# Symmetry Breaking of Actin Networks Governs Confinement Pressure





500 Filaments,

1000 Crosslinkers

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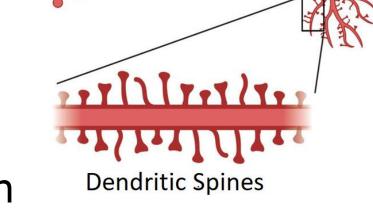
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## Motivation

### Dendritic Spine Morphology

found in numerous biological systems, i.e. the cell cytoskeleton of dendritic spines. Characterizing the biological phenomena that modulate actin

Actin fiber networks are

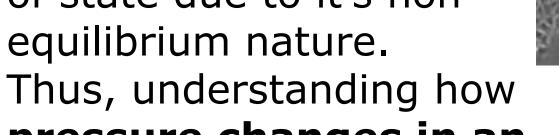


network dynamics can enhance our understanding of dendritic spine growth/shrinkage, which is known to affect fundamental learning

and memory processes.

#### Active Matter

Active matter does not have a specific thermodynamic equation of state due to it's nonequilibrium nature.

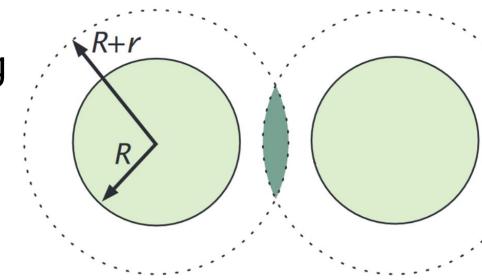


pressure changes in an active system is highly nontrivial.

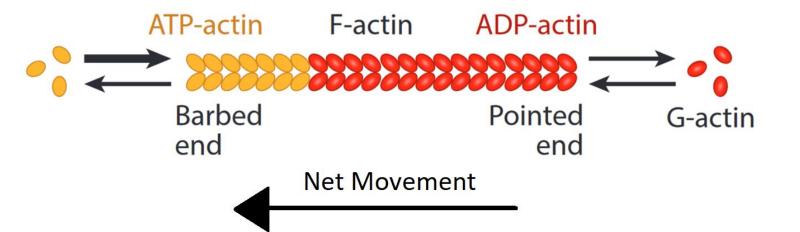
## Biophysical Factors

#### Steric Interactions (Volume Exclusion)

Volume excluding gives rise to depletion (entropic) forces.



#### Treadmilling

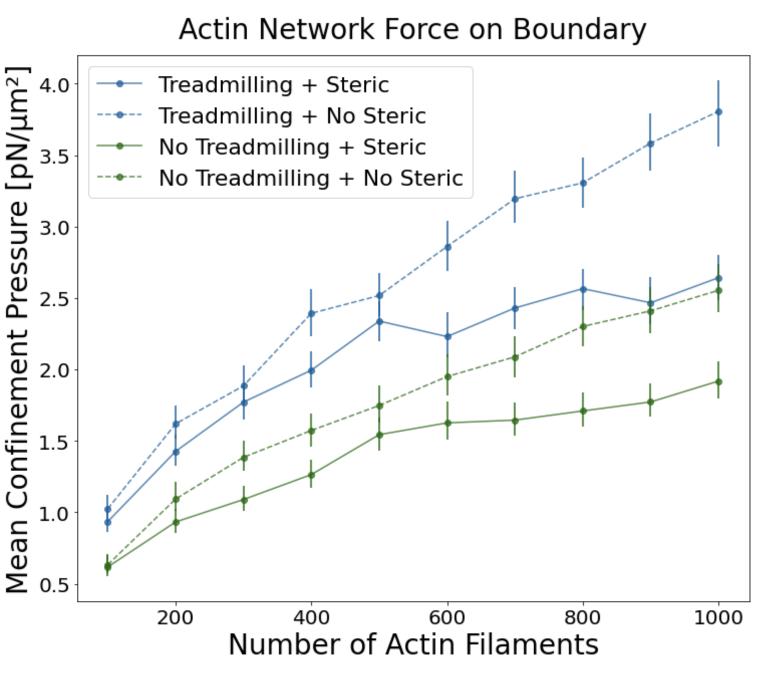


Momomer-filament association/disassociation causes filament to undergo net movement towards the plus end.

#### Crosslinkers

Proteins which are known to affect several physical properties of polymer systems, such as gelation and contraction.





translational symmetry breaking

800 Actin Filaments

— Treadmilling, Steric

----- Treadmilling, No Steric

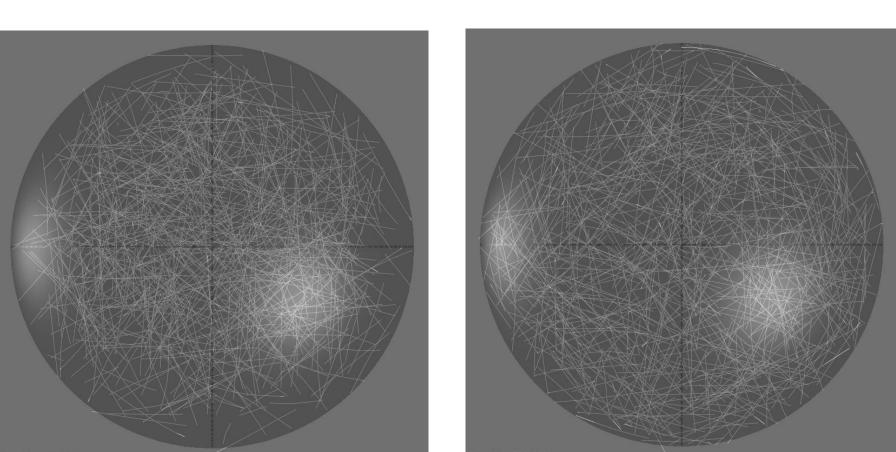
No Treadmilling, Steric

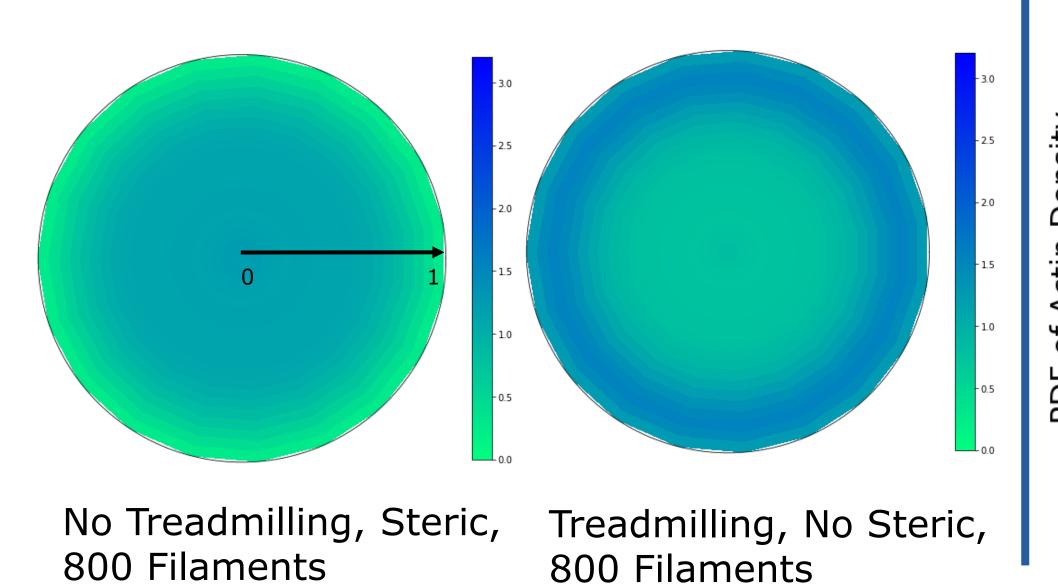
----- No Treadmilling, No Steric

Treadmilling induces

- Treadmilling applies a larger pressure to the boundary
- Depletion forces from steric interactions (volume exclusion) decreases the pressure on the boundary

Treadmilling, No Steric, No Treadmilling, Steric, 800 Filaments 800 Filaments



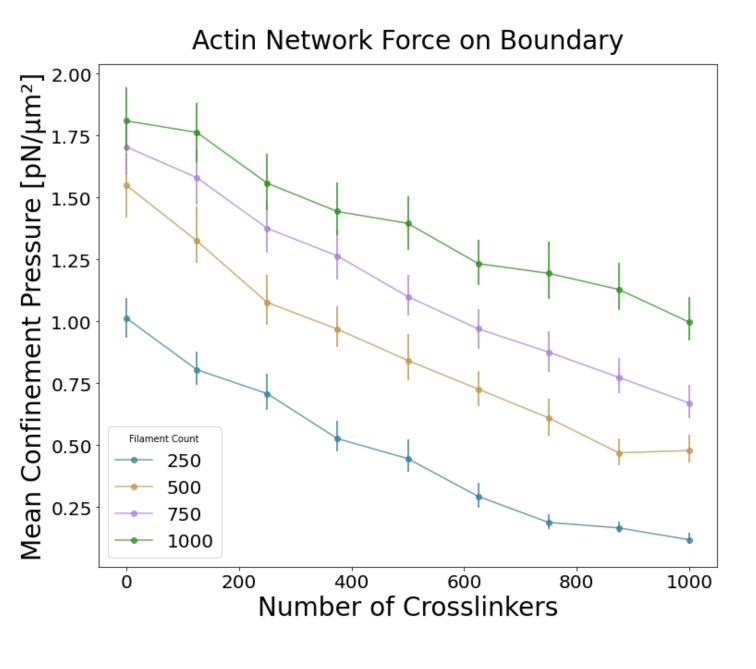


## Crosslinker to Filament Ratio Regulates Cluster Formation

500 Filaments,

250 Crosslinkers

Linear decrease of mean confinement pressure observed when increasing the number of crosslinkers



500 Actin Filaments

R [µm]

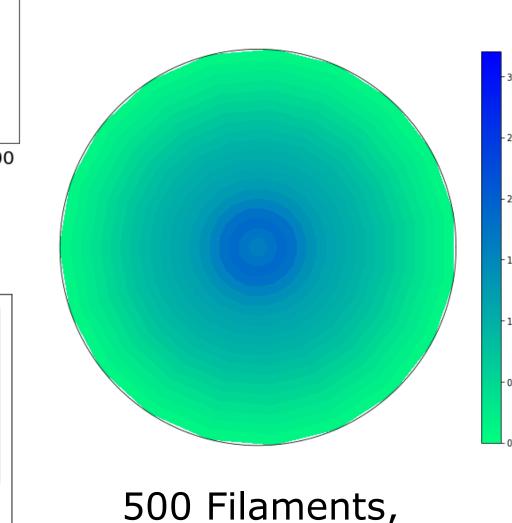
— 0 Crosslinkers

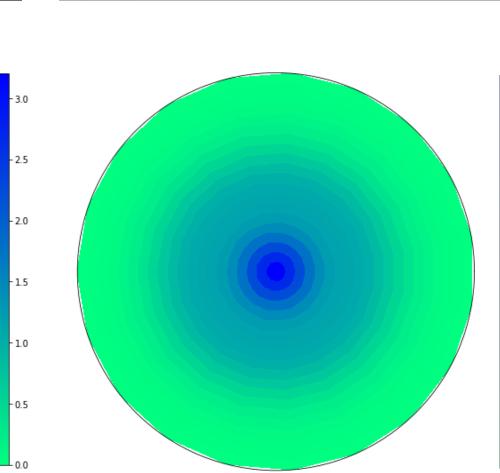
- 250 Crosslinkers

500 Crosslinkers

750 Crosslinkers

— 1000 Crosslinkers





500 Filaments, 250 Crosslinkers

1000 Crosslinkers

500 Filaments,

- Greater ratios of crosslinkers to filaments cause cluster formation towards the center
- After the gelation point, translation symmetry breaking occurs

## Cytosim

Cytosim is used to simulate biologically-relevant cell cytoskeletons and their associated proteins. Filaments, spheres, and solids are represented by a finite number of points. Cytosim uses the Collective Langevin Dynamics equation of motion:

Eq. (1)  $dx = \mu F(x, t) dt + dB(t)$ 

R [μm]

This model considers:

- $\mu$  Mobility coefficients
- $F(\mathbf{x}, t)$  Forces such as bending elasticity, constraints applied by the segment links, etc.
- B(t) The effect of brownian motion

## plus end fixed length bending elasticity **√** minus end

#### Conclusion Confinement Pressure [pN/µm²] **PDF Steric Interactions/** Decrease Minor Perturbation **Volume Exclusion** Towards Boundary Increase **Treadmilling Increasing Crosslinker:** Decrease **Towards Center Filament Ratio**

## Future Work

Varying the:

- Filament length to system size ratio
- Péclet number of the system
- Addition of:
- Branchers (Arp 2/3 complex)
  - Motors



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#### References

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For contact information, technical details, and simulation videos, scan:

