

Mechanics of Model Microtubules

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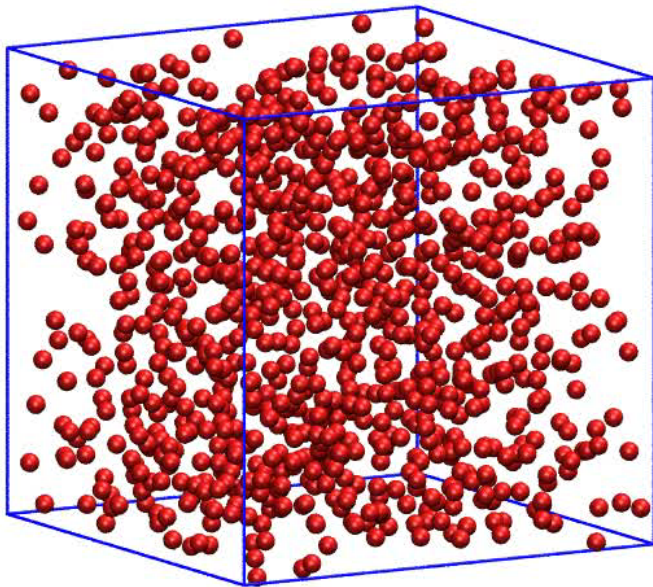
Shengfeng Cheng → Virginia Tech

Funding: Basic Energy Sciences



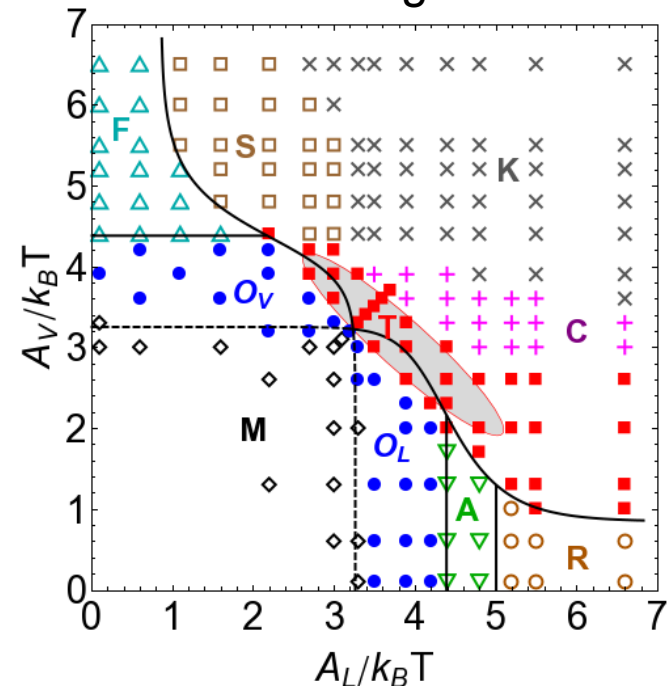
MD Simulations

- standard NVT MD, ~ 1 billion time steps
- 1000 **achiral** wedge monomers
- Starting state: with $A_L = 4.2$ kT and $A_V = 3.0$ kT
- Wedge shown as single sphere
- **Monomers to tubules**



Soft Matter **8**, 5666 (2012).

vary lateral A_L and vertical A_V interaction strengths



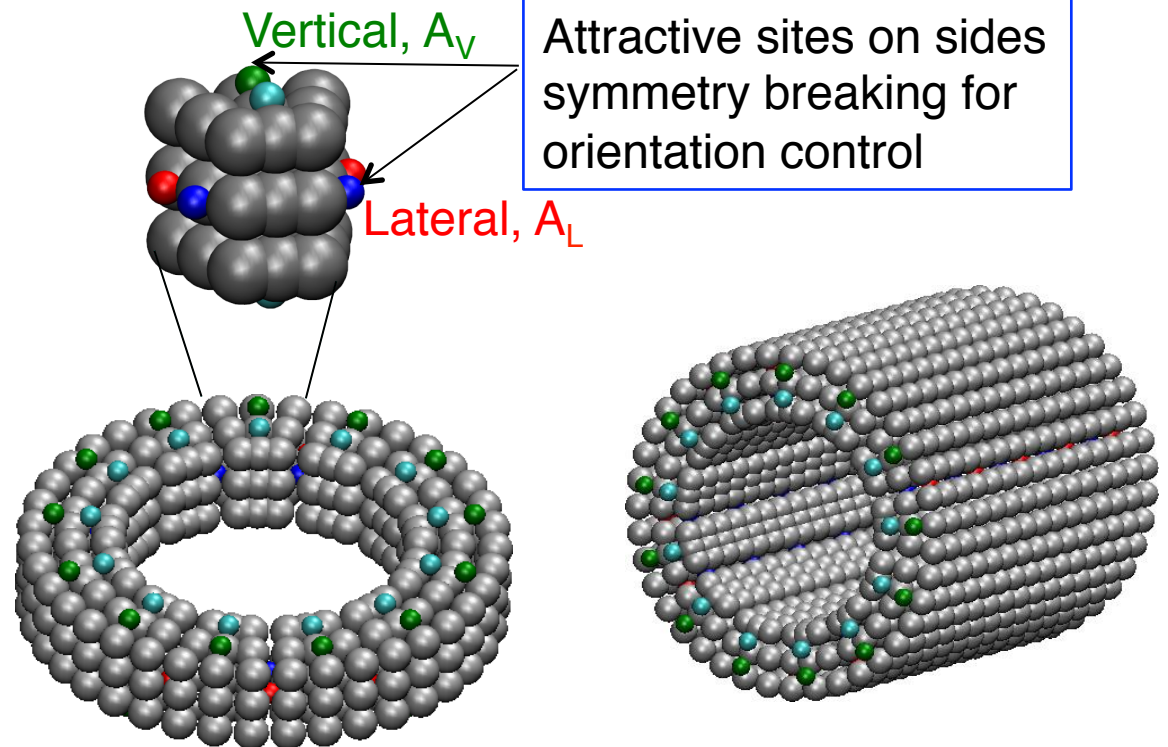
Monomer Model

Wedge monomer

- shape designed to produce rings that stack into cylinders
- 13 wedges/ring
- gray particles define shape
- rigid body
- gray particles interact purely repulsively
- attraction only between sites of same 'color'
- implicit good solvent

monomer

- (rigid) body
- shape (not spherical)
- surface-surface interactions

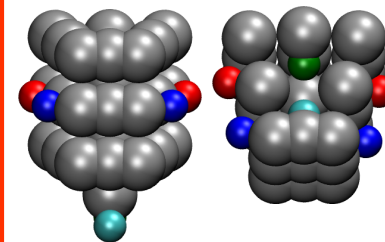
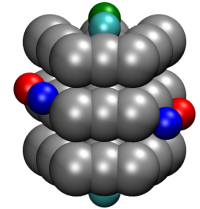


Tubule Pitch

- How control pitch?
 - chiral monomer
 - reduce variation in pitch
- Twist misaligns vertical placement
 - increase A_v ?
 - lock & key

Chirality c

Shift positions of lateral site up/down to produce chirality



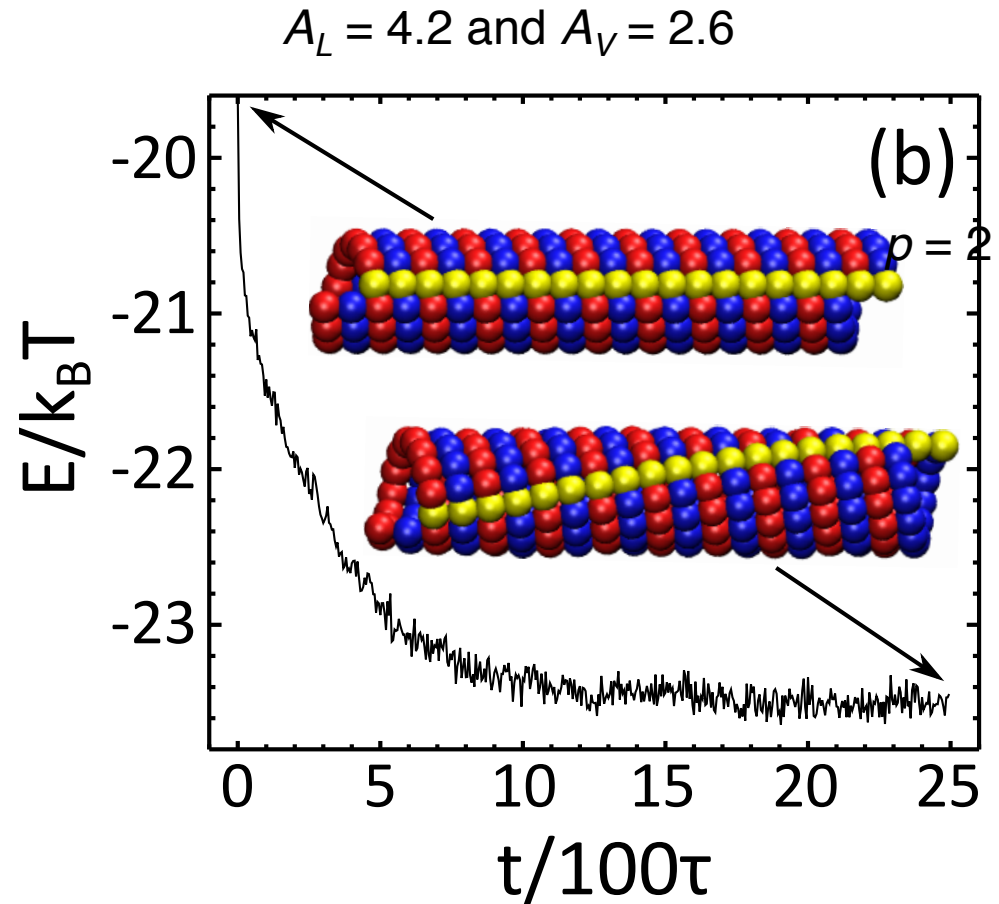
Lock & Key LK

Push central column down



Twist & Energy

- Build tubule with pitch p
- Do MD
- Calculate energy distribution
- Tubule twists
- Energy drops
- Stabilizes structure with pitch



Schematic of Beam Bend

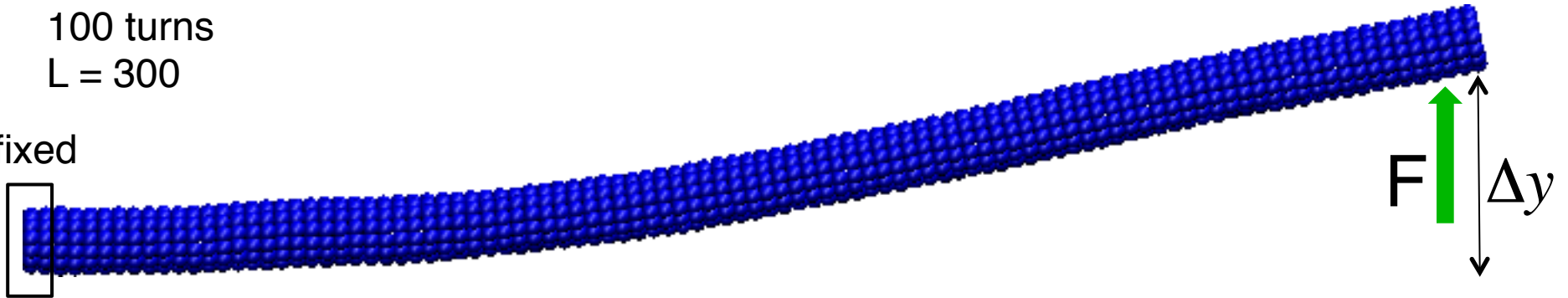
$$A_V=4.2 \ A_L=2.6$$

no chirality

100 turns

$$L = 300$$

fixed

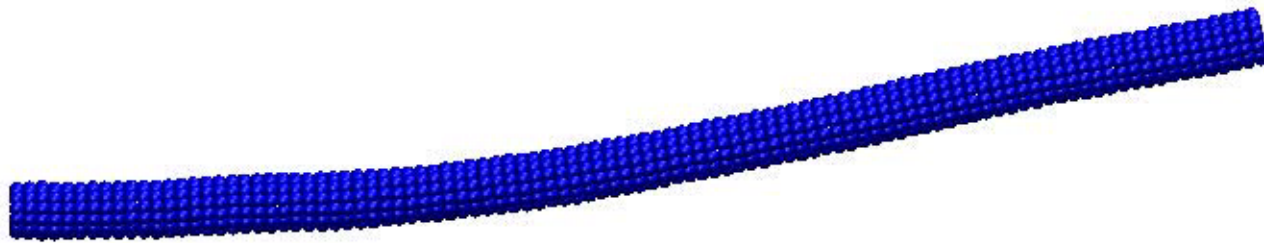


$$\Delta y = F \frac{L^3}{3EI}$$

$$E = F \frac{L^3}{3I\Delta y}$$

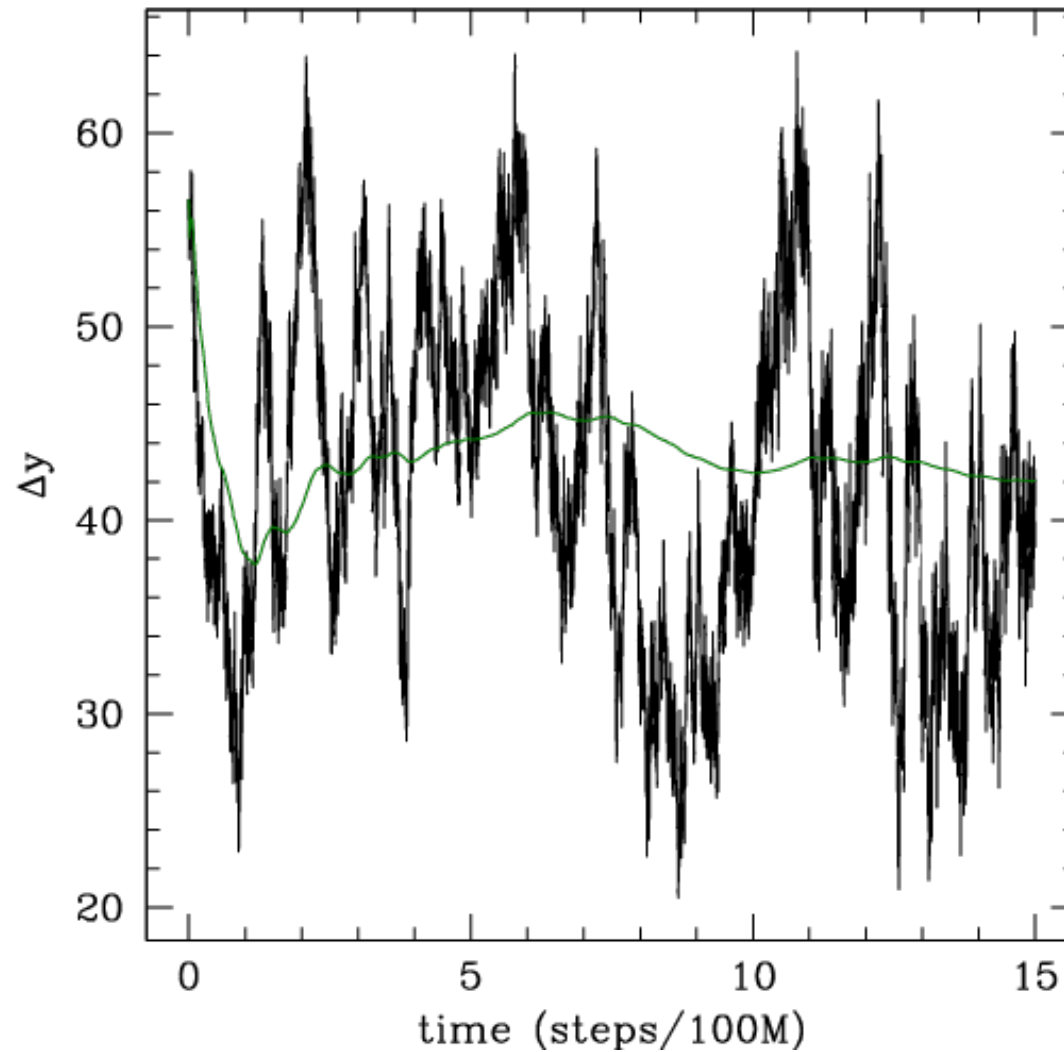


Dynamics



Time dynamics

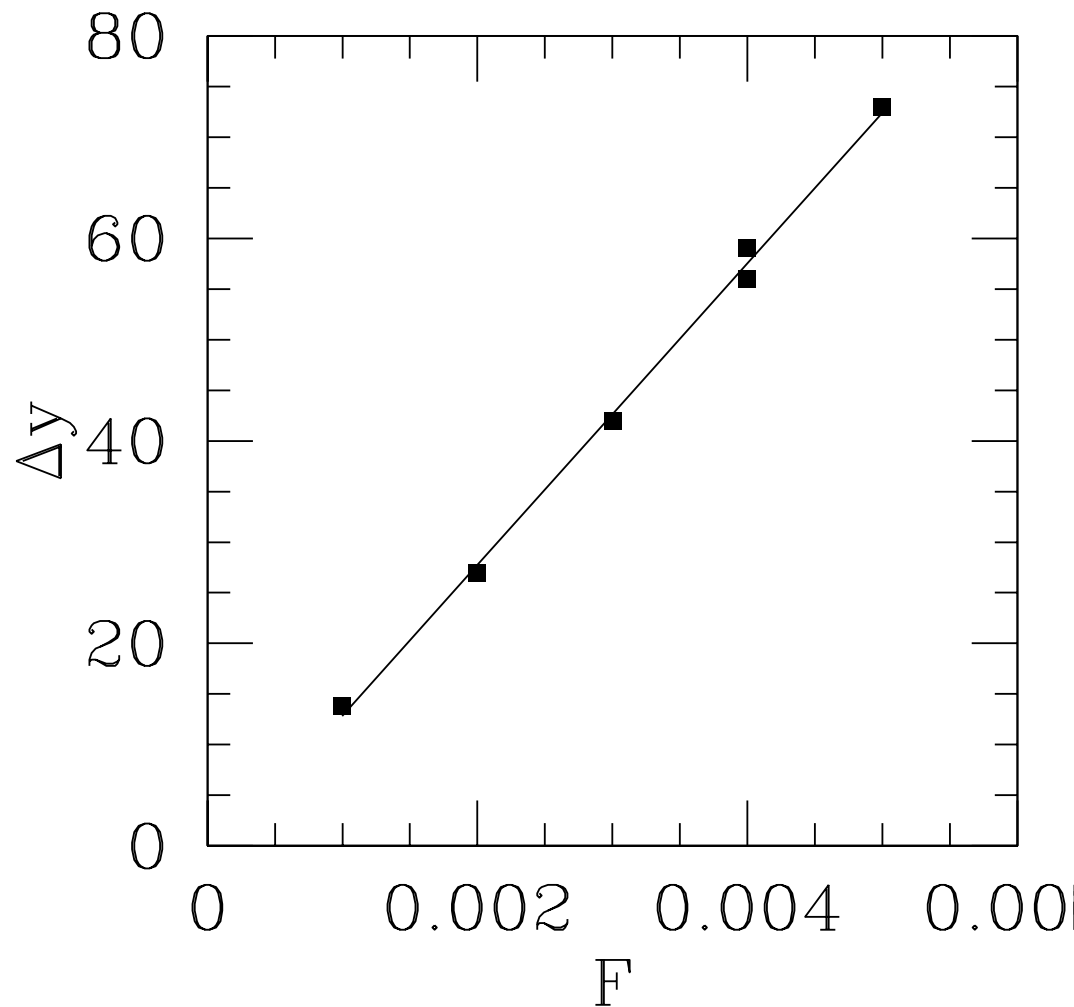
100 turns
chirality = 0
 $F = 0.003$



Bend vs. Force

$A_V=4.2$ $A_L=2.6$
no chirality
100 turns
 $L = 300$

$E = 0.4$



$N = 1000$

$$A_V = 4.2 \quad A_L = 2.6$$

no chirality

1000 turns

$$L = 3000d$$

Take $3d = 4 \text{ nm}$

Then have 4μ long tube.



Summary

Doing micron long tubules within experimental range.

Tubules could be stiffer.

Stiffness is related to 'rigid' body interaction + binding interaction.

Effect of chirality, lock & key ... TBD





Microtubules

monomer is α - β tubulin

13 monomers per turn

24 nm diameter

polymerization/depolymerization

protofilaments \rightarrow sheets \rightarrow tubules

catastrophe

polarity

binding involves GTP/GDP

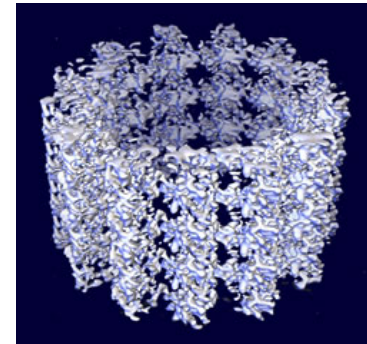
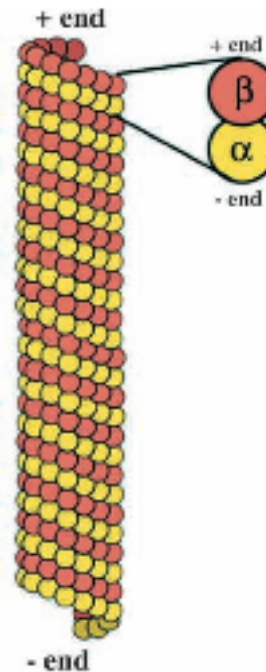
motor proteins walk on MT

tubulin polymers

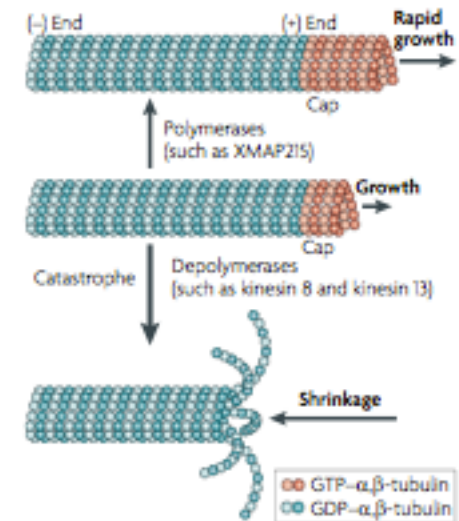
straight: protofilaments, GTP bound

curved: depolymerizing protofilaments/GDP bound

Microtubules are an example of hierarchically assembled structure with many interesting features.



GTP cap



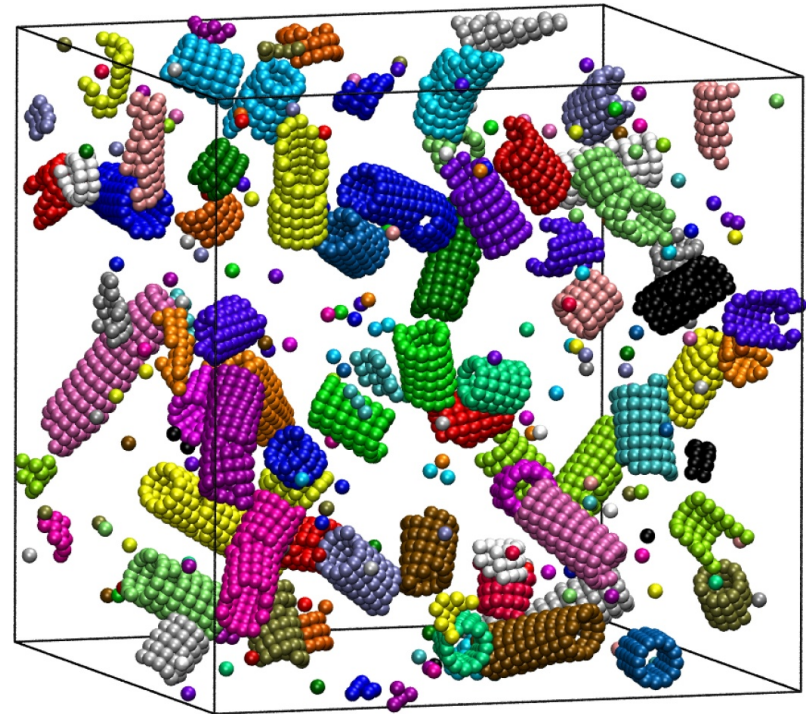
What are the factors that control assembly?

What features are necessary in the monomer?

The monomer is going to be a nanoparticle or a macromolecule (e.g. dendrimer) with large surface area.

- How do monomer shape and structure influence polymer structure (i.e. tubule).
- Where are interaction sites placed?
- How strong should they be?

Proteins as monomers possess a complex set of interactions.

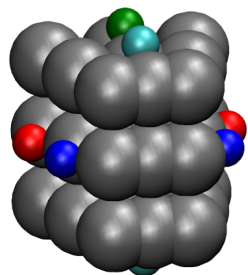


Theoretical Structure Diagram

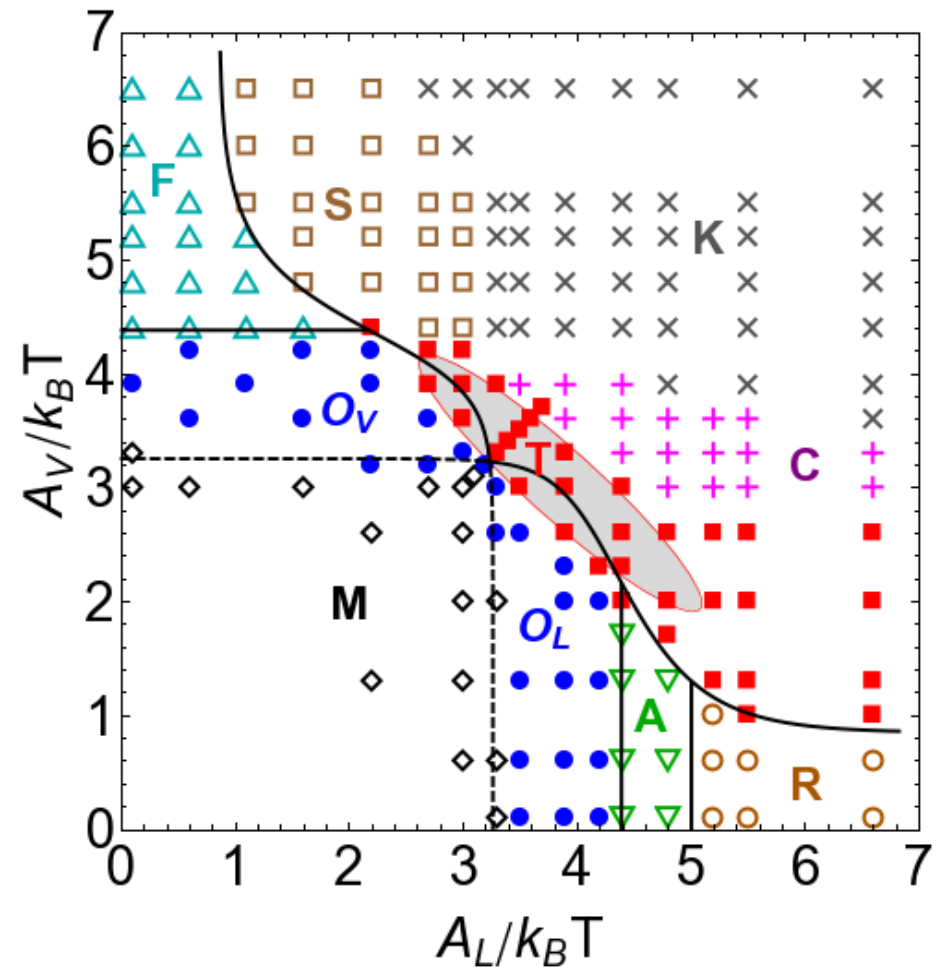
Each wedge is represented by a sphere.
Color represents cluster size.

- M) no assembly
- O) oligomers
- F) filaments
- A) arcs
- R) rings
- S) sheets
- K) kinetically trapped (gunk)
- C) clusters
- T) Tubes

vertical, A_v



Lateral, A_L



Soft Matter, 2012

