



## Gold nanoparticle-assisted polymerase chain reaction: effects of surface ligands, nanoparticle shape and material†

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Recent reports have shown significant improvement of the polymerase chain reaction (PCR) efficiency and specificity with the addition of gold nanoparticles (AuNPs). However, the underlying mechanisms are still unclear and debatable. Here, we report effects of AuNP concentration, shape, material, and surface functionalization on PCR outcome under typical amplification conditions. All nanoparticles were tested with two PCR diagnostic models: (1) the nitrogen fixation (*NifD*) gene from *Azospirillum brasilense* Sp7 bacteria and (2) the polymorphic *ompA* gene encoding major outer membrane protein of *Chlamydia trachomatis*. The optimal concentration of 16 nm citrate-stabilized negatively charged spherical AuNPs (zeta potential  $-29.9$  mV) was determined to be 0.4 nM, in agreement with previous data for quite different PCR systems. Functionalization with poly(vinylpyrrolidone) (PVP) and thiolated PEG (mPEG-SH) essentially did not change the zeta-potential ( $-30$  mV and  $-21$  mV, respectively), but completely suppressed the PCR enhancement. Positively charged AuNPs with poly(diallyldimethylammonium) (PDDA) and cetyltrimethylammonium bromide (CTAB) capping polymers exhibited no specific enhancing effect on PCR or even inhibited the reaction at 5 pM of AuNP-PDDA. Also, no specific PCR enhancement was obtained with CTAB-stabilized gold nanorods. To elucidate the role of the particle material, we performed PCR with gold and silica particles of comparable size and equal negative charges. In contrast to the great PCR enhancement with 16 nm AuNPs, 20 nm silica nanoparticles did not affect the PCR efficiency or inhibit the reaction at concentrations higher than 100 nM. These findings suggest that the nature of the PCR system components, surface stabilizing agents and the thermal conductivity of the nanoparticle material can play important roles in the nanoparticle-mediated PCR.

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### 1. Introduction

The polymerase chain reaction (PCR) has become a standard biomolecular technique, which is commonly used for sensitive and rapid DNA detection. Exponential amplification of PCR allows replication of a single target copy to produce a large

amount of DNA copies.<sup>1–3</sup> However, the PCR can be error-prone and the specificity and efficiency of PCR can be unsatisfactory without proper optimization.<sup>4</sup> As a result of the poor purification of template DNA, primer-template mismatches, spontaneous formation of primer dimers, and non-optimized PCR-mixture ratio and thermocycling conditions, the formation of the target amplicon may be accompanied by non-specific side products called PCR artifacts. To fix this problem, several enhancers have been introduced: (1) various additives in PCR mixture, such as betaine,<sup>5</sup> dithiothreitol,<sup>6</sup> dimethyl sulfoxide,<sup>7</sup> single-stranded DNA-binding protein (SSB);<sup>8</sup> (2) instrumental design, including development of thermocyclers with precise heating/cooling rates;<sup>9,10</sup> (3) optimization of PCR system through optimal Mg<sup>2+</sup> concentration, cycle numbers and proper primer design;<sup>11–13</sup> (4) enzyme modification;<sup>14,15</sup> and (5) new touchdown<sup>16</sup> and nested<sup>17</sup> PCR strategies. Although these tools improve the PCR outcome, yet they are not all-purpose and the optimization protocol can be case dependent.

During the past decade, various nanomaterials have been used for PCR enhancement, including gold nanoparticles (AuNPs),<sup>18</sup> silver nanoparticles and graphene oxide,<sup>19</sup> reduced

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graphene oxide,<sup>19</sup> titanium dioxide and quantum dots,<sup>20</sup> upconversion nanoparticles,<sup>21</sup> fullerenes ( $C_{60}$ ),<sup>22</sup> carbon nanoparticles and nanotubes,<sup>23</sup> carbon nanopowder,<sup>24</sup> magnetic nanoparticles,<sup>25</sup> semiconductor nanomaterials,<sup>26,27</sup> dendrimers,<sup>28</sup> etc. Nevertheless, AuNPs are one of the promising candidates for use as PCR enhancers owing to their controlled geometrical parameters, reproducible synthesis and functionalization protocols. Recently, several hypothetic mechanisms of AuNP-assisted PCR were proposed to account for: (1) the SSB-like mechanism<sup>29</sup> and electrostatic interactions between AuNPs and PCR components;<sup>30</sup> (2) the high surface-to-volume ratio and non-specific adsorption of PCR components;<sup>28–31</sup> and (3) the thermal properties of AuNPs.<sup>32</sup> However, none of the suggested mechanisms can be considered convincing.

Here, we elucidated the role of the particle functionalization with neutral (PVP and mPEG-SH) and cationic (PDDA and CTAB) polymer ligands and role of the nanoparticle material by performing PCR with gold and silica particles of comparable size and charge, other PCR conditions being unchanged. In addition, the nanoparticle shape effect was examined with CTAB stabilized nanorods. All nanoparticles were tested with two PCR diagnostic models: (1) the nitrogen fixation (*NifD*) gene from *Azospirillum brasiliense* Sp7 bacteria; and (2) the polymorphic *ompA* gene encoding major outer membrane protein of *Chlamydia trachomatis*. At an optimal concentration of citrate-stabilized 16 nm negatively charged spherical AuNPs (0.4 nM), the PCR specificity and efficiency was greatly enhanced, whereas the negatively charged PVP- and PEG-stabilized AuNPs did not reveal any enhancing properties. Also, no specific PCR enhancement was observed with positively charged AuNPs and gold nanorods. By contrast to 16 nm negatively charged citrate-stabilized AuNPs, silica nanoparticles with the same charge and comparable 20 nm size have no impact on PCR efficiency. These observations show an important role of the nanoparticle material and surface modification in PCR enhancement, and could be helpful for further mechanistic studies.

## 2. Results and discussion

### 2.1 Characterization of nanoparticles

The following as-prepared and surface-functionalized AuNPs were used: negatively charged citrate-stabilized spherical gold nanoparticles (AuNP); nanoparticles capped with neutral

polymers PVP and mPEG-SH (AuNP-PVP and AuNP-PEG-SH, respectively) and cationic polymers CTAB and PDDA (AuNP-CTAB and AuNP-PDDA, respectively); gold nanorods (AuNR-662, 662 stands for the longitudinal plasmon resonance maximum wavelength) with aspect ratio of about 2.3; and spherical silica nanoparticles (SiO<sub>2</sub>NP). The acronym definitions and chemical structures of polymeric ligands are given in ESI file (Section S1).<sup>†</sup> Fig. 1 shows illustrative TEM images of citrate-stabilized AuNPs, gold nanorods, and silica nanospheres. Extinction spectra of nanoparticles are shown in Fig. S1 (ESI).<sup>†</sup> Table 1 summarizes the values of the zeta-potentials  $\zeta$  (mV), absorption maximum wavelengths of as-prepared NPs in aqueous solution (for nanorods, the longitudinal resonance is indicated); the average TEM and DLS sizes; the nanoparticle molar concentrations normalized to the extinction  $A_{\max} = 1$ ; and the optimal concentrations and maximal specificity and efficiency of different PCR additives used in our PCR systems.

It follows from Table 1 that the average TEM and DLS diameters of citrate NPs ( $16 \pm 0.8$  nm and  $17 \pm 1.2$  nm, respectively) have no significant difference within the standard deviation (SD) of about 1 nm. By contrast, after functionalization with PVP, PEG-SH, CTAB, and PDDA polymeric ligands, the hydrodynamic diameters increase significantly by 6.8, 10.4, 3.6, and 7.4 nm. We consider these data as strong evidence for successful attachment of ligands. The second evidence comes from drastic change in colloidal stability of NPs before and after functionalization. Fig. S2 (ESI)<sup>†</sup> shows extinction spectra and photos of citrate-stabilized and ligand-stabilized AuNPs before and after addition of 0.1 M NaCl salt. As expected, the addition of salt to the citrate-stabilized AuNPs results in immediate aggregation which is evident from changes in suspension color and extinction spectrum. By contrast, the addition of the same amount of salt to the ligand-stabilized AuNPs does not induce any aggregation phenomena accompanied by color and spectral changes. What is more, the ligand-stabilized NPs do not aggregate even in salt environment under strong temperature variations during PCR cycles (see below, Fig. 4).

### 2.2 Mechanisms of AuNP enhancing effect

Effect of nanoparticles on non-optimized PCR system was investigated by performing PCR with gold and silica particles of comparable size and charge, without changing the typical PCR

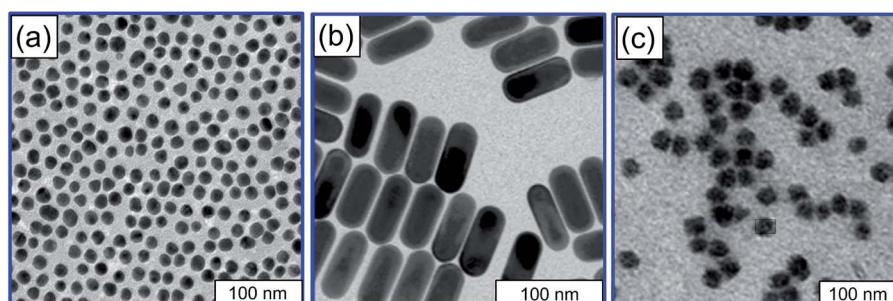


Fig. 1 Representative TEM images of as-prepared AuNPs (a), gold nanorods (b), and silica nanoparticles (c).



Table 1 Parameters of nanoparticles used as additives in PCR systems

Sample	Zeta potential (mV)	TEM size (nm)	DLS size (nm)	$\lambda_{\max}$ (nm)	Concentration (nM)	Optimal concentration	Maximal efficiency	Maximal specificity
AuNP	$-29.9 \pm 3$	$16 \pm 0.8$	$17.1 \pm 1.2$	520	4	0.4 nM	$4.25 \pm 0.33$	$5.7 \pm 0.43$
AuNP-PVP	$-30 \pm 3$	$16 \pm 1.0$	$22.8 \pm 1.8$	522	4	0.4 nM	$0.82 \pm 0.07$	$1.07 \pm 0.11$
AuNP-PEG-SH	$-20.5 \pm 3$	$16 \pm 1.0$	$26.4 \pm 1.3$	522	4	4 nM	$0.95 \pm 0.08$	$1.09 \pm 0.12$
AuNP-CTAB	$+10.2 \pm 1$	$25 \pm 1.2$	$28.6 \pm 2.3$	524	4	1 fM	$1.69 \pm 0.14$	$1.03 \pm 0.14$
AuNP-PDDA	$+30.5 \pm 2$	$16 \pm 1.0$	$23.4 \pm 1.7$	522	2.8	1 fM	$2.69 \pm 0.13$	$1.5 \pm 0.11$
AuNR-662	$+44.7 \pm 2$	$d = 25.1 \pm 1.5, L = 70.2 \pm 7.8$	$63.3 \pm 6.8$	657	0.26	1 fM	$0.9 \pm 0.07$	$1.07 \pm 0.13$
SiO <sub>2</sub> NP	$-31.2 \pm 3$	—	$20.0 \pm 2.5$	—	490	1 nM	$0.97 \pm 0.08$	$1.07 \pm 0.12$

conditions such as reagent concentrations and amplification profile.

To evaluate the overall ability of AuNPs to enhance PCR in terms of specificity and yield (efficiency), two PCR models were examined: (1) nitrogen fixation (*NifD*) gene from *A. brasiliense* Sp7 bacterial culture (*A. brasiliense* DNA detection, 710 bp amplicon), and (2) chromosomal *ompA* gene of *C. trachomatis* (157 to 160 bp target sequences for the variable domain 2 of MOMP). For brevity, we will refer to PCR models 1 and 2, respectively.

First, an optimal concentration of 16 nm citrate-stabilized AuNPs was found to be 0.4 nM by adding AuNPs in various concentrations to both PCR systems (Fig. S3, ESI†). This result is in good agreement with previous reports for quite different PCR models.<sup>29,32</sup> Fig. 2a illustrates the specificity of AuNP-assisted PCR performed with model 1. Numerous nonspecific bands (smears) are observed for duplicate lanes 1 and 2 without nanoparticles, performed by using DNA template mixed with

heterogeneous DNA fragments from similar microorganisms, whereas the addition of citrate-stabilized AuNPs at an optimal 0.4 nM concentration completely inhibits all non-target bands. Fig. 2b exemplifies the efficiency of AuNP-assisted PCR, when no significant bands can be observed without nanoparticles (duplicate lanes 1 and 2) in the case of a low DNA template concentration, performed with ten-fold diluted initial template DNA. It should be emphasized, that even for this unoptimized system, the addition of AuNPs at optimal 0.4 nM concentration results in a great PCR yield enhancement. To the best of our knowledge, this is the first demonstration of the yield-enhanced AuNP-PCR for *A. brasiliense* Sp7 (*NifD*) gene. Additionally, we have observed the enhancing effect of AuNPs on two-round error-prone PCR system by taking the PCR product as a template for the second round of PCR amplification (Fig. S5, ESI†). Moreover, we have shown that AuNPs could be used to improve PCR specificity while amplifying the long PCR product (Fig. S6, ESI†) which is in good agreement with a recent study by Zhou *et al.*<sup>33</sup> Finally, in the third test with the model 1 (Fig. 2c), the AuNPs were added after amplification step to verify whether the AuNPs affect the amplification step themselves (as in Fig. 2a and b) or the post-amplification PCR outcome, just before gel electrophoresis. As shown on Fig. 2c, there was no effect of AuNPs added after amplification step without nanoparticles and the smeared nonspecific bands look similarly for lanes 1 and 2. This gives strong evidence for the first assumption – citrate-stabilized AuNPs enhance the PCR outcome just during the amplification step.

One of hypothetical mechanisms of nano-PCR<sup>29</sup> is that AuNPs exhibit SSB-like behavior related to greater affinity of single-stranded DNA-binding protein for ssDNA compared to dsDNA. However, this explanation remains unanswered the following question: how many AuNPs per one PCR tube are needed to perform efficient PCR amplification caused by an SSB-like moiety alone? Below we provide simple estimations based on our experimental conditions. A typical PCR mixture, used in our experiments with optimal concentration 0.4 nM of AuNPs,<sup>29</sup> contained 2.5 U of Taq DNA-polymerase, equal to 10 ng or 3.5 nM of enzyme (the activity of 250 000 U Taq-DNA polymerase is equal to 4 ng or 43 fmol of 94 kDa  $M_w$  protein). A rough estimation shows that the molar ratio between Taq DNA-polymerase and AuNPs is about 1 : 10. The number of DNA templates, which exponentially grows in each of 35 PCR cycles,

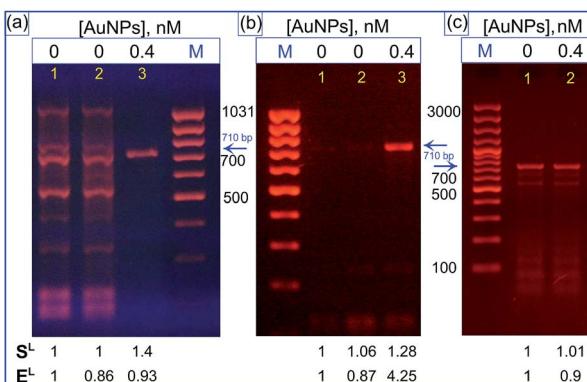
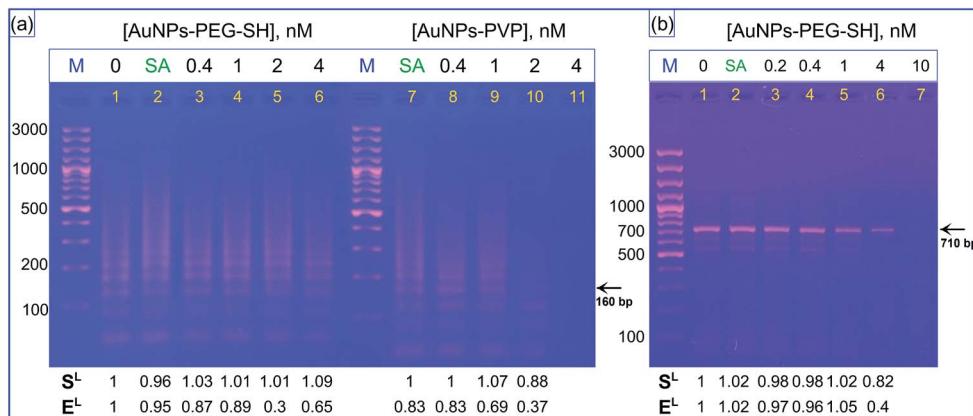


Fig. 2 Effect of citrate-stabilized AuNPs on PCR amplification of 710 bp region of *NifD* gene from *A. brasiliense* Sp7 (model 1). Lanes M stand for DNA markers. (a) Illustration of the AuNP-assisted PCR specificity. The duplicate smeared lanes 1 and 2 were obtained without AuNPs, lane 3 corresponds to addition of 0.4 nM AuNPs. (b) Illustration of the AuNP-assisted PCR high-yield efficiency. No significant bands were observed without AuNPs (duplicate lanes 1 and 2), whereas the addition of 0.4 nM AuNPs results in clear detection of the target 710 bp band. (c) Lanes 1 and 2 correspond to PCR without AuNPs and with AuNPs added after amplification step. The normalized quantities of PCR specificity ( $S^L$ ) and efficiency ( $E^L$ ) for each lane are indicated on the bottom panels.





**Fig. 3** (a). Effect of AuNPs capped with neutral polymers mPEG-SH and PVP on amplification of a 160 bp target from *C. trachomatis*. Symbols SA stand for addition of the capping polymers alone at a concentration corresponding to their concentration in 0.4 nM AuNP solution, symbol M designates DNA markers. No specific enhancement was obtained for both polymers. (b) Effect of AuNP-PEG-SH particles on amplification of a 710 bp target from *A. brasiliense* Sp7 (model 1). With an increase in nanoparticle concentration, a gradual inhibition of all bands is observed, up to visual disappearance of the target band at 10 nM concentration. The normalized quantities of PCR specificity ( $S^L$ ) and efficiency ( $E^L$ ) for each lane are indicated on the bottom panels.

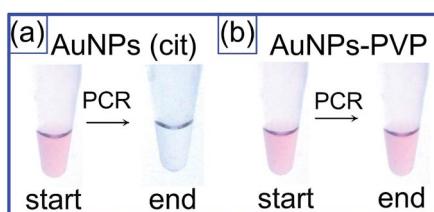
is at least one order greater than the number of DNA polymerase molecules per one PCR tube. This means that the molar ratio DNA : AuNPs is about 1 : 1, which is quite insufficient to ensure AuNP action as an SSB-like substances.<sup>34</sup> Consequently SSB-like behavior of gold nanoparticles seems to be not relevant to AuNP-assisted PCR enhancement.

To investigate a possible role of surface functionalization, we used AuNPs stabilized with two neutral polymer ligands such as PVP (AuNPs-PVP) and mPEG-SH (AuNPs-PEG-SH). According to Table 1 data, the both polymers do not change the sign and, in some extent, the value of nanoparticle zeta-potential. Surprisingly enough, the enhancing effect of AuNPs-PVP and AuNPs-PEG-SH on PCR was negligible (Fig. 3).

In particular, non-specific bands were present even at high concentrations of nanoparticles, and the PCR tubes remained pink colored after amplification (Fig. 4). In some cases, the addition of a high nanoparticle amount resulted in complete inhibition of PCR (see, *e.g.*, lanes 11 and 7 in Fig. 3a and b, respectively). We note that all these experiments included reagent controls with free polymers added to PCR tubes to verify the polymer effect alone (see lanes SA in Fig. 3). Thus, the functionalization of AuNP surface with neutral polymers, which does not change the initial negative charge of AuNPs, results in complete suppression of the PCR enhancement. This observation can be attributed to the steric hindrance for biomolecular

PCR components by protective PVP or mPEG-SH layers. In particular, these stabilizers prevent the nanoparticle aggregation after PCR amplification (Fig. 4b). By contrast, the stabilization of AuNPs with small citrate ions results in great enhancing PCR outcome (Fig. 2a and b), but does not prevent nanoparticle aggregation at the end of PCR (Fig. 4a).

The next set of experiments was aimed at understanding the role of AuNP functionalization with positively charged ligands. These experiments were motivated by a recent study,<sup>30</sup> revealing a crucial role of the particle charge in PCR enhancing. Specifically, the optimal enhancing concentration of positively-charged PDDA-capped AuNPs was shown<sup>30</sup> to be 1.54 pM, which is 260 times lower than the optimal 400 pM concentration of negatively charged citrate-stabilized AuNPs.<sup>29</sup> The goal of our experiment was twofold: (1) to verify the enhancing role of PDDA ligand for PCR models that differs from those investigated in ref. 30; (2) to examine the enhancing role of positive nanoparticle charge with CTAB ligand, which differs from PDDA. Clearly, the affirmative results for both points would expand the reported observations<sup>30</sup> and confirm a universal role of the positive AuNP charge in PCR enhancing. However, our experiments with models 1 and 2 gave negative answers for both questions. Fig. 5a demonstrates the presence of non-target bands for very small (0.001, 0.01 pM) and high (10, 100 pM) concentrations of positively charged CTAB-stabilized AuNPs. Furthermore, for the same experimental model, no significant enhancement of PCR outcome was observed with PDDA-capped AuNPs (Fig. 5b) at variation of nanoparticle concentration from 0.001 pM to 5 pM. Finally, when we used the positively charged CTAB-stabilized gold nanorods, the results were the same: no PCR enhancing or even complete inhibition. This illustrates insignificant role of the nanoparticle shape, at least for the PCR models examined here. Similar negative results were obtained for second PCR model, *i.e.* with nitrogen fixation (*NifD*) gene from *A. brasiliense* Sp7 (Fig. S4, ESI†). Thus, our experiments with two PCR models and with two cationic polymers did not



**Fig. 4** Photos of test-tubes with citrate- (a) and PVP-stabilized (b) AuNPs before and after PCR amplification.



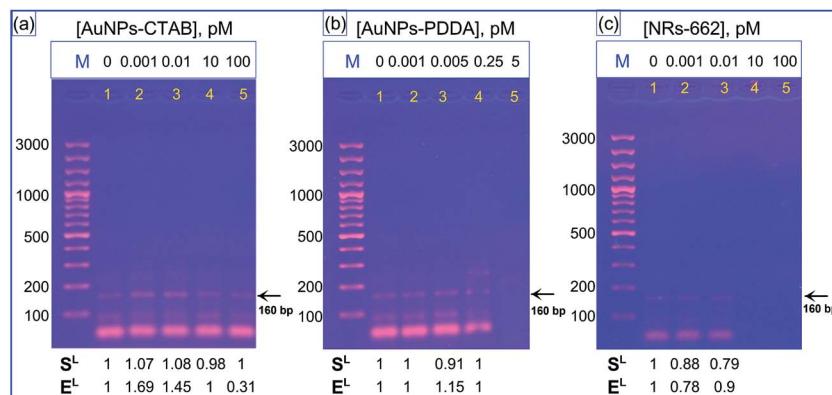


Fig. 5 Effect of AuNPs-CTAB (a), AuNPs-PDDA (b) and AuNRs-662 (c) on PCR amplification of a 160 bp target from model 2. The symbol M stands for DNA markers. Thus, the effect of ligand-capped AuNPs is not significant as compared with that for citrate-stabilized AuNPs. The normalized quantities of PCR specificity ( $S^L$ ) and efficiency ( $E^L$ ) for each lane are indicated on the bottom panels.

confirm any efficient PCR enhancing reported previously<sup>30</sup> for a different PCR model and positively charged AuNPs with PDDA ligand.

The precise reasons for observed discrepancy between our results and previously reported ones are unclear at present. Furthermore, the observed effects at trace 0.001 AuNP concentrations cannot be explained in terms of existing mechanistic models. In any case, we can conclude that the enhancing role of positively charged AuNPs is not ubiquitous. Instead, it can be ligand-dependent and PCR model-dependent.

Another proposed mechanism of AuNP-assisted PCR is related to high thermal conductivity of AuNPs as compared to water.<sup>32</sup> This can change the amplification conditions in local environment around gold nanoparticles, thus changing the

local conditions in “nanoreactor”. On the other hand, it has been supposed<sup>30</sup> that Au nanoparticles in PCR mixture act as small nanoreactors, in which the local concentration of primers and templates is increased owing to electrostatic interactions between AuNPs and PCR components. In this model, the particle charge is important, irrespective of the particle material. To verify these models, we fabricated silica nanospheres with the 20 nm size and  $-31.2$  mV zeta-potential close to those for citrate-stabilized AuNPs (16 nm and  $-29.9$  mV, respectively). If the nanoparticle charge plays a crucial role, one would expect a similar PCR enhancement for both particle types. If the material heat properties are important, one would expect quite different PCR outcomes. Fig. 6 shows that the second alternative agrees with experimental data. Indeed, we observed dramatic difference in PCR results for gold and silica nanoparticles with close parameters except for their heat properties. Of course, the chemical structure and physical properties of the nanoparticle surface should also be considered for ultimate conclusions.

### 3. Conclusion

In conclusion, we have studied the effects of AuNP concentration, charge, shape and material on the PCR efficiency and specificity. For two new PCR systems that were not studied previously, we have obtained high specific and efficient enhancement of target amplicons at optimal citrate-stabilized AuNP concentration about 0.4 nM. This observation agrees with previous reports for other PCR models and confirms a universal nature of underlying enhancement mechanisms at least for citrate-stabilized AuNPs.<sup>29,32-36</sup> Functionalization of citrate-stabilized AuNPs with neutral PVP and PEG-SH polymers results in elimination of enhancing properties, although the particle zeta-potentials remain the original sign and values. This phenomenon can be related to steric hindrance effects that prevent a proper interaction of PCR molecular components with nanoparticles. Functionalization of AuNPs with cationic polymers PDDA and CTAB leads to expected reverse of the particle charge sign. However, in contrast to previous report on positive

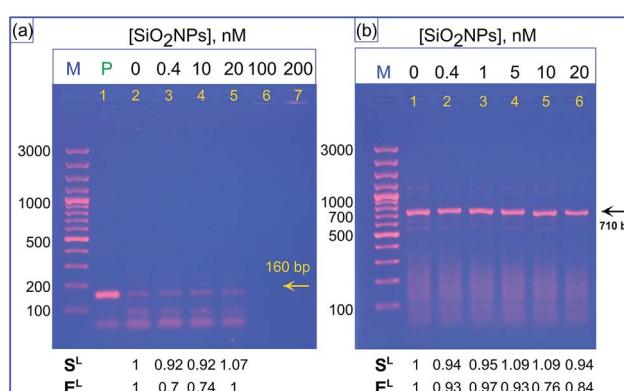


Fig. 6 (a) Effect of silica nanoparticles on amplification of a 160 bp target from PCR model 2. The symbol M stands for DNA markers, the symbol P designates a positive control with a purified DNA of genovar E as a template. No single distinct bands were observed at all nanoparticle concentrations from 0.4 to 200 nM (lanes 3–7). (b) Effect of silica nanoparticles on amplification of a 710 bp target from PCR model 1. For all nanoparticle concentrations from 0.4 to 20 nM (lanes 2–6), the PCR outcomes are similar to those obtained without nanoparticles (lane 1). Thus, for both PCR models, the effect of negatively charged silica nanoparticles is not significant as compared with that for citrate-stabilized AuNPs. The normalized quantities of PCR specificity ( $S^L$ ) and efficiency ( $E^L$ ) for each lane are indicated on the bottom panels.

enhancing properties of PDDA capped particles, we have obtained non-distinguishable enhancing effect. The particle shape plays negligible role as demonstrated by comparison of PCR outcomes with CTAB-coated gold nanospheres and nanorods. We have performed also PCR amplifications with two gold and silica particles of similar size and charge to show that the gold particles greatly enhance PCR results whereas silica counterparts do not. This implies a possible positive role of metallic heat properties in local environment of PCR nanoreactor. To summarize, further studies are needed to understand the underlying physicochemical mechanisms of nanoparticle-enhanced PCR, including the role of nanoparticle size, charge, surface functionalization, and the dynamic ratio between nanoparticle concentration and concentration of PCR system components.

## 4. Experimental section

### 4.1 Materials

All chemicals were obtained from commercial suppliers and used without further purification. Cetyltrimethylammonium bromide (CTAB, >98.0%), L-ascorbic acid (AA, >99.9%), sodium borohydride ( $\text{NaBH}_4$ , 99%), tetraethyl orthosilicate (TEOS), aqua ammonia (28–30%), L-arginine (98%), cyclohexane (anhydrous, 99.5%), thiolated poly(ethylene glycol) (mPEG-SH,  $M_w$  = 5000), poly(vinyl pyrrolidone) (PVP,  $M_w$  = 55 000), poly-diallyldimethylammonium chloride (PDDA, 20% solution) were purchased from Sigma-Aldrich. Hydrogen tetrachloroaurate trihydrate ( $\text{HAuCl}_4 \cdot 3\text{H}_2\text{O}$ ) and silver nitrate ( $\text{AgNO}_3$ , >99%) were procured from Alfa Aesar. Absolute ethanol (ET0016, 99.99%) was obtained from Scharlau Chemie. Sodium hydroxide ( $\text{NaOH}$ ), potassium carbonate ( $\text{K}_2\text{CO}_3$ ) and sodium chloride ( $\text{NaCl}$ ) were purchased from Reachim (Russia). Hydrochloric acid (HCl, 37 wt% in water) was purchased from Vekton (Russia). Tris, ( $\text{C}_4\text{H}_{11}\text{NO}_3$ ); ethylenediaminetetraacetic acid, disodium salt (EDTA); boric acid ( $\text{HBO}_3$ ); LE-2 agarose (Biotechnology Grade), ethidium bromide (EtBr), were purchased from Helicon (Russia). Phosphate buffer in tablets (PB; pH 7.4), 0.01 M was purchased from Biolot. MassRuler Low Range DNA Ladder, 6× loading dye, Taq DNA polymerase, dNTP mix were purchased from Fermentas, ThermoFisher Scientific. Ultrapure water was obtained from a Milli-Q Integral 5 system (Millipore) and used in all experiments.

### 4.2 Synthesis of nanoparticles

Spherical AuNPs with 16 nm average diameter were fabricated by reduction of  $\text{HAuCl}_4$  with sodium citrate as described by Grabar *et al.*<sup>37</sup> and were used as a template for subsequent capping with various surface ligands. Briefly, 25 mL of 38.8 mM sodium citrate was added quickly to boiled 250 mL of 1 mM water solution  $\text{HAuCl}_4$ , which resulted in a change in solution color from pale yellow to deep red.

To obtain polymer-capped AuNPs,<sup>38</sup> the as-prepared AuNPs were mixed with appropriate amount of each surface agent such as mPEG-SH, PVP or PDDA respectively, incubated for some time with subsequent centrifugation and resuspension in MQ

water. A detail description of ligand capping procedures are given in ESI file.<sup>†</sup>

Gold nanorods (AuNRs) with length and diameter about 70 nm and 25 nm, respectively, were fabricated according to Ratto *et al.*<sup>39</sup> with minor modifications.<sup>40</sup> First, gold seed particles were prepared by adding aqueous ice-cold sodium borohydride (10 mM, 0.1 mL) to a mixed aqueous solution of CTAB (0.1 M, 1 mL) and  $\text{HAuCl}_4$  (10 mM, 0.025 mL) and were vigorously stirred for about 2 minutes. After sequential addition of 28  $\mu\text{L}$  of 100 mM AA and 12  $\mu\text{L}$  of two-hours-aged gold seeds to a growth solution (5 mL of 0.1 M CTAB, 250  $\mu\text{L}$  of 0.01 M  $\text{HAuCl}_4$ , and 100  $\mu\text{L}$  of 4 mM  $\text{AgNO}_3$ ), the mixture was incubated for 24 h at 25 °C. Finally, 16  $\mu\text{L}$  of 10 mM ascorbic acid was added in three portions every 24 h. AuNRs were allowed to grow overnight without stirring at 30 °C. Then, as-prepared AuNRs were repeatedly centrifuged and redispersed overnight in 1 mM CTAB.

Highly monodisperse  $\text{SiO}_2$ NPs with 20 nm TEM average diameter were synthesized following the multistep seed-mediated growth technique in an aqueous solution of L-arginine.<sup>41</sup> Briefly, 9.1 mg of L-arginine was added to 6.9 mL of water in a standard 20 mL scintillation vial under magnetic stirring. Then, 0.45 mL of cyclohexane was accurately added to the top of the solution and the mixture was heated to 60 °C. Further, 0.55 mL of TEOS was added to the top layer of cyclohexane and the mixture was allowed to react for 20 h. A key tip for synthesizing high-quality monodisperse  $\text{SiO}_2$ NPs is to keep the cyclohexane and water parts unmixed, ensuring very slow addition of TEOS to the reaction mixture.

### 4.3 Characterization

Extinction spectra of as-prepared and ligand-capped nanoparticles solutions were measured with a Specord BS-250 and Specord S-300 spectrophotometers (Analytik, Jena, Germany). Transmission electron microscopy (TEM) images were obtained using Libra-120 transmission electron microscope (Carl Zeiss, Jena, Germany) at the Simbioz Center for the Collective Use of Research Equipment in the Field of Physical-Chemical Biology and Nanobiotechnology at the IBPPM RAS. The average diameters of as-prepared and polymer-capped AuNPs and silica nanoparticles and their zeta-potentials were measured with a Zetasizer Nano ZS device (Malvern, UK).

### 4.4 DNA samples

**Model 1.** *A. brasiliense* Sp7 genomic DNA. *Azospirillum brasiliense* strain Sp7 (IBPPM 150) from the Collection of Rhizosphere Microorganisms (IBPPM RAS, Saratov, Russia)<sup>42</sup> was used to prepare the DNA template. The target gene was a nitrogen fixation (*nifD*) gene for *Azospirillum* species (GenBank accession number: M64344 (ref. 43)). The experimental synthetic primers (Syntol, Russia) are listed in Table 2. Bacteria were cultured at 37 °C in a liquid malate-salt medium (MSM) as described previously.<sup>44</sup> The starting cell density of the 18 h cultures for DNA extraction was estimated by absorbance measurements on a Specord S-300 spectrophotometer at 660 nm. 1 mL of the bacterial suspension ( $2 \times 10^7$  cells per mL) after centrifugation



Table 2 Oligonucleotide primers used in PCR experiments

Primer name	Sequence (5'-3')	Length (bp)	Amplicon size (bp)	Ref.
<b>Primers used for <i>A. brasiliense</i> Sp7 DNA</b>				
nifD-up	GAGGCTATTGGCTATGACTG	21	710 (sample 1)	43
nifD-do	CGAGATCATGCCGTCGGC	20		
<b>Primers used for <i>C. trachomatis</i> DNA</b>				
momp-fw1	TTCAATTAGTTGGATTGTTGG	23	157–160 (sample 2)	45
momp-fw2	TCAACTTAGTGGCTTATTCGG	22		
momp-fw3	TCAATTAGTGGGGTTATTCGG	22		
momp-rv1	CACATTCCCAGAGAGCTGC	19		
momp-rv2	CACATTCCCACAAAGCTGC	19		
momp-rv3	CGGACTCCCACAAAGCTGC	19		
momp-rv4	GCACTCCCACAAAGCTGC	18		
F1	CGGTATTAGTATTGCGCTTG	23	1156 (sample 3)	46
B11	CGGAATTGTGCATTACGTGAG	22		

(Minispin, Eppendorf) at 5000g for 10 min was suspended in TE buffer (Tris HCl, 10 mM, EDTA, 1 mM). A genomic DNA purification kit (Thermo Scientific, Lithuania) was used for DNA extraction following the manufacturer's instruction. The kit is based on selective detergent-mediated DNA precipitation of crude lysate from different sample sources, including bacterial cells. The extracted DNA samples were further resuspended into 100  $\mu$ L of MQ water. The extracted DNA concentration and purity were estimated by UV-spectrophotometry on a Specord BS-250 spectrophotometer by  $A_{260}/A_{280}$  and  $A_{260}/A_{230}$  ratio, respectively, as indicative of nucleic acids purity ( $A_{260}/A_{280} \geq 1.8$  and  $A_{260}/A_{230} = 2$ ).<sup>4</sup> DNA quality was examined in 1% agarose gel electrophoresis. The DNA samples were stored at  $-20^{\circ}\text{C}$  until use in PCR.

**Model 2 and model 3.** *C. trachomatis* genomic DNA. The autoclaved *C. trachomatis* positive DNA from clinical samples (*C. trachomatis* DNA) were received from FSBSI Saratov SRVI, the Department Zoo- and Zoo-anthroponotic diseases and were used as DNA templates without further purification. The target was chromosomal *ompA* gene of *C. trachomatis* genovars (GenBank accession numbers: M58938, DQ064281, AF352789, AF063196, AY535104 (ref. 45 and 46)). The experimental synthetic primers (Syntol, Russia) are listed in Table 2. The DNA samples were stored at  $-20^{\circ}\text{C}$  until use in PCR.

#### 4.5 PCR and gel electrophoresis

All PCR reactions were conducted in triplicate with a T-100 Thermal Cycler (Bio-Rad, USA). PCR reagents were mixed to a final 30  $\mu$ L reaction volume in 200  $\mu$ L thin-walled tubes (Scientific Specialties, Inc., USA). Full PCR mixture in absence of nanoparticles was used as blank in all experiments. Pure double-distilled water (PCR-grade) as template was used for negative control in all experiments.

Each PCR tube with sample 1 DNA (*A. brasiliense* Sp7) as a template contained: 3  $\mu$ L 10 $\times$  PCR buffer, 0.6  $\mu$ L 10 mM dNTPs, 3  $\mu$ L 25 mM MgCl<sub>2</sub>, 0.5  $\mu$ L 5 U  $\mu$ L<sup>-1</sup> Taq DNA polymerase, 2  $\mu$ L template containing 120–200 ng DNA, 50 pmol of each primer (0.3  $\mu$ L nifD-up and 0.3  $\mu$ L nifD-do primer), and 15  $\mu$ L of appropriate NPs (varying in concentrations and particle

type) and filled up to the 30  $\mu$ L final volume with MQ water. The PCR procedure was as follows: 5 min at  $94^{\circ}\text{C}$  for pre-denaturation, followed by 35 cycles of 30 s at  $94^{\circ}\text{C}$ , 30 s at  $52^{\circ}\text{C}$ , and 30 s at  $72^{\circ}\text{C}$ . Then the cycling was terminated after 10 min final elongation step at  $72^{\circ}\text{C}$ .

Each PCR tube with sample 2 DNA (*C. trachomatis*) as a template contained: 3  $\mu$ L 10 $\times$  PCR buffer, 0.6  $\mu$ L 10 mM dNTPs, 3  $\mu$ L 25 mM MgCl<sub>2</sub>, 0.5  $\mu$ L 5 U  $\mu$ L<sup>-1</sup> Taq DNA polymerase, 2  $\mu$ L template containing 120–200 ng DNA, a multiplex-broad-spectrum PCR primer mix containing 1  $\mu$ L multiple forward primer (equimolar mix of 10 pmol momp-fw1, momp-fw2, momp-fw3) and 1.2  $\mu$ L multiple reverse primer (equimolar mix of 10 pmol momp-rv1, momp-rv2, momp-rv3, momp-rv4), and 15  $\mu$ L of appropriate NPs (varying in concentrations and particle type) and filled up to the 30  $\mu$ L final volume with MQ water. The PCR procedure was as follows: 2 min preheating step at  $94^{\circ}\text{C}$  followed by 38 cycles of amplification (30 s at  $94^{\circ}\text{C}$ , 30 s at  $55^{\circ}\text{C}$ , and 30 s at  $72^{\circ}\text{C}$ ) and a final 10 min elongation step at  $72^{\circ}\text{C}$ . Purified DNA of genovar E as a template was used as positive control in all experiments.

Each PCR tube with sample 3 DNA (*C. trachomatis*) as a template contained: 3  $\mu$ L 10 $\times$  PCR buffer, 0.6  $\mu$ L 10 mM dNTPs, 1.8  $\mu$ L 25 mM MgCl<sub>2</sub>, 0.15  $\mu$ L 5 U  $\mu$ L<sup>-1</sup> Taq DNA polymerase, 1  $\mu$ L template containing 120–200 ng DNA, 50 pmol of each primer (1  $\mu$ L F1 and 1  $\mu$ L B11), and 15  $\mu$ L of AuNPs in serial dilutions and filled up to the 30  $\mu$ L final volume with MQ water. The PCR procedure was as follows: 2 min preheating step at  $92^{\circ}\text{C}$  followed by 35 cycles of amplification (45 s at  $94^{\circ}\text{C}$ , 45 s at  $55^{\circ}\text{C}$ , and 60 s at  $72^{\circ}\text{C}$ ) and a final 10 min elongation step at  $72^{\circ}\text{C}$ . Purified DNA of genovar E as a template was used as positive control in all experiments.

After the PCR amplification, all the samples were stored at  $4^{\circ}\text{C}$  before gel-electrophoresis. The PCR products were analyzed by horizontal electrophoresis system (SE 1, Helicon) with a voltage set at 125 V for 65 min (Elf-4 power supplier, DNA-technology) with EtBr staining. Briefly, 3.5  $\mu$ L of each PCR product was mixed with 0.5  $\mu$ L 6 $\times$  loading dye prior to loading in the wells of an 1.5% agarose gel with EtBr containing in 1 $\times$  TBE buffer (10 mM Tris, 10 mM boric acid, 1 mM EDTA, pH 8.0).



The fluorescence of the bands was visualized by UV trans-illumination (312 nm, Vilber Lourmart), the gel electrophoreograms were obtained with a Canon 350D digital camera equipped with an orange filter to minimize the UV-lamp background lighting.

#### 4.6 PCR product quantification

The performance of PCR additives was quantified according to recent report<sup>30</sup> through calculations of two densitometric quantities, which are termed specificity and yield efficiency. The gel electrophoreogram images were analyzed with an open ImageJ software and the output data were represented as the means and standard deviations ( $n = 3$ ). A detail description of calculations is given in ESI file.<sup>†</sup> In brief, the ratio of the densitometric value of the specific target band to that of all bands amplified by PCR was defined as the specificity of amplification. By this definition, if there are no smears in PCR products, the maximal value of specificity equals 1. The ratio of the densitometric intensity of a specific target band to that for 500 bp of DNA marker was used as a measure of the yield efficiency. For brevity, we use the term "efficiency". If the obtained efficiency is  $\geq 1$ , the PCR additive is considered as high efficient for PCR optimization. In all figures, molecular weight markers (M) represented the same SM0383 (ThermoFisher Scientific). The optimal concentration of PCR additives was defined as a concentration that maximizes the brightest of specific target band.

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