Supplementary information

# Electrostatic interaction driven gold nanoparticle

## assembly on three-dimensional triangular pyramid DNA

## nanostructures

Jaejung Song,<sup>a</sup> Sungmin Park,<sup>b</sup> Sehwan Kim,<sup>b</sup> Kyuhyun Im,<sup>c</sup> and Nokyoung Park<sup>\*b</sup>

<sup>a</sup> School of Interdisciplinary Bioscience and Bioengineering, Pohang University of Science &

Technology (POSTECH), San 31, Hyojadong, Namgu, Pohang 790-784, Republic of Korea.

<sup>b</sup> Department of Chemistry, Myongji University, Yongin, Gyeonggi-do 449-728, Republic of

Korea, E-mail: pospnk@mju.ac.kr; Tel: +82-31-330-6188, Fax: +82-31-335-7248.

<sup>c</sup> Samsung Advanced Institute of Technology, Samsung Electronics, Yongin, Gyeonggido 446-

712, South Korea

Scheme S1. Schematic drawing of X-shaped DNA (X-DNA) structures and oligonucleotide

sequences<sup>#</sup> of the X-DNA building blocks for the trigonal-pyramid DNA nanostructure.<sup>##</sup>



Building block	Strand	Segment 1	Segment 2
X	$\begin{array}{c} X_{11} \\ X_{12} \\ X_{13} \\ X_{14} \end{array}$	5'-(GT) <sub>20</sub> 5'-Phos-TGA CCC TT 5'-Phos-ATC GGT AT 5'-Phos-GCA TCT GG	CGA CCG ATG AAT AGC GGT CAG ATC CGT ACC TAC TCG CGA GTA GGT ACG GAT CTG CGT ATT GCG AAC GAC TCG CGA GTC GTT CGC AAT ACG GCT GTA CGT ATG GTC TCG CGA GAC CAT ACG TAC AGC ACC GCT ATT CAT CGG TCG
X <sub>2</sub>	X <sub>21</sub>	5'-(T) <sub>4C</sub>	CGA CCG ATG AAT AGC GGT CAG ATC CGT ACC TAC TCG
	X <sub>22</sub>	5'-Phos-TTG CAC CT	CGA GTA GGT ACG GAT CTG CGT ATT GCG AAC GAC TCG
	X <sub>23</sub>	5'-Phos-GGA TCA TC	CGA GTC GTT CGC AAT ACG GCT GTA CGT ATG GTC TCG
	X <sub>24</sub>	5'-Phos-AAG GGT CA	CGA GAC CAT ACG TAC AGC ACC GCT ATT CAT CGG TCG
X <sub>3</sub>	X <sub>31</sub>	5'-(AT) <sub>20</sub>	CGA CCG ATG AAT AGC GGT CAG ATC CGT ACC TAC TCG
	X <sub>32</sub>	5'-Phos-CTG TTG GA	CGA GTA GGT ACG GAT CTG CGT ATT GCG AAC GAC TCG
	X <sub>33</sub>	5'-Phos-ATA CCG AT	CGA GTC GTT CGC AAT ACG GCT GTA CGT ATG GTC TCG
	X <sub>34</sub>	5'-Phos-AGG TGC AA	CGA GAC CAT ACG TAC AGC ACC GCT ATT CAT CGG TCG
X <sub>4</sub>	X <sub>41</sub>	5'-(GA) <sub>20</sub>	CGA CCG ATG AAT AGC GGT CAG ATC CGT ACC TAC TCG
	X <sub>42</sub>	5'-Phos-CCA GAT GC	CGA GTA GGT ACG GAT CTG CGT ATT GCG AAC GAC TCG
	X <sub>43</sub>	5'-Phos-GAT GAT CC	CGA GTC GTT CGC AAT ACG GCT GTA CGT ATG GTC TCG
	X <sub>44</sub>	5'-Phos-TCC AAC AG	CGA GAC CAT ACG TAC AGC ACC GCT ATT CAT CGG TCG

<sup>#</sup>Same color sequences represents the complementary sequences between each other.

<sup>##</sup> 5'-Phos represents the phosphorylated 5' end of the oligonucleotide



**Figure s1.** Colloidal stability of TANAs at a time of day 1, week 1 and week 2 after mixing the DNA structures and AuNPs. (a) Absorption spectrum of TANA obtained by mixing TP-DNA and AuNP in a ratio of 1:1, (b) Absorption spectrum of TANA obtained by mixing TP-DNA and AuNP in a ratio of 1:2.

**Scheme S2.** Schematic drawing of hexa-branched DNAs which have three partially complementary sticky ends (red letters) and other three non-complementary free ends (black letters) and oligonucleotide sequences of the hexa-DNA building blocks for the DNA cubic.



Building block	Strand	Segment 1	Segment 2
H <sub>1</sub>	H <sub>11</sub>	5'-CTA ATC CGC ACA	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA
	H <sub>12</sub>	5'-TAT CGA CCA TGC	TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA
	H <sub>13</sub>	5'-ATG GCA ACT ATA	TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA
	H <sub>14</sub>	5'-Phos-GCT AGA GTC GTT	TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA
	H <sub>15</sub>	5'-Phos-ACC TTA GAC TCT	TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA
	H <sub>16</sub>	5'-Phos-TTC GTA TGG ACA	TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
H <sub>2</sub>	$\begin{array}{c} H_{21} \\ H_{22} \\ H_{23} \\ H_{24} \\ H_{25} \\ H_{26} \end{array}$	5'-TTC CTC ATG CGA 5'-AGC CTT GTA ACT 5'-Phos-AGA GTC TAA GGT 5'-Phos-GGT AAC ATT CCG 5'-CGG TTA CGA TTC 5'-Phos-ATT CGC TAC TCA	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
H3	H <sub>31</sub>	5'-GGT AAC TAC TGT	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA
	H <sub>32</sub>	5'-Phos-CGG AAT GTT ACC	TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA
	H <sub>33</sub>	5'-Phos-ATT GCC AGT TAG	TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA
	H <sub>34</sub>	5'-CTA TTC GGT CAA	TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA
	H <sub>35</sub>	5'-AAT TGC CTA GGT	TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA
	H <sub>36</sub>	5'-Phos-GTT AGC GAA ACT	TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
H <sub>4</sub>	H <sub>41</sub>	5'-CGG TTT CTA ACG	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA
	H <sub>42</sub>	5'-Phos-AAC GAC TCT AGC	TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA
	H <sub>43</sub>	5'-ATT GGC TAC TCA	TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA
	H <sub>44</sub>	5'-CCT AAG TTT CGA	TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA
	H <sub>45</sub>	5'-Phos-CTA ACT GGC AAT	TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA
	H <sub>46</sub>	5'-Phos-TTG TCC TAA TCG	TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
H₅	H <sub>51</sub>	5'-Phos-TGT CCA TAC GAA	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA
	H <sub>52</sub>	5'-ATT GAC CAG ACG	TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA
	H <sub>53</sub>	5'-AAG TGG ATT ACC	TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA
	H <sub>54</sub>	5'-Phos-GGT AAT CTC AAG	TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA
	H <sub>55</sub>	5'-Phos-TCT TGC ATG GCA	TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA
	H <sub>56</sub>	5'-AGT AAC TGC GTT	TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
H <sub>6</sub>	$\begin{array}{c} H_{61} \\ H_{62} \\ H_{63} \\ H_{64} \\ H_{65} \\ H_{66} \end{array}$	5'-Phos-TGA GTA GCG AAT 5'-CGT TGA ATC CTA 5'-Phos-TGC CAT GCA AGA 5'-Phos-ATT CGT AGA CTC 5'-GTT ACC ATA ACG 5'-TTT GCG ATC TAC	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
H <sub>7</sub>	H <sub>71</sub>	5'-Phos-AGT TTC GCT AAC	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA
	H <sub>72</sub>	5'-Phos-GAG TCT ACG AAT	TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA
	H <sub>73</sub>	5'-Phos-CCT ATG AAC GTT	TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA
	H <sub>74</sub>	5'-AAG GAT CGG TAT	TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA
	H <sub>75</sub>	5'-CTA AGG TTT CAC	TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA
	H <sub>76</sub>	5'-TTA GTT CCG ATC	TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA
Hs	H <sub>81</sub>	5'-Phos-CGA TTA GGA CAA	TCG CTG ACG TTG CAG ACA TCA CGT TGA CGC TGT CGA
	H <sub>82</sub>	5'-Phos-CTT GAG ATT ACC	TCG ACA GCG TCA ACG TGA AAC GTG AAG CGT CTG CGA
	H <sub>83</sub>	5'-AGC CAA ATG GTT	TCG CAG ACG CTT CAC GTT GAG CAC AGA CGT TGA CGA
	H <sub>84</sub>	5'-GCT TTC CAT AGA	TCG TCA ACG TCT GTG CTC GCA GCG TCT TAA CGT CGA
	H <sub>85</sub>	5'-Phos-AAC GTT CAT AGG	TCG ACG TTA AGA CGC TGC AGA CGT TCA GGA CTA CGA
	H <sub>86</sub>	5'-TTG ACC TAA GGA	TCG TAG TCC TGA ACG TCT TGT CTG CAA CGT CAG CGA

#### **Experimental details**

**General.** DNA materials were purchased from IDT and used without further purification. Water was triply distilled using a Millipore filtration system. Solutions of HAuCl<sub>4</sub> and NaBH4 were freshly prepared in distilled water.

**Synthesis of DNA nanostructures.** Triangular DNA structure has been synthesized by hybridization of four X-shaped DNA followed by ligation. X-shaped DNAs have been fabricated by mixing equimolar amount of four partially complementary oligonucleotide sequences dispersed in pH 8.0, 1xTE buffer(see Table in Scheme S1) followed by annealing step. The detailed procedure followed previous report.<sup>1</sup> Four X-shaped DNAs fabricated in separate tubes were mixed in equimolar amount. The mixture was incubated in 35°C for 10 minutes and the temperature has been decreased to 16°C by 1°C in stepwise for every 5 minute. After completing the annealing, the buffer has been exchanged to DI water using Amicon centrifugal filter (3kD Mw cutoff). The hybridized DNA structures have been tightened by ligating X-DNAs. The ligation has been performed by adding 3 units of T4 DNA ligase to 1 nmole of total X-DNA amount at 16°C for 8 hrs. The synthesized triangular DNA nanostructures have been concentrated using Amicon centrifugal filter and resuspended in DI water to have an appropriate concentration.

**Preparation of the citric acid-capped gold nanoparticle.** An aqueous solution of 99.999% hydrogen tetrachloroaurate hydrate (12.5 ml, 10 mM) was added to 250 ml D.I. water on a stirring hot plate. To the rapidly-stirred boiling solution, 7.5 ml of 50 mM aqueous solution of sodium citrate tribasic dihydrate was quickly added. The color of the solution changed from yellow to purple and finally became red within 5 min. The mixture was removed from heat

when the solution had turned deep red and then cooled for 30 min at room temperature. After cooling, the reaction solution was dialyzed using Amicon ultra 100 kDa Mw cutoff centrifugal filters for purification.

**Ligand exchange to the positive-charged gold nanoparticle.** AuNP surface was dispersed in aqueous solution that contains lipoic acid derivative molecules with quaternary ammonium, which was reported previously, to enhance the colloidal stability and change the surface charge of AuNPs.<sup>2</sup> Lipoic acid-derived quaternary ammonium ion (1 ml, 10 mM) solution was mixed to 1 ml of 100 nM AuNP solution and stirred at room temperature. After 10 h, the reaction solution was dialysed three times using Amicon ultra 100 kDa Mw cutoff centrifugal filters for purification.

**Assembly of AuNPs on a DNA pyramid.** The complex has been made by simple mixing and incubation of positive-charged AuNPs and triangular DNA pyramid in a specific molar ratio at room temperature. The assembled complex of AuNPs and DNA nanostructures have been confirmed by measuring the plasmonic absorption peak shift using Agilent Technologies 8453 UV-Vis Spectrophotometer G1103A.

**AFM imaging.** All the Atomic Force Microscope (AFM) experiments were performed by tapping-mode XE-100 Park AFM system (Korea) with a liquid cell at room temperature. Silicon cantilevers (NSC36, Ti / Pt, 1 N/m) were utilized for imaging. The sample (5  $\mu$ l) was adsorbed on a freshly cleaved mica plate pretreated with 1x Tris-acetate-EDTA (TAE) buffer for 5 min at room temperature and then washed three times with a buffer solution containing 1x TAE and 10mM MgCl<sub>2</sub>.

### Reference

1. N. Park, J. S. Kahn, E. J. Rice, M. R. Hartman, H. Funabashi, J. Xu, S. H. Um and D. Luo, *Nature protocols*, 2009, **4**, 1759-1770.

2. H. Jin, J. Nam, J. Park, S. Jung, K. Im, J. Hur, J. Park, J. Kim and S. Kim, *Chemical Communications*, 2011, **47**, 1758-1760.