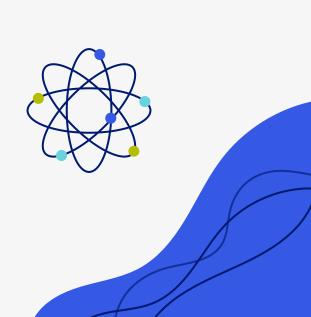
Analyzing the Structure of Bicelles with Small Angle Neutron Scattering

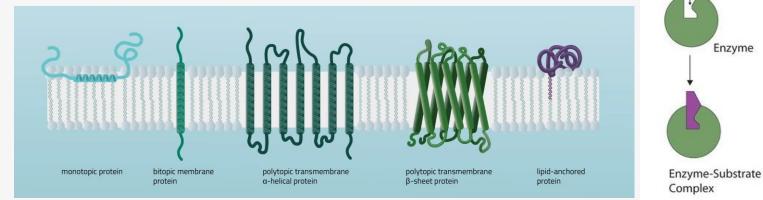
Xael Shan





Why do we care?

Membrane Proteins very important to study, but difficult!



Substrate

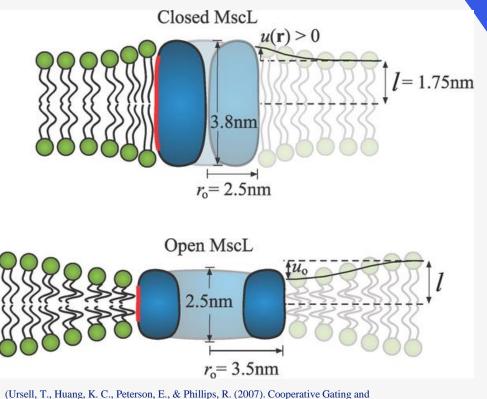
Active site

Enzyme

(Membrane Proteins - Definition, Types & Functions. (n.d.). Cube Biotech. https://cubebiotech.com/knowledge/membrane-protein-stabilization/membrane-proteins/)

But why study the membrane?

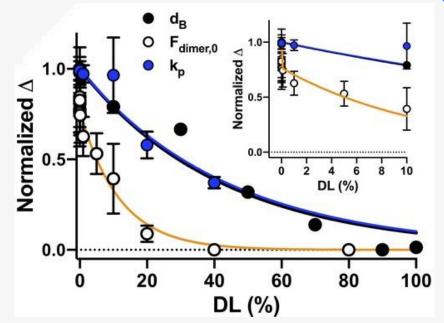
Studying the environment allows us to consider how it impacts the sample



(Ursell, T., Huang, K. C., Peterson, E., & Phillips, R. (2007). Cooperative Gating and Spatial Organization of Membrane Proteins through Elastic Interactions. *PLoS Computational Biology*, *3*(5), e81. https://doi.org/10.1371/journal.pcbi.0030081)

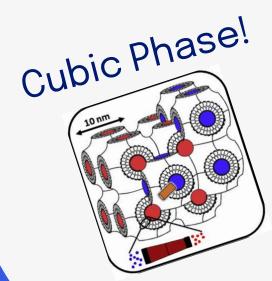


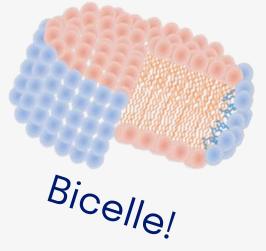
Studying the environment allows us to consider how it impacts the sample



(Chadda, R., Bernhardt, N., Kelley, E. G., Teixeira, S. C., Griffith, K., Gil-Ley, A., Öztürk, T. N., Hughes, L. E., Forsythe, A., Krishnamani, V., Faraldo-Gómez, J. D., & Robertson, J. L. (2021). Membrane transporter dimerization driven by differential lipid solvation energetics of dissociated and associated states. *eLife*, *10*. https://doi.org/10.7554/elife.63288)

Model systems





(Model Membranes / Columbus Lab. (n.d.). Columbus Lab. https://www.columbuslabs.org/model-membranes)

Nano Disk!

(Membrane Proteins - Definition, Types & Functions. (n.d.). Cube Biotech. https://cube-biotech.com/knowledge/membrane-proteinstabilization/membrane-proteins/)

(L. D., & Caffrey, M. (2020). Structure and functional characterization of membrane integral proteins in the lipid cubic phase. *Journal of Molecular Biology*, *432*(18), 5104–5123. https://doi.org/10.1016/j.jmb.2020.02.024)

Bicelle + Lipid Model x=0.5 x=1 Detergent **Bicelle** Lipid Bilayer Micelle Lipid Xeff = [lipidin bicelle - lipidfloating] x=2 detergentin bicelle -

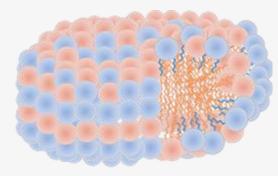
detergent floating]

(Rieth, M. D. (2018). The best of both worlds: A new lipid complex has micelle and bicelle-like properties. *bioRxiv (Cold Spring Harbor Laboratory)*. https://doi.org/10.1101/437327)

Bicelle + Lipid Model

Lipid

(Cadwell et al, 2018) (Katsaras et al , 2005)

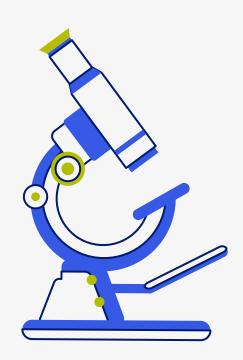


IDEAL MODEL!

(Model Membranes / Columbus Lab. (n.d.). Columbus Lab. https://www.columbuslabs.org/model-membranes)

DMPC

DHPC

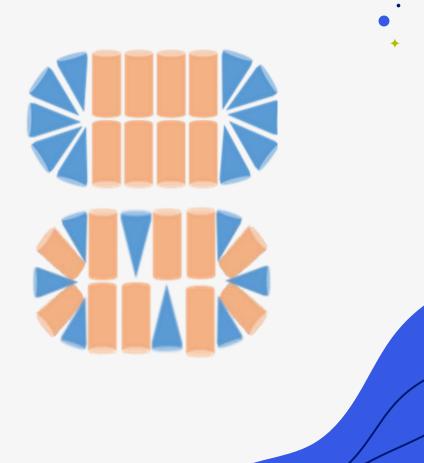


Research Question

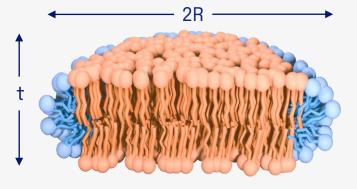
Is the ideal model correct? What is the true segregation of bicelles? Quantify?

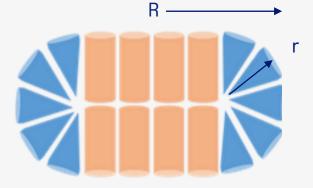


Fully Segregated? Fully Mixed?



Radius versus q_{eff}



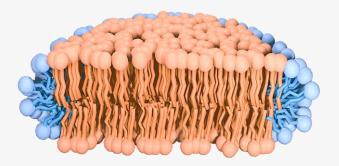


Only true for ideal model:

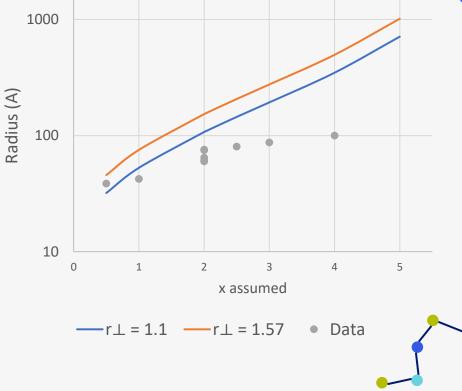
Predicts composition -> size

$$R_{v} = \frac{X_{v}r}{4} \left(\pi + \sqrt{\pi^{2} + 32/3X_{v}} \right) + r$$

Radius versus Lipid Ratio



Xeff = <u>[lipidin bicelle - lipidfloating]</u> [detergentin bicelle detergentfloating]



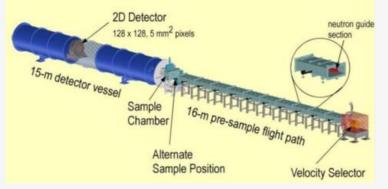
Ideal wrong!

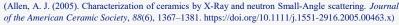
Next steps:

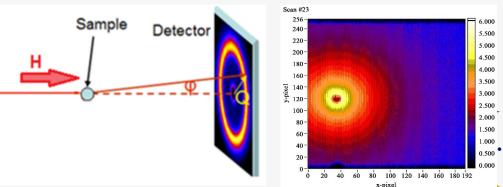
- Pick specific x value
- Continue doing analysis
- Quantify segregation

Small Angle Neutron Scattering (SANS)

- Two Instruments: NG3 and NG7
- Nuclear Generator -> Neutrons
- Long flight path = Thinner beam
- Hits Sample, Scatters
- Hits Detector



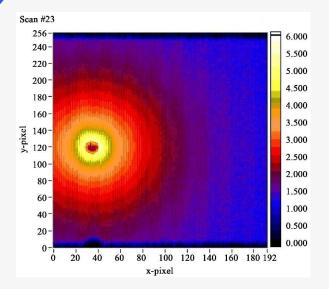


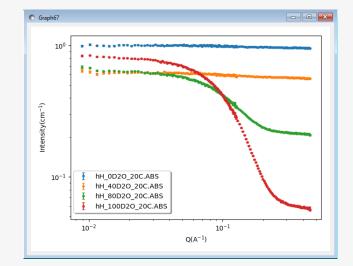


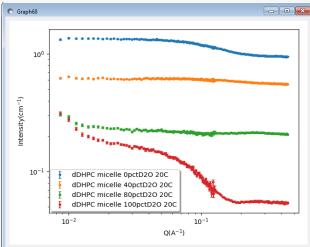
(Searching for Magnetic Skyrmions in Powdered B20 Materials with SANS / NIST. (2023, May 2). NIST. https://www.nist.gov/ncnr/2018-summer-schoolfundamentals-neutron-scattering/coursematerials/experiment-handouts-0jc

(Blach, T., & Holl, H. (2019). Small angle neutron scattering investigation of compressed bentonite plugs. *Journal of Minerals and Materials Characterization and Engineering*, 07(05), 230–260. https://doi.org/10.4236/jmmce.2019.75018)

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Why SANS?

The special thing about SANS:

- Nanoscale info + molecular composition = <u>*small*</u> angle scattering
- Contrast!





ρ = Scattering Length DensityAll Materials have scatteringlengths!



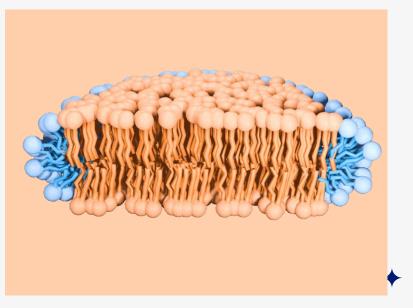


Make invisible

dDMPC contrast hDHPC

100D20 to match with dDMPC

Deuterate the Head W/ 100D20

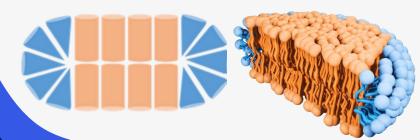


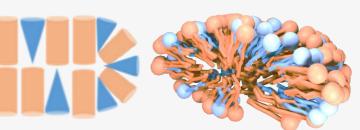
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Segregated and Fully Mixed SLDs

Segregated	Value
SLD Core	7.4
SLD Face	6.2
SLD Rim	3.5

Fully Mixed	Value	
SLD Core		5.5
SLD Face		5.7
SLD Rim		5.9

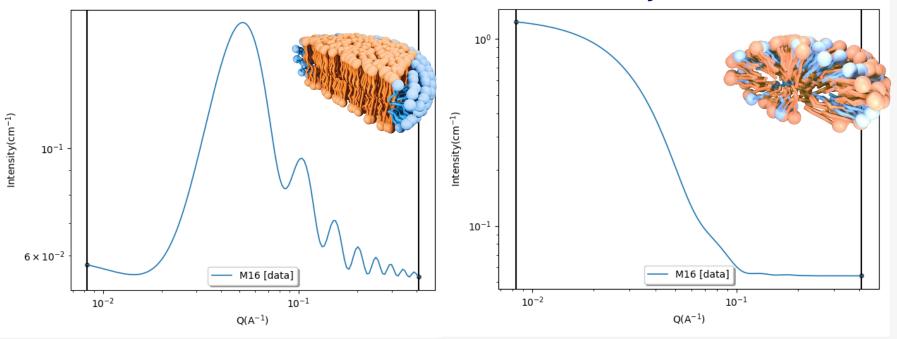






Fully Segregated

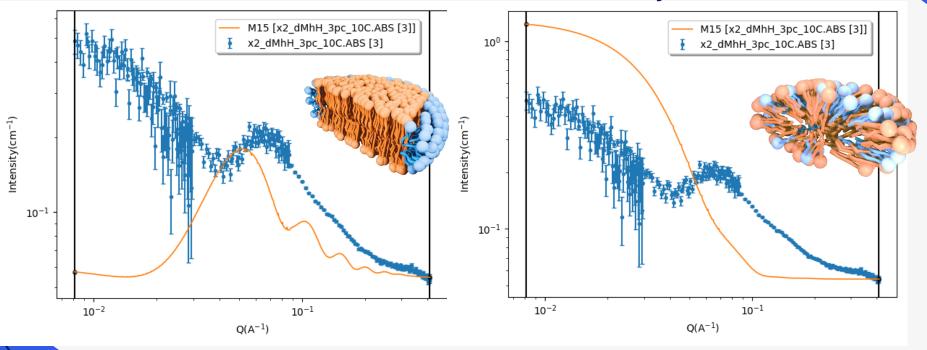
Fully Mixed

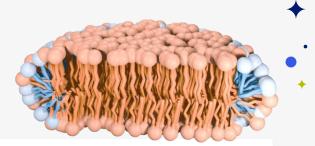


Plot Model w/ Data

Fully Segregated

Fully Mixed

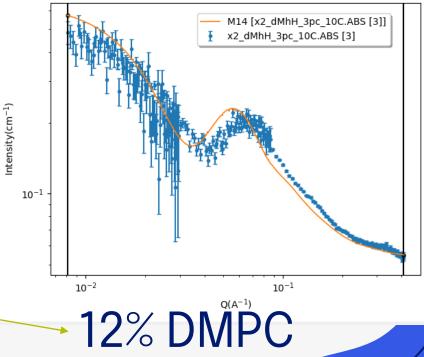




Final Fitting

dDMPC/hDHPC and dtail Lipids

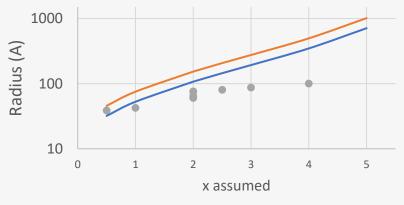
0.0273
0.054
57.4
8.3
11.8
30.4
7.3
6.13
1.44
6.4



Conclusions



- Around 12% mixing in rim
- Mixing increase as we increase x



 $r \perp = 1.1$ $r \perp = 1.57$ • Data

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Continued research needs to be made to get conclusive numbers:



SPECIAL THANKS

My Mentors: Elizabeth Kelley, Paul Butler Coordinators of SURF: Julie Borchers, Leland Harringer, Susana Teixeira, Cara O'Malley

To the people at NCNR and sharktank room members...

And to YOU who sat through this!



