
Engineering Polymer-Protein Interactions to Control the Polymersome Protein Corona

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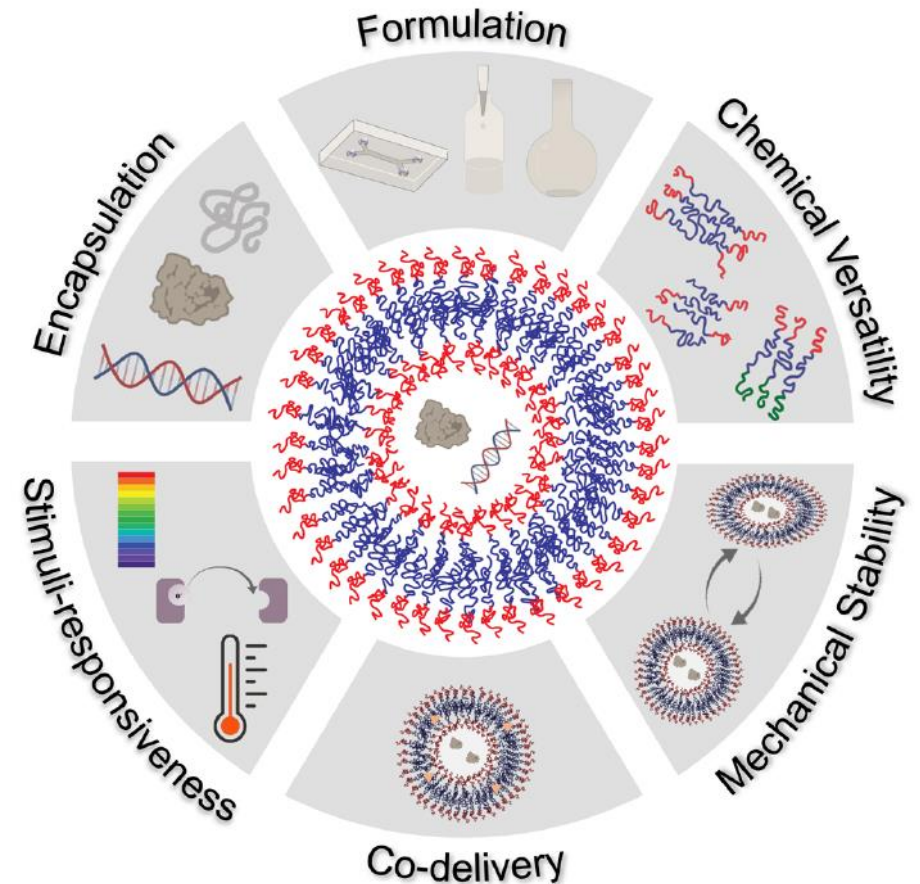


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Polymersomes are versatile drug delivery vehicles

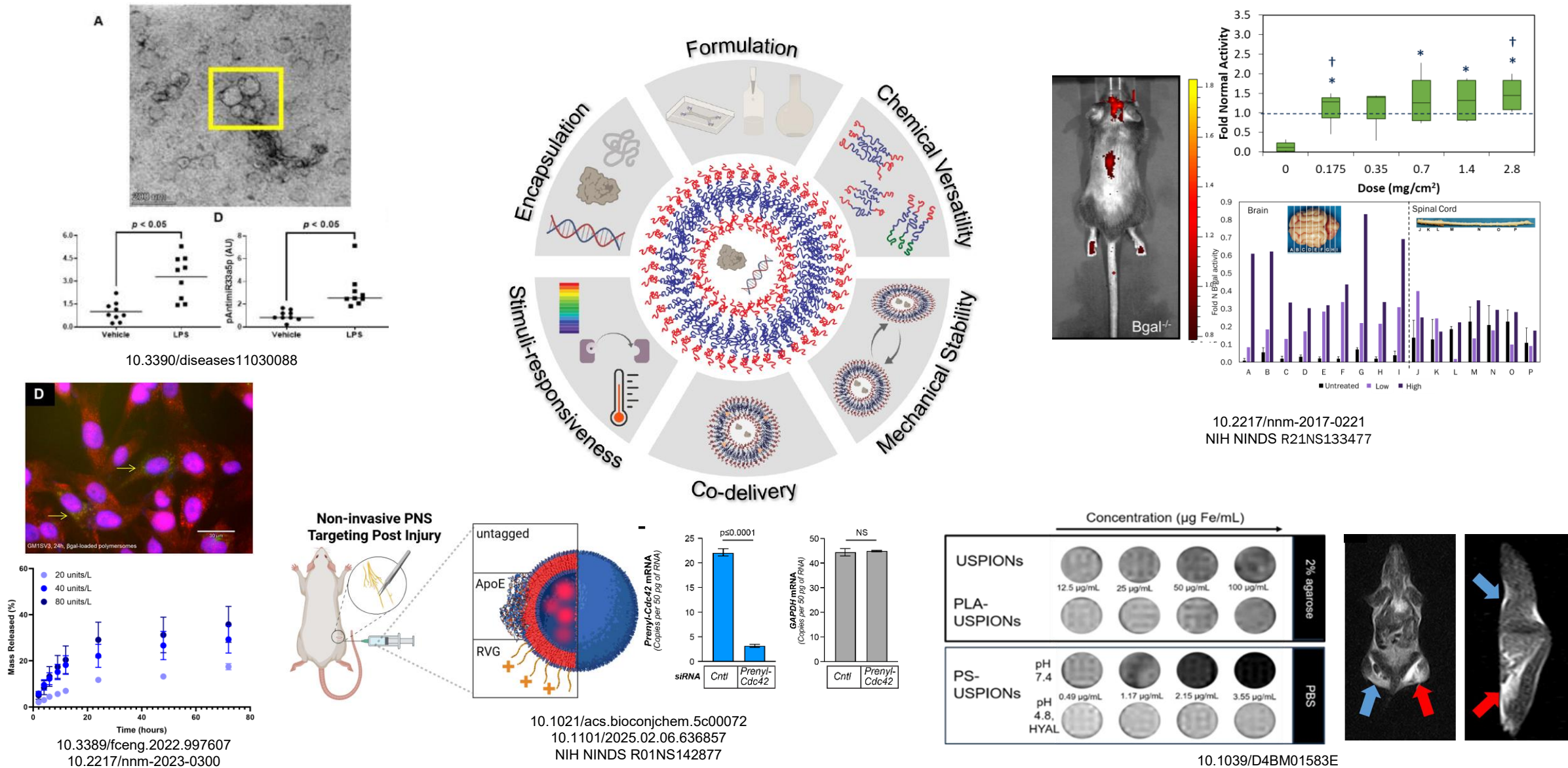
Polymersomes are nanoparticle vesicles developed from the formation of amphiphilic block copolymers

- + Self assemble into vesicles under with hydrophilic fraction between 25 and 45%
- + Clinically approved polymers
- + Tunable properties
- + High MW payloads or multiple payloads

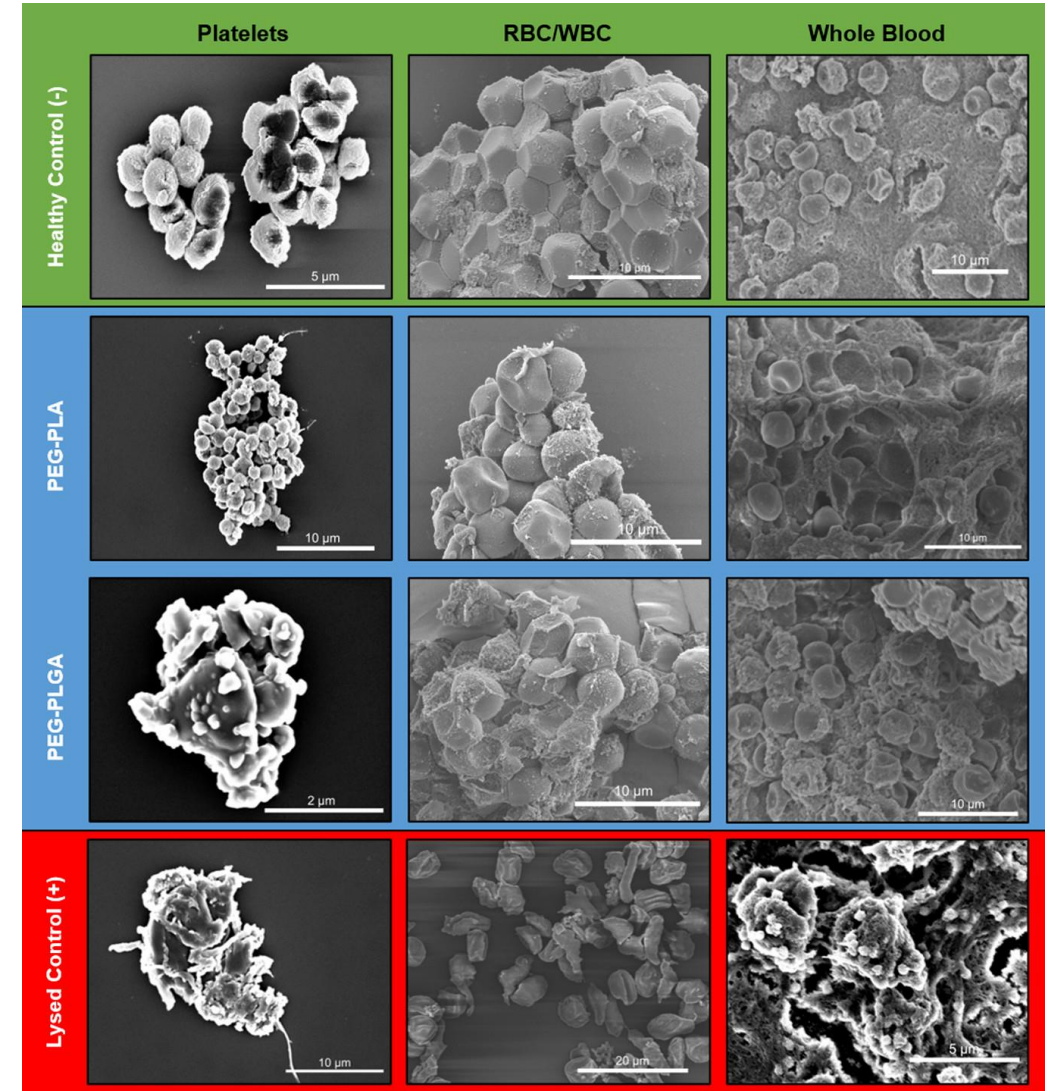
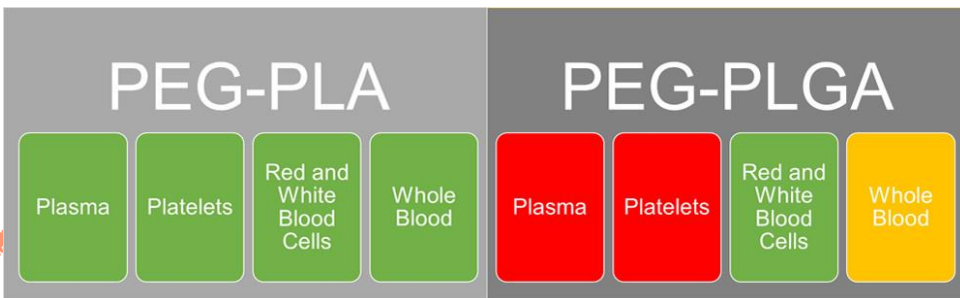
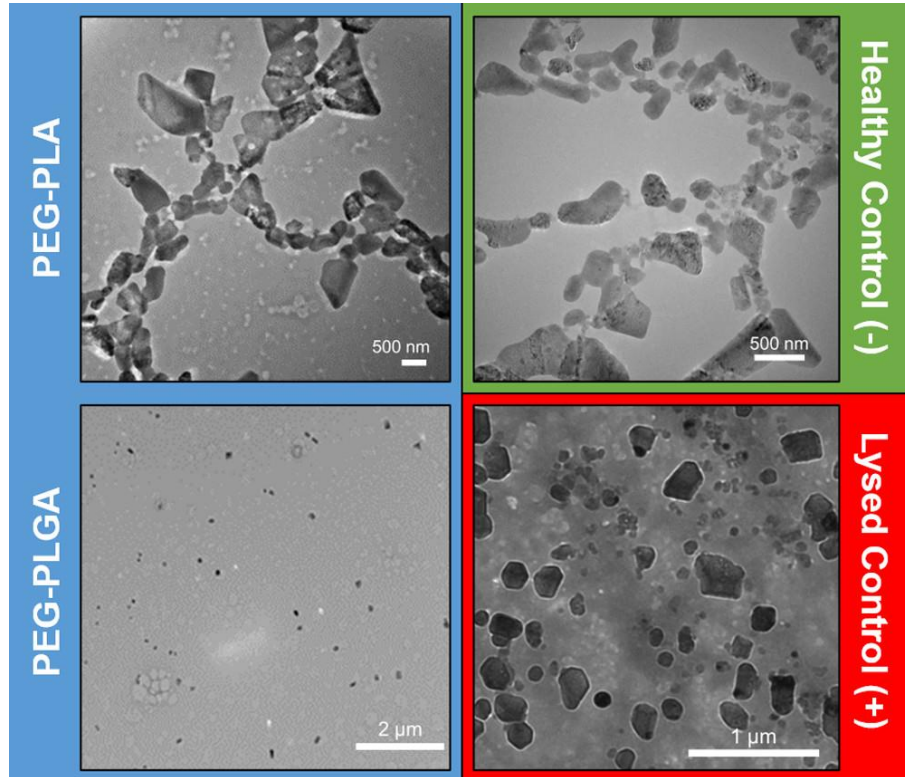
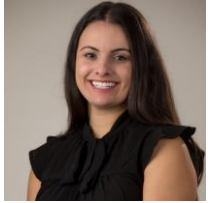


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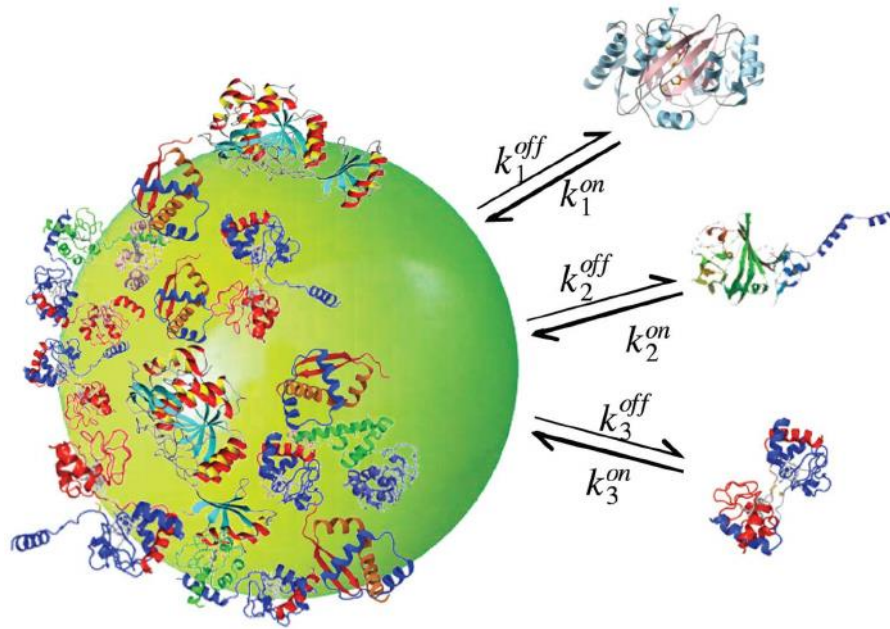
Polymersomes are versatile drug delivery vehicles



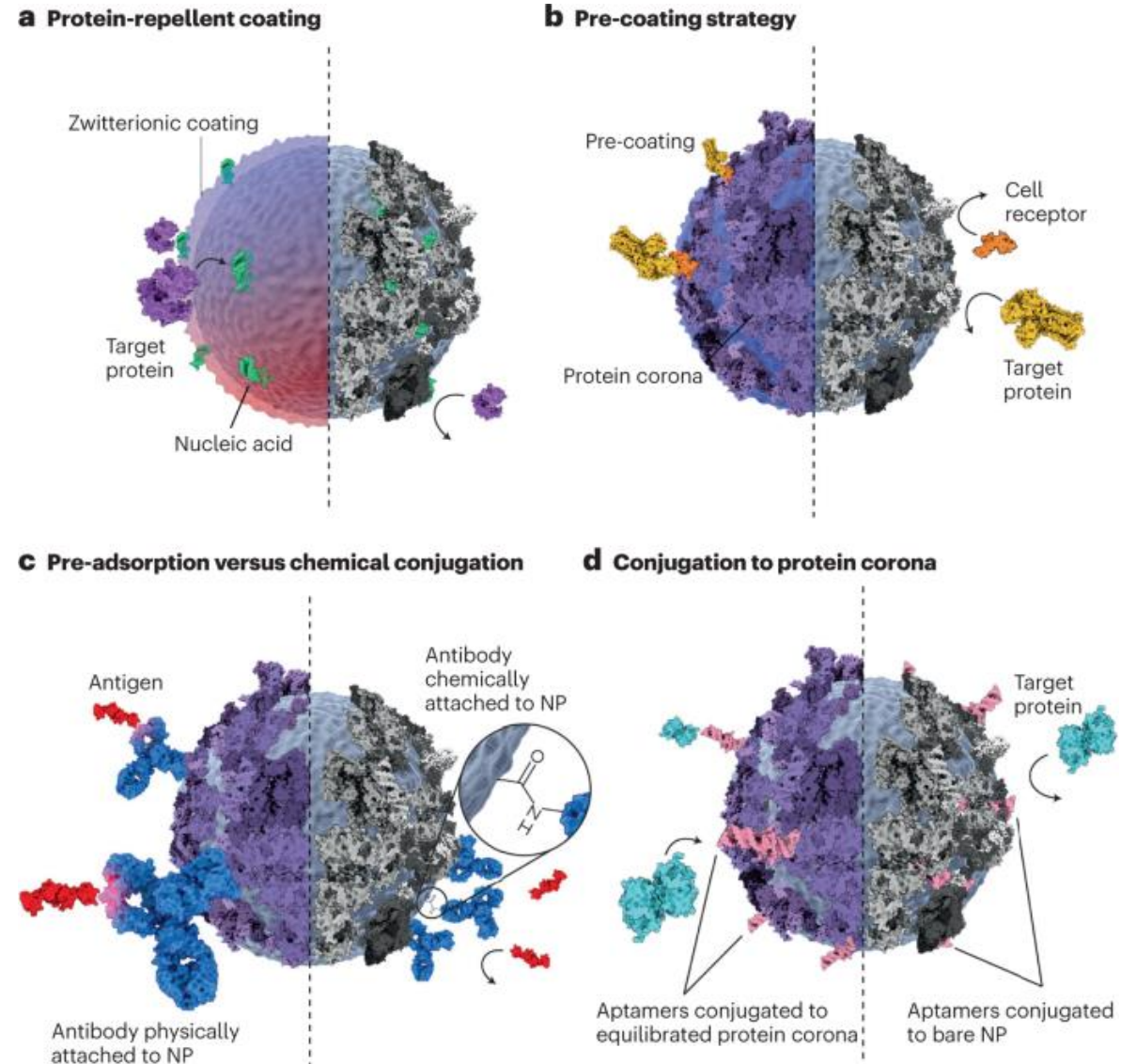
Polymersomes can be effective but are also synthetic.



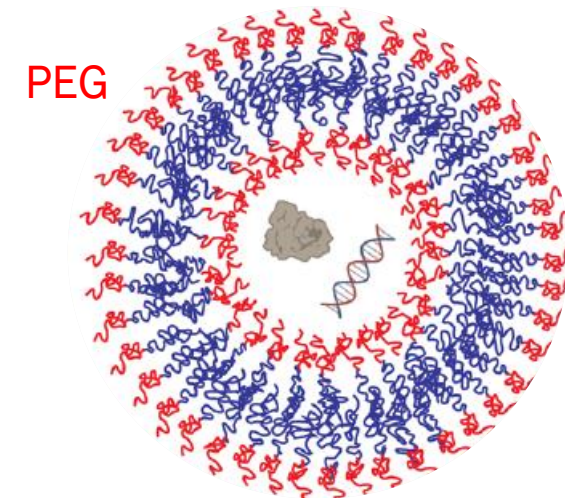
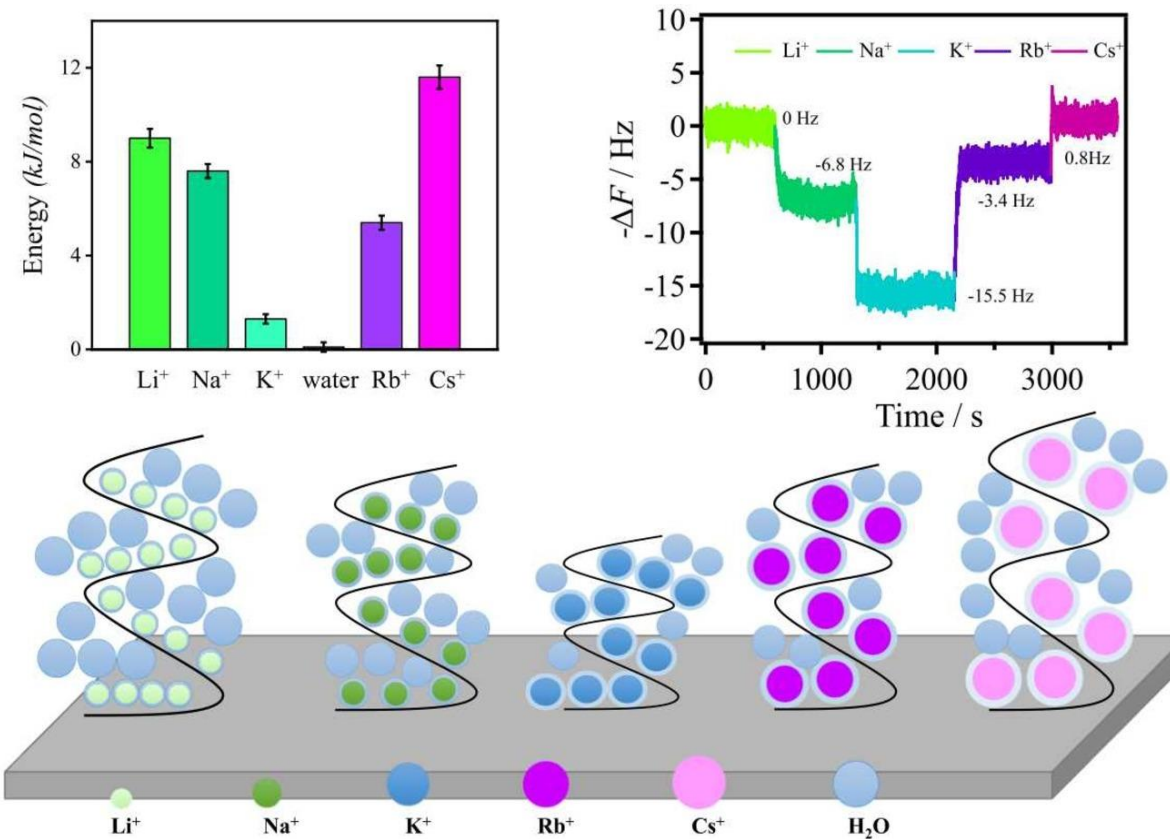
A protein corona forms around polymersomes upon injection.



Biomolecular identity of nanoparticles changes upon injection, which impacts the cellular interactions.



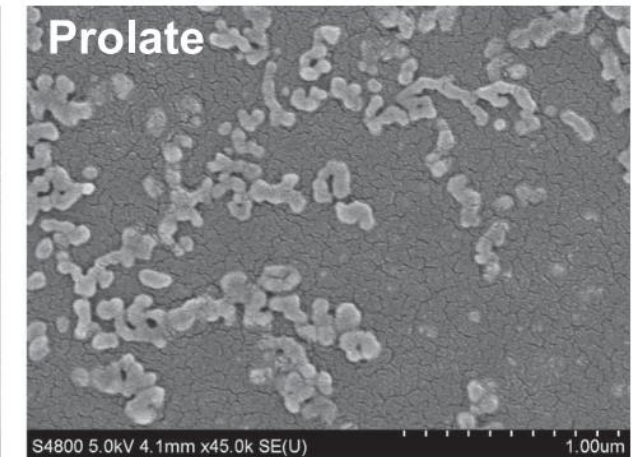
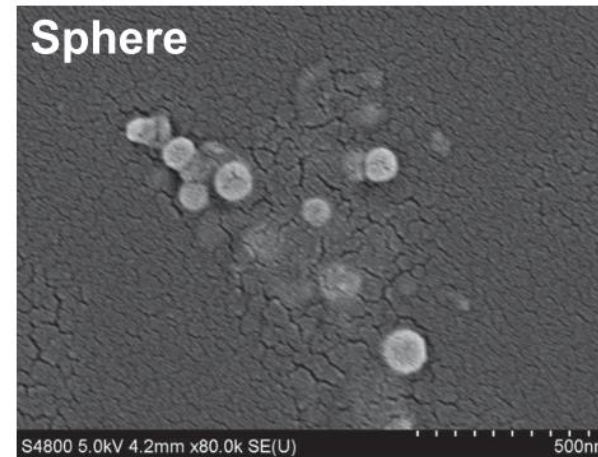
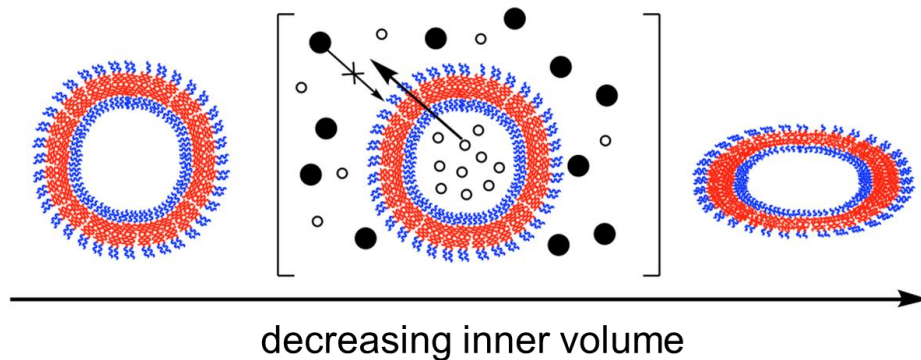
Cations impact polyethylene glycol conformation and hydration.



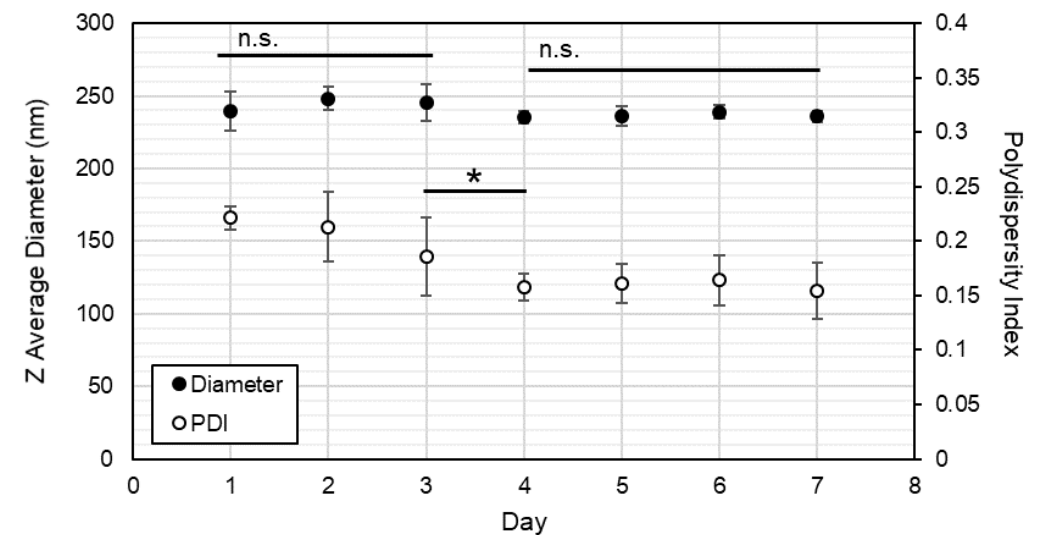
Yu, et al. Colloids and Surfaces A, 2024.



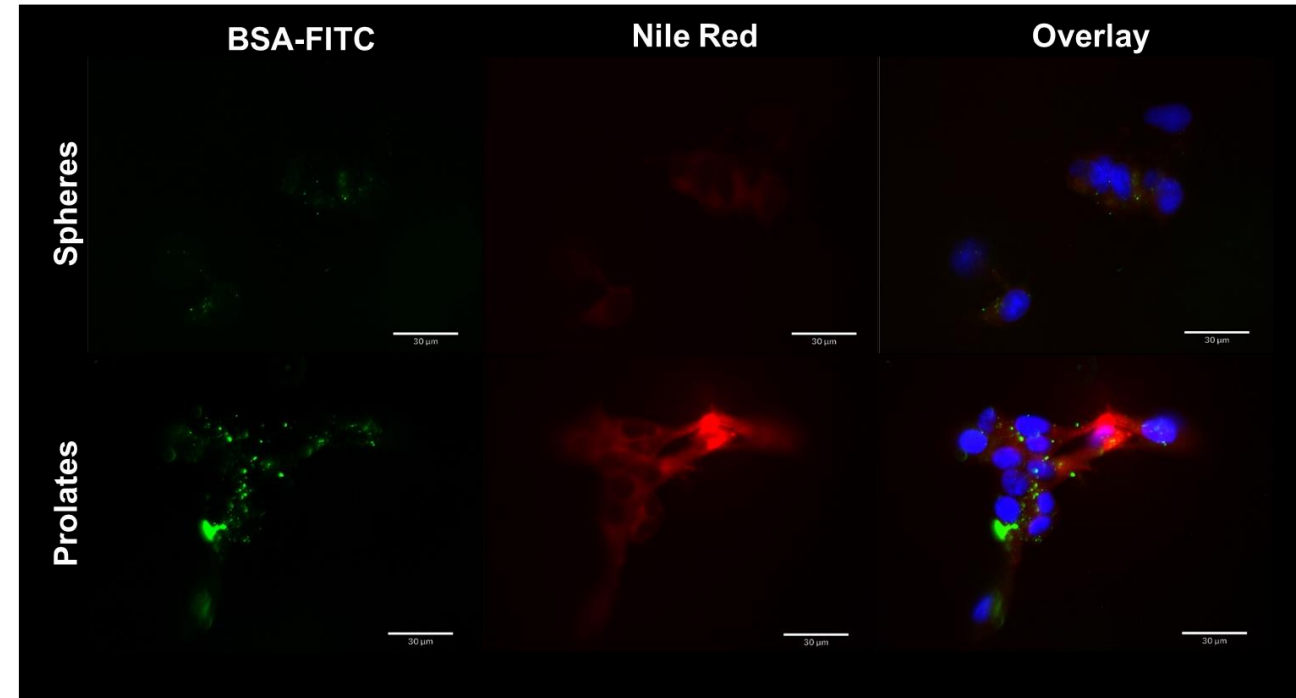
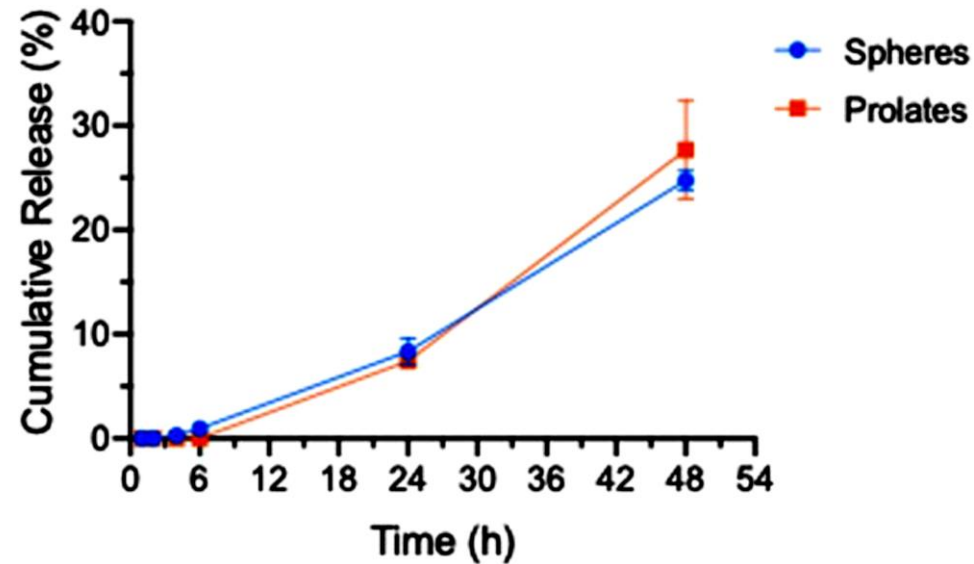
PEGPLA Polymersomes are Modulated using Osmotic Pressure



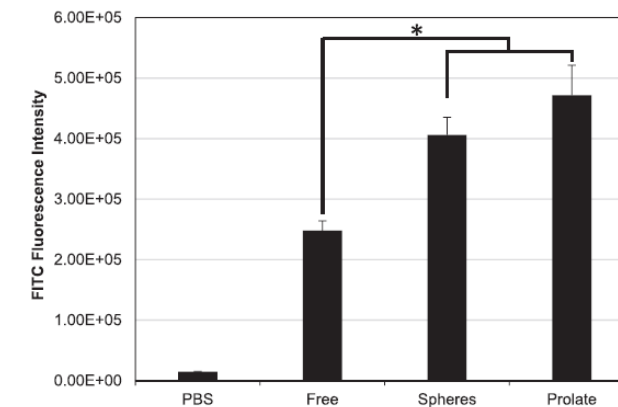
NaCl	Diameter $\pm$ SD (nm)	Zeta Potential $\pm$ SD (mV)	PDI $\pm$ SD
-	145.5 $\pm$ 6.1	-41.6 $\pm$ 1.7	0.05 $\pm$ 0.02
+	190.9 $\pm$ 1.3	-19.3 $\pm$ 1.6	0.12 $\pm$ 0.01



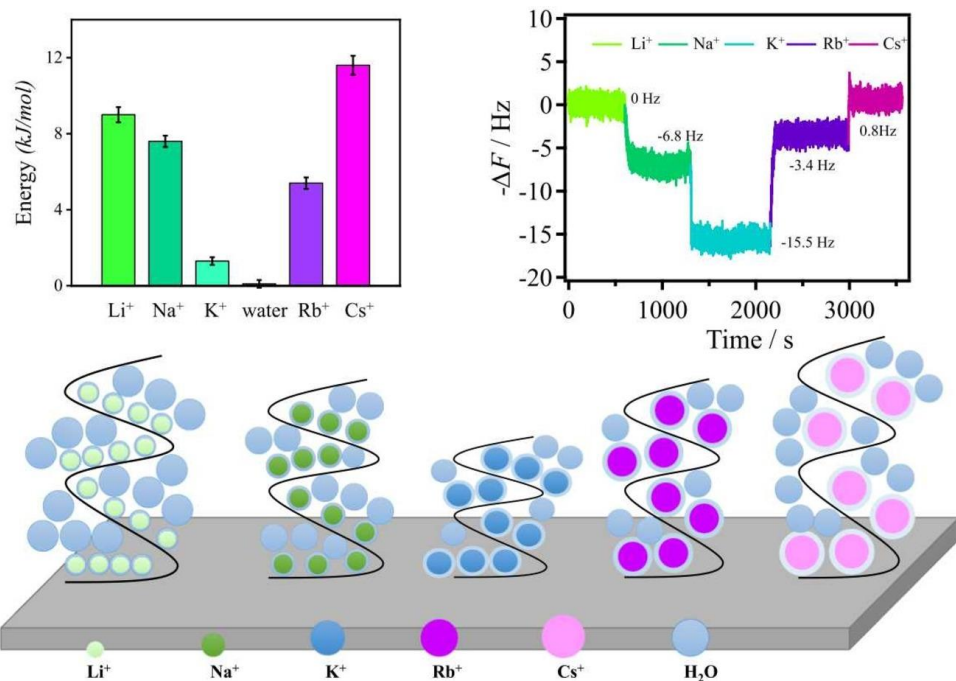
Prolates encapsulate and deliver more payloads to cells.



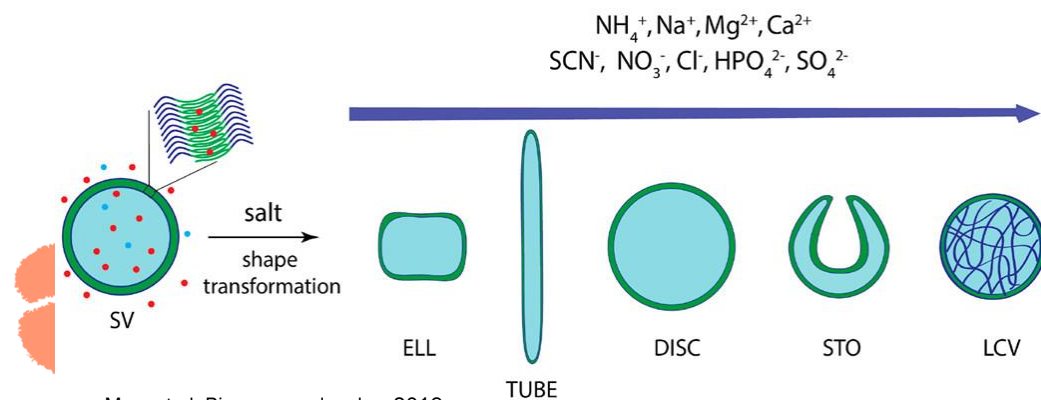
NaCl	Shape	Nile Red		BSA-Fluorescein	
		EE $\pm$ SD (%)	Loading Content (μ g dye/mg polymer)	EE $\pm$ SD (%)	Loading Content (μ g dye/mg polymer)
-	Sphere	5.2 $\pm$ 0.4	1.7 $\pm$ 0.004	13.9 $\pm$ 1.2	46.5 $\pm$ 4.0
+	Prolate	10.9 $\pm$ 0.3	3.6 $\pm$ 0.1	23.6 $\pm$ 9.4	78.5 $\pm$ 31.4



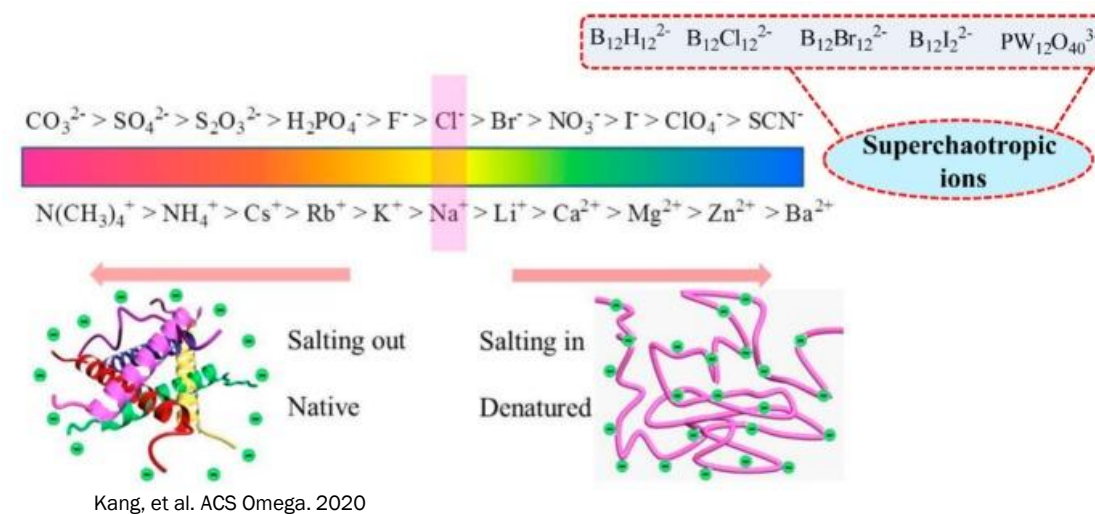
Chaotropic salts impact PEG and protein conformation.



Yu, et al. Colloids and Surfaces A, 2024.



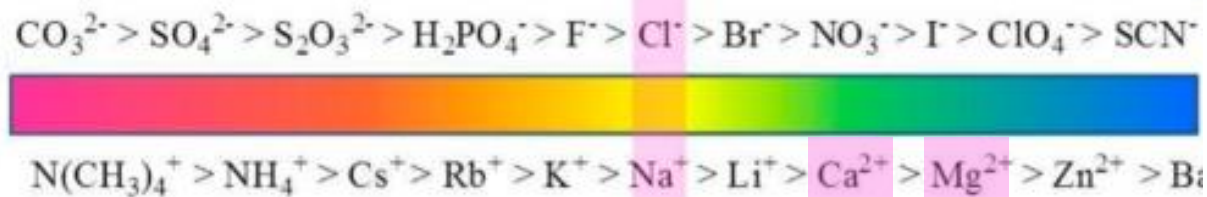
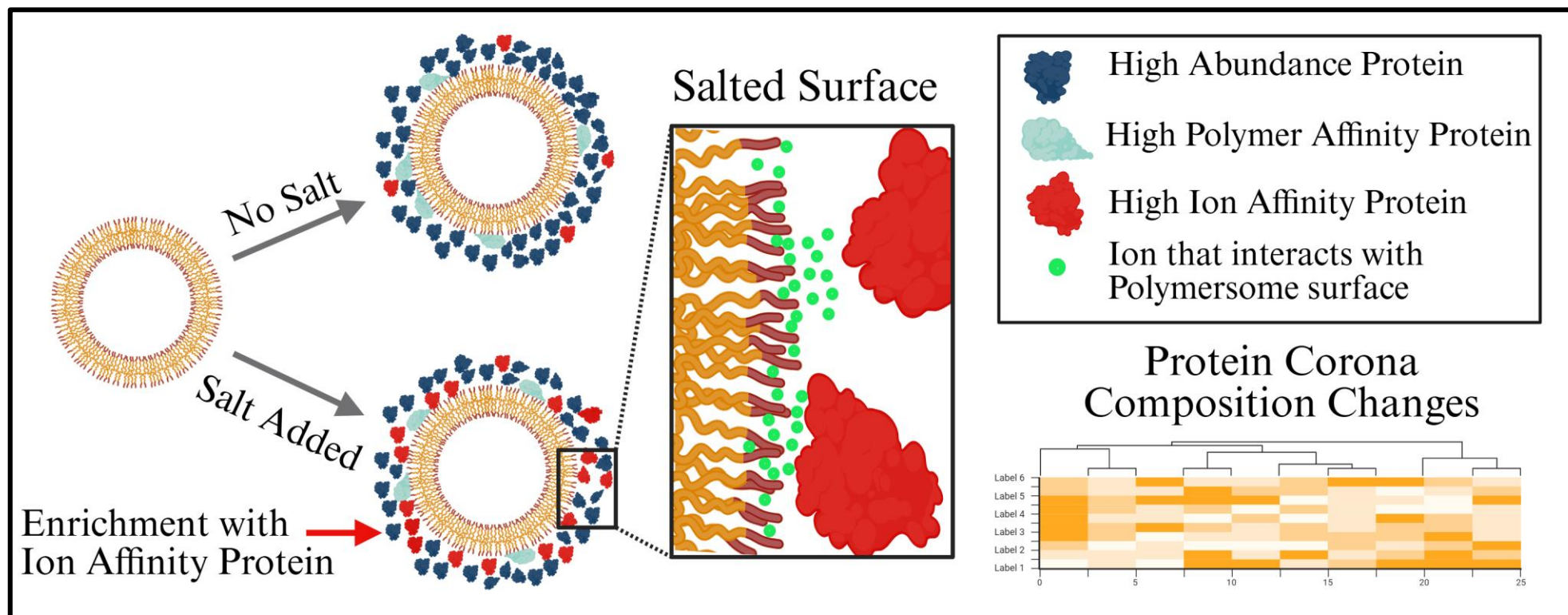
Men, et al. Biomacromolecules, 2019



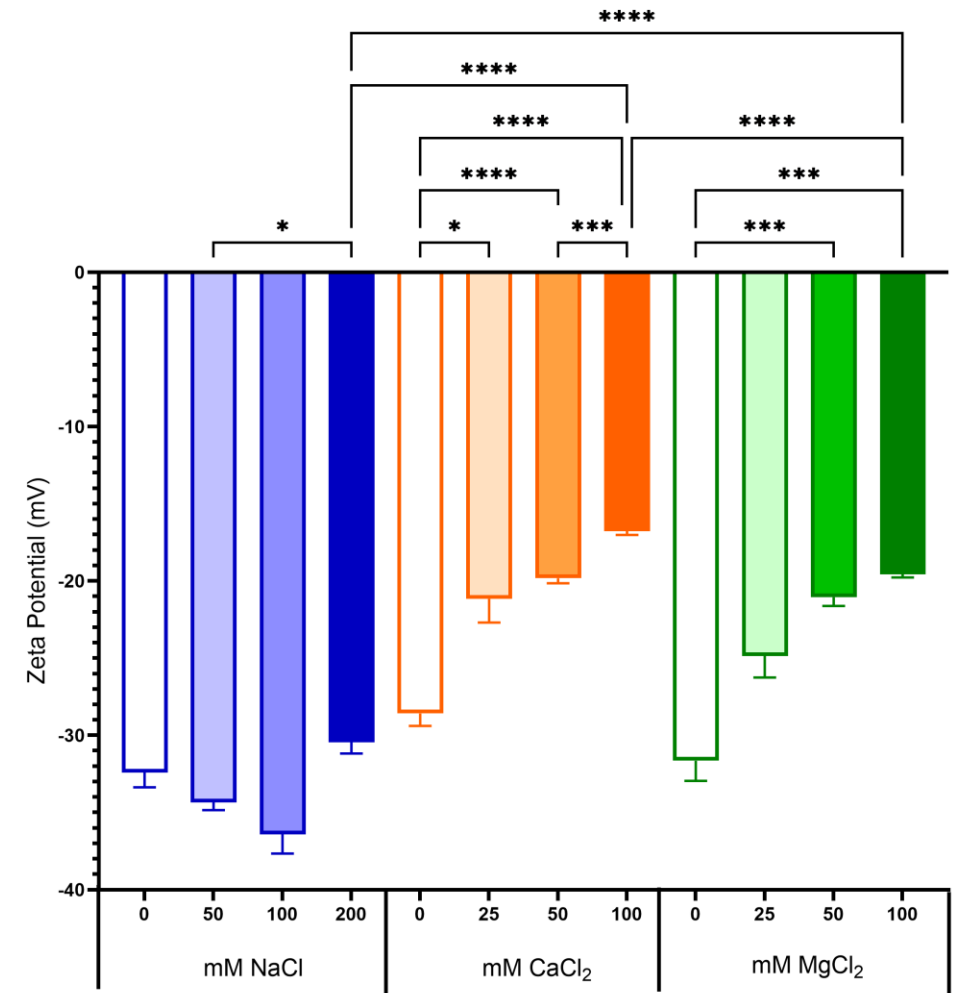
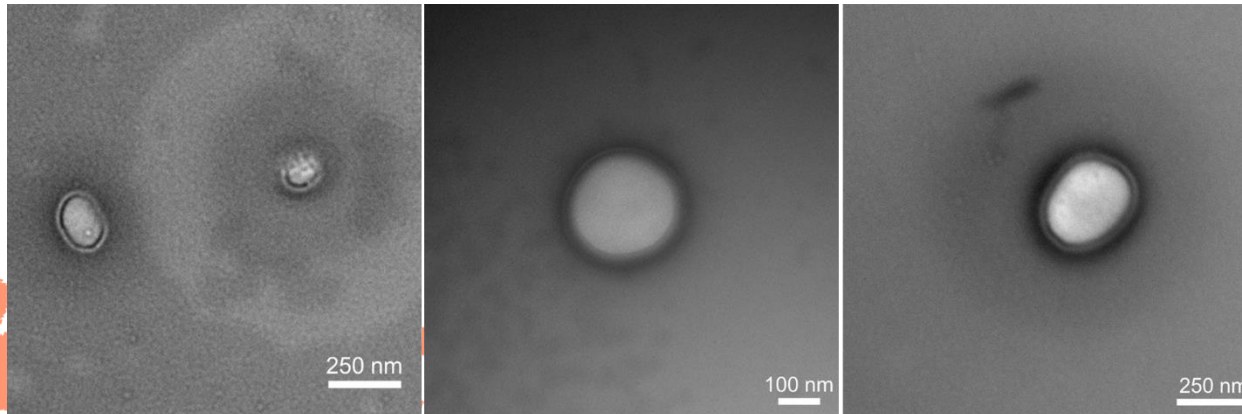
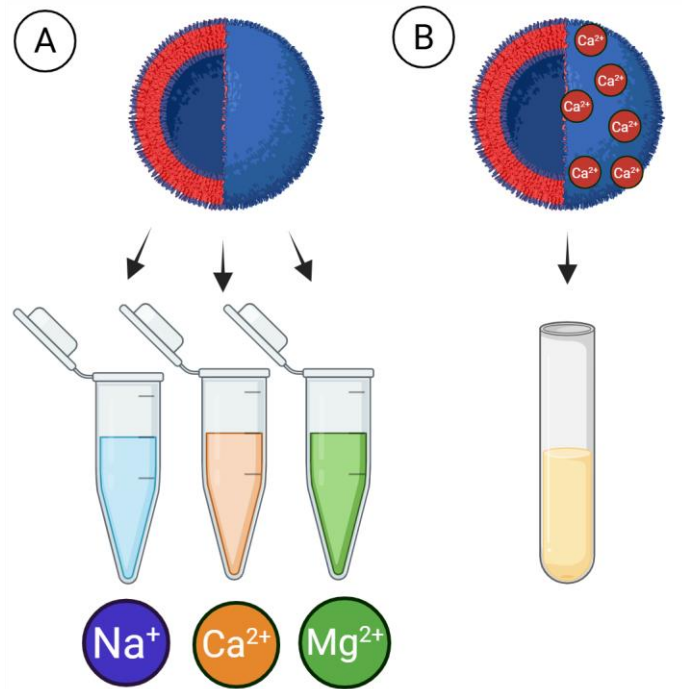
Hofmeister series salts change protein and PEG conformation.

Was the shape change improving uptake or was uptake improved due to an altered protein corona?

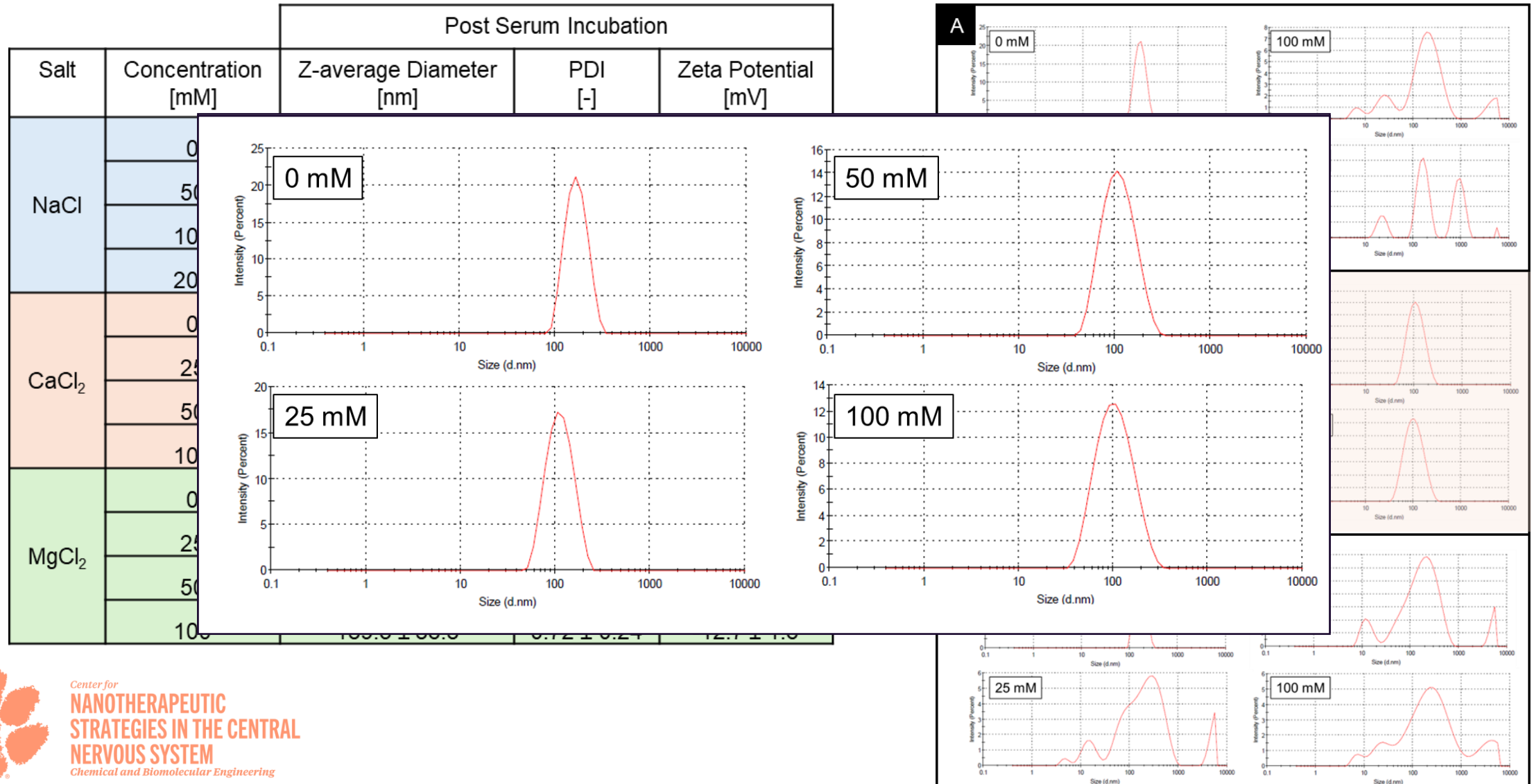
Chaotropic salts impact PEG and protein conformation.



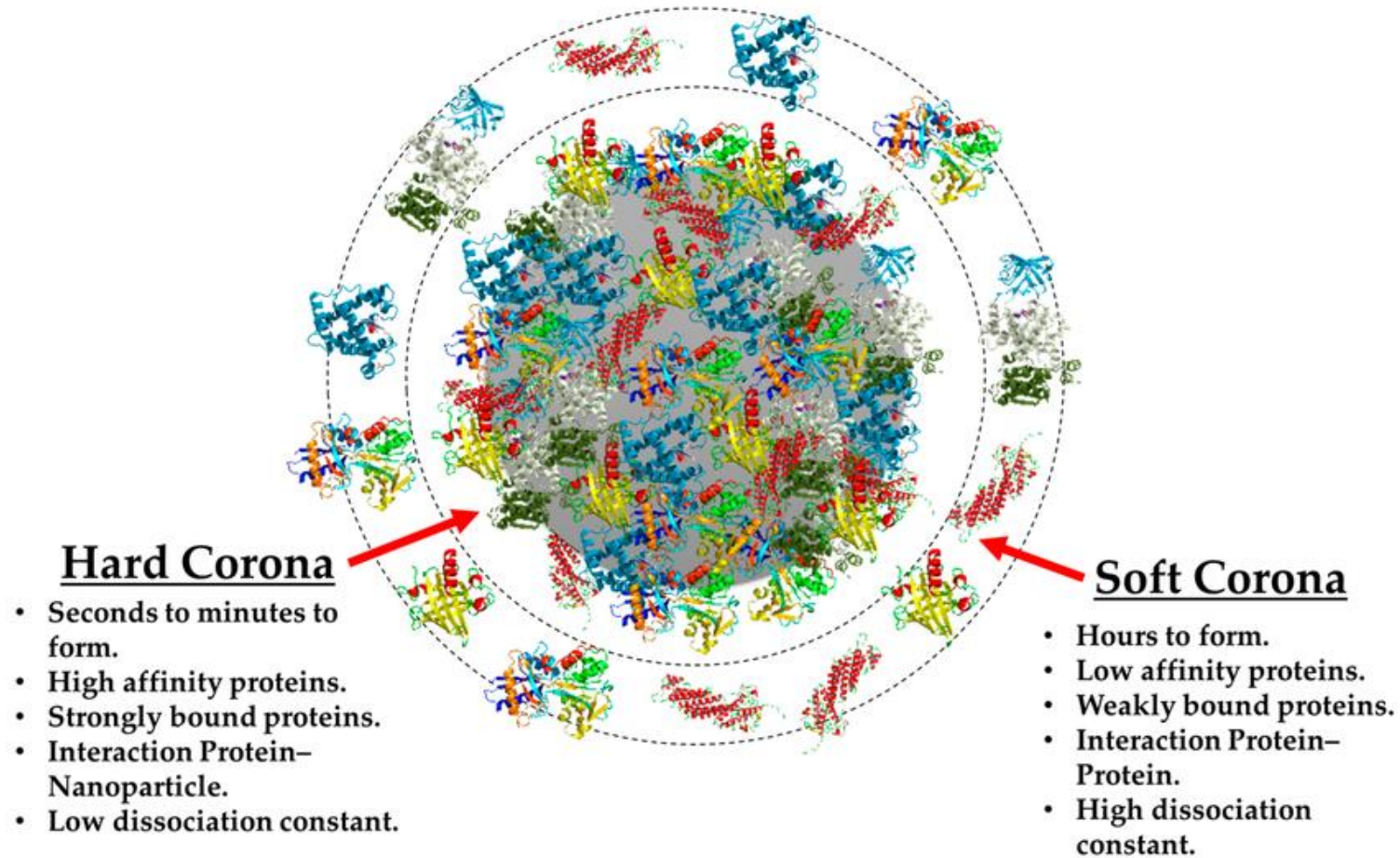
Different salts had varying affinities for PEG-b-PLA polymersomes.



Only Ca^{2+} formed a predominantly hard protein corona.

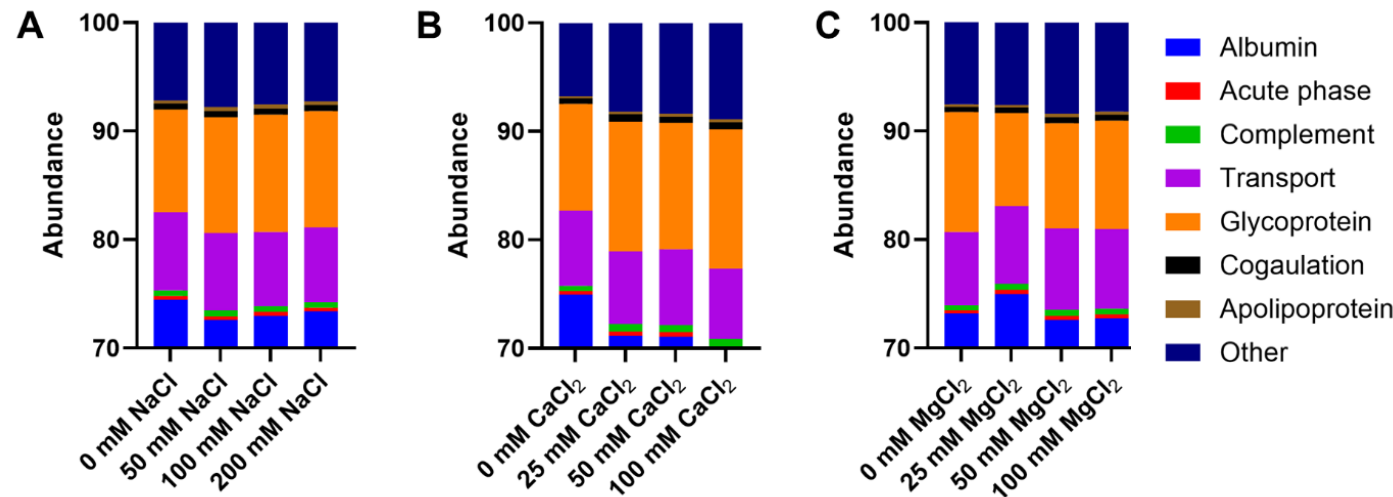
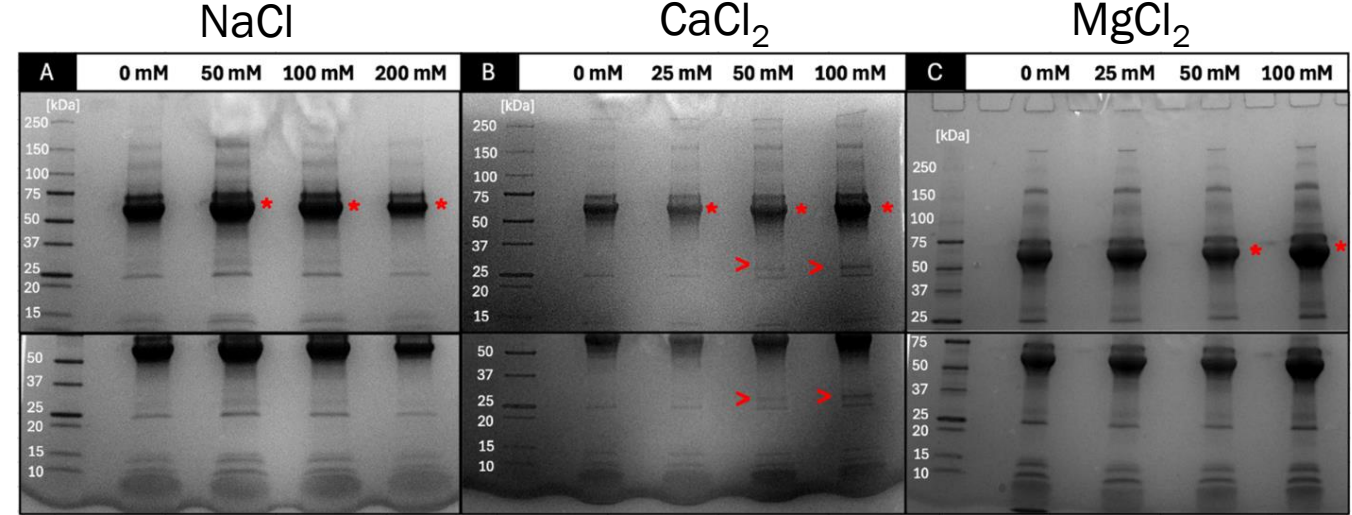
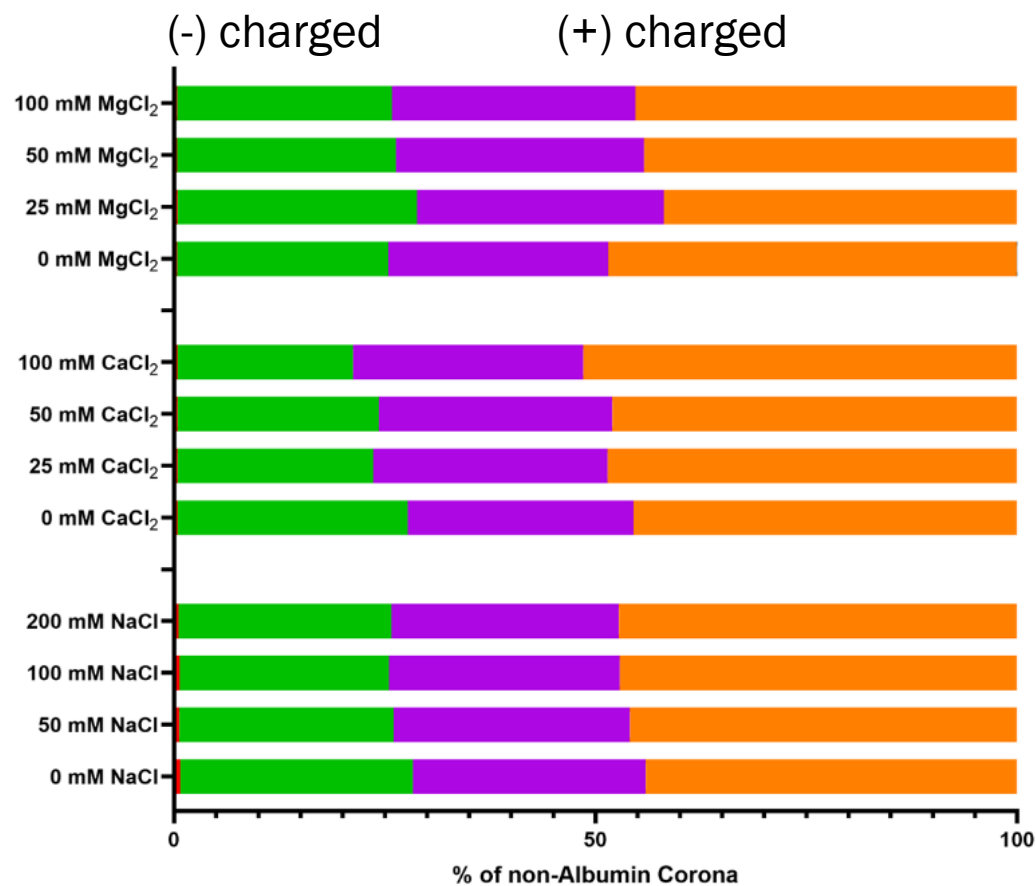


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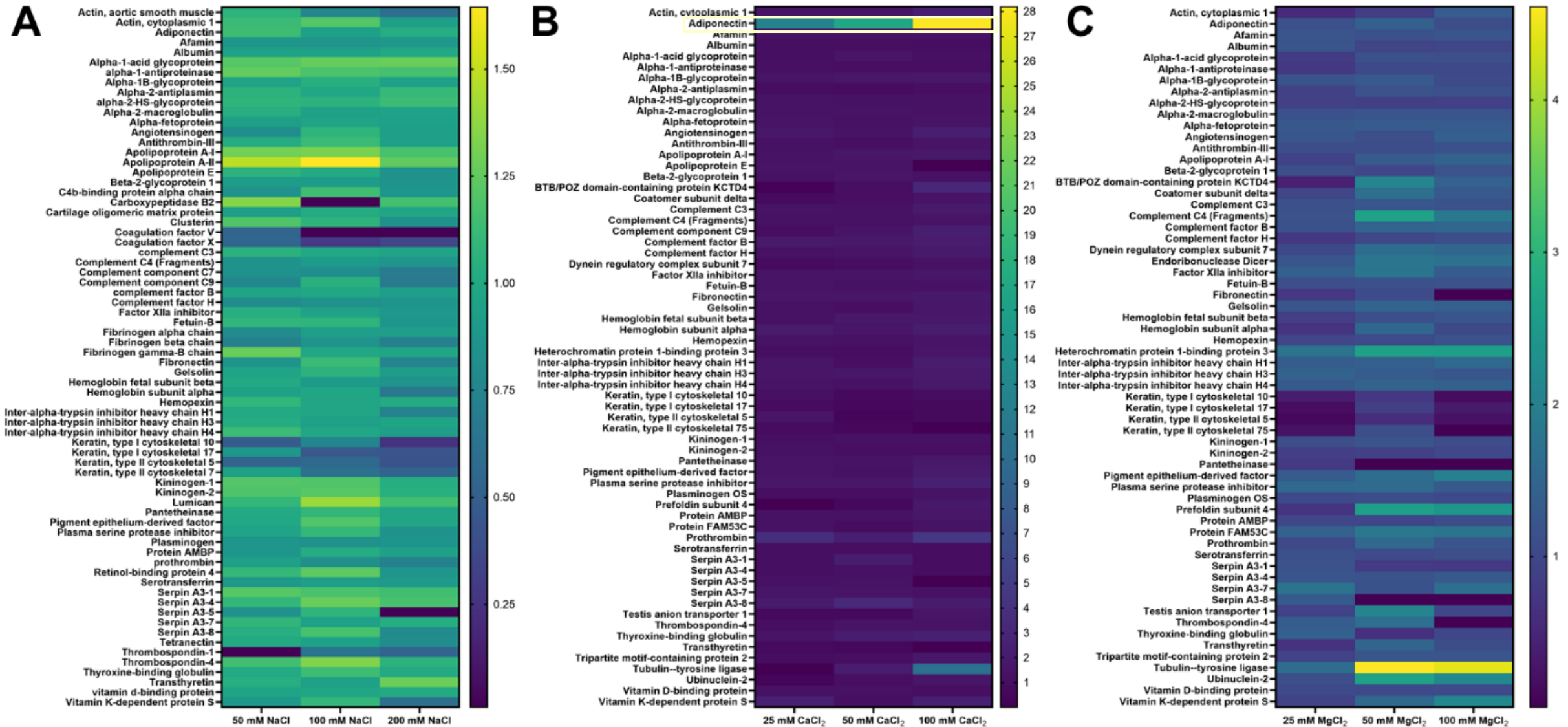


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Strongly bound cations had a dramatic impact on composition.



Protein corona composition changed amongst salts.



20 most abundant proteins changed after incubation with CaCl₂.

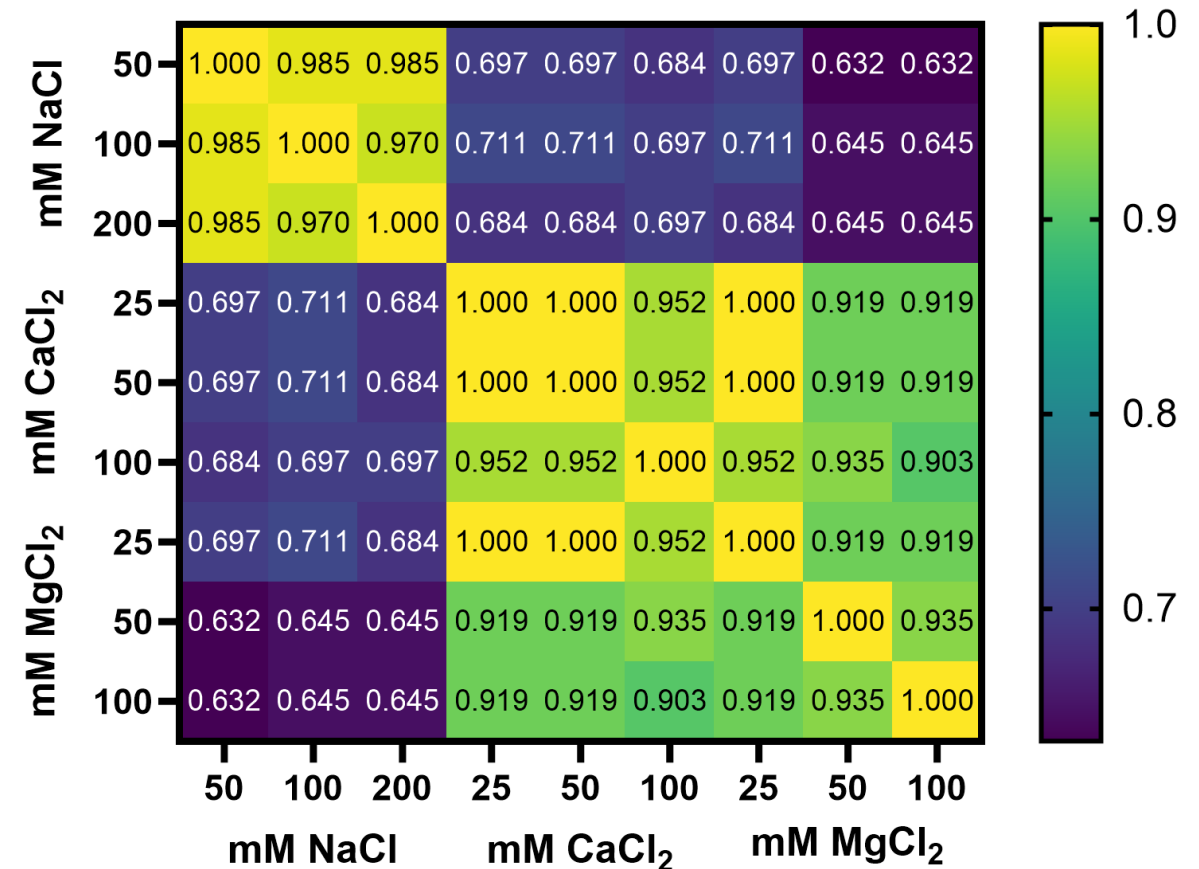
Protein Rank	0mM CaCl ₂	25mM CaCl ₂	50mM CaCl ₂	100mM CaCl ₂
1	Albumin OS	Albumin OS	Albumin OS	Albumin OS
2	Alpha-2-HS-glycoprotein OS	Alpha-2-HS-glycoprotein OS	Alpha-2-HS-glycoprotein OS	Alpha-2-HS-glycoprotein OS
3	Serotransferrin OS	Serotransferrin OS	Serotransferrin OS	Serotransferrin OS
4	Alpha-1-antiproteinase OS	Alpha-1-antiproteinase OS	Alpha-1-antiproteinase OS	Alpha-1-antiproteinase OS
5	Alpha-2-macroglobulin OS	Alpha-2-macroglobulin OS	Alpha-2-macroglobulin OS	Alpha-2-macroglobulin OS
6	Alpha-fetoprotein OS	Alpha-fetoprotein OS	Alpha-fetoprotein OS	Alpha-fetoprotein OS
7	Vitamin D-binding protein OS	Fetuin-B OS	Vitamin D-binding protein OS	Complement C3 OS
8	Fetuin-B OS	Complement C3 OS	Fetuin-B OS	Vitamin D-binding protein OS
9	Complement C3 OS	Vitamin D-binding protein OS	Complement C3 OS	Adiponectin OS
10	Serpin A3-1 OS	Serpin A3-1 OS	Serpin A3-1 OS	Serpin A3-1 OS
11	Apolipoprotein A-I OS	Inter-alpha-trypsin inhibitor heavy chain H4 OS	Apolipoprotein A-I OS	Apolipoprotein A-I OS
12	Inter-alpha-trypsin inhibitor heavy chain H4 OS	Apolipoprotein A-I OS	Inter-alpha-trypsin inhibitor heavy chain H3 OS	Inter-alpha-trypsin inhibitor heavy chain H4 OS
13	Antithrombin-III OS	Adiponectin OS	Adiponectin OS	Prothrombin OS
14	Plasminogen OS	Prothrombin OS	Alpha-1B-glycoprotein OS	Alpha-1B-glycoprotein OS
15	Alpha-2-antiplasmin OS	Antithrombin-III OS	Antithrombin-III OS	Angiotensinogen OS
16	Kininogen-2 OS	Plasminogen OS	Plasminogen OS	Antithrombin-III OS
17	Hemopexin OS	Alpha-1B-glycoprotein OS	Hemopexin OS	Plasminogen OS
18	Alpha-1B-glycoprotein OS	Hemopexin OS	Kininogen-2 OS	Hemopexin OS
19	Protein AMBP OS	Kininogen-1 OS	Angiotensinogen OS	Protein AMBP OS
20	Alpha-1-acid glycoprotein OS	Angiotensinogen OS	Alpha-2-antiplasmin OS	Alpha-2-antiplasmin OS

Adiponectin: Ca²⁺ binding motif
Angiotensinogen: Ca²⁺ in renin-angiotensin system

Conclusions

- Salt ions can change the shape of **polymersomes** while maintaining their properties.
- Divalent chaotropic salts bind more tightly to **PEG**, impacting the isoelectric point of proteins found in the protein corona.
- **Calcium** incubation leads to the most **biologically specific** protein corona changes.

Protein corona composition can be engineered on the surface of PEG-b-PLA polymersomes.



Thank you for your time and attention!

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