ $\mu$ l DNA template could amplify enough  
42 specific PCR product to combine with the specific probes, so that there were not adequate free  
43 probes to prevent from strong aggregation of AuNPs after salt induction. The result implied the  
44 extent of amplification dominated the sensitivity of this colorimetric DNA detection methodology.  
45 In addition, 0.1 $\mu$ l of DNA used in visual detection method had consistent results with 7 $\mu$ l of DNA  
46 used in PCR, which indicated the visual detection of *L. monocytogenes* was more seventy fold  
47 sensitive than gel electrophoresis.  
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### 53 **Discussion**

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55 *L. monocytogenes* spreads widely in fresh meat, aquatic product, vegetables, and ubiquitous and  
56 capable of growth at refrigeration temperature, which may contaminate foods at any point of  
57 production, distribution or storage. Nowadays, several methods have been developed to identify *L.*  
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3 *monocytogenes* infection. PCR has been a simple and rapid detection method for detection *L.*  
4 *monocytogenes*. In fact, the signal transduction gene, *vicK* has been identified as a molecular  
5 marker for detection of *Staphylococcus aureus* in our previous research based on comparative  
6 genome and sequences alignment analysis<sup>26</sup>. With the similar strategy, 226-bp sequence (*lmo0202*  
7 *hly*) has been identified as a novel molecular marker for detection of *L. monocytogenes* by means  
8 of bioinformatics in our previous research<sup>20</sup>, and tested in the biological experiment in this study.

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11 NaCl concentration is an important contributor to affect the color of AuNPs colloidal solution.  
12 Although single strand DNA probes adsorb on AuNPs to help them more stable and prevent their  
13 aggregation, the increase in NaCl concentration will destroy gradually the double electrostatic  
14 layer of AuNPs, and once NaCl concentration is over its critical value, AuNPs will aggregate  
15 unavoidably, and the solution turns blue. Certainly, the critical concentration of NaCl can vary  
16 according to various target, probes and AuNPs.

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19 Except for NaCl concentration, the concentration of the probes also affects the AuNPs state. In  
20 the presence of the target, a very high probe concentration will not only hybrid with target, and the  
21 rest of probes can absorb the surface of AuNPs to prevent aggregation leading to a false negative  
22 result. On the other hand, in the absence of the target the low concentration probes absorb AuNPs,  
23 and excess AuNPs will aggregate after salt induction leading to a false positive result.  
24 Consequently, the optimal probe concentration was found to be 10  $\mu\text{M}$  in the total assay volume.

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27 The denaturation and annealing steps are deemed necessary before the addition of the AuNPs to  
28 increase the specificity of the assay, this because that the extent of PCR targets combination with  
29 the species-specific probe is the key of the successful assay, and increasing the time of the  
30 denaturation and annealing steps also increases the percentage of positive results. In our opinion,  
31 increasing the time of denaturation and annealing might increase the probability of the probes  
32 annealing specifically to species-specific PCR products in *L. monocytogenes*, hence, after adding  
33 the species-specific probes to PCR products, the mixture was subjected to denaturation at 95 °C  
34 for 2 min, annealing at 53 °C for 6min, and then cooling to room temperature for 10 min in our  
35 experiment.

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38 Excellent sensitivity, specificity, and speed have made molecular assays an attractive alternative  
39 to culture or enzyme immunoassay methods. Wang et al reported the sensitivity of detection of *L.*  
40 *monocytogenes* by cross-priming amplification of *lmo0733* gene is 2.5 pg DNA per reaction<sup>19</sup>;  
41 Liu et al described the sensitivity of a novel paper-based microfluidic enhanced  
42 chemiluminescence biosensor for detection of *L. monocytogenes* is  $6.3 \times 10^{-2}$  pmol/l of DNA<sup>27</sup>. In  
43 our experiment, the sensitivity of detection *L. monocytogenes* is 1.304 fg/ $\mu\text{l}$  of DNA templates, it  
44 means the sensitivity of the novel detection developed in our study is 1.304 fg DNA per reaction,  
45 and  $1.98 \times 10^{-2}$  pmol/l, all of these suggest visual detection of *Listeria monocytogenes* using  
46 unmodified gold nanoparticles based on *lmo0202hly* gene in our study has more sensitivity than  
47 others.

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50 To our knowledge, this is the first approach employing probes for visual detection of *L.*  
51 *monocytogenes* using unmodified AuNPs. The new method requires only addition of AuNPs with  
52 the solution after PCR. Surface modification of AuNPs is not required. Although it still requires a  
53 PCR step, post-PCR analysis such as gel electrophoresis is eliminated in this method. As a  
54 consequence of its simplicity, the methodology should find wide application in assays performed  
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in small-scale corporations or in developing countries, where only inexpensive equipment and nonhighly trained personnel need to be available.

## Conclusion

In this work, we took full advantage of interaction AuNPs with nucleic acids and developed a low-cost, facile, sensitive method for visual detection of *L. monocytogenes* with the unmodified AuNPs. The assay described is easily read with the naked eye. In comparison with other methods for detection of *L. monocytogenes*, the method is more attractive because of its high sensitivity, low cost, ready availability and simple manipulation. This is the first application of the unmodified AuNPs-based biosensing platform for detection of *L. monocytogenes*.

## Acknowledgement

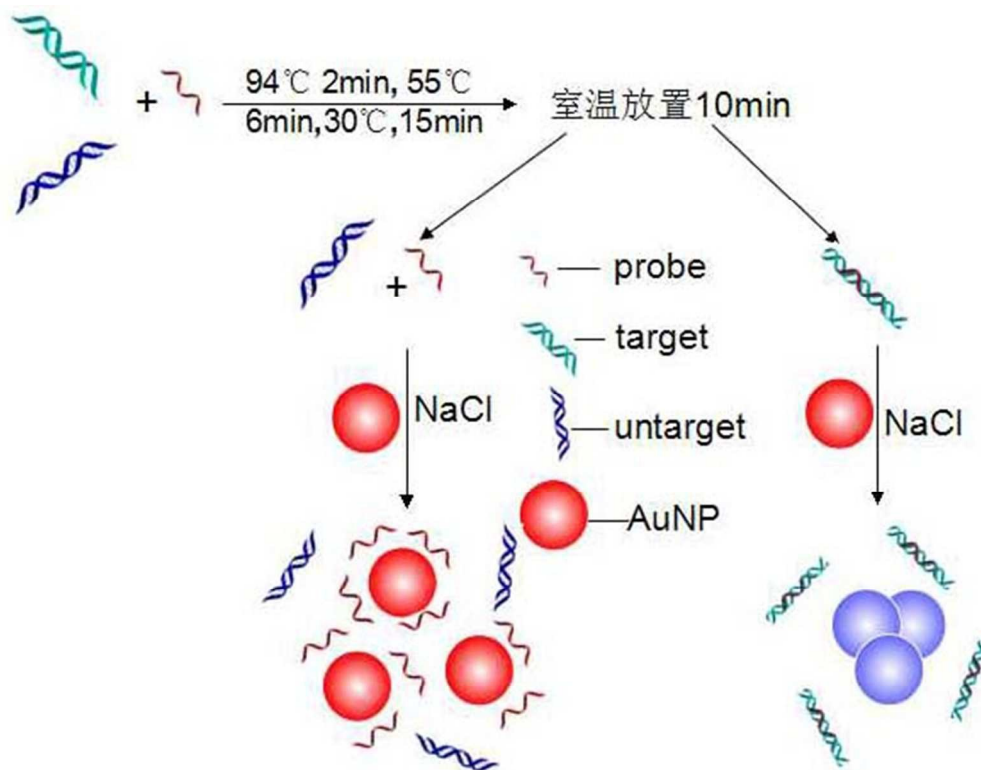
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Scheme 1. Schematic illustration for visual detection of MCMV based on unmodified AuNPs.  
156x121mm (96 x 96 DPI)

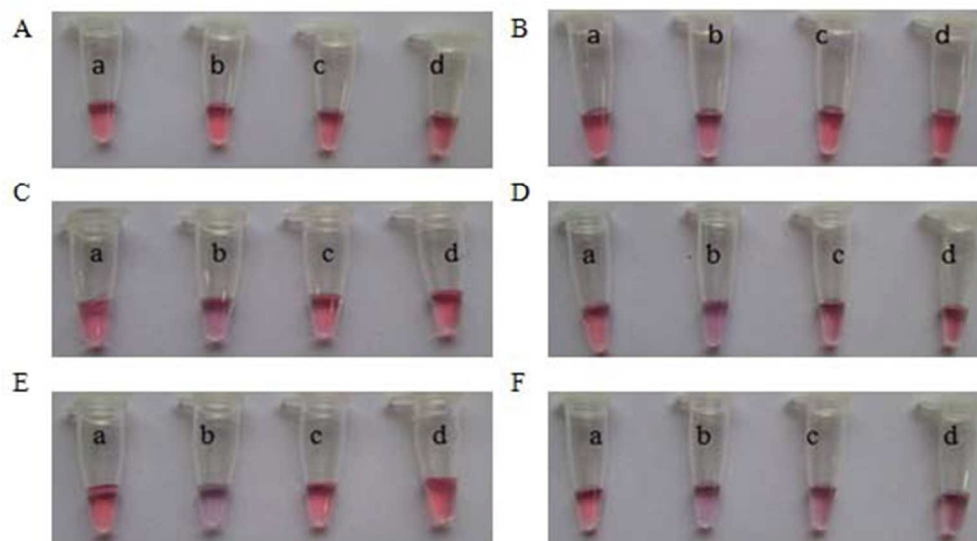


Fig. 1 Critical NaCl concentration for the color change of existing status of AuNPs collides. 35  $\mu$ l of mixtures (containing 0.1  $\mu$ l DNA or distilled water, 1  $\mu$ l of 10  $\mu$ M probes, 30  $\mu$ l of AuNPs, and some distilled water) was subjected to denaturation at 95  $^{\circ}$ C for 2 min, annealing at 53  $^{\circ}$ C for 6min, and then cooling to room temperature for 10 min before adding different concentration of NaCl induction for 90 min. (a) distilled water as control; (b) 226-bp PCR of *L. monocytogenes*; (c) 476-bp RT-PCR of MCMV RNA; (d) 289-bp PCR of *Staphylococcus aureus*; (A) 40 mM NaCl; (B) 50 mM NaCl; (C) 60 mM NaCl; (D) 62 mM NaCl; (E) 65 mM NaCl; (F) 70 mM NaCl.

192x106mm (72 x 72 DPI)

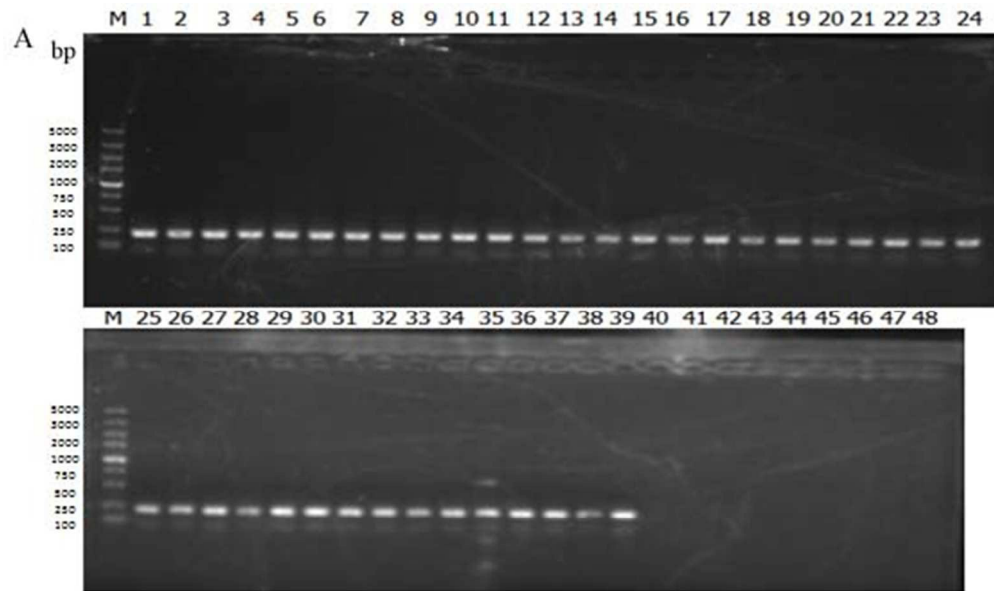


Fig. 2 Specificity of visual detection of *L. monocytogenes*. (A) Electrophoresis of PCR products from different microorganisms and control. M was the DNA ladder; Lane 1-39 represented *L. monocytogenes*; 40-47 were *L. ivanuii*, *L. innocua*, *L. seeligeri*, *L. grayi*, *Listeria welshimeri*, *Salmonella choleraesuis*, *Salmonella typhimurium*, *Serratia marcescens*, respectively; lane 48 was distilled water ; (B) Different results of PCR combination with a species-specific probe after salt induction. Lane 1 was distilled water and used as control; lane 2 was *L. monocytogenes*; lane 3-10 represented *L. ivanuii*, *L. innocua*, *L. seeligeri*, *L. grayi*, *L. welshimeri*, *Salmonella choleraesuis*, *Salmonella typhimurium*, *Serratia marcescens*, respectively.  
196x121mm (72 x 72 DPI)



14 (B) Different results of PCR combination with a species-specific probe after salt induction. Lane 1 was  
15 distilled water and used as control; lane 2 was *L. monocytogenes*; lane 3-10 represented *L. ivanuii*,  
16 *L. innocua*, *L. seeligeri*, *L. grayi*, *Listeria welshimeri*, *Salmonella choleraesuis*, *Salmonella typhimurium*,  
17 *Serratia marcescens*, respectively.  
18 172x31mm (72 x 72 DPI)

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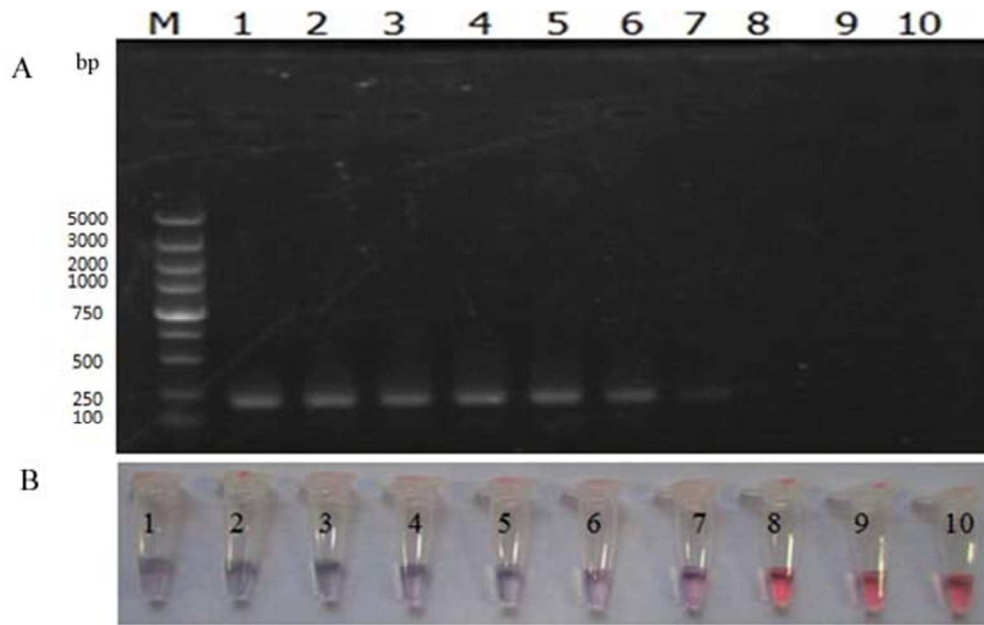


Fig.3 Sensitivity of visual detection of DNA of *L. monocytogenes*. (A) M: 5000 bp DNA ladder; lane 1: 1.304 ng/ $\mu$ l; lane 2 : 130.4 pg/ $\mu$ l; lane 3: 13.4 pg/ $\mu$ l; lane 4: 1.304 pg/ $\mu$ l; lane 5:130.4 fg/ $\mu$ l; lane 6: 13.4 fg/ $\mu$ l; lane 7: 1.304 fg/ $\mu$ l; lane 8: 0.1304 fg/ $\mu$ l; lane 9:0.01304 fg/ $\mu$ l; Lane 10: distilled water.  
207x128mm (72 x 72 DPI)