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# Interaction between functionalized gold nanoparticles in physiological saline

Shada A Alsharif<sup>a</sup>, Liao Y Chen<sup>a,b</sup>, Alfredo Tlahuice-Flores<sup>a</sup>, Robert L Whetten<sup>a</sup>, and Miguel Jose Yacaman<sup>a</sup>

<sup>a</sup>Department of Physics and Astronomy, University of Texas at San Antonio, One UTSA Circle, San Antonio, TX 78249

# Abstract

Interactions between functionalized noble-metal particles in aqueous solution are central to applications relying on controlled equilibrium association. Herein we obtain the potentials of mean force (PMF) for pair-interactions between functionalized gold nanoparticles (AuNPs) in physiological saline, based upon > 1000-ns experiments in silico of all-atom model systems under equilibrium and nonequilibrium conditions. Four types of functionalization are built by coating the globular Au<sub>144</sub> cluster each with 60 thiolate groups: GS-AuNP (glutathionate), PhS-AuNP (thiophenol), CyS-AuNP (cysteinyl), and *p*-APhS-AuNP (para-aminothiophenol), which are, respectively, negatively charged, hydrophobic (neutral-nonpolar), hydrophilic (neutral-polar), and positively charged at neutral pH. The results confirm the behaviour expected of neutral (hydrophilic or hydrophobic) particles in dilute aqueous environment, but the PMF curves demonstrate that the charged AuNPs interact with one another in a unique way — mediated by  $H_2O$  molecules and electrolyte (Na<sup>+</sup>,Cl<sup>-</sup>) — in a physiological environment. In the case of two GS-AuNPs, the excess, neutralizing Na<sup>+</sup> ions form a mobile (or 'dynamic') cloud of enhanced concentration between like-charged GS-AuNPs, inducing a moderate attraction (~ 25-kT) between them. And, to a lesser degree, for a pair of p-APhS-AuNPs, the excess, neutralizing Cl<sup>-</sup> ions (less mobile than Na<sup>+</sup>) also form a cloud of higher concentration between the two like-charged *p*-APhS-AuNPs, inducing weaker yet significant attractions (~ 12-kT). In the combination of one GS- and one p-APhS-AuNP, the direct, attractive Coulombic force is completely screened out while the solvation effects give rise to moderate repulsion between the two unlike-charged AuNPs.

Due to their intermediacy in size, nanoparticles (NPs) possess unique mesoscopic properties that neither microscopic nor macroscopic bodies have. They are therefore widely investigated for various purposes. For examples, some NPs have been studied in the pursuit of nanomedicine.<sup>1–7</sup> In particular, gold nanoparticles (AuNPs) are being exploited for drug delivery and bioimaging,<sup>8–15</sup> which can involve solutions at high AuNP concentrations and controlled, reversible association of NPs.<sup>16</sup> In light of the current studies, it is of essential relevance to accurately determine the interactions between functionalized AuNPs in aqueous environment approximating physiological conditions. While *in vitro* and *in vivo* experiments

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<sup>&</sup>lt;sup>b</sup>Liao.Chen@utsa.edu

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have the final say, *in silico* studies, complementing *in vitro/vivo* experiments, can give atomistic insights of details that cannot be gained by *in vitro/vivo* means alone. For example, *in silico* experiments<sup>17–20</sup> have been employed to study the diffusion of functionalized AuNPs and to study how an AuNP interacts with biomolecules and cell membranes.<sup>21–28</sup> In this paper, we present *in silico* studies of AuNPs coated with hydrophobic, neutral hydrophilic, positively charged, and negatively charged organic/biological groups that lead to the determination of AuNP-AuNP interactions in physiological saline. The selected AuNP size is one that has been in continuous use in biological investigations over the past decade.<sup>29, 30</sup> Our main objective is to elucidate atomistic details of how functionalized AuNPs interact with one another under near-physiological conditions.

It is interesting to note that our *in silico* study of all-atom models concludes that likecharged AuNPs in physiological saline attract one another significantly over a distance a few times the NP diameter. This counter-intuitive behavior is in line with the long-studied attractions between like-charged colloids in electrolyte solutions.<sup>31–57</sup> However, oppositely charged colloids have been rarely studied.<sup>58</sup>

We study five atomistic model systems (see Table 1) each having a pair of AuNPs in a box of physiological saline (150-*mM* NaCl; 1-*bar*; 298-*K*). An example is shown in Fig. 1. A single globular AuNP comprises a compact 144-atom metal core protected by a surface layer of 60 thiolate groups (shown in Fig. 2) either the negatively charged tripeptide L-glutathionates (GS-); the hydrophobic thiophenols (PhS-), the hydrophilic (neutral but polar) amino-acid L-cysteinates (CyS-), or the positively charged aminothiophenols (*p*-APhS-). The ligand structures are illustrated in the supplemental information (Fig. S1). The atomic structure of the inorganic core, including the sulfur-gold bonding, was taken from Refs. 59–61. System I consists of two GS-AuNPs; System II, two PhS-AuNPs; System III; two CyS-AuNPs; System IV, two *p*-APhS-AuNPs; System V, one GS-AuNP and one *p*-APhS-AuNP. All share a common dimension of 2.0-nm diameter (enclosing the inorganic core Au<sub>144</sub>S<sub>60</sub>).

All the inter- and intra-molecular (other than Au-Au and S-Au) interactions were represented with the CHARMM36 force field,<sup>62, 63</sup> to which we add the following van der Waals (vdW) parameters for Au:  $\sigma = 1.66$  Å,  $\varepsilon = -0.10585kcal/mol$ . Water was represented with the TIP3P model.<sup>64</sup> Parameters for the interactions involving sulfur and gold atoms were taken from Ref. 65. The *in silico* experiments, namely, molecular dynamics (MD) simulations, were implemented using NAMD<sup>66</sup> as adapted for steered molecular dynamics (SMD).<sup>67, 68</sup> Periodic boundary conditions were applied in all dimensions. The cut-off distance applied to the vdW interactions is 1.2 nm, with a switching distance of 1.0 nm and a pair list distance of 1.4 nm. Particle-mesh Ewald was employed to compute the electrostatic interactions in an exact manner. Langevin MD was implemented with a time-step of 1.0 fs for short range interactions and 4.0 fs for long range interactions. The Langevin damping was chosen as 5.0/ps. The temperature was maintained at T=298-K and pressure at 1.0-bar.

The building procedure of System I is summarized as follows: (1) Starting from the widely accepted<sup>59–61</sup> structure of  $Au_{144}(SR)_{60}$  cluster, [R = H or CH<sub>3</sub>] we build a GS-AuNP by bonding one moiety (HS-less glutathione) to each of the 60 S atoms on the surface of the cluster. The mutual spatial arrangements of the 60 glutathionate groups were attained by

minimizing the interaction energy of the GS-AuNP while fixing the coordinates of all the Au and S atoms to their initial values. Then we equilibrate the GS-AuNP for 100-ns at 298

Au and S atoms to their initial values. Then we equilibrate the GS-AuNP for 100-ns at 298 K in vacuum, first with the Au and S constrained, and then for another 100-ns, without constraints on any of the atoms. It should be noted that the Au-S core was found to be stable, preserving the icosahedral symmetry, which indicates the validity of the interaction parameters we used. (2) We replicate the GS-AuNP and place two of them in a box of water. We then neutralize the system with 120 Na<sup>+</sup> ions and salinate the system with 150 mM NaCl. Thus we have System I. The procedures for building Systems II–V are essentially identical to those for System I and, therefore, not explicitly stated here. Some details are provided in the supplementary information (Figs. S2 and S3).

# Equilibrium molecular dynamics

Following initial optimizations, we executed 100-ns equilibrium MD runs for each system. During the last 50-ns of the MD run, we computed the root-mean-square-deviation of the 144-Au atoms from their initial structure (excluding the overall diffusion and rotation of the NP) to check the validity of the interaction parameters we used. Note that none of the atoms were fixed or constrained during the equilibrium MD run. All atoms were freely subject to the stochastic dynamics on the basis of atomistic interactions of the system modeled by the CHARMM36 force field.

# Nonequilibrium steered molecular dynamics

We followed the multi-sectional scheme of Ref. 70, conducting SMD runs for each of the Systems I–V in order to compute the PMF<sup>71–76</sup> as a function of the distance r between the centres-of-mass of the pairs of AuNPs. In each case, we steered (pulled) the pair of AuNPs toward each other at the same pulling speed  $v_d = 1.0 nm/ns$  in every case. The pulling was along the z-axis and over sections of 0.1-nm each in width. Namely, the center-of-mass (COM) coordinates of the innermost 12-Au core (three degrees of freedom) were controlled as functions (0, 0,  $z_i \pm v_d t$ ) of time t along a forward-reverse pulling path in the *i*<sup>th</sup> section while all other 0.57 million degrees of freedom were left freely determined by the stochastic dynamics of the system. Here  $z_i$  and  $z_{i+1}$  are, respectively, the z-coordinates of the two end points of the *i*<sup>th</sup> section over which we pulled the centre-of-mass of the inner Au core forward and backward to sample four (eight in some cases) forward-reverse pulling paths in each given section. The work done to the system, along these pulling paths (shown in supplementary Figs. S5 to S9), were used in the Brownian-dynamics fluctuation-dissipation theorem<sup>77</sup> to compute the free-energy difference (reversible work or PMF) as a function of the distance, i.e. the pairwise separation of AuNPs. Namely, the free-energy difference between two states:

$$G(z) - G(z_A) = -k_B T \ln \left( \frac{\langle \exp[-W_{A \to Z}/2k_B T] \rangle_F}{\langle \exp[-W_{Z \to A}/2k_B T] \rangle_R} \right). \quad (1)$$

Here  $W_{A\to B}$  is the work done to the system along a forward path when the NP pair was steered from A to Z.  $W_{Z\to A} = W_{B\to A} - W_{B\to Z}$  is the work Z for the part of a reverse path when the NP pair was pulled from Z to A. kB is the Boltzmann constant and T is the

absolute temperature.  $z_A$  and  $z_B$  are the z-coordinates of the COM of one of the two NPs of the pair at the end states A and B of the system, respectively. Note that the pulling was symmetrical between the two NPs of each pair.

Stability of the globular inorganic  $Au_{144}S_{60}$  core was clearly established in all five cases. None of the four different surfactants has a substantial effect on the arrangement, as the icosahedral symmetry is maintained, including the 30-fold Au-S-Au-S-Au staple geometry. Figure S4 shows quantitatively that the *rms*-deviation (from the initial Au-atom coordinates) remains within 0.02-nm, confirming the validity of the parameters used to characterize the Au-S and Au-Au interactions, which are not part of the standard CHARMM force field. Our main finding is that like-charged AuNPs in physiological saline do not repel but rather attract one another, via electrolyte ion-mediated attractions, starting from distances around a few times the AuNP diameter. Fig. 3 shows the PMF between two negatively charged GS-AuNPs (-60e charge each), and similarly the PMF between two positively charged *p*-APhS-AuNPs (+60e charge each). These PMF curves establish that the direct Coulombic repulsion between two like-charged AuNPs are very effectively screened by the physiological saline that provides 120 extra counter ions (120 Na<sup>+</sup> ions, for GS-AuNPs; 120 Cl<sup>-</sup> ions, for pAPhS-AuNPs). Some of the counter ions reside at the surface of these AuNPs, as illustrated in Fig. 4, reducing its effective charge to a small fraction of 60e. The free counterions (those not residing at the AuNP surfaces) gather in between the two like-charged AuNPs, as much as electrons gather in between the two protons of an  $H_2$  molecule (on the Å length-scale), mediating attraction between two like-charged AuNPs.

The origin of the difference between the GS-AuNP pair and the *p*-APhS-AuNP pair, as indicated in Fig. 4, is that the Cl<sup>-</sup> ions are less mobile than Na<sup>+</sup> ions and, therefore, are more effective in neutralizing the positively charged pAPhS-AuNP. This implies a reduced effective charge in *p*-APhS-AuNP, versus GS-AuNP. Consequently, the attraction between *p*-APhS-AuNPs is weaker than that between GS-AuNPs. Note that the distances corresponding to the minima of the PMFs for both like-charged particle pairs in Fig. 3 coincide with 2 times the distances corresponding to the maxima of the surrounding charge densities in Fig. 4.

By contrast, oppositely-charged AuNPs in physiological saline do not attract one another. The PMF for the unlike-charged pair of AuNPs is also shown in Fig. 3, which clearly indicates a small but significant effective repulsion between the two. In this case, the direct Coulombic attraction, between a positively charged *p*-APhS-AuNP and a negatively charged GS-AuNP (see supplementary Fig. S10), is totally screened out. Instead, the long-range interaction between them is repulsive due to the solvation effects. <sup>78</sup>

Fig. 3 further shows that the neutral hydrophilic AuNPs — the PMF is for a pair of such AuNPs, CyS-AuNPs — tend to separate from one another. This behaviour is easily rationalized in terms of the drive to maximize the total area exposed to aqueous solution.

Finally, the neutral hydrophobic AuNPs tends to coagulate together. Fig. 3 also shows the PMF for a pair of (neutral) thiophenol-coated AuNPs, indicating the long range and deep,

broad free-energy minimum associated with the classic hydrophobic effect, i.e. the reduction in the total (non-polar) surface-area exposed to aqueous solution.<sup>78</sup>

## Conclusions

Extensive atomistic *in silico* experiments (equilibrium MD and nonequilibrium SMD simulations), based on realistic molecular structures for all solution components, lead to the conclusion that the interactions between functionalized AuNPs in physiological saline are fundamentally distinct from the case of vacuum and other environments. On the length-scale of a few times the AuNP diameter, the direct Coulombic interactions are completely screened out. Like-charged AuNPs do not repel and opposite-charged AuNPs do not attract one another. The water molecules and the Na<sup>+</sup> and Cl<sup>-</sup> ions play essential roles to mediate effective attractions between a pair of like-charged AuNPs (positive or negative) as well as a pair of hydrophobic AuNPs. This fact also explains the experimental observation that Thiol passivated AuNPs are stable for long time in sharp contrast with nonpassivated particles. Solvation effects cause both the opposite-charged pair and the neutral-hydrophilic pair to dissociate. These predicted characteristics of various functionalized AuNPs are expected to impact further research of AuNPs in biological applications.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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#### Figure 1.

In silico box of physiological saline with two AuNPs each coated with 60 glutathiones. The system dimensions are 10nm×10nm×20nm, consisting of 191355 atoms. Waters are in licorice (sticks) and the NPs and ions in vdw (balls) representations respectively. All colored by element names (Na<sup>+</sup>, yellow, Cl<sup>-</sup>, grayish blue, O, red, C, cyan, N, blue, and H white). Au and S are not visible in this graph. All graphics in this paper were rendered with VMD<sup>69</sup>.



#### Figure 2.

144-AuNPs coated with 60 aminothiophenols (top left), glutathiones (top right), cysteines (bottom left), and thiophenols (bottom right). Gold and sulphur atoms are in VDW representation (large spheres) in gold and light yellow colors respectively. The ligands are in CPK (ball-and-stick) representations colored with element names (O, red, C, cyan, N, blue, and H white).





Reversible work for bringing two NPs together. Vertical axis: Potential of mean force. Horizontal axis: Distance between the centers of mass of the two NPs involved.



#### Figure 4.

The ion-charge density distributions around isolated AuNPs: the positively charged *p*-APhS-AuNP (blue, sign reversed) and the negatively charged GS-AuNP (red). Each curve includes contribution from all the  $Cl^-$  and  $Na^+$  ions.

#### Table 1

Details of systems considered in this study.

System	NP1 Charge	NP2 Charge	Interaction	Range of attraction
I	GS-AuNP -60e	GS-AuNP -60e	-25 kT	3.7 to 4 nm
II	PhS-AuNP 0e	PhS-AuNP 0e	-15 kT	2.7 to 4 nm
III	CyS-AuNP 0e	CyS-AuNP 0e	repulsive	
IV	p-APhSAuNP +60e	<i>p</i> -APhSAuNP +60e	-12 kT	3 to 3.7 nm
v	<i>p</i> -APhSAuNP +60e	GS-AuNP -60e	repulsive	