

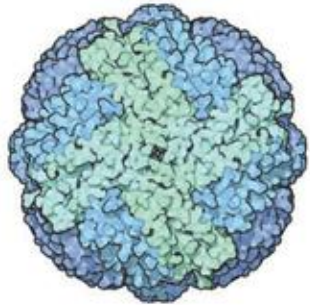
Design of self-assembling protein nanomaterials as next-generation vaccine scaffolds

 INSTITUTE FOR
Protein Design

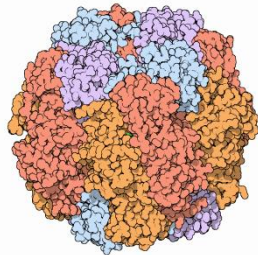
Neil King | March 15, 2016
UNIVERSITY *of* WASHINGTON

Protein self-assembly enables specialized functions; our goal is to design new self-assembling molecular machines

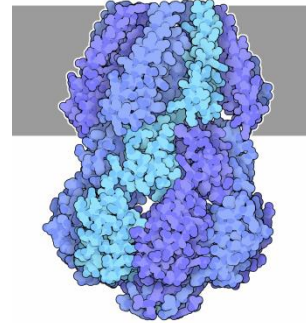
Storage containers



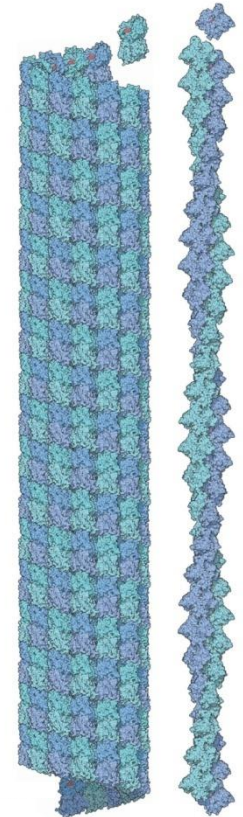
Catalysts



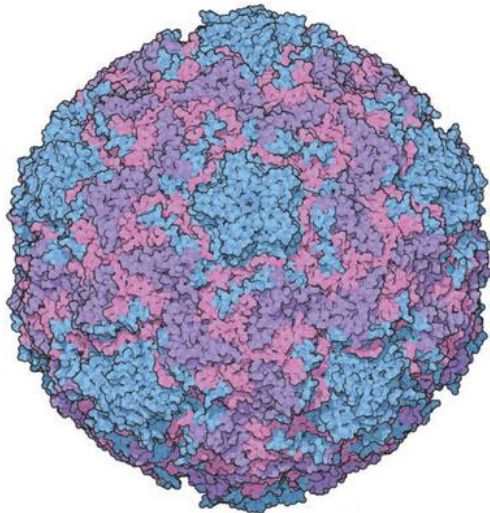
Signal Transducers



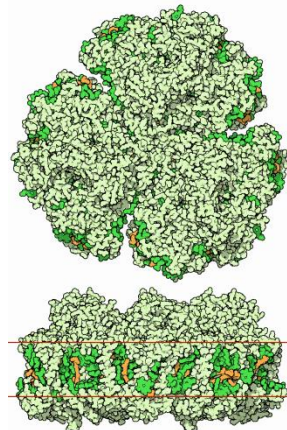
**Cellular scaffolding,
tracks for transport
and motility**



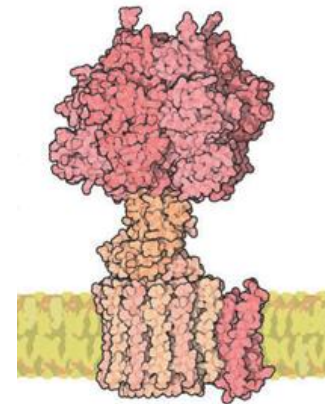
Delivery vehicles



Energy Converters

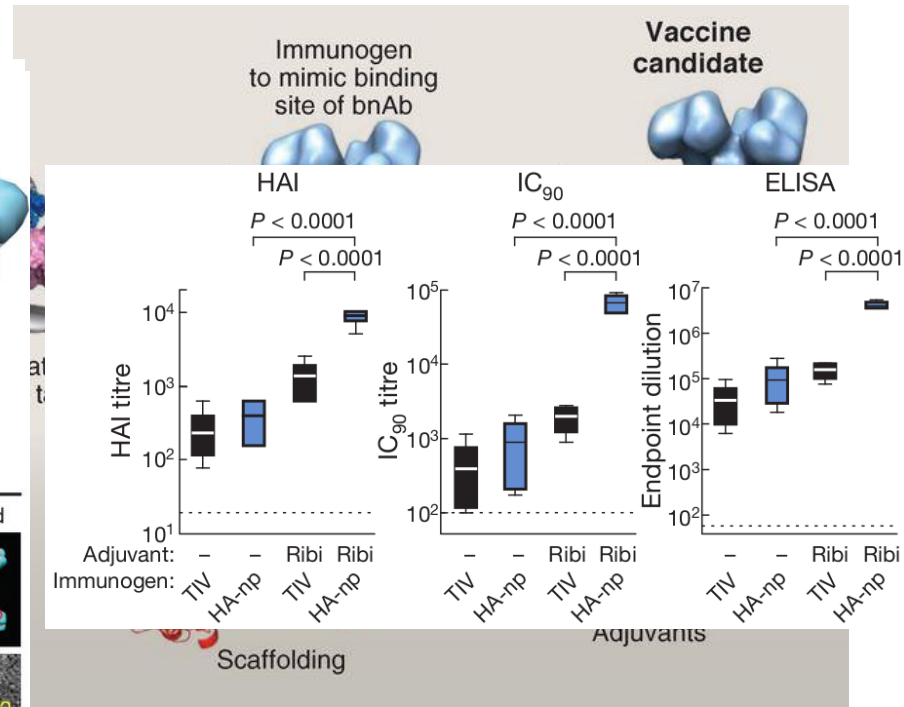
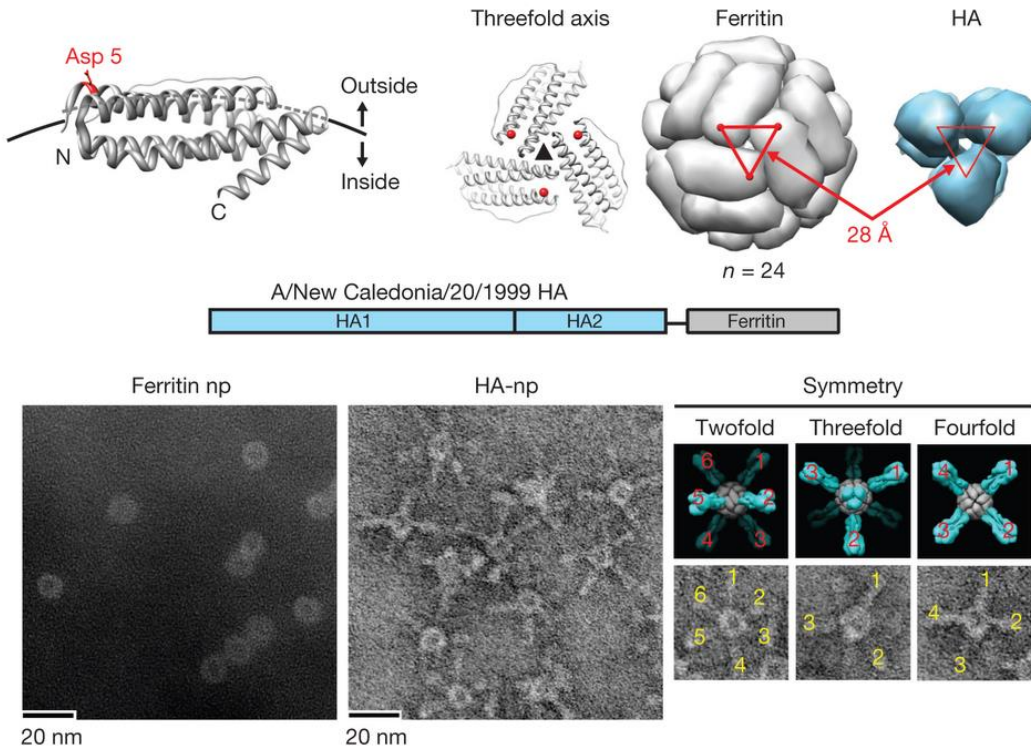


Molecular motors



Custom-designed self-assembling protein nanomaterials could facilitate new approaches to next-generation vaccine design

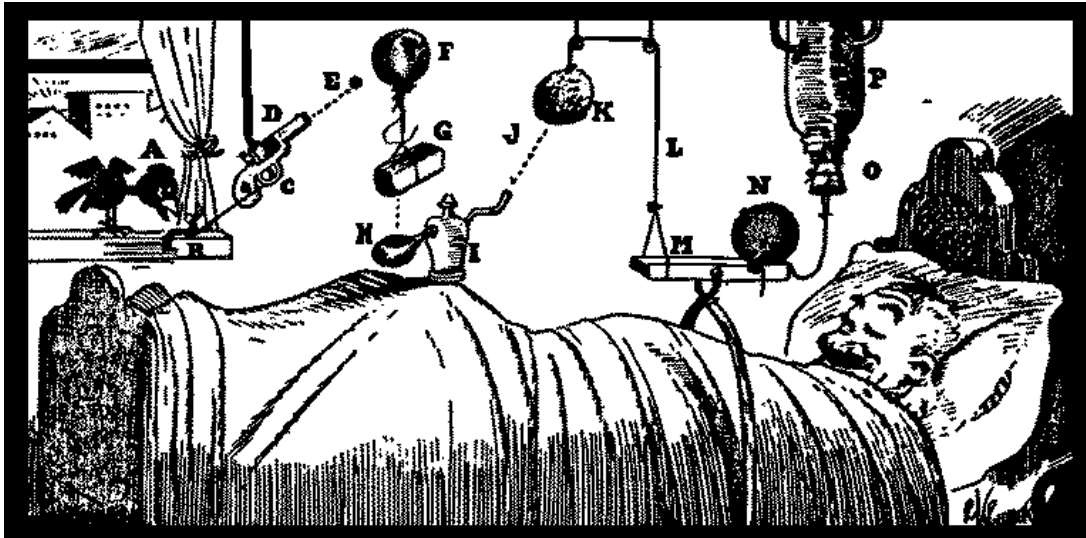
Self-assembling influenza nanoparticle vaccines elicit broadly neutralizing H1N1 antibodies



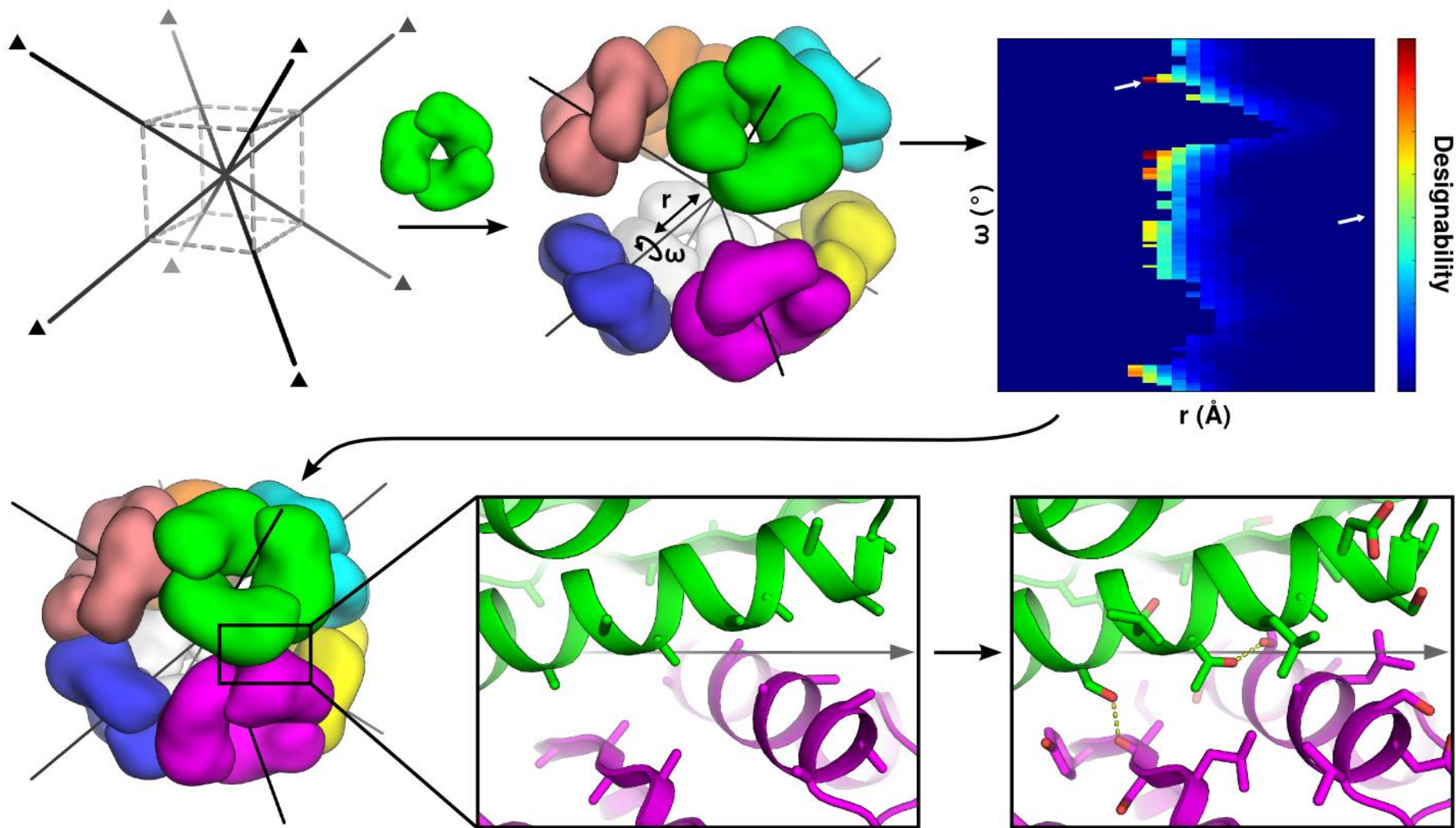
Koff WC, et al. (2013) *Science* **340**:1232910.

Kanekiyo M, et al. (2013) *Nature* **499**:102-6.

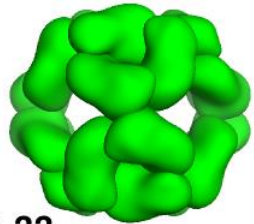
Design allows parts (proteins) to be built for a specific purpose



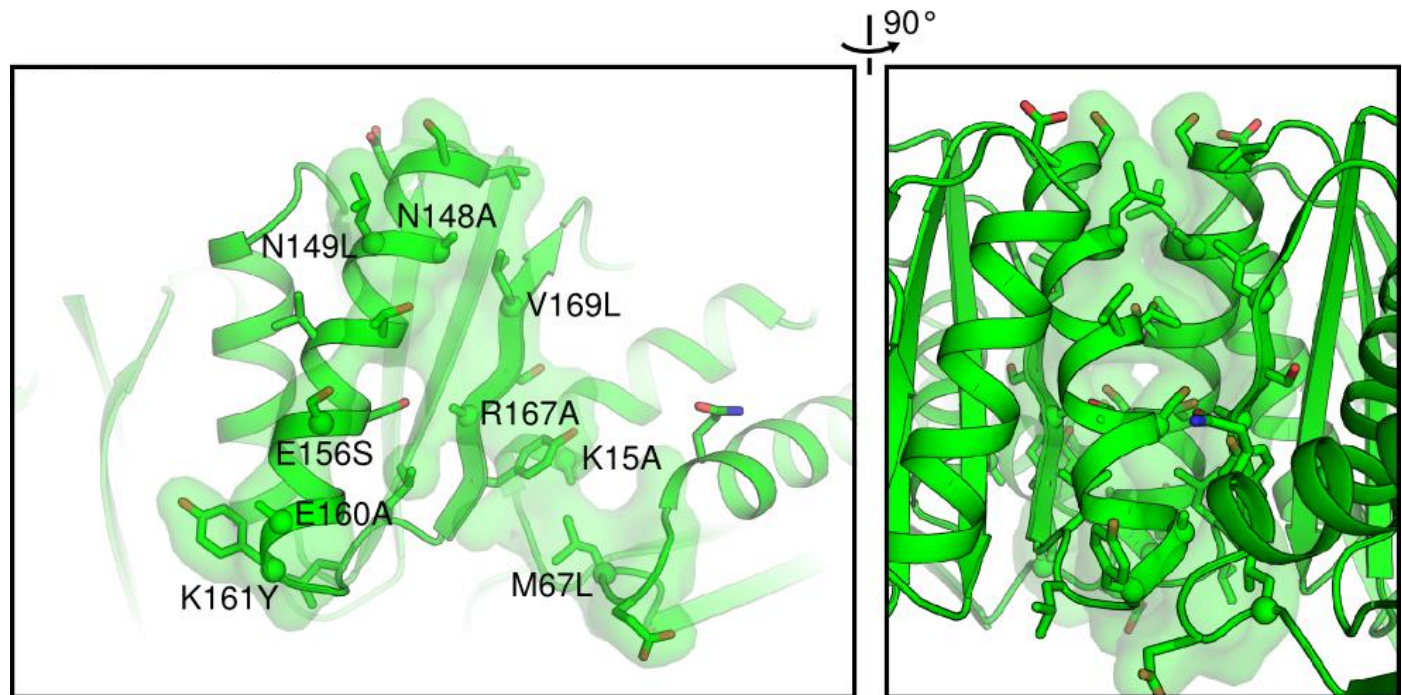
We have developed a general computational method for designing new self-assembling protein nanomaterials



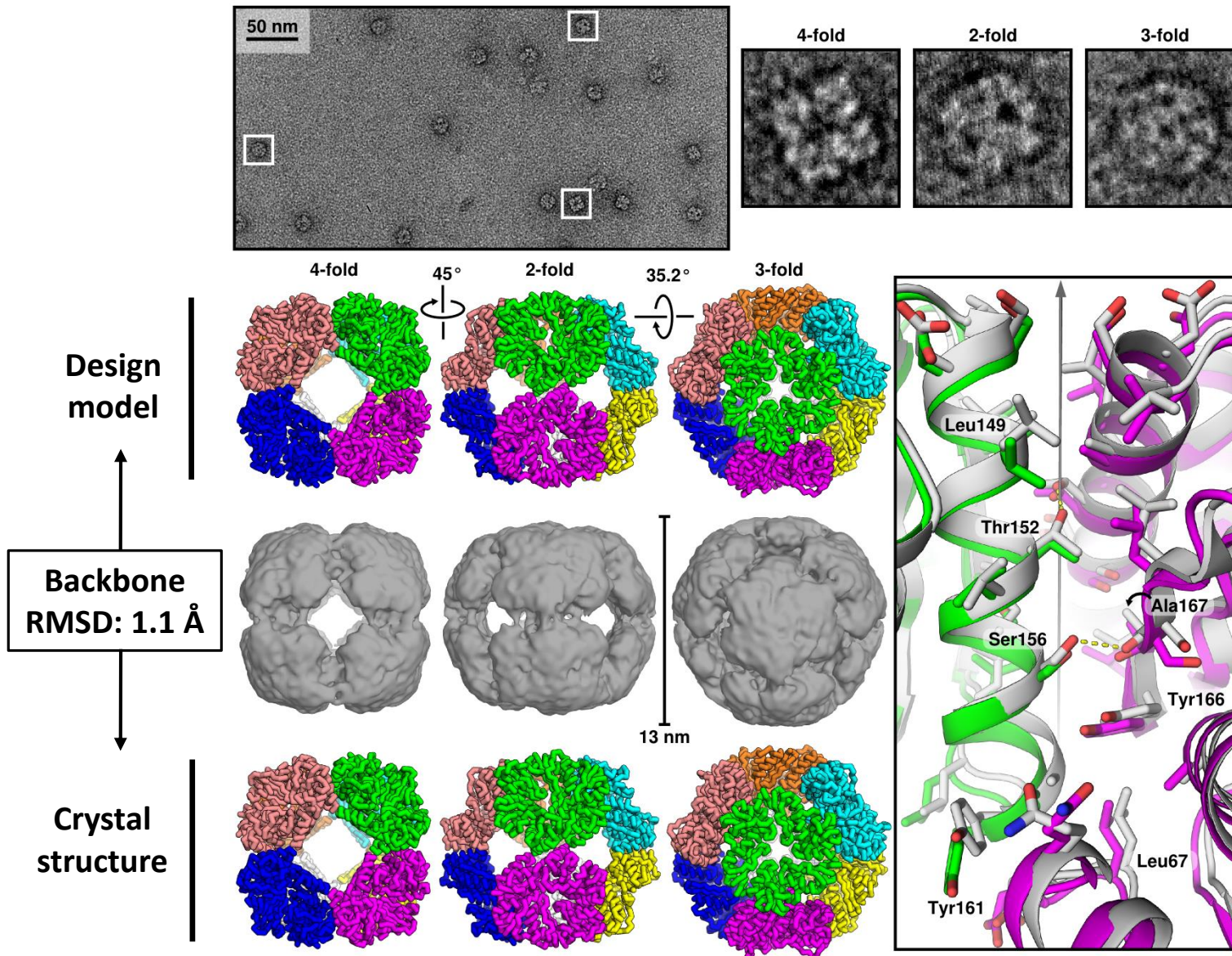
The designed interfaces have features resembling natural protein-protein interfaces



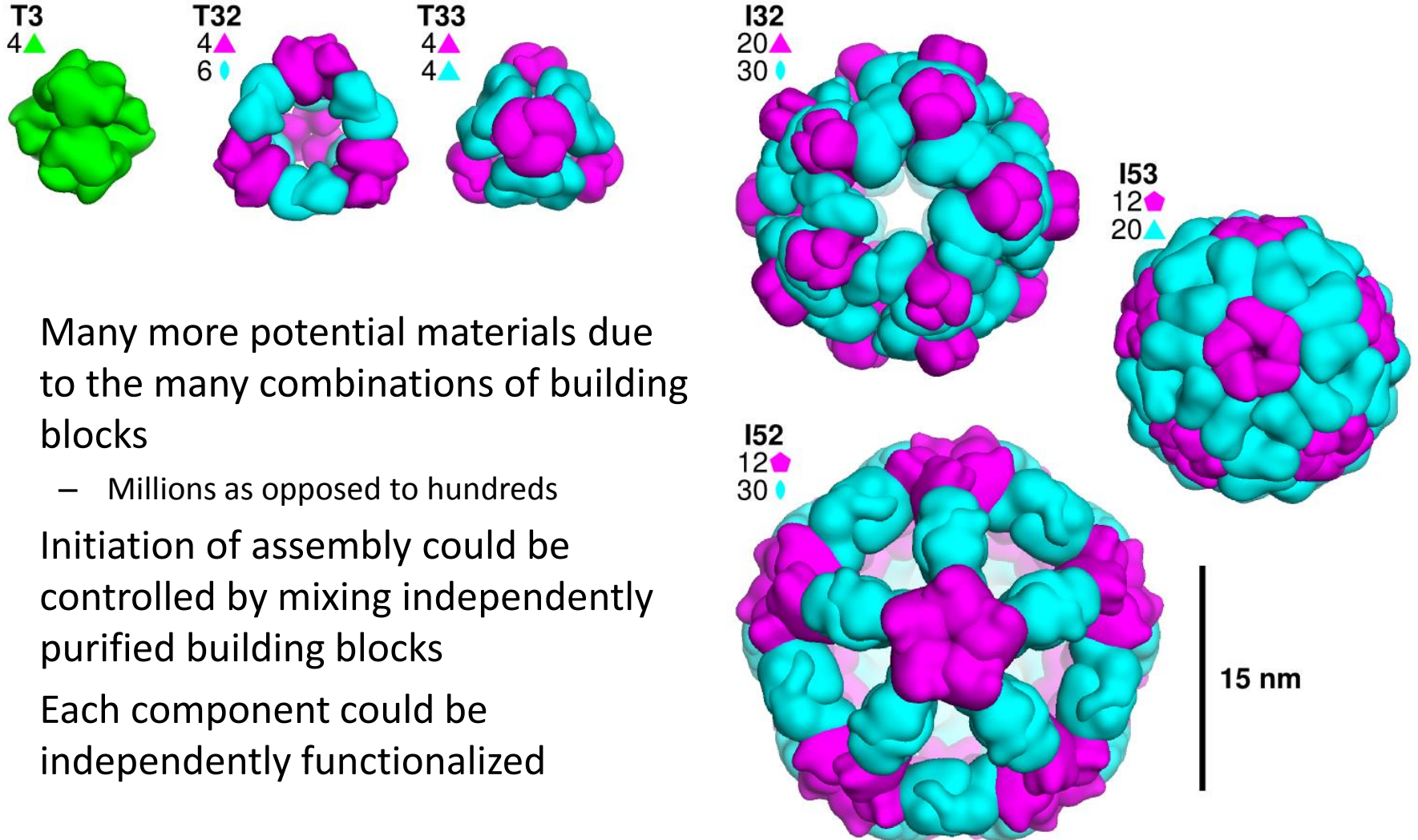
O3-33
Octahedral symmetry
8▲ (24 subunits), 480 kDa



The method enables the design of novel protein nanomaterials with atomic-level accuracy

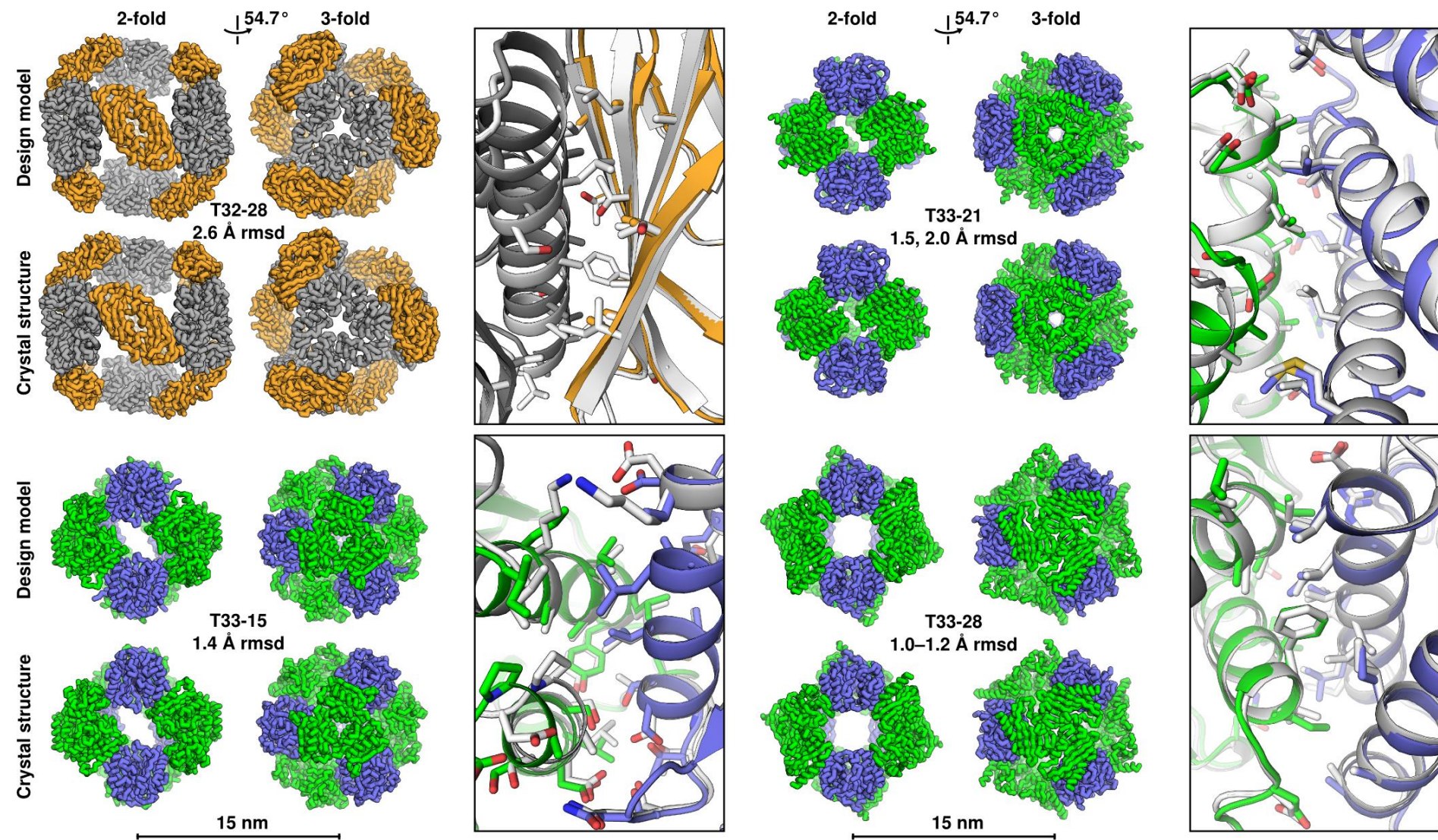


“Two-component” materials should be much more versatile for various applications

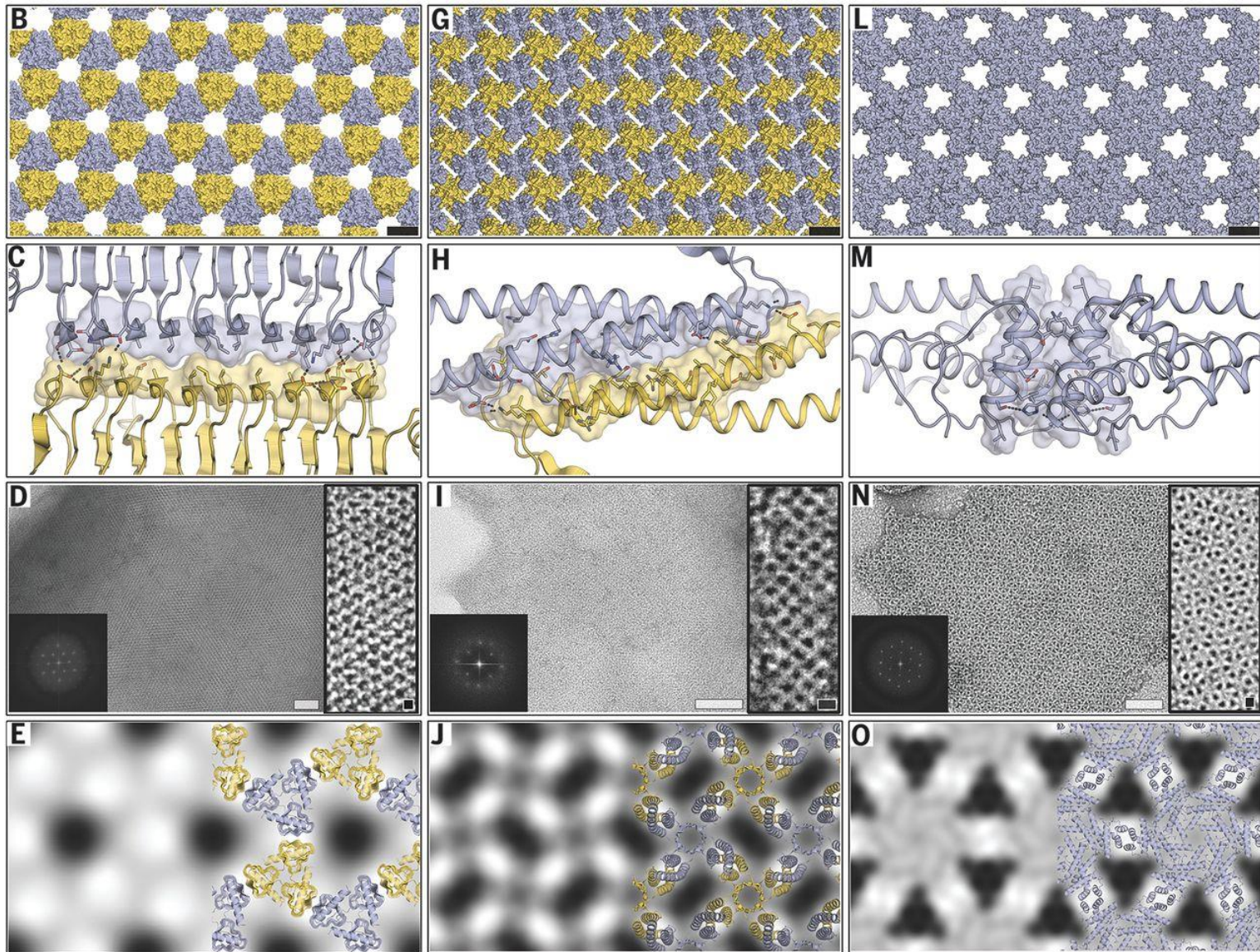


- Many more potential materials due to the many combinations of building blocks
 - Millions as opposed to hundreds
- Initiation of assembly could be controlled by mixing independently purified building blocks
- Each component could be independently functionalized

We extended the method to accurately design two-component co-assembling nanomaterials



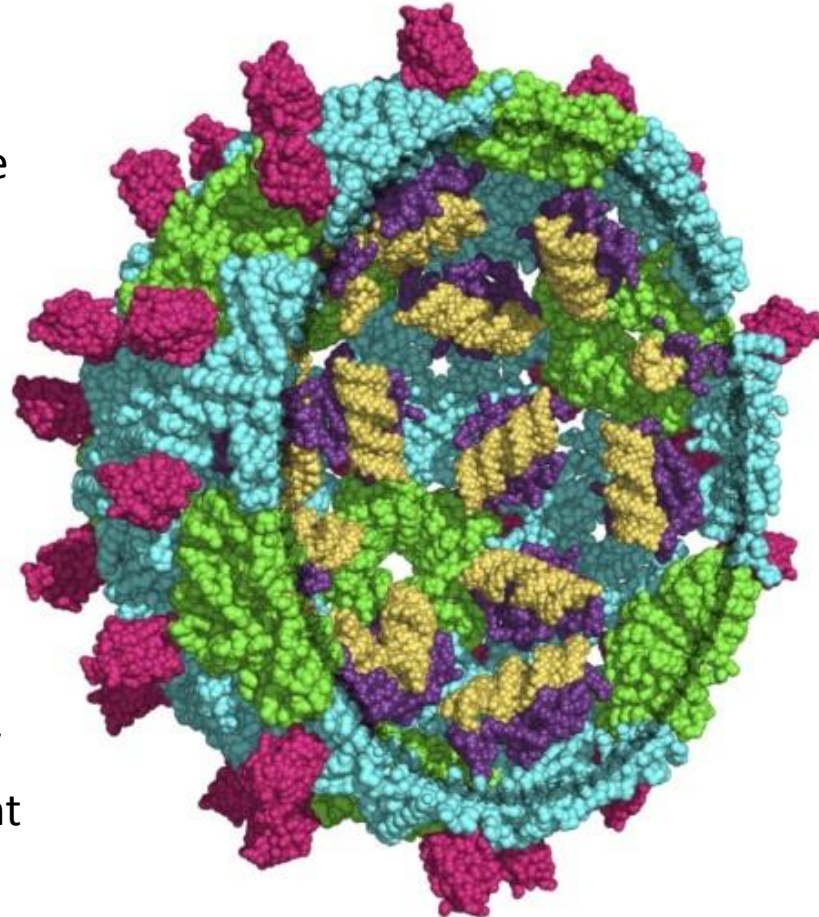
A wide variety of symmetric architectures can be designed using the approach (e.g., 2D layers)



Targeted drug delivery and nanoparticle vaccine design: twin applications for designed protein cages

Features:

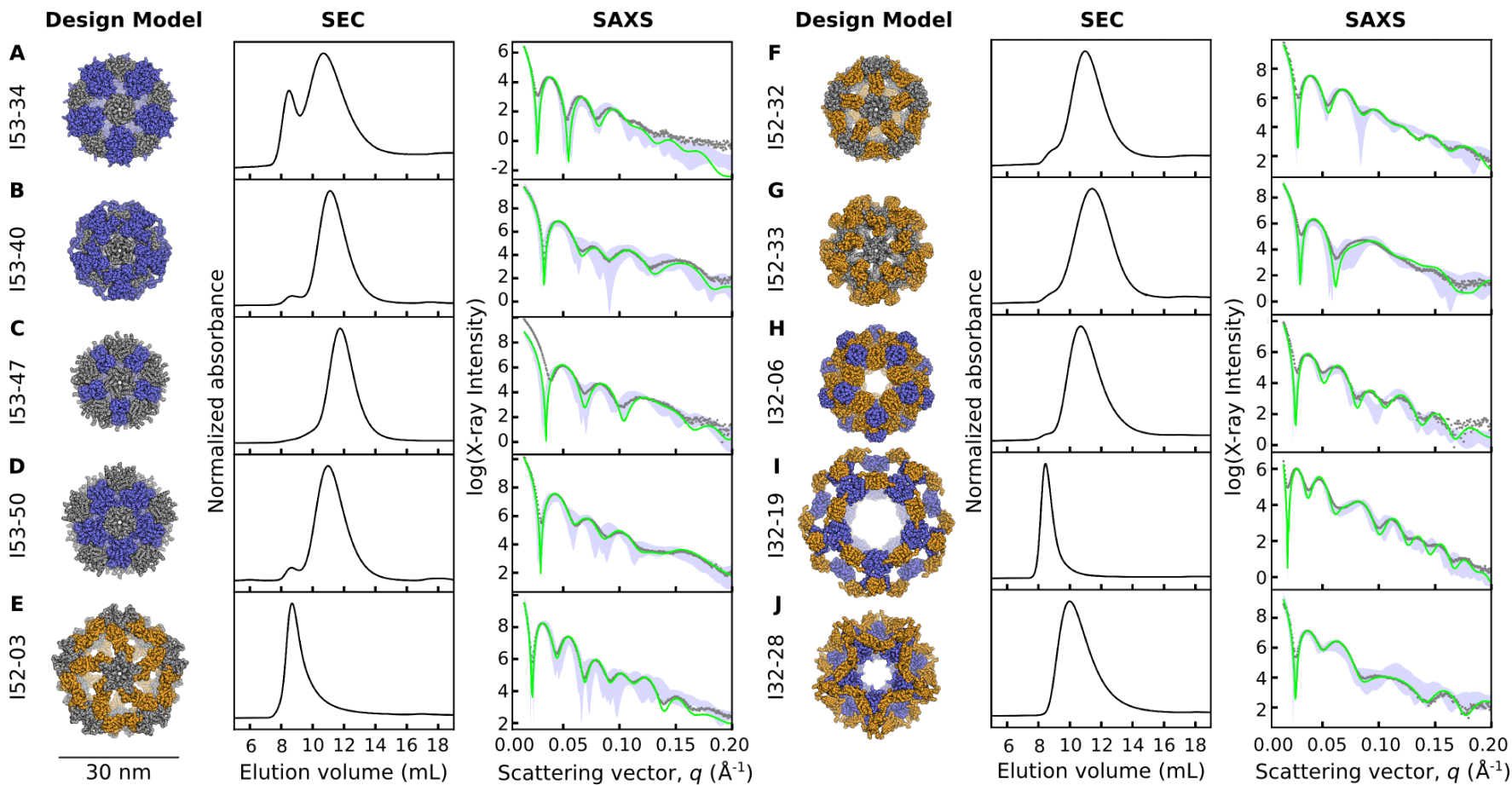
- ● Two-component cage for encapsulation & multivalent display
- Targeting domain or antigen/epitope
- Drug- or adjuvant-loading domain/residue/polymer
- Small molecule/xNA/protein drug/adjuvant



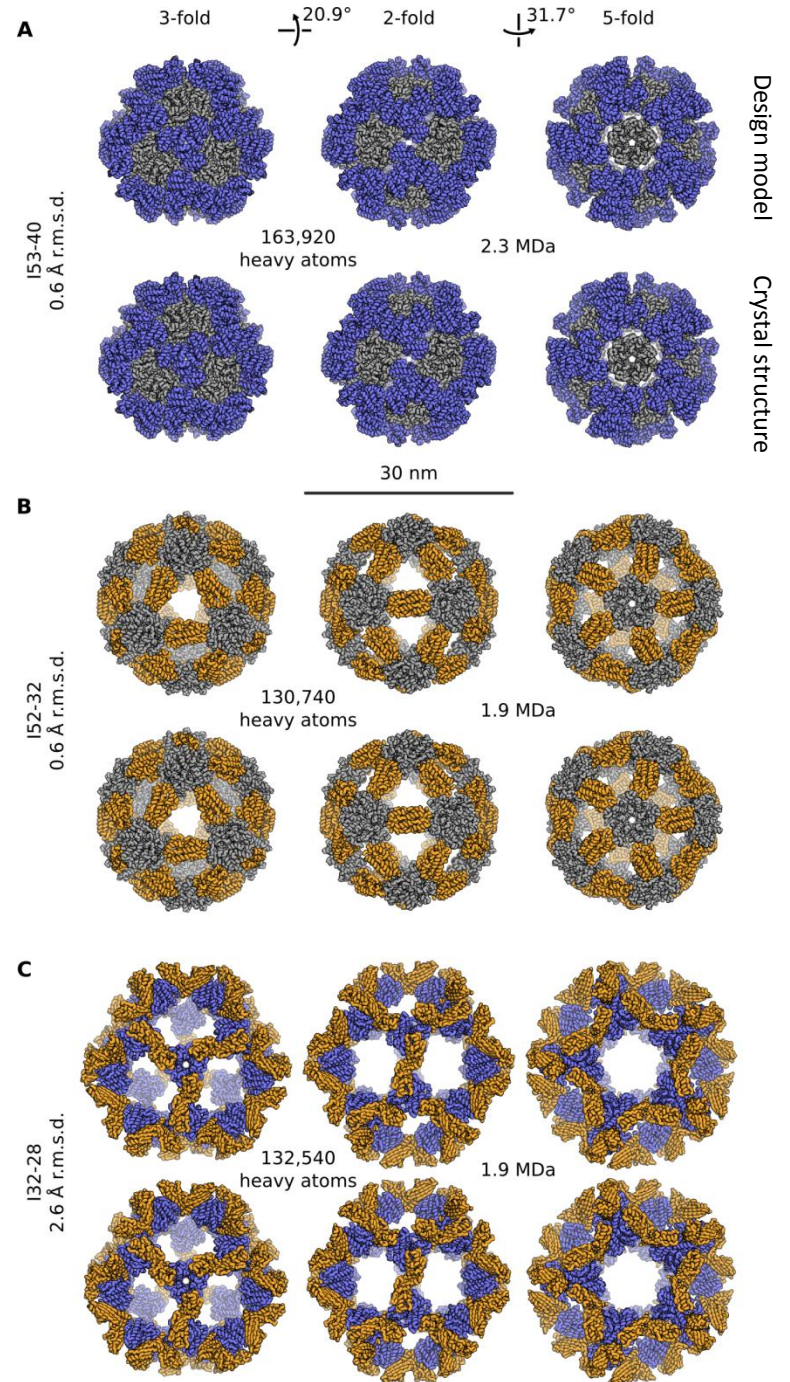
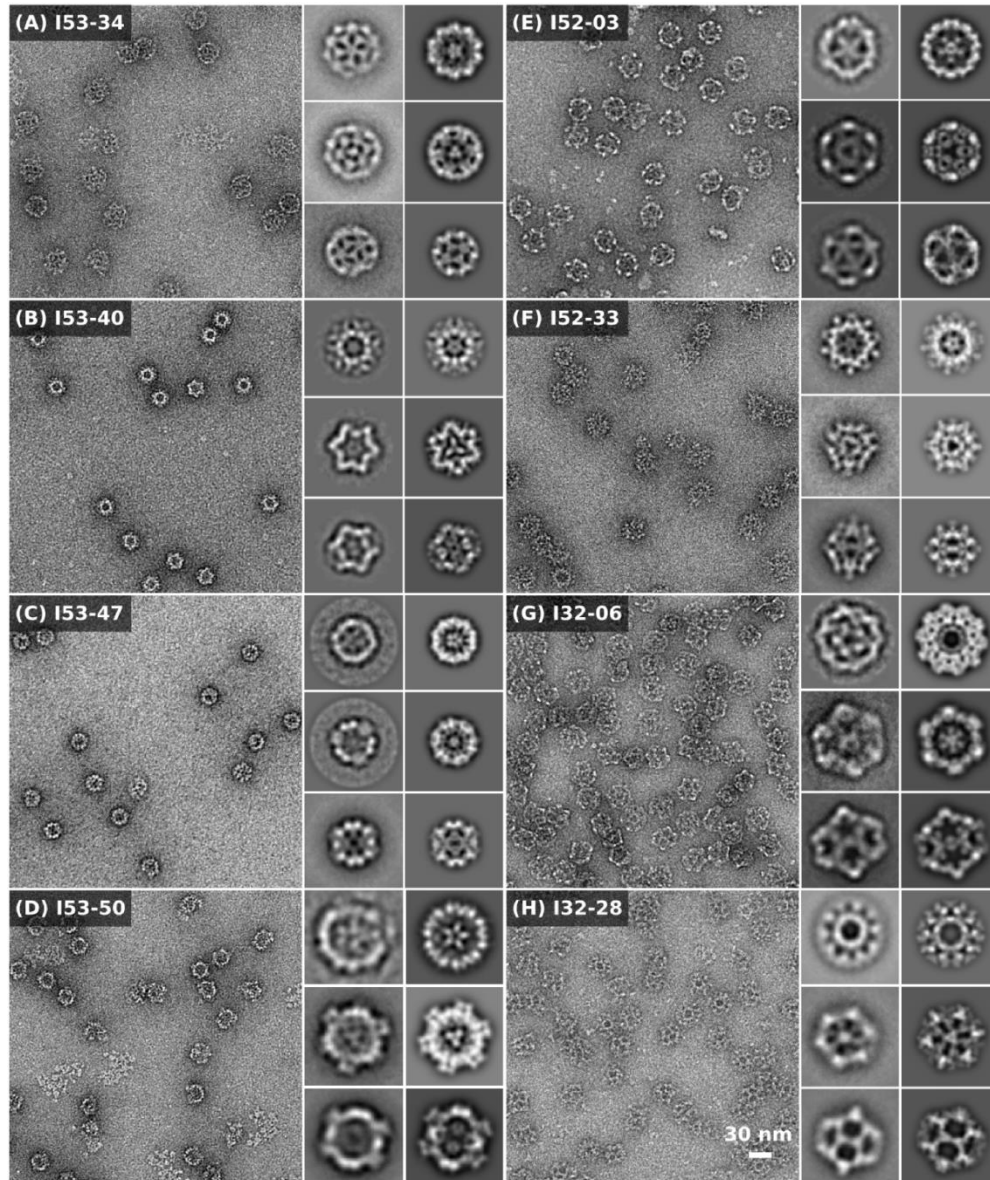
Additional potential features:

membranes, immune evasion/stimulation, environmental responsiveness, endosomal escape/subcellular localization, allostery, etc.

We have recently designed 12 two-component icosahedra, 10 of which are well-behaved, with large packaging capacities

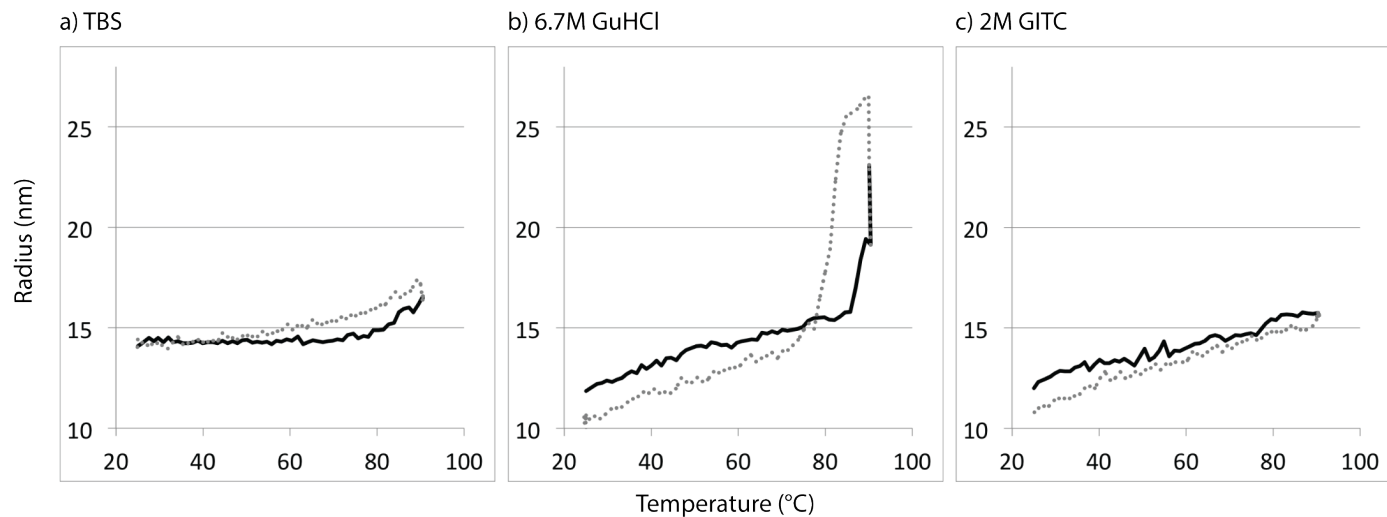


The 120-subunit, megadalton-scale structures were designed with atomic-level accuracy

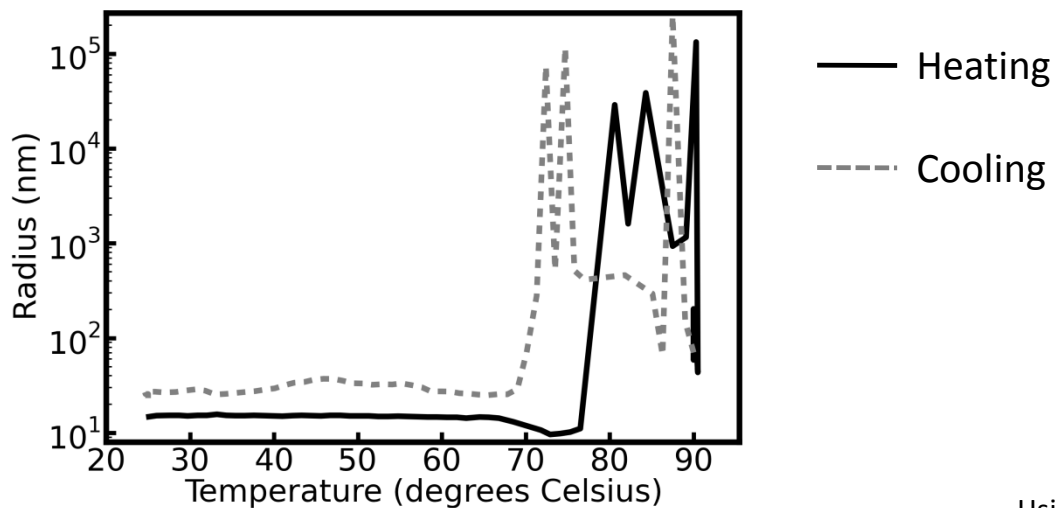


The nanoparticles are highly resistant to thermal stress

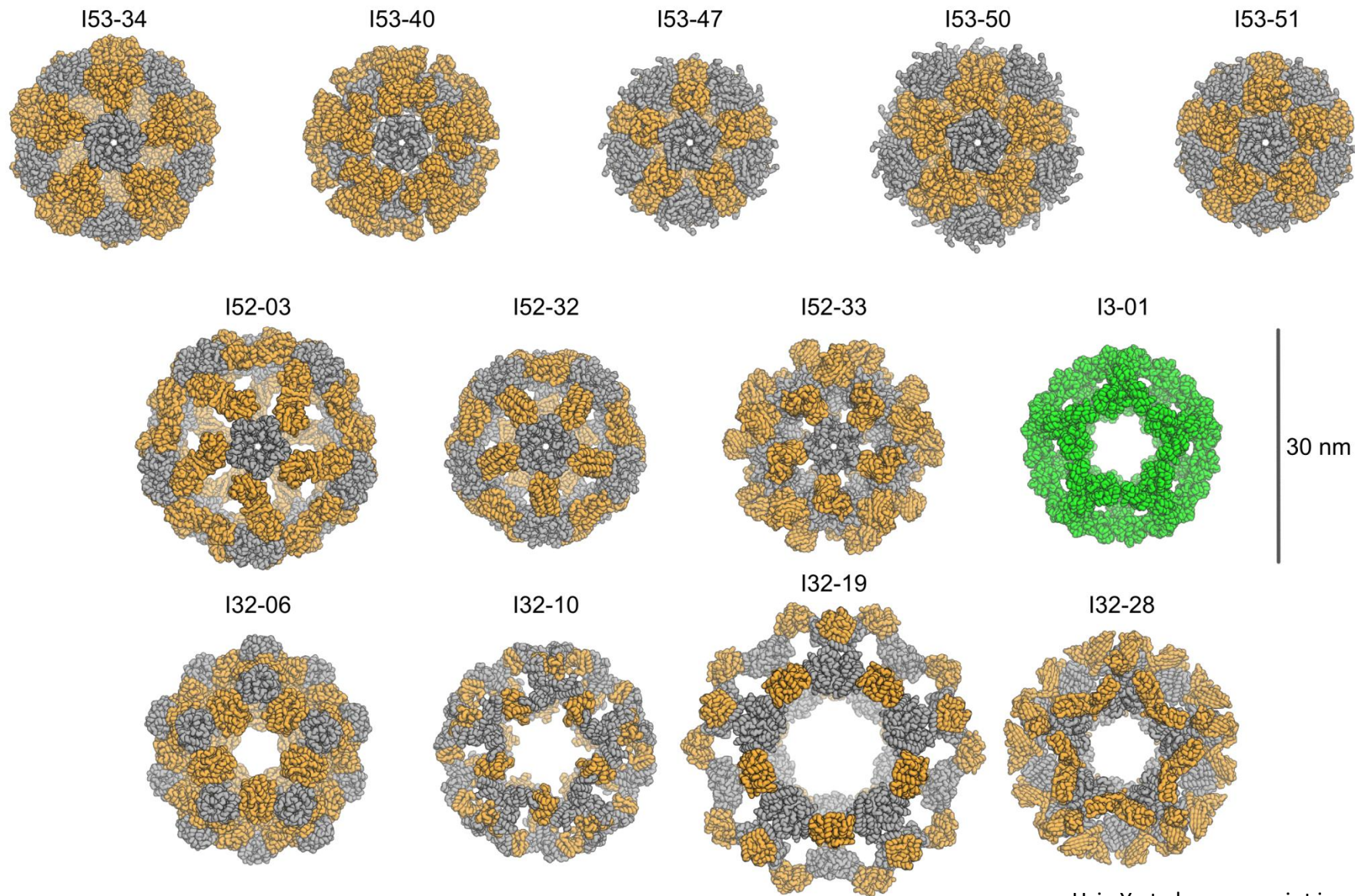
I3-01



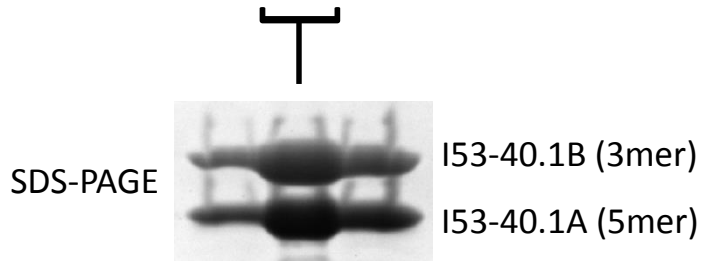
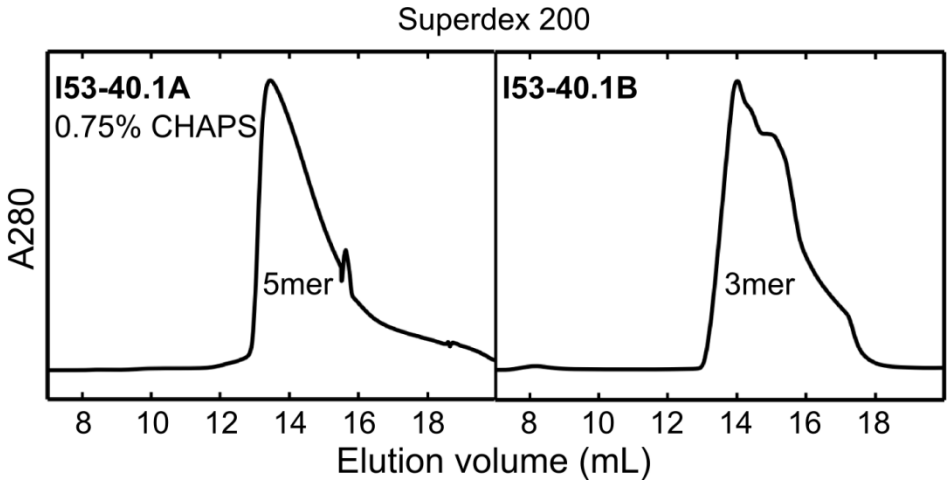
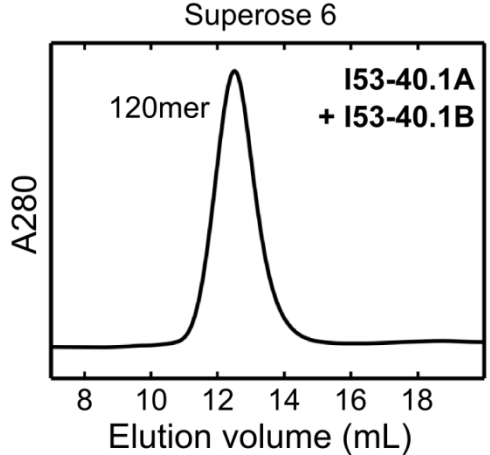
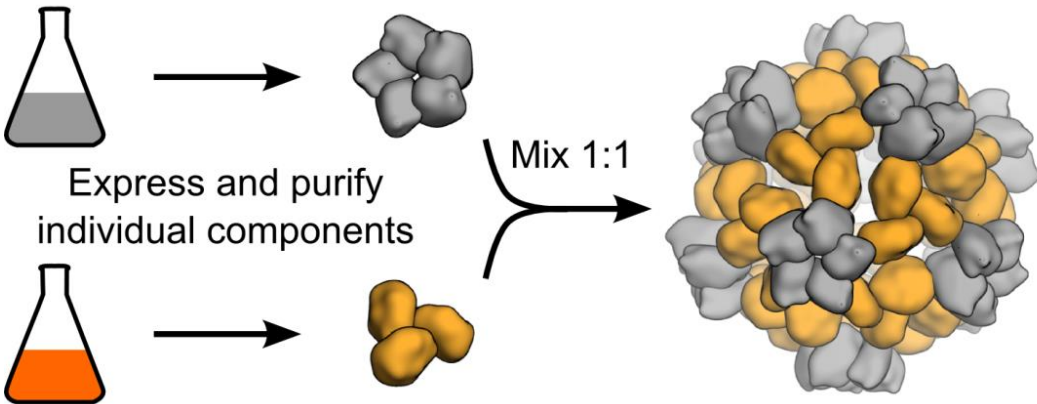
I53-50



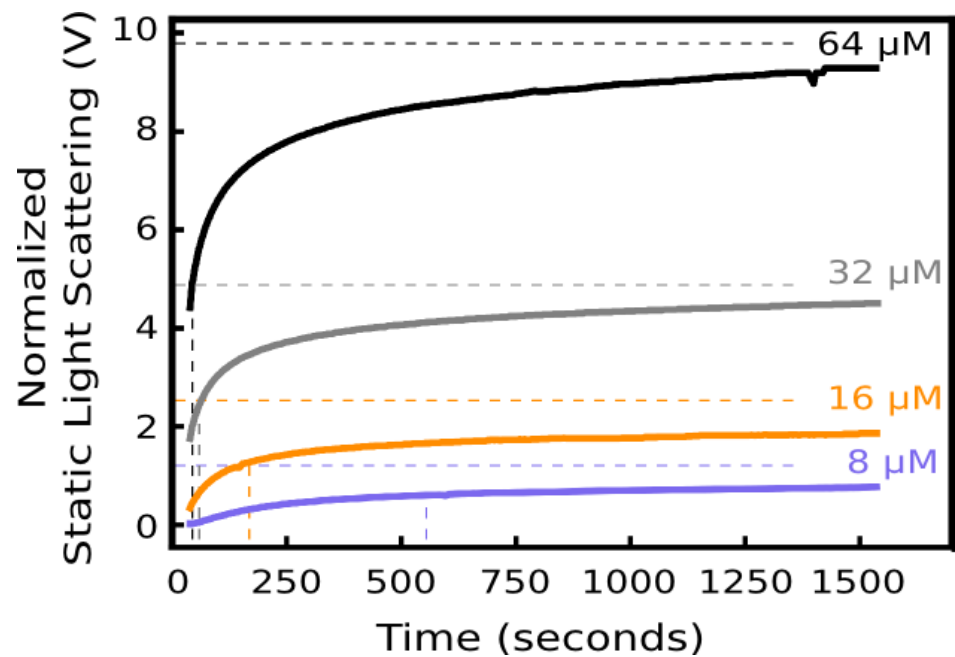
Our current crop of icosahedral nanoparticles



Mixing independently purified components enables simple, efficient, and controlled *in vitro* assembly



Assembly occurs on the timescale of seconds to minutes



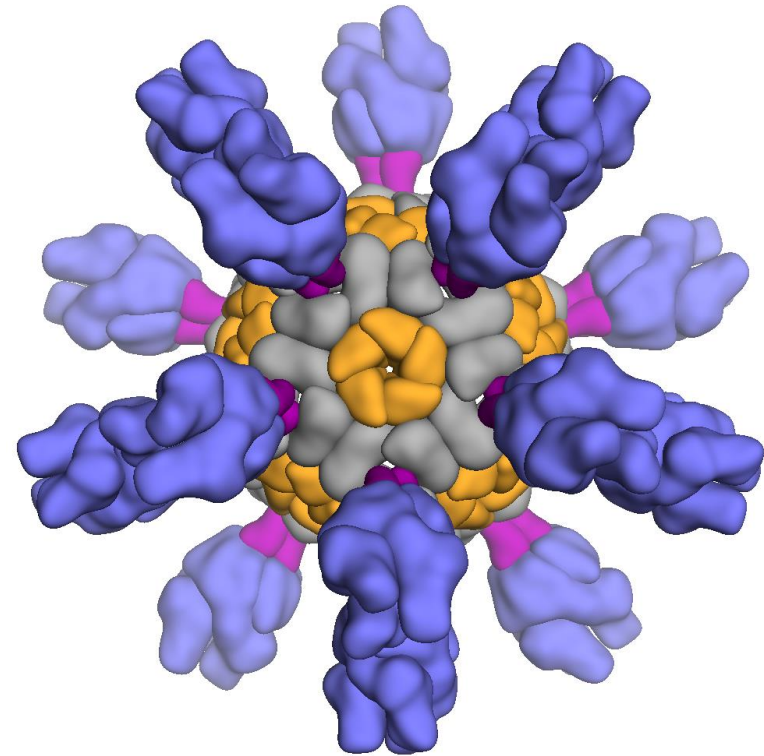
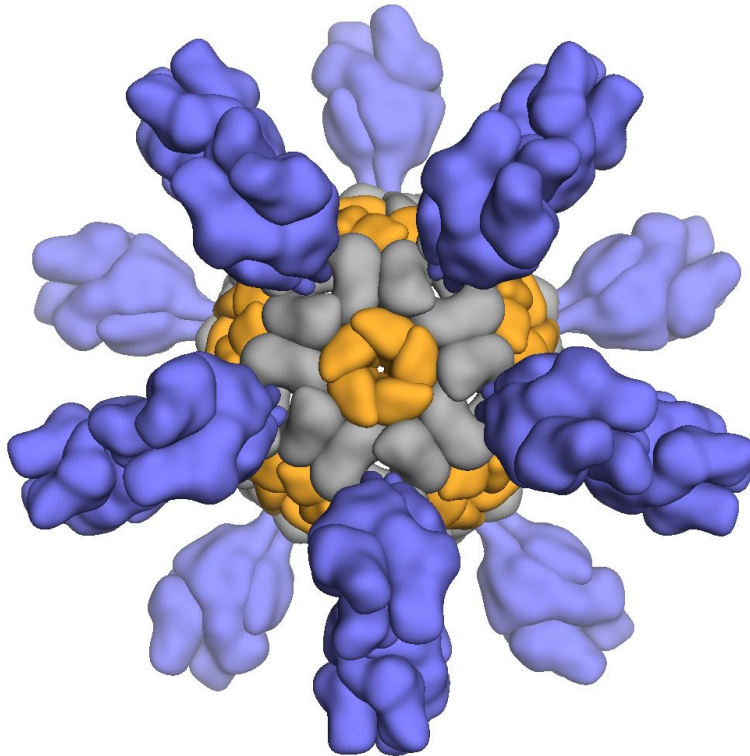
Two-component protein nanoparticles: a versatile platform for multivalent display

Component A

Component B

Functional domain

Adaptor protein complex



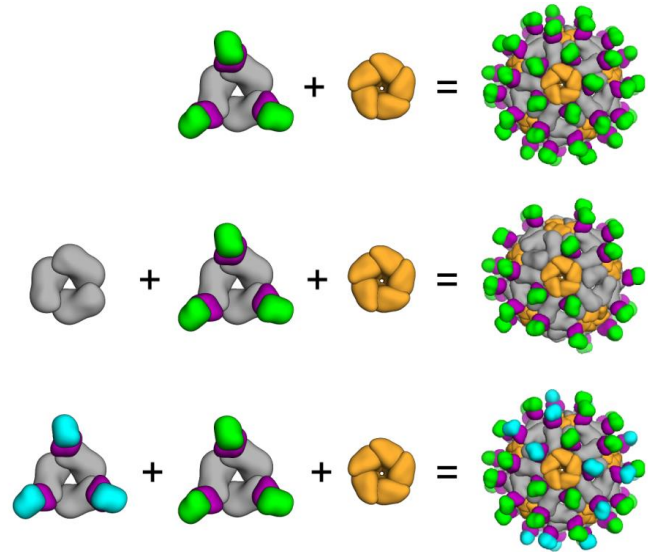
Functional domain expressed as a genetic fusion to cage component

Adaptor protein mediates attachment to nanoparticle components

Two-component protein nanoparticles: a versatile platform for multivalent display

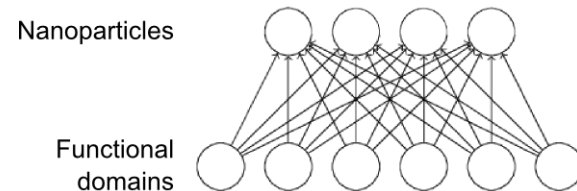
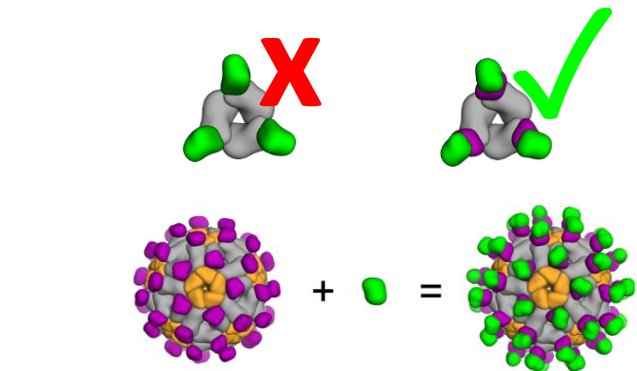
In vitro assembly

- Expression, purification, and quality control can be performed independently on distinct building blocks
- Antigen valency/copy number can be controlled by including unmodified components during *in vitro* assembly
- Distinct antigens/costimulatory proteins can be scaffolded in defined ratios



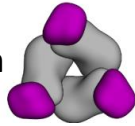
Adaptor-mediated labeling

- Complex antigens that fail as genetic fusions to nanoparticle subunits can be labeled with a small adaptor tag
- Post-assembly labeling
- Rapid prototyping of functional domain/nanoparticle combinations

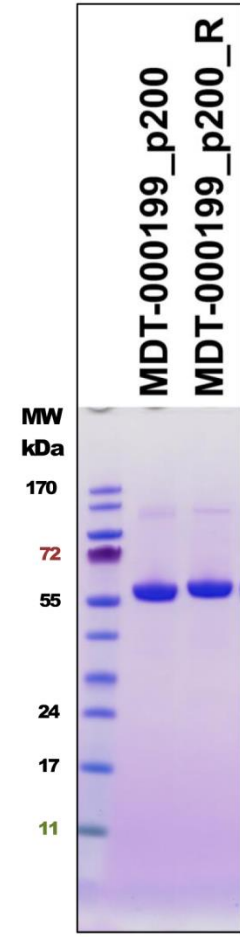
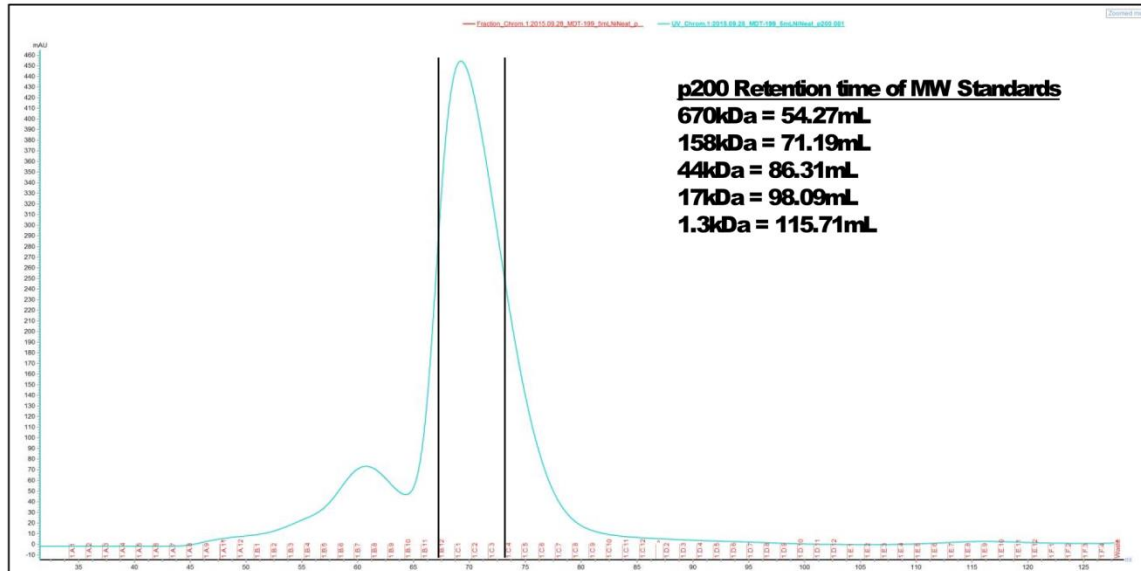


An anti-CD20 scFv-I53-50A fusion protein can be produced in good yield

scFv-trimer genetic fusion

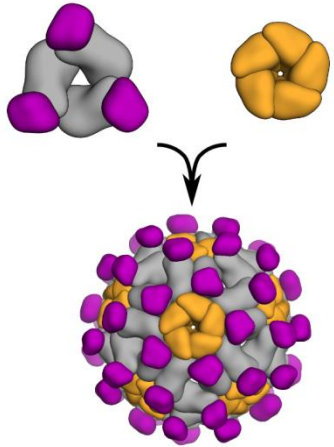


Superdex 200 16/600 pg

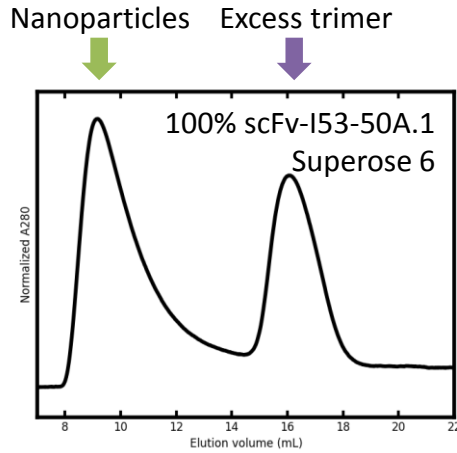


In vitro assembly allows control over scFv valency

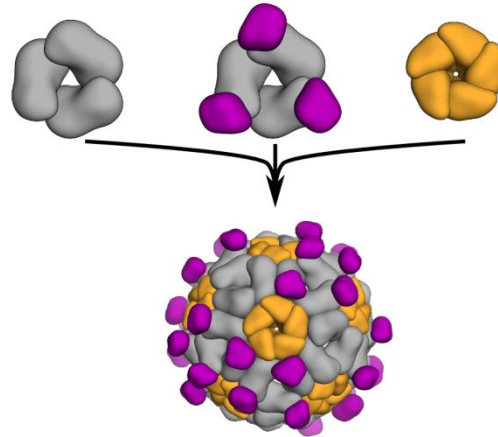
100% scFv-trimer



Full-valency nanoparticle

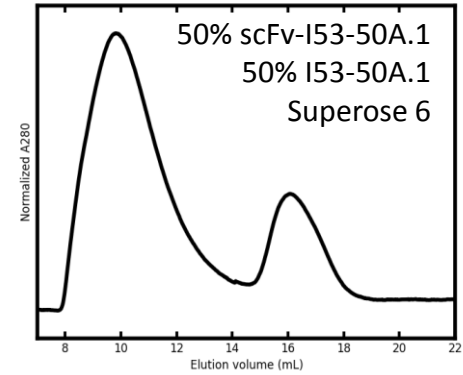


50% 50% scFv-trimer

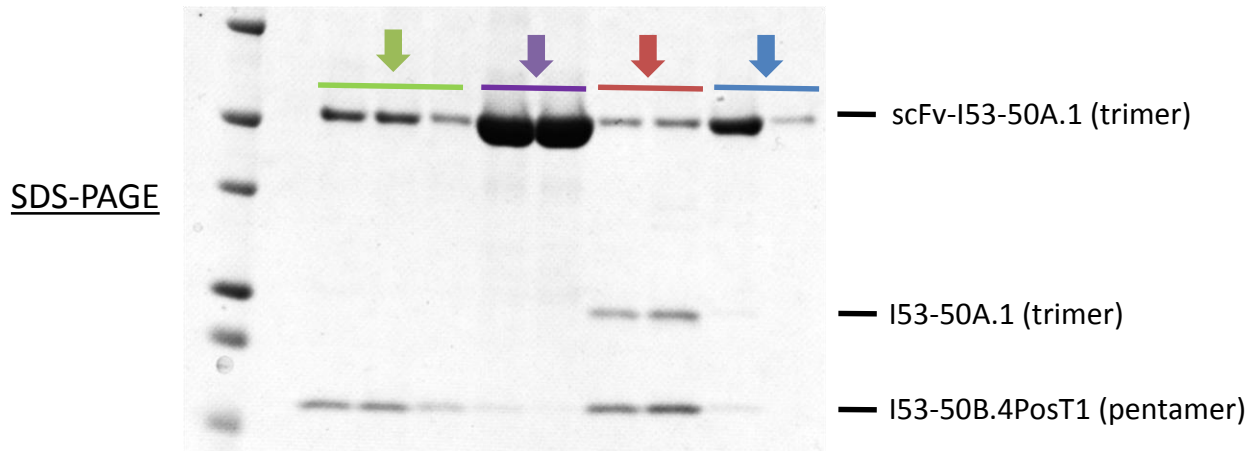


Half-valency nanoparticle

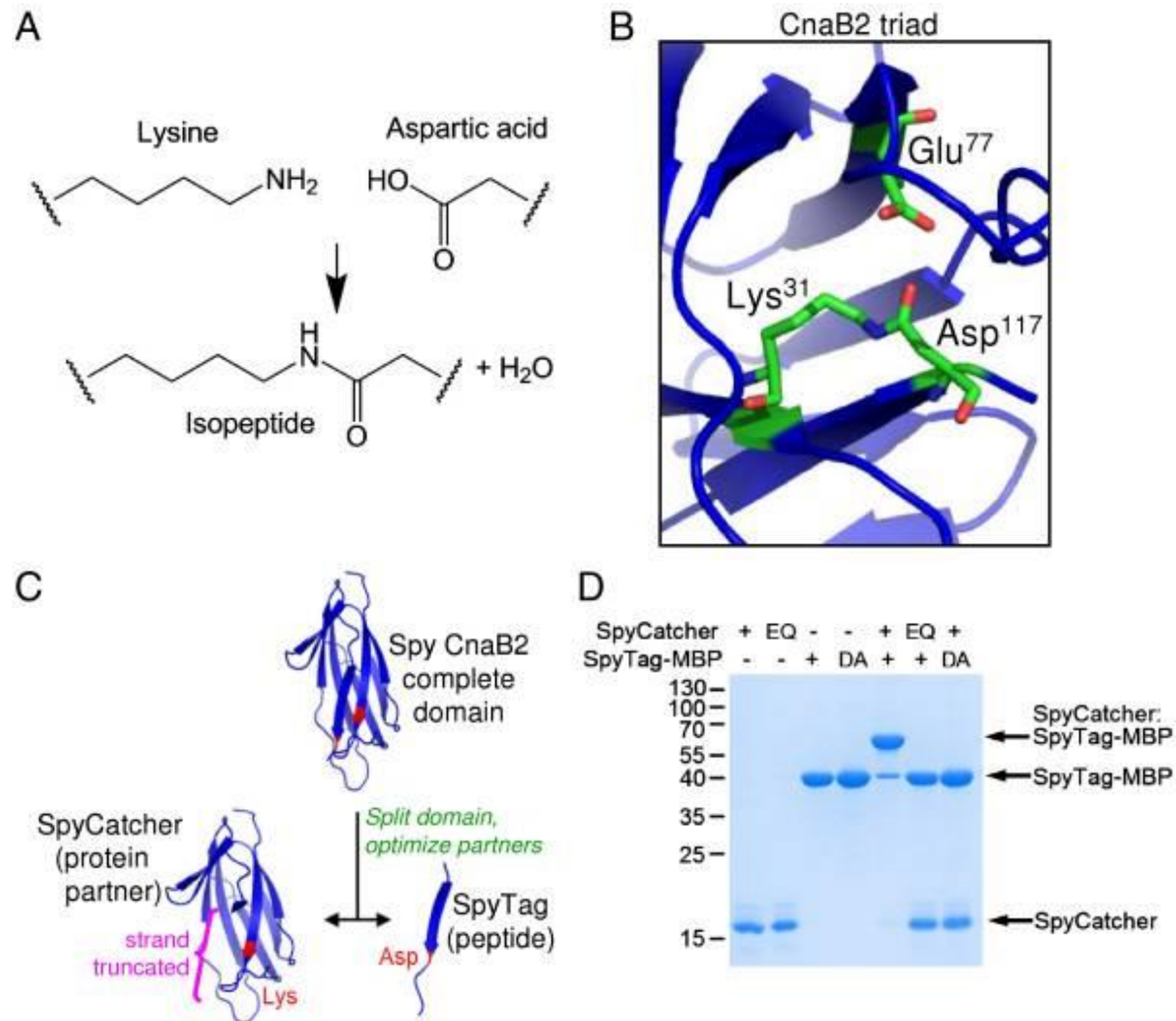
Nanoparticles Excess trimer



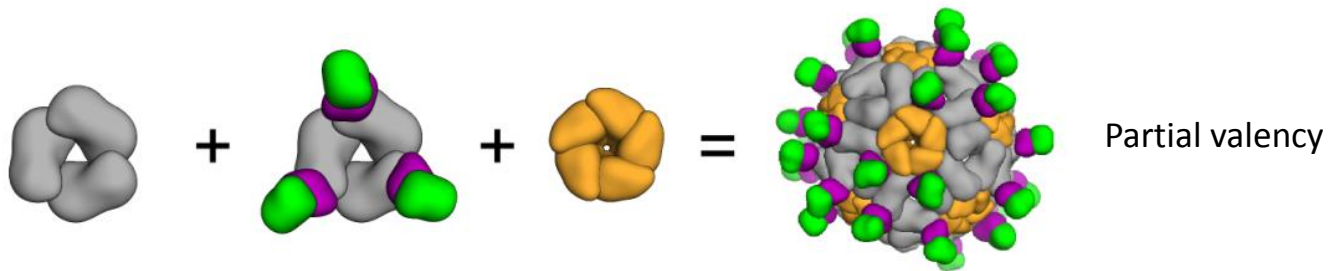
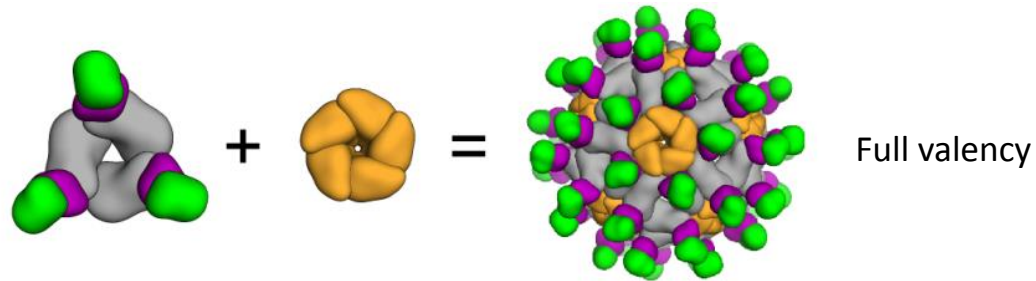
100% scFv-trimer 50% scFv-trimer



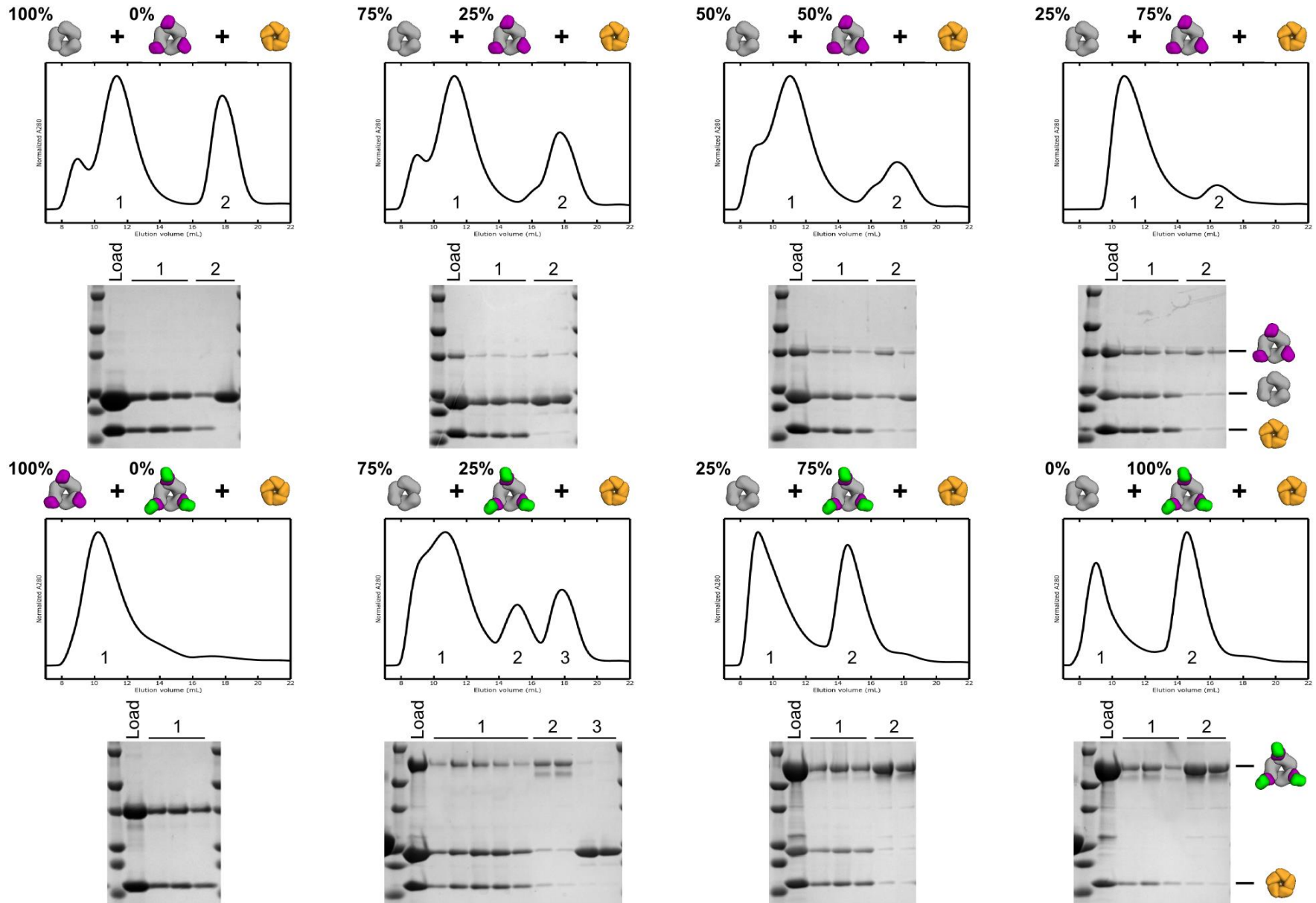
SpyCatcher-SpyTag is a molecular adaptor capable of selective and stable labeling



Conjugating SpyTag-GFP to purified SpyCatcher-I53-50A enables *in vitro* assembly of nanoparticles with variable GFP valencies



Conjugating SpyTag-GFP to purified SpyCatcher-I53-50A enables *in vitro* assembly of nanoparticles with variable GFP valencies



Summary and future directions

Summary:

- We have developed a general computational approach to designing self-assembling protein nanomaterials with atomic-level accuracy
- We have recently designed and experimentally validated 120-subunit icosahedral nanoparticles with sizes and molecular weights comparable to small viruses
- We have demonstrated the multivalent display of complex proteins (e.g., scFvs, viral envelope glycoproteins) on the nanoparticles using both direct genetic fusion and molecular adaptors
- The designed nanoparticles boost the immunogenicity of a multivalently displayed peptide antigen comparably to RNA-containing bacteriophage particles in mice

Future directions:

- Need to obtain additional immunogenicity data on nanoparticles bearing antigens of interest
- Further modify antigen-bearing nanoparticles to co-package adjuvants to increase/tailor immune response
- Explore possibilities afforded by two-component nanoparticles to display multiple antigens or combinations of antigens and costimulatory proteins

Acknowledgements

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