Below is an oral presentation I gave with my research group for SOURCE 2023. The primary goal of this presentation was to tell the story of our research in such a way that a broad undergraduate audience could understand. To create this presentation we started with a poster that we had brought to the Biophysical Society Meeting 2023 as it would contain nearly all the information we felt was relevant to the project. The only thing this poster was missing was adequate background for a non-professional audience. In adding that background some finer details about the model were cut. We made this decision with the belief that our audience would be more interested in contextualizing results in terms of the biological system. To deliver this broad background effectively our presentation is mostly image and video based using cartoon models to illustrate behavior leading up to clearly labeled plots to indicate results. This is the fourth official revision with each one having significant changes from its predecessor. Not to mention the many practice attempts and script changes. We gave this presentation at SOURCE 2023 in front of physics majors, chemistry majors, and biology majors which matched our expectations for a broad audience. The presentation went very well and I am very fortunate for having that experience of professional public speaking.

Computational Models of Self Organization of Neuronal Cytoskeleton

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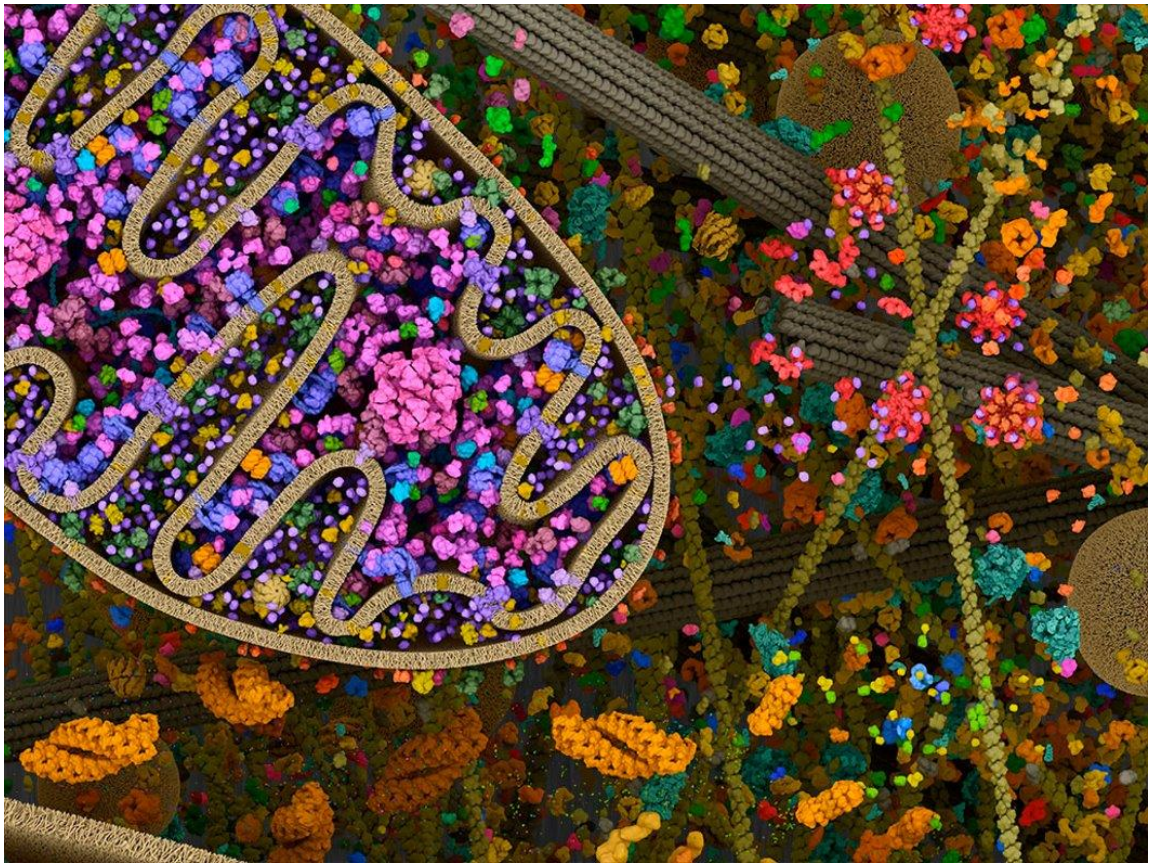
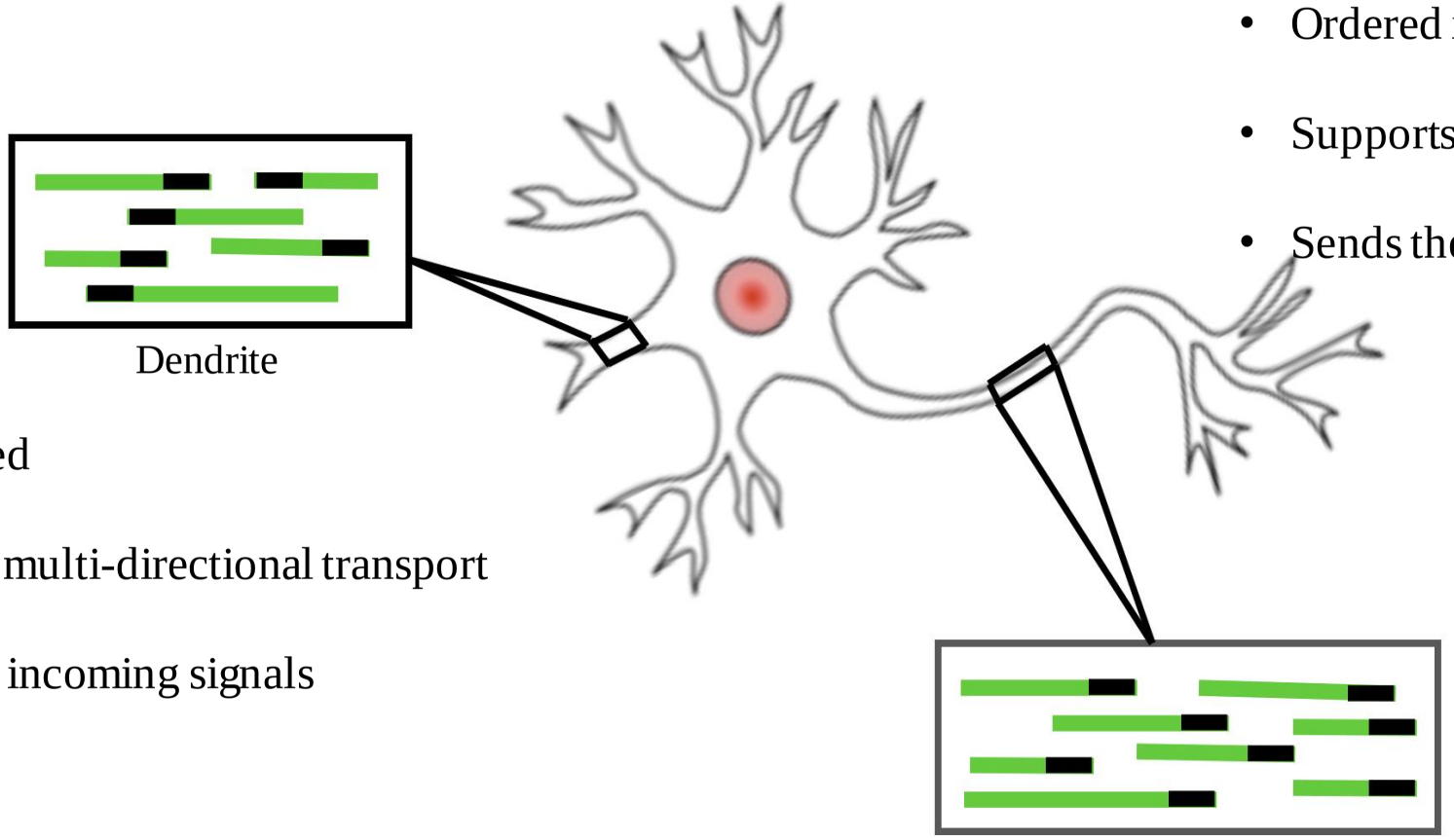
CWU Computational Biophysics Lab

Physics of Cell Biology



- Crowded
- Highly viscous
- Intricately-timed dynamic processes
- Physical principles determine biological function

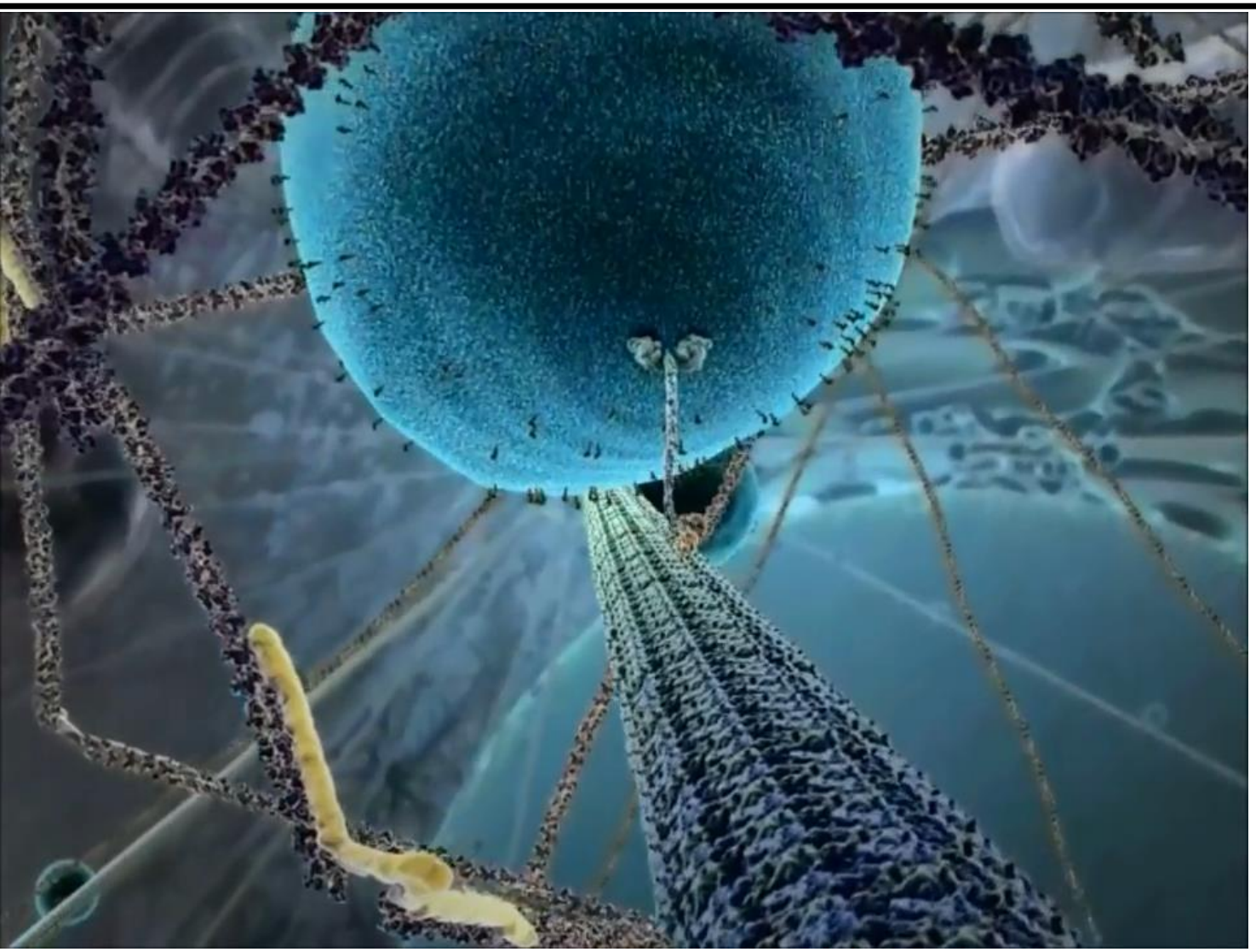
Microtubule Organization in the Neuron



- Unordered
- Supports multi-directional transport
- Receives incoming signals

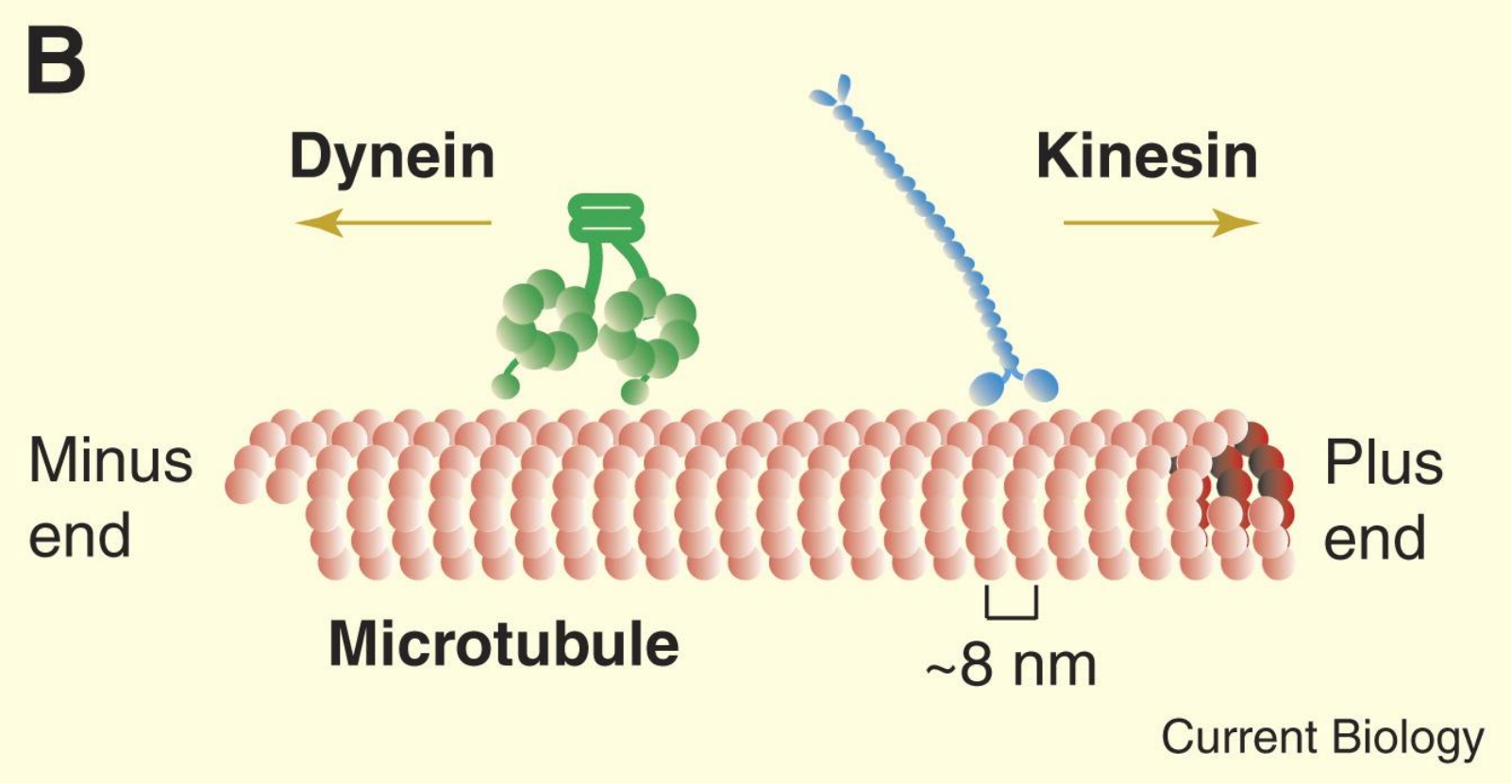
- Ordered in healthy neurons
- Supports one-way transport
- Sends the outgoing signals

The Role of Motor Proteins



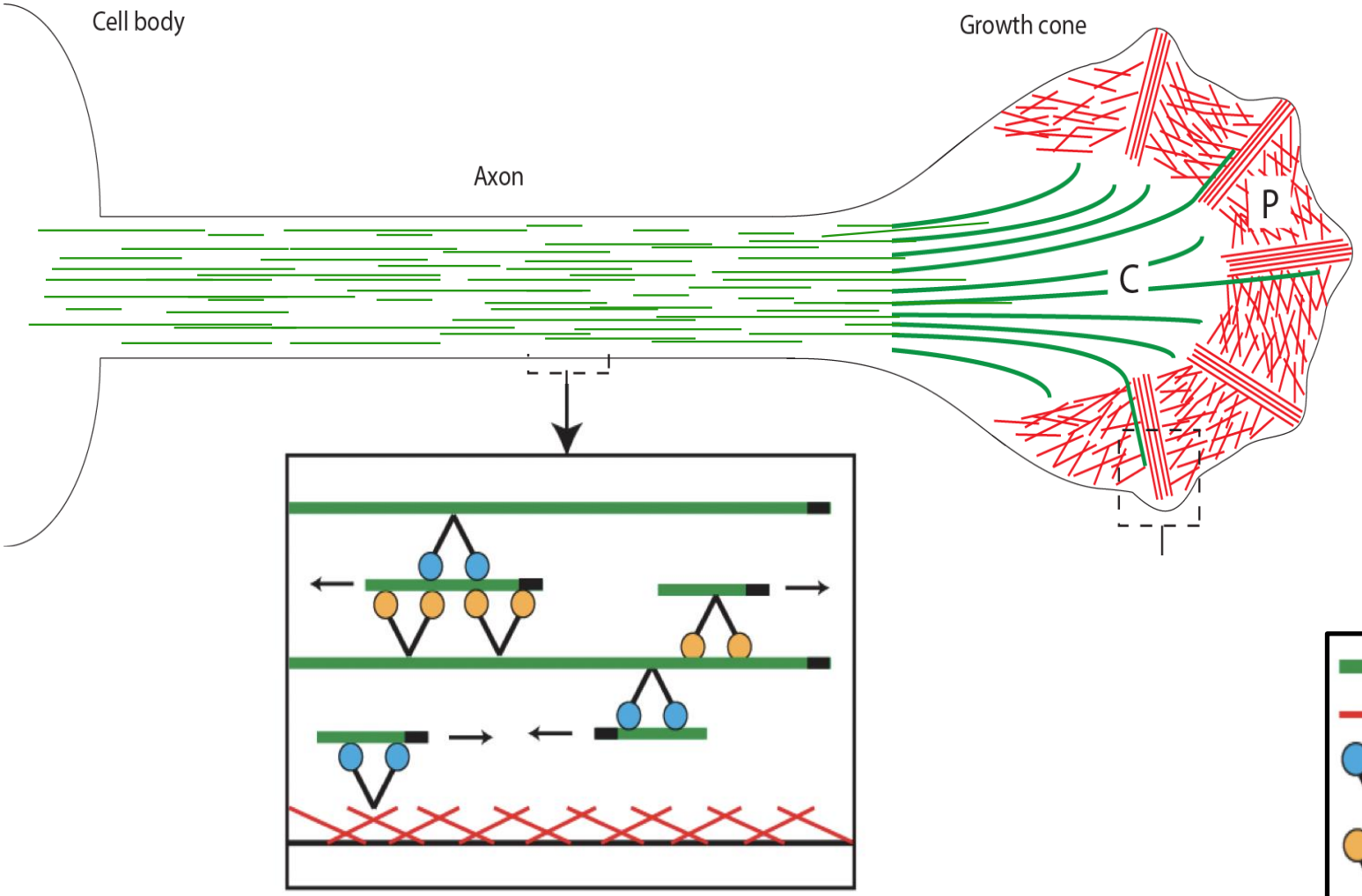
- A motor protein carries cargo along a microtube
- The motion of the “feet” has been smoothed; on the cellular level it is random with a tendency towards forward motion

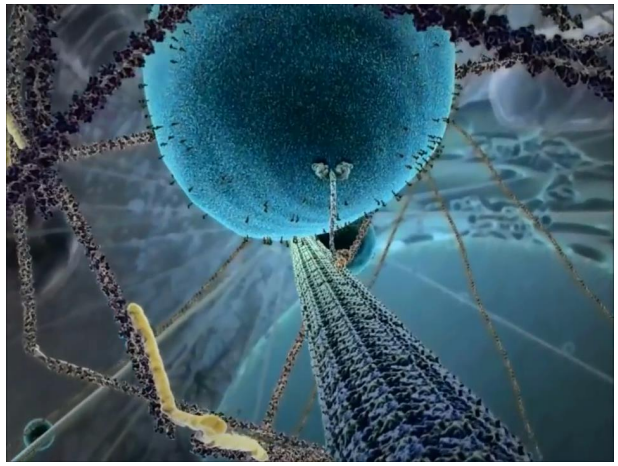
The Type of Motor Proteins



Each motor has a preferred direction to walk on the microtube. Kinesin walks in the same direction the microtube points while Dynein walks in the opposite direction.

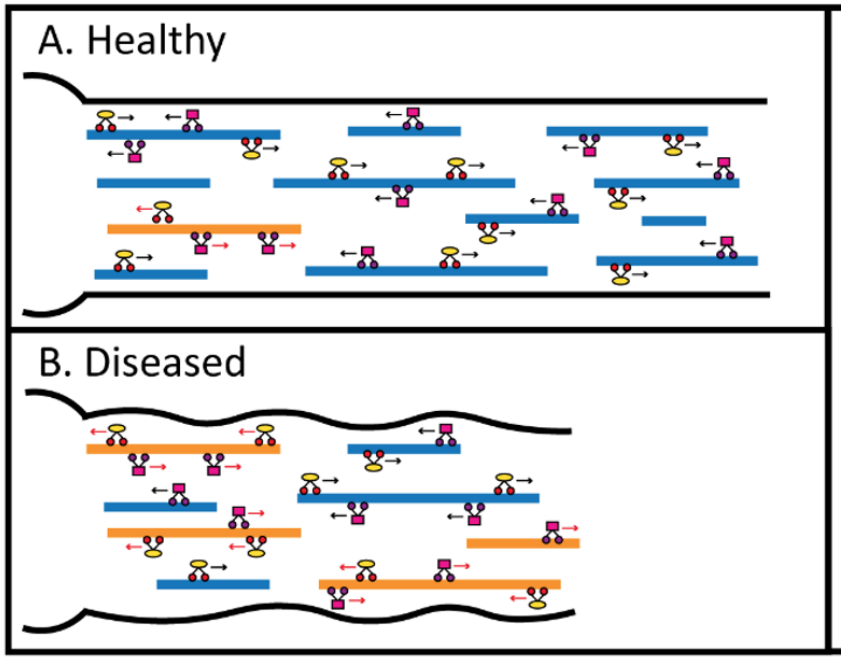
Microtubule Organization in the Axon

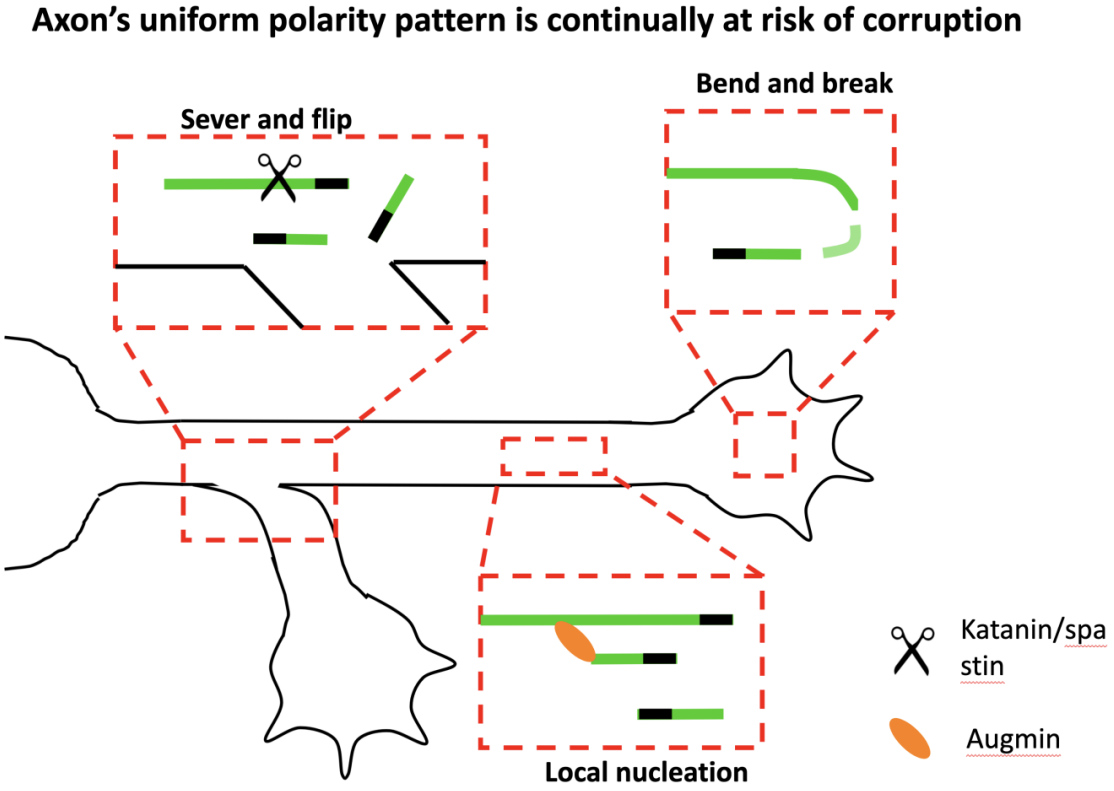




Microtubule (black tip = +-end)
Actin filament
Minus-end motor (e.g. dynein)
Plus-end motor (e.g. kinesin)

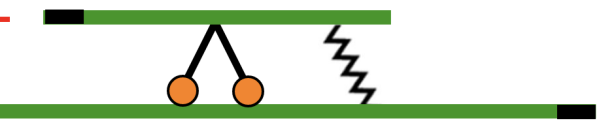
Corruption of the Polarity Pattern

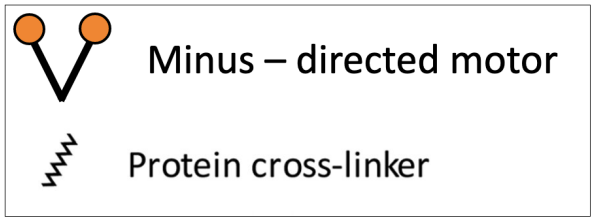




Damage to the polarity pattern may arise from a variety of sources. Significant local damage to the polarity pattern impacts cargo transport and creates traffic jams of molecular motor proteins.

Computational Model for Forces on Microtubes





nber of attached cytoplasmic dynein motors, N_d:

$$\frac{dN_d}{dt} = r_{d,on} - r_{d,off}N_d$$

achment
nds on # of
able sites)

Detachment
(Force-dependent)

Balance of forces:

$$F_{d+} = F_{d-} + N_x\gamma v + \xi L v$$

Motor forces
(+ end leading)

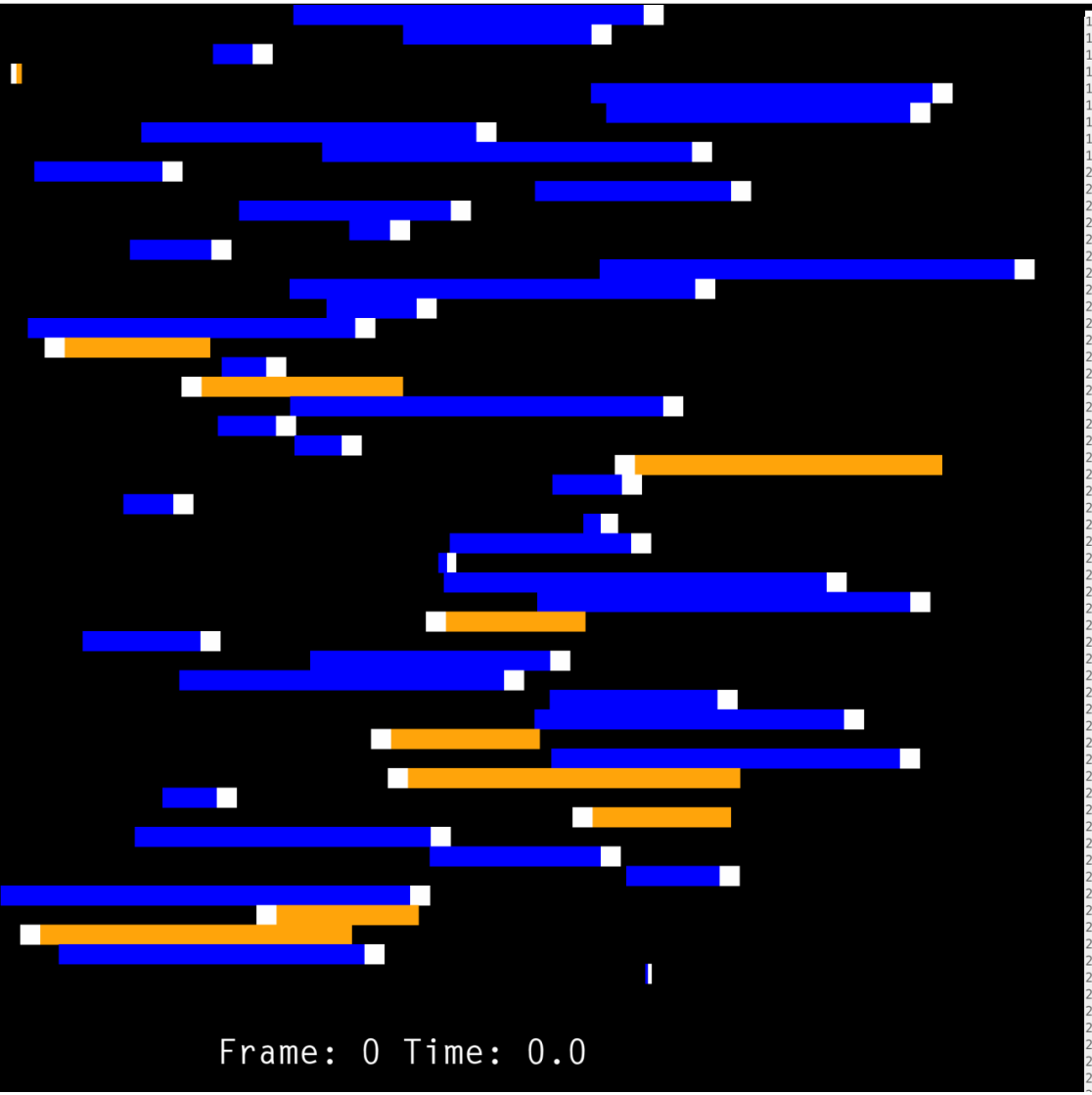
Motor forces
(- end leading)

Resistance from
cross-linkers

Filament drag
force

- Net 0 force assumption due to high fluid viscosity
- Essentially objects do not move except when under active forces

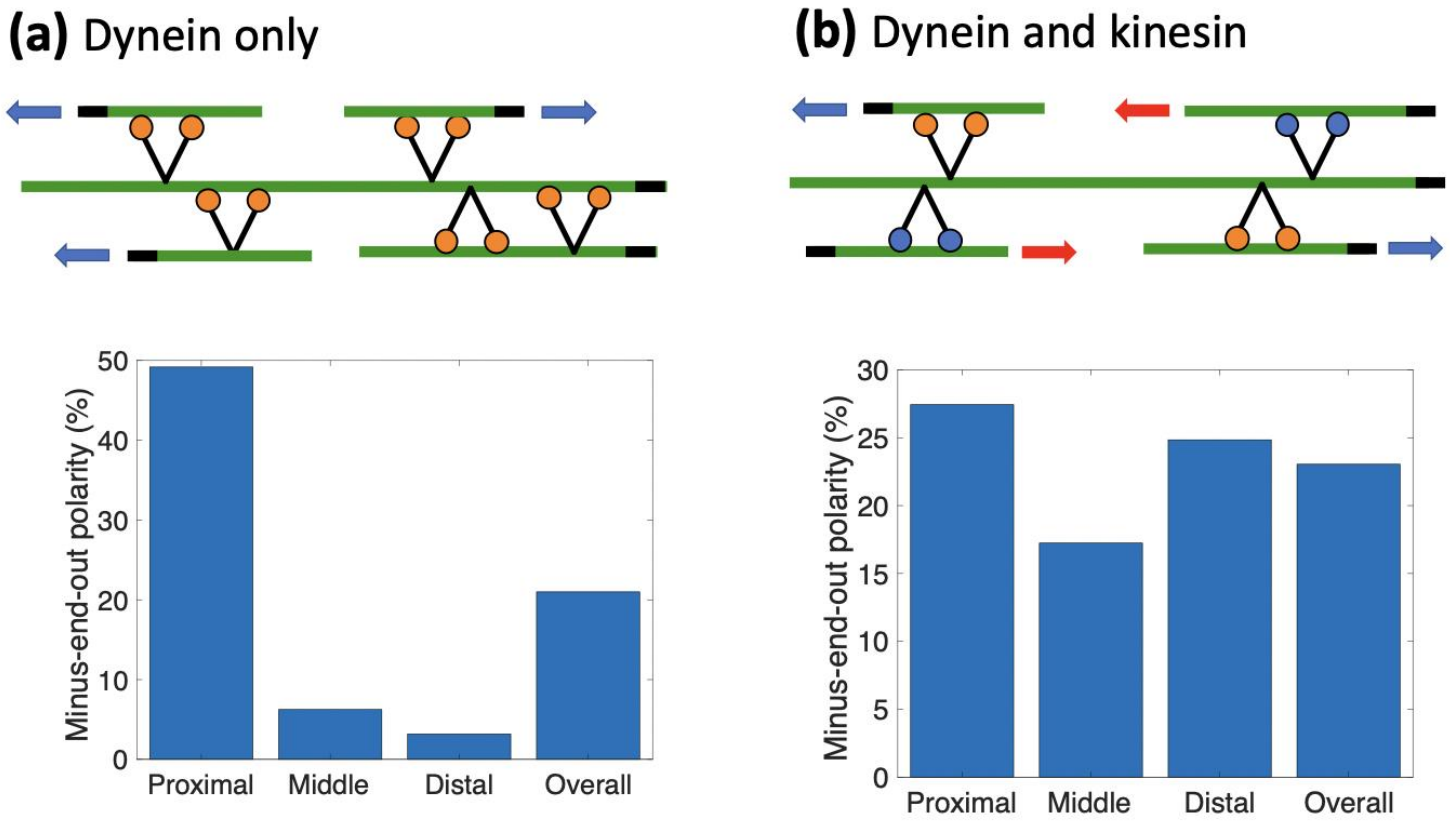
Computational Model for Forces on MTs




```
% Update position at time i+1 based on position and velocity at time i

A(i+1,2,j)=A(i,2,j)+A(i,3,j)*dt; % Translocate MT based on vel calculated for this time step
A(i+1,11,j)=A(i,11,j); % Default is for orientation to stay the same. Several possibilities later for this to flip.
% Apply conditions for if MT reaches a boundary
if (A(i+1,2,j)<0) % MT has been cleared from axon into cell body
    'MT cleared';
    % Replaced cleared MT with a new short growing MT, random
    % location and orientation
    A(i+1,2,j)=Laxon*rand([1,1,1]); % random location
    A(i+1,3,j)=0.0; % Initial velocity (v)
    A(i+1,4,j)=0.0; % Initial force (Fdp)
    A(i+1,5,j)=0.0; % Initial force (Fdm)
    A(i+1,6,j)= 0.0; % Initial dynein attachment number, forward pulling (Ndp)
    A(i+1,7,j)=0.0; % Initial dynein attachment number, backward pulling (Ndm)
    A(i+1,8,j)=0.0; % Initial cross-linker attachment number, parallel MTs (Nxpar)
    A(i+1,9,j)=0.0; % Initial cross-linker attachment number, anti-parallel MTs (Nxanti)
    A(i+1,10,j)=0.1; % Short initial length, 0.1micron for newly nucleated MT

    % Fraction of plus-out MTs at new MT's location
    loc=round(A(i+1,2,j));
    Fmin=Polarity(loc+1,3);
    if(rand<Fmin) % Newly nucleated MT has random orientation
        A(i+1,11,j)=1; % minus-end-out
        'minus out new MT';
    else
        A(i+1,11,j)=0; % plus-end-out
        'plus out new MT';
    end


    A(i+1,12,j)=1; % dynamic and growing
elseif(A(i+1,2,j)>Laxon) % MT hits distal end
    A(i+1,2,j)=Laxon; % Can't grow further
    A(i+1,12,j)=0; % Switches to stable
end

% Update length of dynamic MTs
if(A(i,12,j)==0)
    A(i+1,10,j)=A(i,10,j);
elseif(A(i,12,j)==1)
    A(i+1,10,j)=A(i,10,j)+V_MTPoly*dt;
    if(A(i+1,10,j)>Laxon)
        A(i+1,10,j)=Laxon;
    end
elseif(A(i,12,j)==2)
    A(i+1,10,j)=A(i,10,j)-V_MTdePoly*dt;
end

% Apply conditions for if MT length shrinks to zero, nucleate new MT
if(A(i+1,10,j)<0)

    A(i+1,3,j)=0.0; % Initial velocity (v)
    A(i+1,4,j)=0.0; % Initial force (Fdp)
    A(i+1,5,j)=0.0; % Initial force (Fdm)
    A(i+1,6,j)= 0.0; % Initial dynein attachment number, forward pulling (Ndp)
    A(i+1,7,j)=0.0; % Initial dynein attachment number, backward pulling (Ndm)
    A(i+1,8,j)=0.0; % Initial cross-linker attachment number, parallel MTs (Nxpar)
    A(i+1,9,j)=0.0; % Initial cross-linker attachment number, anti-parallel MTs (Nxanti)
    A(i+1,10,j)=0.1; % Short initial length, 0.1micron for newly nucleated MT

    %
```

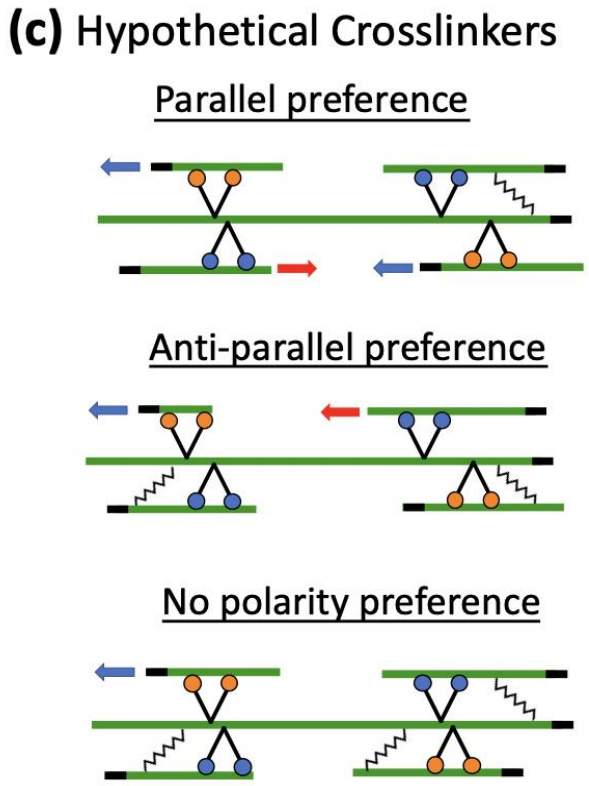
Result:

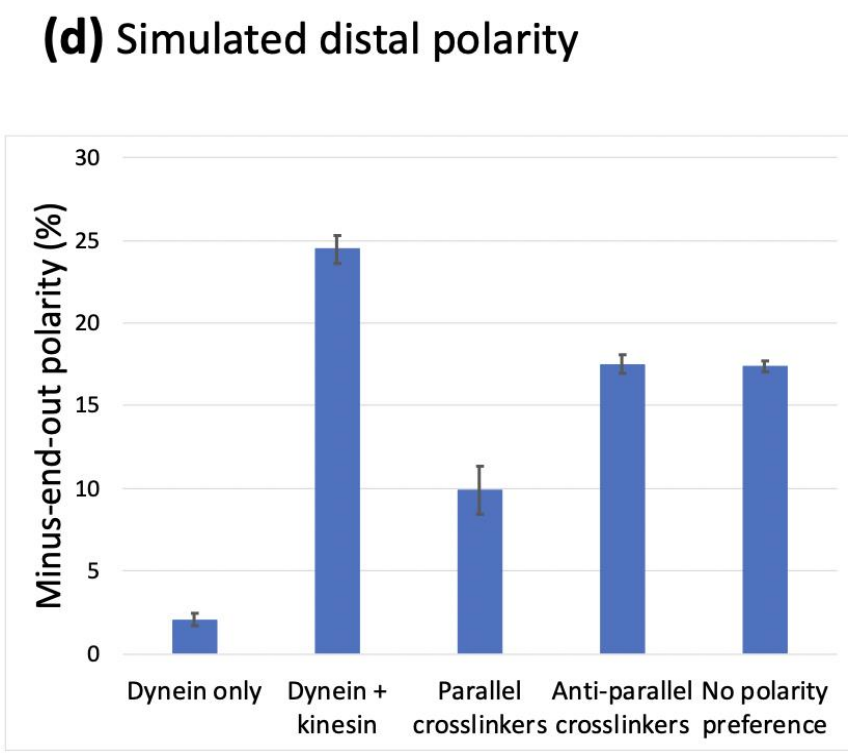
Kinesin prevents Dynein-based polarity sorting



Result:

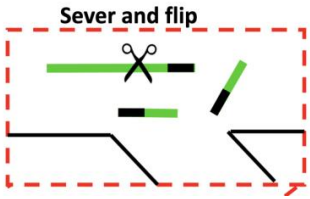
Certain types of crosslinkers impact polarity sorting more than others

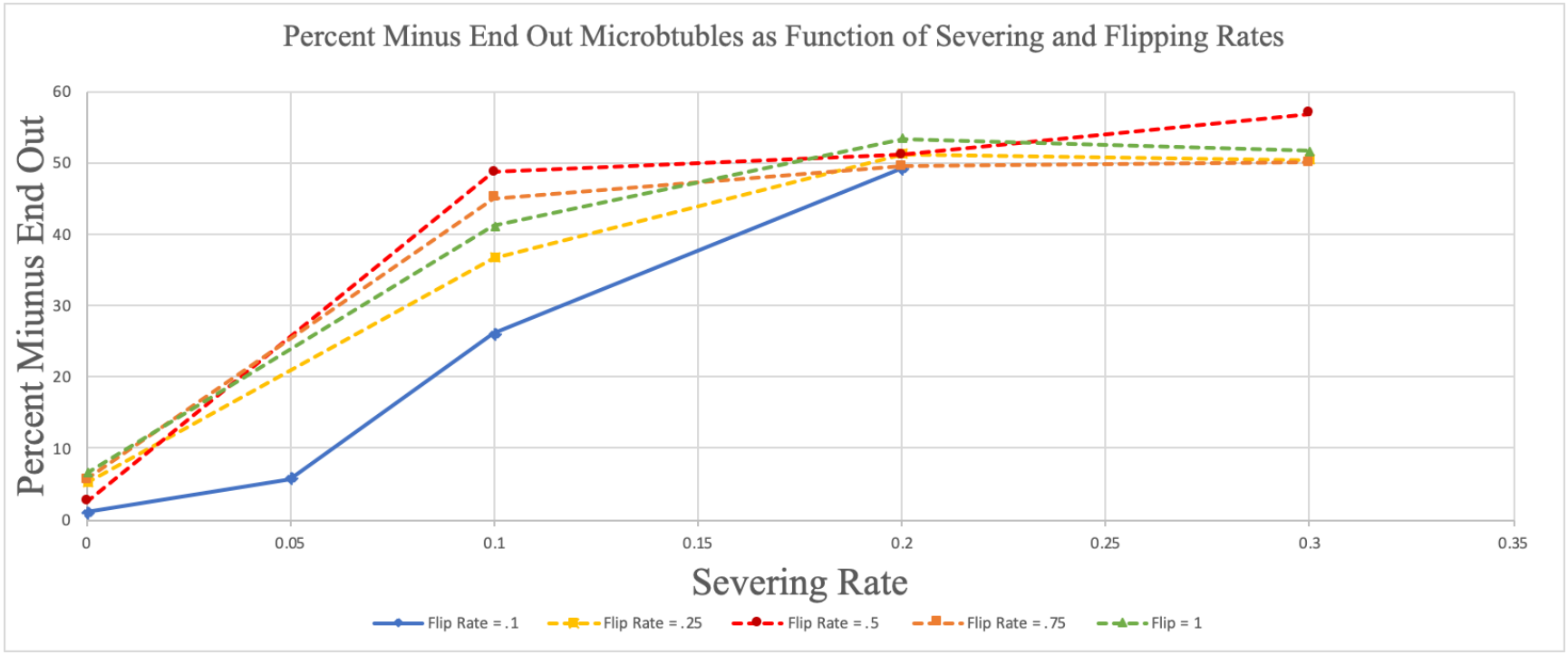




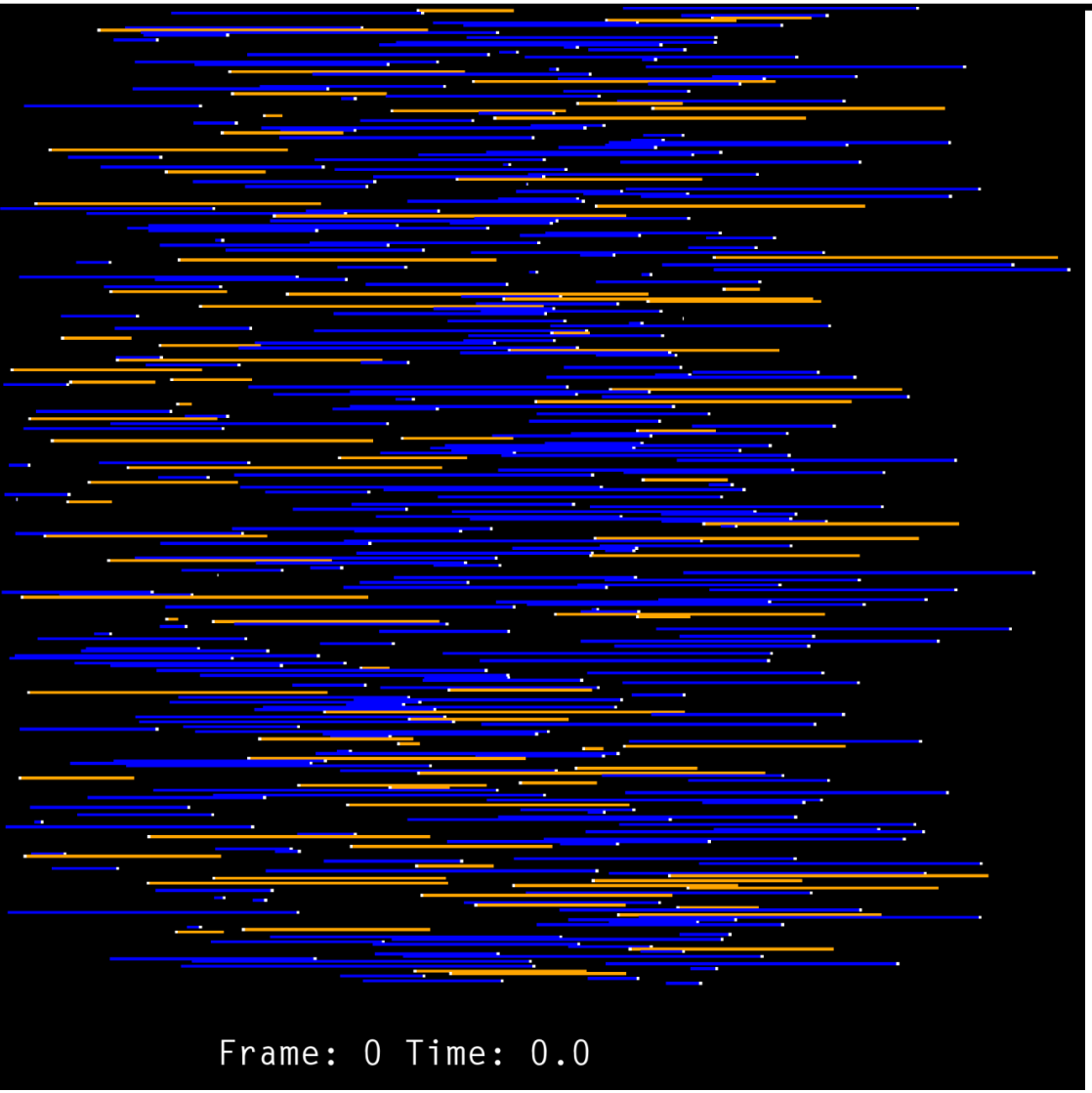
Result:

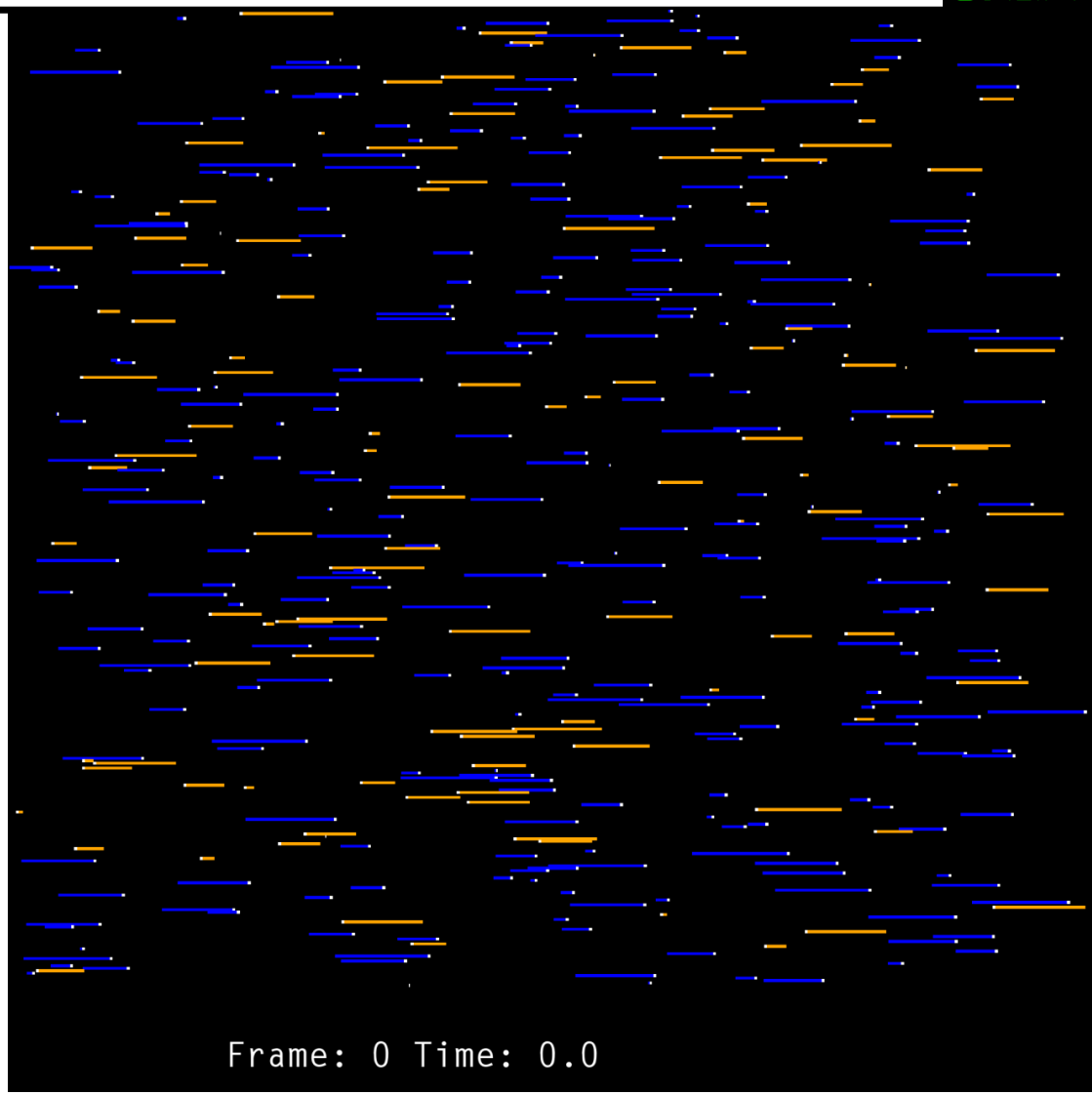
Severing and flipping are important in moderation for polarity sorting





Result:





Conclusion

- Our models show how severing and flipping impact polarity sorting, being necessary but only to an extent
- We can visualize this model to gain further intuition and demonstrate experimentally testable predictions

Next Steps

- Experimental test of model predictions.
  Related study: Rao et al., 2017.

- Investigate the impact of polarity flaws on neuronal function.
  Related study:
  Eckel et al., 2022.

Acknowledgements

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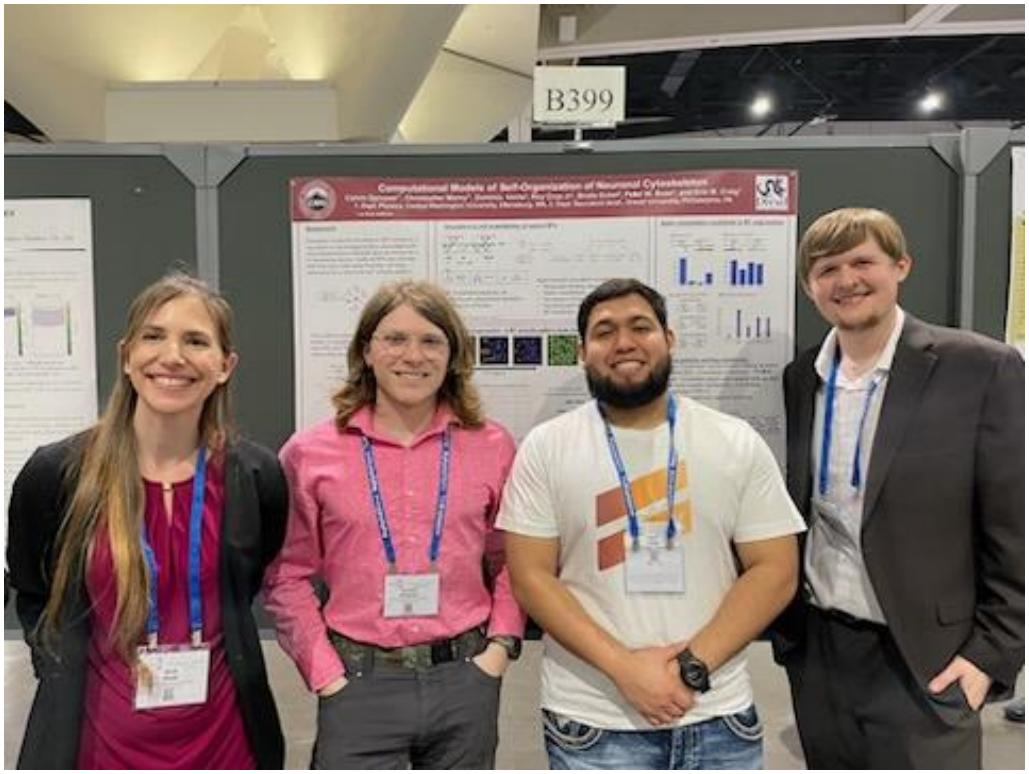

References

1. Rao, A.N. and Baas, P.W. (2017), Polarity Sorting of Microtubules in the Axon, Trends in Neuroscience, 41(2):77-88.
2. Craig et al. (2017), Polarity sorting of axonal microtubules: a computational study, Mol. Biol. Cell, 28(23):3271–3285.
3. Rao et al. (2017), Cytoplasmic dynein transports axonal microtubules in a polarity-sorting manner, Cell Reports, 19:2210-2219.
4. Eckel et al. (2022). Microtubule polarity flaws as a treatable driver of neurodegeneration., Brain Research Bulletin, 192:208-215.

See more animations here!