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Research Article

One-Pot Polymerase Chain Reaction with Gold Nanoparticles for Rapid and Ultrasensitive DNA Detection

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ABSTRACT

We report a novel method for rapid, colorimetric detection of a specific deoxyribonucleic acid (DNA) sequence by carrying out a polymerase chain reaction in the presence of gold nanoparticles functionalized with two primers. Extension of the primers when the target DNA is present as a template during the polymerase chain reaction process affords the complementary sequences on the gold nanoparticle surfaces and results in the formation of gold nanoparticle aggregates with a concomitant color change from red to pinkish/purple. This method provides a convenient and straightforward solution for ultrasensitive DNA detection without any further post-treatment of the polymerase chain reaction products being necessary, and is a promising tool for rapid disease diagnostics and gene sequencing.

KEYWORDS

deoxyribonucleic acid (DNA) detection, polymerase chain reaction (PCR), Au nanoparticle, colorimetric

1. Introduction

The area of molecular diagnostics is at the forefront of modern bioanalytical research. The ability to detect trace amounts of target deoxyribonucleic acid (DNA) sequences in a highly selective manner is of ever-growing interest for rapid detection of disease [1–11]. Herein, we report a method for rapid detection of DNA with ultrahigh sensitivity and specificity based on a novel strategy involving a combination of a polymerase chain reaction (PCR) and Au nanoparticles (Au NPs). Au NPs, with their extraordinarily high extinction coefficient, show controllable optical properties which are strongly dependent on the particle size and interparticle distance. The aggregation of

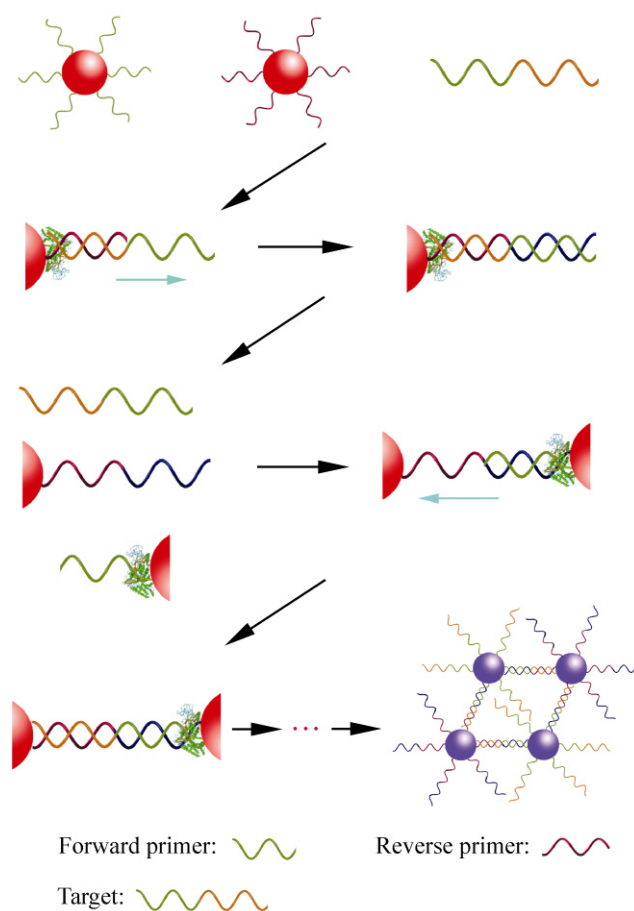
Au NPs causes a significant shift in the extinction spectrum as the solution color changes from red to purple [2, 8, 9]. The clearly distinguishable color change facilitates a very simple sensor readout which can be performed by the naked eye. The use of controllable aggregation of Au NPs was pioneered by Mirkin et al., harnessing DNA hybridization to induce an assembly of particles modified with single-stranded DNA (ssDNA) [12, 13]. PCR is a revolutionary technique in molecular biology and medical sciences, possessing both high sensitivity and high selectivity [14, 15]. The minimum amount of the target sequence required in PCR could be as few as 1000 copies, which is far below the detection limit of other technologies. The high selectivity of PCR can be realized by different

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design of forward and reverse primer sequences which result from different target sequences. However, post-treatment of the PCR products by processes such as gel electrophoresis is time-intensive, and does not allow the rapid detection of target DNA. Real-time PCR—as a more advanced technique—is used to amplify and simultaneously quantify a targeted DNA molecule, which enables both detection and quantification of one or more specific sequences in a DNA sample. However, for real-time PCR methods based on a fluorescent intercalator, dyes such as SYBR Green will bind to all double-stranded DNA (dsDNA) PCR products, including nonspecific PCR products. This will potentially interfere with, or prevent, accurate quantification of the intended target sequence [16].

Our concept for rapid and ultrasensitive detection of specific DNA sequences is based on amplified Au NP aggregation via PCR as shown in Scheme 1. In detail, 13 nm Au NPs are functionalized with two kinds of thiolated DNA sequences, forward primer and reverse primer. The forward primer is the same as part of the target DNA sequence and the reverse primer is complementary to the other part of the target DNA sequence (for sequence information, see the Experimental section). Note that the two primers have minimal base pairing between each other, so that no aggregation of Au NPs is observed before PCR reaction, as shown by the UV–visible absorbance peak at 519 nm. During the PCR process, extension of the reverse primer by rTaq DNA polymerase with the target DNA as a template will produce the complementary sequences on the Au NP surfaces, and result in the formation of a polymeric network of Au NPs. Aggregation of the Au NPs causes a red shift of the UV–visible absorbance, which enables ultrasensitive real-time DNA detection when UV–visible spectroscopy is employed during the PCR procedure. In addition, the concomitant color change from red to pinkish/purple can be observed by the naked eye. This method provides a convenient and straightforward solution for ultrasensitive DNA detection without any further post-treatment of the PCR products being required.



Scheme 1 Schematic illustration of the Au NPs involved in the PCR process

2. Experimental

2.1 Reagents and apparatus

Synthetic 5'-modified forward and reverse primers were obtained from Shanghai Sangon Co., Ltd. PCR kits were purchased from Takara Co., Ltd. Hydrogen tetrachloroaurate (HAuCl_4) and sodium citrate were purchased from Sigma–Aldrich and used without further purification. Ultrapure water (18.2 M Ω) was produced by Millipore Academic. UV–visible spectra were recorded on a PerkinElmer Lambda 25 UV–Vis Spectrometer. Transmission electronic microscopy (TEM) images were obtained on a Tecnai G2 F20 S-Twin TEM.

2.2 DNA primer design and synthesis

The primer sequences used in experiments were:

forward primer, 5'-SH-C6-GACAGGCCCGAAGGA ATAGA-3'; reverse primer, 5'-SH-C6-CTCTCTCTCC ACCTTCTTCT-3'; target: 5'-GACAGGCCCGAAGGA ATAG AAGAAGAAGGTGGAGAGAGAG-3'; random non-target sequence, 5'-CACGGTGCTCTGAGTTCA ACCTGGCCTACGAGCCGGATCA-3'.

2.3 Particle synthesis and functionalization

Au NPs with an average diameter of 13 nm were prepared by citrate reduction of HAuCl₄ according to the Ref. [17]. The NP size was confirmed by TEM. The concentration of Au NPs was determined by UV-visible absorbance at 519 nm (extinction coefficient = $3.64 \times 10^8 \text{ L}\cdot\text{mol}^{-1}\cdot\text{cm}^{-1}$). Functionalization of Au NPs with thiolated forward and reverse primers was performed by addition of DNA to the Au NPs solution. The mixture was then left at room temperature to incubate overnight. To remove any unbound DNA, the particles were centrifuged, and the pellet was collected and resuspended in ultrapure water. Centrifugation was repeated until all unbound DNA was removed from the mixture.

2.4 Primer extension via PCR

The PCR reaction was performed in a 25 μL mixture containing 1 \times Standard Taq Reaction Buffer (50 mmol/L KCl, 10 mmol/L Tris-HCl pH 9.0 at 25 $^\circ\text{C}$, and 1.5 mmol/L MgCl₂), 2 mmol/L each of four deoxynucleotidetriphosphates (dNTPs), 5 ng of target sequences, and 0.5 U of rTaq DNA polymerase. About 20 pmol each of the Au-primer conjugates instead of the pure primer were used in PCR. The PCR reaction was performed in a Bio-Rad PCR system (Bio-Rad, USA). The amplifying procedure of the PCR was: (1) 2 min at 94 $^\circ\text{C}$; (2) 5, 10, 15, 20, 25, and 30 cycles of 30 s at 94 $^\circ\text{C}$, 30 s at 55 $^\circ\text{C}$, and 30 s at 72 $^\circ\text{C}$; (3) 3 min at 72 $^\circ\text{C}$.

2.5 TEM imaging

Samples for TEM analysis were prepared by drying a drop of the PCR mixture on amorphous carbon-coated copper grids.

3. Results and discussion

As a proof-of-concept experiment, HIV gp140 exon

was selected as the detection target sequence for the sensor system. After 30 cycles of PCR, the UV-visible spectrum of the reaction mixture showed a red-shift from 519 nm to 526 nm (Fig. 1(a)). A clear color change from red to purple was also observed (Fig. 1(b)). The color change from red to purple indicates the association of Au NPs via hybridization of extended primer DNA. The aggregation of Au NPs was further confirmed by TEM. As shown in Fig. 1(c), massive agglomeration was observed after PCR, in contrast to the randomly dispersed NPs present before PCR.

The concentration of Mg²⁺ in the PCR mixture solution is crucial to our detection system. rTaq DNA polymerase cannot function without appropriate amounts of Mg²⁺ present in the reaction mixture. However, the presence of high concentrations of Mg²⁺ will induce the aggregation of Au NPs in the solution. Hence, the appropriate concentration of Mg²⁺ in the PCR reaction mixture was systematically determined using a series of Mg²⁺ concentrations (0, 0.3, 0.6, 0.9, 1.2, 1.5, 1.8, and 2 mmol/L). rTaq showed optimal enzyme activity with an Mg²⁺ concentration around 1.5 mmol/L. Usually, such a high concentration of Mg²⁺ will result in the agglomeration of Au NPs. Through a careful ligand-exchange process for Au NP surface functionalization [18–20], highly stable Au NPs were obtained with about 300 copies of DNA per Au NP, which are able to resist agglomeration in 1.5 mmol/L Mg²⁺. Meanwhile, 1.5 mmol/L Mg²⁺ is ideal for maximum rTaq enzyme activity.

To evaluate the effect of Mg²⁺ and rTaq on the Au NP behavior during PCR, control experiments were carefully performed to exclude the possibility of any influence of Mg²⁺ and rTaq on the PCR results. Without the target DNA in the PCR mixture but with the same concentrations of Mg²⁺ and rTaq in the reaction mixture, we did not observe the absorbance peak shift and color change of the reaction mixture after PCR.

The specificity of our system was studied with a non-target DNA. We harnessed a randomly designed sequence (5'-CACGGTGCTCTGAGTTCAACCTGG CCTACGAGCCGGATCA-3') for our purpose. After the same PCR procedure, no UV-visible absorbance peak shift was observed. Nor did we witness any color change in the PCR tube. However, as with most techniques involving PCR, the specificity of our system



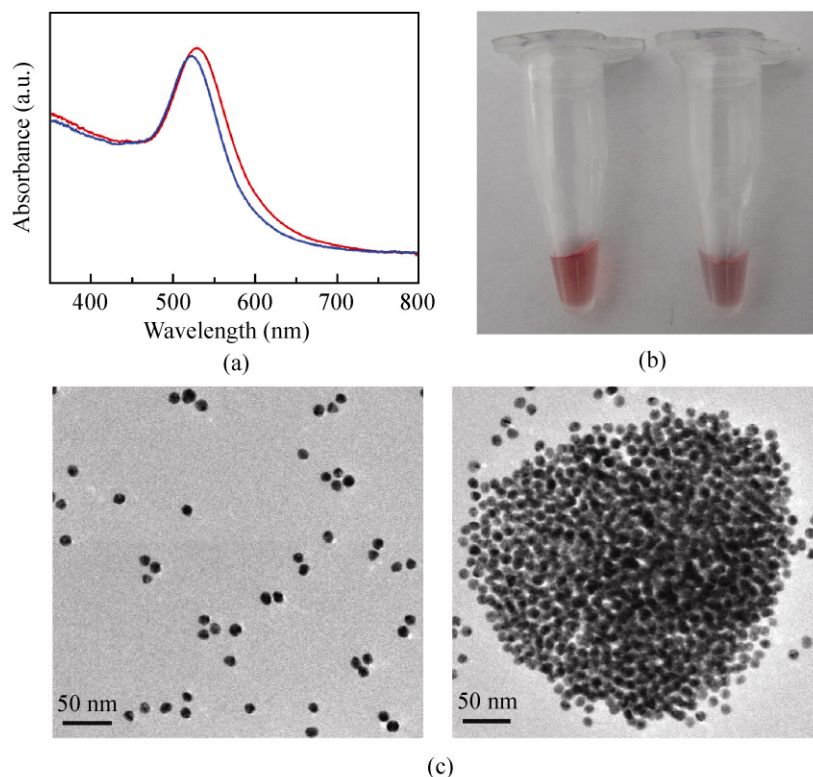


Figure 1 (a) UV–visible spectra of the PCR mixture before (blue curve, absorbance peak at 519 nm) and after (red curve, absorbance peak at 526 nm) PCR. (b) Photographs of the mixture before (left) and after (right) PCR. (c) TEM images of the mixture before (left) and after (right) PCR. Note that Au NPs are randomly distributed before PCR whereas a state of massive agglomeration is observed after PCR

relies largely on the PCR technique itself. To achieve a high specificity, careful design of the primer sequences is essential.

In a further set of experiments, we studied the correlation between the number of PCR cycles and the shift in the spectral maximum position in order to determine the shortest detection time. The degree of Au NP aggregation is expected to be influenced by the PCR products which serve as the linker molecules for forming the Au NP agglomeration. More cycles of the PCR procedure will lead to more PCR amplification which provides more linker molecules, and will result in a larger red-shift of UV–visible spectrum [21]. As expected, the spectral shift increased when the cycle number N was increased from 5 to 20 in our experiments (Fig. 2). As shown in Fig. 2, there is a 2.3 nm red-shift of the absorbance peak of the PCR solution when 5 cycles were run. Therefore if UV–visible spectroscopy is employed for real time detection, the detection can be finished after 5 cycles

of PCR, that is, in less than 15 min.

However, the spectral shift ceased to increase after 20 cycles. This may possibly indicate that PCR has reached full capacity after 20 cycles and no more

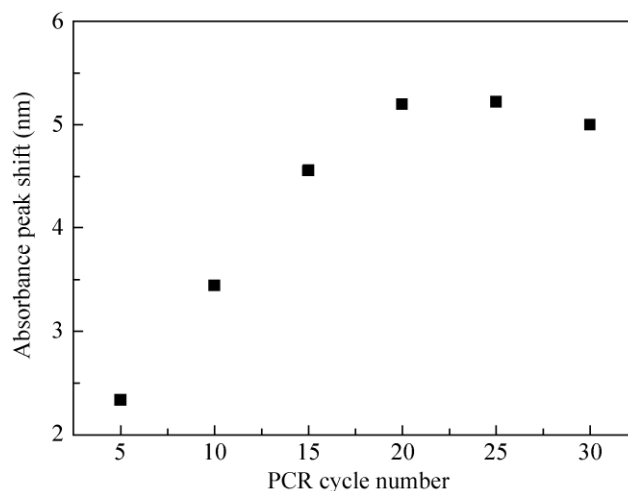


Figure 2 Correlation of the UV–visible absorbance peak shift with number of PCR cycles

linker molecules are produced after 20 cycles of PCR. It is also possible that more linker molecules are still produced after 20 cycles, but cease to contribute to the aggregation and therefore aggregation of the Au NPs reaches the maximum possible extent after 20 cycles.

It is important to point out that the UV–visible absorbance peak shift may possibly be affected by a number of other factors besides the number of PCR cycles.

The plasmonic interaction between Au NPs is largely dependent on the size and interparticle distance of Au NPs. For Au NPs of a fixed size, the length of target DNA sequences plays an important role in determining the shift in the position of the spectral maximum [22–23]. Use of longer target sequences will result in a larger distance between Au NPs and hence, weaker plasmonic interactions and a smaller spectral shift. However, if the target sequence selected is too short, it is not ideal for application in DNA detection because the steric hindrance of the 3 nm sized rTaq polymerase adsorbed on Au NPs makes it difficult for short DNA to hybridize. We speculate that the target sequence should be at least 20 nucleotides (nt) (~6.8 nm) in length to overcome the steric hindrance caused by rTaq polymerase. It turns out that the detection of a 40 nt target in our experiment is feasible.

We also investigated the sensitivity of our detection system. A series of solutions containing various concentrations of target DNA sequences (5000, 500, 50, and 5 pmol/L) were utilized in our experiments. The results are shown in Fig. 3. For the 5 pmol/L target DNA sequence test, we observed a 2 nm red-shift of the Au NP absorbance peak after 20 cycles of PCR. A larger spectral shift was observed with target concentrations above 50 pmol/L (3 nm for 50 pmol/L, 5 nm for 500 pmol/L, and 6 nm for 5000 pmol/L). Thus, the detection sensitivity of our system using 13 nm Au NPs is on the pmol/L level when using UV–visible spectroscopy. With the naked eye, we can detect target DNA concentrations of 500 pmol/L as the color of the Au NP solution changed from red to purple. That is, we can detect 0.1 femtomole of target DNA with the help of UV–visible spectroscopy and 10 femtomoles of target DNA by the naked eye. Considering the Au NP size effect, the sensitivity of our system could be

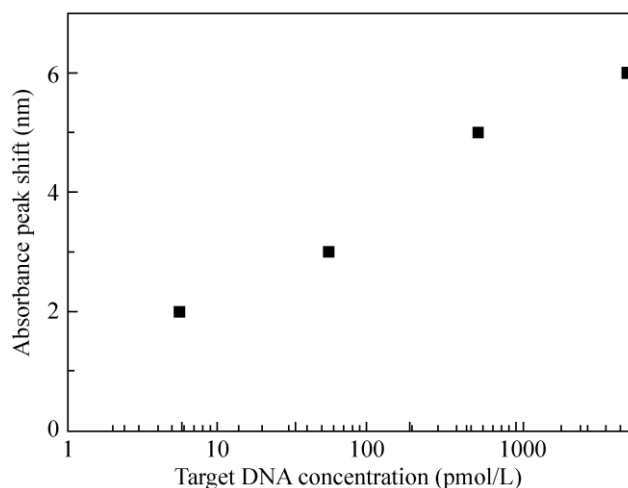


Figure 3 The correlation of UV–visible absorbance peak shift with initial target DNA concentration

further improved by employing larger Au NPs because the plasmonic interaction between larger Au NPs has been shown to be significantly stronger than for 13 nm Au NPs [24, 25].

So far, we have carefully studied the specificity, sensitivity and time response of our proposed Au NP/PCR system. The above results show that this novel method is rapid and ultrasensitive with high specificity for DNA detection. However, there is one concern that should be addressed: if we obtained the target DNA strands in the same concentration by PCR beforehand, what would happen when DNA primer-functionalized Au NPs were added? Would the same shift of the absorbance spectrum be observed? We carefully carried out the necessary control experiment. As shown in Fig. 4, there is a negligible shift in the absorbance spectrum (~1 nm) of the Au NPs solution before and after mixing with the same concentration of the target DNA strands; this is much smaller than that (6 nm) for the one-pot Au NP-involved PCR system. This demonstrates that the one-pot PCR with Au NPs is more sensitive in DNA detection than the two-step method.

4. Conclusions

We have demonstrated a novel method for rapid and ultrasensitive DNA detection. Adding primer-conjugated Au NPs into the PCR system gives a convenient and straightforward method for ultrasensitive



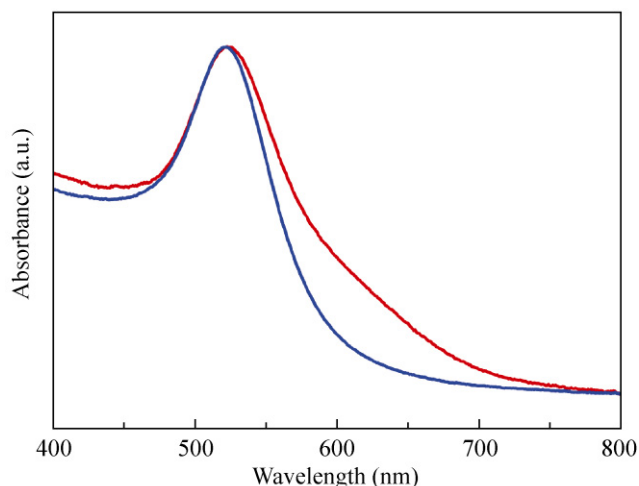


Figure 4 The UV–visible spectra of Au NPs without the PCR product (blue curve) and Au NPs with the PCR product (red curve)

DNA detection without any post-treatment of the PCR products being necessary. This rapid, ultrasensitive DNA detection method holds promise in disease diagnostics and gene sequencing.

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