

The Hong Kong University of Science and Technology

Department of Mathematics

PhD THESIS EXAMINATION

Uncovering the Dynamics of Molecular Systems: Structure, Simulation, and Signal

By

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<u>ABSTRACT</u>

The primary challenge in understanding biological systems lies in deciphering the complex, multiscale relationships between structure and function. From molecular assemblies to whole tissues, biological structures give rise to dynamic behaviors that are nonlinear, high-dimensional, and often emergent. This thesis presents a unified framework for exploring biological dynamics through four interconnected research directions. Initially, we address protein-ligand binding by predicting ligand poses under spatial and physical constraints, generating biologically realistic conformations that capture molecular interaction dynamics. Building on this, we reconstruct long-timescale biomolecular conformational changes from short simulations, bridging the gap between limited sampling and functional timescales. Next, we analyze single-molecule fluorescence (smFRET) data to reveal hidden dynamic states and transitions from noisy time-series, providing insights into transient molecular behaviors. Finally, we develop a structural sparsity framework using inverse scale space to extract key features from high-dimensional biological data, enabling interpretable models. Together, these directions form a cohesive approach to modeling, learning, and interpreting biomolecular dynamics across spatial, temporal, and statistical dimensions.

> Date : 15 May 2025, Thursday Time : 3:00 pm Venue : Room 4472 (Lifts 25/26)

<u>Thesis Examination Committee</u> :		
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(Open to all faculty and students)

The student's thesis is now being displayed on the reception counter in the General Administration Office (Room 3461).