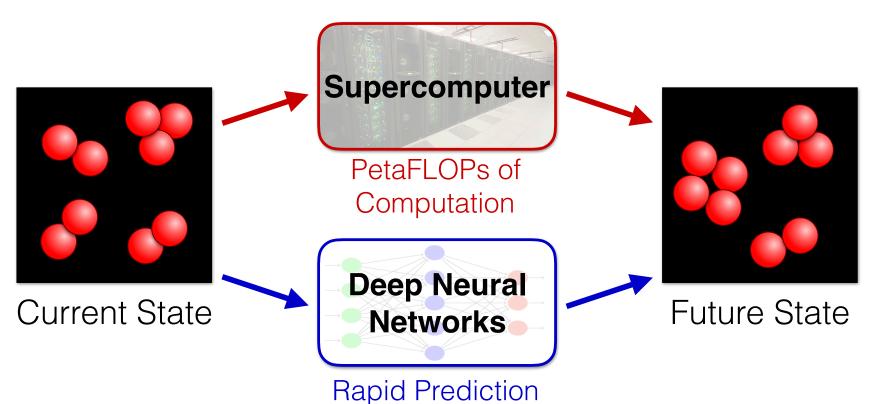
# Stanford Future Prediction in Brownian Dynamics Simulations Using Deep Neural Networks University

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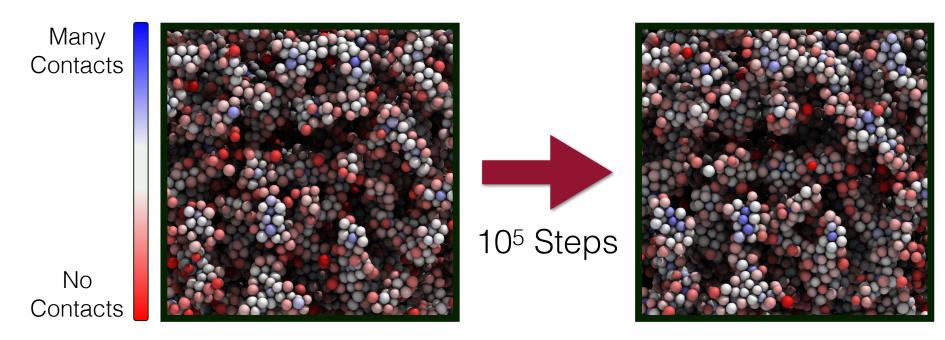
#### **Motivation**

- Scientists utilize molecular dynamics (MD) and Brownian dynamics (BD) to study dynamic or non-equilibrium properties of materials.1
- The principle challenge of MD/BD simulations is the prohibitive computational cost to simulate phenomena that occur at long time scales.
- The goal is to predict the future states of simulations using deep neural networks and bypass explicit computation of every time step:



## **Data & Features**

- 14,400 simulation trajectories list of particle positions (x,y,z) at a given time step — generated using the LAMMPS Molecular Dynamics Simulator.<sup>2</sup>
- Simulation undergoes slow arrested phase separation with different system parameters (volume fraction and interparticle attraction strength).3



Images rendered from simulation snapshots undergoing slow phase separation, separated by 10<sup>5</sup> time steps. Particle colors indicate number of contacts

- Each input and output snapshot contains (x, y, z) positions of 8,788 particles: i.e. 26,364 floating point numbers.
- Particle motion is mildly stochastic and largely deterministic; particles with few contacts are free to diffuse while particles with many contacts move less.

#### Model

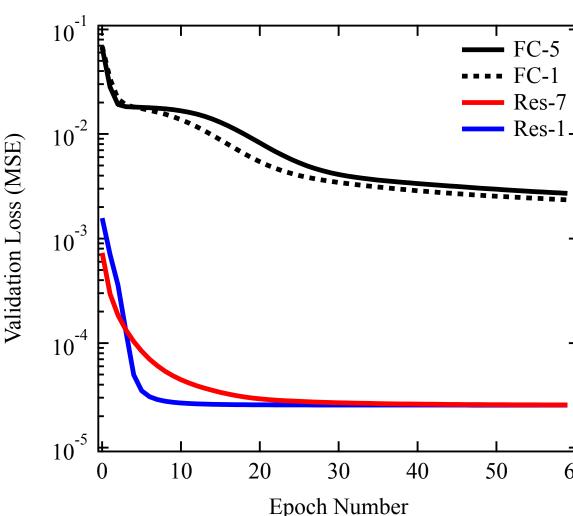
#### Model Architecture Input Trajectory **Output Layer** Flatten and Compute $(8788 \times 3)$ $(8788 \times 3)$ Normalize Displacement Residual **Preprocess Blocks**

- Mean-squared error (MSE) for loss function.
- Use Adam optimizer with L2 regularization.
- Scan over architectures and hyperparameters.

## Residual Block<sup>4</sup> Transfer input positions via residual connection Input positions Output positions Neural network Displacements predicts displacement (26364)(26364)(26364)

## Results

- Our model accurately predicts particle positions with small MSE.
- Residual networks result in significantly smaller loss compared to simple plain networks with only fully connected layers.



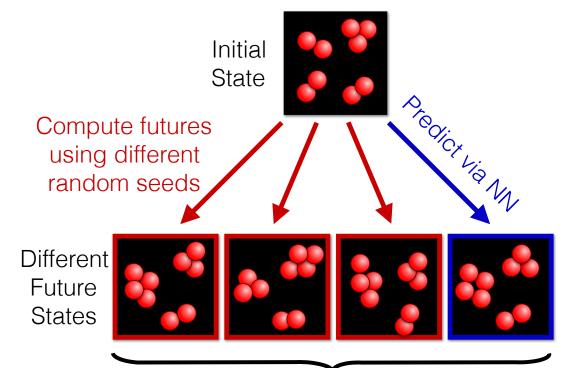
	Mean-Squared Error		
	Training	Validation	Test
FC-1	2.41E-03	2.35E-03	2.34E-03
FC-5	2.81E-03	2.71E-03	2.74E-03
Res-1	2.61E-05	2.55E-05	2.48E-05
Res-3	2.60E-05	2.55E-05	2.48E-05
Res-5	2.60E-05	2.56E-05	2.49E-05
Res-7	2.60E-05	2.55E-05	2.48E-05

Plain (fully connected) Nets: FC-(# of Layers) Residual Nets: Res-(# of Residual Blocks)

## **Discussion and Conclusions**

#### Bayes Error Quantification

- Different random seeds in simulations result in different realizations of future states.
- Mean-squared difference of particle positions are compared across simulated and predicted outcomes.



Simulations	1.90E-05	8.90E-06
Res-7	2.48E-05	1.11E-05
• Residu	ıal netw	orks are

**Mean-Squared Error** 

SD

Compare positions to quantify degree of randomness (signal-to-noise ratio)

- able to predict future positions with randomness comparable to actual simulations
- L2 regularization ( $\lambda = 0.0001$ ) suppresses overfitting.
- Learning rate  $\alpha = 0.00005$ , achieves slow training but small loss.
- Batch size 512 achieves a balance of speed and performance.
- Residual block size with 3 fully-connected hidden layers each with O(1000) neurons provide accurate displacement predictions.

## Conclusions and Future Work

- Successfully developed a deep neural network for predicting future states in Brownian Dynamics simulations.
- Utilizing a residual network structure to focus on displacement significantly improves performance.
- Future work should visualize and interpret what and how each residual block is computing displacements.
- Future work should extend current architecture for variable number of particles.

## **Acknowledgements and References**

- The author would like to thank Ahmad Momeni and the CS 230 teaching staff for helpful discussion and guidance.
- [1] Chen, Jim C., and Albert S. Kim. "Brownian dynamics, molecular dynamics, and Monte Carlo modeling of colloidal systems." Advances in colloid and interface science 112.1-3 (2004):
- [2] Erban, Radek. "From molecular dynamics to Brownian dynamics." Proceedings of the Royal Society A: Mathematical, Physical and Engineering Sciences 470.2167 (2014): 20140036.
- [3] Zia, Roseanna N., et al. "A micro-mechanical study of coarsening and rheology of colloidal gels: Cage building, cage hopping, and Smoluchowski's ratchet." Journal of Rheology 58.5 (2014): 1121-1157.
- [4] He, Kaiming, et al. "Deep residual learning for image recognition." Proceedings of the IEEE conference on computer vision and pattern recognition. 2016.