Transfer of molecular recognition information from DNA nanostructures to gold nanoparticles

Thomas G.W. Edwardson, Kai Lin Lau, Danny Bousmail, Christopher J. Serpell and Hanadi F. Sleiman*

Corresponding author: hanadi.sleiman@mcgill.ca

Department of Chemistry and Centre for Self-assembled Chemical Structures, McGill University, 801 Sherbrooke Street West, Montreal, QC H3A 0B8, Canada

Abstract

DNA nanotechnology offers unparalleled precision and programmability for the bottom-up organisation of materials. The approach relies on pre-assembling a DNA scaffold, typically containing hundreds of different strands, and using it to position functional components. A particularly attractive strategy is to employ DNA nanostructures not as permanent scaffolds, but as transient, reusable templates to transfer essential information to other materials. To our knowledge, this approach, akin to top-down lithography, has not been examined. Herein, we report a molecular printing strategy, which chemically transfers a discrete pattern of DNA strands from a 3D-DNA structure to a gold nanoparticle. We show that the particles inherit the DNA sequence configuration encoded in the parent template with high-fidelity. This provides control over the number of DNA strands, their relative placement, directionality and sequence asymmetry. Importantly, the nanoparticles produced exhibit the site-specific addressability of DNA nanostructures, and are promising components for energy, information and biomedical applications.

Introduction

Since the power of DNA as a chaperone for the assembly of gold nanoparticles (AuNPs) was first exemplified, ^{1,2} it has been successfully applied to the 1D, 2D and 3D organization of AuNPs. ^{3,4} DNA-mediated AuNP assembly typically employs poly-functionalized nanoparticles, and has yielded a range of functional AuNP superlattices. ⁵⁻¹⁰ On the other hand, the construction of discrete nanoparticle assemblies of arbitrary geometry requires the use of DNA scaffolds to precisely position the nanoparticles. ¹¹⁻²⁰ Often, complex DNA origami scaffolds are used to organize a few nanoparticles. ²¹⁻²⁵ Importantly, the information is solely provided by the scaffold, while the nanoparticle is either isotropically or mono-functionalized. DNA nanostructures have previously been employed as templates for the growth of nanowires and inorganic surface patterning. In these instances the overall shape of the structures is transferred to the substrate but the complex DNA sequence information encoded in the template is lost. ²⁶

Recently, the ability to impart particles themselves with the geometries and valences that we take for granted in molecular self-assembly has been proposed as an alternative approach, ^{27,28} and it may open doors to higher levels of control over nanoparticle assembly with a minimum number of DNA strands. With this in mind, the creation of AuNPs which possess anisotropic DNA patterns has been highly sought after. ^{28,29} For example, controlled spacing between two or three DNA strands on the AuNP surface has been achieved using 1D DNA templates. ³⁰⁻³² Microsphere clustering with in situ polymerization has been used to create DNA patches on polymer colloids. ³³ Surface immobilization has also been used, creating Janus particles with localized patches of DNA strands. ³⁴⁻³⁷ The combination of surface techniques with a stepwise approach exploiting electrostatic repulsion has produced geometrically defined DNA-AuNPs with a controlled number of up to six strands. ³⁸ More recently, amphiphilic polymers have been used to allow regioselective DNA functionalization on gold nanoparticles. ³⁹ While these methods have produced

AuNPs with a significant measure of anisotropic regioselectivity, the ability to position many DNA strands with different sequences into geometrically controlled patterns has remained elusive.

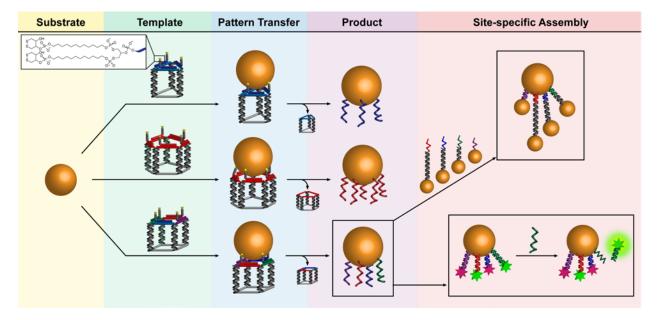


Figure 1 – An overview of the template-guided pattern transfer. An isotropic gold nanoparticle substrate is treated with a DNA nanostructure template, with controlled size, shape and sequence asymmetry (represented here as colors). The reactive arms presented by the DNA nanostructures bind covalently to gold nanoparticles, rendering a permanent pattern of DNA strands on the nanoparticle surface. The template structure is then removed, producing nanoparticles which have inherited molecular recognition information from the parent template. These information-rich nanoparticles are unique building blocks, which can be addressed site-specifically with different components, such as AuNPs with different spacer arms (top right) or different combinations of fluorescent probes (bottom right). They are also capable of dynamic assembly processes such as strand displacement (bottom right).

Herein, we describe a general method to transfer DNA strand motifs from a parent three-dimensional DNA template to gold nanoparticles, Figure 1. We show that the nanoparticle substrates acquire the DNA sequence information encoded in the parent template with high fidelity. Importantly, this produces nanoparticles which exhibit one of the defining properties of DNA nanostructures themselves: site-specific addressability. To our knowledge, this represents the first example of the direct 'printing' of a DNA pattern onto gold nanoparticles from a 3D nanoscaffold, providing arbitrary control over the number of strands, their relative placement, directionality and sequence asymmetry. This creates directionally functionalized AuNPs,²⁷ which are very promising components for the self-assembly of complex DNA programmed architectures. This also represents a first step in chemically copying the complex information contained in DNA nanostructures to another material, offering a potential approach to the problem of cost and scalability of DNA-based technologies. The process is related to the lithographic replication of electron beam patterns onto other materials, but here, the molecular information of the template is retained in the replication process.

Results and Discussion

The goal of our approach is to produce a robust patterning of DNA strands on the nanoparticle surface. This requires anchoring groups appended to the oligonucleotide which can bind strongly to the gold surface and remain so after downstream processing and manipulation of the samples.²⁹ With this in mind, a novel DNA-conjugate was synthesized,^{40,41} terminated with two cyclic disulfide moieties to allow AuNP-DNA conjugation (Supplementary S-IIIa-f).

The bis-disulfide-DNA (**Dx**, where x refers to the unique sequence) binds to AuNPs more efficiently and with faster binding kinetics than simple cyclic disulfide (**CD**) terminated DNA (Supplementary S-IIIh). The extending C12 portion in **Dx** may help to orient the disulfide moieties away from the highly charged DNA backbone, thus reducing electrostatic repulsion which is a key factor in the binding of DNA strands to gold nanoparticles. Furthemore, AuNPs functionalized with **Dx** were considerably more stable than the cyclic disulfide DNA (**CD**), as evidenced by a displacement assay using 1,4-dithiothreitol (DTT), see Supplementary S-IIIi. These binding and stability data suggested that the **Dx** conjugates are well-suited to use in the patterning experiments to follow.

The criteria we have set for the DNA nanostructure used as a template are as follows; it has the potential for 1) geometric variation, 2) introduction of sequence asymmetry, 3) positioning of different numbers of **Dx** and 4) can be removed easily after pattern transfer. We have shown previously that DNA minimal 'clip-by-clip' based structures can be efficiently used to site-specifically position DNA-amphiphiles and DNA-polymer conjugates. Here, the same strategy is employed to precisely arrange gold binding DNA conjugates for pattern transfer to a nanoparticle.

The first step in the patterning process is to prepare the template, in this case a 3D DNA scaffold decorated with **Dx** for transfer to the AuNP substrate. As such, the organization of different numbers of **Dx** on the DNA scaffold was carried out to produce a range of different letters for the printing process. (Figure 2a) The single-stranded regions of the cubic scaffold **Cb** are 20 nucleotides (nt) in length. Based on previous designs, only the central 14nt were used for the binding of the **Dx** which have a spacer of 5nt to orient them away from the crowded corners of the structure. This design also lowers the melting temperature of this region to facilitate removal of the template after pattern transfer. The structures shown in Figure 2b were assembled by one-pot thermal annealing of appropriate molar equivalents of

the component DNA strands in a buffer solution containing 100mM Na⁺. Quantitative yields of the desired products **Cb-Da**_x (x=1-4, cubes with 1-4 **Da** strands positioned on one face) eliminated the need for purification of the DNA nanostructure prior to AuNP patterning. (Figure 2b)

With the **Da** decorated cubic scaffolds **Cb-Da**_x in hand, we proceeded to investigate their ability to bind to AuNPs and transfer the desired number of **Da** strands. Based on geometry calculations (Supplementary S-Va) 10nm AuNPs (**NP**¹⁰) were used, as they represent the best size match for the cubic scaffold. Gold nanoparticles were incubated with **Cb-Da**_x at a 1:1 molar ratio for 16 hours at room temperature. The resulting hybrid DNA cage-nanoparticle assemblies, **NP**¹⁰-**Cb-Da**_x, (Figure 2c) were analyzed by agarose gel electrophoresis (AGE) with reference to a control AuNP sample. (Figure 2d)

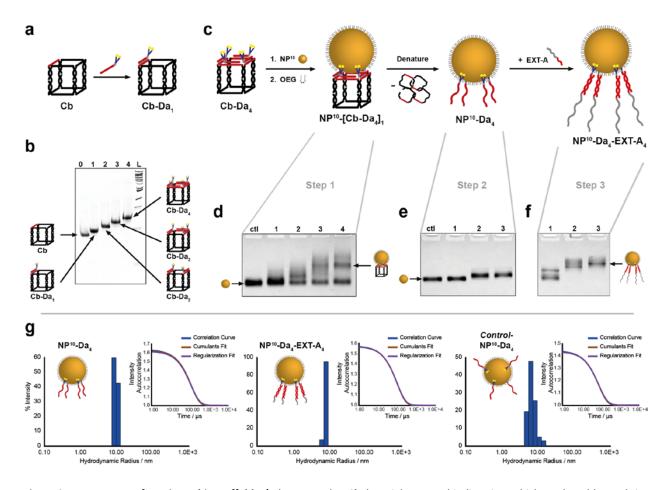


Figure 2 – Pattern transfer using cubic scaffold. a) The DNA cube, Cb, has eight ssDNA binding sites which can be addressed site specifically to organize Da molecules. b) Native Polyacrylamide Gel Electrophoresis (PAGE) analysis of Da addition to the cube scaffold: Lane 0, Cb; Lane 1, Cb-Da₁; Lane 2, Cb-Da₂; Lane 3, Cb-Da₃; Lane 4, Cb-Da₄; Lane L, 25-700bp DNA ladder c) Exemplary reaction scheme showing the binding of Cb-Da₄ to NP¹⁰ to produce the template bound NP¹⁰-[Cb-Da₄], which is treated with octaethylene glycol disulfide (OEG) for surface passivation before removal of the template by denaturing to produce NP¹⁰-Da₄, which can then hybridize to four EXT-A strands to produce the structure NP¹⁰-Da₄-EXT-A₄ (drawn to approximate scale, see Supplementary S-Va for geometry calculations). d) AGE analysis of products obtained at step 1 for the Cb-Da_x variants: Lane ctl, NP¹⁰; Lane 1, Cb-Da₁ + NP¹⁰; Lane 2, Cb-Da₂ + NP¹⁰; Lane 3, Cb-Da₃ + NP¹⁰; Lane 4, Cb-Da₄ + NP¹⁰. e) AGE analysis at step 2 for denatured product bands: Lane ctl, NP¹⁰; Lane 1, NP¹⁰-Da₂; Lane 2, NP¹⁰-Da₃; Lane 3, NP¹⁰-Da₄. f) AGE analysis at step 3 after addition of EXT-A for resolution of products: Lane 1, NP¹⁰-Da₂-EXT-A₂; Lane 2, NP¹⁰-Da₃-EXT-A₃; Lane 3, NP¹⁰-Da₄-EXT-A₄. g)

Dynamic light scattering (DLS) measurements of the templated product NP^{10} - Da_4 at steps 2 and 3 in comparison to a control sample in which four Da are positioned randomly on the nanoparticle reveals lower polydispersity, consistent with regionselectivity.

Electrophoretic mobility data for the reaction mixtures of cubes **Cb-Da**_x with 10-nm particles **NP**¹⁰ (*step* 1) are shown in Figure 2d. While the binding of **Cb-Da**₁ and **Cb-Da**₂ (cubes with one or two **Da** conjugates) to **NP**¹⁰ is inefficient, both **Cb-Da**₃ and **Cb-Da**₄ show a product exhibiting mobility consistent with the **NP**¹⁰-[**Cb-Da**_x] complex (see below for elucidation). Increasing conjugation yield is observed with increasing number of **Da** positioned on the cube, suggesting a cooperative effect due to the spatial organization on the template. (see below for the identity of the minor product of lower gel mobility in Figure 2d, lane 4) Control experiments using cubes without **Dx** strands revealed no interaction with AuNPs, confirming that binding was mediated by the **Dx** organized on the DNA cube (Supplementary Figure SF8).

To further stabilize the structures for downstream processing, passivation of the remaining surface of the AuNPs with a stable ligand was carried out. (Figure 2c) Carboxyl terminated octaethylene glycol disulfide (**OEG**) chains were used to cover the AuNP surface. Interestingly, after this stage the patterned AuNPs were found to be remarkably stable in sodium, and even in magnesium containing buffers. This passivation step could be carried out before or after removal of the template scaffold with no observable difference to the assemblies (see Supplementary Figure SF10). The samples were then run on AGE and the product bands excised and isolated by electroelution. The template scaffold was removed by disrupting all DNA hybridization under denaturing conditions (3M urea, 1xTBE), followed by centrifugation to isolate AuNPs. This process afforded purified NP¹¹O-Dax, see step 2 in Figure 2, with average isolated yields of 35% with respect to NP¹¹O. In principle, the DNA strand constituents of the cube may be isolated at this stage, and the cube re-assembled for repeated use as a transient template.

To determine the number of **Dx** strands transferred to the AuNP, and thus the fidelity of the patterning process, extension strands (**EXT-X**) were used to provide greater gel mobility differences, see *step 3* in Figure 2. Importantly, these also probe the addressability of the **Dx** strands bound to the AuNP surface. The **EXT-X** strands have a 17nt region of complementarity with each **Dx** and an extending single-stranded DNA tail of 43nt.

The gold nanoparticle containing 4 transferred DNA strands, NP¹⁰-Da₄ was incubated with an excess of EXT-A (Figure 2f, lane 3). The resulting sample exhibits decreased mobility, revealing that the Da strands transferred to the AuNP surface remain available for duplex formation and retain practical hybridization kinetics. The process was repeated for NP¹⁰-Da₃ and NP¹⁰-Da₂ in which the cube is bound to the nanoparticle via three or two DNA strands, and higher fidelity pattern transfer was observed with higher numbers of Da on the scaffold. In contrast, a non-templated control sample, prepared by incubating 4 molar equivalents of Da with 10nm AuNPs displayed different behavior. Hybridization of this structure to EXT-A showed a statistical distribution of products with between one and four Da strands bound to the AuNP (Supplementary Figure SF9). Dynamic light scattering (DLS) analysis of NP¹⁰-Da₄ showed a structure with less than 10% polydispersity, compared to the non-templated control with more than

20% polydispersity, corroborating the electrophoresis results and suggesting regioselective control via the cube template (see Supplementary S-Ve for further details).

DNA prismatic cages can be readily varied with respect to their geometry. Each geometry should transfer a different pattern of DNA strands to the nanoparticle, with different number and spacing of the **Dx** on the AuNP surface (Figure 3a). The preparation of Janus particles as a strategy to introduce anisotropy to spheres has been used to produce a range of asymmetrically functionalized AuNPs. ^{34,35,37,39,46} Although hemispherical separation can be obtained, the number and placement of DNA strands on the AuNP is not precisely controlled.

To investigate this geometric control, triangular (**TP**) and pentagonal prism (**PP**) templates were synthesized in the same manner as for the cube, Figure 3b. The scaffolds were decorated with **Da** strands to produce **TP-Da**₃ and **PP-Da**₅ which were incubated with 10nm AuNPs and then passivated with **OEG**. (Figure 3a)

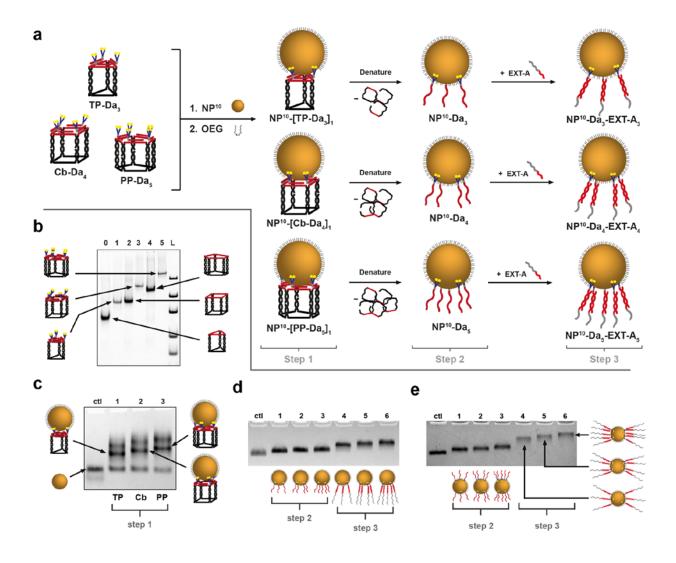


Figure 3 – Introduction of geometric diversity via template. a) General scheme for the patterning of AuNP using the DNA scaffolds TP-Da₃, Cb-Da₄ and PP-Da₅ to produce tri-, tetra- and pentavalent DNA-AuNPs NP¹⁰-Da₃, NP¹⁰-Da₄ and NP¹⁰-Da₅. b) Native Polyacrylamide Gel Electrophoresis (PAGE) analysis of Da decorated prisms: Lane 0, TP; Lane 1, TP-Da₃; Lane 2, Cb; Lane 3, Cb-Da₄; Lane 4, PP; Lane 5, PP-Da₅; Lane L, 75-300bp DNA ladder. c) AGE analysis of crude products at step 1: Lane ctl, NP¹⁰; Lane 1, TP-Da₃ + NP¹⁰; Lane 2, Cb-Da₄ + NP¹⁰; Lane 3, PP-Da₅ + NP¹⁰. The lower mobility band is assigned to a single gold nanoparticle bound to two DNA prisms, see Figure 3e. d) AGE analysis of purified patterned products NP¹⁰-Da₃, NP¹⁰-Da₄ and NP¹⁰-Da₅ and the addition of EXT-A strands: Lane ctl, NP¹⁰; Lane 1, NP¹⁰-Da₃; Lane 2, NP¹⁰-Da₄; Lane 3, NP¹⁰-Da₅.; Lane 4, NP¹⁰-Da₃-EXT-A₃; Lane 5, NP¹⁰-Da₄-EXT-A₄; Lane 6, NP¹⁰-Da₅-EXT-A₅. e) Products which are derived from the binding of two prisms to one AuNP, with and without EXT-A strands: Lane 0, NP¹⁰(DEG); Lane 1, NP¹⁰-Da₆; Lane 2, NP¹⁰-Da₈; Lane 3, NP¹⁰-Da₁₀; Lane 4, NP¹⁰-Da₆-EXT-A₆; Lane 5, NP¹⁰-Da₈-EXT-A₈; Lane 6, NP¹⁰-Da₁₀-EXT-A₁₀.

The prism-AuNP constructs were analyzed by AGE and a decrease in electrophoretic mobility was seen with increasing DNA scaffold size, consistent with the binding of the prisms to the AuNPs. In each case the second bands, highlighted in Figure 3c, were determined to be 1:1 complexes, e.g. NP^{10} -[TP-Da₃]₁. The band of lowest mobility was assigned as the 2:1 product, e.g. NP^{10} -[TP-Da₃]₂, based on gel mobility and titration assays (Figure 3e and Supplementary Figure SF11).

After purification and removal of the prism strands by denaturation, **EXT-A** was added to resolve the number of strands transferred to the AuNP. In Figure 3d, electrophoretic mobility patterns reveal that 3, 4 and 5 strands were robustly transferred by **TP-Da₃**, **C-Da₄** and **PP-Da₅**, respectively. Additionally, hydrodynamic radii (R_h) of the patterned products, determined by DLS, Supplementary S-Ve, revealed a steady increase with increasing number of strands on the nanoparticles, which is accentuated by the addition of extension strands **EXT-A**. Polydispersities were lower than non-templated gold nanoparticle conjugates (Supplementary Figures SF14-18 and Table ST5). These data, obtained for each step of the patterning process, demonstrate that well-defined numbers of DNA strands can be transferred to the AuNPs by geometric variation of the parent template.

Finally, we were interested in the potential of this strategy to transfer strands of different sequences to the nanoparticle. This confers particles with the ability for site-specific addressability and allows for anisotropic functionalization, thus producing building blocks with great potential for the self-assembly of advanced materials.^{27,28} A cubic scaffold with eight unique ssDNA binding sites **Cb-Da-d** was assembled and decorated with four unique gold binding conjugates; **Da**, **Db**, **Dc**, and **Dd**, on one face. (Figure 4a) Incubation with 10nm AuNPs followed by **OEG** surface passivation, isolation of the major product, **NP**¹⁰-**[Cb-Da-d]**₁, and subsequent removal of the cubic scaffold, produced sequence asymmetric **NP**¹⁰-**Da**₁-**Db**₁-**Dc**₁-**Dd**₁ (**NP**¹⁰-**Asym**). To confirm the correct four strand DNA sequence code had been transferred to the AuNP, stepwise addition of four unique **EXT** strands, complementary to **Da**, **Db**, **Dc** and **Dd** (**EXT-A** to **EXT-D**) was carried out as before. In Figure 4b, the stepwise binding and site selective addressability of **NP**¹⁰-**Asym** is shown. In lane 3 the addition of **EXT-A** produces a particle with one extension strand bound and **Db**, **Dc** and **Dd** available for further hybridization, **E1**. The subsequent addition of the remaining three extension strands reveals a ladder of decreasing mobility as each additional **EXT** binds to its specific site on **NP**¹⁰-**Asym**. This confirmed that the asymmetric pattern had been correctly transferred to the AuNP and the **Dx** are individually addressable due to their unique sequences.

A control sample of AuNP incubated with one molar equivalent of each **Dx** strand (**Da-Dd**) without organization on a template was prepared under the same conditions as the patterned sample. This should produce a statistical mixture of number and identity of **Dx** on the AuNP. Stepwise addition of the **EXT** strands **A-D** reveals this in Figure 4c. This highlights the control over information transfer possible through the use of the DNA nanoscaffold template.

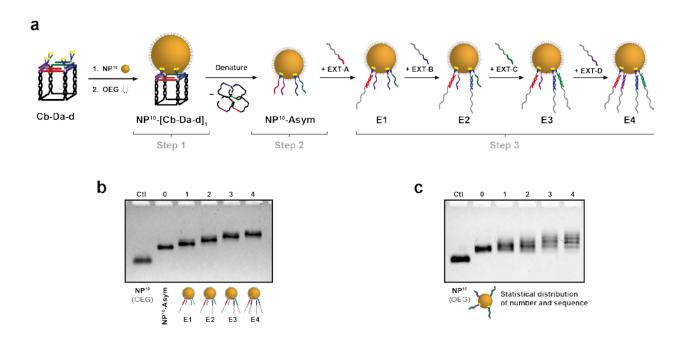


Figure 4 – Introduction of sequence asymmetry to patterning. a) General strategy for the production of sequence specifically patterned AuNPs using asymmetric DNA scaffolds Cb-Da-d to transfer one of each Dx strand to AuNP. After removal of cube template, each Dx strand on NP¹⁰-Asym can be targeted individually to produce E1-4. b) AGE analysis of purified NP¹⁰-Asym showing the site-specific binding of each unique EXT strand: Lane ctl, NP¹⁰; Lane 0, NP¹⁰-Da-d; Lane 1, E1; Lane 2, E2; Lane 3, E3.; Lane 4, E4 c) AGE analysis of template free control where one equivalent of each Da-Dd was incubated with NP¹⁰, the addition of specific EXT strands reveals the statistical distribution of number and sequence: Lane ctl, NP¹⁰(OEG); Lane 0, NP¹⁰-Asym; Lane 1, NP¹⁰-Asym + EXT-A; Lane 2, NP¹⁰-Asym + EXT-A/B; Lane 3, NP¹⁰-Asym + EXT-A/B/C; Lane 4, NP¹⁰-Asym + EXT-A/B/C/D.

To further characterize the asymmetrically functionalized nanoparticles, NP^{10} -Asym, transmission electron microscopy (TEM) was used. Firstly, to visualize the asymmetric pattern and regioselectivity by TEM, polyvalent 6nm AuNPs (NP^6) bearing complementary sequences (a'-d') to the patterned NP^{10} -Asym were used as site-selective probes. The parent particle NP^{10} -Asym was incubated with an excess of NP^6 -Dx', each with a sequence complementary to one of the strands on the nanoparticle (16 hours, room temperature, 1xTAMg) to produce the structures **T1-T4** shown in Figure 5a. To aid microscopy analysis, assemblies were purified by AGE prior to TEM analysis, see Supplementary S-Vf. TEM revealed that the desired satellite structures were successfully generated, with yields decreasing for the more

complex assemblies **T3** and **T4**. The side products observed are likely due to the ability of the polyvalent particles to act as crosslinks between **NP**¹⁰-**Asym** particles. Furthermore, inter-particle distances were consistent with the expected values for **T1** and **T2**, however **T3** and **T4** showed mostly collapsed structures, possibly due to surface drying effects and Mg²⁺ mediated aggregation between the particles in such close proximity.

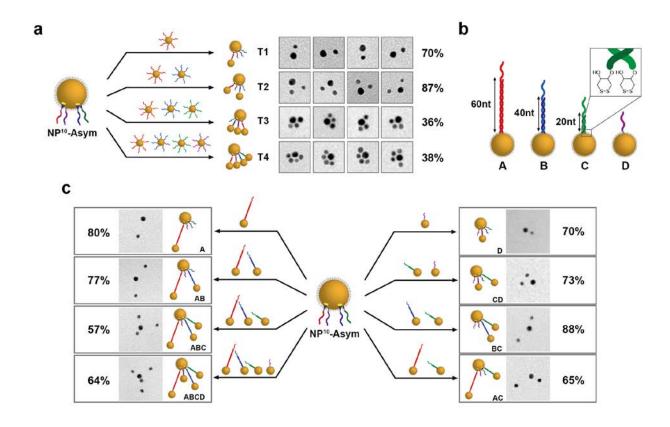


Figure 5 – Microscopy analysis of patterned AuNP self-assembly. a) TEM images show the site-specific addressability of the structure NP¹⁰-Asym by hybridization of polyvalent NP⁶ (superscripts 10 and 6 are the nanoparticle diameters 10 and 6 nm), labelled with complementary strands Da', Db', Dc' or Dd', the structure of these satellite structures shows the anisotropic nature of the patterning (for additional TEM images and statistics see supplementary S-Vg). b) Four different dsDNA arm lengths were created to allow each unique sequence (A-D) to be visualized via TEM. In each case the ssDNA binding portion is 17nt with sequences A, B and C having dsDNA spacers of 20, 40 and 60bp respectively. D has only the ssDNA 17nt binding region with no dsDNA portion. c) A range of anisotropic satellite structures can be accessed from the patterned particle NP¹⁰-Asym and four unique monofunctionalized NP⁵ with differing arm lengths A-D. This produces inter-particle separations of 30.4, 23.6, 16.8 and 10.0nm for A, B, C and D respectively. Additional experimental details and statistical analysis can be found in Supplementary S-Vf-h.

Although these assemblies highlight the sequence-specific addressability of the parent particle, **NP**¹⁰-**Asym**, inherited from the asymmetric cube template, an improved design was developed to improve microscopy characterization. This uses rigid dsDNA arms of differing lengths on monofunctionalized nanoparticles to probe each addressable site on the **NP**¹⁰-**Asym**. Complementary sequences to the four

strands on the nanoparticles had different arm lengths, denoted **A-D** in Figure 5b. These should produce inter-particle distances of 30.4, 23.6, 16.8 and 10.0nm. Assemblies were carried out with an excess of the 5nm monoconjugates **A-D**, and purified by AGE for TEM. Figure 5c shows representative images of some of the anisotropic satellite structures, which can be obtained from **NP**¹⁰-**Asym** and the four unique arm lengths. These assemblies exhibited less collapsed structures and misassembled products than the structures **T1-T4**. This is consistent with the longer, more rigid, dsDNA arms distancing the particles from each other, and the DNA mono-conjugated nanoparticles. Additionally, inter-particle distances were measured for each of the assemblies and were found to correlate well with the expected values. A full statistical overview can be found in the Supplementary information, S-Vh. Overall, this method represents a simple, modular strategy to give access to many unique architectures using a small number of building blocks. More importantly, these experiments also reveal the potential of the information-rich patterned particles, **NP**¹⁰-**Asym**, to undergo selective DNA-mediated self-assembly, showcasing their viability for the production of unique DNA-AuNP assemblies.

To further investigate the sequence specificity imparted to the **NP**¹⁰-**Asym** particles a fluorescence assay was employed. A 'barcoded' particle with four different dye-labelled DNA strands was generated. Two fluorophores were used; 6-carboxyfluorescein (6-FAM) and cyanine 3 (Cy3), which have discernible excitation/emission properties. We synthesized four 5' dye- labelled DNA sequences a' – d'; **Cy3-a'**, **Cy3-b'**, **6-FAM-c'** and **6-FAM-d'**. Incubation of 1.5 molar equivalents of fluorophore to each unique binding site on **NP**¹⁰-**Asym** was carried out at room temperature for 12 hours. This was followed by three centrifugation and washing cycles, under native conditions, to remove any traces of unbound dyes.

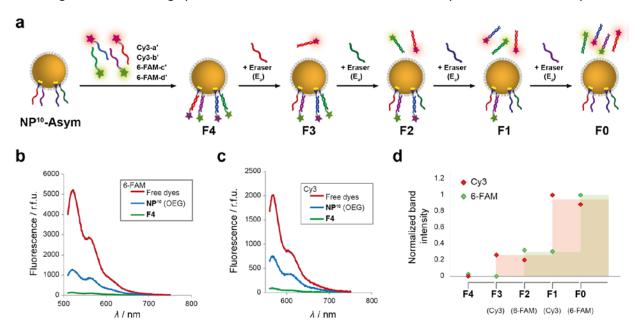


Figure 6 – Fluorescence investigation of asymmetric patterning. a) Scheme showing the preparation of a fluorophore 'barcode' on the sequence asymmetric NP¹⁰-Asym using Cy3 and 6-FAM labelled oligonucleotides. b) Fluorescence emission spectra of 6-FAM free in solution, in the presence of NP¹⁰(OEG) and when bound to the AuNP in structure F4 (Excitation; 490nm) c) Spectra for Cy3 free in solution, in the presence of NP¹⁰(OEG) and when bound to the AuNP in structure F4 (Excitation; 545nm) d)

Histogram produced from AGE analysis of sequential strand displacement of dyes from **F4** to finally produce **F0.** The order of eraser strands added was: E_a , E_c , E_b then E_d . This corresponds to alternating removal of the two dyes starting with Cy3. The resulting increase in fluorescence at each step was seen to be dye specific, through the sequence specific eraser strands. (refer to Supplementary S-Vi for additional details).

Steady-state fluorescence spectra of the nanoparticle bearing all four dye-labelled DNAs, **F4** in Figure 6a, were measured, and compared to control samples with the same concentration of free dyes in solution, and in the presence of unbound AuNP, **NP**¹⁰(**OEG**). Firstly, a decrease in fluorescence is seen in the presence of **NP**¹⁰(**OEG**) compared to the free dyes, suggesting some non-specific binding. For the sample **F4**, a large decrease in emission is observed when the dye-labelled DNAs are bound to the nanoparticle by hybridization to their surface bound complements. The fluorescence quenching observed is likely caused by both the AuNP as well as the close proximity of the dyes to each other.⁴⁷

The efficient fluorescence quenching of the AuNP bound dyes allowed a displacement assay to be monitored directly. Four specific eraser strands, E_a to E_d , were designed to remove the dye-tagged strands one-by-one from F4 by strand displacement, utilizing a 10nt toehold region for specificity and rapidity. Figure 6d shows the cumulative addition of the different eraser strands, which will release the dyes from the AuNP sequentially, giving an increase in fluorescence emission of a specific dye at each step. For example, E_a is added to E4 to liberate E4 and produce the structure E4 at this step an increase in E4 fluorescence is observed with negligible change in the 6-FAM fluorescence. The following displacement steps were carried out by alternating between the two dyes, in each case an increase in fluorescence for the specific displaced dye was seen. These results further support that four DNA strands with unique sequences have been transferred to the AuNP by the patterning method and present a platform with unique properties well-suited to biosensing. In particular, this method can result in a very large and diverse set of 'barcoded' gold nanoparticles from a minimum number of dyes of different colours, allowing efficient multiplexing and direct detection in bioanalyte sensing platforms.

Conclusion

In summary, we have developed a method to encode gold nanoparticles with complex DNA strand patterns comprising number, geometry and positioning of different unique strands. The use of DNA nanostructures as transient templates for the creation of patterned AuNPs is a general strategy which could be applied to a variety of inorganic nanoparticles. For example, the application of our approach to non-spherical particles could be used as a strategy for furthering complexity. This method also presents the advantage of potential recycling of the template or solid-support immobilization for high throughput, scalable generation of patterned structures. From a DNA nanotechnology standpoint this represents an interesting avenue to explore for the role of DNA scaffolds in nanofabrication, as a potential way to reduce costs by using DNA nanostructures in a transitory fashion. As well as allowing the transfer of different geometrical patterns to the AuNP, it is possible that the size ratio between the different DNA scaffolds and the AuNP could be used to further control the surface spacing and angle between DNA strands. In principle, the transfer should produce a chiral arrangement of the DNA strands

on the gold nanoparticles, useful for the development of chiral plasmonic structures; we are currently working to test this stereospecificity.

As well as applications in colloidal assembly and biocomputing,⁴⁹ we predict that this method could also be applied to surfaces to guide nanoelectronic or photonic circuitry or create robust anchors for positioning other functional components such as aptamers and enzymes. It is becoming well understood that surface structure and ligand placement are key factors to the cellular uptake of nanoparticles *in vitro* and their fate *in vivo*. This patterning creates a platform for exploring site-specific multivalent ligand-receptor or cell-surface interactions in combination with nucleic acid functionalized nanoparticles, for regulation of gene expression.⁵⁰

Methods

Preparation of DNA nanostructure patterned AuNPs.

A generic protocol was developed and applied for all prism/sequence variants, here \mathbf{Cb} - $\mathbf{Da_4}$ is provided as an example. In a typical patterning experiment, BSPP coated AuNPs (20 picomoles) were incubated with \mathbf{Cb} - $\mathbf{Da_4}$ (20 pmol [3D construct]) (BSPP: bis(p-sulfonatophenyl)phenylphosphine). Final sample concentrations were ca. 300nM in a tris (90mM), boric acid (90mM), EDTA (2mM) and NaCl (100mM) buffer (1xTBEN, pH 8.3) with 2mM BSPP. Following Incubation at room temperature overnight, 20,000 equivalents of OEG (ca. 0.2M in 1xTBEN) was added to stably passivate the surface. After a 30 minute incubation with OEG at room temperature, samples were centrifuged (12000 x g) at 4°C, the supernatant was removed and fresh buffer added. This process was repeated at least twice before loading the samples on gel. Gel bands were assigned, see Figures 2 and 3, and the desired band carefully excised. Samples were extracted by electroelution in 1xTBE buffer and subsequently concentrated by centrifugation (15000 x g) at 30°C and removal of supernatant. The 'pellet' was then washed with 3M urea in 0.625xTBE three times. This was followed by three washes with sterile de-ionized water. Finally, sample solutions were quantified by absorption spectroscopy, monitoring absorbance at 450nm. Further experimental details, protocols and characterization data can be found in the Supplementary Information.

Ackowledgements

The authors acknowledge the Natural Sciences and Engineering Research Council of Canada (NSERC), the Canada Foundation for Innovation (CFI), the Centre for Self-Assembled Chemical Structures (CSACS) and the Canada Research Chairs Program for financial support. T.G.W.E thanks the Canadian Institutes of Health Research (CIHR) for a Drug Development Training Program (DDTP) scholarship. D.B. thanks NSERC for a Bionano scholarship and C.J.S thanks NSERC for a Banting Postdoctoral Fellowship. H.F.S. is a Cottrell Scholar of the Research Corporation.

Author Contributions

H.F.S. and T.G.W.E. designed the project. T.G.W.E. primarily contributed to the production of experimental results. K.L.L. carried out gold nanoparticle synthesis, TEM analysis, preparation of pattered AuNPs for Fluorescence studies and aided in data interpretation. D.B. carried out all DLS

experiments. C.J.S. synthesized clip strands TC3-AB, PC4-AB and PC5-AB. All authors have agreed to all the content of the manuscript.

Competing Financial Interests

The authors declare no competing financial interests.

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