

Understanding polymerization kinetics and self-assembly in PISA and Coacervate systems

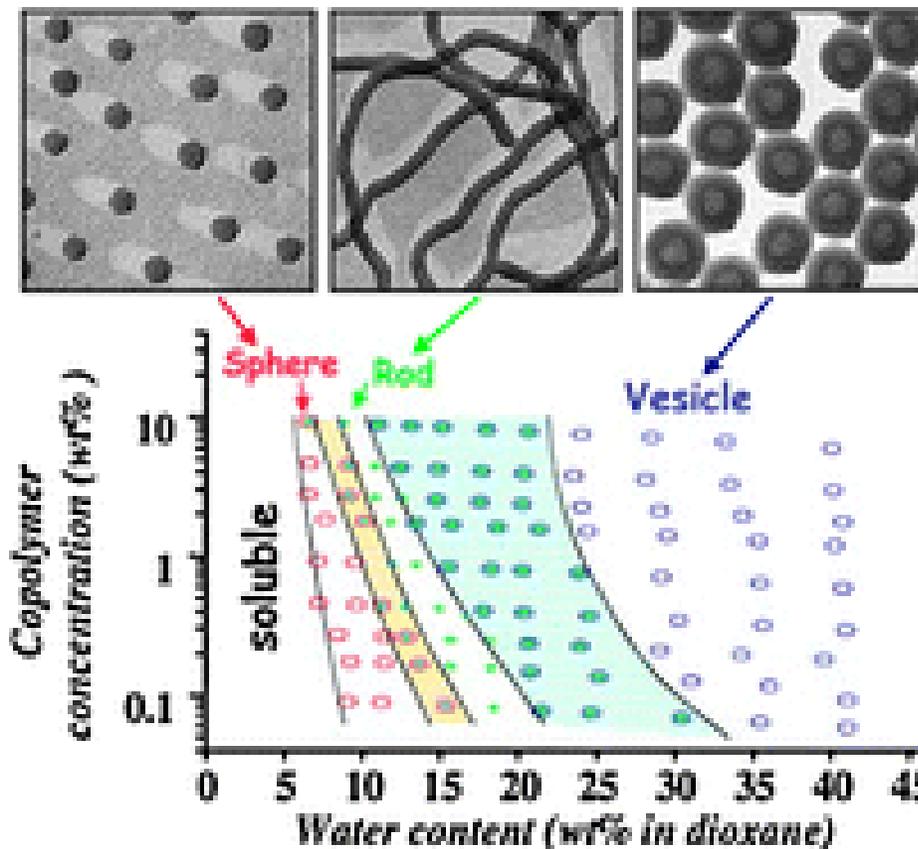
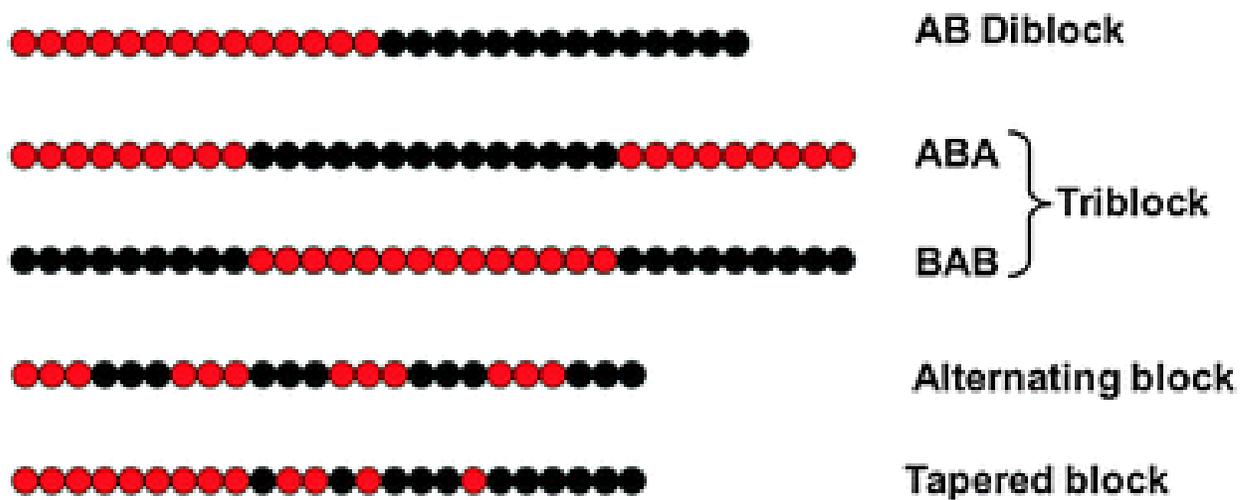
Riya Singh

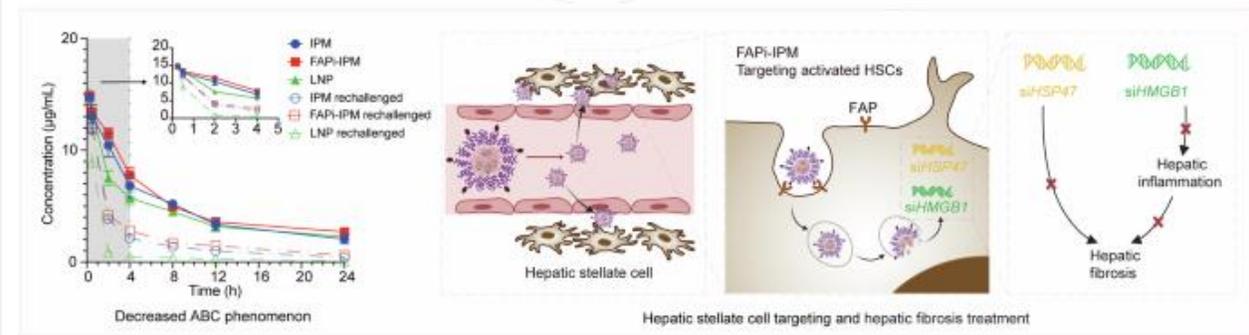
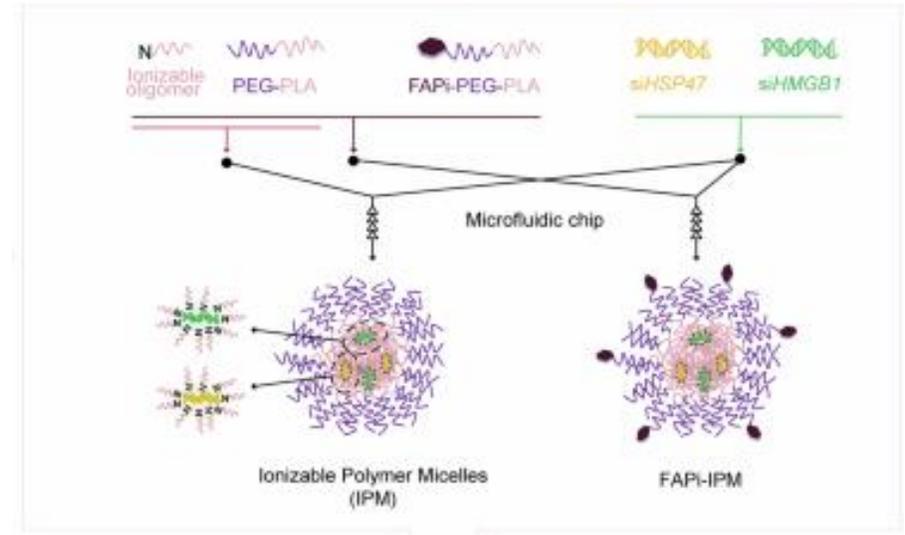
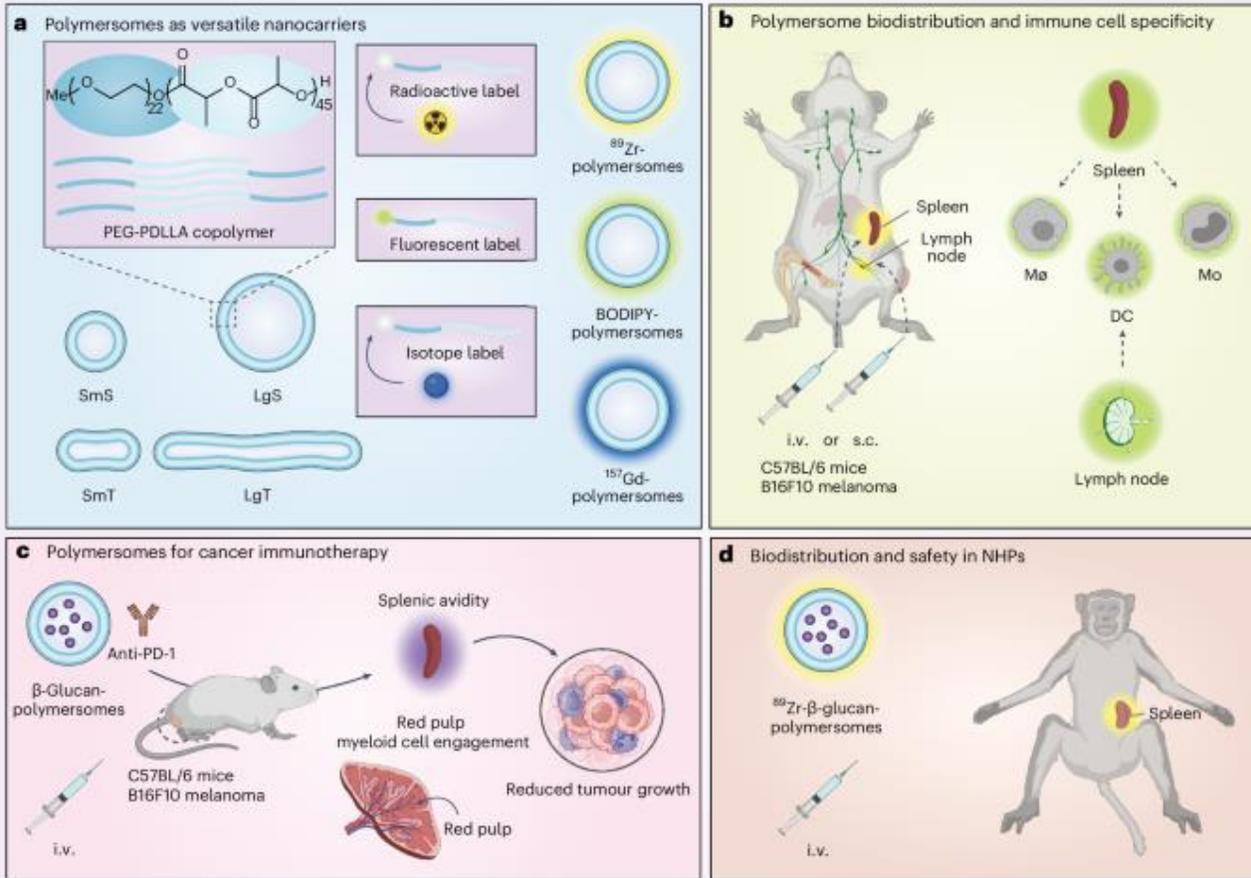
Ph.D. Candidate

Advisor : Prof. Joe Patterson

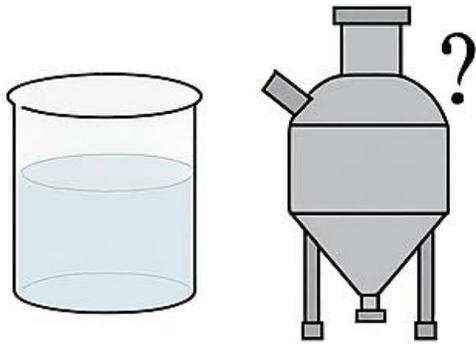
Department of Chemistry

University of California, Irvine



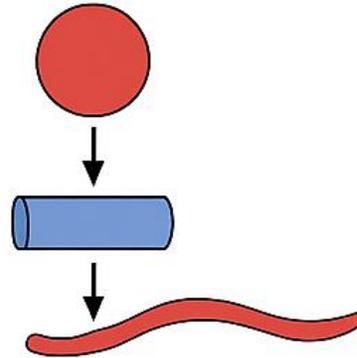


Scalability & Reproducibility



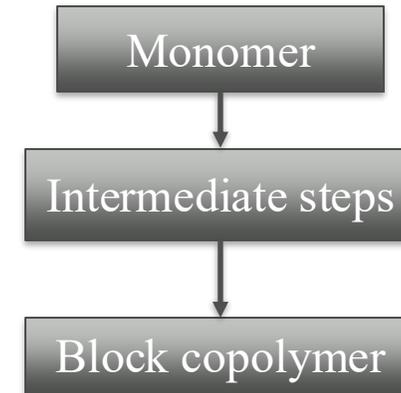
- Difficult to scale or reproduce on a larger scale.
- Batch to batch variability in morphology.

Morphological control



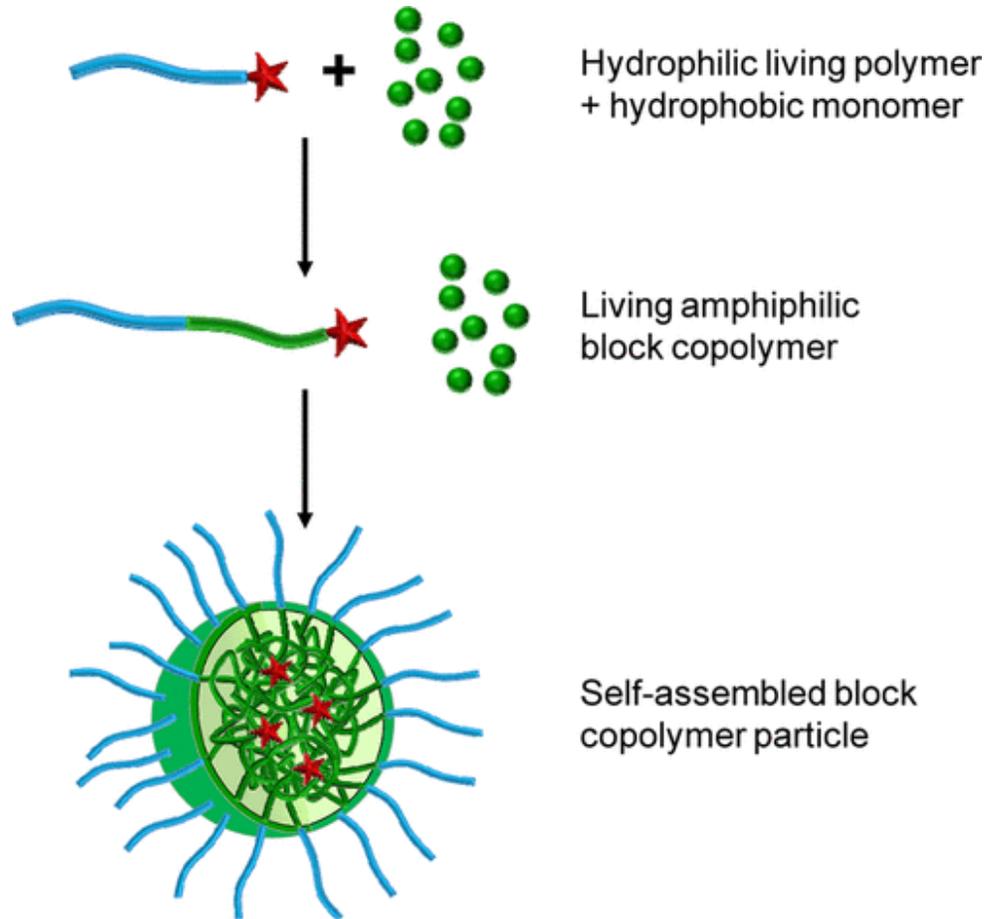
- Spherical micelles dominate thermodynamically
- Kinetic pathways for rods or worms are narrow.
- Difficult to get anisotropic structures.

Complex chemistry & Limited tunability

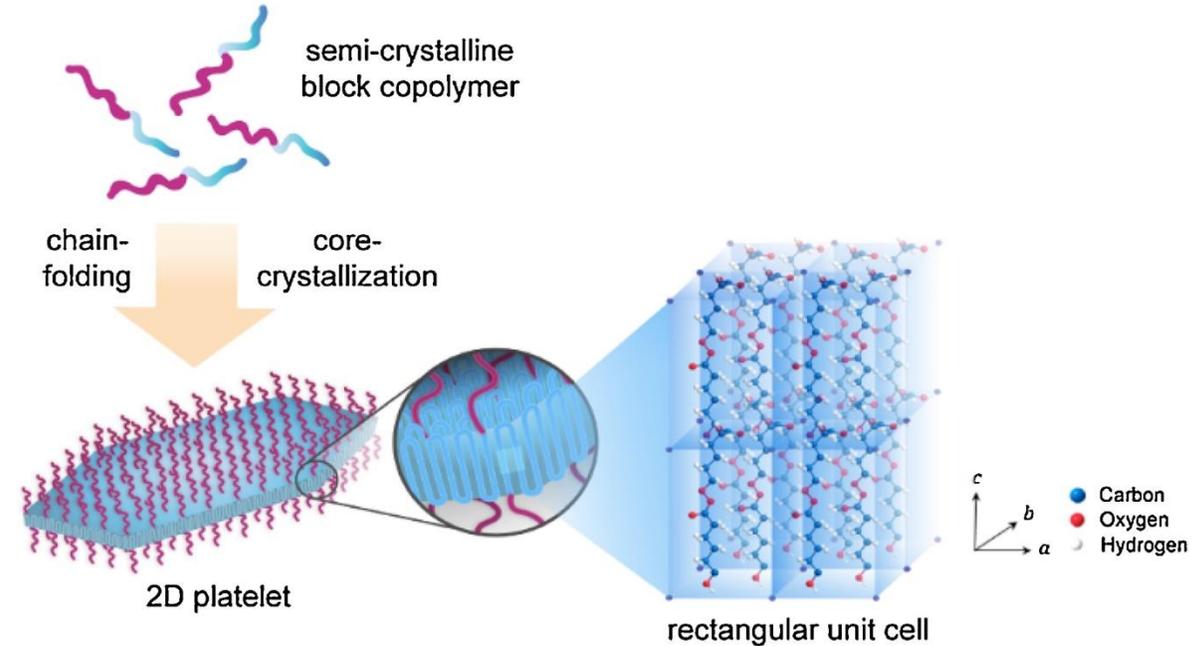


- Controlled polymerizations demand multiple steps.
- Functional monomers often require tedious synthesis.

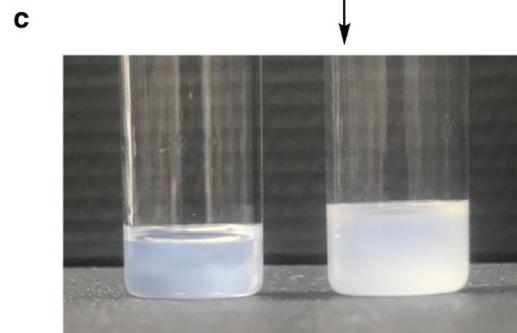
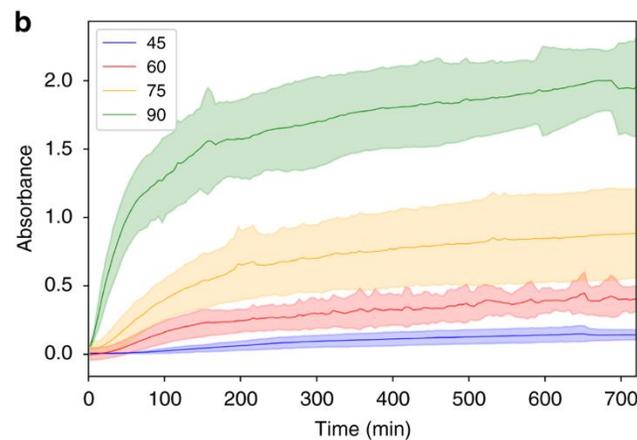
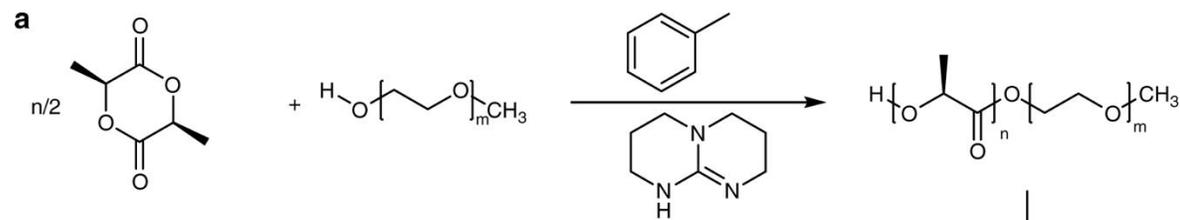
Polymerization-Induced Self-Assembly (PISA)



Crystallization-Driven Self-Assembly (CDSA)

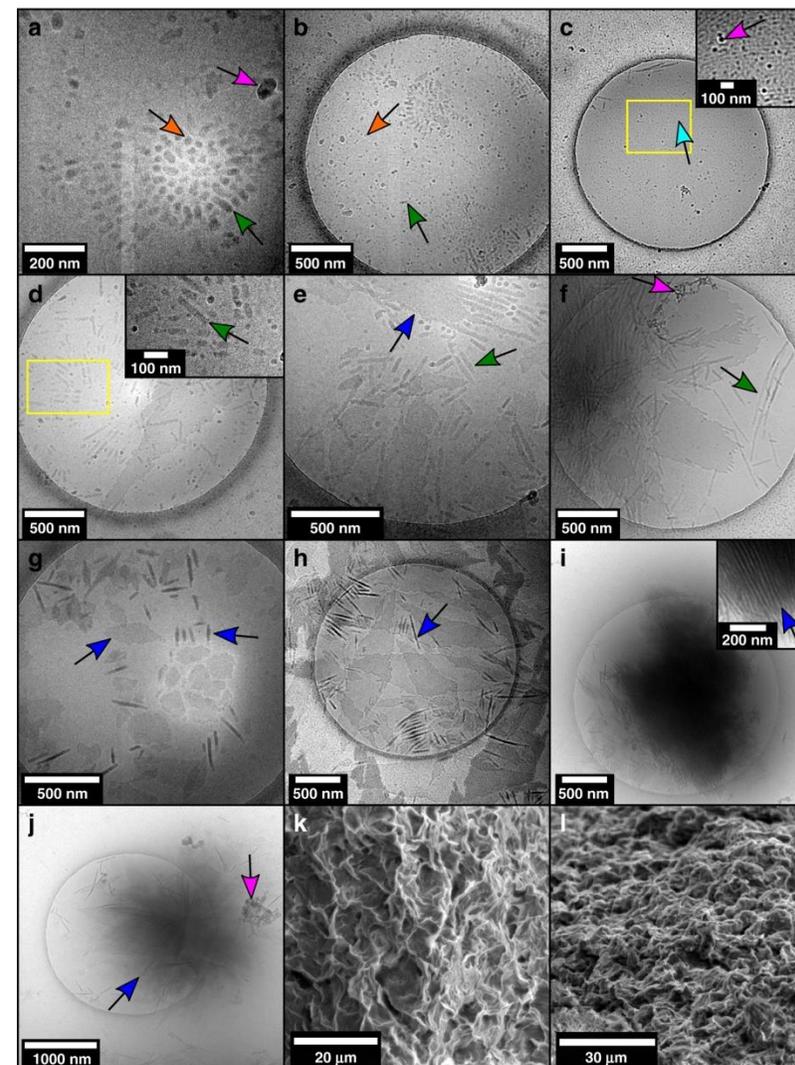


ROPI-CDSA scheme and self-assembly kinetics

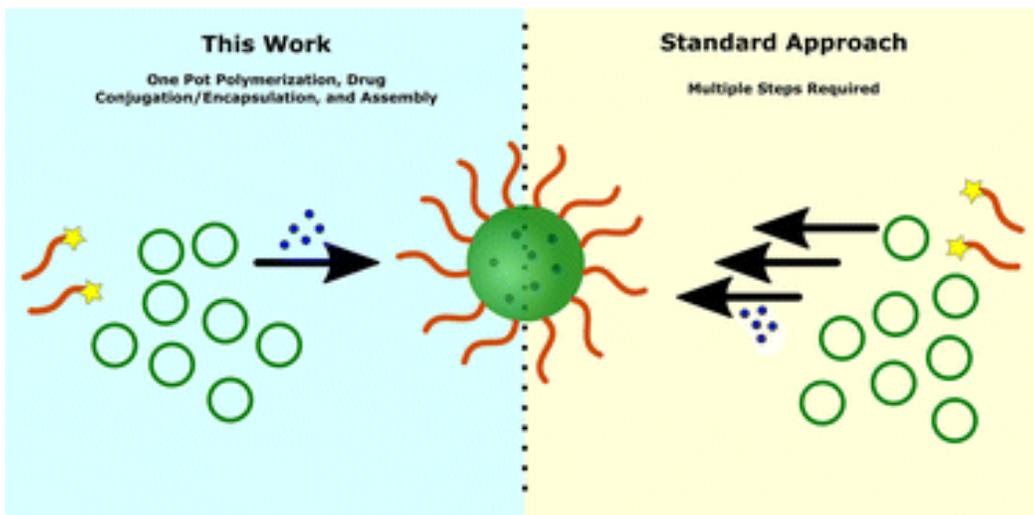
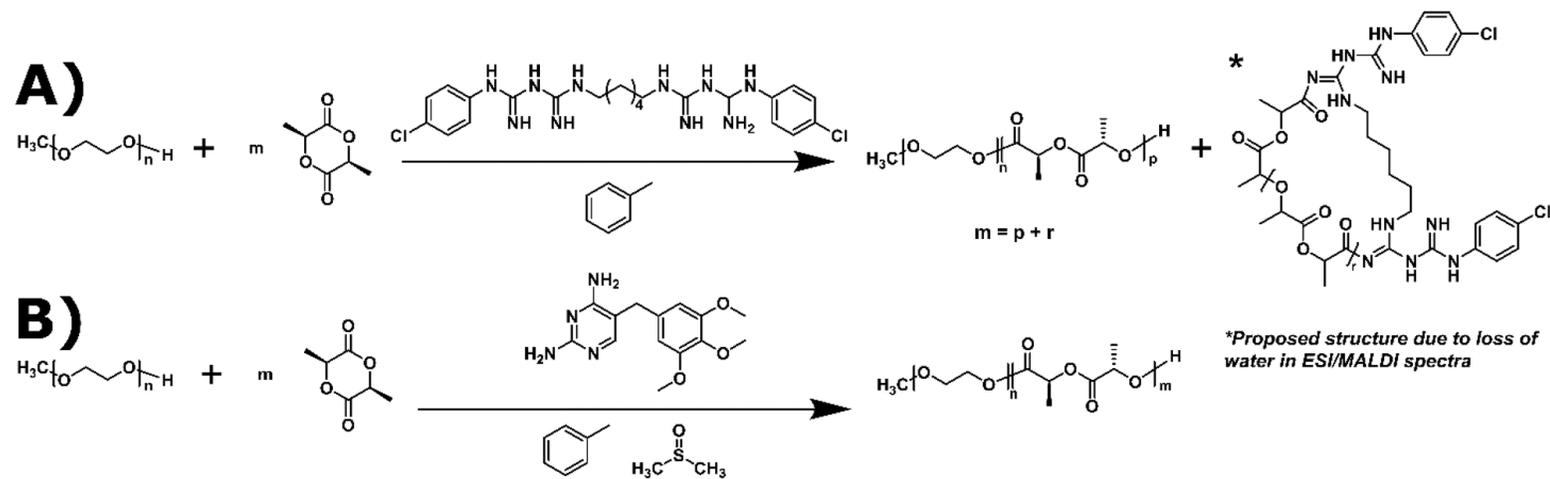


Legend:

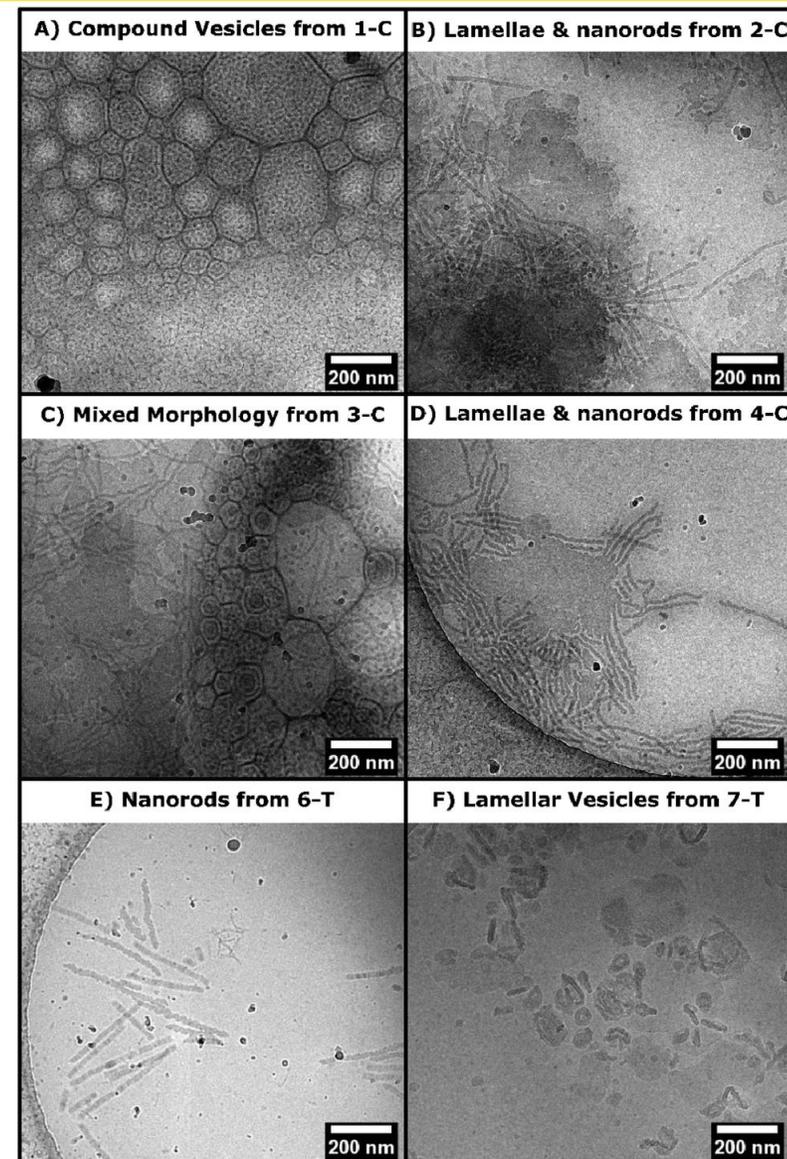
- Spheres
- Rods
- Fibers
- Lamellae
- Ice Contamination

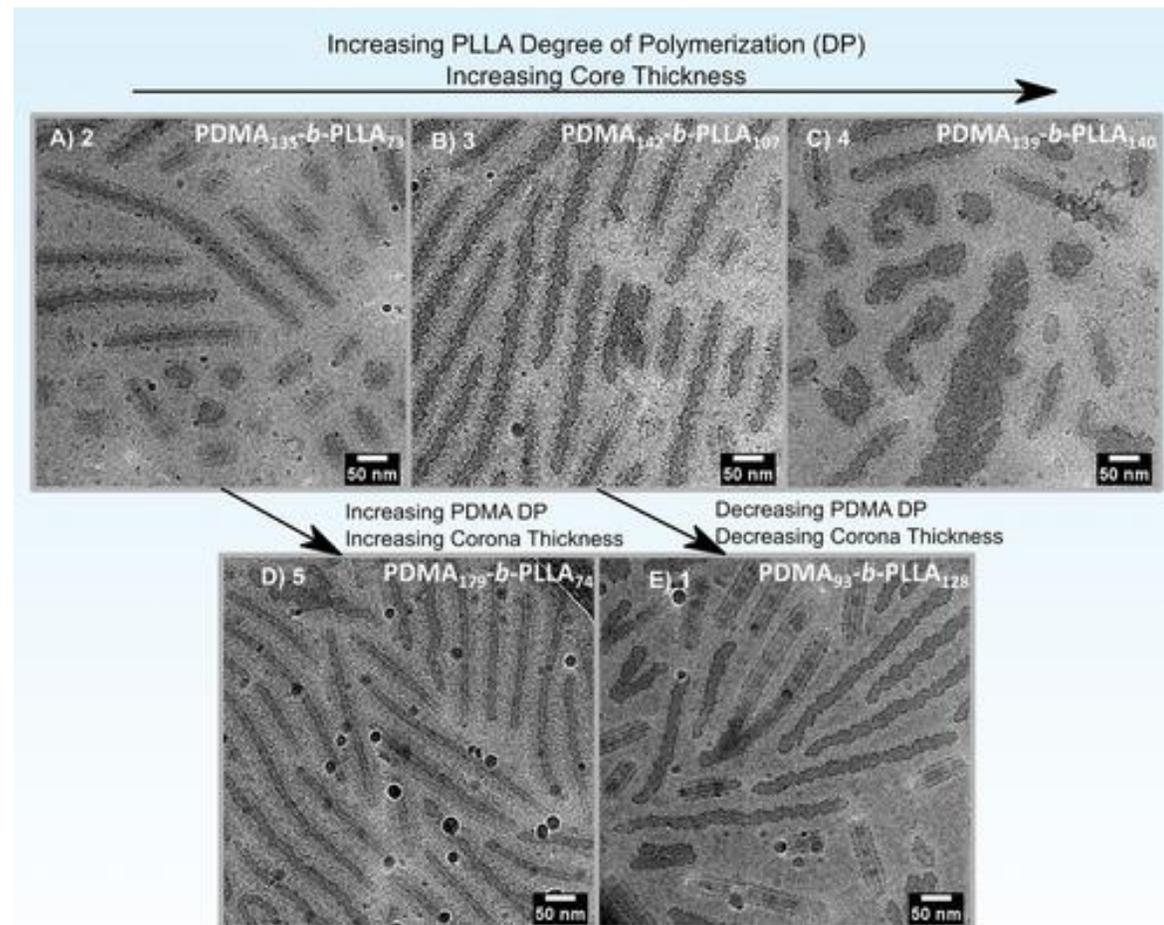
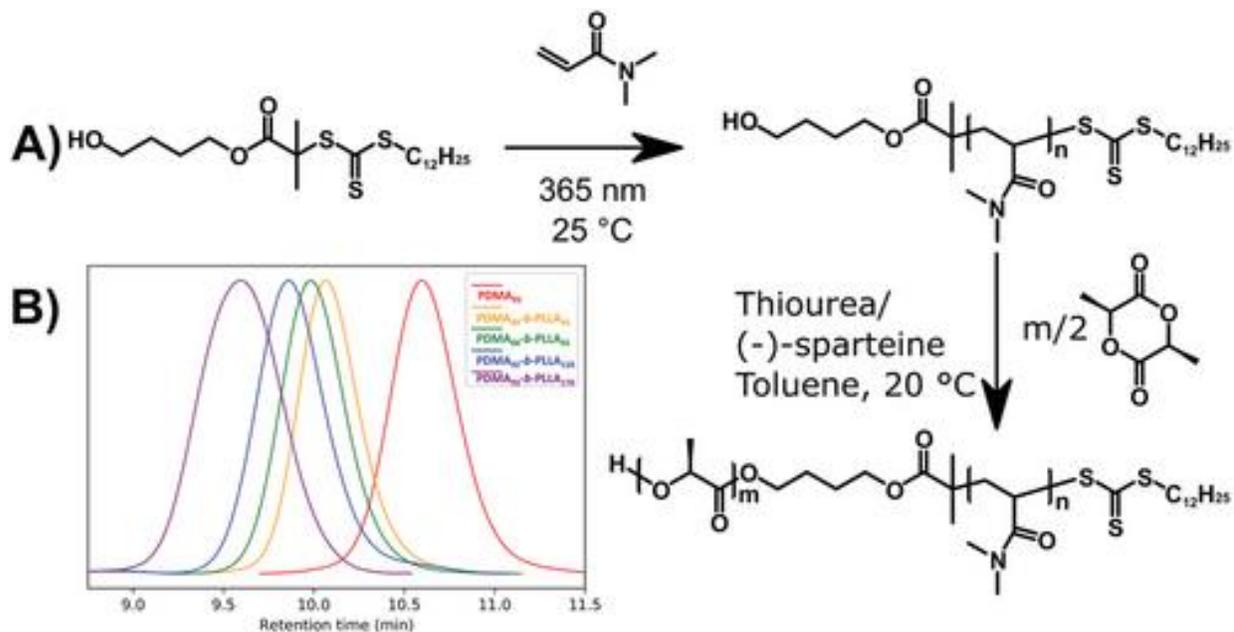


Nat. Commun. 2020, 11, 4690.

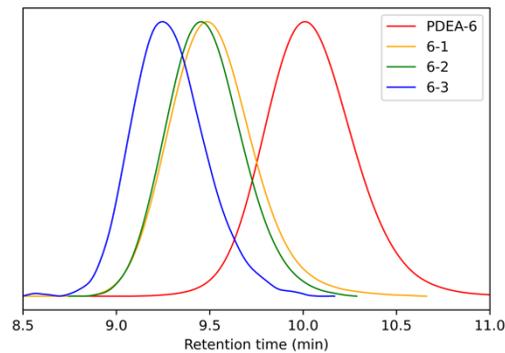
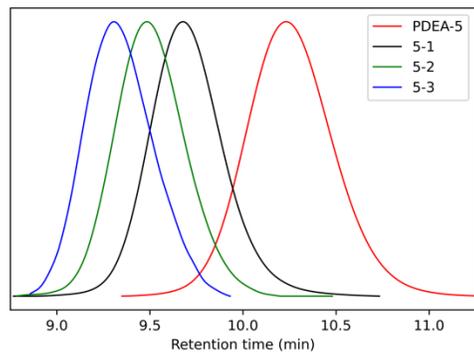
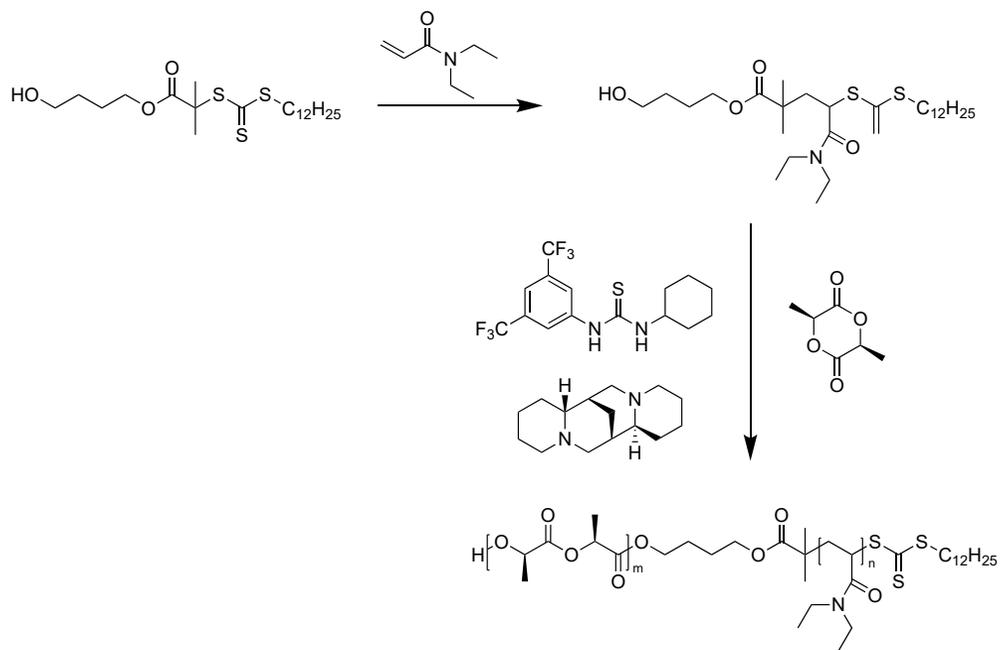


RSC Appl. Polym. 2024, 2, 238–247.



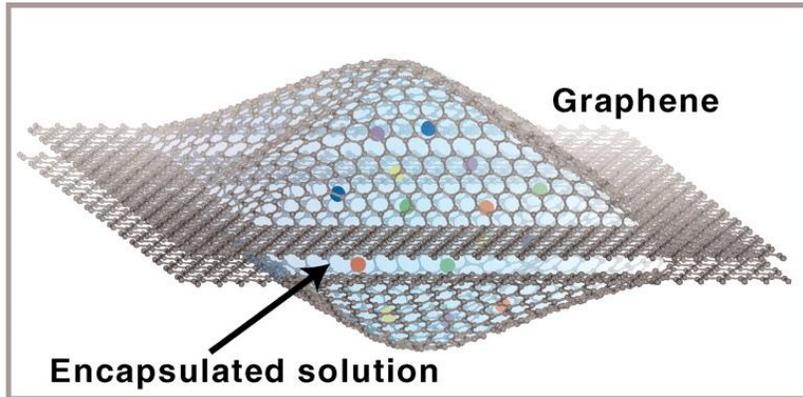


Hurst, P. J., Yoon, J., **Singh, R.**, Abouchaleh, M. F., Stewart, K. A., Sumerlin, B. S., & Patterson, J. P., *Macromol. Rapid Commun.*, 2024, 45(12)



Sample ID	Target PDEA DP	PDEA Conversion (%)	Target PLLA DP	PLLA Conversion (%)	\bar{D}	M _n (g/mol)	Polymer Structure	Turbid
PDEA-5	150	95	-	-	1.07	18200	PDEA ₁₄₃	N/A
5-1	150	96	100	84	1.07	24400	PDEA ₁₄₄ - <i>b</i> -PLLA ₈₄	Yes
5-2	150	94	150	84	1.07	27000	PDEA ₁₄₁ - <i>b</i> -PLLA ₁₂₆	Yes
5-3	150	96	200	92	1.07	31600	PDEA ₁₄₄ - <i>b</i> -PLLA ₁₈₄	Yes
PDEA-6	200	98	-	-	1.08	24900	PDEA ₁₉₆	N/A
6-1	200	97	100	77	1.10	30200	PDEA ₁₉₄ - <i>b</i> -PLLA ₇₇	Yes
6-2	200	99	150	81	1.08	34000	PDEA ₁₉₈ - <i>b</i> -PLLA ₁₂₂	Yes
6-3	200	99	200	83	1.08	37100	PDEA ₁₉₈ - <i>b</i> -PLLA ₁₆₆	Yes

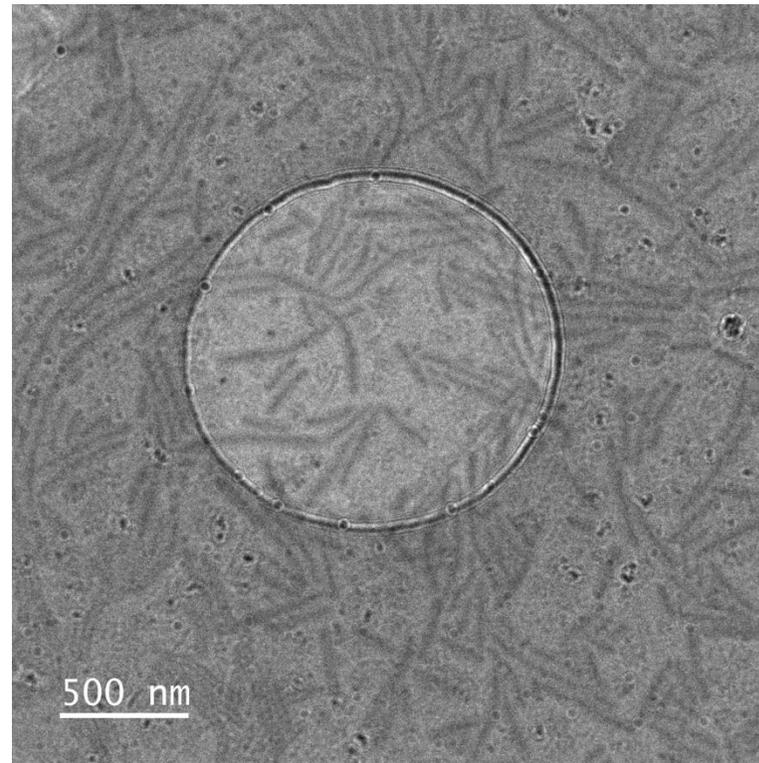
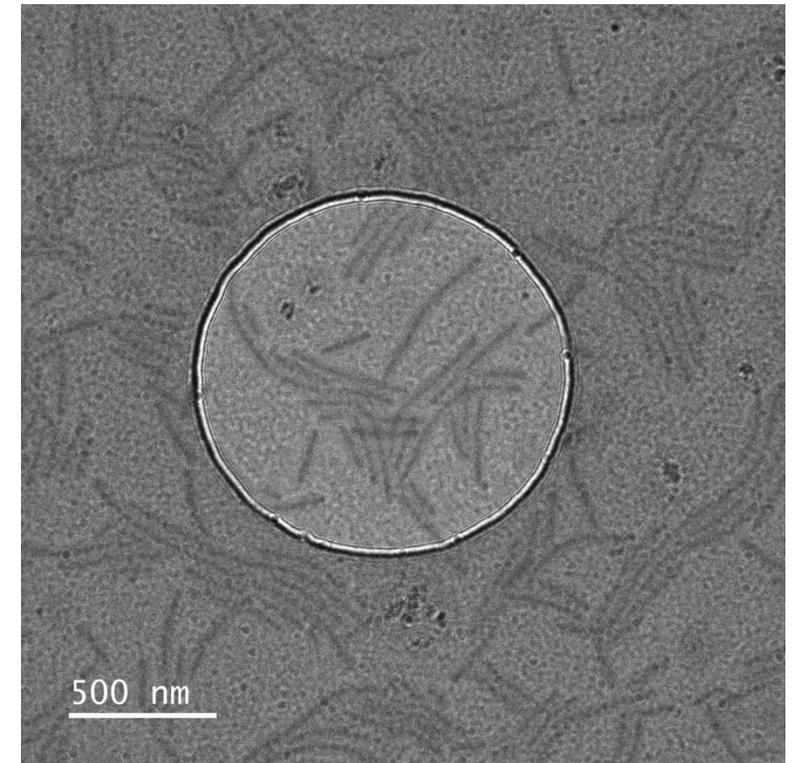
Graphene Liquid-Cell
Transmission Electron Microscopy
(GLC-TEM)



Liquid

PDEAA₂₀₀-b-PLLA₁₀₀

Dry

Average diameter = 43 ± 9.1 nmAverage diameter = 28.4 ± 3.1 nm

- Study the self-assembly behavior of these polymers directly in their native solvent environment using graphene liquid cell transmission electron microscopy (GLC-TEM).
- Overcome limitations of conventional TEM, which captures dried-state artifacts rather than true solution morphologies.
- Capture in-situ dynamics of block copolymer self-assembly and crystallization, allowing time-resolved observation of structural evolution.
- Focus on lactone-based polymers synthesized via PI-CDSA and other related biodegradable block copolymers.

Acknowledgement

Thanks to: Prof. Joe Patterson & Patterson lab members



Thank you for your time and attention!

Questions?