

# *Approches mathématiques pour la simulation multi-échelle des matériaux*

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based on a series of works by

F. Legoll, T. Lelièvre, G. Stoltz and collaborators



European Research Council  
Established by the European Commission



# Numerical simulation of coarse-grained stochastic dynamics

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*Work also supported by ANR Funding ANR-14-CE23-0012 (“COSMOS”)*

# Dissipative Particle Dynamics and the like

- Simulation of shock waves at the atomistic level requires **very large** systems
- Coarse-graining through stochastic dynamics which is Galilean invariant  
→ friction using **relative velocities** (consistence with **hydrodynamics**)
- **Dissipative Particle Dynamics with conserved energy (DPDE)**
  - can be used in **nonequilibrium** situations
  - replace a molecule or some group of atoms by a **mesoparticle**
  - consistent thermodynamics
  - input: static properties (ab-initio), dynamical parameters
- **Collaboration** with J.-B. Maillet (CEA/DAM) and J. Brennan (Army Research Lab); 2 PhD students (A.-A. Homman and G. Faure)

# Dissipative particle dynamics with conserved energy

- **Coarse-graining** interpretation:

- a (fragment of a) molecule is replaced by a mesoparticle
- $(q_i, p_i)$  describes the center of mass of the  $i$ th mesoparticle
- missing degrees of freedom described by an **internal energy**  $\varepsilon_i$

- Evolution at constant total energy  $\mathcal{H}(q, p, \varepsilon) = V(q) + \sum_{i=1}^N \frac{p_i^2}{2m_i} + \sum_{i=1}^N \varepsilon_i$

- **Microscopic state law**: entropies  $s_i = s_i(\varepsilon_i)$ , internal temperature defined from the entropy as

$$T_i(\varepsilon_i) = \frac{1}{s'_i(\varepsilon_i)}$$

- Simplest case: harmonic internal degrees of freedom,  $T(\varepsilon) = \varepsilon/C_v$

J. Bonet Avalos and A. Mackie, *Europhys. Lett.* **40**, 141-146 (1997)

P. Español, *Europhys. Lett.* **40** 631-636 (1997)

# Equations of motion

$$\begin{cases} dq_i = \frac{p_i}{m_i} dt \\ dp_i = -\nabla_{q_i} V(q) dt + \sum_{i \neq j} -\gamma_{ij} \chi^2(r_{ij}) v_{ij} dt + \sigma_{ij} \chi(r_{ij}) dW_{ij}, \\ d\varepsilon_i = \frac{1}{2} \sum_{j \neq i} \chi^2(r_{ij}) \left( \gamma_{ij} v_{ij}^2 - \frac{\sigma_{ij}^2}{2} \left( \frac{1}{m_i} + \frac{1}{m_j} \right) \right) dt - \sigma_{ij} \chi(r_{ij}) v_{ij} \cdot dW_{ij} \end{cases}$$

where  $W_{ij} = -W_{ji}$ ,  $\chi$  is a cut-off function and  $v_{ij} = \frac{p_i}{m_i} - \frac{p_j}{m_j}$

## Invariant measures

$$\rho(dq dp d\varepsilon) = f(\mathcal{H}(q, p, \varepsilon)) g\left(\sum_{i=1}^N p_i\right) \exp\left(\sum_{i=1}^N s_i(\varepsilon_i)\right) dq dp d\varepsilon,$$

- Fluctuation-dissipation relation

$$\sigma_{ij} = \sigma, \quad \gamma_{ij} = \frac{\sigma^2 \beta_{ij}(\varepsilon_i, \varepsilon_j)}{2}, \quad \beta_{ij}(\varepsilon_i, \varepsilon_j) = \frac{1}{2k_B} \left( \frac{1}{T_i(\varepsilon_i)} + \frac{1}{T_j(\varepsilon_j)} \right)$$

# Numerical integration of DPDE: our key contribution

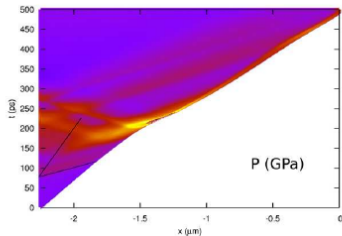
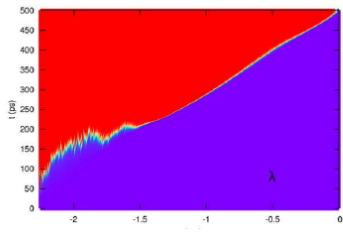
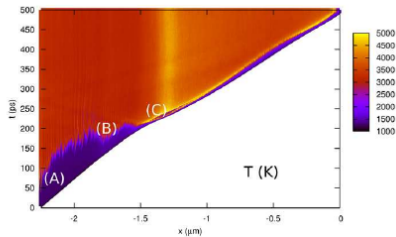
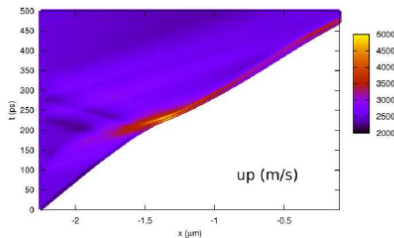
- “Naive” schemes lead to internal energies  $\varepsilon_i < 0$ : **simulation stopped!**
  - this happens more often for small heat capacities
  - this will necessarily happen at some point for large systems
- **Stable and accurate integration schemes?**
  - **Splitting strategy**: Hamiltonian part vs. elementary stochastic dynamics
  - elementary stochastic dynamics **reduce to a dynamics on  $v_{ij}$  only**
  - superimpose a **Metropolis correction** for discretizations of these reduced dynamics<sup>1</sup>, even in the nonequilibrium setting considered
- **Pro/cons of this integrator:**
  - automatically corrects for **negative internal energies** (stabilization)
  - parallelization/threadability limited → dedicated schemes for that<sup>2</sup>

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<sup>1</sup>G. Stoltz, *J. Comput. Phys.* (2017)

<sup>2</sup>A.-A. Homman, J.-B. Maillet, J. Roussel and G. Stoltz, *J. Chem. Phys* (2016)

# Detonation waves in nitromethane



Particle velocity, temperature, progress variable, pressure

J.-B. Maillet, G. Vallverdu, N. Desbiens and G. Stoltz, *Europhys. Lett.* (2011)

# Orders of magnitude of current simulations

- **At CEA/DAM**

- number of particles  $N \sim 10^6 - 10^8$
- number of cores: several thousands, with vectorized/threadable code ExaSTAMP
- number of steps  $10^6$ , timestep  $\Delta t \sim 1 - 5 \times 10^{-15}$
- CPU time: a few  $\mu\text{s}/N/\text{ts}$  for simple LJ potential,  $\times 10 - 100$  for more complicated one

- **At Army Research Lab (Aberdeen Proving Ground)**

- number of particles up to  $N = 1,126,926,339$
- machines: Thunder (USAF), Stampede2 (Texas Advanced Computing Center), Trinity/KNL (Los Alamos)
- 3,000 to 8,900 nodes (Intel Xeon Phi 7250 KNL or E5-2699v3); between 4.3 and 27 PFLOPS/s (in double precision)
- simulation time 0.5 ns

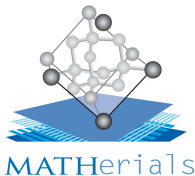


# Adaptive Multilevel Splitting algorithms for rare event simulations

Tony Lelièvre

Ecole des Ponts ParisTech and INRIA

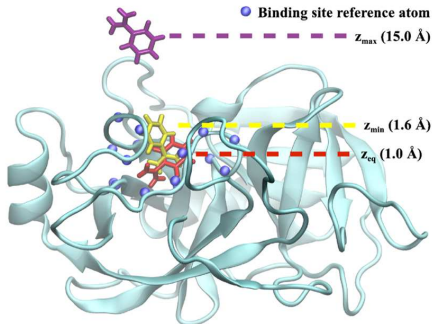
Joint work with C.-E. Bréhier, F. Cérou, M. Gazeau, L. Goudenège,  
A. Guyader, C. Mayne, M. Rousset and I. Teo



# Motivation 1: Simulations of biological systems

## Unbinding of a ligand from a protein

### Trypsin with various conformational states of benzamidine



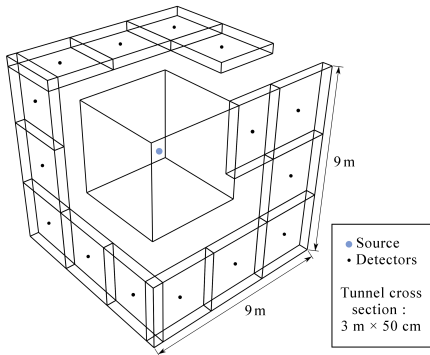
Elementary time-step for the molecular dynamics =  $10^{-15}$  s

Dissociation time  $\simeq 0.02$  s

**Challenge:** bridge the gap between timescales

## Motivation 2: Radiation protection

### Monte Carlo particle transport



### Concrete tunnel with a neutron source

How to compute the neutron flux at the detector ?

**Challenge:** the flux is very small

## Mathematical setting: rare event computation

Consider a stochastic process  $(X_t)_{t \geq 0}$  and two stopping times  $\tau_A$  and  $\tau_B$ . Objective: simulate and compute the probability of the event  $\{\tau_B < \tau_A\}$  when  $\mathbb{P}(\tau_B < \tau_A)$  is very small ( $10^{-8}$  to  $10^{-18}$ ).

Basic idea of **splitting technique**: find intermediate events which are easier to simulate:

$$\{\tau_{z_1} < \tau_A\} \supset \{\tau_{z_2} < \tau_A\} \supset \dots \supset \{\tau_{z_{\max}} < \tau_A\} \supset \{\tau_B < \tau_A\}$$

and simulate the successive *conditional events*: for  $k = 1, 2, \dots$ ,

$$\{\tau_{z_q} < \tau_A\} \text{ knowing that } \{\tau_{z_{q-1}} < \tau_A\}$$

where  $\tau_z = \inf\{t, \xi(X_t) > z\}$  for a well chosen real valued **importance function**  $\xi$ .

**Adaptive feature**: build the intermediate levels  $(z_i)_{i \geq 1}$  on the fly.

→ **Adaptive Multilevel Splitting** algorithm [C erou, Guyader, Stoch. Annal. Appl., (2007)]

## Numerical results

Example 1: In collaboration with the group of K. Schulten (C. Mayne and I. Teo), AMS is currently implemented in the NAMD code. We have studied the unbinding event of benzamidine from trypsin.

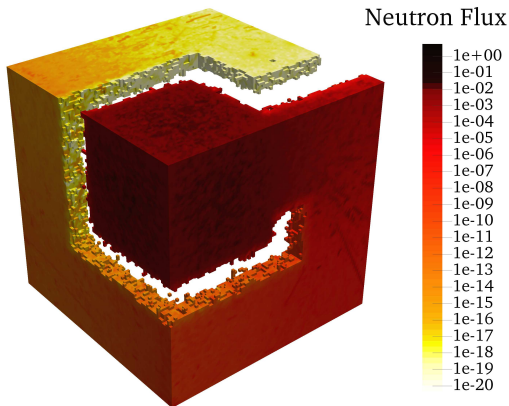
**Estimated dissociation rate:**  $k_{off} = (260 \pm 240)s^{-1}$  which is in the same order of magnitude as the experimental rate  $(600 \pm 300)s^{-1}$ .

**Overall simulation time:**  $2.3\mu s$  which is 4 orders of magnitude shorter than than the estimated dissociation time.

MD setup: about 70 000 atoms, CHARMM36 force field, NPT conditions (298 K).

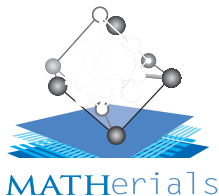
## Numerical results

Example 2: In collaboration with CEA (Eric Dumonteil, Cheikh Diop and Henri Louvin), AMS is currently implemented in the Tripoli code.



## References

- C.-E. Bréhier, M. Gazeau, L. Goudenège , T. Lelièvre and M. Rousset, *Unbiasedness of some generalized Adaptive Multilevel Splitting algorithms*, Annals of Applied Probability 26(6), 2016.
- F. Cérou, A. Guyader, T. Lelièvre and D. Pommier, *A multiple replica approach to simulate reactive trajectories*, Journal of Chemical Physics 134, 2011.
- I. Teo, C. Mayne, K. Schulten and T. Lelièvre, *Adaptive multilevel splitting method for molecular dynamics calculation of benzamidine-trypsin dissociation time*, Journal of Chemical Theory and Computation 12(6), 2016.



# Multiscale computations based on MsFEM: model reduction and goal-oriented a posteriori error estimation

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*Joint works with Ludovic Chamoin (LMT Cachan)*



# Problem of interest (SPE 10,

<http://www.spe.org/web/csp/index.html>)

$$-\operatorname{div}[A_\varepsilon(\mu, x)\nabla u^\varepsilon(\mu, x)] = f(x) \quad \text{in } \Omega, \quad u^\varepsilon = 0 \quad \text{on } \partial\Omega$$

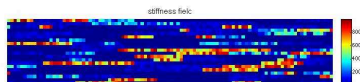
where  $\mu$  is a parameter. We take

$$A_\varepsilon(\mu, x) = \lambda(\mu, x)A_\varepsilon(x), \quad \lambda(\mu, x) = \mu + (1 - \mu)\lambda_c(x)$$

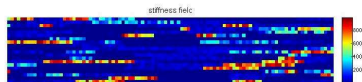
Depending on the value of  $\mu$ , the central channel is present or not.  
Very large contrast in  $A_\varepsilon$ :  $10^6$



Representation of  $\lambda_c(x)$



$A_\varepsilon(\mu, x)$  for  $\mu = 1$



$A_\varepsilon(\mu, x)$  for  $\mu = 0.1$

# Model reduction approaches

$$-\operatorname{div}[A_\varepsilon(\mu, x)\nabla u^\varepsilon(\mu, x)] = f(x) \quad \text{in } \Omega, \quad u^\varepsilon = 0 \quad \text{on } \partial\Omega$$

- **Direct MsFEM approach:** for each new  $\mu$ ,
  - compute the MsFEM basis functions:

$$(\star) \quad -\operatorname{div}[A_\varepsilon(\mu, x)\nabla \phi_i^\varepsilon(\mu, x)] = 0 \quad \text{in } K, \quad \phi_i^\varepsilon(\mu, \cdot) = \phi_i^0 \quad \text{on } \partial K$$

- solve the global problem on  $\operatorname{Span}\{\phi_i^\varepsilon(\mu, \cdot), \quad 1 \leq i \leq I\}$ .

Too expensive!

- **Our approach:** model reduction (**PGD approach**) on  $(\star)$ :

$$\phi_i^\varepsilon(\mu, x) \approx \phi_i^0(x) + \sum_{j=1}^J \psi_j^\varepsilon(x) \alpha_j(\mu)$$

for (hopefully) a small number  $J$  of terms. The decomposition is built **iteratively** (greedy algorithm).

# Proper Generalized Decomposition (Ladevèze, Chinesta, Nouy, ...)

Idea to compute  $w(x, \mu)$ :

- represent the solution as a linear combination of tensor products of small-dimensional functions:

$$w(x, \mu) = \sum_{j \geq 1} \psi_j(x) \alpha_j(\mu)$$

- look iteratively for the best tensor product: once some approximation

$$w_{n-1}(x, \mu) = \sum_{j=1}^{n-1} \psi_j(x) \alpha_j(\mu)$$

has been computed, improve it by considering

$$w_n(x, \mu) = w_{n-1}(x, \mu) + \psi_n(x) \alpha_n(\mu)$$

# PGD within MsFEM

- For a given parameter  $\mu_0$ , perform a MsFEM computation and adapt the discretization parameters ( $H$ ,  $h$  and oversampling).  
This discretization will be kept unchanged.

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- Perform a PGD approach on  $\phi_i^\varepsilon(\mu, \cdot)$ , solution to

$$-\operatorname{div}[A_\varepsilon(\mu, x)\nabla\phi_i^\varepsilon(\mu, x)] = 0 \text{ in } K, \quad \phi_i^\varepsilon(\mu, \cdot) = \phi_i^0 \text{ on } \partial K$$

It amounts to writing

$$(\star) \quad \phi_i^\varepsilon(\mu, x) \approx \phi_i^{\varepsilon, J}(\mu, x) = \phi_i^0(x) + \sum_{j=1}^J \psi_j^\varepsilon(x)\alpha_j(\mu)$$

# PGD within MsFEM

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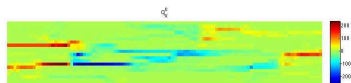
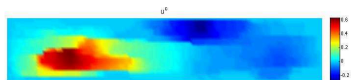
$$(\star) \quad \phi_i^\varepsilon(\mu, x) \approx \phi_i^{\varepsilon, J}(\mu, x) = \phi_i^0(x) + \sum_{j=1}^J \psi_j^\varepsilon(x)\alpha_j(\mu)$$

- For each new  $\mu$ :
  - evaluate the basis functions  $\phi_i^{\varepsilon, J}(\mu, x)$  using  $(\star)$
  - solve the global problem on  $\operatorname{Span}\left\{\phi_i^{\varepsilon, J}(\mu, \cdot), 1 \leq i \leq I\right\}$ .
  - estimate the error

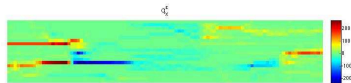
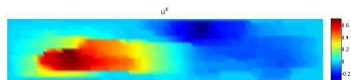
Alternative strategy: PGD on global problem followed by MsFEM discretization

# Numerical results (crude discretization)

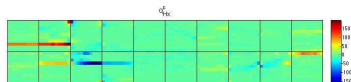
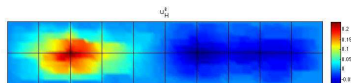
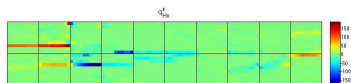
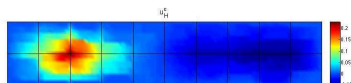
$\mu = 1$  (initial permeability field)



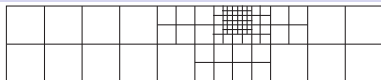
$\mu = 0.1$  (central channel removed)



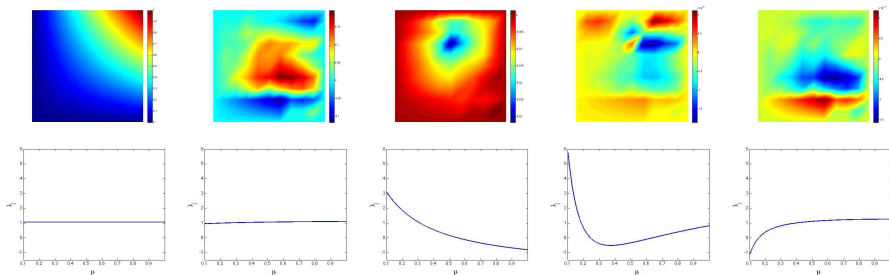
MsFEM solution (no oversampling):



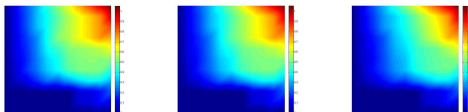
# PGD approach for the computation of $\phi_i^\varepsilon(\mu, \cdot)$



PGD modes  $\psi_j^\varepsilon(x)$  (top) and  $\alpha_j(\mu)$  (bottom),  $j = 1, \dots, 5$ :

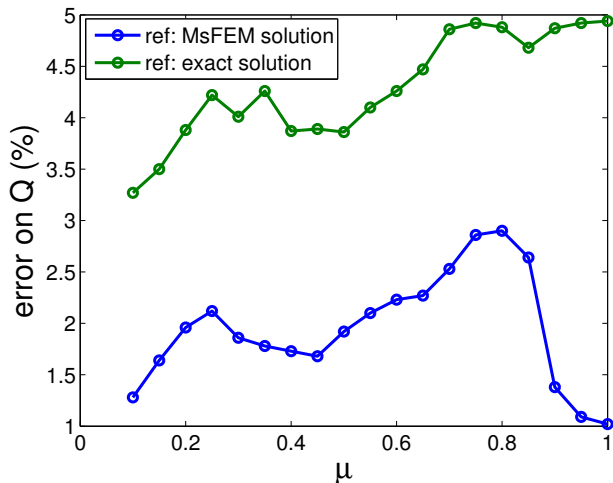


MsFEM basis functions  $\phi_i^0(x) + \sum_{j=1}^J \psi_j^\varepsilon(x) \alpha_j(\mu)$  ( $\mu = 1, 0.5$  and  $0.1$ ):



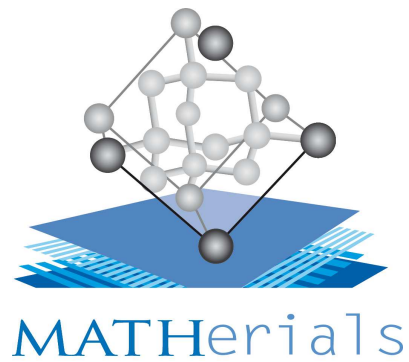


# Error estimation (identical MsFEM discretization for any $\mu$ )



The error remains under 5% for all  $\mu$ .

<https://www.rocq.inria.fr/mathematics/>



Support from ONR and EOARD is gratefully acknowledged.