

Statistical Aspects of a Multiscale Computational Scheme for Complex Fluids

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Statistical Investigations in Multiscale Computing

Time-Parallel Continuous-Kinetic-Molecular (tp-CKM):

- ▶ **Offline assessment** of quality of **mesoscale representation** frameworks
- ▶ **Online assessment** of consistency between **microscale** variables and **mesoscale PDF**
- ▶ Estimation of **secular errors** in microscale-mesoscale comparisons

Note that statistical model **validation** and **verification** is done **on-the-fly!**

Elements of tp-CKM (Mitran & Young 2011)

Molecular: ensemble of detailed **trajectories** of **microscale** variables $\{\mathbf{Y}^{(j)}(t)\}_{j=1}^N$; $N \sim 10^4$ realizations

Kinetic: approximate propagation of **mesoscale PDF** $\tilde{\psi}(\mathbf{y}, t)$ of microscale variables

Continuum: evolution of **macroscale** field variables

- ▶ **closure** requires some **local statistic** of **microscale** variables $\tau(t) = \mathbf{F}(\tilde{\psi}(\cdot, t))$ (dependence on macroscale spatial variable suppressed)

Motivation for method is when **statistics** of **microscale** can be much **slower** than their nominal time scale.

- ▶ rupture, cellular “blebbing,” . . .

My particular interest is primarily on the **mesoscale** and its **interaction** with the **microscale**.

Mesoscale Representation of PDF: Challenges

Microscale variable \mathbf{Y} is **high-dimensional** ($\sim 3M - 5$ for polymer with M monomers)

- ▶ require some **simplification** to permit efficient **propagation** and **sampling**
- ▶ but rich enough to capture important **statistical structure** (correlations, ...)
- ▶ **parametric** approaches (fitting Gaussian or Gibbs-Boltzmann distribution) may or may not be appropriate for desired **non-equilibrium** applications

But **microscale data** is relatively **abundant** ($N \sim 10^4$ realizations).

Mesoscale Representation of PDF: One Implementation

Expansion into **radial basis functions**

$$\tilde{\psi}_{\mathbf{Y}}(\mathbf{y}, t) \equiv \sum_{\ell=1}^L c_{\ell}(t) B \left(\|\mathbf{y} - \mathbf{y}^{(\ell)}\| \right)$$

where

- ▶ $\{\mathbf{y}^{(\ell)}\}_{\ell=1}^L$ are points **interpolating** between mean of $\tilde{\psi}(\mathbf{y}, t_n)$ at **beginning** of macro time step and mean of estimated $\tilde{\psi}(\mathbf{y}, t_{n+1})$ at **end** of macro time step ($L \sim 10^2$)
 - ▶ uses **Jordan-Kinderlehrer-Otto (JKO)** gradient flow formalism for Fokker-Planck equation
- ▶ $B(y)$ is a **radial basis function**, taken as **Gaussian**

$$B(y) = \frac{\exp(-y^2/(2h^2))}{\sqrt{2\pi h^2}}$$

with h a fixed width parameter

- ▶ $c_{\ell}(t)$ are **coefficients** computed through JKO and spline interpolation

Mesoscale Representation of PDF: Research Directions

Systematic **offline assessment** of **quality** of **mesoscale** representation **framework**

- ▶ direct **confrontation** with **microscale** simulation **data** via statistical **hypothesis testing**
- ▶ statistical challenges:
 - ▶ **high-dimensional**
 - ▶ **hypothesis** for PDF framework does **not** have **specific parameters**
- ▶ explore and develop **multivariate nonparametric hypothesis tests**
- ▶ formulate test based on **metric involving statistic** $\tau(t) = \mathbf{F}(\tilde{\psi}(\cdot, t))$ of interest to **macroscale?**
- ▶ ascertain **adequate level** of **statistical resolution** via tests for **statistically significant improvement** of goodness-of-fit of data in richer frameworks

Compare against **alternative mesoscale representation** frameworks

- ▶ **Markov chain** models
- ▶ **Normal modes**

Statistical Comparison: Microscale \rightarrow Mesoscale

From simulated ensemble $\{\mathbf{Y}^{(j)}(t)\}_{j=1}^N$, want to compare its fidelity with predicted mesoscale PDF $\tilde{\psi}(\mathbf{y}, t)$.

Currently, one way this is done is by projecting the microscale data onto the vector space **spanned by the same radial basis functions** as used to represent the mesoscale, to obtain **PDF estimated by microscale data**:

$$\hat{\psi}_{\mathbf{Y}}(\mathbf{y}, t) \equiv \sum_{\ell=1}^L \hat{c}_{\ell}(t) B\left(\|\mathbf{y} - \mathbf{y}^{(\ell)}\|\right)$$

- ▶ only $\{\hat{c}_{\ell}\}_{\ell=1}^L(t)$ depend on the **data**

Comparison of **microscale** and **mesoscale** data is done by appropriate vector space norm $\|\mathbf{c} - \hat{\mathbf{c}}\|$ with

$$\mathbf{c} = \{c_{\ell}\}_{\ell=1}^L, \quad \hat{\mathbf{c}} = \{\hat{c}_{\ell}\}_{\ell=1}^L.$$

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More recent developments (**Mitran**):

- ▶ **Bayesian hypothesis testing** for **Gaussian** parametric model,
- ▶ **lattice-Boltzmann** type representation

Research direction:

- ▶ Develop new **online parametric hypothesis** tests for how well **microscale data** is actually **consistent** with **proposed mesoscale representation** when **not Gaussian** or comparably simple

Systematic Biases in Microscale-Mesoscale Comparisons

The microscale-mesoscale comparisons may not detect a **small systematic error** which **accumulates**.

Ideas:

- ▶ Statistical test for **correlations** in errors
- ▶ Estimate accumulated error through “**vectorial**” **addition** of errors at each time step
- ▶ Use mesoscale theory to **estimate amplification** and accumulation of **error**

Work in Progress

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General goals:

- ▶ **Stimulate** new directions of **statistical** research **motivated** by **multiscale computation**
- ▶ **Incorporate** more **sophisticated statistical** ideas into multiscale computation

References on tp-CKM

S. Mitran, J. Young, “Multiscale Computation of Cytoskeletal Mechanics During Blebbing,” pp. 345-372, in *Cellular and Biomolecular Mechanics and Mechanobiology*, Springer, A. Gefen (ed.), 2011.

- ▶ See in particular **Sec. 3 for description of method**
- ▶ download from
<http://mitran.web.unc.edu/publications/>

S. Mitran, “Time parallel kinetic-molecular interaction algorithm for CPU/GPU computers,” *Proc. Comp.Sci.* **1**:745-752, 2010.

- ▶ Elements of combining **parareal** and **multiscale** computing

Sorin Mitran will be visiting this KITP program June 11-29.