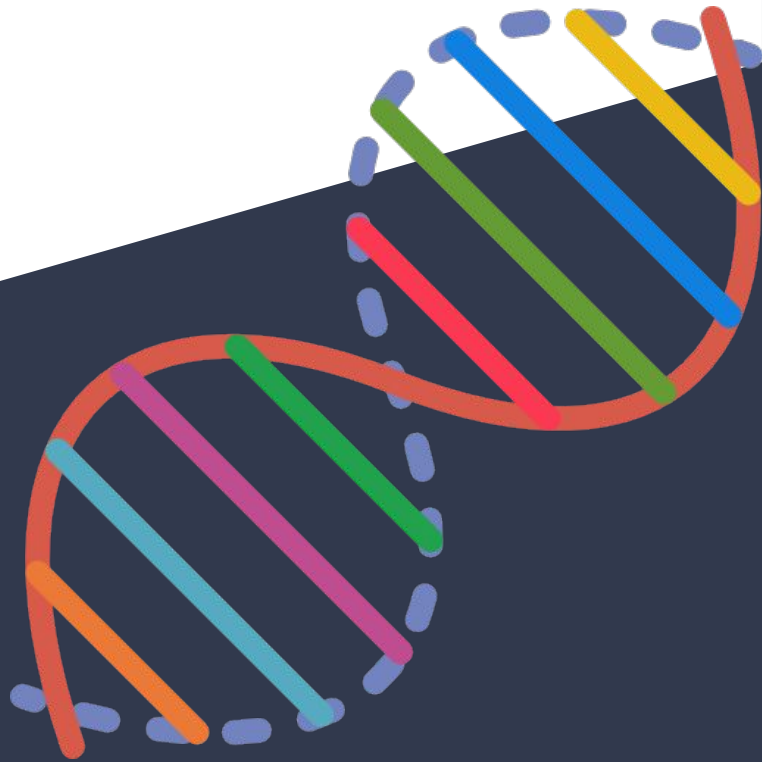
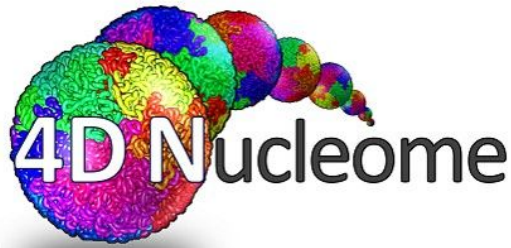
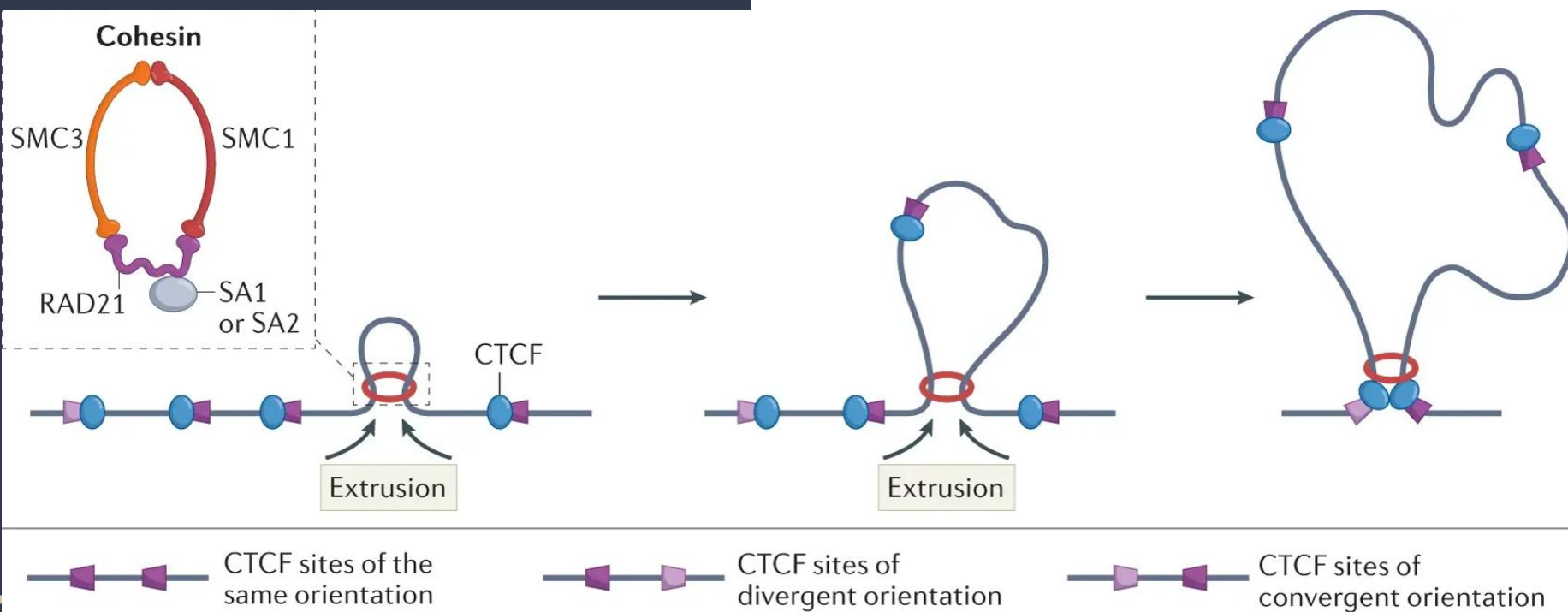


# Utilizing Machine Learning to Identify Time Asymmetry of DNA Loop Extrusion

Anna Du



# Loop Extrusion

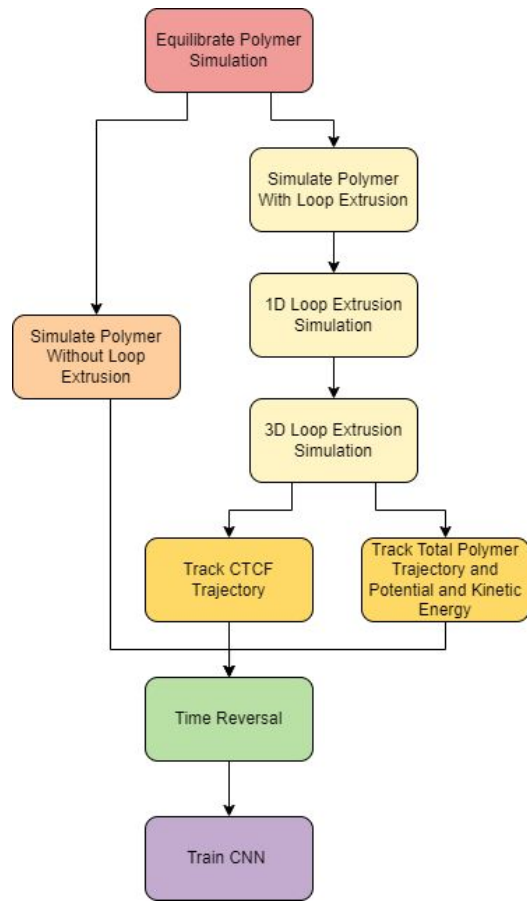


(Zhang, 2022)

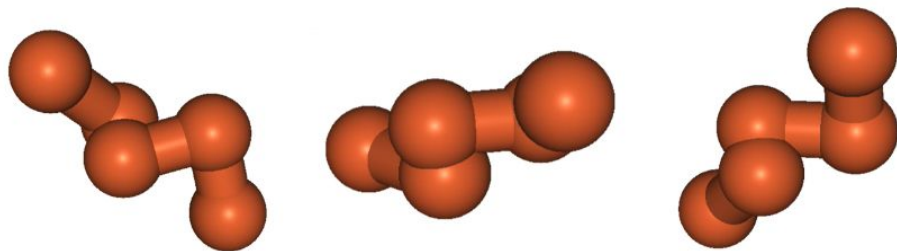
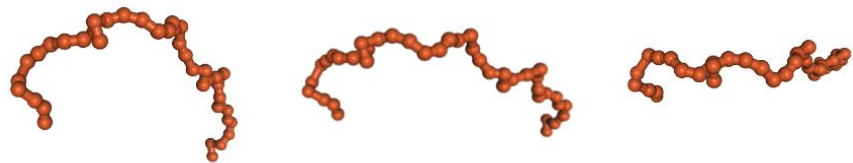
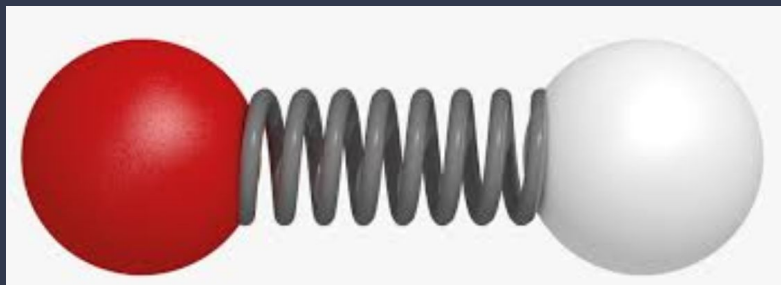
# Goal

Detect loop extrusion in living systems by comparing “movies” of DNA with loop extrusion that are played forward and backward in time.

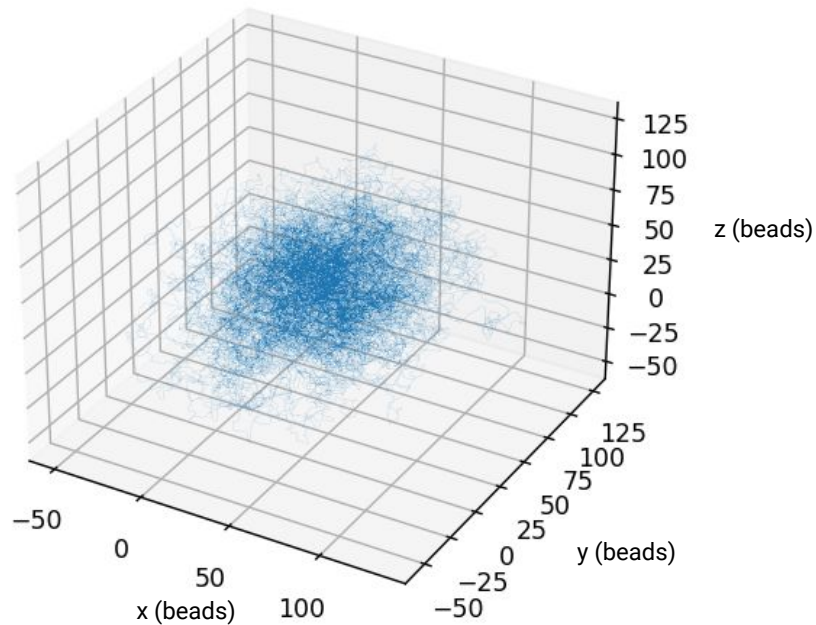
# Workflow



# Polychrom Synthetic Dataset

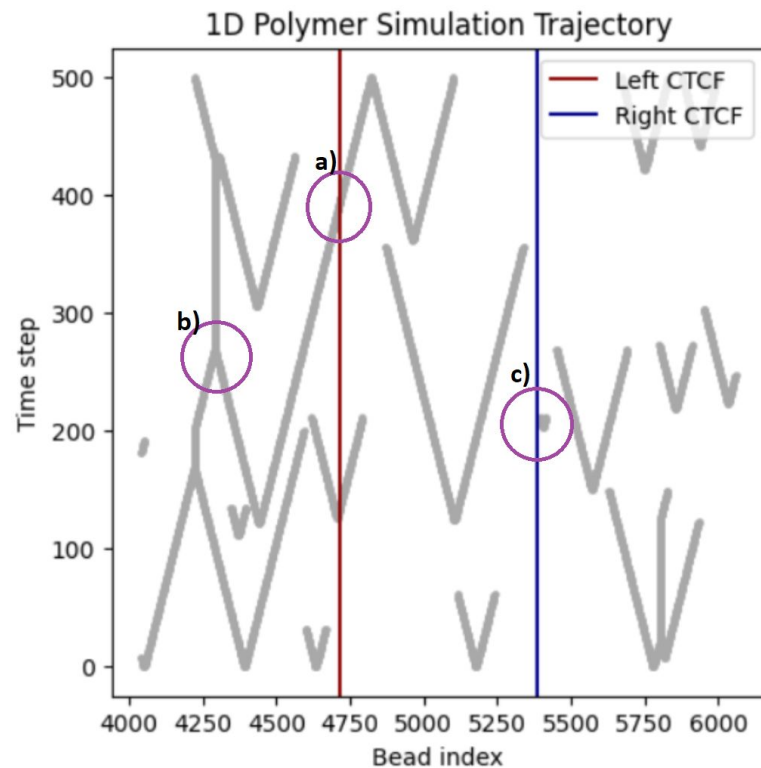


# Equilibrated Polychrom Simulations

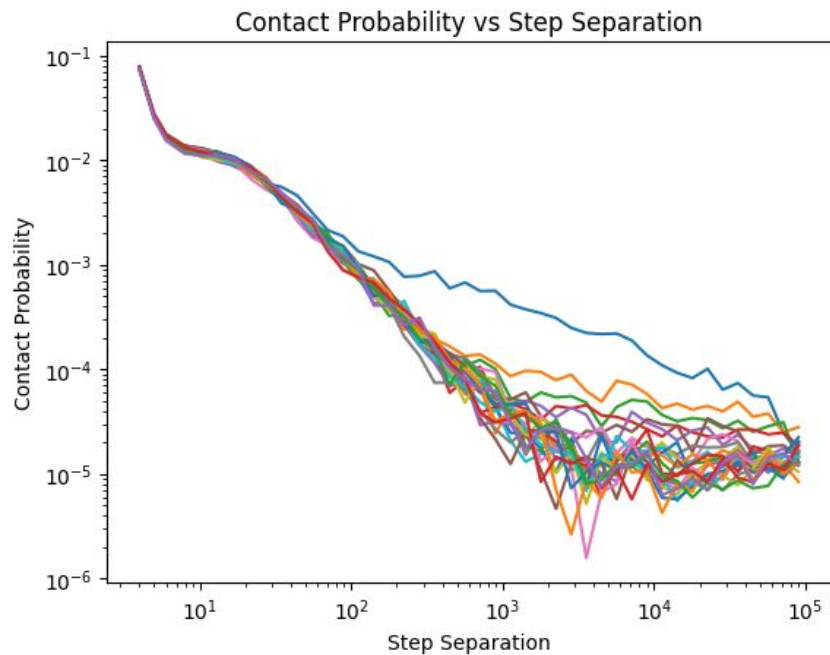


Representation of the Polychrom generated polymer using  
Matplotlib

# 1D Loop Extrusion Simulations

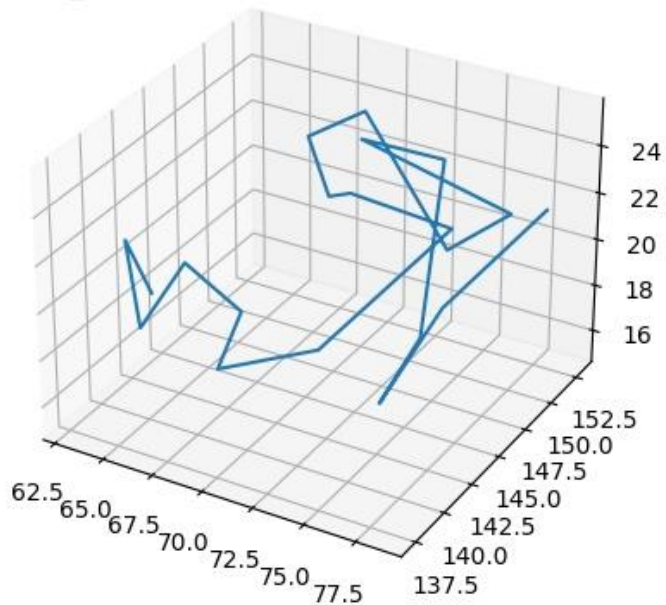


# 3D Loop Extrusion Simulations



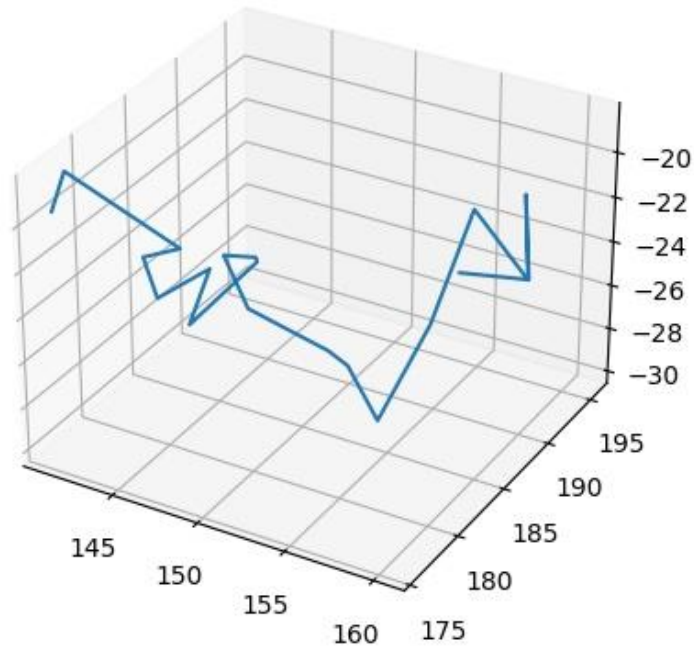
# CTCF Location Tracking

ctcf\_index==2693.0 and iteration ==0.0



framerate=1nTraj=122\_leftCTCF.csv

ctcf\_index==673.0 and iteration ==0.0



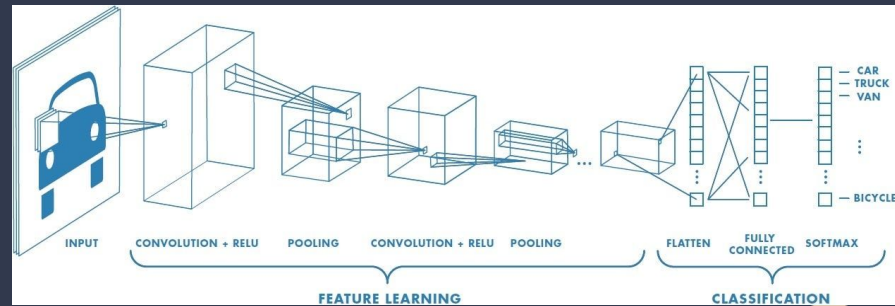


# Time Reversal



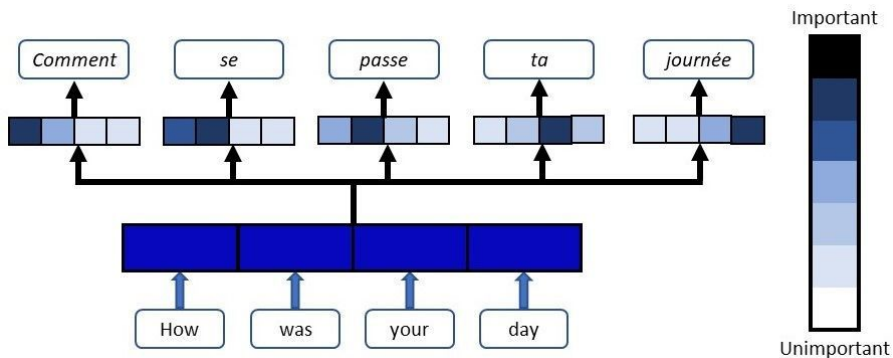
- Create a synthetic dataset for assessing the potential for loop extrusion detection
- DNA motion movies exhibit distinct behaviors when played forward compared to when played backward due to the influence of loop extrusion
- Quantitative method to distinguish between reversed and non-reversed DNA movement.

# CNN



- Time-dependence relates to forward and backward propagation in neural networks.
- Create more efficient pathways to generate numerical training data both time dependent and multi-variant.
- Seek correlations both forward and backward through time.
- Help observe changes in genetic systems that can form a potential mutation event, which could form cancer, diabetes, or other diseases.

# Alternative Strategies



- Transformers can be used as an alternative model to run simulations
  - Utilizes attention to focus on the entire sequence as a whole
  - Requires fewer training data to form a more efficient and accurate model
  - Handle sequences of variable lengths

# Current Applications

- Used as an alternative to live imaging cells on the sub-nuclear level
- Understand the fundamental nature of the structure of sub-nuclear arrangements of genetic material
- Identify targeted therapies for cancers and diseases, based on driver and passenger mutation

# Acknowledgement

- MIT PRIMES organizers and coordinators
- Professor Leonid Mirny, Mr. Henrik Pinholt, and Dr. Aleksandra Galitsyna from the MIT Mirny Lab

# References

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Thank You!