



Accelerated Microstructure Evolution Predictions Using Machine Learning Based Reduced-Order Models

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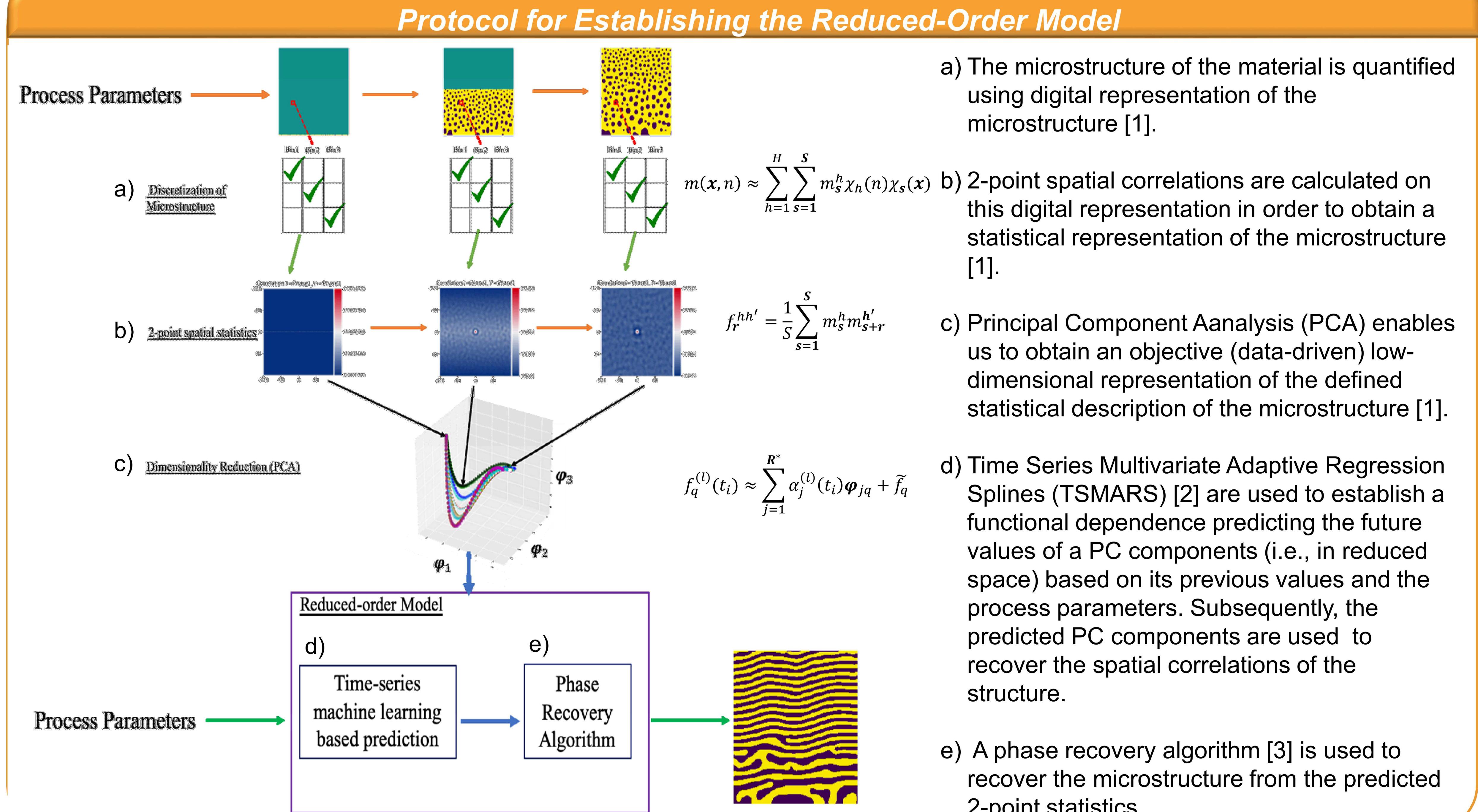
Executive Summary

Problem: The numerical nature of phase-field models causes for these model to be ill-suited for scenarios where a large number of candidate microstructures need to be evaluated to identify an optimal solution (i.e., inverse-type problems and high-throughput scenarios).

Solution: We developed high-fidelity surrogate mesoscale models that accurately predict microstructure evolution without the need to numerically solve the computationally intensive phase-field equations. This is achieved by combining low-dimensional statistically representative microstructure representation with machine learning and time-series analysis techniques.

Results: The protocol developed in this work successfully established a reduced-order model that accurately predicts the microstructure evolution with a significantly reduced computational load: our simulations only **required 5 minutes on a regular laptop** while the explicit phase field model simulations **required 90 minutes on 180 processors**.

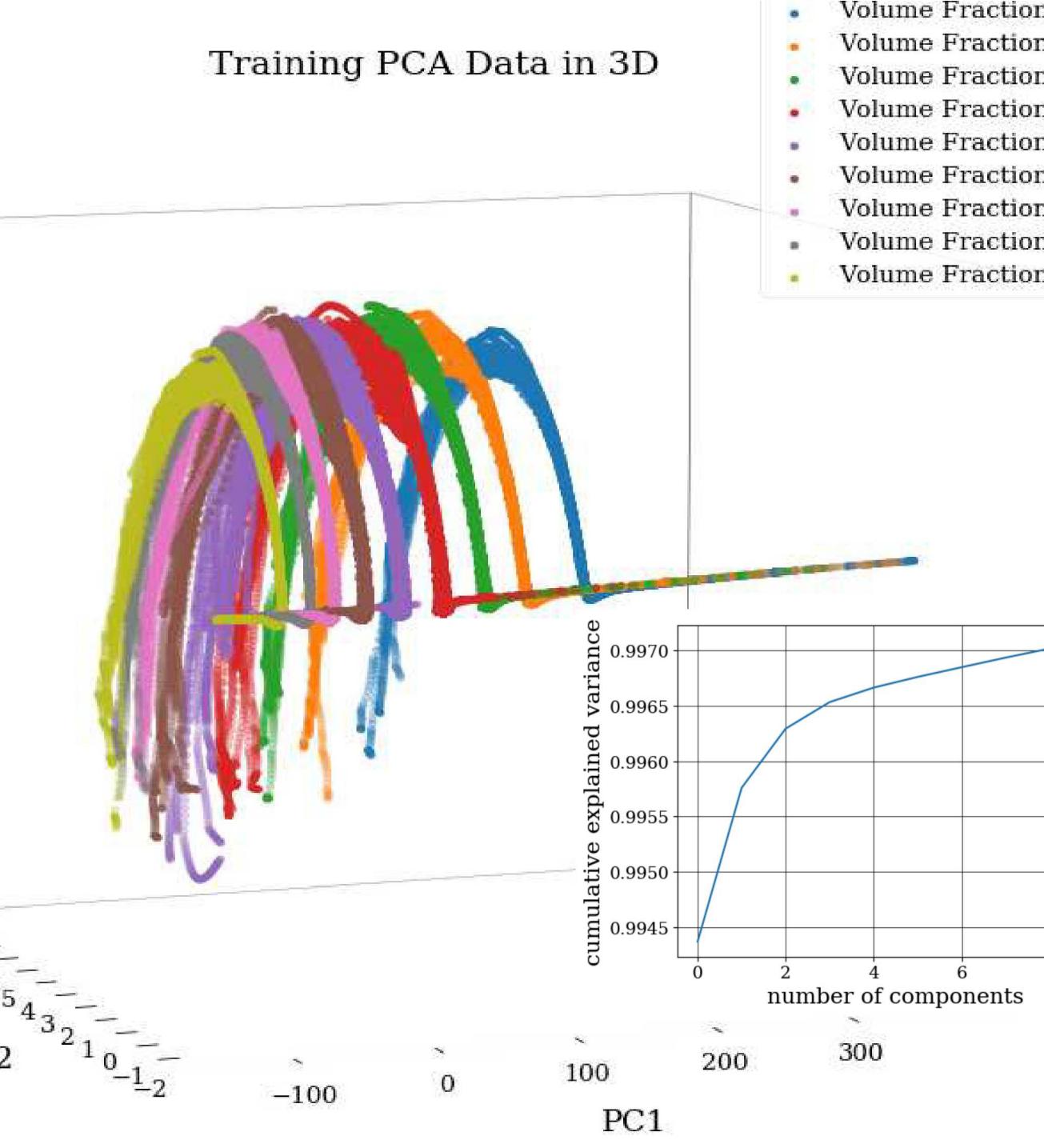
Conclusions: The ability to obtain accurate yet fast phase-field-based microstructure evolution results will enable unbiased automated explorations of vast model spaces to identify the optimal process and material parameters for any desired evolutionary process.



Case Study: Spinodal Decomposition

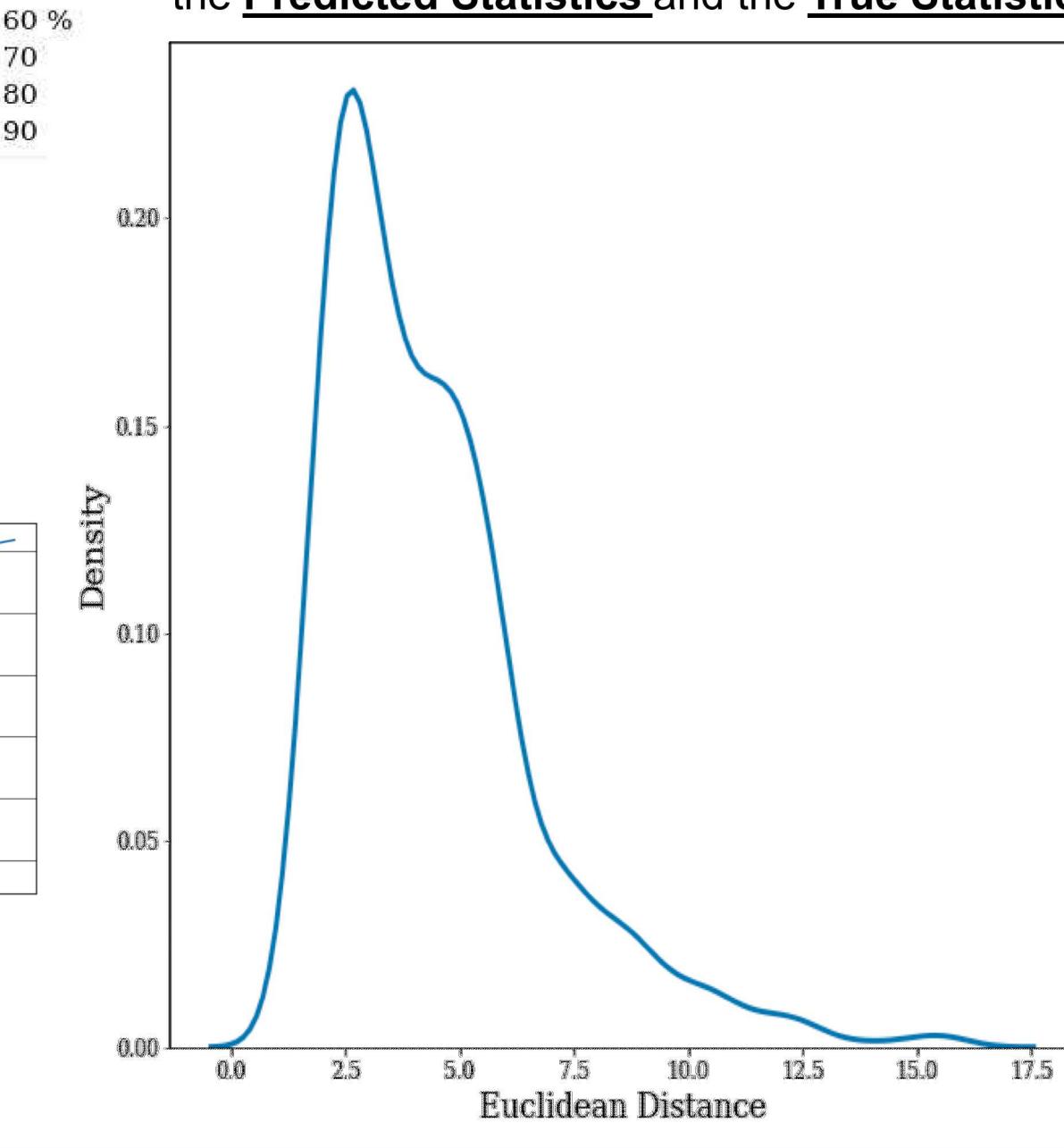
Building Reduced-Order Model for Spinodal Decomposition:

- 9 different volume fractions, sampled every 10%
- 100 Latin hypercube sampled mobilities sampled over 4 orders of magnitudes $\rightarrow [.01, 100]$ for each volume fraction sampled
- Each sampled mobility and volume fraction combination was run for 50,000,000 timesteps and 100 “snapshots” (or timesteps) of the evolution of the microstructure per simulation (101 including the initial microstructure) were captured.
- Each simulation required 90 minutes and 180 processors.**
- Total of 900 simulations.
- Trained the TSMARS surrogate model starting at timestep 50 for 40 timesteps.**



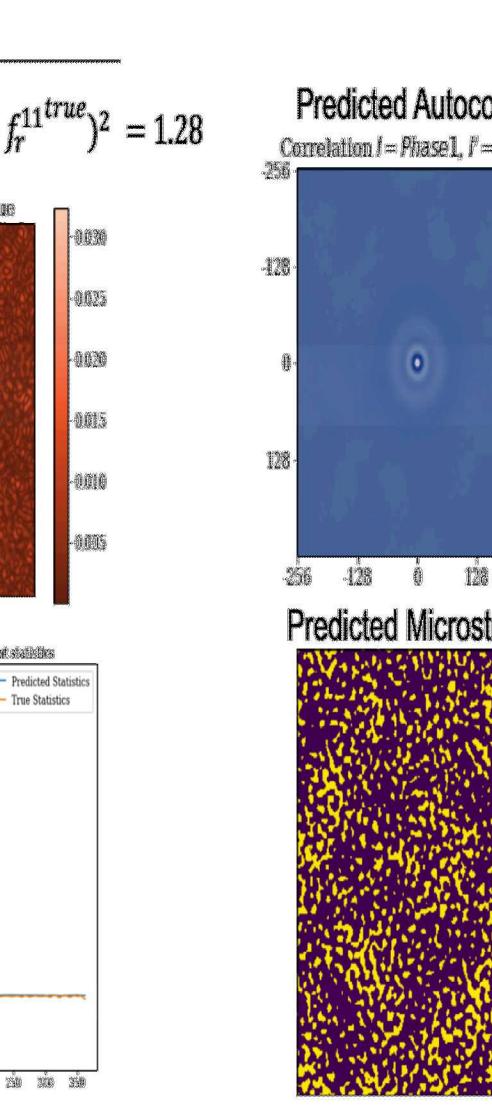
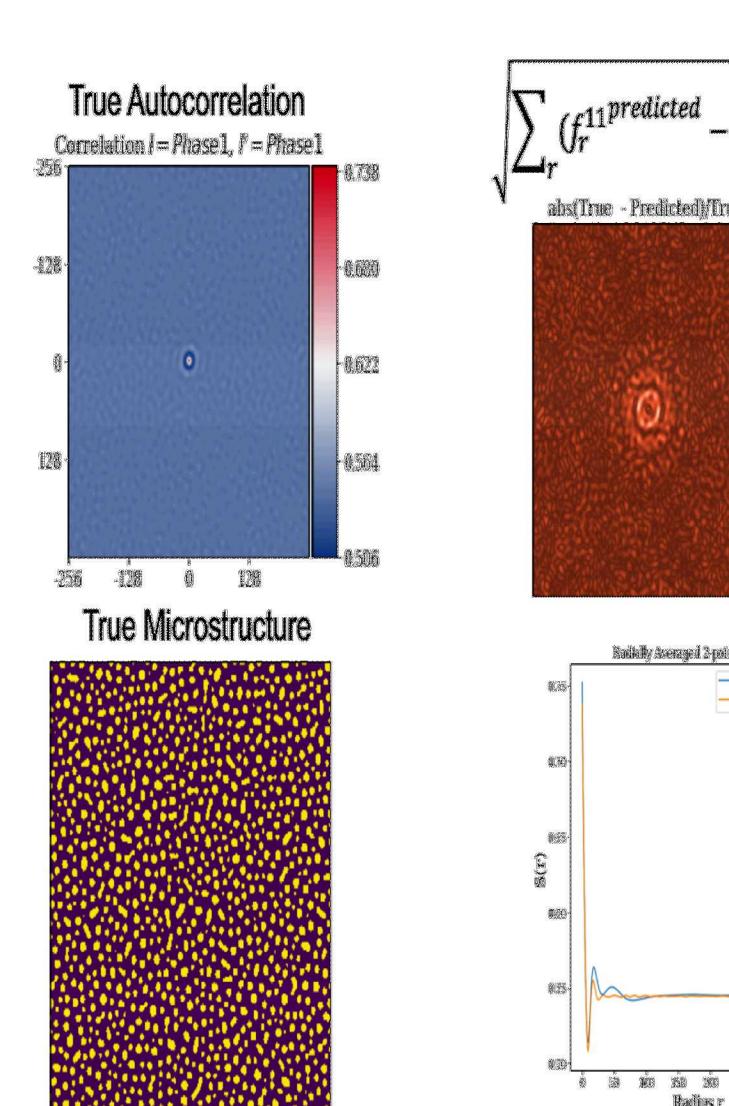
Training Set Results: Predictions at timestep 100

Density Plot of the Euclidean Distance between the Predicted Statistics and the True Statistics



Examples:

a) Initial Values:
 $V_{f_A} = 10\% V_{f_B} = 90\% mob_A = .027 mob_B = .029$



b) Initial Values:
 $V_{f_A} = 40\% V_{f_B} = 60\% mob_A = .021 mob_B = 7.39$

