



# Time-dependent Non-Equilibrium Molecular Dynamics

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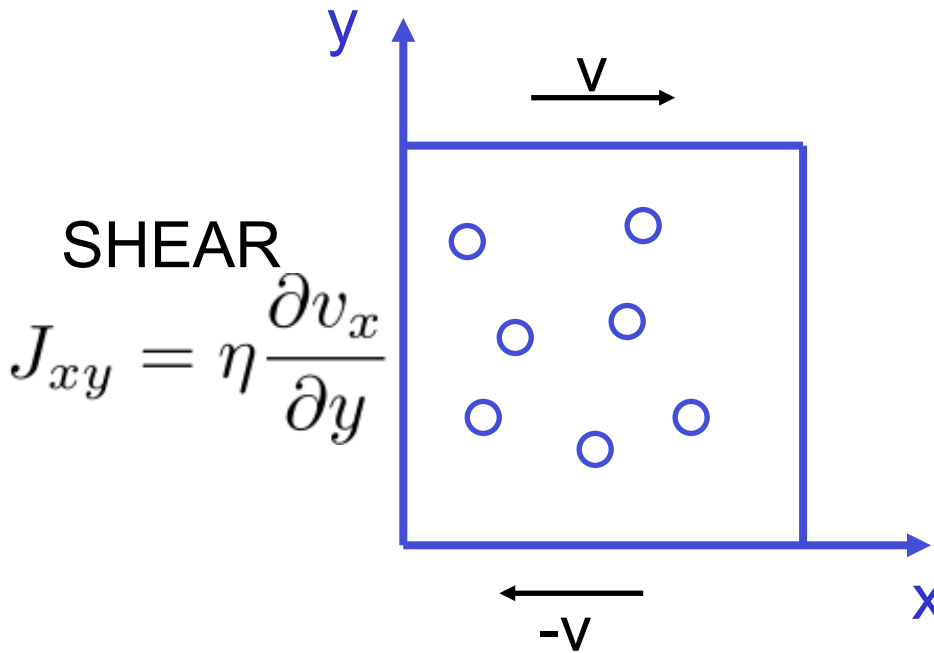
With **S. Meloni (CINECA)**, **S. Orlandini (CINECA)**, **M. Mareschal (ZCAM)**, **C. Pierleoni (Sapienza)**, **M. L. Mugnai (Austin)**

- **t-dependent averages in NEMD (Onsager-Kubo)**
- **sampling an initial condition ensemble by MD**
- **beyond macroscopic Hydrodynamics: Convective cells and relaxation of an interface**

# What we mean by Non-Equilibrium

TRANSPORT

HYDRODYNAMICS

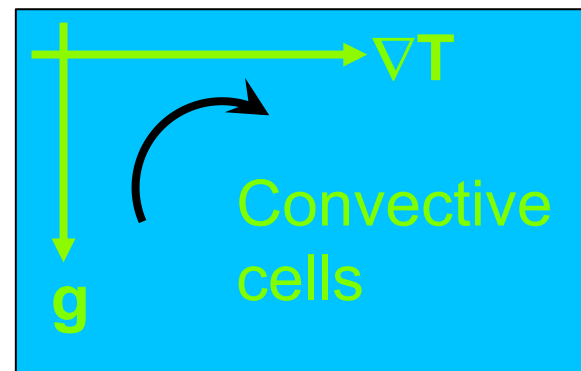
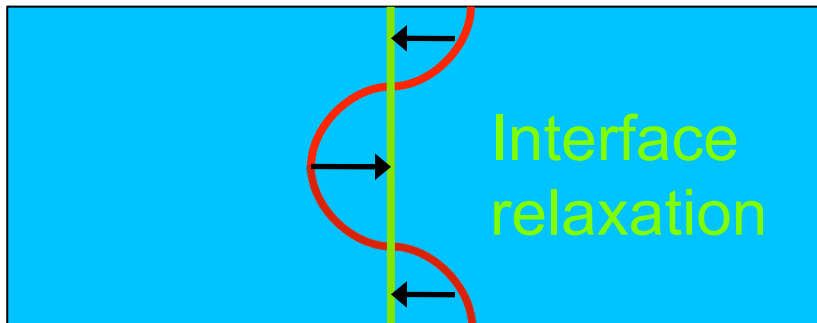


$$\rho(\vec{x}, t) = \langle \hat{\rho}(\vec{x}) \rangle_{ne}$$

$$\hat{\rho}(\vec{x}) = \sum_i \mu_i \delta(\vec{x} - r_i)$$

$$\vec{v}(\vec{x}, t) = \frac{\langle \vec{\hat{p}}(\vec{x}) \rangle_{ne}}{\rho(\vec{x}, t)}$$

$$\vec{\hat{p}}(\vec{x}) = \sum_i \vec{p}_i \delta(\vec{x} - r_i)$$



# NEMD (1)

- Assume that a given, time-dependent external local field  $\psi(x, t)$  is coupled to our system via a suitable local property

$$A(x|\Gamma) = \sum_{i=1, N} A_i(\Gamma) \delta(r_i - x)$$

- The total Hamiltonian of the system is  $\mathcal{H}(\Gamma, t) = \mathcal{H}_0(\Gamma) + \mathcal{H}_p(\Gamma, t)$

$\mathcal{H}_0$  standard equilibrium Hamiltonian

$$\begin{aligned} \mathcal{H}_P(\Gamma) &= - \int dx A(x|\Gamma) \underline{\psi(x, t)} \\ &= -g \chi(t) \sum_i A_i \phi_i \quad \boxed{\psi(x, t) = g\phi(x)\chi(t)} \end{aligned}$$

## PERTURBED SYSTEM:

- Equations of motion

$$\begin{cases} \dot{r} = \frac{\partial H_0}{\partial p} + \frac{\partial H_p}{\partial p} = \frac{p}{\mu} - g \frac{\partial h_p}{\partial p} \chi(t) \\ \dot{p} = -\frac{\partial H_0}{\partial r} - \frac{\partial H_p}{\partial r} = F + g \frac{\partial h_p}{\partial r} \chi(t) \end{cases}$$

- Liouville equation

$$\frac{\partial m}{\partial t} = iLm = iL_0m + iL_p m \equiv \{\mathcal{H}_0, m\} + \{\mathcal{H}_p, m\}$$

with  $m(\Gamma, t) = S^\dagger m(\Gamma, 0)$

$$S^\dagger(t) = S_0^\dagger(t) + \int_{-\infty}^t d\tau S^\dagger(t - \tau) iL_p(\tau) S_0^\dagger(\tau)$$

an observable  $J$  of the system evolves with  
 $\hat{J}(t) \equiv \hat{J}(\Gamma(t)) = S(t)\hat{J}(\Gamma(0))$

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with  $m(\Gamma, t) = S^\dagger m(\Gamma, 0)$

$m_0(\Gamma) = m(\Gamma, 0)$  is a given initial distribution which can be sampled by MD if it comes from a stationary state (in particular but not necessarily an equilibrium one)

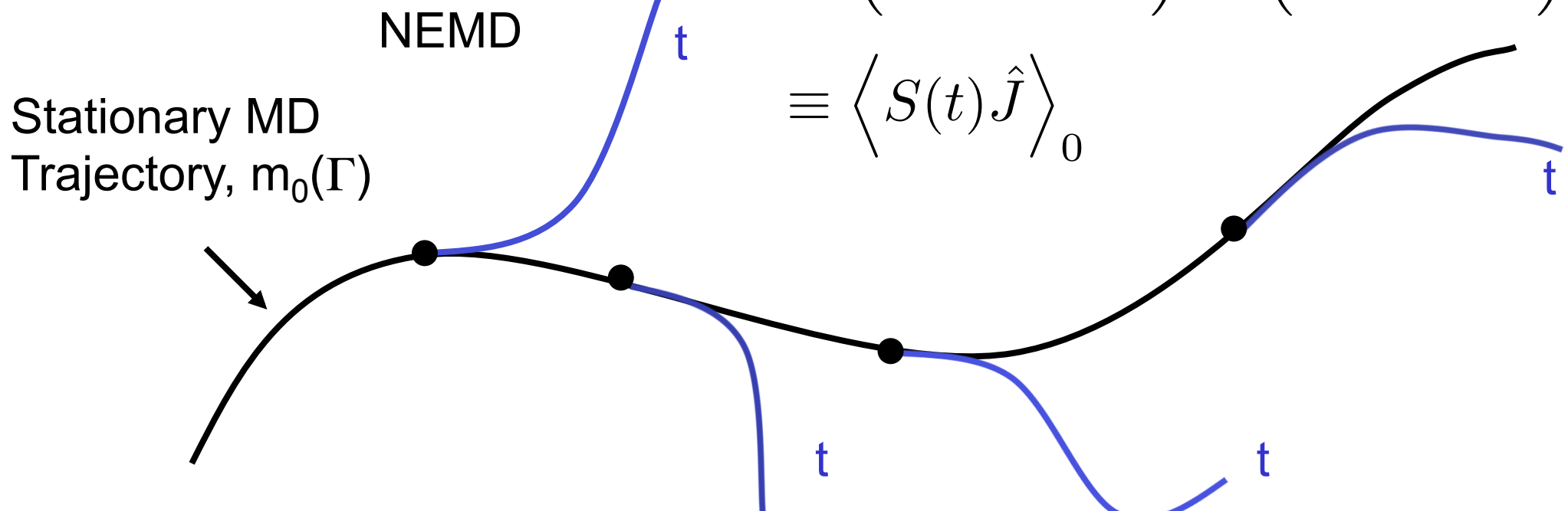
The time-dependent NE average of a given observable

$$\frac{d\hat{J}}{dt} = -iL\hat{J} \quad , \quad \hat{J}(t) = S(t)\hat{J}$$

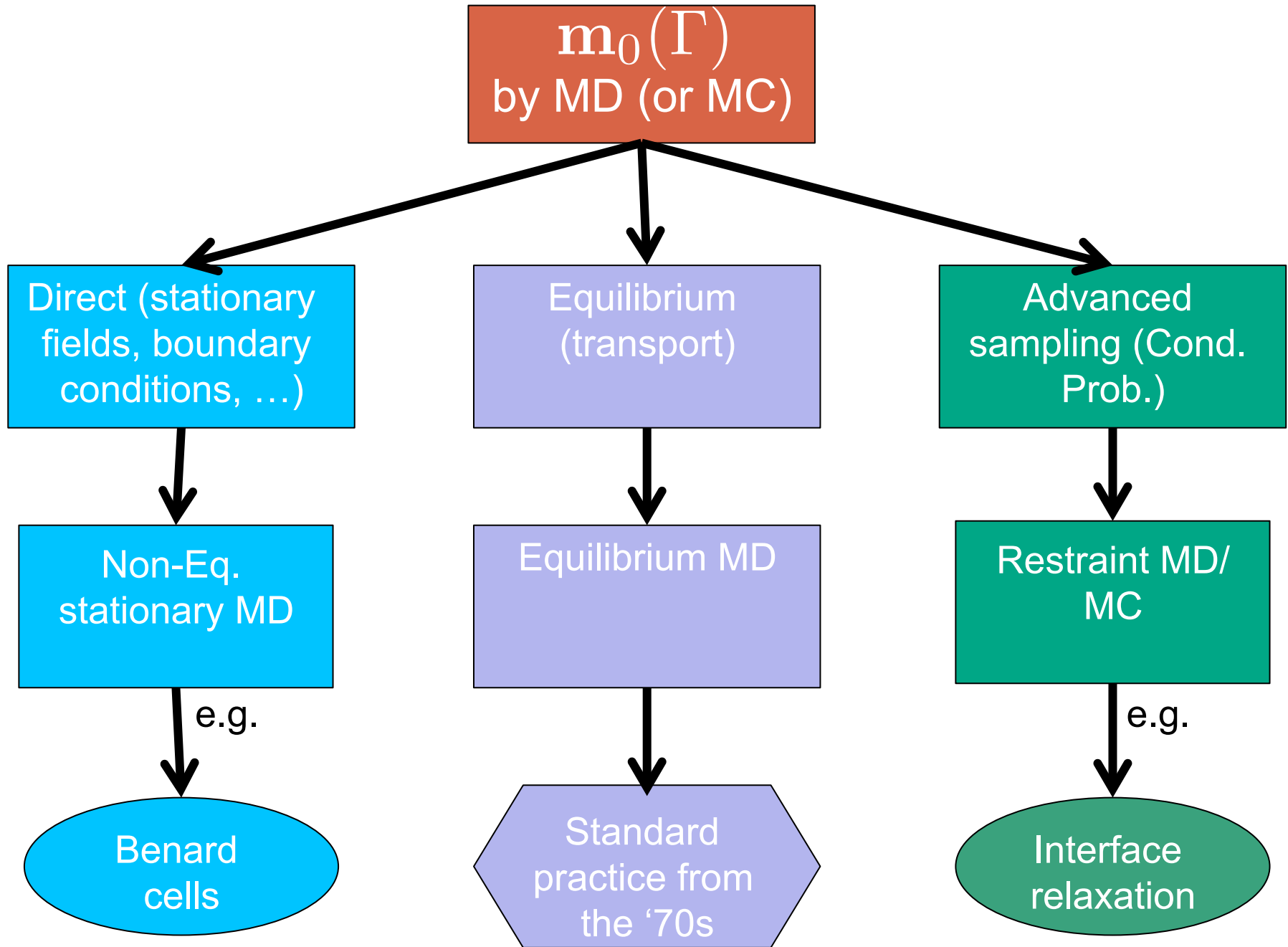
can be obtained as follows

(Onsager-Kubo equation)

$$\begin{aligned} J(t) &= \langle \hat{J} \rangle_{NE}^t = \int d\Gamma \hat{J}(\Gamma) m(\Gamma, t) \\ &\equiv \left( \hat{J}, S^\dagger(t) m_0 \right) = \left( S(t) \hat{J}, m_0 \right) \\ &\equiv \langle S(t) \hat{J} \rangle_0 \end{aligned}$$

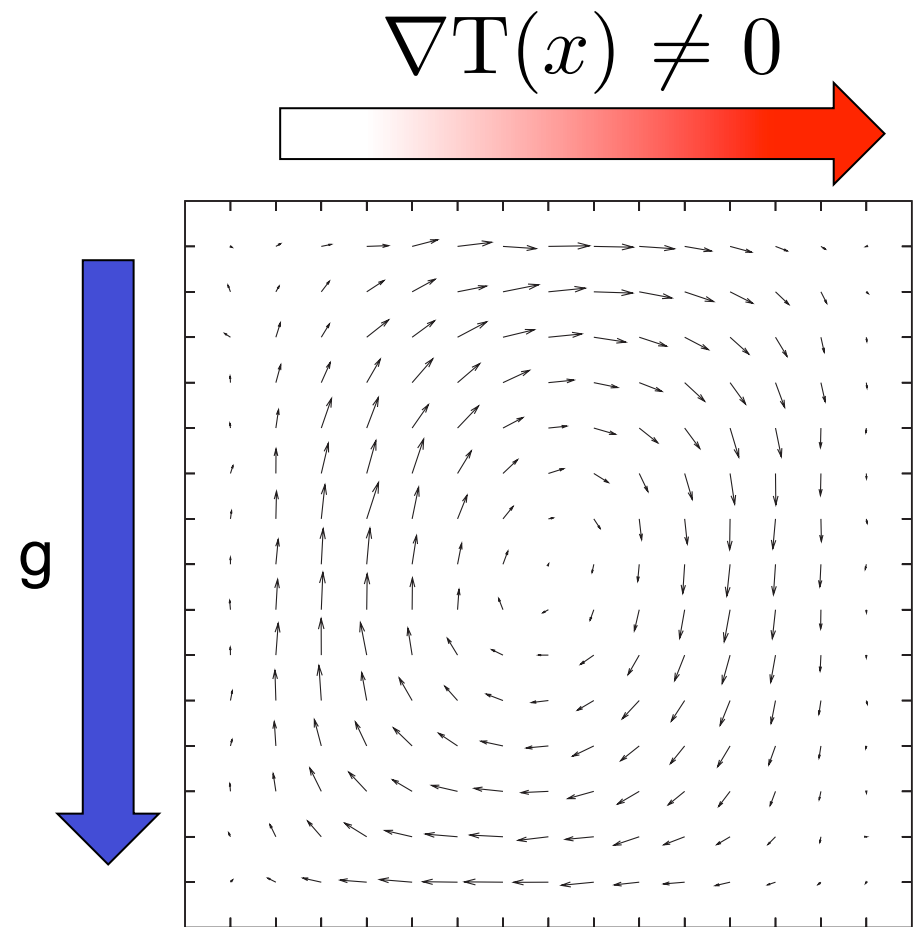


# Sampling of the initial distribution



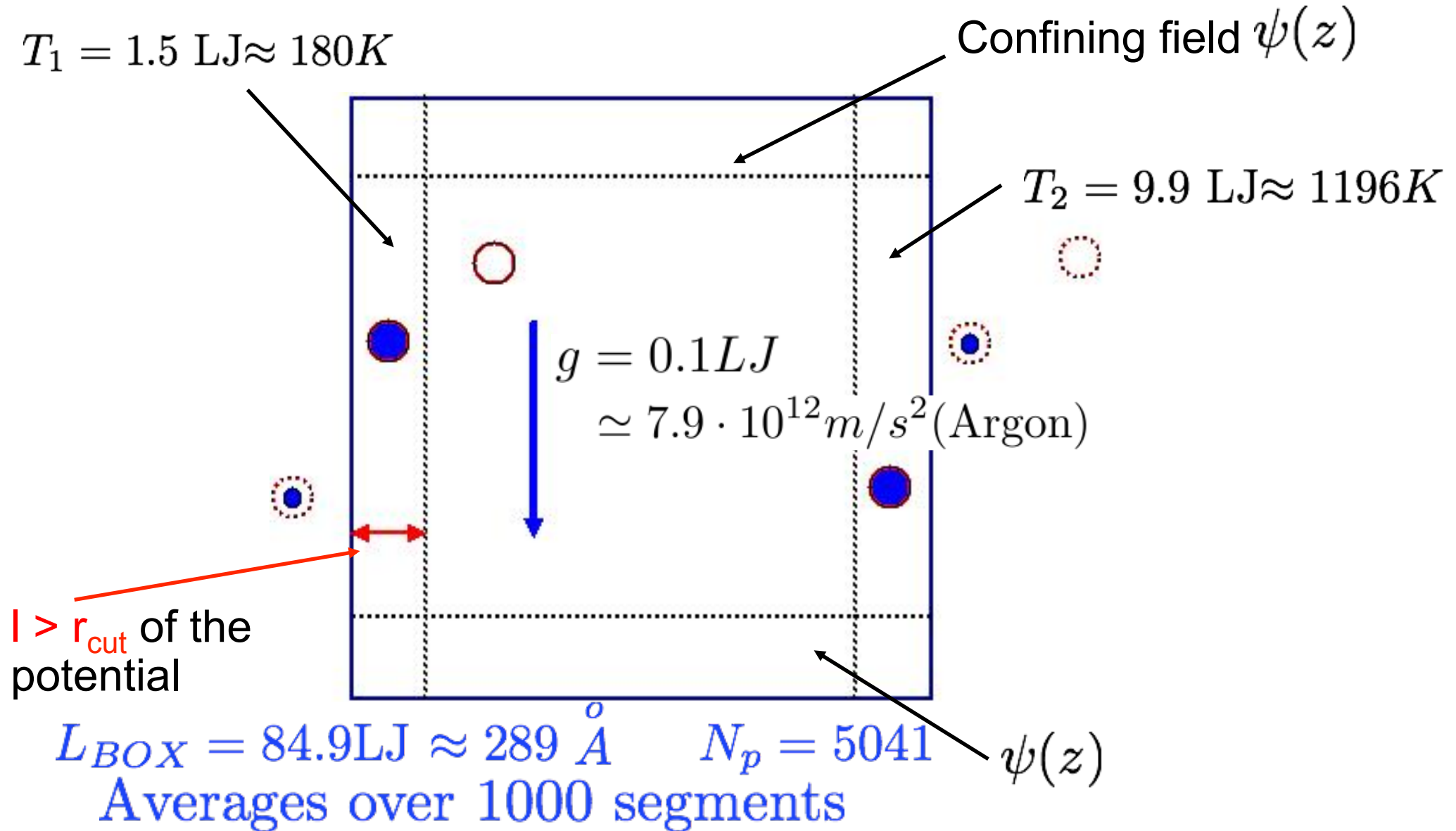
# Convection

- Sampling  $m_0(\Gamma) = m(\Gamma, t = 0)$ : standard stationary Non-Equilibrium MD with  $\nabla T(x) = \text{const.}, g = 0$
- Sampling  $J(x, t)$ : standard segments of MD, starting at  $t = 0$ , with  $\nabla T(x) = \text{const.}, g = \text{const.}$





# Convection



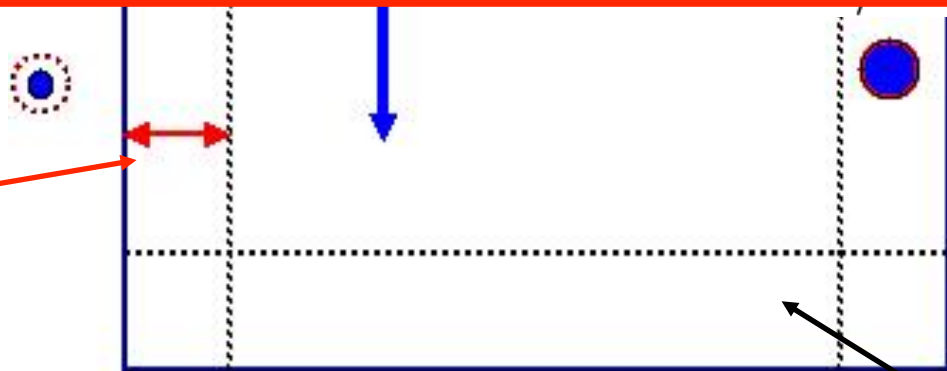
# Convection Cells

$$T_1 = 1.5 \text{ LJ} \approx 180 \text{ K}$$

Confining field  $\psi(z)$

The particles interact via a WCA potential (a LJ truncated and shifted at the minimum). A completely repulsive potential has only solid and fluid states and the thermodynamic conditions are such that the system is everywhere fluid.

1196K



$l > r_{\text{cut}}$  of the potential

$$L_{\text{BOX}} = 84.9 \text{ LJ} \approx 289 \overset{\circ}{\text{A}} \quad N_p = 5041$$

Averages over 1000 segments

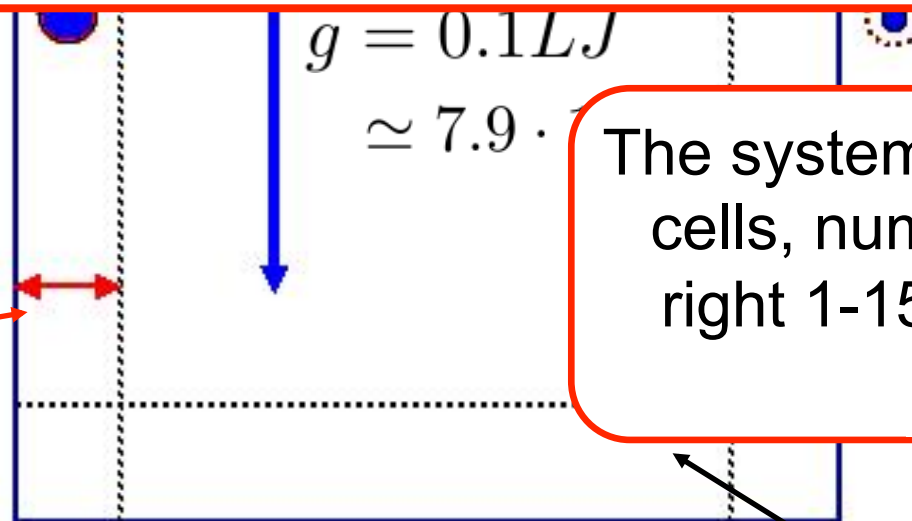
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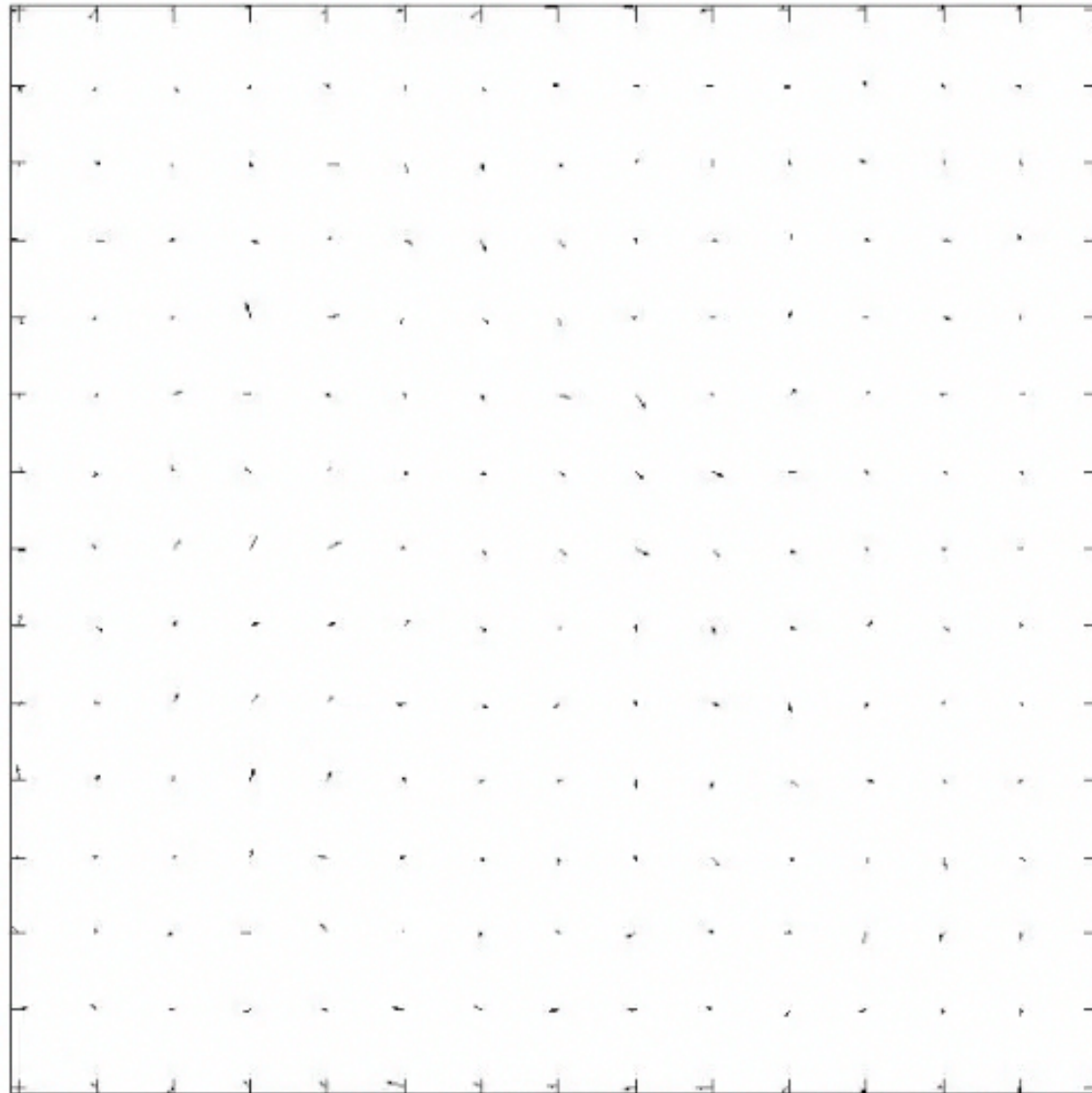


The system is divided in 15\*15 cells, numbered from left to right 1-15. A generic cell is  $(n_x, n_y)$

$$L_{BOX} = 84.9LJ \approx 289 \text{ \AA} \quad N_p = 5041$$

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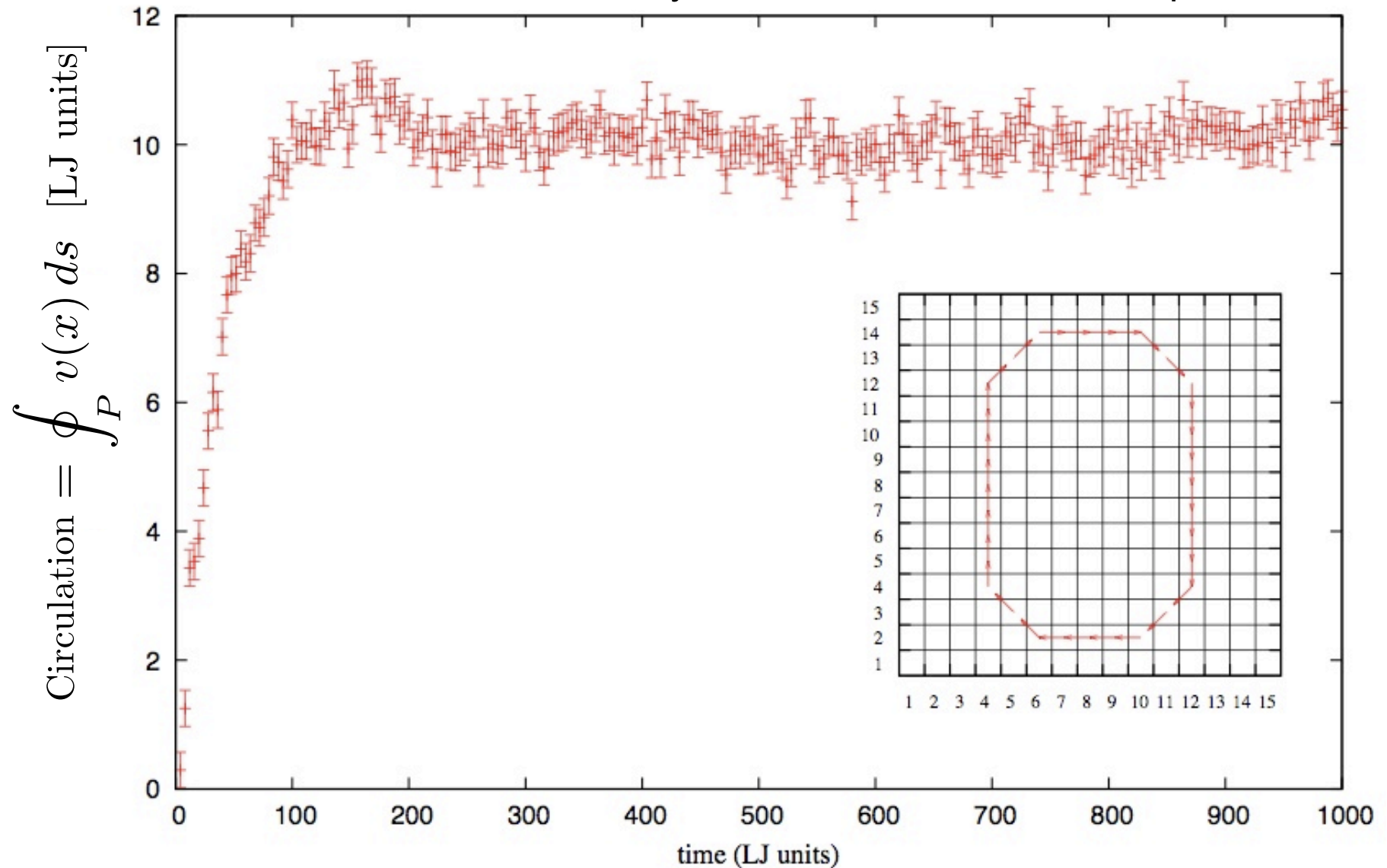
# Establishing Convective Cells



$t=0.0000$

# Establishing Convective Cells

Circulation of the velocity field. The inset shows the path.

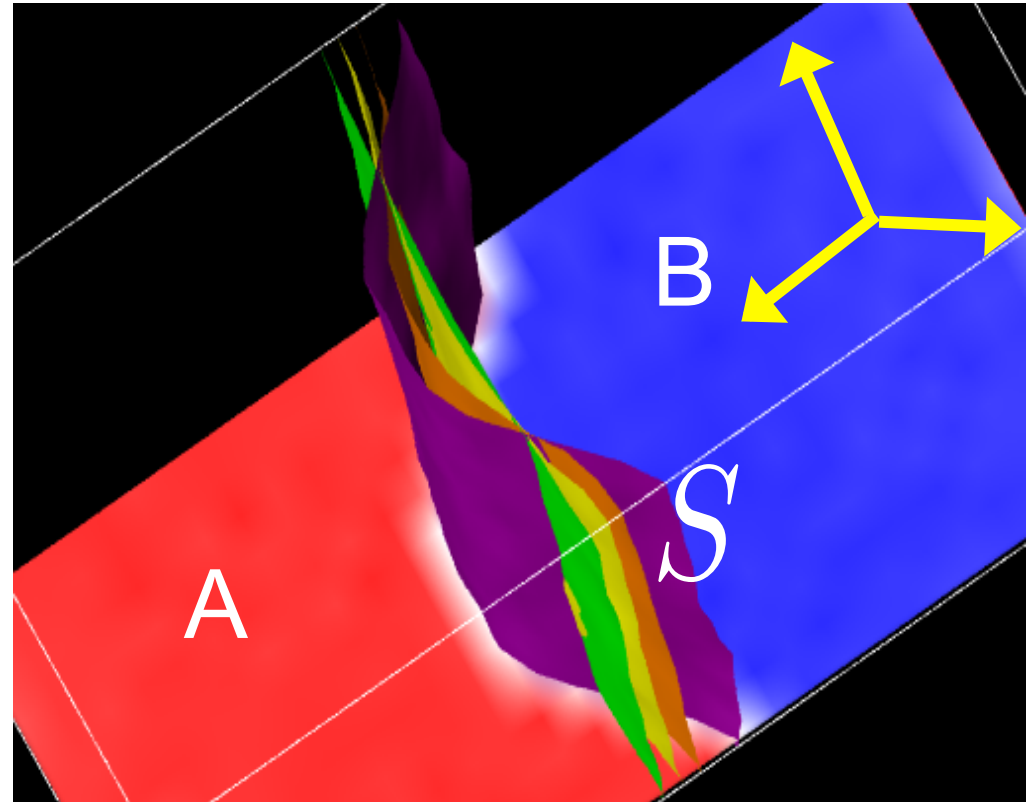


# Interface Relaxation

- Sampling

$m_0(\Gamma) = m(\Gamma | \Delta\rho(x) = 0, x \in S)$ ,  
 where  $\Delta\rho(x) = \rho^A(x) - \rho^B(x)$ :  
 restrained MD with  $\Delta\rho(x) = 0$   
 for  $x \in S$  and  $S$  a given  
 initial interface

- Sampling  $J(x, t)$ :  
 standard segments of MD with  
 system Hamiltonian



# Hydrodynamics by NEMD

- Field at the grid point  $\vec{x}_\beta$  of a discrete decomposition of the simulation box, with the atoms in the phase space point  $r, p \equiv (\vec{r}_1, \dots, \vec{p}_N)$ :

$$\hat{O}(\vec{x}_\alpha; r, p) = (1/\Omega_\alpha) \int_{\Omega_\alpha} d\vec{x} \sum_{i=1}^N \delta(\vec{x} - \vec{r}_i) O_i(r, p)$$

Where:

- $O_i(r, p) = \mu_i$  for the density
- $O_i(r, p) = p_i$  for the momentum density
- ...

# Hydrodynamics by NEMD

- The field at time  $t$  given initial macroscopic conditions is (Onsager-Kubo)

$$O[x, t | \Delta\rho(x_\alpha) = 0, x_\alpha \in S] = \int dr^0 dp^0 m_0 [r^0, p^0 | \Delta\rho(x_\alpha) = 0, x_\alpha \in S] \hat{O}(x; r(t), p(t))$$

where

$$m_0 [r^0, p^0 | \Delta\rho(x_\alpha; r^0) = 0, x_\alpha \in S] = \frac{\exp[-\beta H(r^0, p^0)] \prod_{x_\alpha \in S} \delta(\Delta\rho(x_\alpha; r^0))}{Z^* P_{\Delta\hat{\rho}}[\Delta\hat{\rho}(x_\alpha; r^0) = 0, x_\alpha \in S]}$$

and  $\begin{pmatrix} r(t) \\ p(t) \end{pmatrix} \equiv \begin{pmatrix} r(t; r^0, p^0) \\ p(t; r^0, p^0) \end{pmatrix}$

unrestrained MD started from a sample of points taken along a restrained MD (next slide)



# Conditional Averages by Restrained MD

- $\delta(\Delta\rho(x_\alpha; r^0))$  is smoothed by a Gaussian and the sampling of the ensemble at  $\Delta\hat{\rho}(\vec{x}_\alpha; r^0) = 0$  on the surface  $S$  is performed by restrained MD

$$\begin{aligned}\langle O(\vec{x}_\beta; r(t)) \rangle_{cond} &= \lim_{k \rightarrow \infty} \frac{\langle O(\vec{x}_\beta; r(t)) \prod_{\vec{x}_\alpha \in S} \exp[-\beta \frac{k}{2} \Delta\hat{\rho}(\vec{x}_\alpha; r^0)^2] \rangle_H}{\langle \prod_{\vec{x}_\alpha \in S} \exp[-\beta \frac{k}{2} \Delta\hat{\rho}(\vec{x}_\alpha; r^0)^2] \rangle_H} \\ &= \lim_{k \rightarrow \infty} \langle O(\vec{x}_\beta; r) \rangle_{H'}\end{aligned}$$

where:

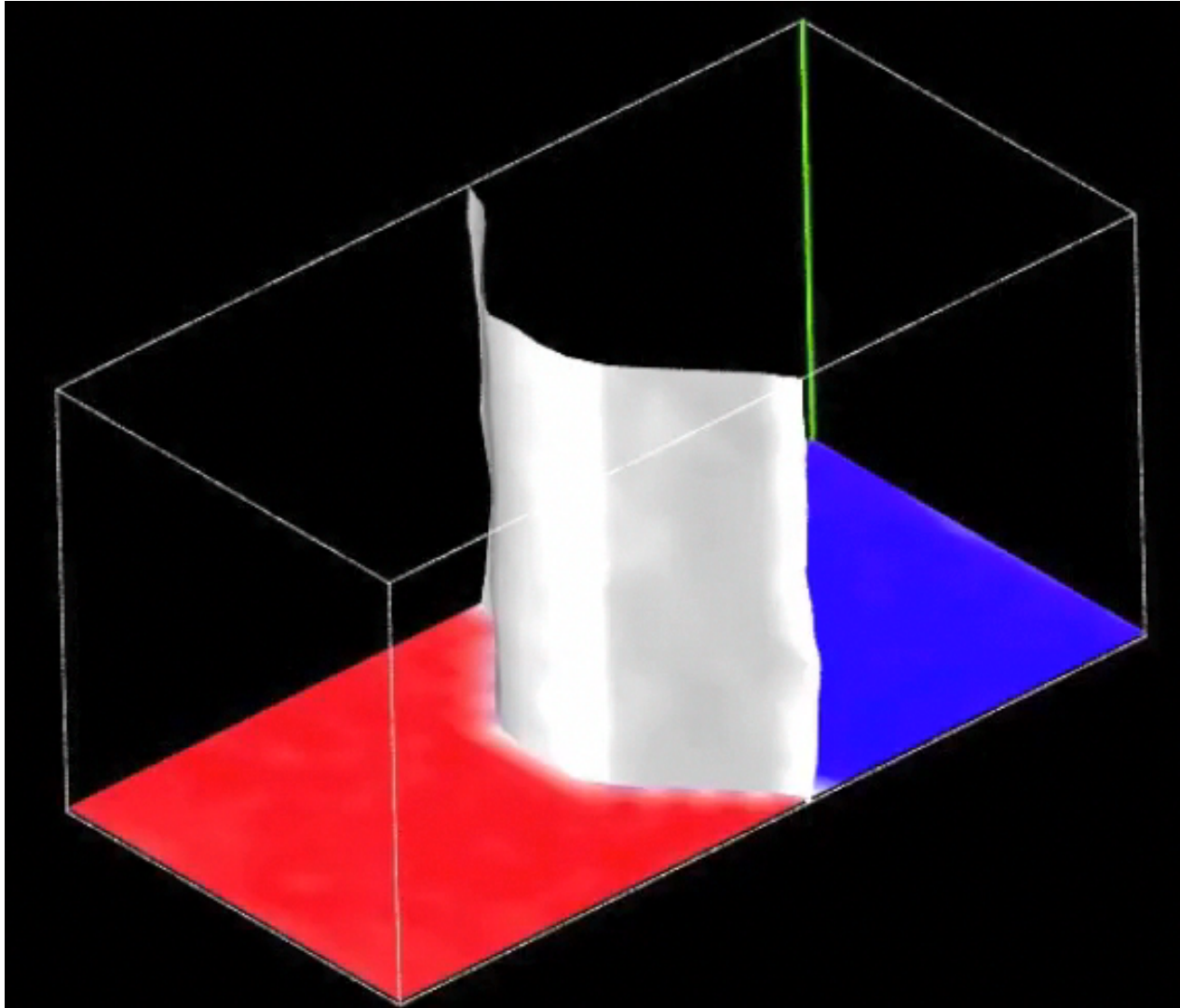
$$H'(r, p) = H(r, p) + \frac{k}{2} \sum_{\vec{x}_\alpha \in S} \Delta\hat{\rho}(\vec{x}_\alpha; r)^2$$

# Simulation Details

- 171500 particles: 88889 particles A, 82611 particles B
- Pair potential:  $u^{AA}(r) = u^{BB}(r) = 4\varepsilon \left( \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^6 \right)$   $u^{AB}(r) = 4\varepsilon \left( \frac{\sigma}{r} \right)^{12}$
- Simulation box:  $\sim(90 \times 45 \times 45) \sigma$
- Average density:  $1.024 \text{ particles} \cdot \sigma^3$
- Temperature:  $1.5 \varepsilon/k_b$
- Simulation time:
  - Restrained MD: 75000 steps
  - Unrestrained MD: 600000 steps
- fields are averaged over (only) 40 unrestrained trajectories (for the moment)

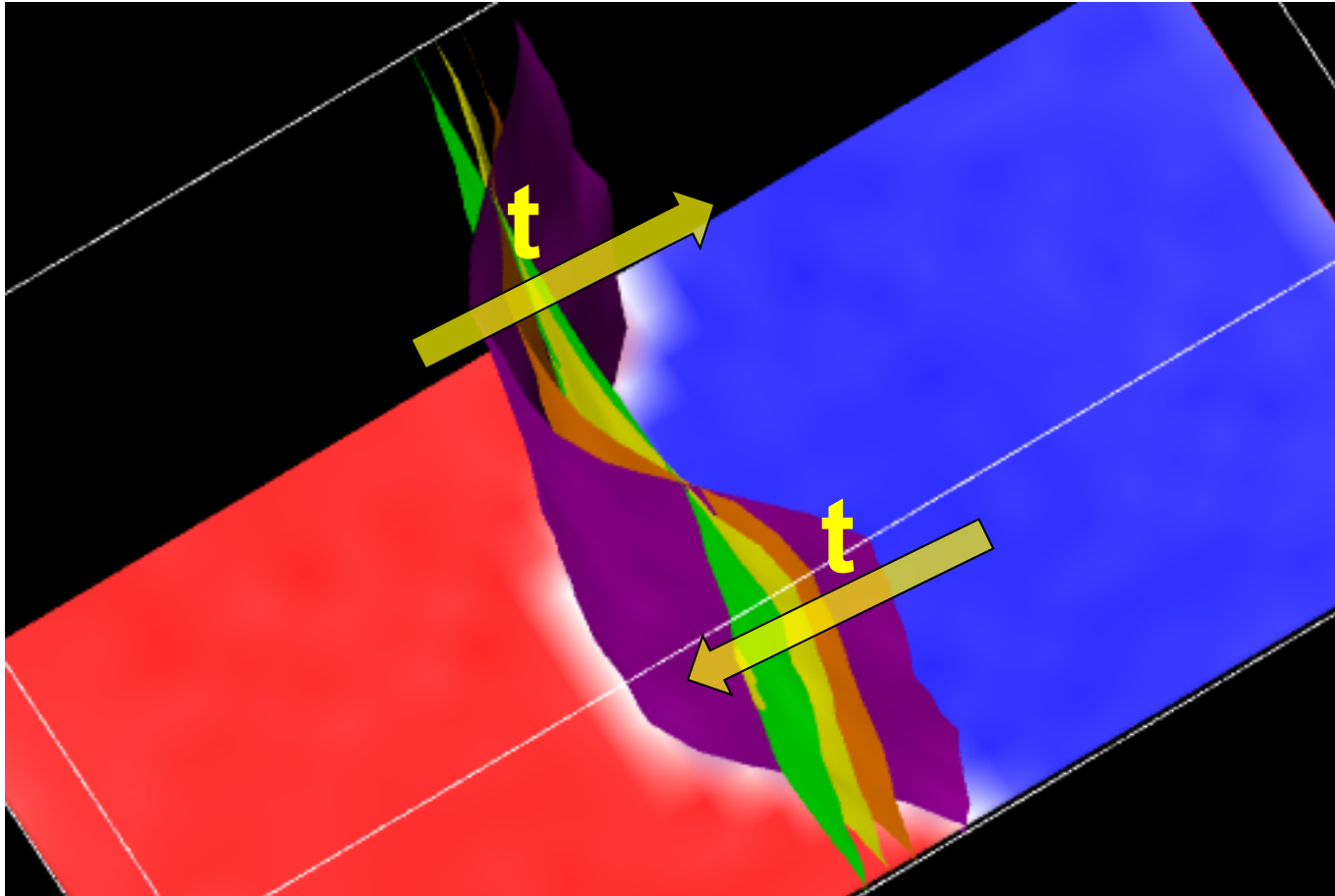
In the fluid domain of pure L-J

$$\Delta\rho(\vec{x}_\alpha; t) = 0$$



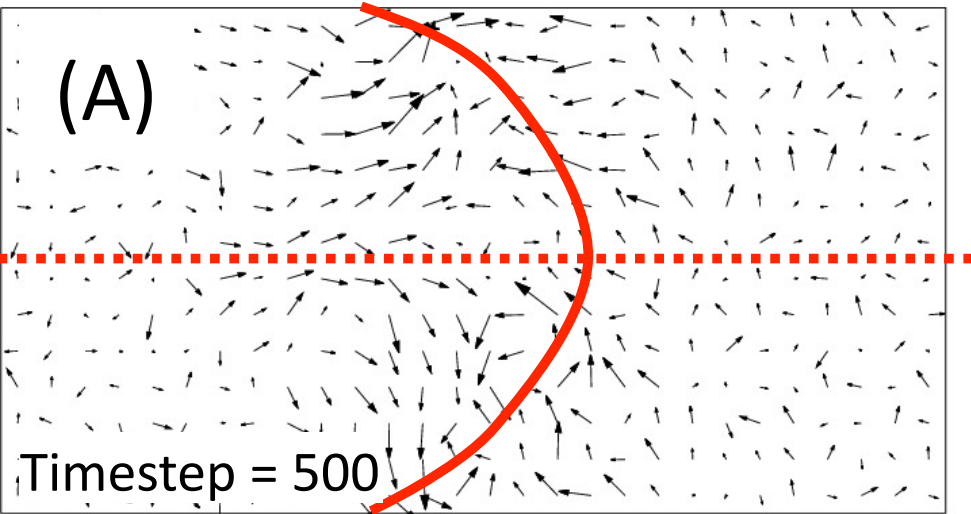
$$v(\vec{x}_{\alpha, \max}) \sim 80 \text{ m/s}$$

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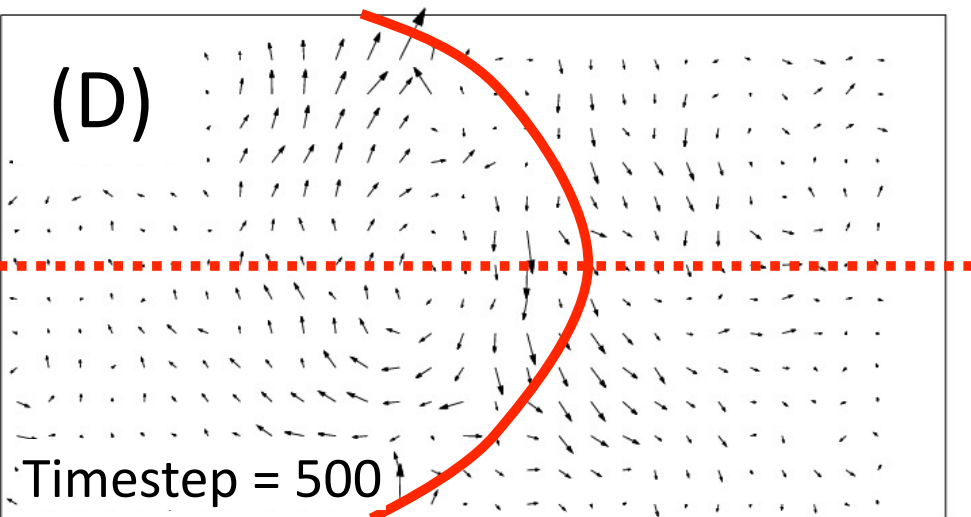


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# Rigorous non-equilibrium ensemble averages vs local time averages



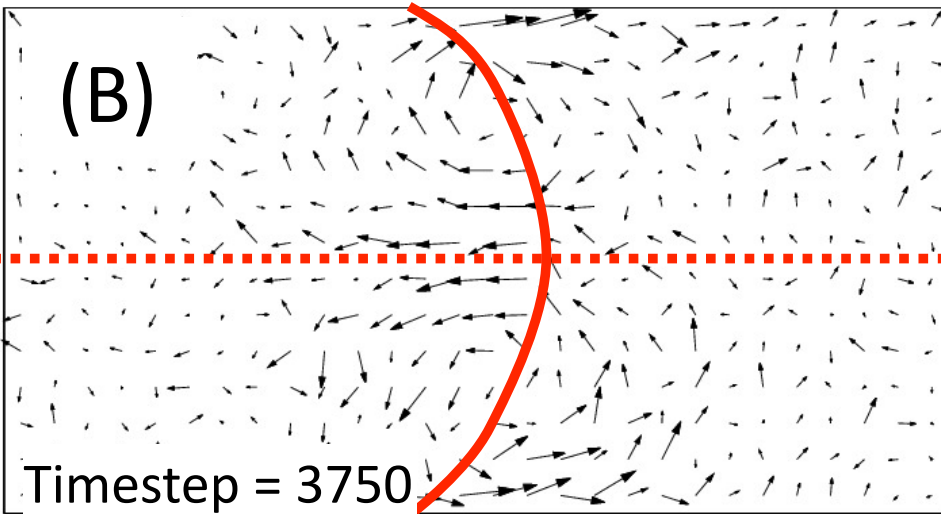
$$\vec{v}(\vec{x}, t) = \frac{\langle \hat{p}(\vec{x}, t) \rangle_{H'}}{\rho(\vec{x}, t)}$$



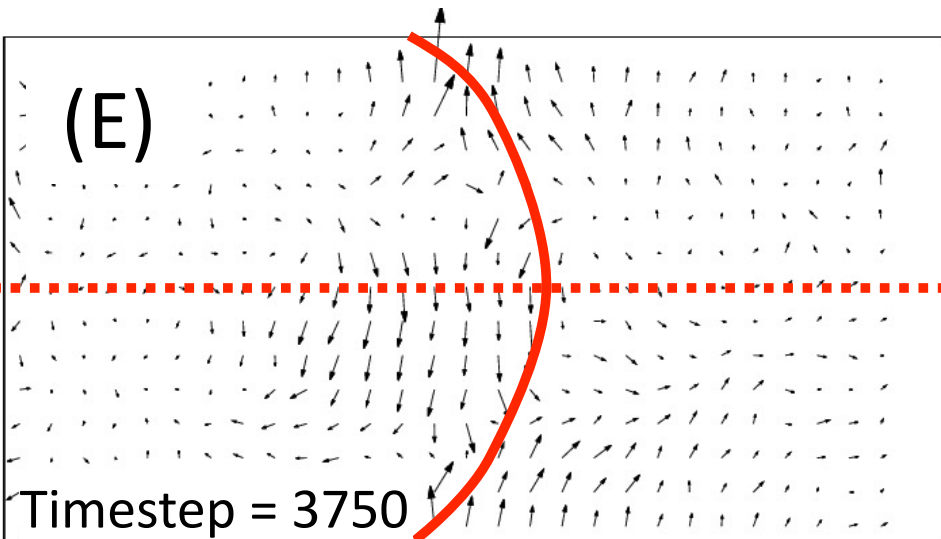
$$\vec{v}_l(\vec{x}, t) = \frac{\frac{1}{2\tau} \int_{t-\tau}^{t+\tau} ds \hat{p}(\vec{x}, s)}{\frac{1}{2\tau} \int_{t-\tau}^{t+\tau} ds \hat{\rho}(\vec{x}, s)}$$

- The surface relaxes to the equilibrium by forming initially a two-tail profile of the velocity field that then stabilizes into a double-roll profile
- The velocity field obtained via the local time average technique violates the symmetry of the problem

# Rigorous non-equilibrium ensemble averages vs local time averages



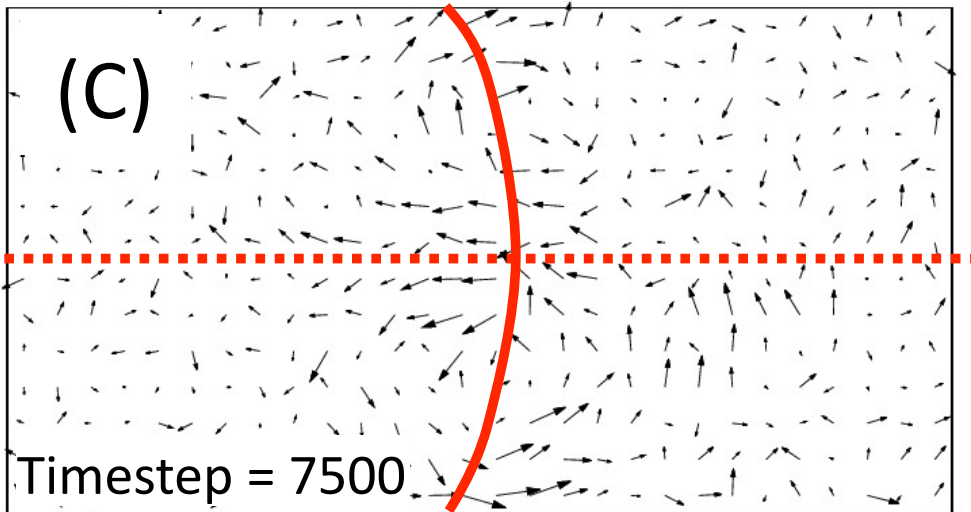
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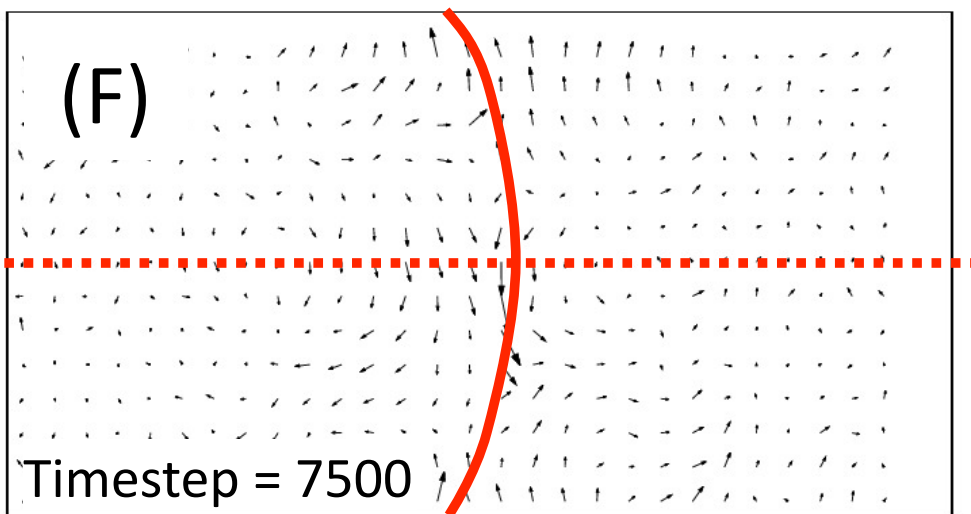
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# Conclusions

- ① It is possible to **compute**, numerically but, otherwise, **rigorously, time-dependent non-equilibrium responses**, i.e. responses in non-stationary regimes. **Local time-averages should be avoided**
- ② Nonequilibrium atomistic dynamics possibly combined with the ability to compute conditional averages (restrained MD) allows to simulate hydrodynamic phenomena *ab initio* (without using phenomenological approximations).
- ③ **Coupling non-equilibrium non-stationary systems** is a **fundamental question of multi-scale approaches**





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Computer resources

