

DNA assembly of nanoparticle superstructures for controlled biological delivery and elimination

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1 DNA assembly mediates nanoparticle interactions with biological systems

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11
12 **DNA assembly of nanomaterials is a powerful approach to build complex nanostructures¹⁻⁴,**
13 **but the biological applications of such structures remain unexplored. Here we describe the**
14 **use of DNA to control the biological delivery and elimination of inorganic nanoparticles by**
15 **organizing them into colloidal superstructures. The individual nanoparticles serve as**
16 **building blocks, whose size, surface chemistry, and assembly architecture dictate overall**
17 **superstructure design. These superstructures interact with cells and tissues as a function**
18 **of their design, but subsequently degrade into building blocks that can escape biological**
19 **sequestration. We demonstrate that this strategy reduces nanoparticle retention by**
20 **macrophages, and improves their in vivo tumor accumulation and whole-body elimination.**
21 **Superstructures can be further functionalized to carry and protect imaging or therapeutic**
22 **agents against enzymatic degradation. These results suggest a new strategy to engineer**
23 **nanostructure interactions with biological systems, and highlight new directions in the**
24 **design of biodegradable and multifunctional nanomedicine.**

25
26 Inorganic nanoparticles can be synthesized in the 1-100nm size range with precise shapes, surface
27 chemistries, and physical properties. This engineering flexibility has enabled their design as novel
28 therapeutics, contrast agents, and integrated systems for the diagnosis and treatment of diseases⁵⁻
29 ⁸. To optimally deliver these nanoparticles to their biological targets with low toxicity, recent
30 studies have focused on understanding the effects of nanoparticle size, shape, and surface

31 chemistry – known as the physicochemical properties – on interactions with cells and tissues⁹⁻¹².
32 While several formulations have been shown to effectively target diseased tissues (e.g. tumors)<sup>13-
33 15</sup>, these designs diverge from those required for mitigating toxicity. Tumor targeting
34 nanoparticles require sufficiently large sizes to reduce clearance and improve retention within
35 tumors^{16,17}, yet such inorganic nanoparticles will remain in the body for a long time because they
36 do not biodegrade¹⁸. This in vivo persistence has raised concerns of chronic toxicity due to the
37 possibility for inorganic nanoparticles to aggregate^{19,20}, generate harmful metabolites^{21,22}, and
38 redistribute to vital organs within the body²³⁻²⁵. Few studies have demonstrated how the
39 physicochemical properties of inorganic nanoparticles can be engineered to mediate both delivery
40 and elimination²⁶. This design bottleneck will stall the clinical translation of these
41 nanotechnologies. Here we explore the use of DNA to organize sub-6nm inorganic nanoparticles, a
42 size that can be cleared through the kidneys, into larger superstructures to mediate their
43 biological delivery and elimination. This strategy combines the engineering flexibility of inorganic
44 nanoparticles with the biodegradability of organic molecules, which should open new avenues to
45 rationally engineer the interactions of inorganic nanoparticles with complex biological systems.

46
47 **Figure 1A** illustrates the principles of using DNA-nanoparticle assembly to engineer colloidal
48 superstructures with different physicochemical properties. First, we used metal-thiol or
49 streptavidin-biotin chemistry to functionalize inorganic nanoparticles with DNA. We then mixed
50 DNA-functionalized nanoparticles together with linker DNA strands containing complementary
51 sequences to initiate their assembly into colloidal superstructures. The architecture of the
52 assembled superstructure was controlled by using both nanoparticle geometry and DNA grafting
53 density, the latter determining the number of connections each nanoparticle makes with other
54 building blocks. Finally, the outer surface of the resulting superstructure was coated with
55 additional ligands to present the appropriate surface chemistries for interfacing with biological
56 systems. This was achieved by assembling nanoparticles with low DNA grafting densities on the
57 outer layer of the superstructure, such that their unsaturated surfaces provide binding sites for
58 ligand attachment.

59 Here, we used a “core-satellite” architecture to build DNA-assembled superstructures
60 where one or multiple layers of satellite nanoparticles surround a central core nanoparticle^{27,28}
61 (**Fig.1B**). Each layer of the core-satellite was encoded by a unique DNA sequence, such that

62 nanoparticles grafted with the specific DNA sequence inserted into the corresponding layer. A
63 linker DNA containing complementary regions to every layer joined the nanoparticles together.
64 Each layer of nanoparticles can be designed with a different composition, size, or surface
65 chemistry (**Fig.1B**). This modularity allowed us to construct superstructures with controlled
66 dimensions and multiple functionalities from relatively simple building blocks. The permutations
67 amongst nanoparticle designs and DNA sequences can also quickly generate superstructures with
68 distinct physiochemical properties. **Figure 1C** shows the use of 2 unique nanoparticle building
69 blocks and 2 DNA sequences (e.g. 2 layers) to give $2^2=4$ unique core-satellite superstructures. The
70 total number of unique superstructures increases exponentially with increasing number of core-
71 satellite layers and nanoparticle designs. For example, the combination of 10 nanoparticle designs
72 in a 3-layer (e.g. 3 DNA sequences) core-satellite would give $3^{10}= 59049$ unique superstructures;
73 the use of “*n*-layer” core-satellites with *m* nanoparticle designs gives n^m unique superstructures,
74 each may interact differently with cells and tissues. This diversity of superstructure candidates
75 allowed us to identify designs with high biological stability, low non-specific biological
76 interactions, and favorable pharmacokinetics for disease targeting.

77 Based on these principles, we generated a sub-library of colloidal superstructures with
78 different hydrodynamic sizes and surface chemistries to study the impact of their design on
79 molecular and cellular interactions. **Figure 2A-B** shows the simplest 2-layer core-satellite
80 structures that were synthesized for these experiments. First, we synthesized 13nm gold
81 nanoparticles and used them as the core by grafting them with thiolated *core* oligonucleotides at a
82 density of ~ 0.12 DNA/nm². This density corresponded to a valency of 80 to 90 DNA strands per
83 particle, allowing them to make a large number of connections with the satellites. DNA grafting
84 density was controlled by varying the DNA-to-nanoparticle grafting stoichiometry and quantified
85 by using a fluorescence depletion assay (**fig.S1**). We then synthesized 3 and 5nm gold
86 nanoparticles as the satellites by coating them with the *satellite* oligonucleotide sequence at a
87 density of ~ 0.05 DNA/nm². This density corresponded to 2 to 3 DNA strands per particle, which
88 was sufficient to stabilize the satellites against aggregation but minimized their probability of
89 cross-linking superstructures into macroscopic aggregates. We note that this low DNA coverage
90 also left the rest of the satellite nanoparticle surface available for further ligand conjugation. A
91 *linker* DNA containing complementary regions to both the *core* and *satellite* sequences was used to
92 join these nanoparticles together. To assemble core-satellites, we first annealed a stoichiometric

93 amount of linker DNAs with the core nanoparticles in a hybridization buffer that was first heated
94 to 60°C for 10 minutes and then kept at 37°C for 2 hours. Linker-hybridized core nanoparticles
95 were then purified by centrifugation and subsequently combined with satellite nanoparticles
96 under similar hybridization conditions. We used a 100X molar excess of satellite nanoparticles per
97 core nanoparticle to further eliminate the probability of superstructure cross-linking. Following
98 core-satellite assembly, colloidal superstructures were back-filled with the polymer poly(ethylene
99 glycol) (PEG) to improve their biological stability and reduce non-specific interactions with
100 biomolecules and cells²⁹. We used 4 different linker stoichiometries (2, 8, 16, 24 linkers per core,
101 see characterization in **fig.S2**), which generated superstructures with different satellite-to-core
102 ratios (**Fig.2A**). We used 3 different lengths of PEG (1, 5, 10kDa) to control overall superstructure
103 surface chemistry and morphology (**Fig.2B**). We also generated 3-layer core-satellite structures in
104 which a third DNA sequence (*satellite2*) hybridizes to an internal region of the linker (see
105 schematic in **fig.S3** and images in **Fig.2C-i**). By grafting this DNA sequence onto other sets of
106 nanoparticles, superstructures with additional satellite layers could be constructed (**Fig.2C ii-iv**
107 and **fig.S4**). Varying these parameters generated a diverse set of superstructures with
108 hydrodynamic sizes ranging from 50-150nm (**fig.S5**). Transmission electron microscopy (TEM,
109 **Fig.2D**) and UV-vis absorbance characterizations (**fig.S6**) demonstrated that these
110 superstructures were monodisperse and colloidally stable in saline.

111 A key question regarding the biological application of colloidal superstructures is whether
112 they can carry and protect pharmaceuticals against biological degradation. We found that
113 therapeutic or imaging agents such as doxorubicin and several fluorescent molecules can be
114 incorporated into superstructures through DNA intercalation or groove binding (**Fig.2E**).
115 Incorporation efficiency was dependent on linker sequence, improving with increasing number of
116 TCG repeats which is a known binding site for doxorubicin³⁰. Other agents such as quantum dots
117 and fluorescein amidite (FAM), which do not intercalate or bind DNA directly, could be
118 incorporated within superstructures as hybridized DNA conjugates (**Fig.2F**). An advantage of
119 using assembly to store these agents is that they are embedded within the superstructure and not
120 exposed on the nanoparticle surface (**Fig.2G**). By selecting the appropriate core and satellite
121 building blocks, superstructures enhanced DNA resistance against nuclease and serum
122 degradation by up to 5-fold relative to non-assembled nanoparticles (**fig.S7**). This improvement in
123 DNA stability effectively protected the superstructures and its payloads from disintegrating in

124 biological solutions. These results provide the first example of using assembly architecture to
125 mediate payload stability, and highlight a novel strategy to build integrated platforms that carry
126 multiple functionalities.

127
128 The potential application of colloidal superstructures as delivery platforms motivated us to
129 further investigate their interactions with cells. We selected J774A.1 macrophage cells as a model
130 cell system because macrophages sequester the majority of in vivo administered nanoparticles³¹.
131 Sequestration of nanoparticles by macrophages not only limits the dose that is available to
132 accumulate at diseased sites but is further associated with immune-toxicity^{32,33}. The ability to
133 control nanoparticle interactions with macrophages could improve disease-specific delivery and
134 reduce toxicity. We measured macrophage uptake by incubating J774A.1 cells in culture media
135 containing gold nanoparticles for 4 hours and then analyzing the total cellular gold content using
136 inductively coupled plasma atomic emission spectroscopy (ICP-AES, **Fig.3A**). To assess the impact
137 of nanoparticle design and assembly on uptake, we first exposed macrophages separately with
138 13nm core nanoparticles, 5nm satellite nanoparticles coated with 1kDa PEG, as well as
139 superstructures assembled using these two components. **Figure 3B** shows that macrophages
140 sequestered 13nm core nanoparticles 7 times more effectively than 5nm satellite nanoparticles
141 coated with PEG 1kDa, consistent with previous findings that macrophage uptake correlates with
142 nanomaterial size and surface charge³⁴. Interestingly, the core-satellite superstructure was 2.5
143 times larger than its core component but resulted in 2-fold lower uptake into macrophages,
144 suggesting that the superstructure displayed a different surface chemistry which inhibited its
145 uptake. These results also motivated us to systematically characterize the impact of building block
146 design and their assembly architecture on macrophage uptake. We used serum-free culture media
147 for these experiments because our results (**fig.S8**) and a previous study³⁵ have shown that DNA
148 coated nanomaterials are taken up by cells through direct interactions with receptors (e.g.,
149 scavenger receptors) on the cell surface rather than through interactions with serum proteins
150 adsorbed on the nanomaterial surface. Here we observed a monotonic decrease in superstructure
151 uptake by macrophages as a function of satellite-to-core ratio (**Fig.3C**), suggesting that satellites
152 inhibited macrophages from interacting with the core. This hypothesis is further supported by the
153 different dose-responses between these nanomaterials; macrophages sequestered DNA coated
154 core nanoparticles in a dose-independent manner suggesting that cells take up such nanoparticles

155 efficiently. In contrast, core-satellite structures exhibited a dose-dependent decrease in cell uptake
156 similar to PEGylated nanomaterials (**fig.S9**). The length of PEG on the satellites also impacted
157 macrophage uptake, where an increase from PEG 1 to 10kDa reduced macrophage uptake by an
158 additional 30% (**Fig.3D**). Interestingly, 5nm nanoparticles were ~2 times more effective than 3nm
159 nanoparticles at mitigating core-satellite uptake by macrophages, implying they provide a denser
160 PEG surface chemistry (**Fig.3E**). Taken together, the optimal superstructure design reduced
161 macrophage uptake by 80% relative to the core nanoparticle, despite being 3 times larger in size.
162 This was achieved by using 5nm nanoparticles coated with PEG 10kDa as satellites at a saturating
163 satellite-to-core ratio. Other parameters such as linker length (**fig.S10**) had relatively little effect
164 on uptake. These results highlight the central role of satellite design and assembly stoichiometry
165 in dictating superstructure interactions with cells. Nanoparticle assembly can reduce macrophage
166 uptake by: 1) burying DNA within the superstructure to decrease their accessibility from cellular
167 interactions, and 2) using nanoparticles as scaffolds to increase the density of PEG coverage above
168 the DNAs.

169 Nanomaterials internalized by macrophages are sequestered within the cells if they are not
170 biodegraded. This contributes to the persistence of inorganic nanoparticles within the body^{18,25}.
171 To investigate how superstructures are processed within macrophages, we washed the cells
172 following their incubation with superstructures and chemically fixed them for visualization under
173 TEM (**Fig.4A**). Electron micrographs reveal that superstructures associated with the extracellular
174 membrane of macrophages both as single entities (**fig.S11**) and as clusters (**Fig.4B-i**). Associated
175 superstructures were eventually internalized by macrophages within vesicles, in which
176 superstructures disassembled into their respective building blocks (**Fig.4B-ii**). We did not observe
177 intact superstructures within macrophages, even in cells fixed immediately following exposure to
178 the superstructures (**fig.S12**), suggesting that the intracellular degradation of superstructures
179 occurred rapidly. In contrast, superstructures incubated in culture medium alone (i.e. without
180 cells) remained largely intact over 8 hours of incubation (**Fig.4C**), indicating that superstructure
181 degradation was intracellularly-mediated. Phagocytic vesicles are known to contain a complex
182 mixture of 40 or more hydrolytic enzymes that are responsible for digesting foreign pathogens or
183 endogenous debris. It is possible for this mixture to quickly hydrolyze the DNA linkages that
184 connect the nanoparticles together^{36,37}. While many nanoparticle formulations have been reported
185 to aggregate under such environments¹⁰, superstructure components remained dispersed

186 following breakdown. These building blocks eventually escaped from the vesicles and distributed
187 throughout the cellular cytoplasm (**Fig.4B-iii**). To test whether this intracellular behaviour is
188 mediated by superstructure assembly, we incubated core and satellite nanoparticles separately
189 with macrophages under identical conditions and then examined their subcellular localization
190 over time under TEM. Here we observed that while both core and satellite nanoparticles were
191 endocytosed within vesicles, some satellite nanoparticles also entered cells via vesicle-
192 independent pathways (**fig.S13-14**). In cells incubated with core nanoparticles, endocytosis
193 resulted in the appearance of nanoparticle clusters that were confined within vesicles and grew in
194 size over time, suggesting core nanoparticles are actively sorted into phagosomes from which they
195 fail to escape. Satellite nanoparticles, in contrast, could be identified within cells as both being
196 confined to vesicles and as individual, discrete nanoparticles within the cytoplasm. **Figure S14**
197 further shows several instances where satellite nanoparticles originally confined within vesicles
198 were released into the cytoplasm or excreted across the plasma membrane (**fig.S14**). These
199 results suggest that, when delivered to the cells alone, the intracellular behavior of nanoparticles
200 is determined by their design. However, the assembly of these nanoparticles into superstructures
201 alters their intracellular behavior.

202 These results prompted us to carry out a parallel experiment, in which we measured
203 changes in total intracellular gold content to assess whether dispersed building blocks could
204 escape from these macrophages following uptake (**Fig.4D**). In cells treated with superstructures,
205 we found a 10 to 40% reduction in intracellular gold content over the course of 8 hours (**Fig.4E**).
206 The extent of this reduction was dependent on satellite design. In contrast, no change in gold
207 content was measurable in cells incubated with core nanoparticles alone (**Fig.4E**), suggesting that
208 measured differences were attributable to the satellites. These differences occurred
209 independently of changes in cell density (**fig.S15**) and plasma membrane permeability (**fig.S16**),
210 and were apparent when cells harvested from different time points were cross-examined under
211 TEM (**fig.S17**). Control experiments further verified that satellites alone escaped macrophage
212 sequestration in a time- and PEG length-dependent manner (**Fig.4F & fig.S18**). While the role of
213 nanoparticle and PEG size in cellular uptake has been widely reported, results herein suggest that
214 these design parameters also define the thresholds for cellular excretion, which has implications
215 for the in vivo clearance and toxicity of nanomaterials. Additionally, molecular assembly
216 techniques may offer a unique approach whereby PEGylated satellite nanoparticles can be used to

217 facilitate therapeutic delivery without contributing significantly to overall in vivo persistence of
218 nanomaterials.

219

220 If small nanoparticles degraded from the superstructures can escape macrophage sequestration,
221 we suspect that they can be designed small enough for renal elimination in vivo. This would
222 decrease the biological persistence of nanoparticles injected into the body and eliminate their
223 risks of chronic toxicity. To test this, we synthesized a panel of satellite building blocks and
224 administered them intravenously into CD1 mice. We housed the mice within metabolic cages,
225 collected their urine for up to 48 hours post-injection, and analyzed the urine for gold content.
226 Urinary excretion was highest for the smallest satellites at 15% of the injected dose and
227 diminished rapidly with nanoparticle size (**Fig.5A**). Building on these results we assembled core-
228 satellites with the smallest satellite nanoparticles to test the ability of superstructures to undergo
229 renal clearance. Urinary excretion efficiency of superstructures mirrored closely with the
230 clearance behavior of their building blocks (**Fig.5B**), suggesting they can be engineered to
231 eliminate from the body unlike larger solid nanoparticles. More importantly, this result
232 underscores an approach to tailor the size and surface chemistry of nanostructures for mediating
233 their delivery while allowing them to clear from the body.

234 Finally, we assessed the potential of using superstructures to target tumors via a passive
235 mechanism. Preliminary results with xenograft tumor models demonstrate that one of our
236 current superstructure formulations accumulated within tumors better than its controls (e.g. core
237 nanoparticle and non-assembled mixture) following systemic administration (**Fig.5C**). Using a
238 previously published procedure for fluorescently-labeling gold nanoparticles³⁸, we chemically
239 conjugated this formulation with near-infrared dyes in order to monitor their distribution in
240 tumor xenograft models in real time. Our characterization shows the structures were not altered
241 during this modification (**fig.S19**). Whole-animal fluorescence imaging showed that this
242 superstructure design increased tumor-specific fluorescence contrast steadily over time (**fig.S20**),
243 achieving a final tumor-over-background ratio of 2.3 ± 0.1 and a signal-over-noise ratio of 5.2 ± 0.5
244 at 24 hours following administration (**Fig.5D** and **fig.S21**). Analysis of the fluorescence images
245 estimated a blood circulation half-life of 5 hours (**fig.S21**). To ensure that superstructures were
246 non-toxic, we collected blood from these animals for biochemistry and hematology analysis, and
247 harvested organs for biodistribution and histology analysis. Results show that, while a large

248 proportion of these superstructures also accumulated in the liver and spleen (**fig.21**), they did not
249 cause acute toxicity and were well-tolerated by the animals at the given dose (**fig.S22-S23**).
250 Together, these results demonstrate promise of using molecularly assembled superstructures for
251 in vivo biomedical applications. As a next step, we are currently preparing a library of
252 nanoparticles to further investigate and understand the effect of design on the rate and efficiency
253 of tumor accumulation and whole-body clearance, similar to our previous study¹³. Our current
254 findings have now defined the required building block designs and assembly architectures to
255 engineer superstructures that can accumulate in tumors and be eliminated from the body.

256
257 In summary, we demonstrated the use of molecular assembly to mediate the biological delivery
258 and elimination of nanoparticles. We showed that colloidal superstructures assembled with the
259 appropriate building blocks and architectures can reduce their uptake and sequestration by
260 macrophages, improve their accumulation into tumors, and facilitate their elimination from the
261 body. The use of DNA assembly to engineer nanodelivery vehicles offers five advantages: 1)
262 accurate and programmable control over nanocarrier architecture, 2) modular construction of
263 complex platforms from simple nanoparticle building blocks, 3) compartmentalization of imaging
264 or therapeutic payloads against biological degradation, 4) new strategies for controlling the
265 release of therapeutics (e.g., DNAzymes), and 5) ability to control the design of multifunctional
266 nanomedicines (e.g. delivery vehicle with PET, MRI, and optical imaging agents or therapeutics).
267 The use of both DNA and nanoparticle technologies together can help translate fundamental
268 nanomaterial design principles that are being discovered into clinically relevant nanomedicine
269 solutions.

270

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280

281 **Author Contributions**

282 W.C.W.C, L.Y.T.C., and K.Z. conceived the idea. W.C.W.C and L.Y.T.C. wrote the paper. L.Y.T.C. and
283 K.Z. designed and performed experiments. All authors analyzed data.

284

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367

368

369 **FIGURE CAPTIONS**

370

371 **Figure 1. Design of colloidal superstructures using DNA-nanoparticle assembly.** (A) DNA-coated
372 nanoparticles were assembled into superstructures using linker DNAs. Nanoparticles on the surface of the
373 superstructure served as “scaffolds” for displaying additional ligands (e.g. PEG), which modulated overall
374 superstructure interactions with cells and tissues. (B) This study focused on the design of “core-satellite”
375 superstructures, in which a central nanoparticle (i.e. *core*) is surrounded by one or multiple layers of
376 satellite nanoparticles (i.e. *layers 2 to n*). Each layer is encoded by a unique DNA sequence capable of
377 connecting to nanoparticles of different size, surface chemistry, or composition. The combination of
378 building blocks determines the overall dimension and functionality of the superstructure. (C) Large
379 numbers of unique superstructures can be generated and screened by combining different building blocks
380 together. An example is shown here, where 2 different nanoparticle designs and 2 unique DNA sequences
381 combine to give 4 unique superstructures.

382

383 **Figure 2. Synthesis and characterization of core-satellite superstructures.** TEM characterization of 2-
384 layer core-satellites as a function of (A) satellite-to-core ratio ($r=2, 8, 16$ and 24) and (B) satellite PEG
385 length (MW=bare, 1, 5 and 10kDa). (C) 3-layered core-satellites were synthesized by introducing a third
386 DNA sequence (*Satellite2*) which inserts into the linker DNA. (i) Attaching this sequence to the linker
387 increased core-satellite separation distance when viewed under TEM. This DNA sequence (*Satellite2*) were
388 also grafted onto other nanomaterials to generate various 3-layer superstructures (ii = 5nm gold
389 nanoparticles, iii = 3nm + 5nm gold nanoparticles, and iv = quantum dots + 3nm gold nanoparticles). Scale
390 bars = 50nm. (D) Core-satellite superstructures appeared colloiddally stable and monodisperse in saline.
391 Scale bars = 100nm. (E) Left: fluorescence spectra of core-satellites with and without doxorubicin
392 incorporation; right: fluorescence images of vials containing superstructures labeled with various DNA-
393 binding dyes (SG=Sybr Gold, PI=Propidium Iodide, S61=Syto61, TP3=TO-PRO-3). (F) Left: fluorescence
394 spectra of core-satellite superstructures with and without FAM incorporation; right: vials of
395 superstructures labeled with or without quantum dot incorporation under UV excitation. (G) Cross-
396 sectional view of a core-satellite showing the positioning of encapsulated payloads.

397

398 **Figure 3. Macrophage uptake of superstructures as a function of design.** (A) Workflow for assessing
399 superstructure uptake into J774A.1 macrophages by ICP-AES. (B) Relative uptake of 13-Core, 5-P1k
400 satellite nanoparticles, and corresponding assembled superstructures by macrophages. (C) Macrophage
401 uptake of superstructures as a function of satellite-to-core ratios. (D) Macrophage uptake of
402 superstructures with varying PEG lengths (satellite-to-core ratio kept constant at 24). (E) Effect of satellite
403 size on superstructure uptake by macrophages. Error bars represent S.E.M. from at least three independent
404 replicates; * $P<0.05$, ** $P<0.01$, *** $P<0.001$.

405

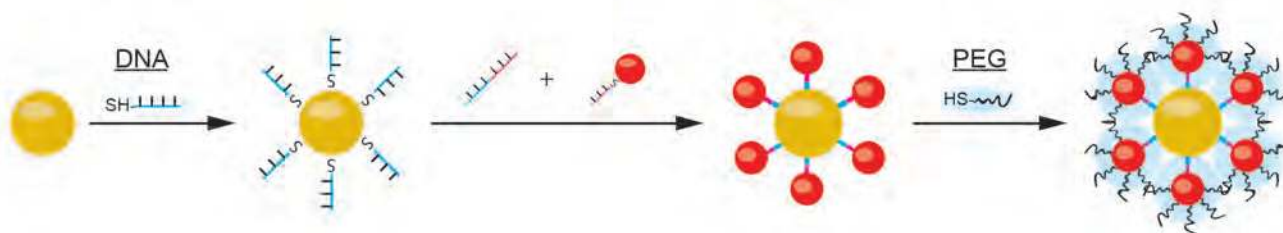
406 **Figure 4. Intracellular processing of superstructures by J774A.1 macrophages.** (A) Overview of
407 subcellular localization of superstructures. Outlined areas are magnified in (B): (i) Intact superstructures
408 first interact with the plasma membrane, and (ii) are internalized by the macrophage within vesicles,
409 where they undergo disassembly. (iii) Individual superstructure components eventually escape from
410 vesicles and were seen distributed throughout the cytosol. The panel on the right depicts similar stages of

411 processing occurring in another macrophage cell. (C) Core-satellite structures show minimal structural
412 disintegration after 8 hours of incubation in culture media containing 10% fetal bovine serum. All scale
413 bars = 100nm. (D) Workflow for assessing superstructure excretion by J774A.1 macrophages by ICP-AES.
414 (E) Time-course changes in intracellular gold content following exposure to various superstructures. (F)
415 Time-course changes in intracellular gold content following exposure to various satellite nanoparticles.

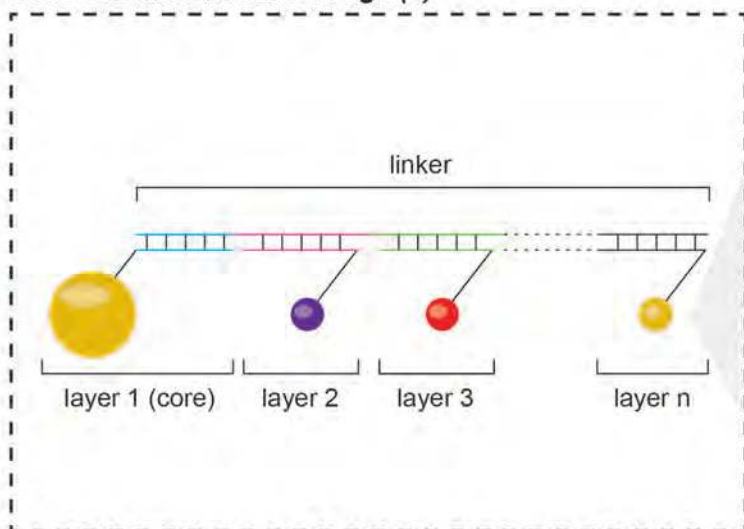
416

417 **Figure 5. Renal elimination and tumour accumulation of superstructures** (A) Total gold content in
418 mice urine up to 48 hours following systemic injection of various superstructure components. (B) Total
419 gold content in mice urine up to 48 hours following systemic injection of superstructures assembled from
420 the smallest satellite nanoparticles in (A). Results show that urinary excretion of superstructures mirrors
421 closely with their respective satellites building blocks. (C) Tumor accumulation of core-satellite
422 superstructures using 5nm-PEG10k satellite nanoparticles at 24 hours post-injection. Injection of core
423 nanoparticles alone and mixtures of unassembled superstructure components were used as separate
424 controls. Error bars represent S.E.M. from three to five independent replicates; *P<0.05, **P<0.01,
425 ***P<0.001. (D) Whole-animal imaging using fluorescently-labeled superstructures show tumor-specific
426 accumulation (white arrow) as well as accumulation in the liver (yellow arrow).

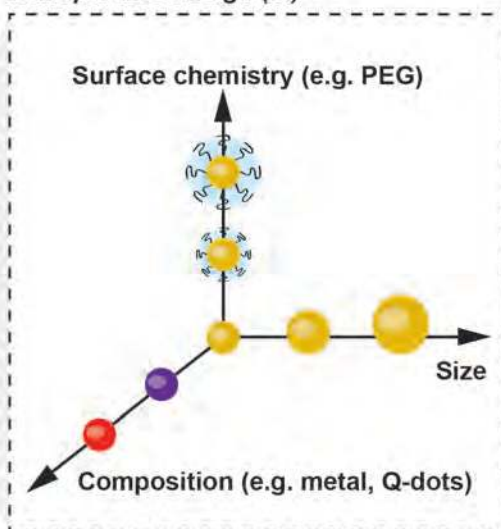
(A)



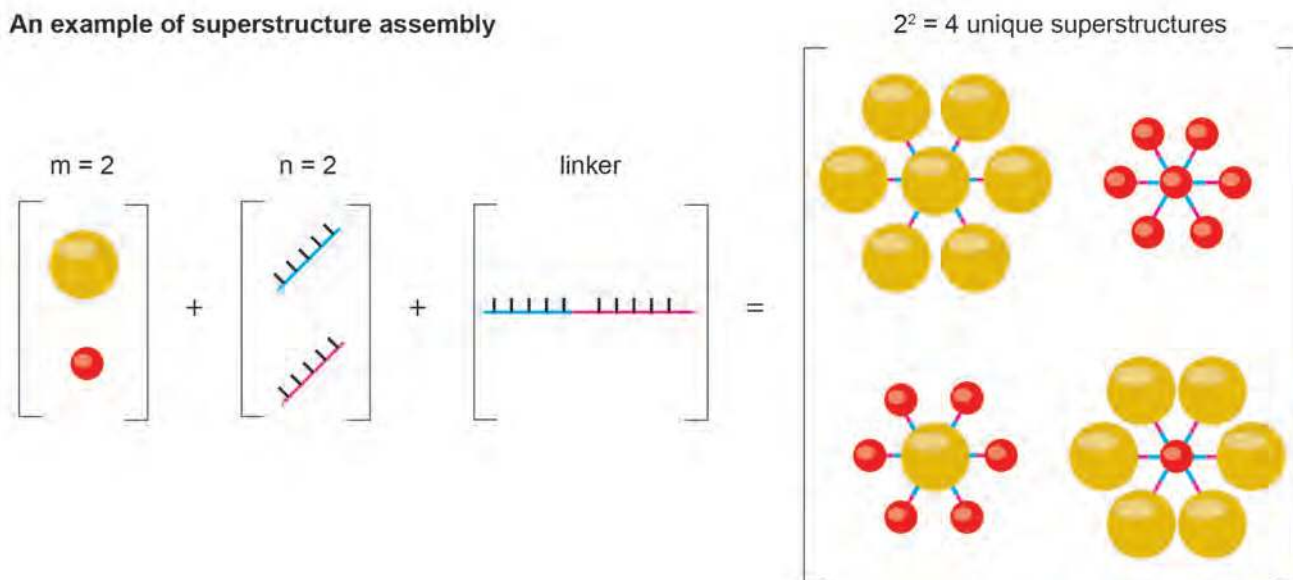
(B) Core-satellite structure design (n)



Nanoparticle design (m)



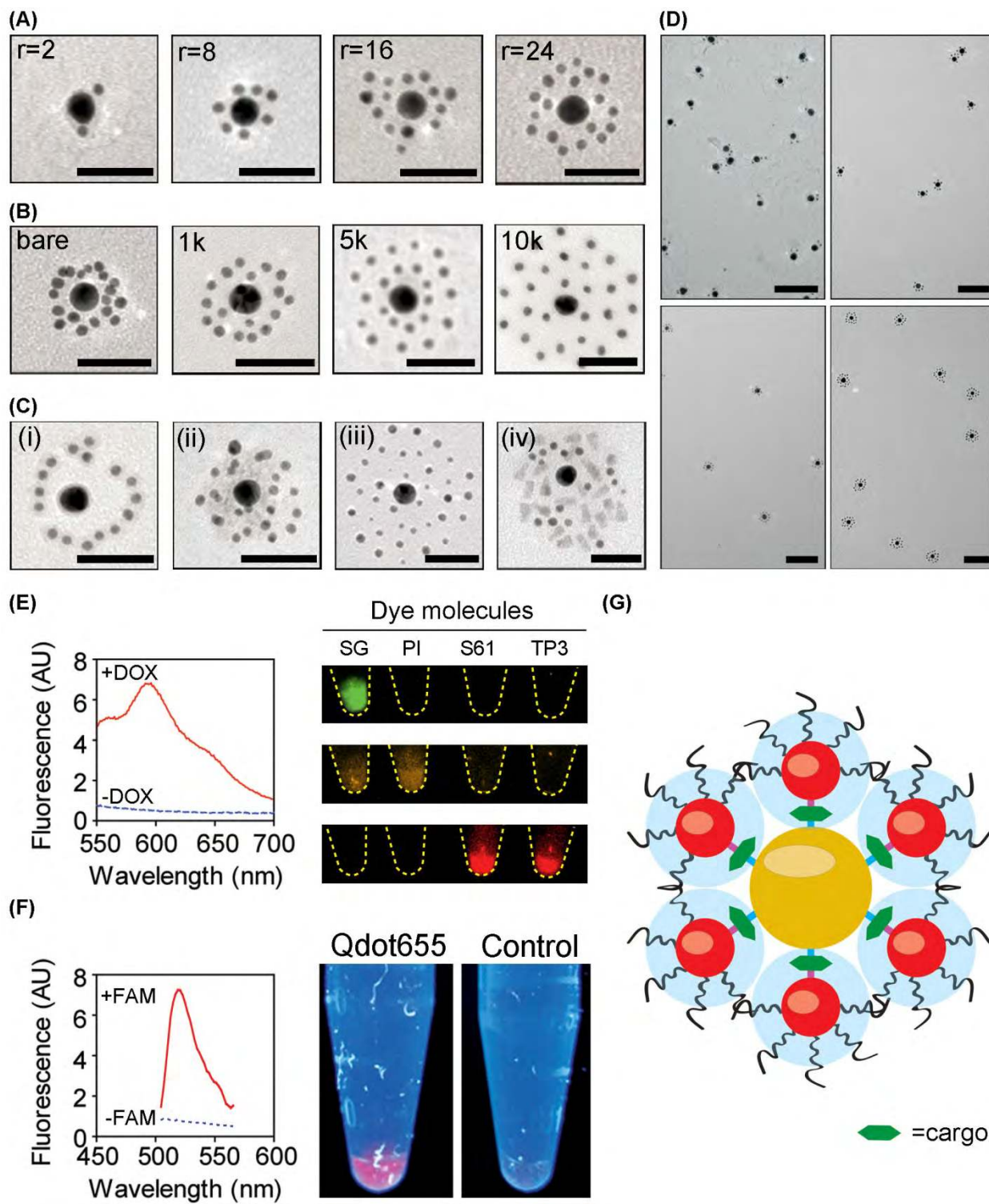
(C) An example of superstructure assembly



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428

Figure 1



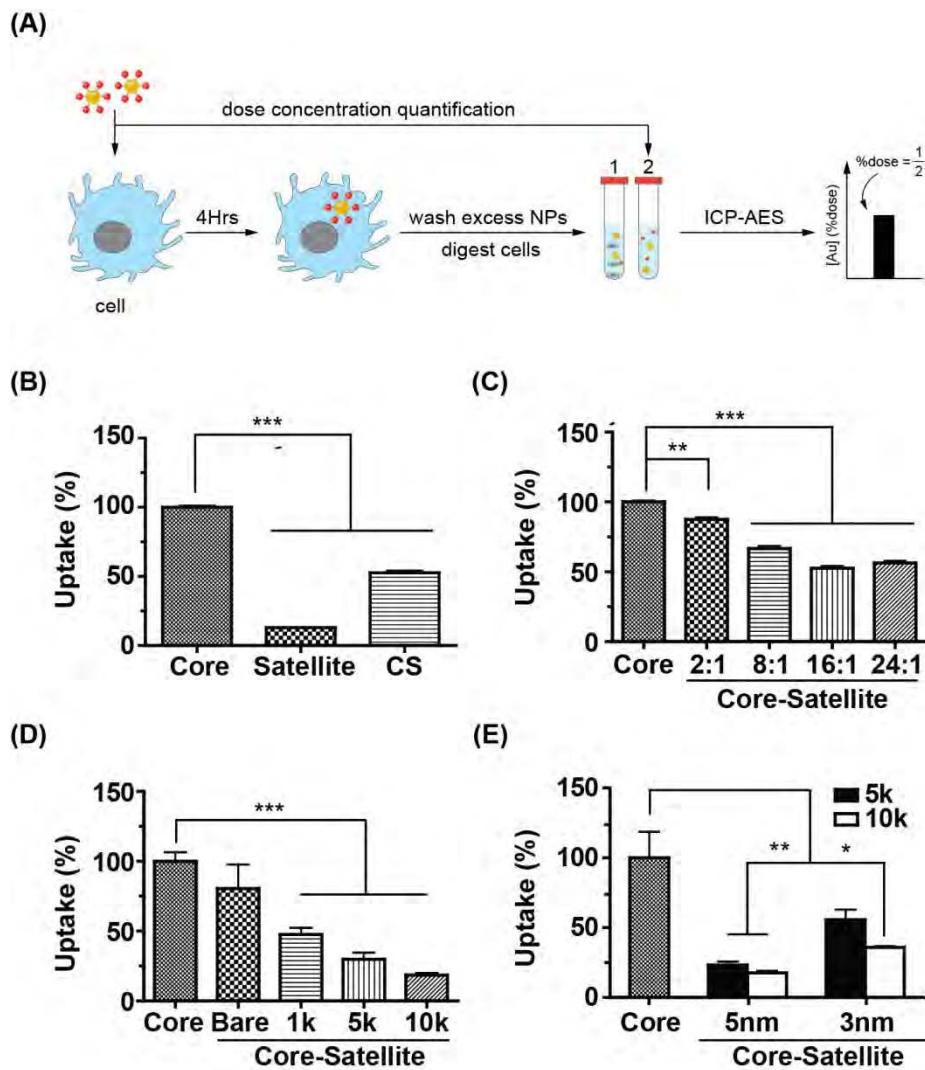
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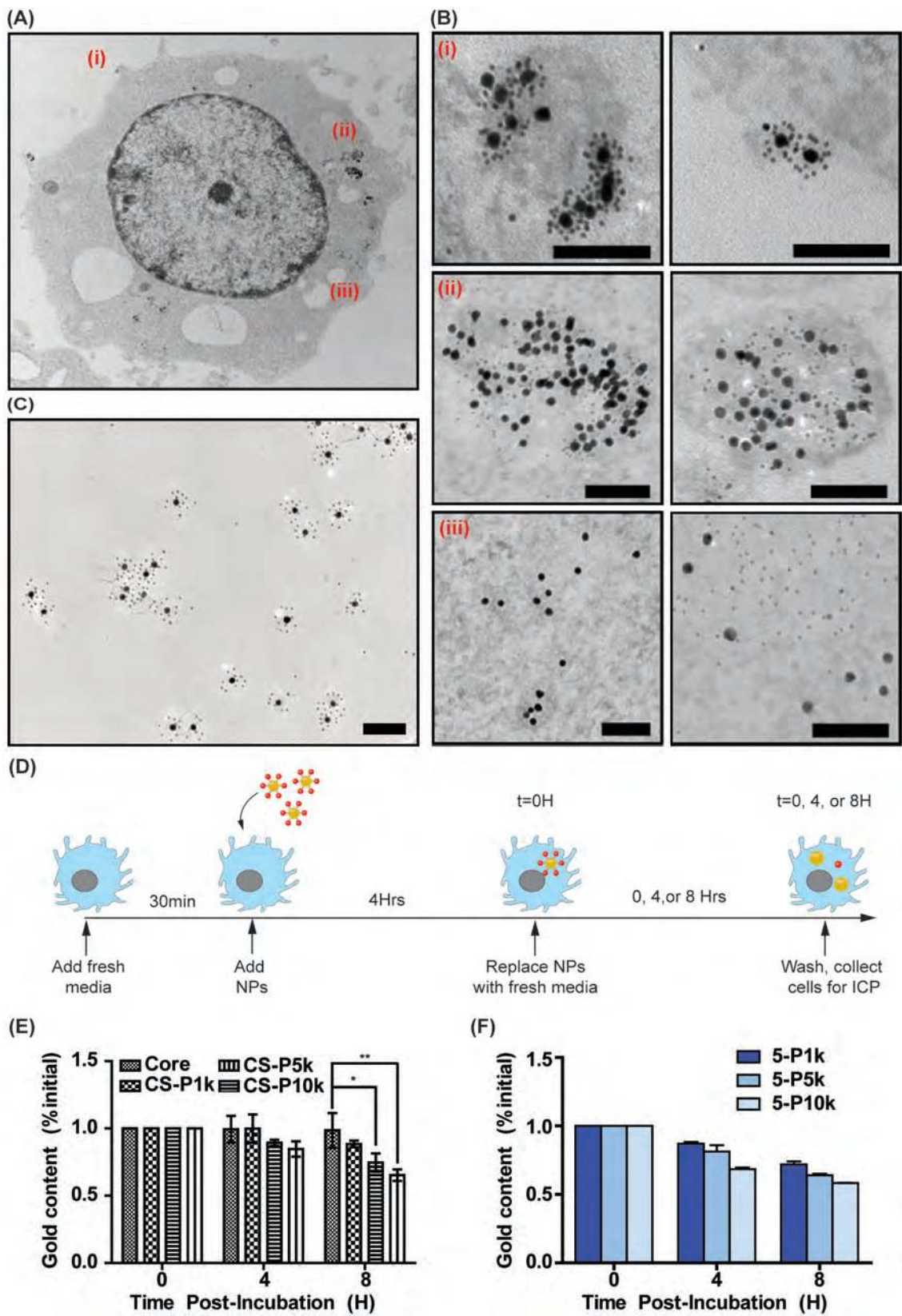
Figure 2



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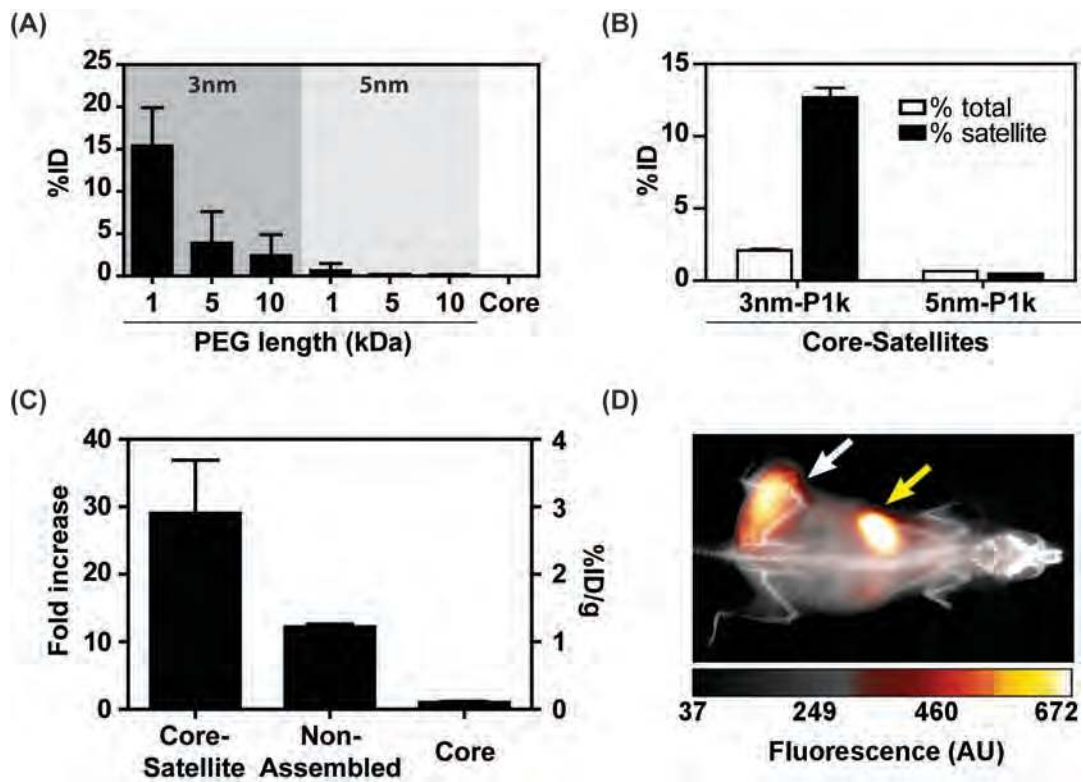
Figure 3



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Figure 4



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Figure 5