

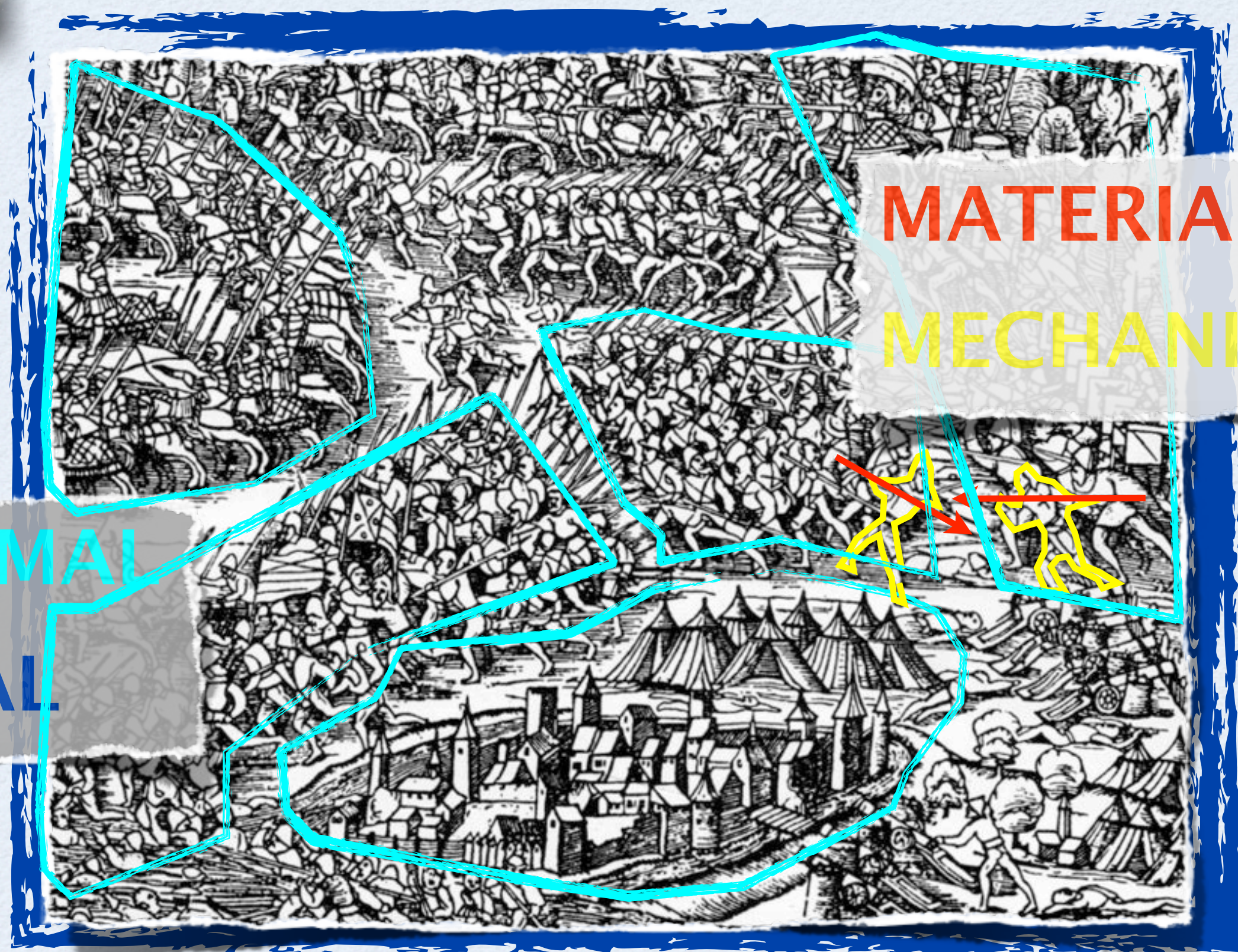
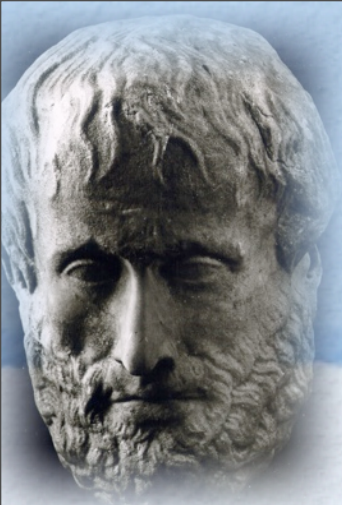
# Adaptive Stochastic and Deterministic Simulations Using Particles

Petros Koumoutsakos



# MULTISCALING as CAUSALITY

adapted from Ulanowicz



**MATERIAL -  
MECHANICAL**

**FORMAL  
FINAL**



# OUTLINE

- INTRODUCTION
- STOCHASTIC
  - R-Leaping + AMR-S
- DETERMINISTIC
  - Particles + Wavelets
- BOUNDARIES
- OUTLOOK



# MULTIPLE SCALES + PARTICLES

Molecular  
Dynamics

-9

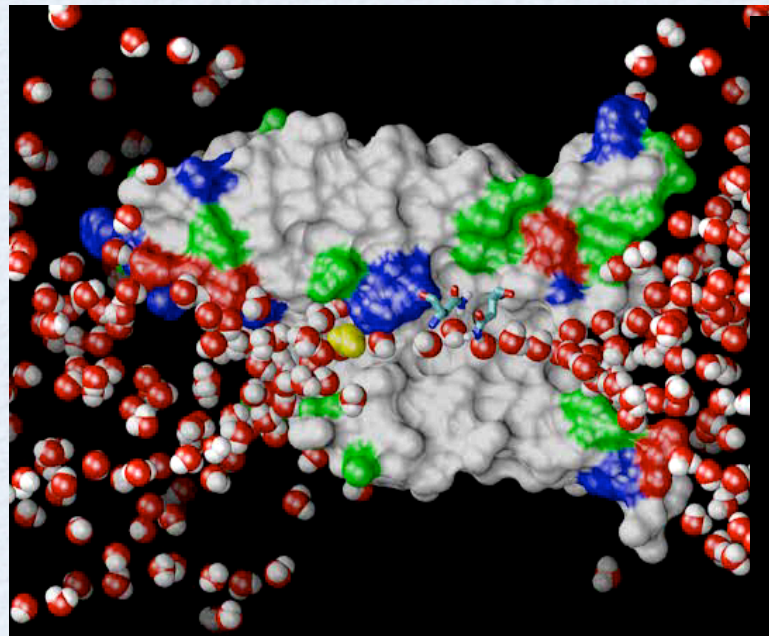
Vortex  
Methods

0

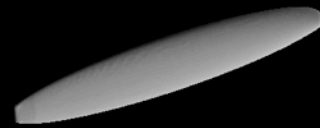
Smoothed Particle  
Hydrodynamics

+9

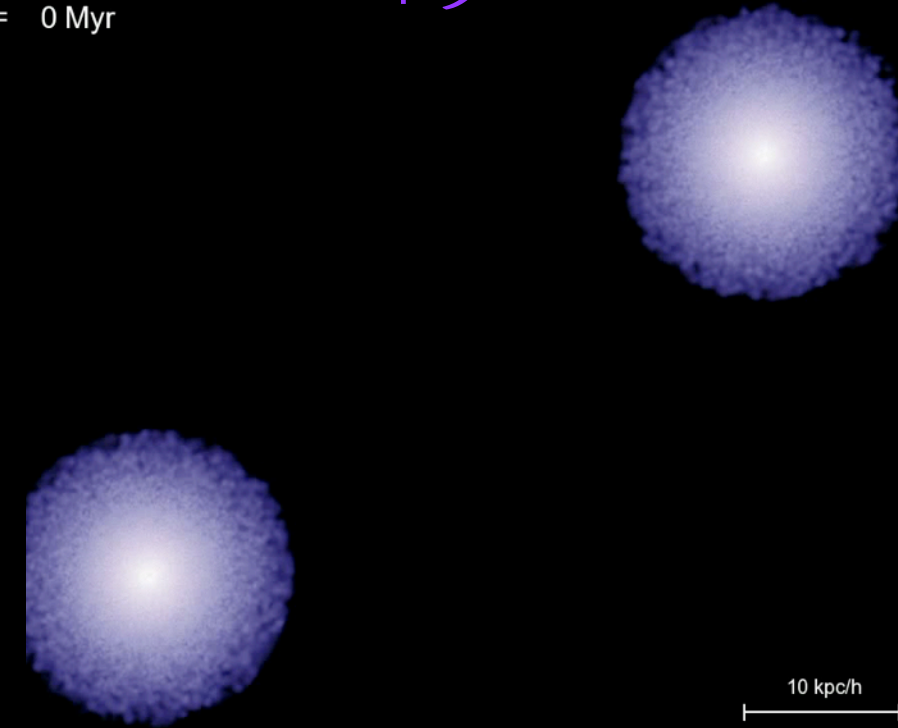
T = 0 Myr



Transport in aquaporins  
Schulten Lab, UIUC



Vortex Dynamics  
Koumoutsakos Lab, ETHZ



Growth of Black Holes  
Springel, MPI - Hernquist, Harvard



# PARTICLES : Lagrangian, Conservation and Other Laws

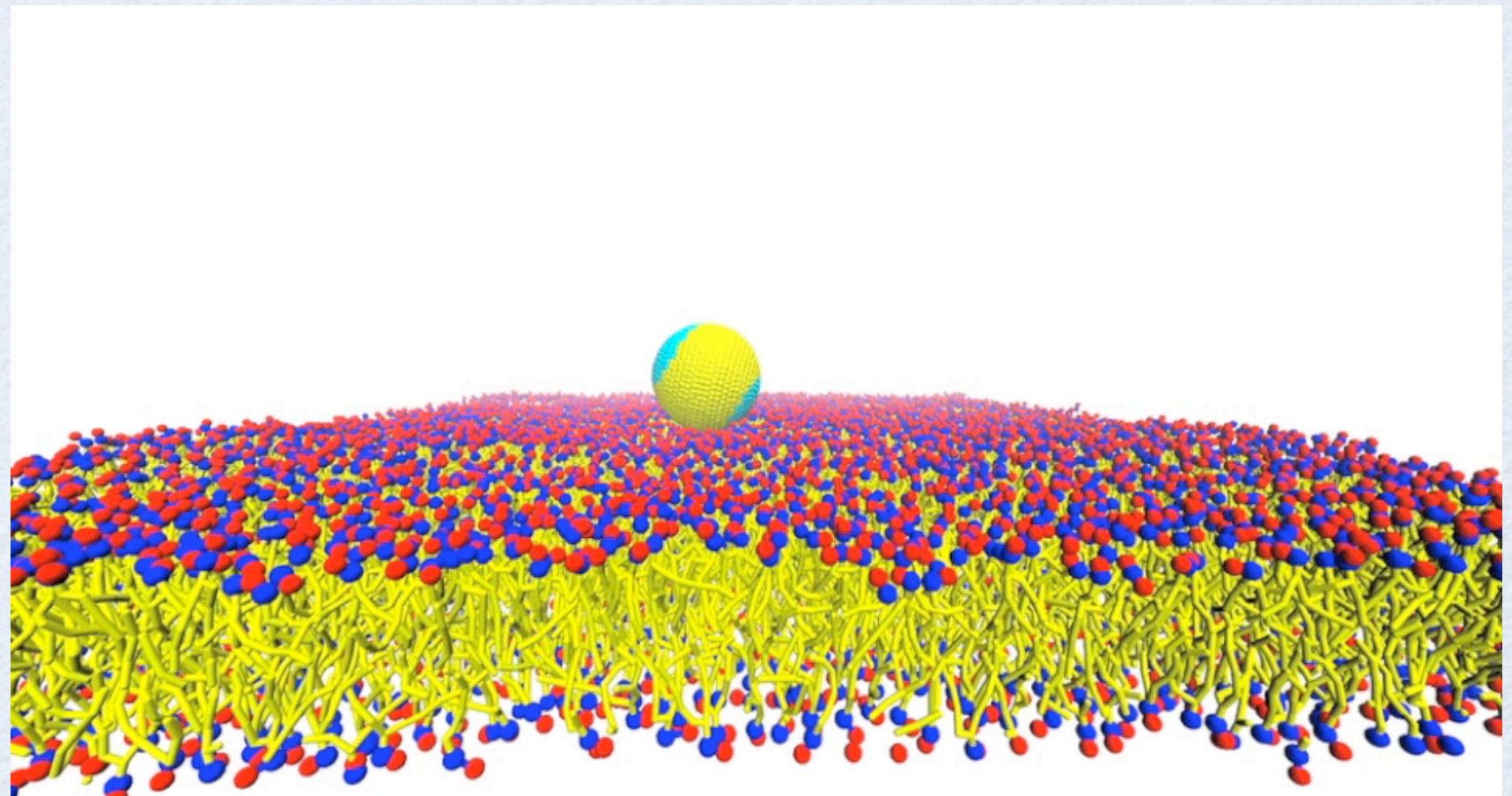
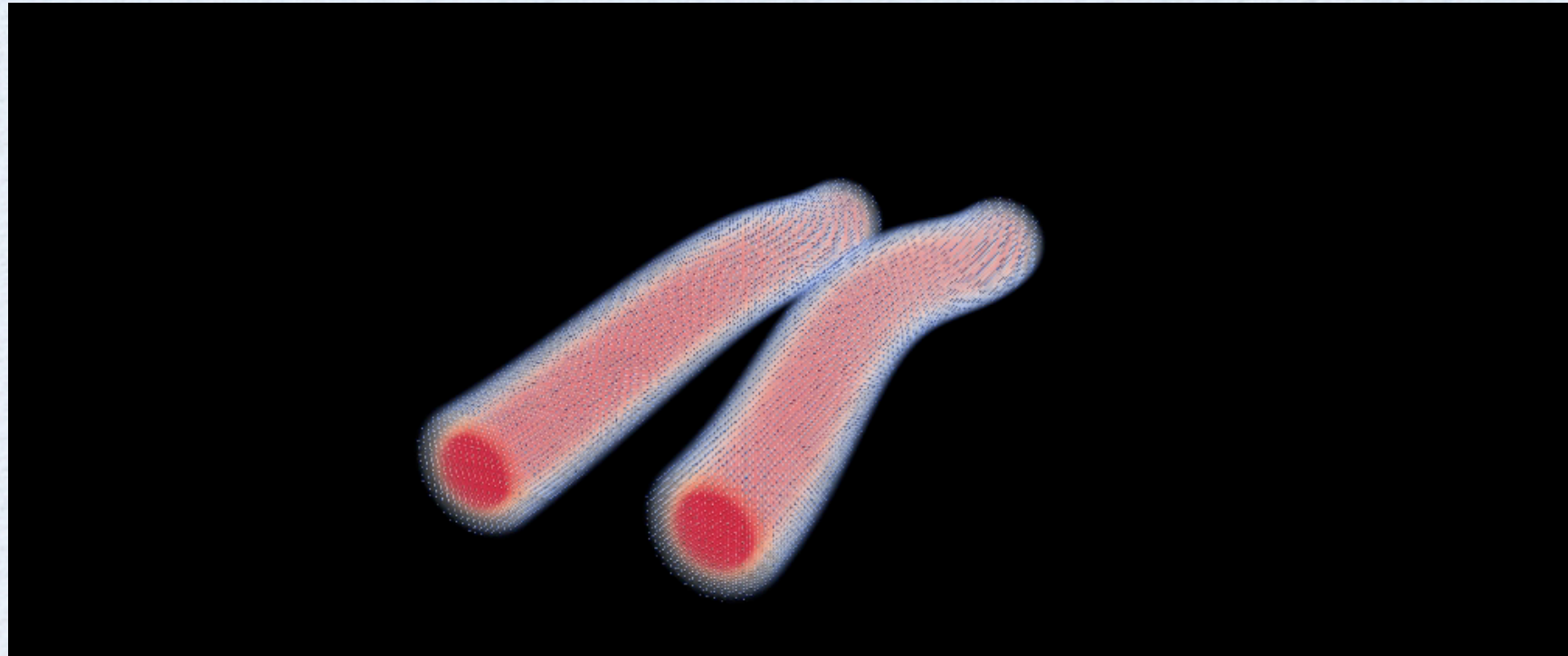
## SPH, Vortex Methods

$$\rho_p \frac{D\mathbf{u}_p}{Dt} = (\nabla \cdot \sigma)_p$$

$$\frac{d\mathbf{x}_p}{dt} = \mathbf{u}_p$$

$$m \frac{d\mathbf{u}_p}{dt} = F_p$$

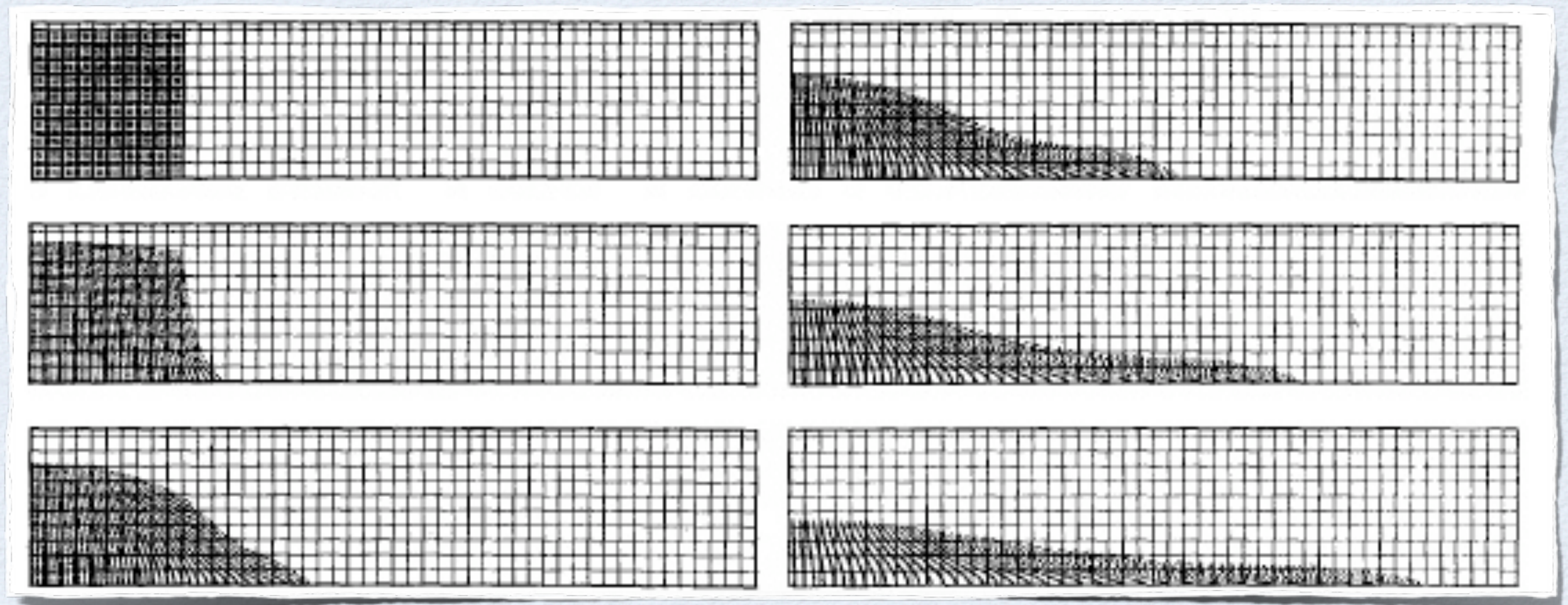
## MD, DPD, CGMD





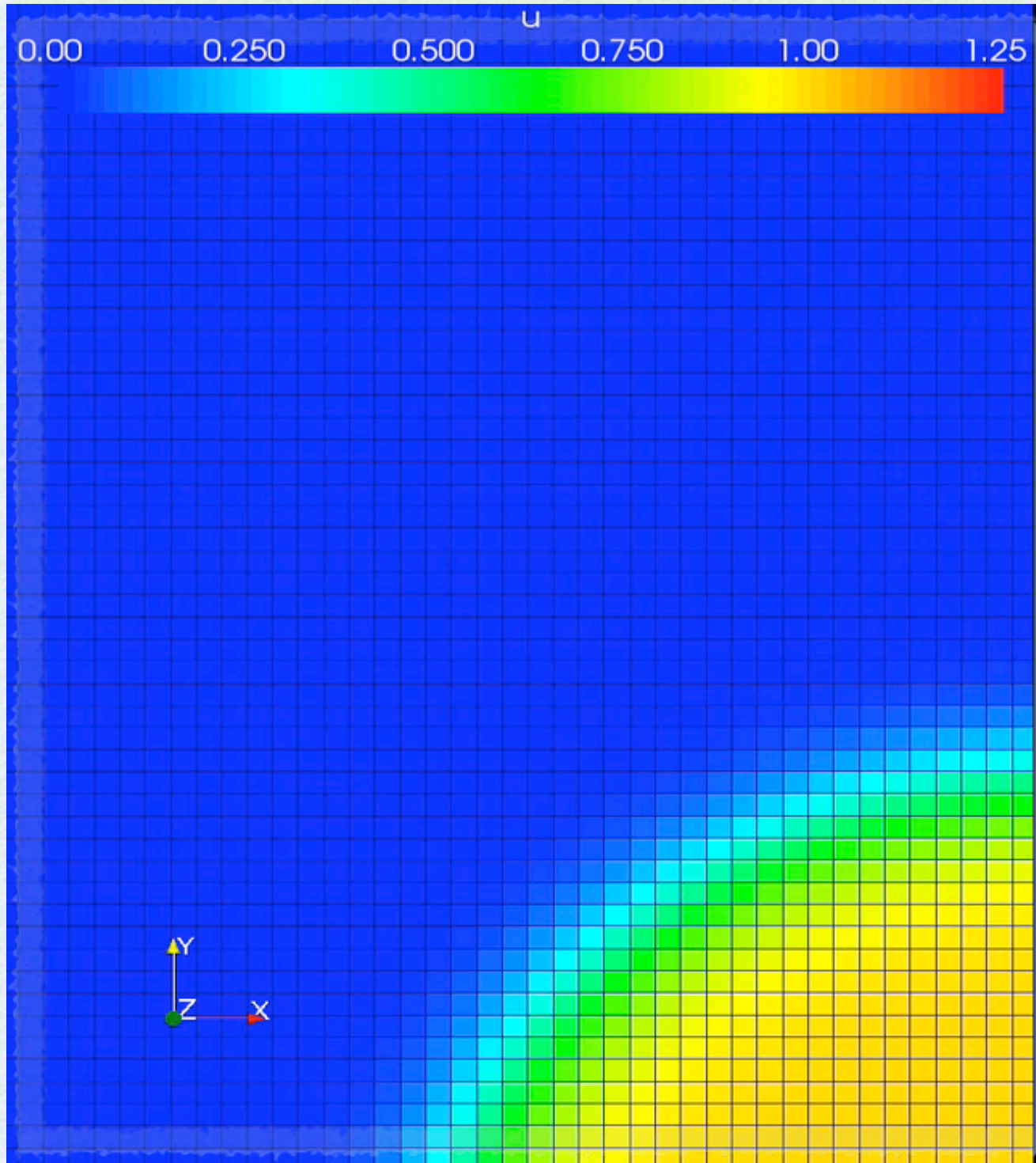
# Particles and Grids

Marker and Cell (MAC) - F.H. Harlow and E.J. Welch

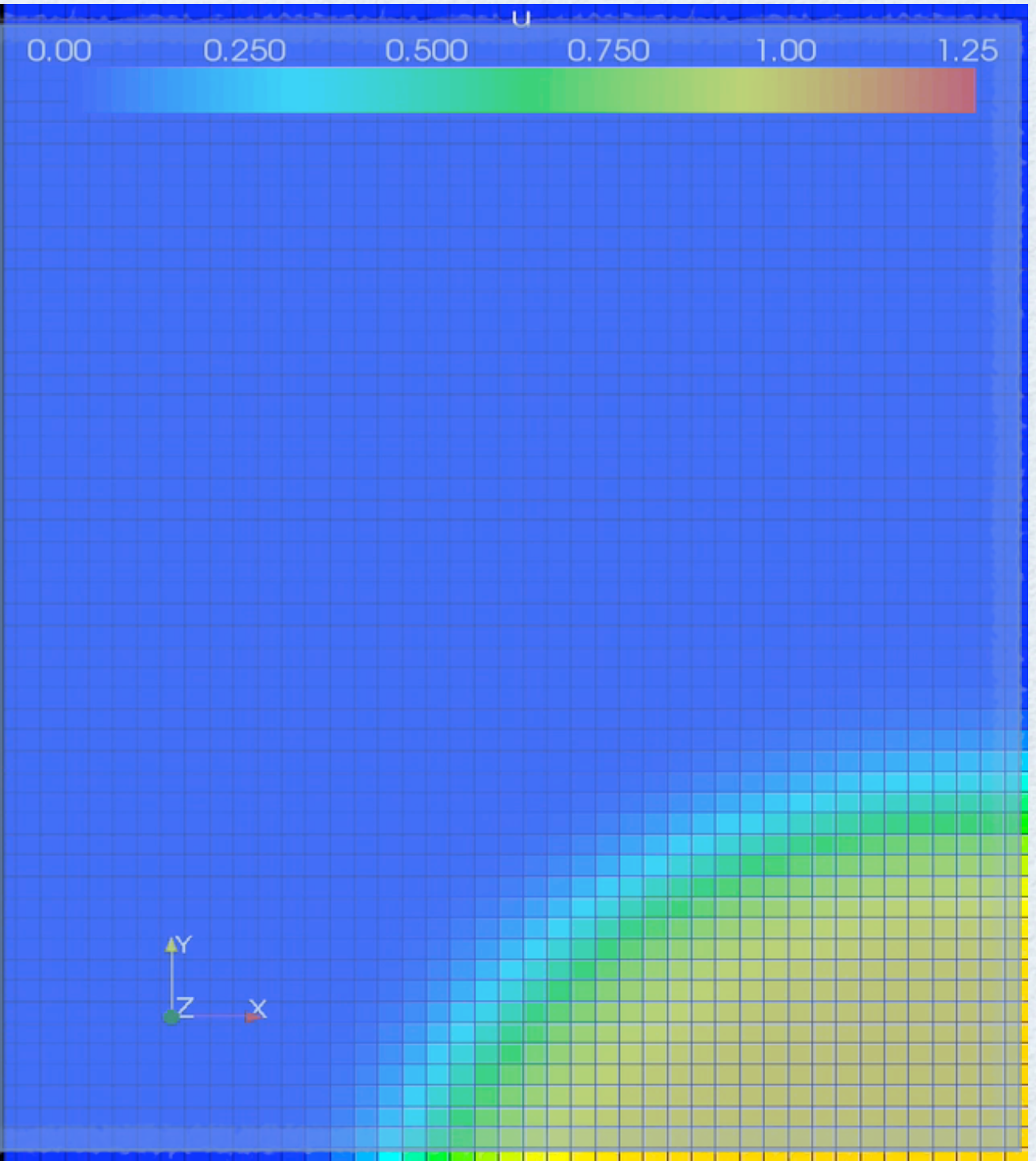


*Numerical Calculation of Time-Dependent Viscous Incompressible Flow of Fluid with Free Surface*, Harlow, Francis H. and Welch, J. Eddie, Physics of Fluids, 1965





**STOCHASTIC**

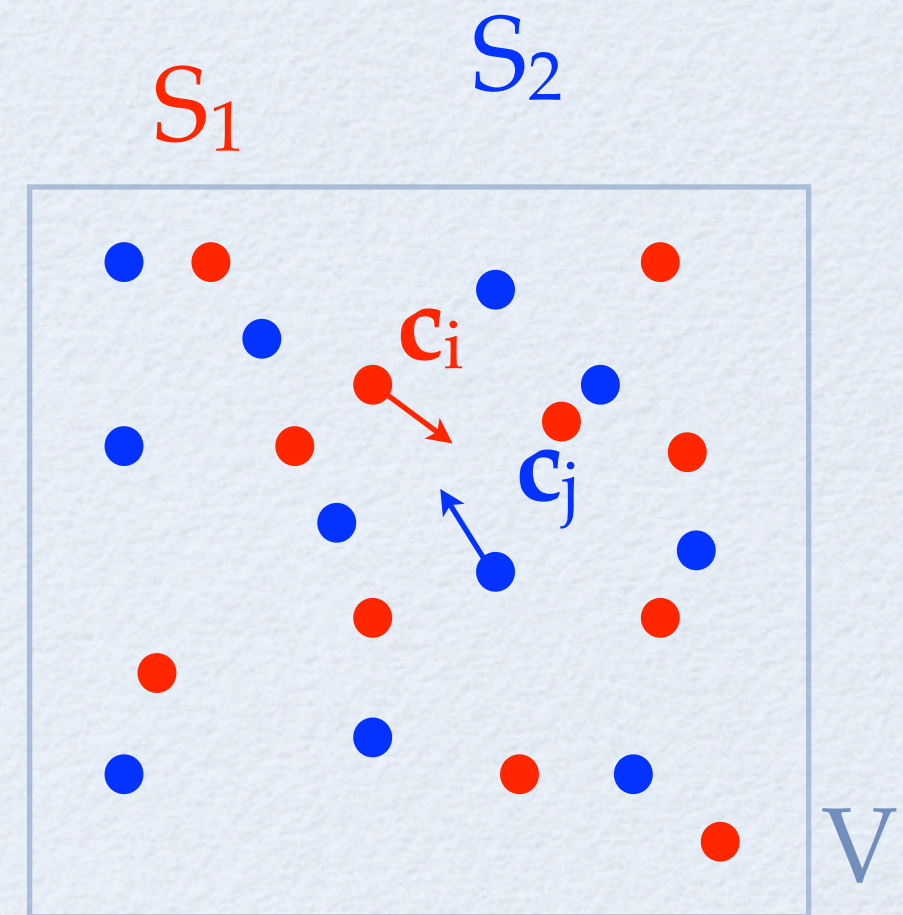


**DETERMINISTIC**



# Chemical kinetics : Set-up

- Well stirred reaction volume  $V$
- Experiment length  $T$
- $N$  different species  $S_1, S_2, \dots, S_N$  in numbers  $X_1, X_2, \dots, X_N$
- random collisions and reactions through  $M$  channels  $R_1, R_2, \dots, R_M$





# Kinetics + Space

## Diffusion in 1-D

### Deterministic

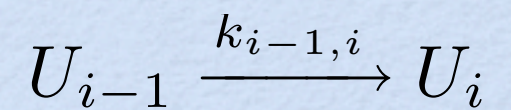
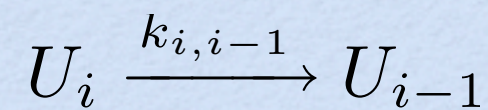
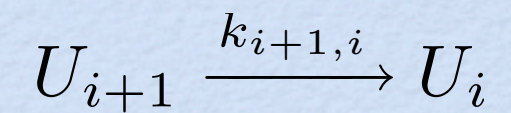
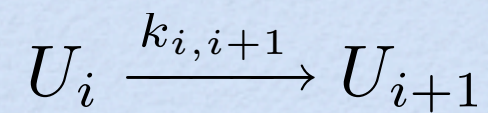
$$\frac{\partial u}{\partial t} = D \frac{\partial^2 u}{\partial x^2}$$

### Stochastic

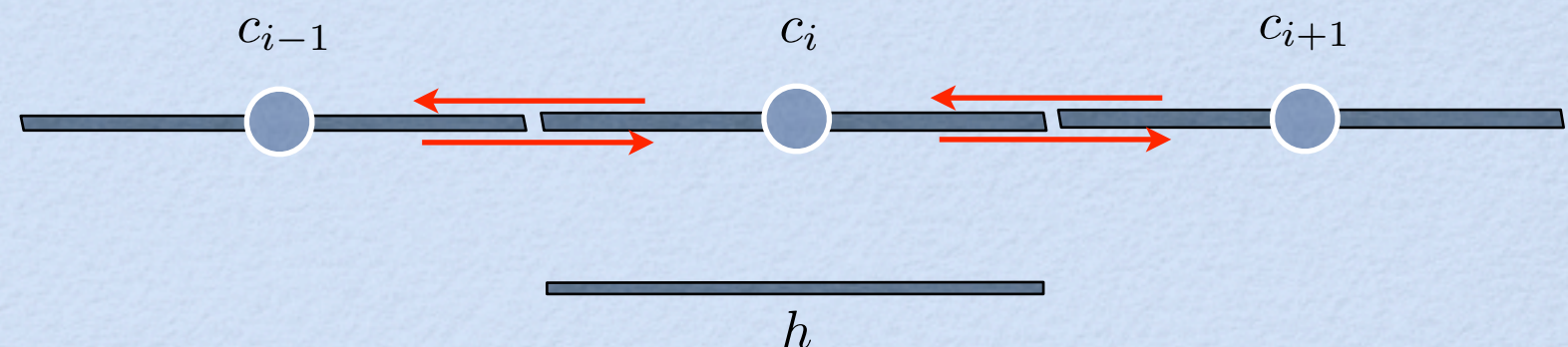
D. Bernstein, *Phys. Rev. E*, 2005.

A species  $U$ , with elements labeled by an index  $i$

**Diffusion as reactions of the form:**



Uniform Cells:  $k_{i,j} = \frac{D}{h^2}$





# SPACE: $10^6 - 10^9$ "Reactions"

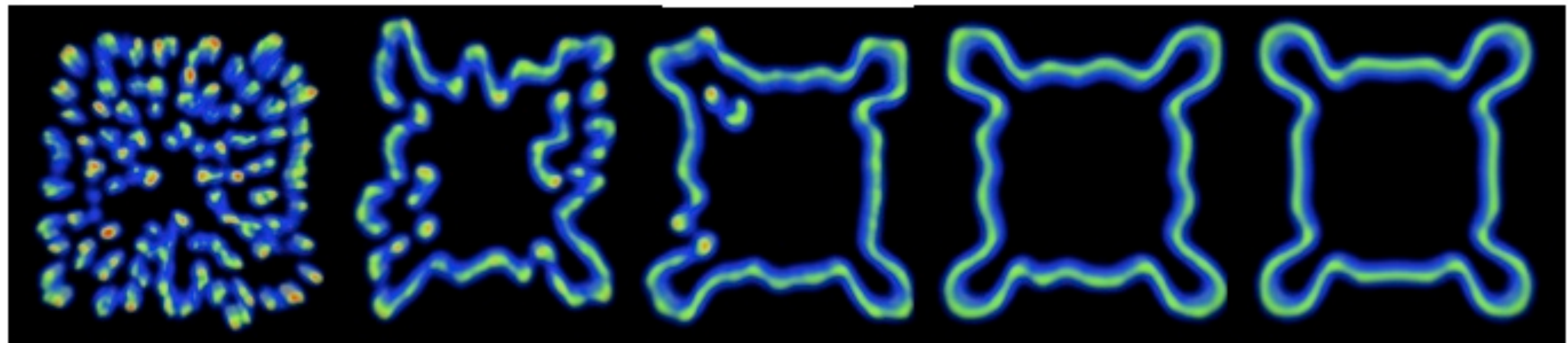
Molecules per grid cell for a 300 x 300 grid

500

1000

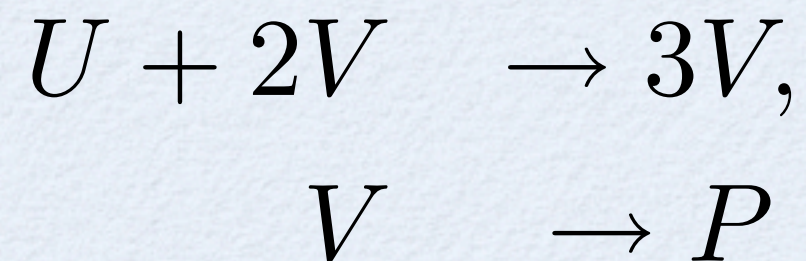
5000

10000



Microscopic scale

Macroscopic scale



$$\frac{\partial u}{\partial t} = d_u \Delta u - uv^2 + F(1 - u),$$

$$\frac{\partial v}{\partial t} = d_v \Delta v + uv^2 - (F + \kappa)v.$$

$$F = 0.04, \kappa = 0.06, t = 1000$$



# Stochastic Simulation Algorithm

Gillespie,  
J. Comp. Phys. 1977

- For  $M$  reactions, time until **any** reaction

$$\tau \sim \mathcal{E}(1/a_0)$$

$$a_0 = \sum_{j=1}^M a_j$$

- Reaction **index** : point-wise distribution

$$p(j = l) = \frac{a_l}{a_0}$$

## ONE STEP

- Sample  $\tau$
- Sample the index  $j$
- Update the  $X_i$ ,  $t=t+\tau$

The SSA simulates  
every reaction event !

Exact but SLOW



# ACCELERATING SSA : $\tau$ leaping

Gillespie,  
J. Chem. Phys. 2001

$\tau$  leaping : several reaction events over one time step

**ASSUMPTION** : reaction propensities  $a_i$  remain essentially constant over  $\tau$ , in spite of several firings

- Over this given  $\tau$ , the number of reaction firings  $K_j^{\mathcal{P}}$  is governed by a Poisson distribution

$$K_j^{\mathcal{P}} \sim \mathcal{P}(a_j \tau)$$
$$\mathbf{X}(t + \tau) = \mathbf{X}(t) + \sum_{j=1}^M K_j^{\mathcal{P}} \boldsymbol{\nu}_j.$$

Cost ~ M Poisson samplings

SPEEDUP ~ 100 X SSA



# $\tau$ -leaping : Consequences

$\tau$  leaping can generate negative populations

- Binomial  $\tau$  leaping : Approximate the unbounded Poisson distributions with Binomial ones

Chatterjee et al.,  
J. Chem. Phys. 2005

- Modified  $\tau$  leaping

Tian & Burrage,  
J. Chem. Phys. 2004

- Critical reactions, i.e. those likely to drive some populations negative, handled by SSA
- Other reactions advanced by  $\tau$  leaping

Cao et al.,  
J. Chem. Phys. 2005



# R-leaping : Accelerate SSA by reaction leaps

**Leaps** : number of firings  $L$  across all reaction channels

- Time increment  $\tau_L$  is Gamma-distributed  $\tau_L \sim \Gamma(L, 1/a_0(\mathbf{x}))$

- In this interval we will have  $K_m$  firings of channel  $R_m$

- with : 
$$\sum_{m=1}^M K_m = L$$

- In R-leaping, as in SSA, the index  $j$  of every firing obeys a point-wise distribution

$$P(j = l) = \frac{a_l(\mathbf{x})}{a_0(\mathbf{x})} \text{ for } l = 1, \dots, M.$$



# R-leaping : One step

- Define L

$$\tau_L \sim \Gamma(L, 1/a_0(\mathbf{x}))$$

- Sample the index j

$$P(j = l) = \frac{a_l(\mathbf{x})}{a_0(\mathbf{x})} \quad \text{for } l = 1, \dots, L.$$

- Number of reactions for channel m

$$K_m = \sum_{l=1}^L \delta_{l,m}$$

- Update species and time :

$$\mathbf{X}(t + \tau_L) = \mathbf{X}(t) + \sum_{j=1}^M K_j \boldsymbol{\nu}_j$$

Auger et al.,  
J. Chem. Phys. 2006



# R-Leaping Theorem

The distribution of  $K_1$  is a binomial distribution :

$$\mathcal{B}(L, a_1(\mathbf{x})/a_0(\mathbf{x}))$$

and for every  $m \in \{2, \dots, M\}$  the conditional distribution of  $K_m$

given the event  $\{(K_1, \dots, K_{m-1}) = (k_1, \dots, k_{m-1})\}$  is

$$K_m \sim \mathcal{B} \left( L - \sum_{i=1}^{m-1} k_i, \frac{a_m(\mathbf{x})}{a_0(\mathbf{x}) - \sum_{i=1}^{m-1} a_i(\mathbf{x})} \right).$$

This result is invariant under any permutation of the indices

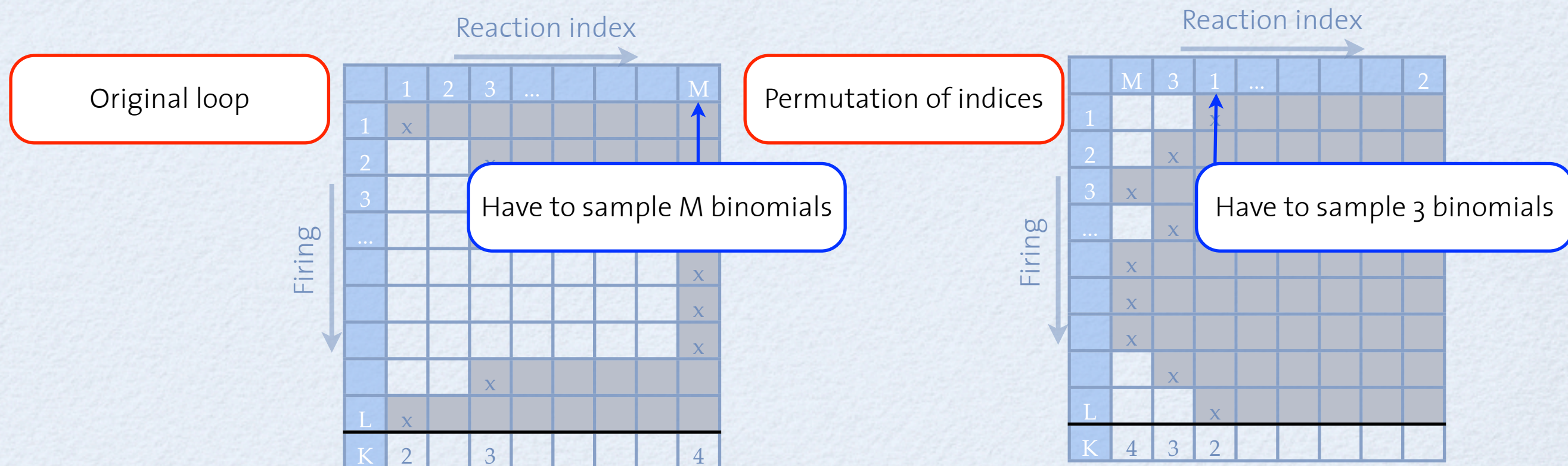


# R-leaping : Sorting Reactions

- *Sampling the  $M K_j$  efficiently*

If  $K_i = k_i, \forall i < m, K_m \sim \mathcal{B} \left( L - \sum_{i=1}^{m-1} k_i, \frac{a_m}{a_0 - \sum_{i=1}^{m-1} a_i} \right)$

- When  $\sum_{i=1}^{m-1} k_i = L$  , sampling is done!
- Minimize the average  $m$  by a **permutation of the indices**, such that propensities are decreasing





# Results

- LacZ/LacY genes expression and enzymatic/transport activities of LacZ/LacY proteins in E. Coli

Kierzek,  
Bioinformatics 2002

- Moderately large system ( $M = 22$ )
- Disparate rates
- Scarce reactants and negative species

|          | Reaction Channel   | Reaction rate         |
|----------|--|-----------------------|
| $R_1$    | $\text{PLac} + \text{RNAP} \rightarrow \text{PLacRNAP}$                    | 0.17                  |
| $R_2$    | $\text{PLacRNAP} \rightarrow \text{PLac} + \text{RNAP}$                    | 10                    |
| $R_3$    | $\text{PLacRNAP} \rightarrow \text{TrLacZ1}$                               | 1                     |
| $R_4$    | $\text{TrLacZ1} \rightarrow \text{RbsLacZ} + \text{PLac} + \text{TrLacZ2}$ | 1                     |
| $R_5$    | $\text{TrLacZ2} \rightarrow \text{TrLacY2}$                                | 0.015                 |
| $R_6$    | $\text{TrLacY1} \rightarrow \text{RbsLacY} + \text{TrLacY2}$               | 1                     |
| $R_7$    | $\text{TrLacY2} \rightarrow \text{RNAP}$                                   | 0.36                  |
| $R_8$    | $\text{Ribosome} + \text{RbsLacZ} \rightarrow \text{RbsRibosomeLacZ}$      | 0.17                  |
| $R_9$    | $\text{Ribosome} + \text{RbsLacY} \rightarrow \text{RbsRibosomeLacY}$      | 0.17                  |
| $R_{10}$ | $\text{RbsRibosomeLacZ} \rightarrow \text{Ribosome} + \text{RbsLacZ}$      | 0.45                  |
| $R_{11}$ | $\text{RbsRibosomeLacY} \rightarrow \text{Ribosome} + \text{RbsLacY}$      | 0.45                  |
| $R_{12}$ | $\text{RbsRibosomeLacZ} \rightarrow \text{TrRbsLacZ} + \text{RbsLacZ}$     | 0.4                   |
| $R_{13}$ | $\text{RbsRibosomeLacY} \rightarrow \text{TrRbsLacY} + \text{RbsLacY}$     | 0.4                   |
| $R_{14}$ | $\text{TrRbsLacZ} \rightarrow \text{LacZ}$                                 | 0.015                 |
| $R_{15}$ | $\text{TrRbsLacY} \rightarrow \text{LacY}$                                 | 0.036                 |
| $R_{16}$ | $\text{LacZ} \rightarrow \text{dgrLacZ}$                                   | $6.42 \times 10^{-5}$ |
| $R_{17}$ | $\text{LacY} \rightarrow \text{dgrLacY}$                                   | $6.42 \times 10^{-5}$ |
| $R_{18}$ | $\text{RbsLacZ} \rightarrow \text{dgrRbsLacZ}$                             | 0.3                   |
| $R_{19}$ | $\text{RbsLacY} \rightarrow \text{dgrRbsLacY}$                             | 0.3                   |
| $R_{20}$ | $\text{LacZ} + \text{lactose} \rightarrow \text{LacZlactose}$              | $9.52 \times 10^{-5}$ |
| $R_{21}$ | $\text{LacZlactose} \rightarrow \text{product} + \text{LacZ}$              | 431                   |
| $R_{22}$ | $\text{LacY} \rightarrow \text{lactose} + \text{LacY}$                     | 14                    |



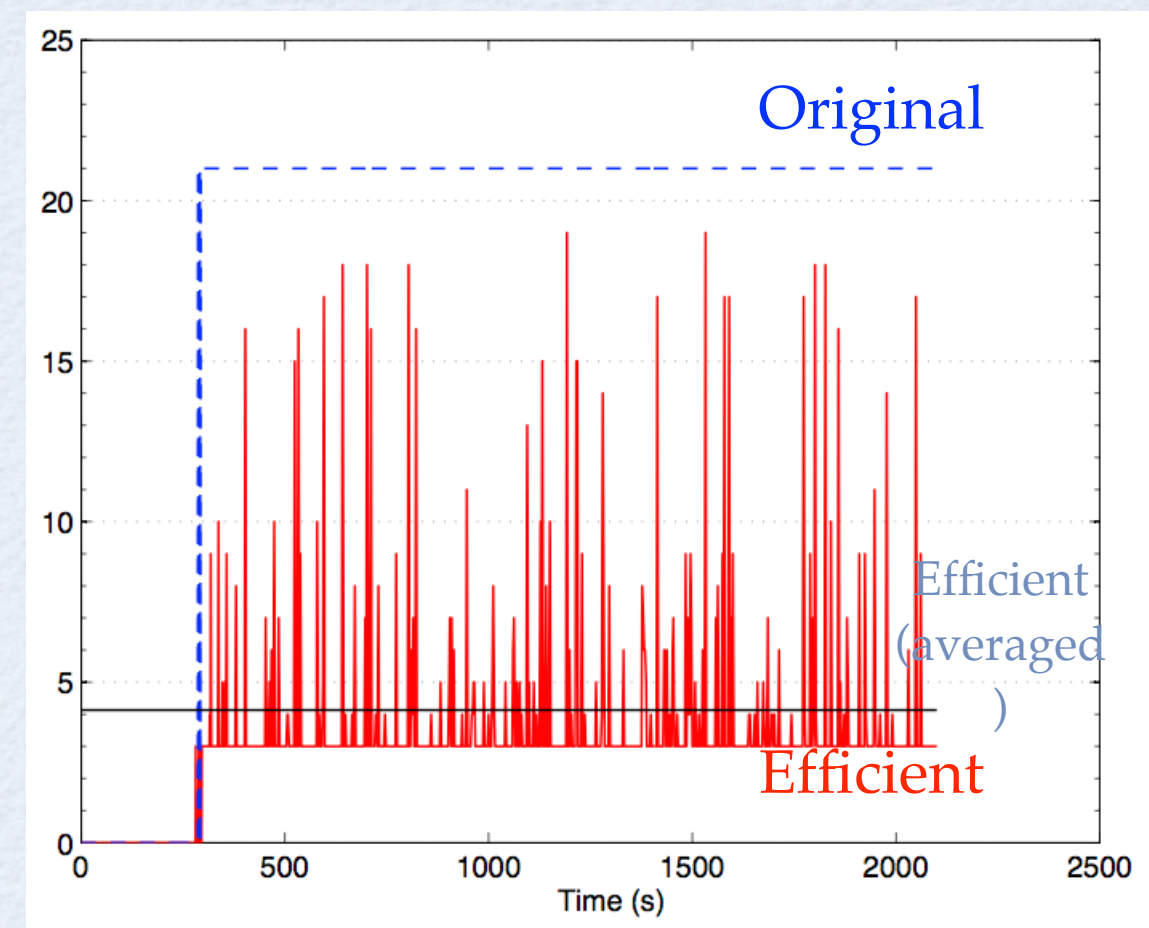
# R-leaping : Sampling the $M K_j$ efficiently

- $M$  can be large ( $\sim 10^2$ ) for bio-chemical systems!
- $M$  can be very large ( $\sim 10^6$ ) for diffusion
- Efficient sampling effectively loops over a fraction of  $M$ .

The larger the system, the bigger the payoff.

The more disparate the reaction rates are,  
the smaller the fraction.

Price to pay: carry out re-ordering often enough

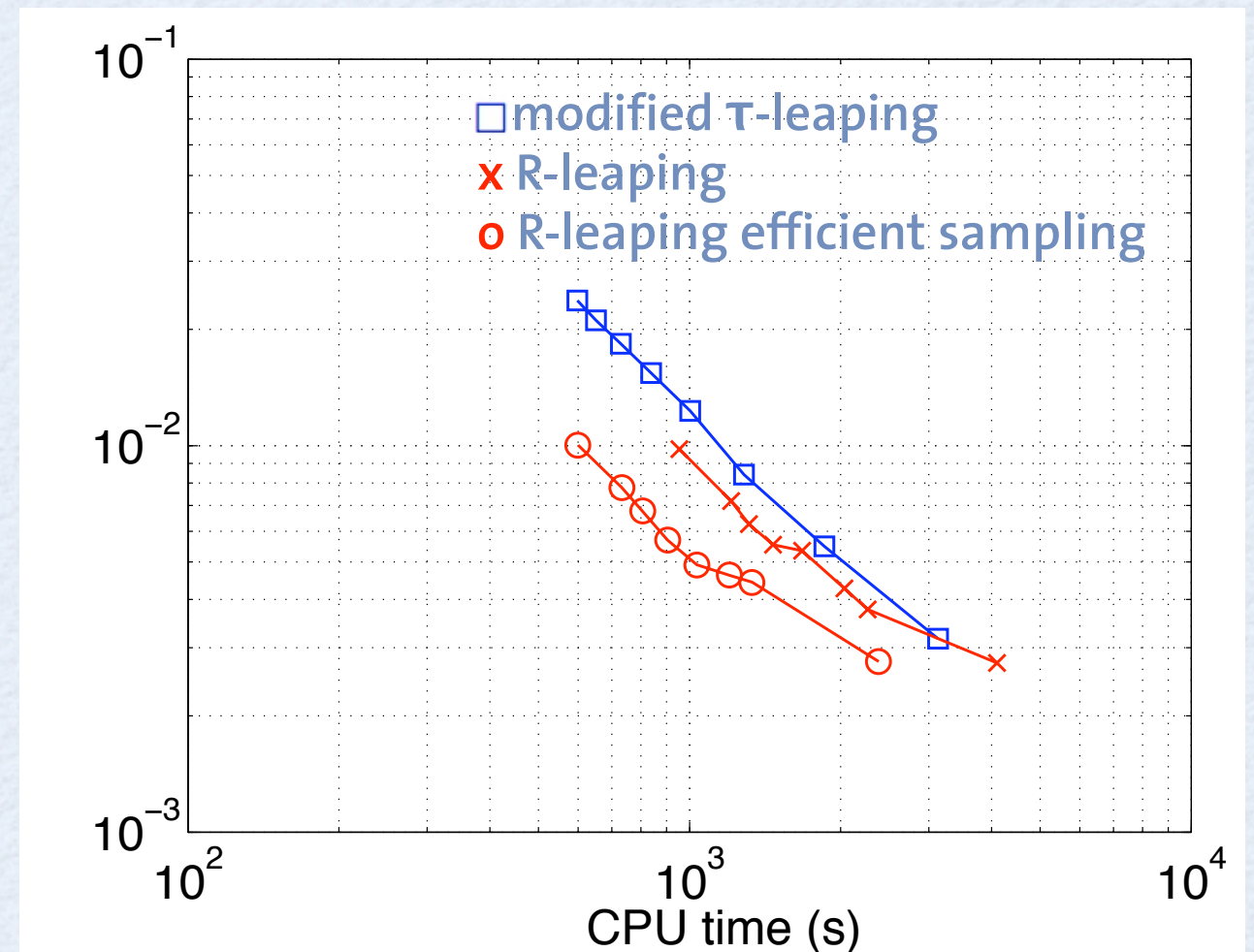
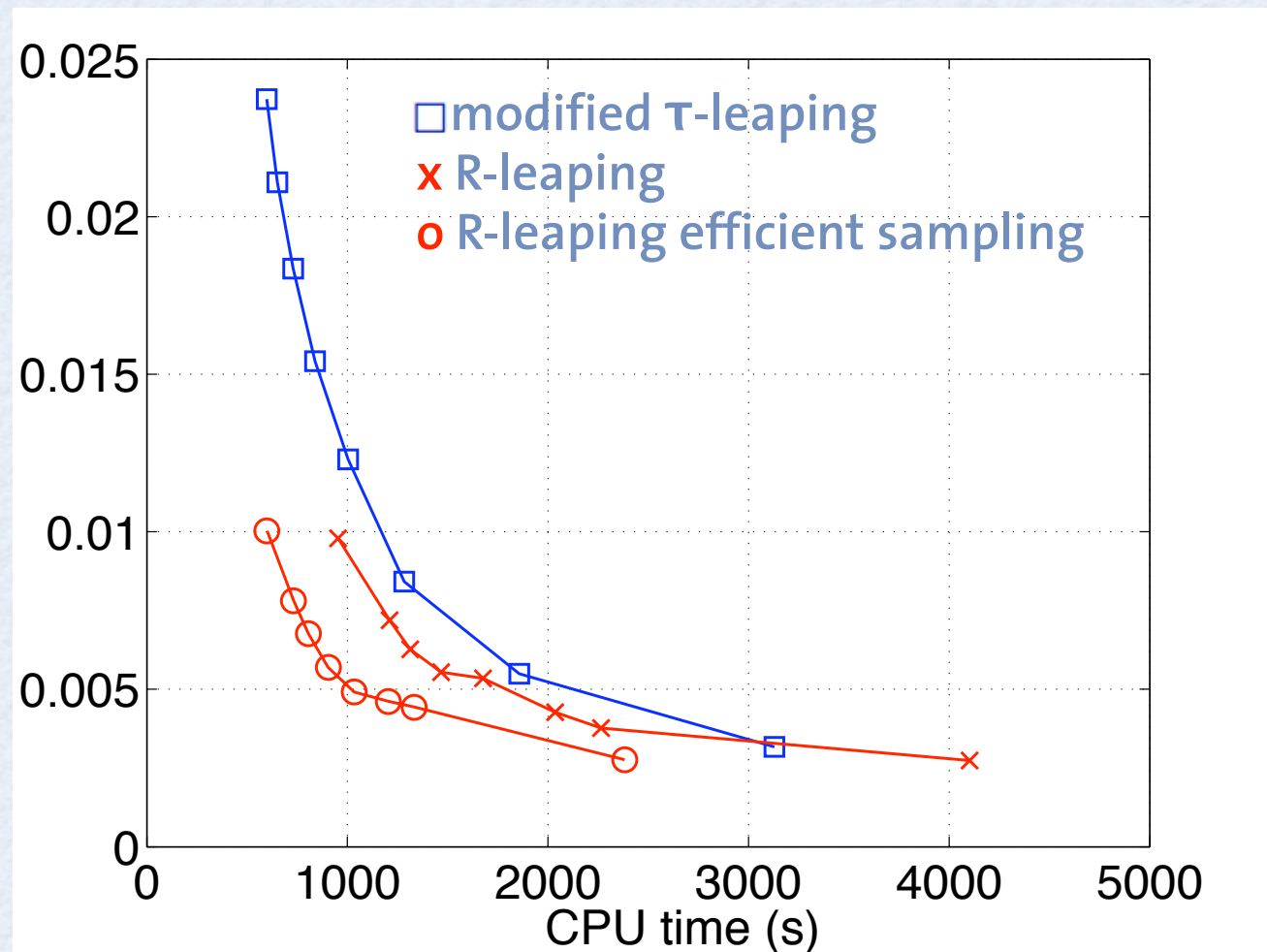


Number of binomial samples per time step  
LacY LacZ activities in E. Coli.,  $M=22$



# Histogram errors vs CPU time

- LacZ/LacY genes expression and enzymatic/transport activities of LacZ/LacY proteins in E. Coli



- $M = 22$  : R-leaping **2X** faster than modified  $\tau$ -leaping!



# Reaction-Diffusion : SSA + AMR

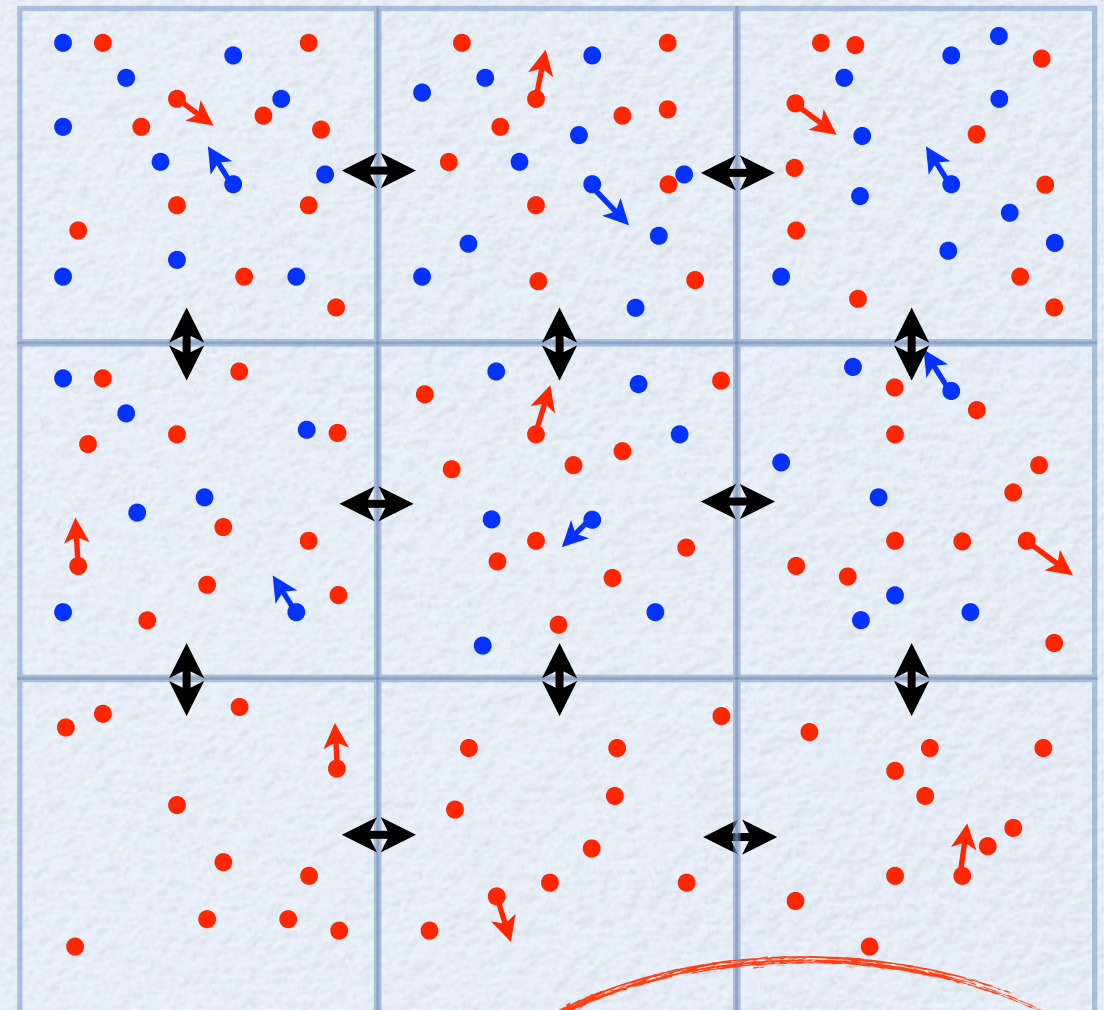
- Collisions and reactions **within each element**

- For homogeneity :

Kuramoto, Prog. Theor. Phys. 1974

$$\frac{\tau_R}{\tau_D} \gg 1$$

- $\tau_R$  mean free time for reactive collisions in a cell
- $\tau_D$  mean time during which a molecule will remain in element.



- For a bimolecular reaction with rate  $k$  and diffusion coefficient  $D$

$$\frac{\hat{\tau}_R}{\hat{\tau}_D} = \frac{D}{h^2 k}$$

**$h$  must be small for the discretization to be valid**



# AMR + STOCHASTIC

- i. PROPENSITIES FROM FV**
- ii. GRID REFINEMENT**
- iii. STOCHASTIC INTERPOLATION**
- iv. INTEGRATE: t/R Leaping**
- v. DATA STRUCTURES : OVERTURE**

B. Bayati, et.al. , Phys. Chem. Chem. Phys., 2008

Bayati B., et.al., J. of Computational Physics, 2011



# I. Propensities from FV schemes

D. Bernstein. Simulating mesoscopic reaction-diffusion systems using the Gillespie algorithm. *Phys. Rev. E*, 2005.

$\mathbf{J} \left( x, y + \frac{2}{3}\delta y \right)$  approximated by three Taylor series:

around the point  $\left( x, y + \frac{2}{3}\delta y \right)$

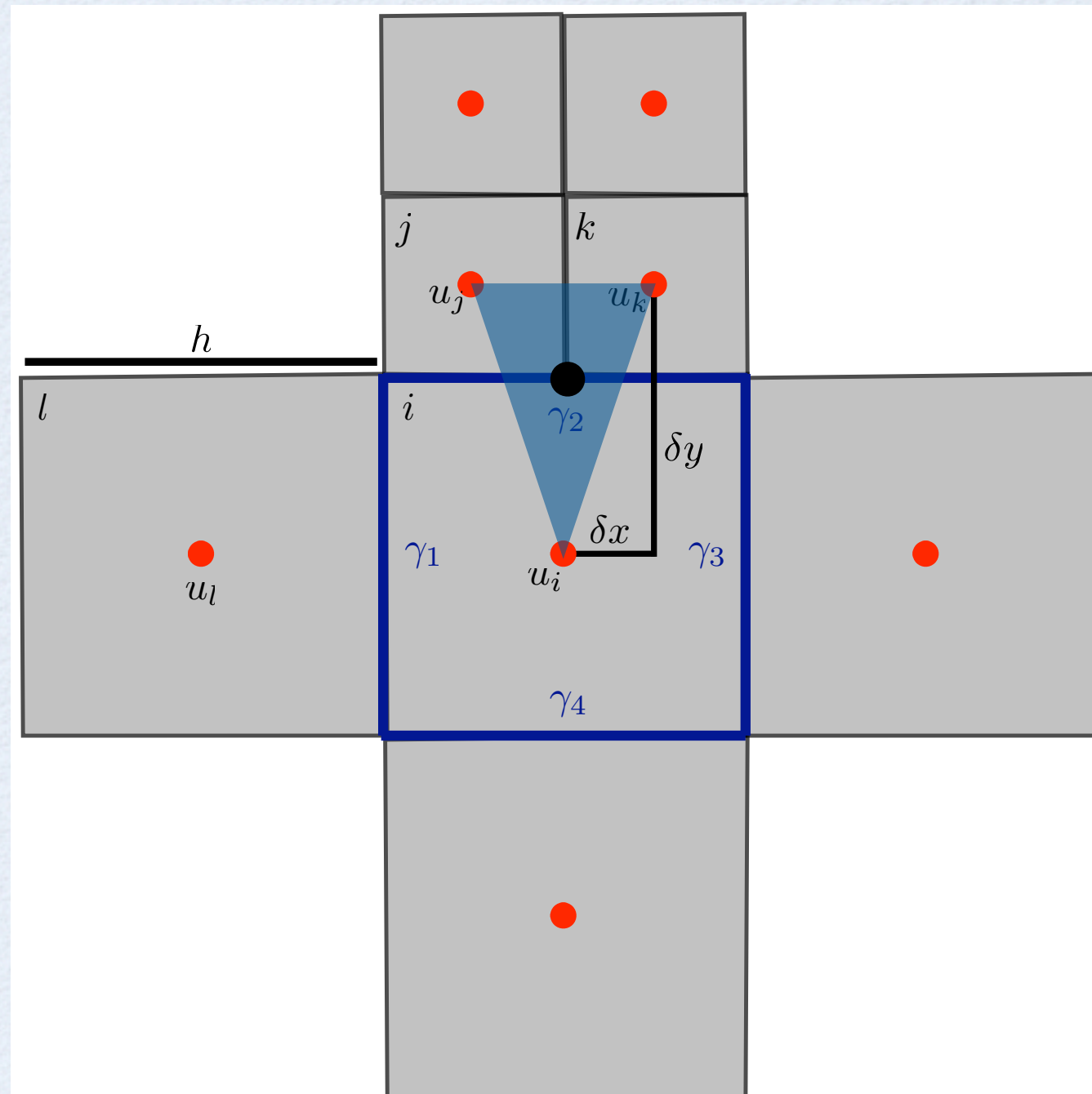
$$\mathbf{J} \left( x, y + \frac{2}{3}\delta y \right) = -\frac{D}{2\delta y} \left( \bar{u}_j^{(s)} + \bar{u}_k^{(s)} - 2\bar{u}_i^{(s)} \right) + \mathcal{O}(h)$$

$$\frac{dU_i^{(s)}}{dt} = \frac{4D}{3h^2} \left( 2(U_j^{(s)} + U_k^{(s)}) - U_i^{(s)} \right) + \mathcal{O}(h).$$

propensities for transitions between  $j$  &  $i$

$$a_{i,j}^D = \frac{4D}{3h^2} (U_j^{(s)} + U_k^{(s)}),$$

$$a_{j,i}^D = \frac{2D}{3h^2} U_i^{(s)}$$

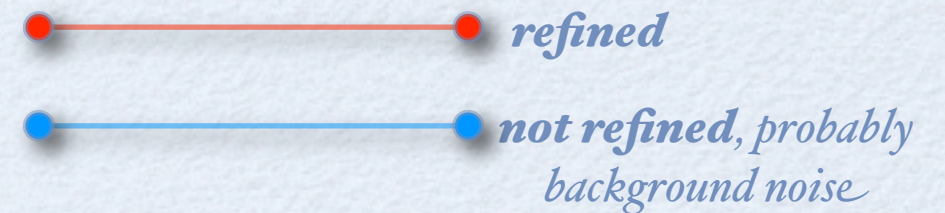




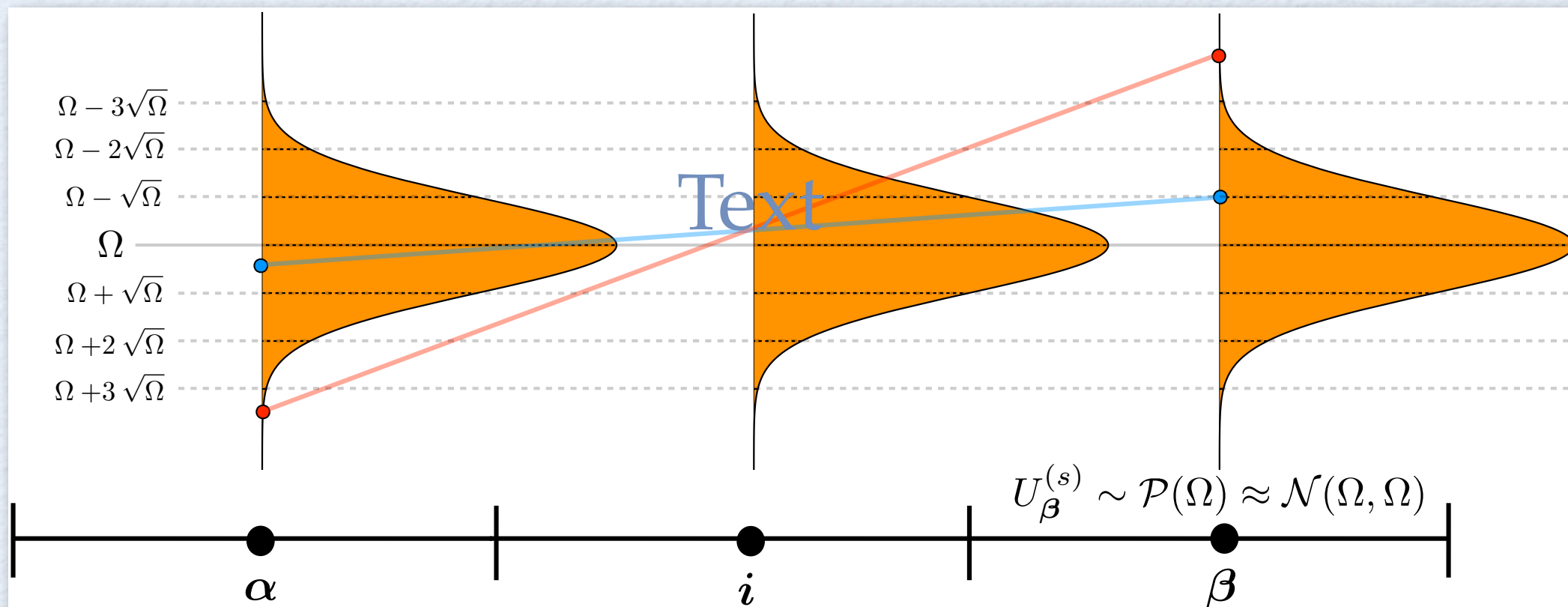
# II. Refinement Criteria

Bell et al., J. Comp. Phys., 2007

Distinguish between gradients and fluctuations,



In equilibrium:



Refine volume element  $i$  if:

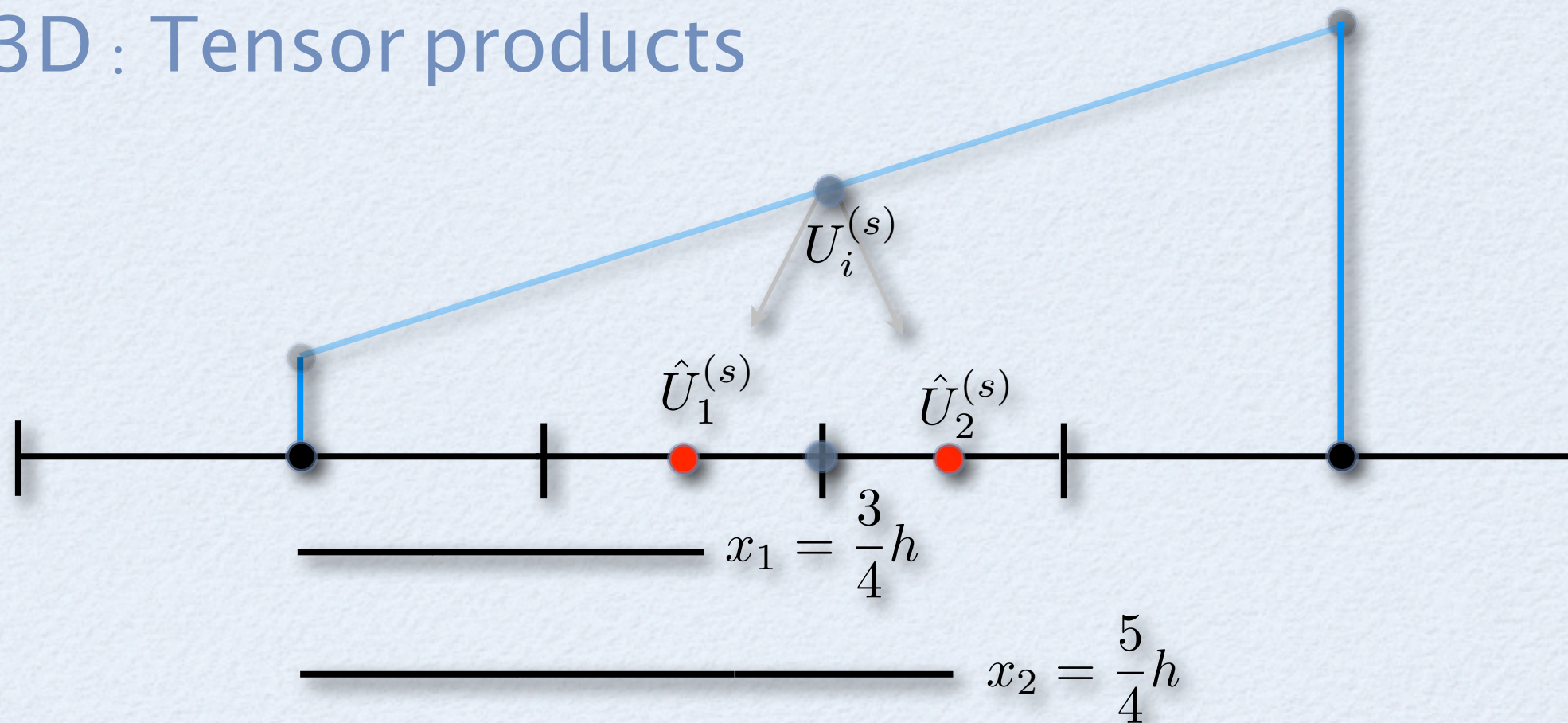
$$|U_{i+1}^{(s)} - U_{i-1}^{(s)}| > 2C\sqrt{\Omega}$$



# III. Stochastic Interpolation

Conservative + Strictly Positive

2D, 3D : Tensor products



**weights:**  $w_1 := \frac{mx_1 + a}{m(x_1 + x_2) + 2a}$

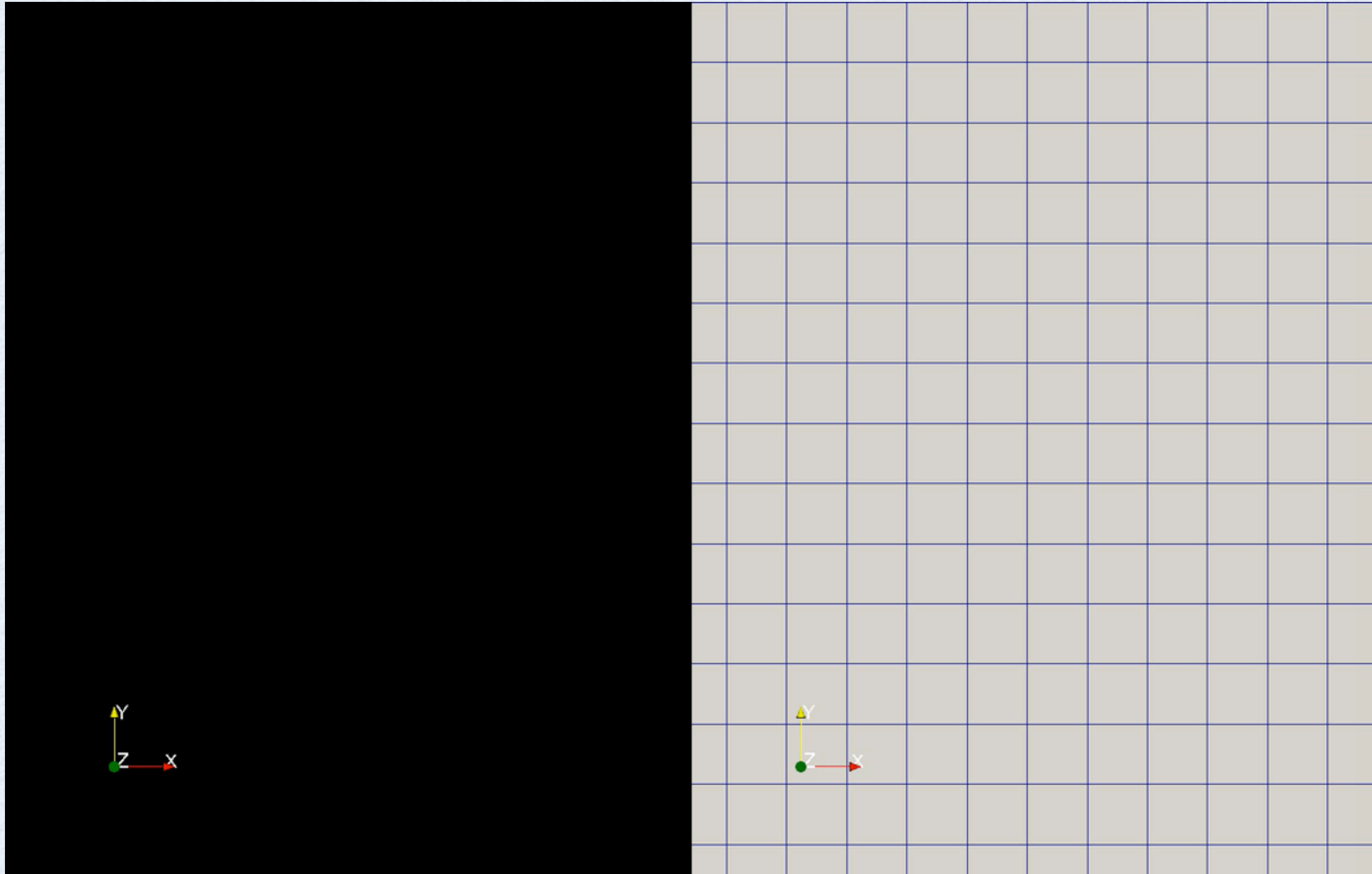
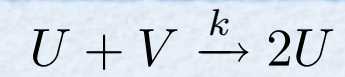
$$w_2 := \frac{mx_2 + a}{m(x_1 + x_2) + 2a}$$

**sampling:**  $\hat{U}_1^{(s)} \sim \mathcal{B}(U_i^{(s)}, w_1)$

$$\hat{U}_2^{(s)} = U_i^{(s)} - \hat{U}_1^{(s)}$$



# Fisher-Kolmogorov Reaction-Diffusion System in 2-D



Fisher,  
Ann. Eugenics 1937

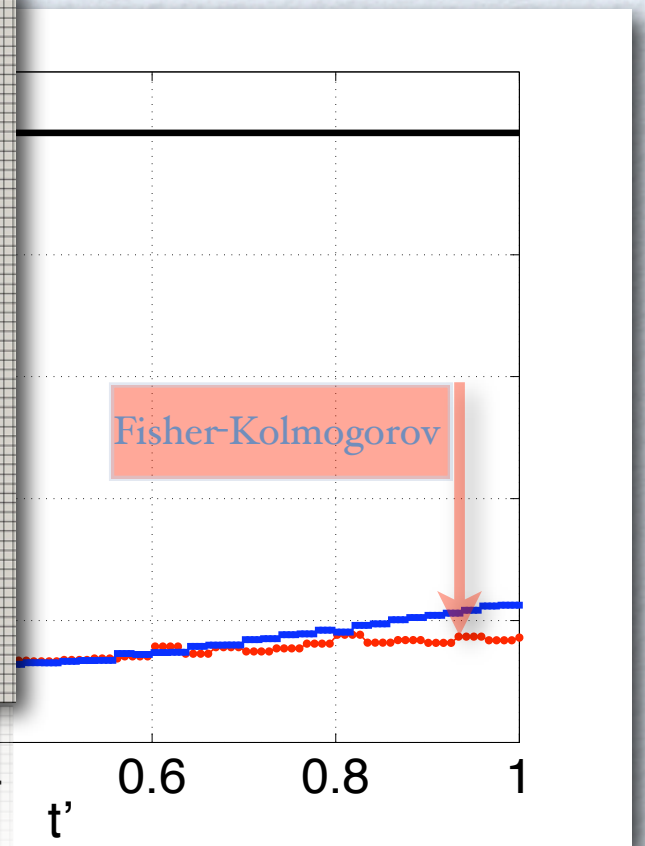
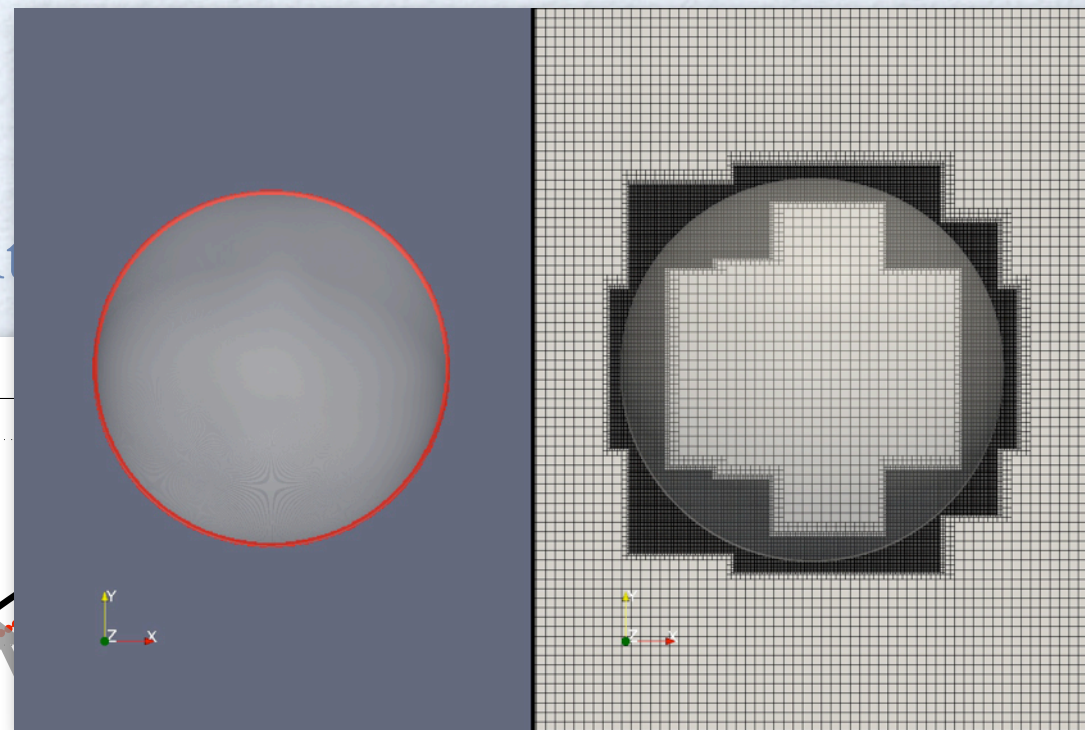
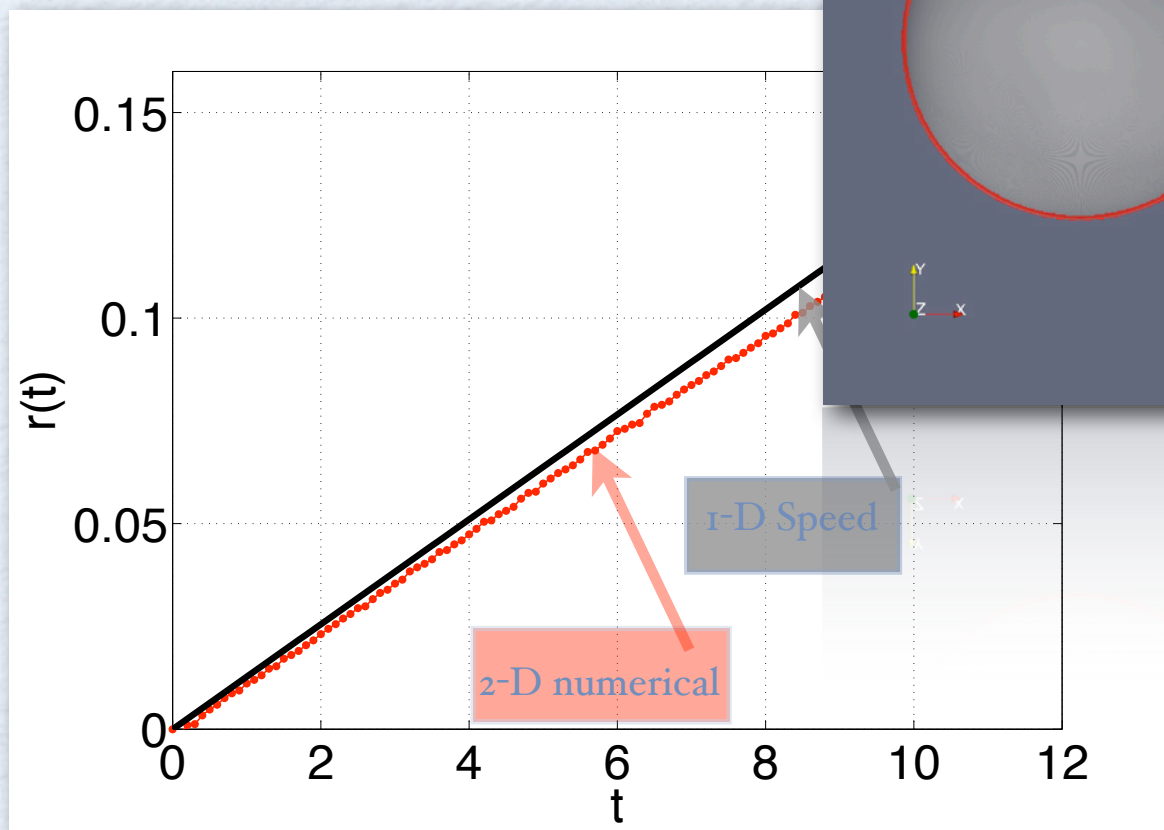


# Wavespeed & Efficiency

Ratio of Cells vs Time

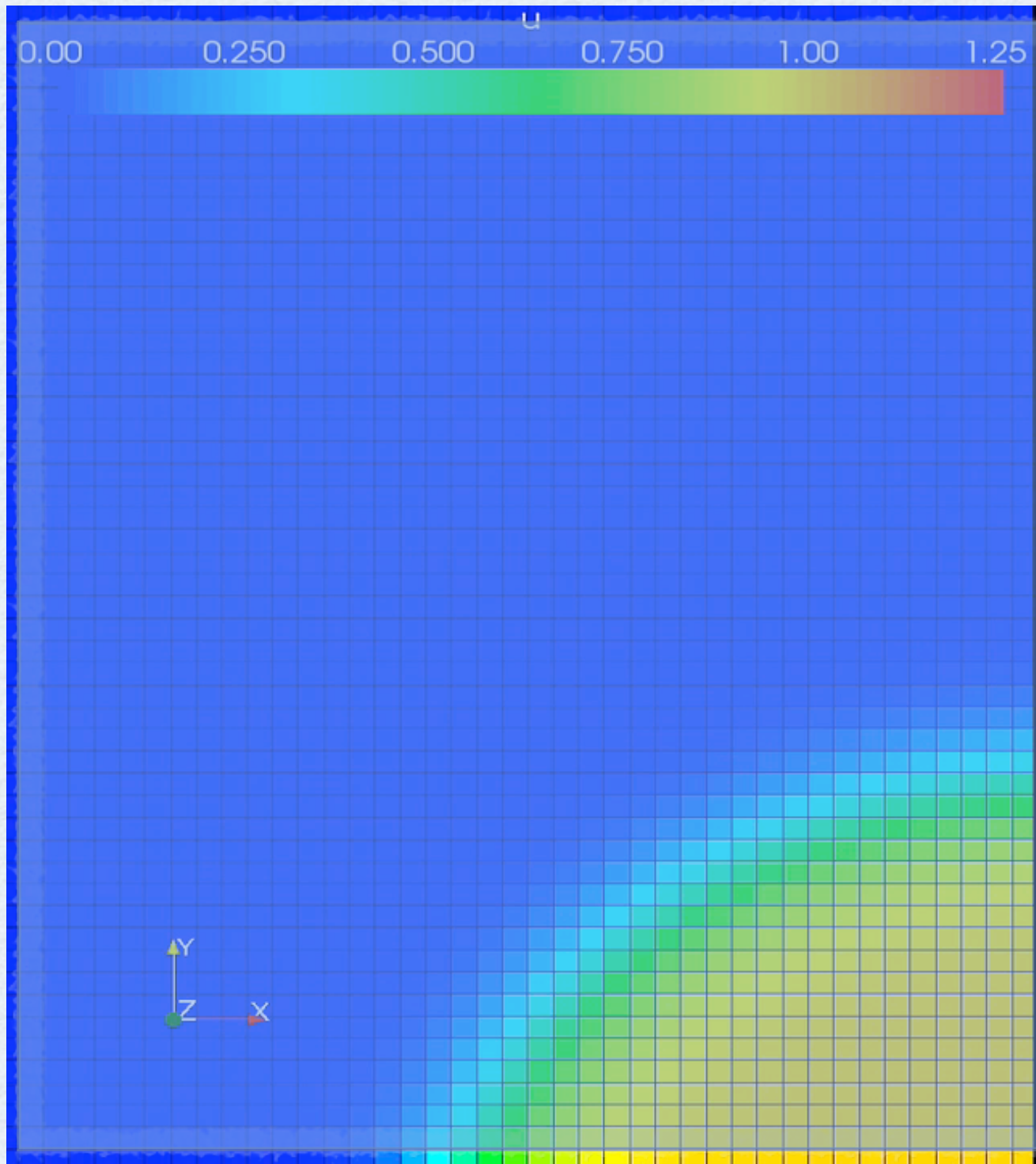
$$E(t') = \frac{L_{AMR}(t')}{L_{h_{min}}}$$

Fisher: Wavefront

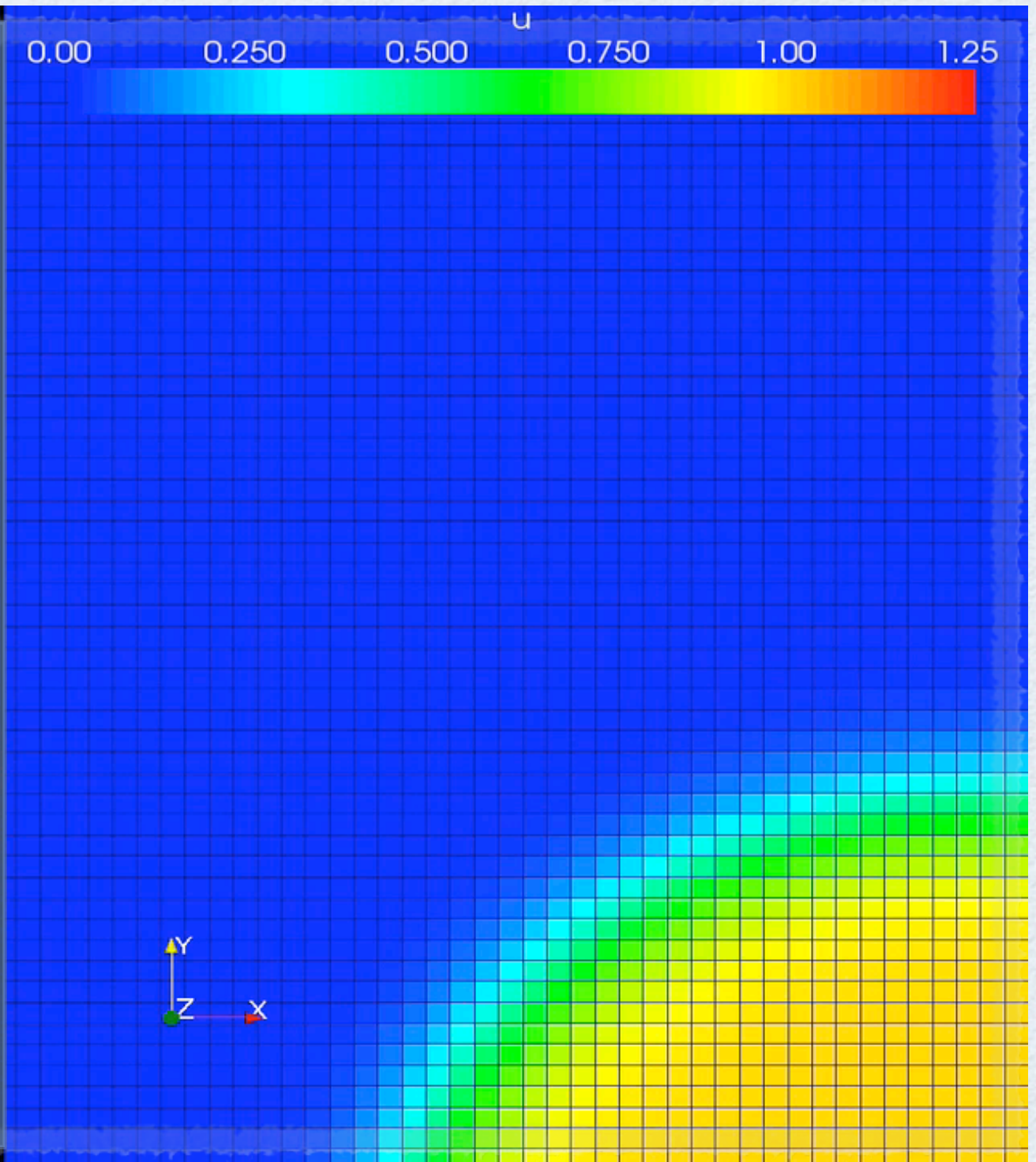


Bayati B., et al., J. of Comp. Phys., 2011





**STOCHASTIC**

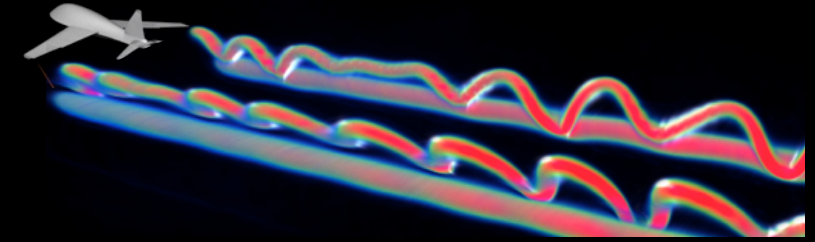


**DETERMINISTIC**



# 16384 Cores - 10 Billion Particles - 60% efficiency

Runs at IBM Watson Center - BLue Gene/L

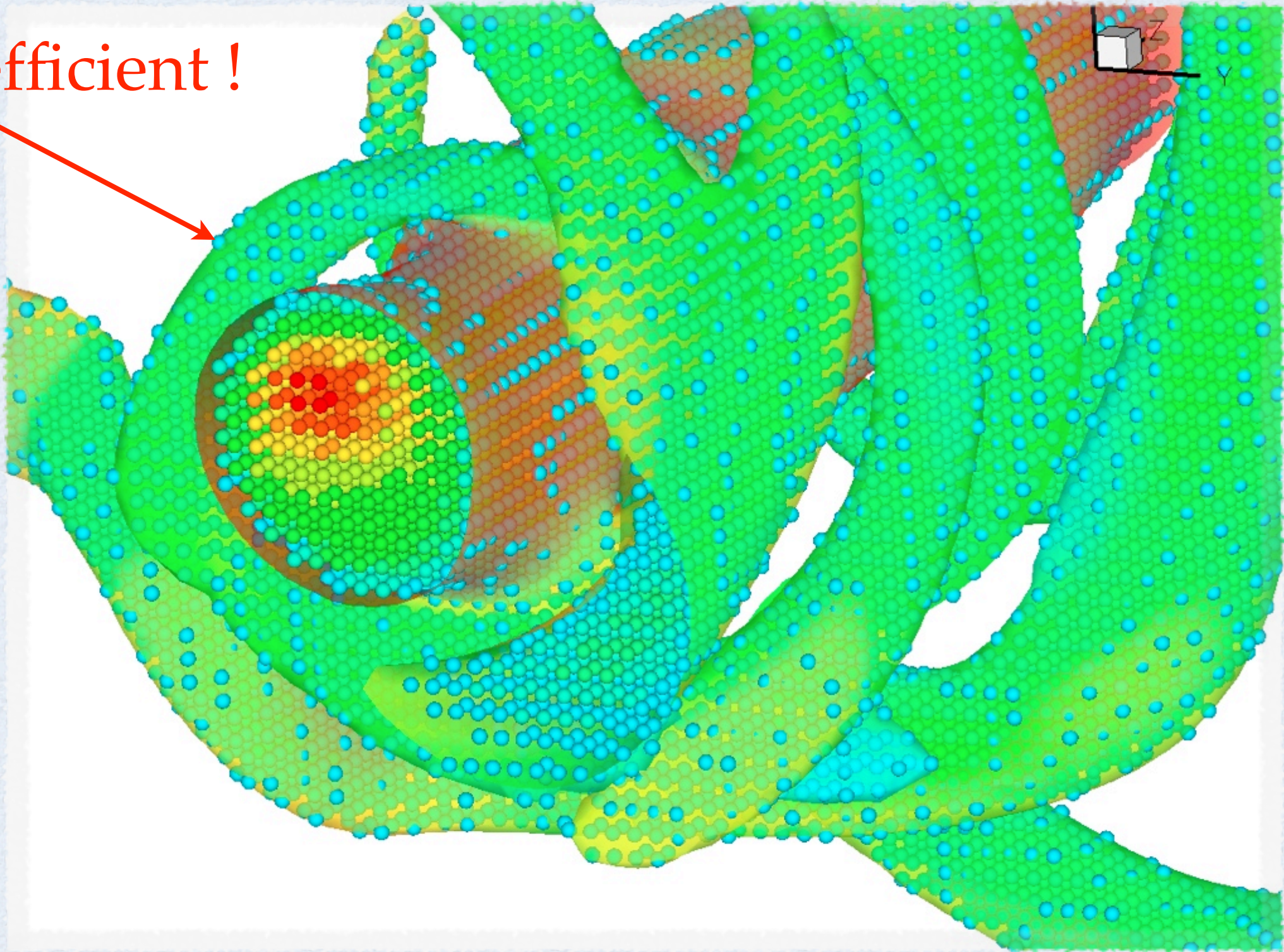


Chatelain P., Curioni A., Bergdorf M., Rossinelli D., Andreoni W., Koumoutsakos P., Billion Vortex Particle Direct Numerical Simulations of Aircraft Wakes, Computer Methods in Applied Mech. and Eng. 197/13-16, 1296-1304, 2008



# PARTICLES ARE ADAPTIVE

yet inefficient !





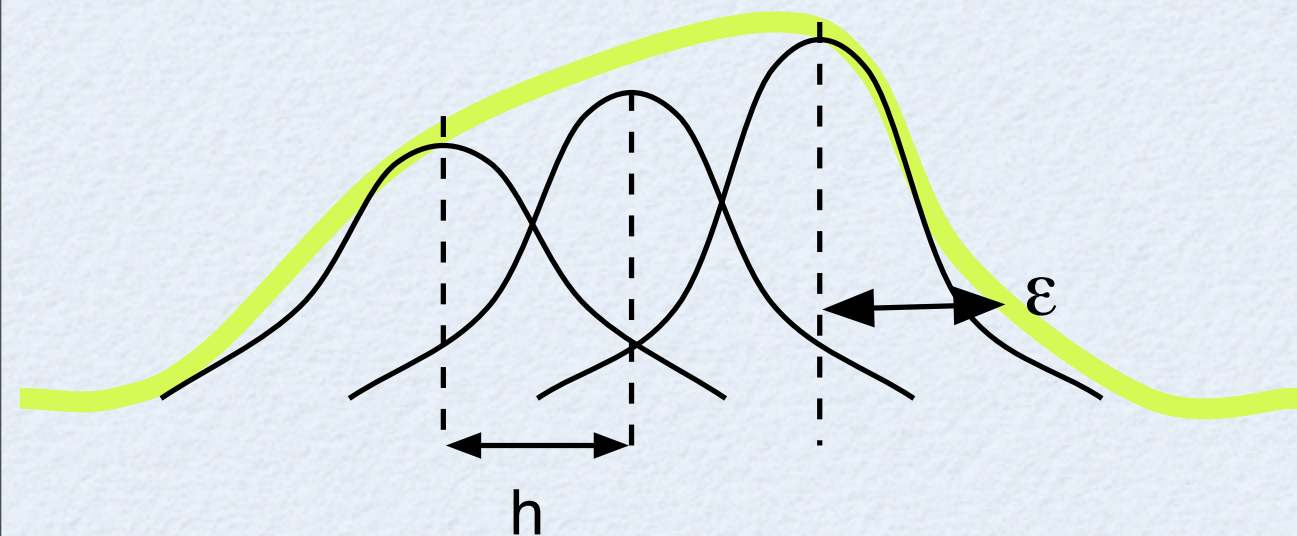
# FUNCTIONS and PARTICLES

## Integral Function Representation

$$\Phi(x) = \int \Phi(y) \delta(x - y) dy$$

## Function Mollification

$$\Phi_\epsilon(x) = \int \Phi(y) \zeta_\epsilon(x - y) dy$$



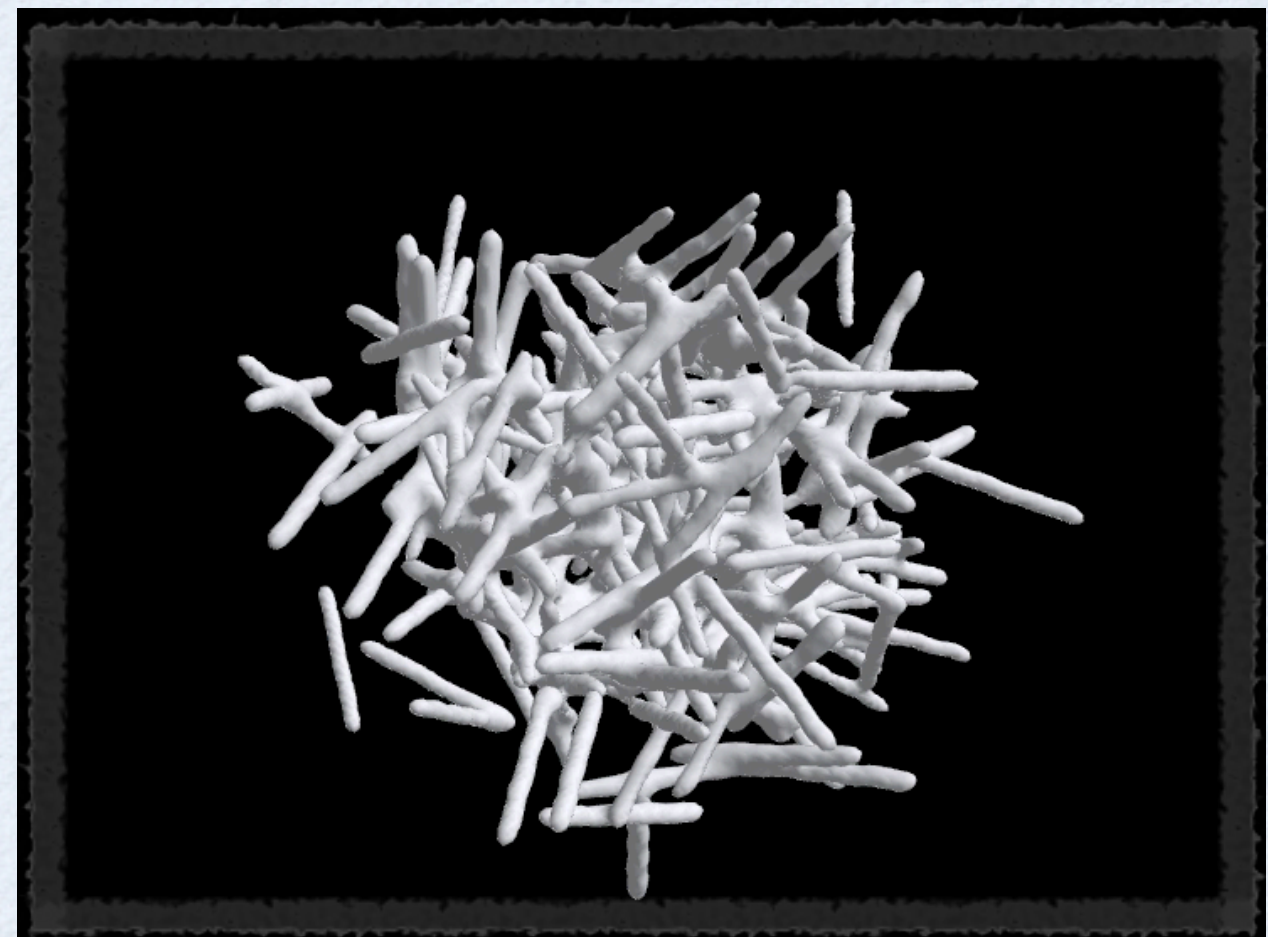
Particles are “mesh” free

## Point Particle Quadrature

$$\Phi^h(x, t) = \sum_{p=1}^{N_p} h_p^d \Phi_p(t) \delta(x - x_p(t))$$

## Smooth Particle Quadrature

$$\Phi_\epsilon^h(x, t) = \sum_{p=1}^{N_p} h_p^d \Phi_p(t) \zeta_\epsilon(x - x_p(t))$$

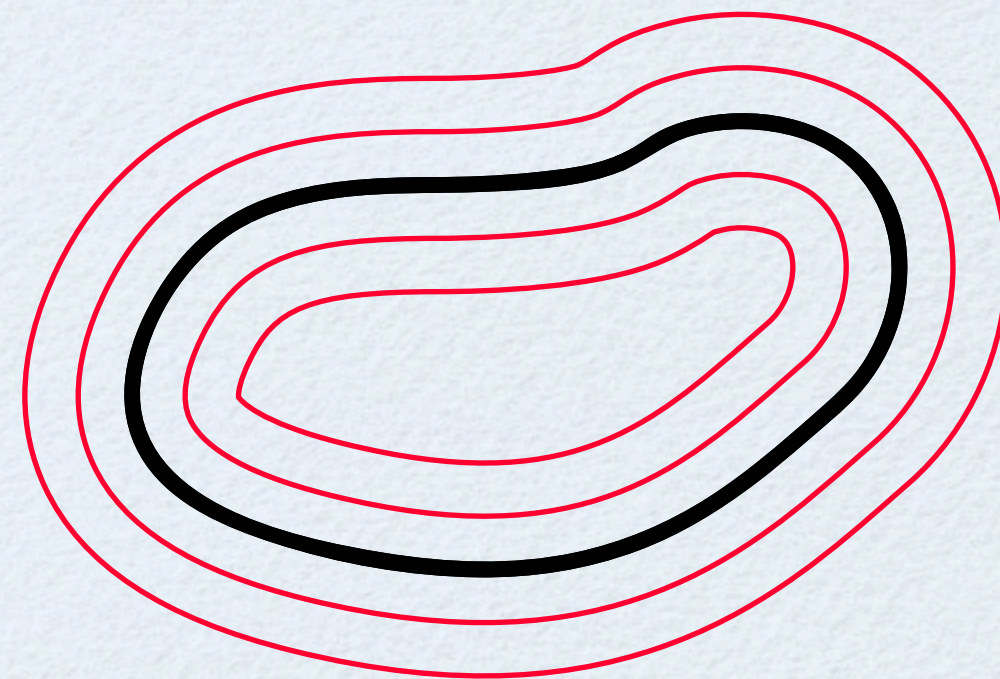




# SURFACES AS LEVEL SETS

$$\Gamma(t) = \{ \mathbf{x} \in \Omega \mid \phi(\mathbf{x}, t) = 0 \}$$

$$|\nabla \phi| = 1$$



## EVOLVING THE LEVEL SETS

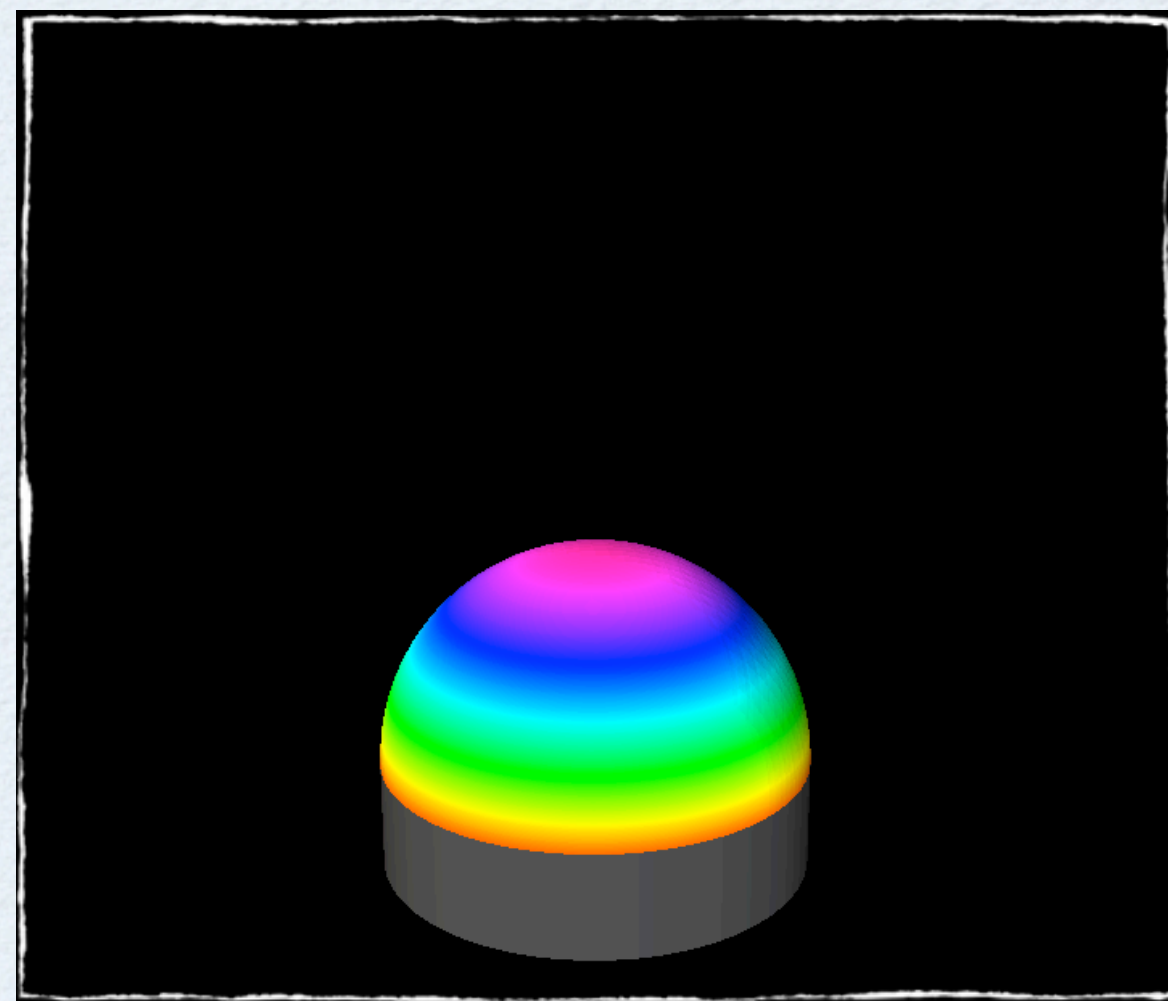
$$\frac{\partial \Phi}{\partial t} + \mathbf{u} \cdot \nabla \Phi = 0$$

## PARTICLE APPROXIMATION

$$\Phi_\epsilon^h(x, t) = \sum_{p=1}^{N_p} h_p^d \Phi_p(t) \zeta_\epsilon(x - x_p(t))$$

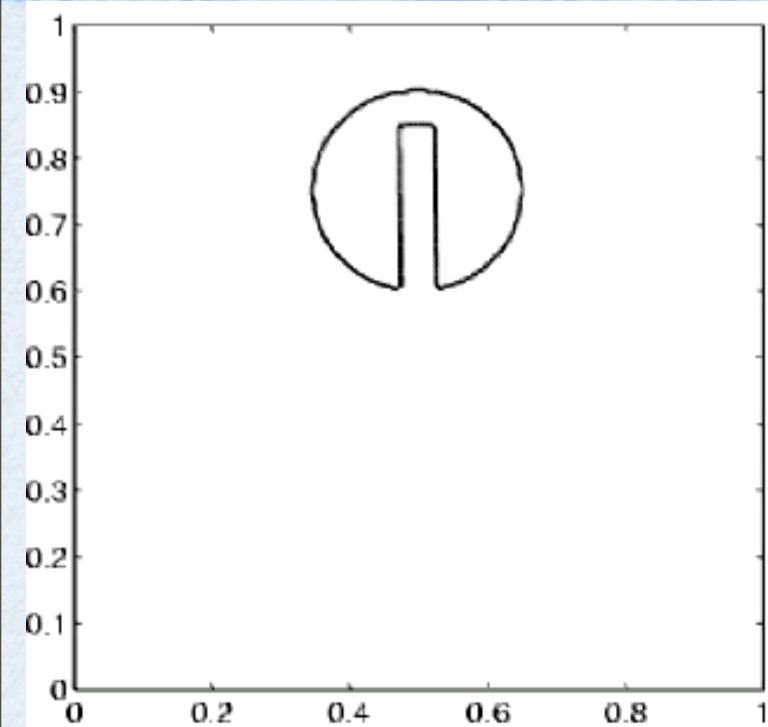
### Lagrangian Surface Transport

$$\frac{dx_p}{dt} = \mathbf{u}_p \quad \frac{D\Phi_p}{Dt} = 0$$





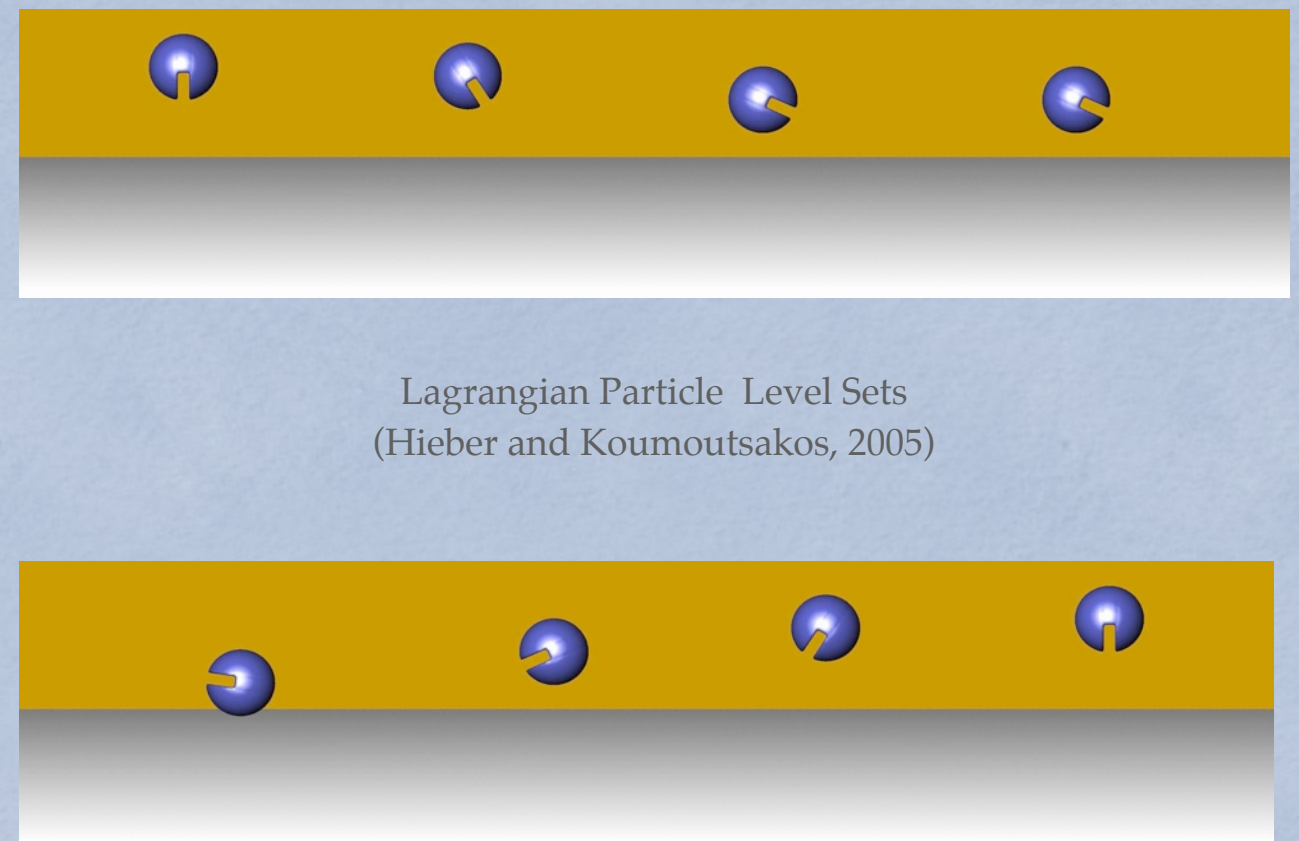
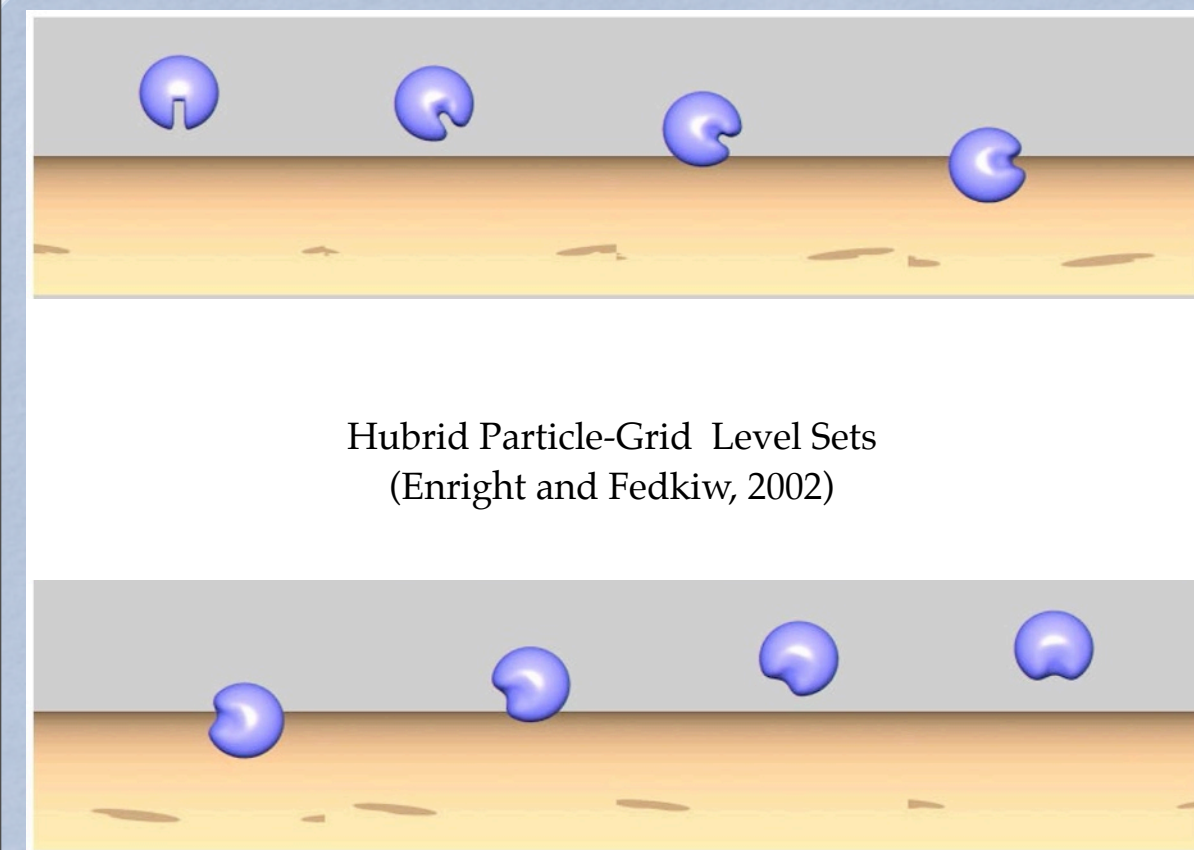
# Lagrangian vs Eulerian Descriptions



- **PARTICLE LEVEL SETS** exact for rigid body motion

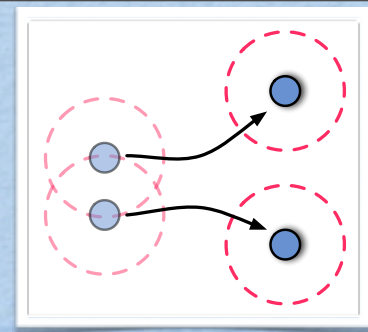
$$\Phi(\mathbf{x}, t) = \Phi_0(\mathbf{x} - \mathbf{u}t)$$

Lagrangian Particle methods  
good for **linear** advection





# LAGRANGIAN DISTORTION



- loss of **overlap** -> loss of **convergence**

Particles follow flow trajectories - **Location distortion**

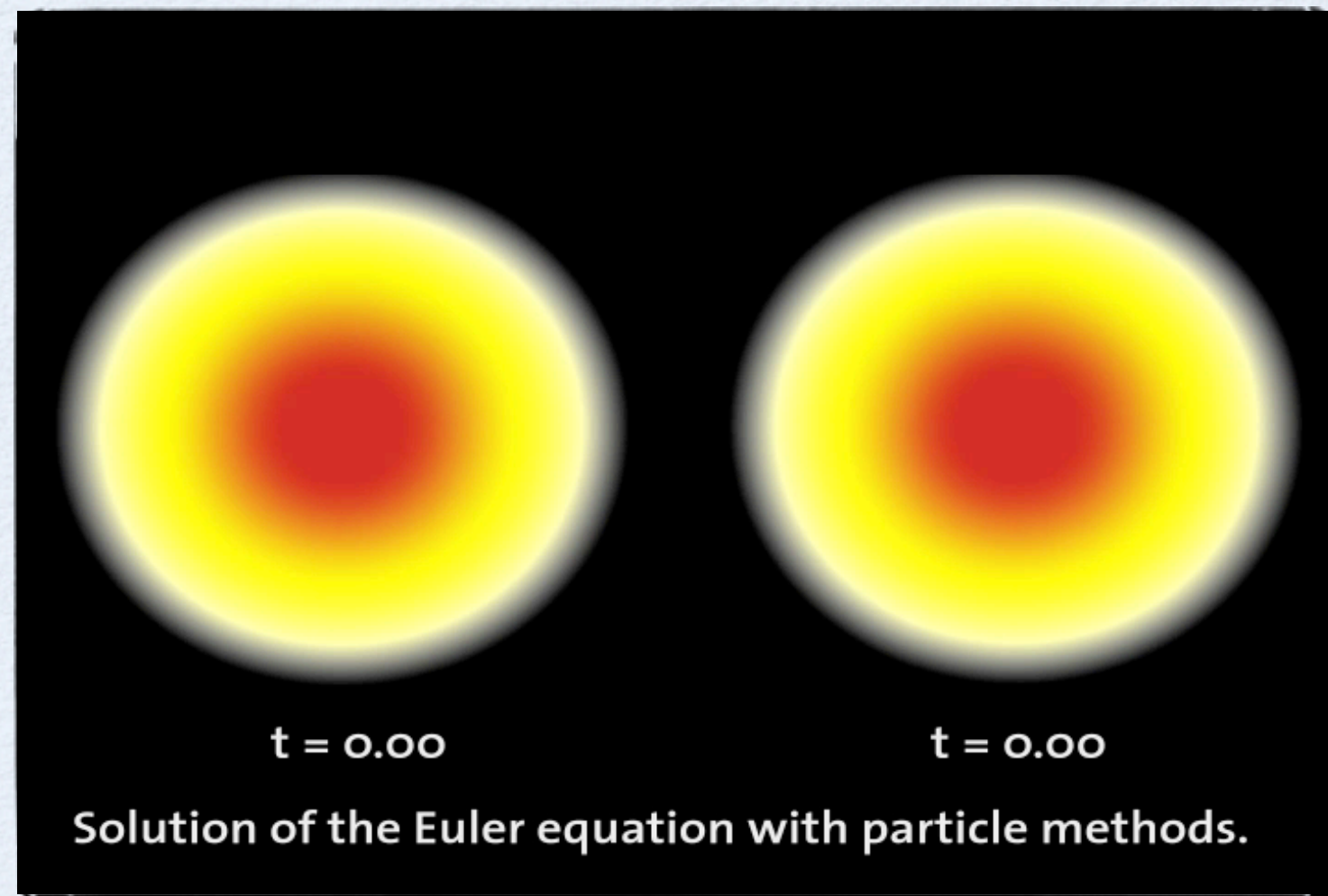
## EXAMPLE :

Incompressible 2D Euler Equations

$$\omega = \nabla \times \mathbf{u} \quad \nabla \cdot \mathbf{u} = 0$$

$$\frac{D\omega}{Dt} = 0$$

There is an **exact** axisymmetric solution





# SMOOTH PARTICLES MUST OVERLAP

## Integral Function Representation

$$\Phi(x) = \int \Phi(y) \delta(x - y) dy$$

## Function Mollification

$$\Phi_\epsilon(x) = \int \Phi(y) \zeta_\epsilon(x - y) dy$$

$$\int \zeta x^\alpha dx = 0^\alpha \quad 0 \leq \alpha < r$$

## TOTAL ERROR

$$\begin{aligned} \|\Phi - \Phi_\epsilon^h\| &\leq \|\Phi - \Phi_\epsilon\| + \|\Phi_\epsilon - \Phi_\epsilon^h\| \\ &\leq (C_1 \epsilon^r) + C_2 \left(\frac{h}{\epsilon}\right)^m \|\Phi\|_\infty \end{aligned}$$

## Point Particle Quadrature

$$\Phi^h(x, t) = \sum_{p=1}^{N_p} h_p^d \Phi_p(t) \delta(x - x_p(t))$$

## Smooth Particle Quadrature

$$\Phi_\epsilon^h(x, t) = \sum_{p=1}^{N_p} h_p^d \Phi_p(t) \zeta_\epsilon(x - x_p(t))$$

Need  $h/\epsilon < 1$  for accuracy

**PARTICLES MUST OVERLAP**



# Are Particle Methods Grid Free ?

## How to fix it ?

- Modify the smoothing kernels (SPH - Monaghan)
- Re-distribute particles with Voronoi Meshes (ALE - Russo)
- Re-initialise particle strengths (WRKPM - Liu, Belytchko)

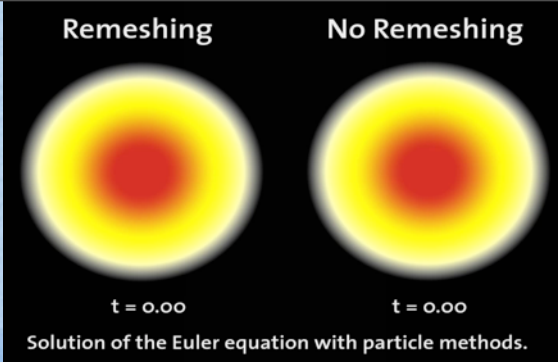
DOES NOT WORK  
EXPENSIVE - UNSTABLE  
EXPENSIVE

**REMESHING** : Re-project particles on a mesh

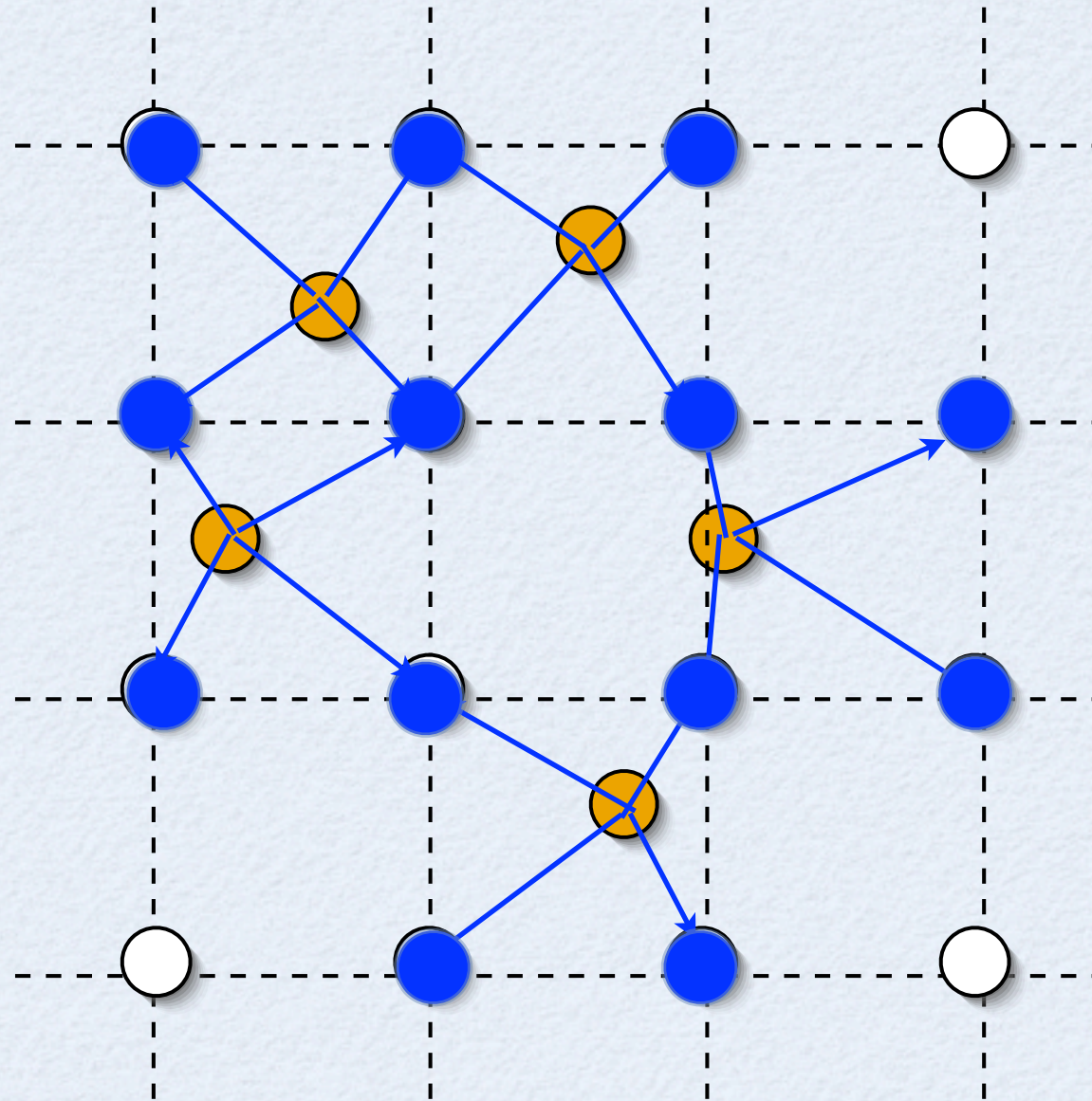
- NO MESH-FREE particle methods
- Can use all the “tricks” of mesh based methods
- High CFL
- Multiresolution & Multiscaling
- .....



# Particle Remeshing



Koumoutsakos, JCP, 1997



**Moment Conserving Interpolation :**  $Q_p^{\text{new}} = \sum_{p'} Q_{p'} M(j h - x_{p'})$



# REMESHED PARTICLE METHODS

1. ADVECT : Particles -> Large CFL

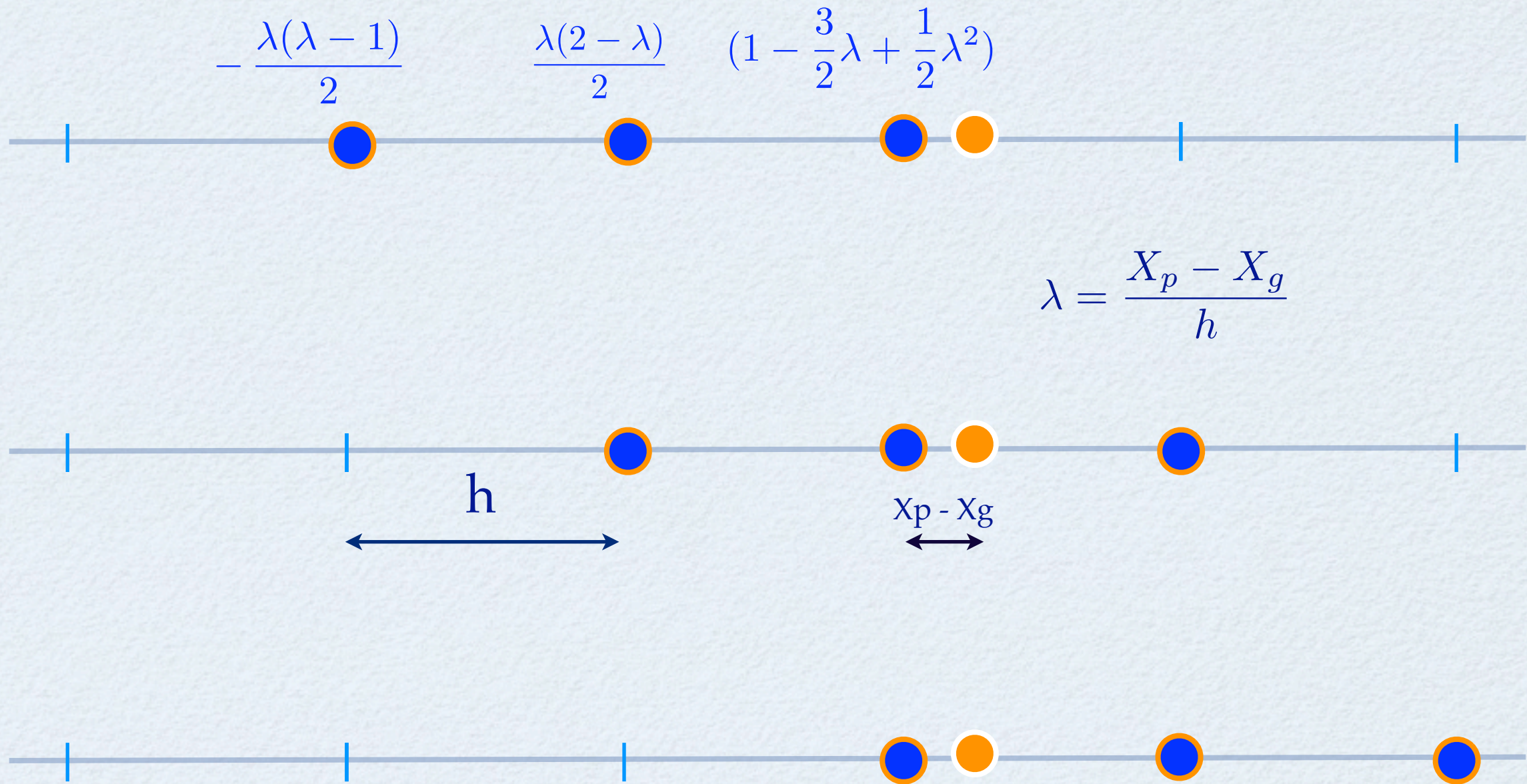
2. REMESH : Particles to Mesh -> Gather/Scatter

3. SOLVE : Poisson/Derivatives on Mesh -> FFTw/Ghosts

4. RESAMPLE : Mesh Nodes BECOME Particles



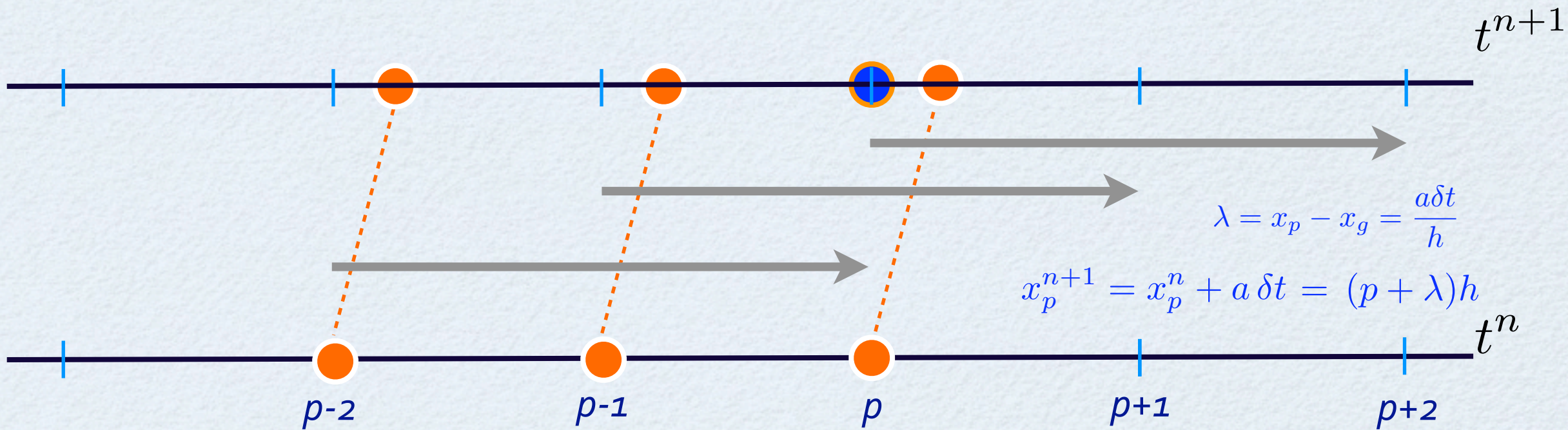
# Remeshing Stencils



Bergdorf et. al., MMS,2005  
**Cottet et.al., CRAS, 2008**



$$u_p^{n+1} = -\frac{\lambda(\lambda-1)}{2}u_{p-2}^n + \frac{\lambda(2-\lambda)}{2}u_{p-1}^n + \left(1 - \frac{3}{2}\lambda + \frac{1}{2}\lambda^2\right)u_p^n$$



$$\frac{\partial u}{\partial t} + a \frac{\partial u}{\partial x} = 0$$

$$u_p = u(x_p)h$$

$$\begin{aligned} \frac{du_p}{dt} &= 0 \\ \frac{dx_p}{dt} &= a \end{aligned} \quad + \text{REMESH}$$

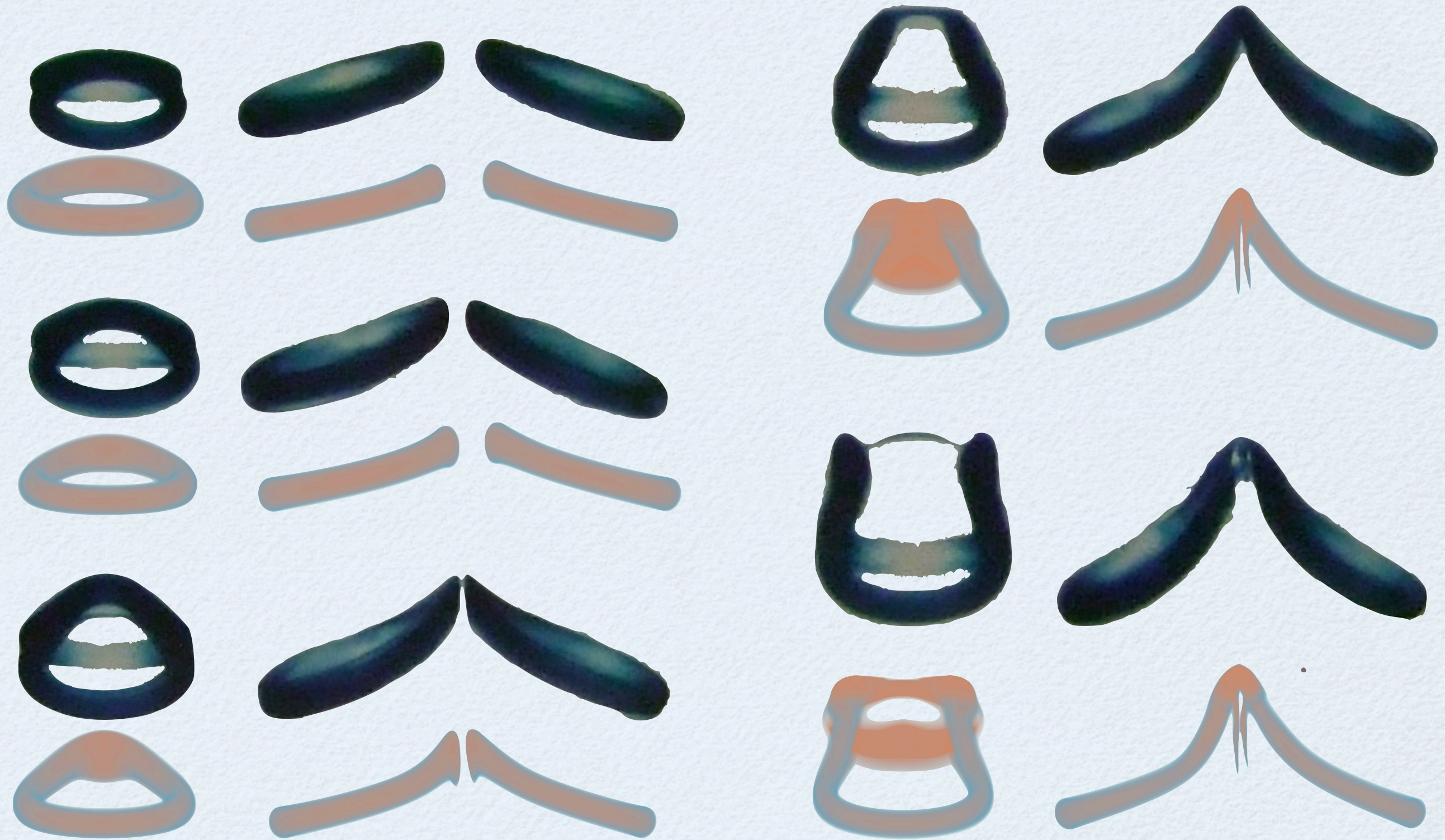
$$u_p^{n+1} = u_p^n - \frac{\lambda}{2}(3u_p^n - 4u_{p-1}^n + 4u_{p-2}^n) + \frac{\lambda^2}{2}(u_p^n - 2u_{p-1}^n + u_{p-2}^n)$$

**Euler Advect + One-sided Remesh = Beam-Warming FD**

**Euler Advect + Central Remesh = Lax - Wendroff FD .....**



# VORTEX RING COLLISION, $Re = 1800$



Experiments : P. Schatzle & D. Coles (1986)

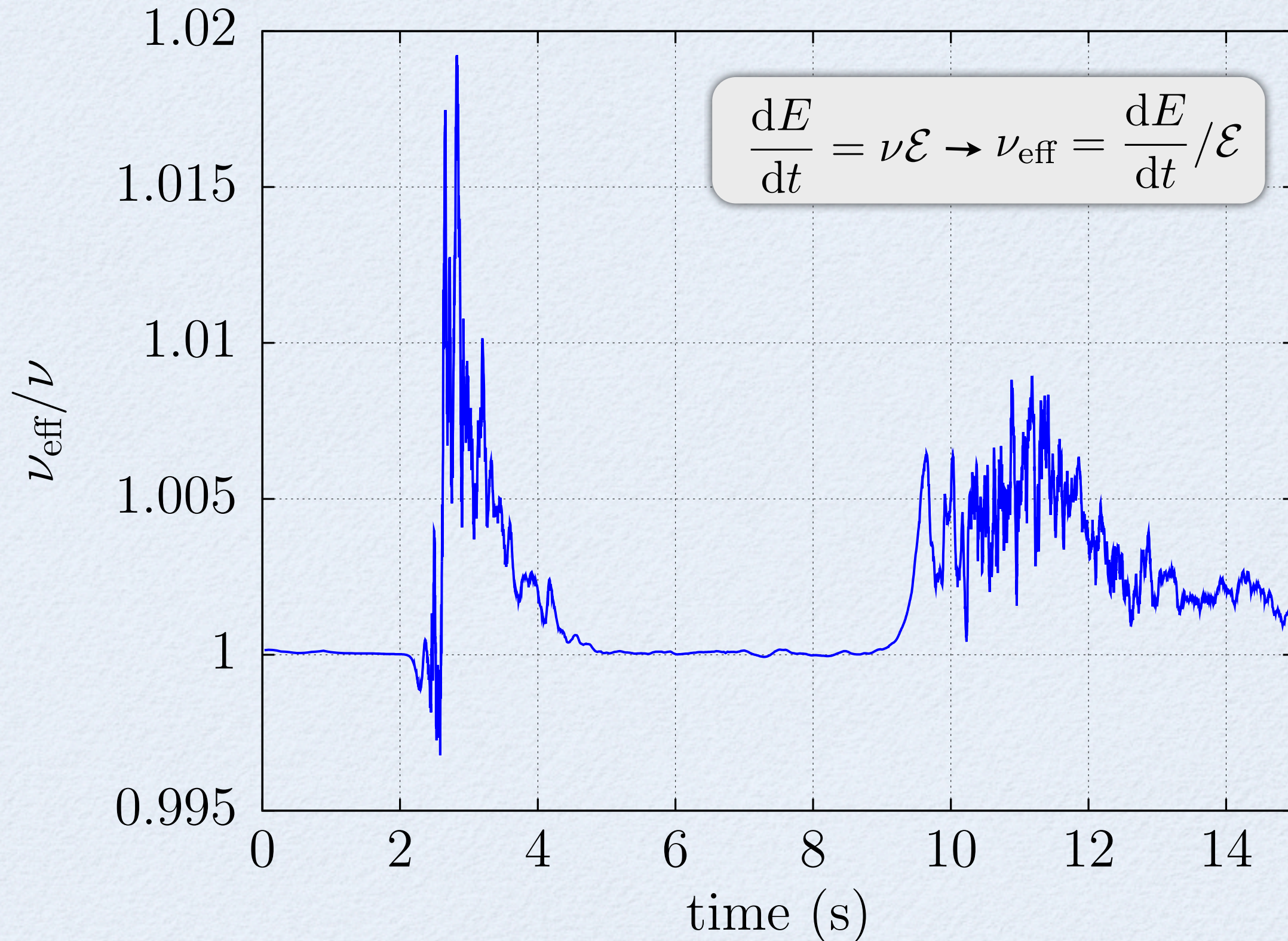


# VORTEX DYNAMICS at High Re



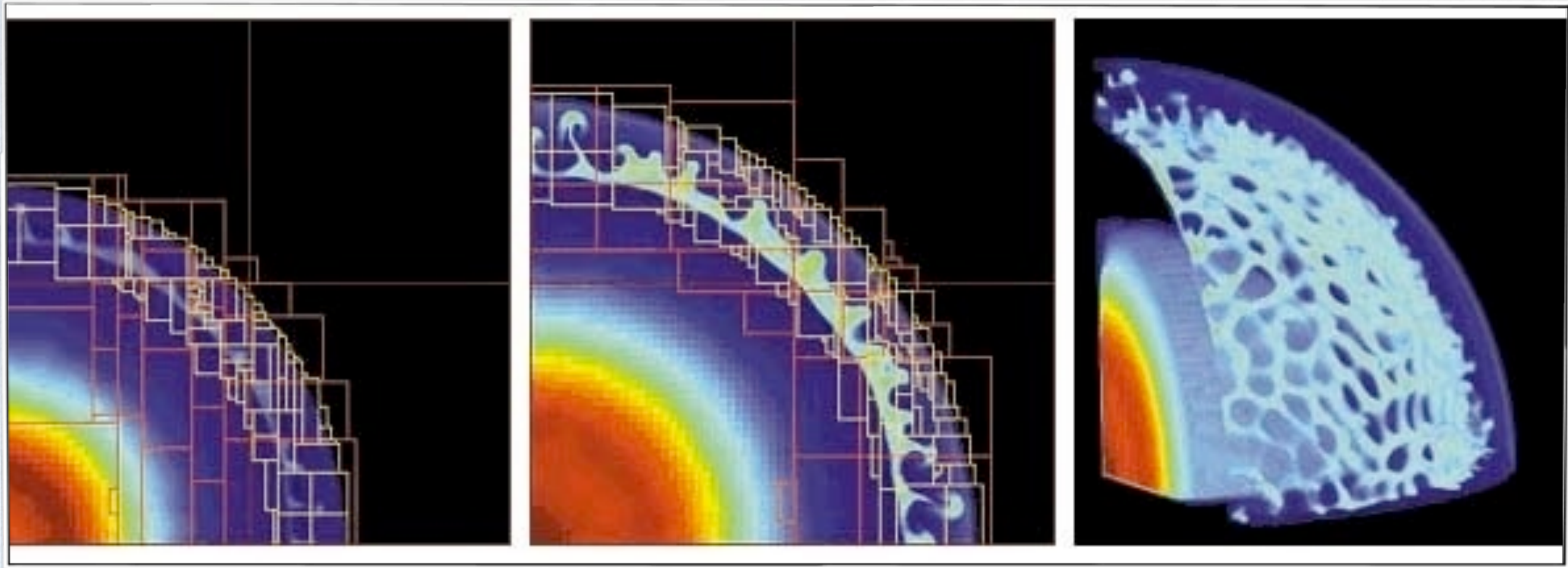


# Evolution of error in effective viscosity





# Adaptive Mesh Refinement



- Support of unstructured grids
- Different mesh orientations
- Low compression rate (Gradient, curvature)
- No explicit control on the compression error

Berger, Colella, J. Comp. Phys., 1989





M. Bergdorf, P. Koumoutsakos. A Lagrangian Particle-Wavelet Method.  
**Multiscale Modeling and Simulation**: A SIAM Interdisciplinary Journal, 5(3), 980-995, 2006

# PARTICLETS : Particles and Wavelets



# WAVELET PARTICLE METHOD

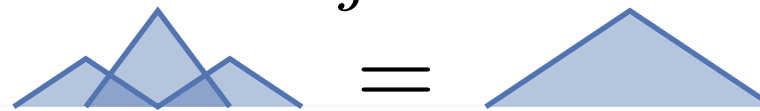
While particles are on grid locations

mollification kernel  $\longleftrightarrow$  basis/scaling function

Multiresolution analysis (MRA)  $\{\mathcal{V}^l\}_{l=0}^L$  of particle quantities

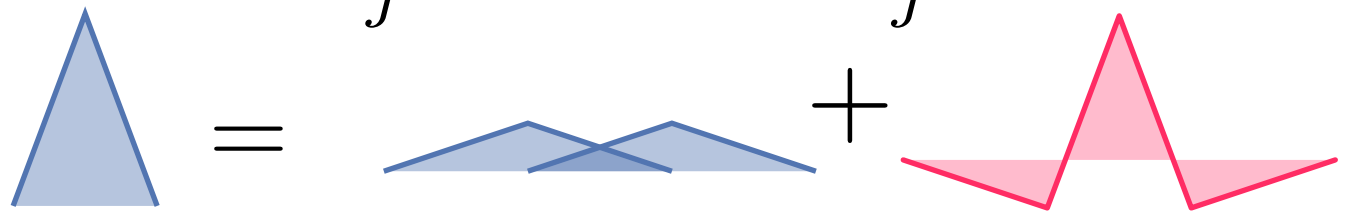
Refineable kernels  
as basis functions of  $\mathcal{V}^l$

$$\zeta_k^l = \sum_j h_{j,k}^l \zeta_j^{l+1}$$



Wavelets as basis functions of the  
complements  $\mathcal{W}^l$

$$\zeta_k^{l+1} = \sum_j \tilde{h}_{j,k}^l \zeta_j^l + \sum_j \tilde{g}_{j,k}^l \psi_j^l$$



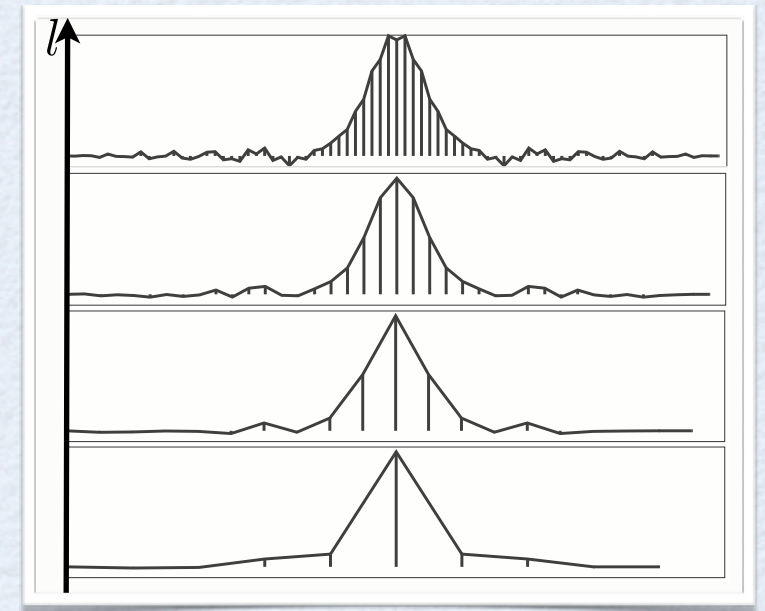


# Multiresolution function representation:

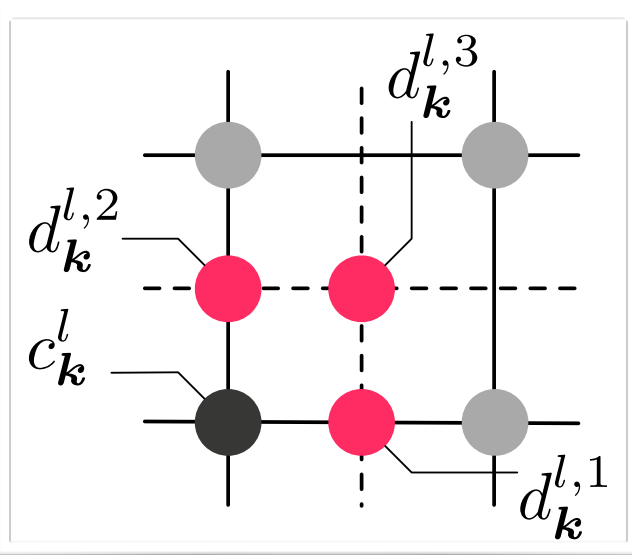
**Analysis** (collocation):  $d_k^l \sim | \text{fine} - \text{Prediction}(\text{coarse}) |$

$$q^L = \sum_k c_k^0 \zeta_k^0 + \sum_{l < L} \sum_k d_k^l \psi_k^l$$

GROUND LEVEL
WAVELETS  
DETAIL COEFFICIENTS



Each wavelet is associated with a specific grid point/particle (2D)



**Compression / Adaptation:**

**Discard** insignificant detail coefficients:  $|d_k^{l,m}| < \epsilon$

**Compressed** function representation:

$$\|q^L - q_{\geq}^L\| < \epsilon \rightarrow \text{Adapted grid}$$



# PARTICLETS : REMESHED PARTICLES + WAVELETS

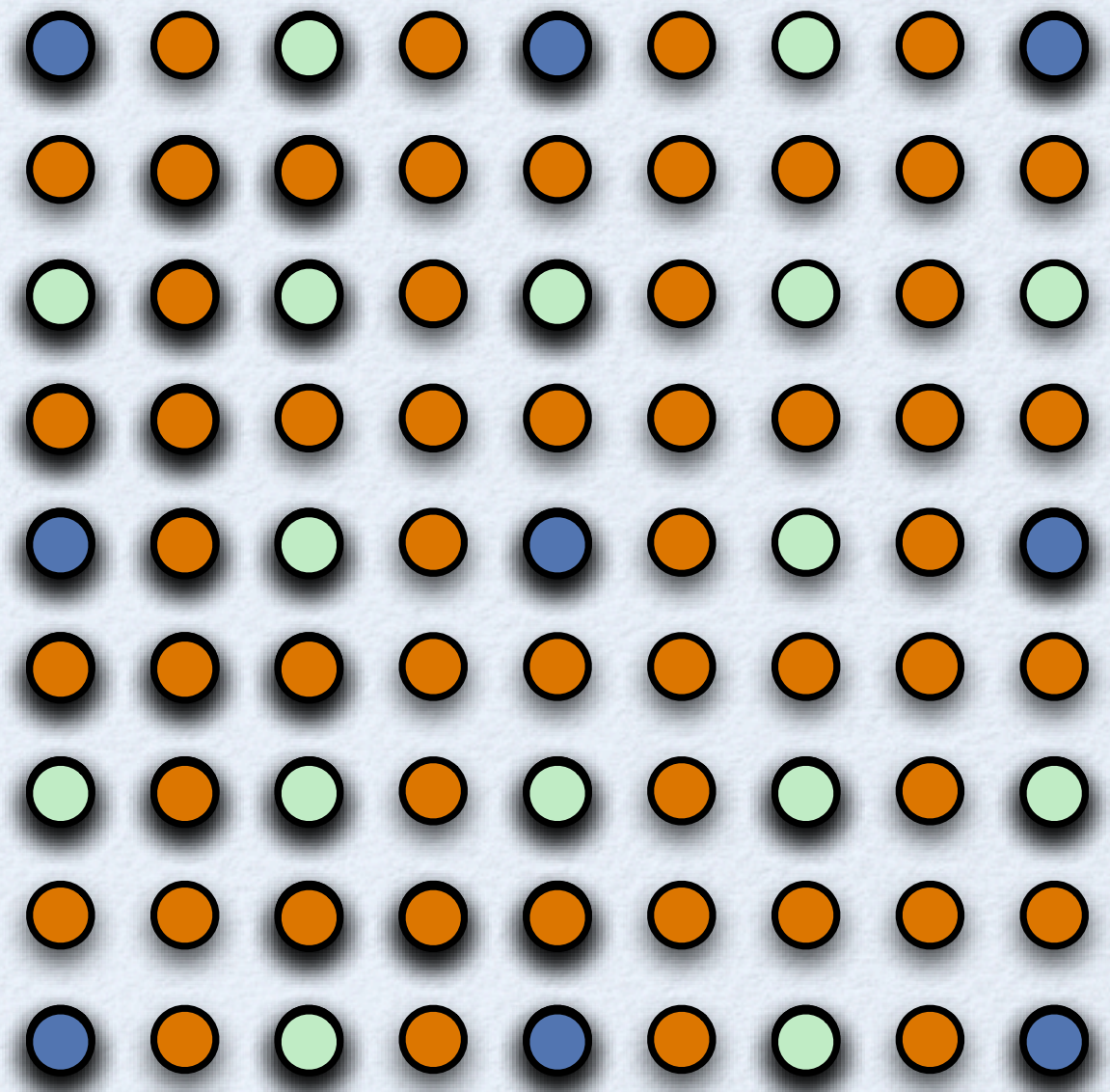
$$q^L = \sum_k c_k^0 \zeta_k^0 + \sum_{l < L} \sum_k d_k^l \psi_k^l$$

“ground” level  $\nearrow$   $c_k^0$

detail coefficients  $\nearrow$   $d_k^l$

wavelets  $\nearrow$   $\psi_k^l$

1. Remesh
2. Wavelets - Compress/Adapt
3. Convect
4. Wavelets Reconstruct
5. GOTO 1





# Wavelet-adapted grids

PDE:

$$\frac{\partial \phi}{\partial t} + \mathbf{u} \cdot \nabla \phi = 0$$

Spatial Differences = filtering operations:

$$F(c_k^l) = \sum_{j=s_f}^{e_f-1} c_{k+j}^l \beta_j^l, \quad \beta_j^l \text{ function of } \{c_m^l\}$$

**GHOSTS** : **easy to compute** – (locally) uniform filtering of the grid

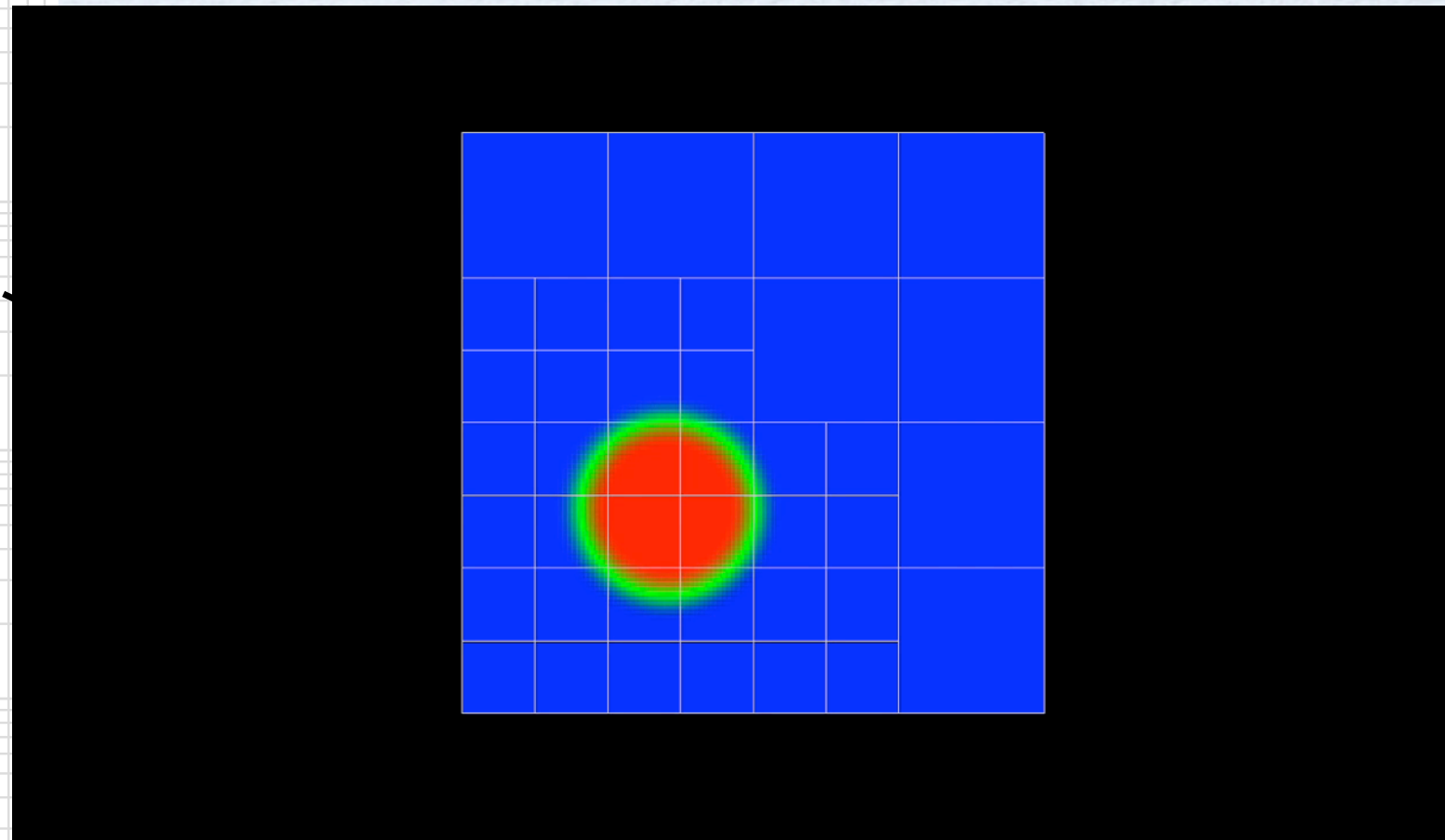
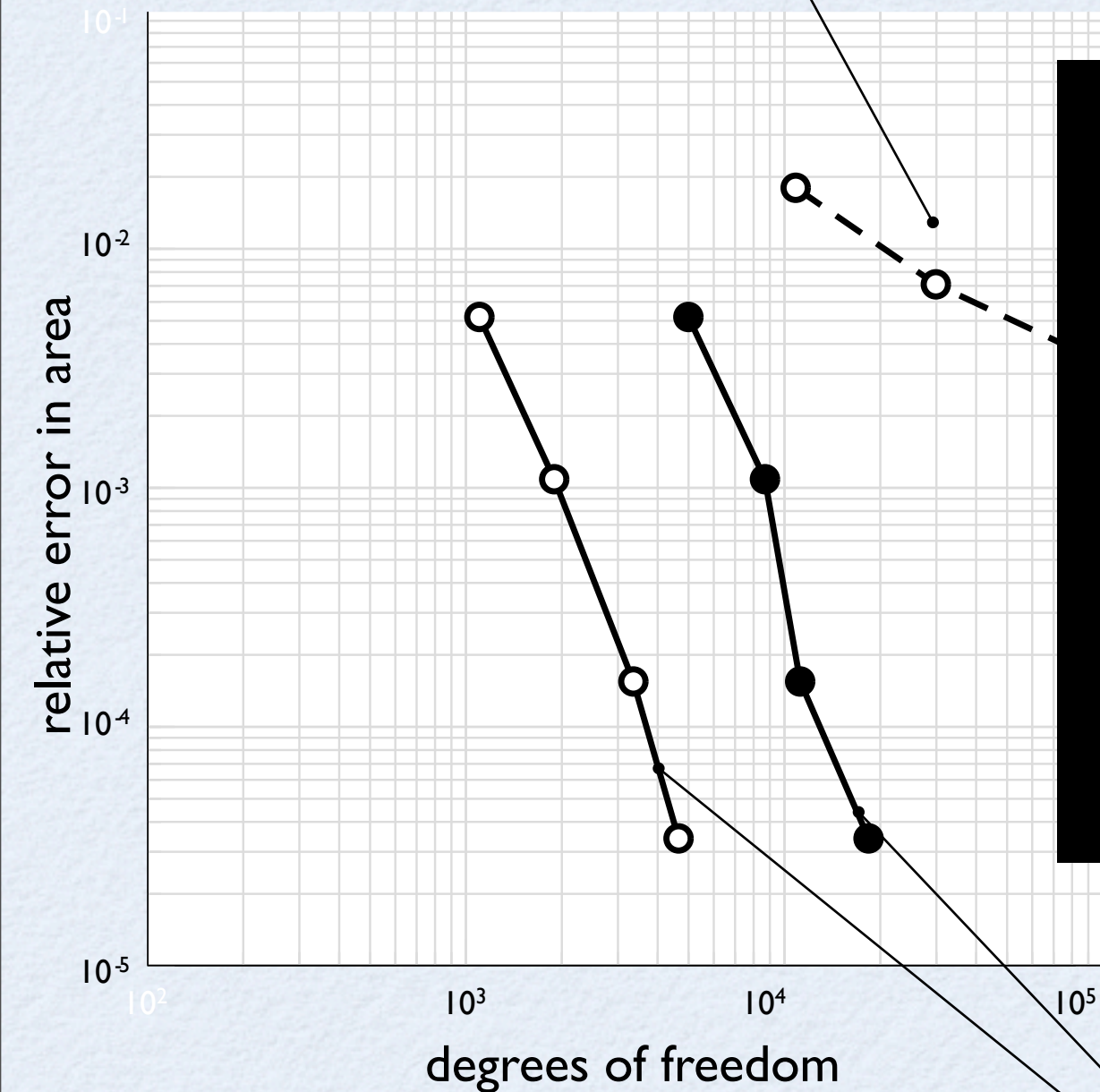


# MULTIRESOLUTION LEVEL SETS

M. Bergdorf, P. Koumoutsakos. A Lagrangian Particle-Wavelet Method, Multiscale Modeling and Simulation: A SIAM Interdisciplinary Journal, 5(3), 980-995, 2006

Enright, Fedkiw et al, 2002

dof = # grid points + aux. particles at t=0.0



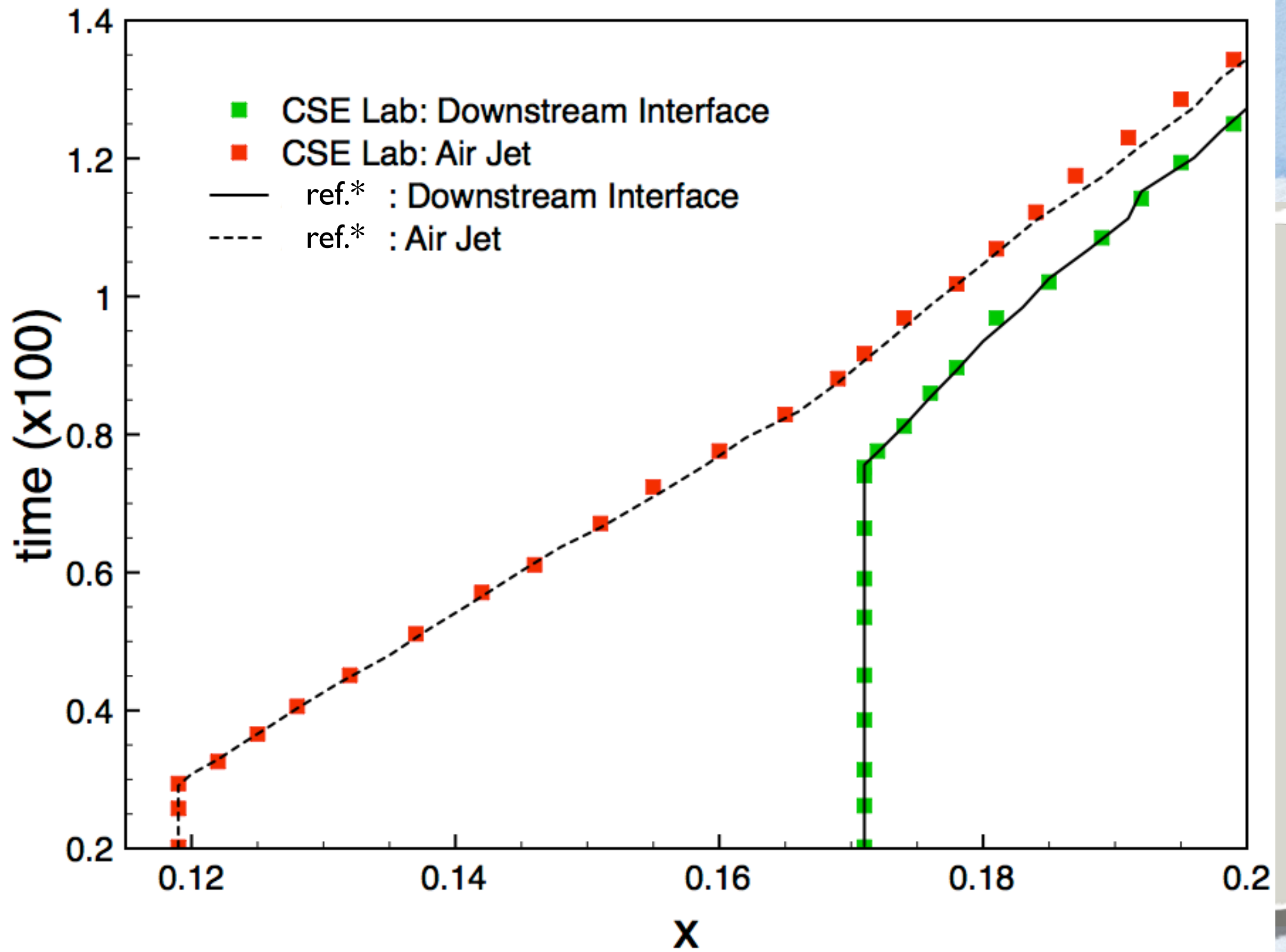
$CFL_{max} \sim 40$

Present Method

dof = # active gp/particles at t=0.0

dof = # active gp/particles at final time

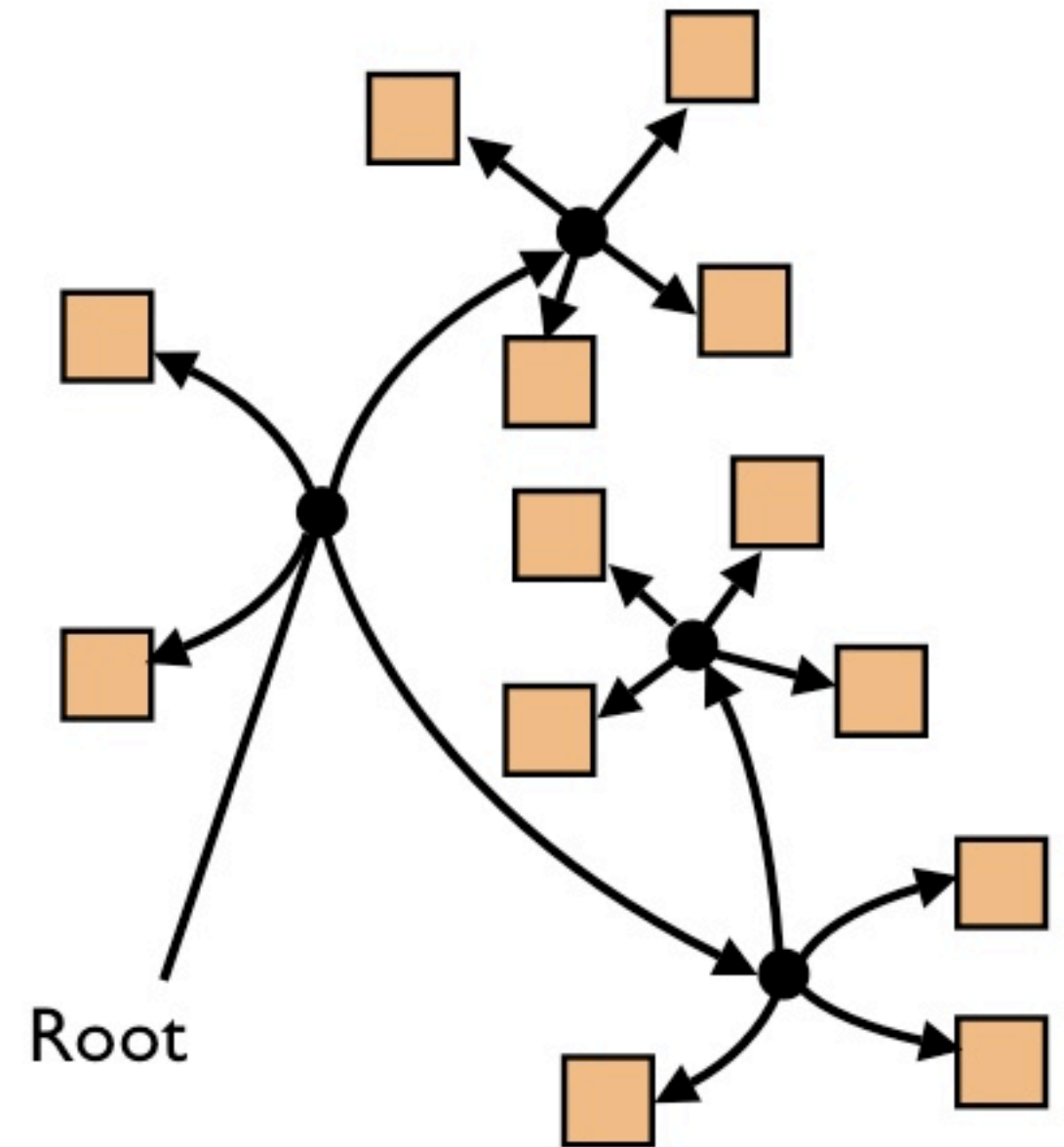
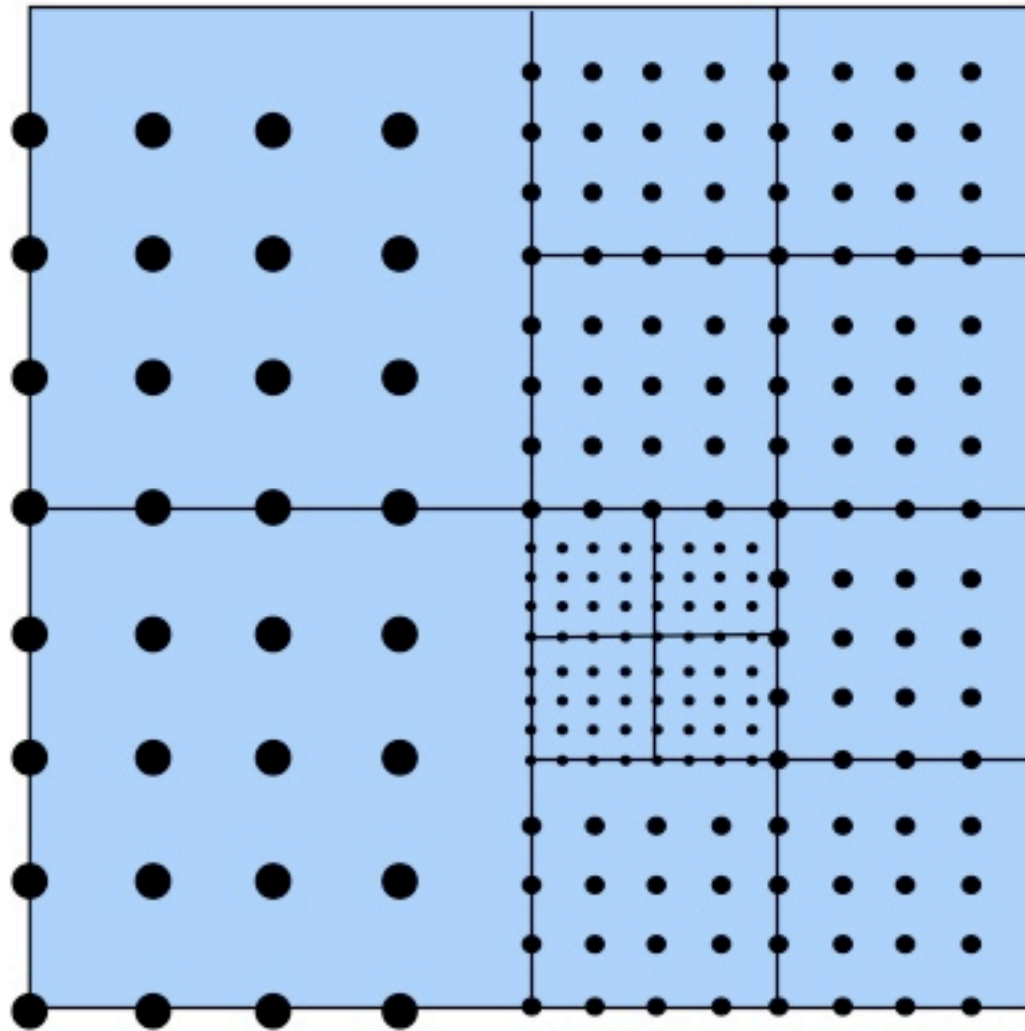




\*Hu, Khoo, Adams and Huang, 2006



# Multi-core: **Block**ed Grid



- Neighbors look-up: less memory indirections
- Less #ghosts
- Within a block: random access



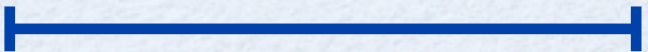
# Multiresolution + MultiCore + GPU

- DISTRIBUTED TASKS

1.task parallel,ghost computing  $\rightarrow$  **multi-core**

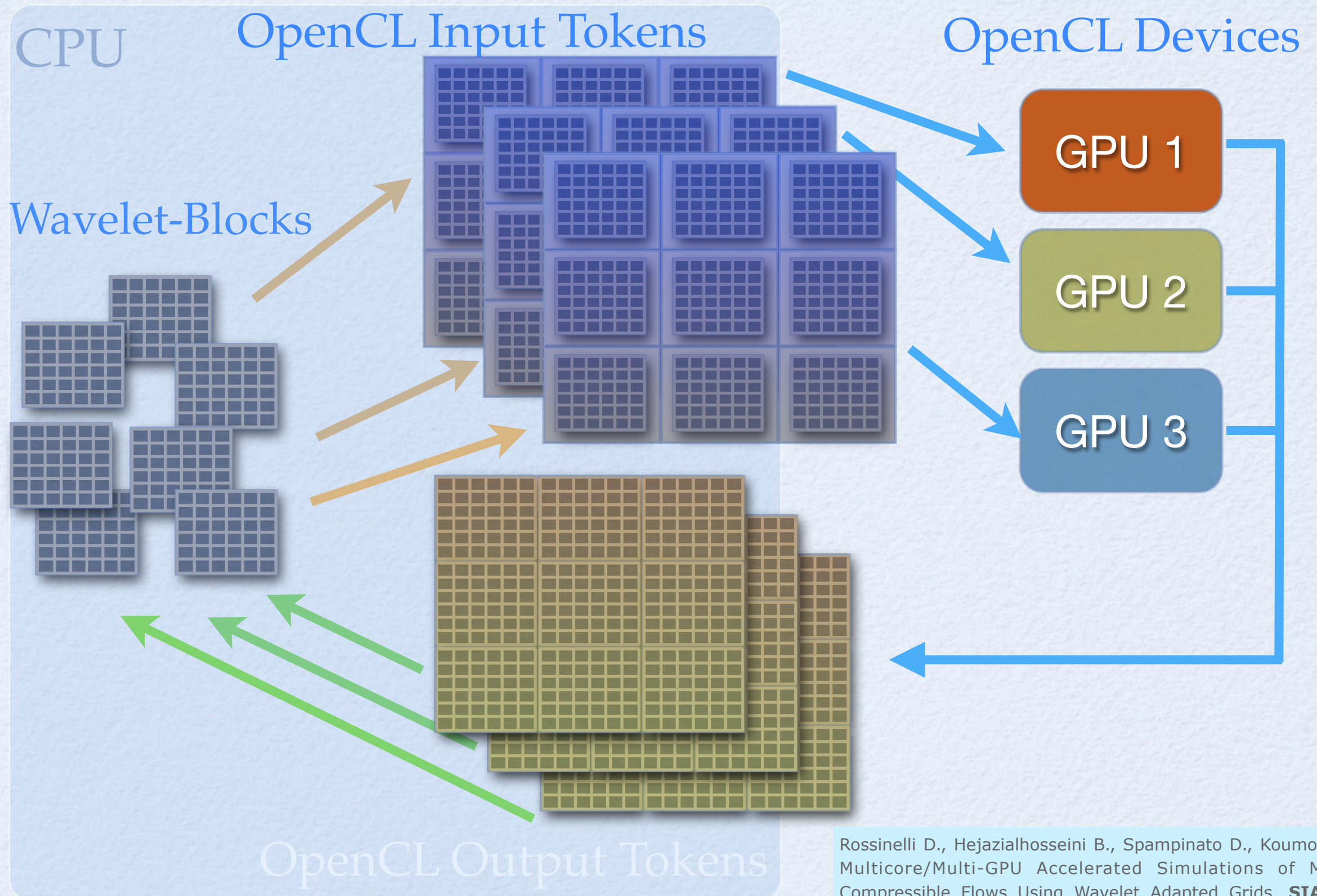
2.fine-grained data parallelism for RHS  $\rightarrow$  **GPUs**

$$q^{\text{new}} = q^{\text{old}} + \delta t \mathbf{F} (q^{\text{old}}, \nabla q^{\text{old}})$$

  
**OpenCL/GPUs**



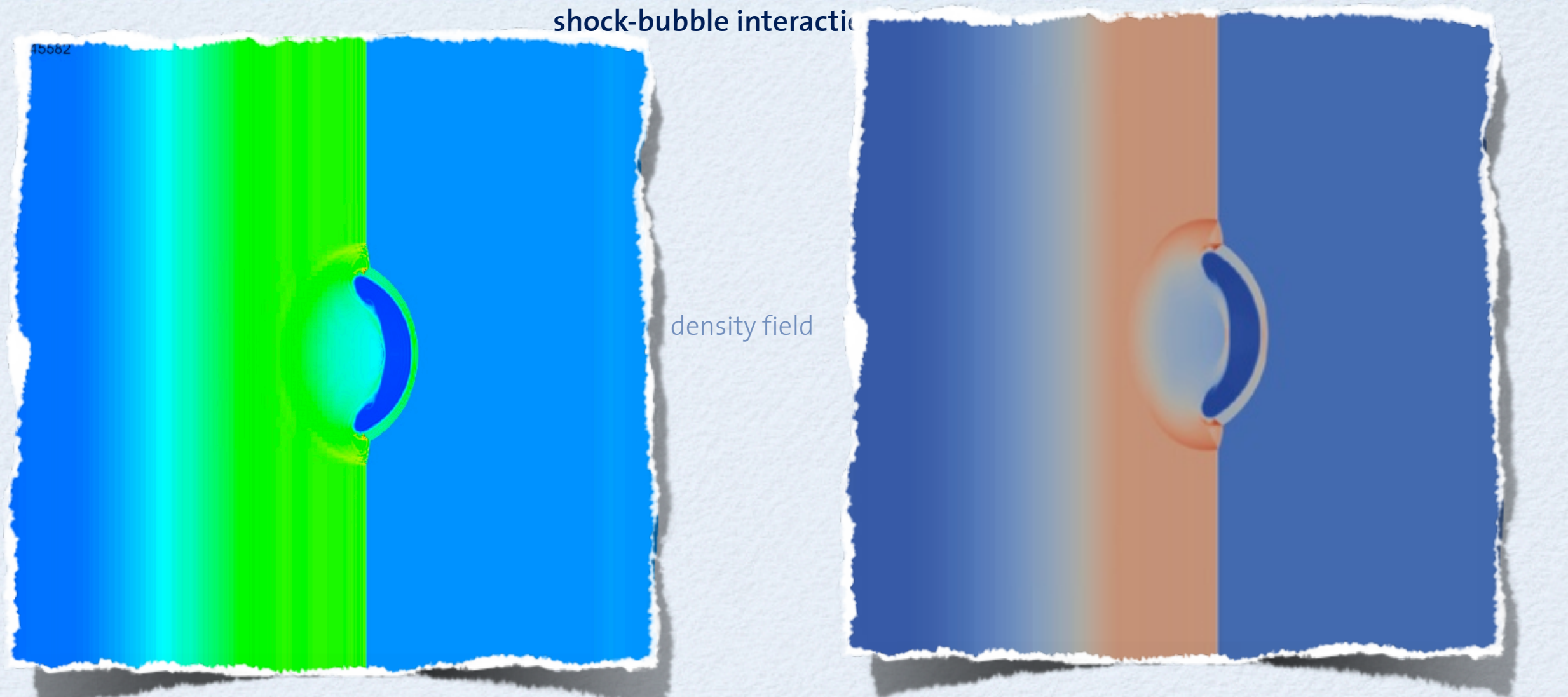
# Wavelet Blocks on GPUs



Rossinelli D., Hejazialhosseini B., Spampinato D., Koumoutsakos P., Multicore/Multi-GPU Accelerated Simulations of Multiphase Compressible Flows Using Wavelet Adapted Grids, **SIAM J. Sci. Comput.**, 33, pp. 512-540, 2011



# A comparison of CHOMBO vs MRAG



**Chombo:**  
**91 min, 230 MB**

single-phase  
2nd order PPM

**MRAG** (home grown, swiss quality stuff):  
**56 min, 244 MB (+ 1 GPU: 7 min)**

multi-phase  
3rd order WENO scheme

Rossinelli D., Hejazialhosseini B., Spampinato D., Koumoutsakos P., Multicore/Multi-GPU Accelerated Simulations of Multiphase Compressible Flows Using Wavelet Adapted Grids, **SIAM J. Sci. Comput.**, 33, pp. 512-540, 2011





# BOUNDARIES + ALGORITHMS



# TIME : FLOW AVeraging integratOR

Tao, Owhadi, & Marsden, Multiscale Model. Simul., 2010.

Stiff ODEs:

$$\dot{\mathbf{u}} = \mathbf{G}(\mathbf{u}) + \frac{1}{\epsilon} \mathbf{F}(\mathbf{u})$$

$\epsilon \ll 1$

with the *legacy* integration scheme:

$$\bar{\mathbf{u}}_{t+\tau} = \Phi_{\tau}^{1/\epsilon}(\bar{\mathbf{u}}_t)$$

Small time-step:  $\tau \ll \epsilon$

*FLAVOR*:

$$\bar{\mathbf{u}}_{t+h} = \left( \Phi_{\frac{h}{M}-\tau}^0 \circ \Phi_{\tau}^{1/\epsilon} \right)^M (\bar{\mathbf{u}}_t)$$

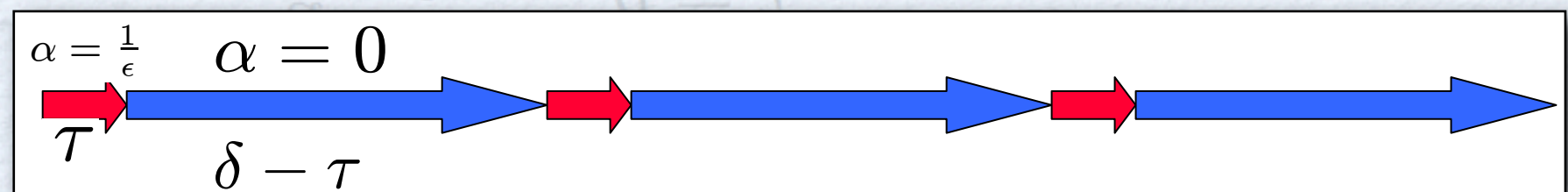
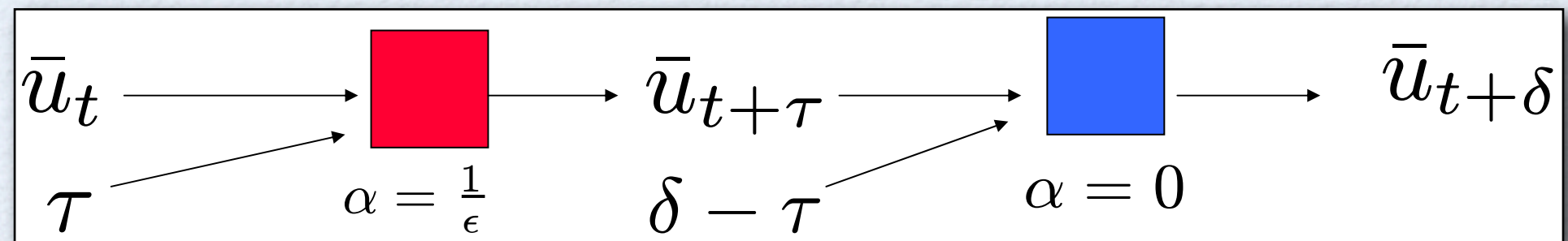


Large time-step:  $\delta = \frac{h}{M}$

$M$  regulates accuracy of large time-step

$$0 < \tau \ll \epsilon \ll \delta \ll 1$$

turn on/off large coefficients:





# FLAVOR + Stiff Stochastics

SSA written as

$$(\mathbf{X}_n, t_n) = \left( \Phi^{(1/\epsilon)} \right) (\mathbf{X}_{n-1}, t_{n-1})$$

Define FLAVOR-SSA:

$$(\mathbf{X}_n, t_n) = \left( \Phi^{(\xi/\epsilon)} \circ \Phi^{(1/\epsilon)} \right) (\mathbf{X}_{n-1}, t_{n-1})$$

*amounts to rescaling  
the total propensity  
every other  
iteration.*

$$\hat{a}_0(t, \xi) := \frac{\xi}{\epsilon} \sum_i \tilde{a}_i^{(fast)}(t) + \sum_j a_j^{(slow)}(t)$$

$\delta = \delta(\xi) \sim \mathcal{E} \left( \frac{1}{\hat{a}_0} \right)$   
 $\xi \in [0, 1]$

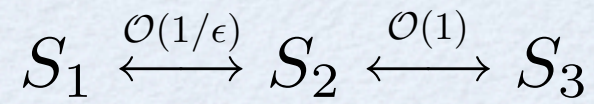
$\xi = 0 \implies$  Largest Speedup

$\xi = 1 \implies$  SSA

**OPTIMAL**  
*value?*

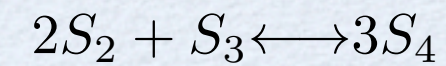
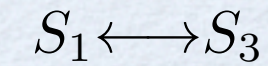
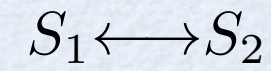
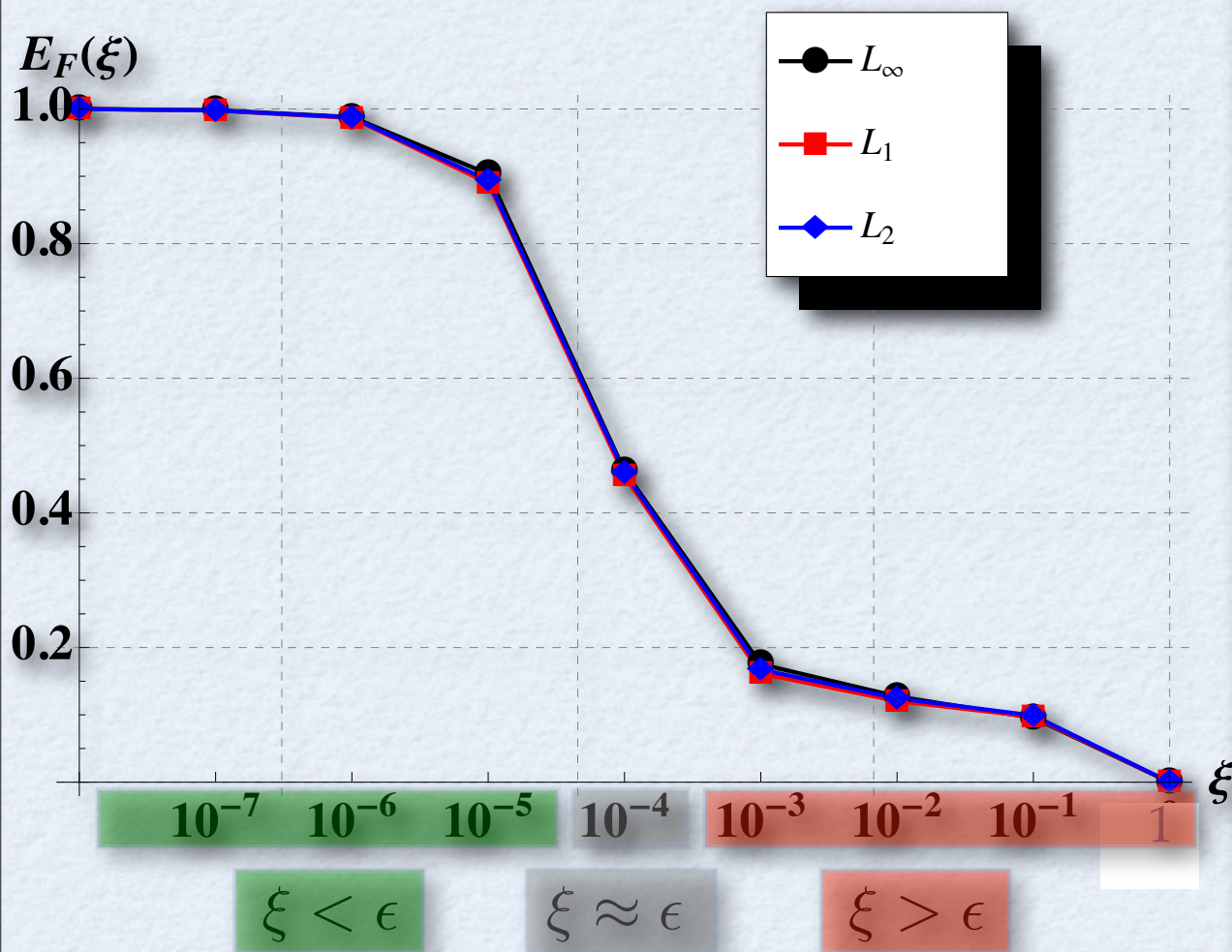


# FLAVOR -S : A Cutoff Phenomenon

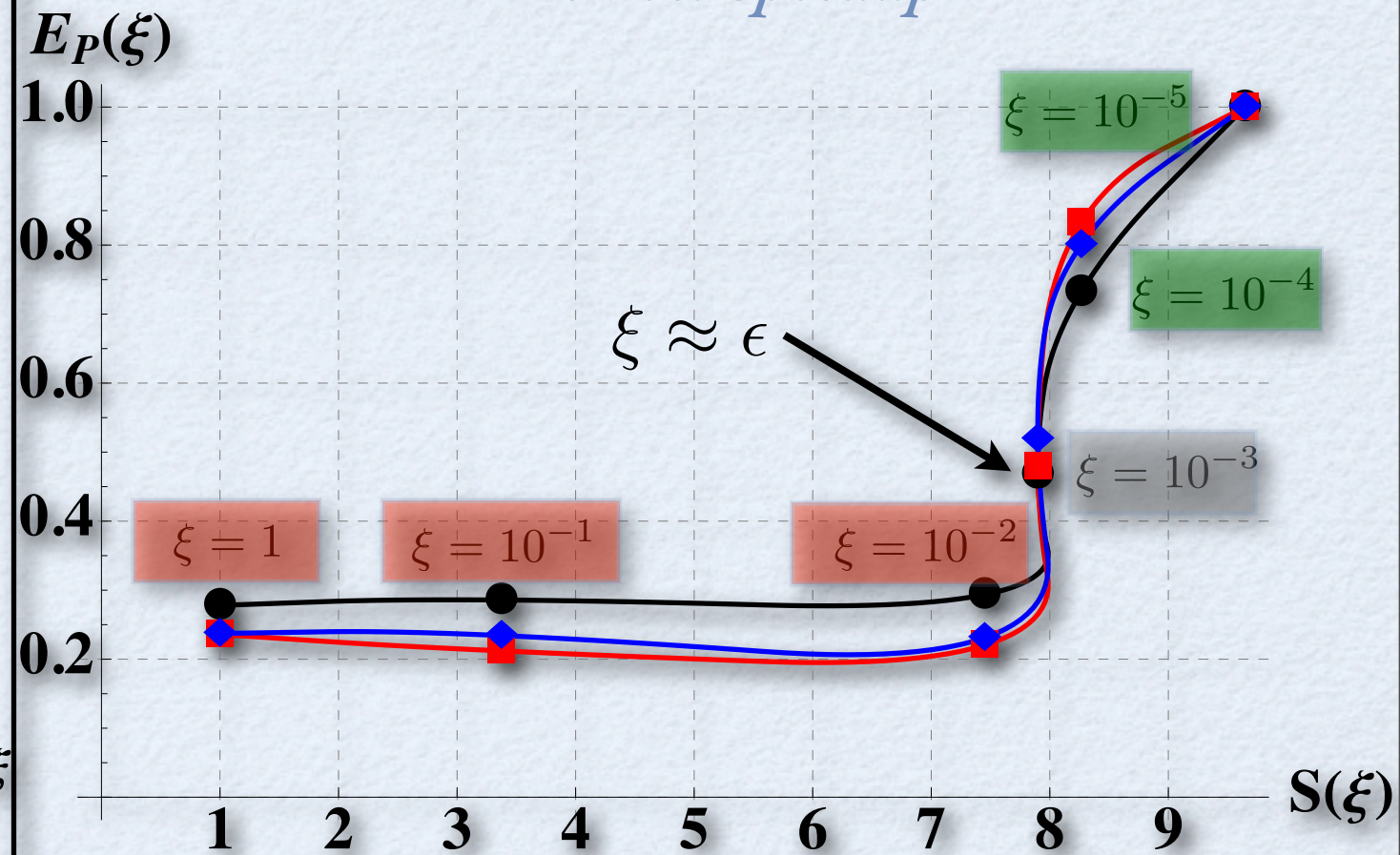


$$\epsilon = 10^{-4}$$

*Error vs.  $\xi$*



*Error vs. Speedup*



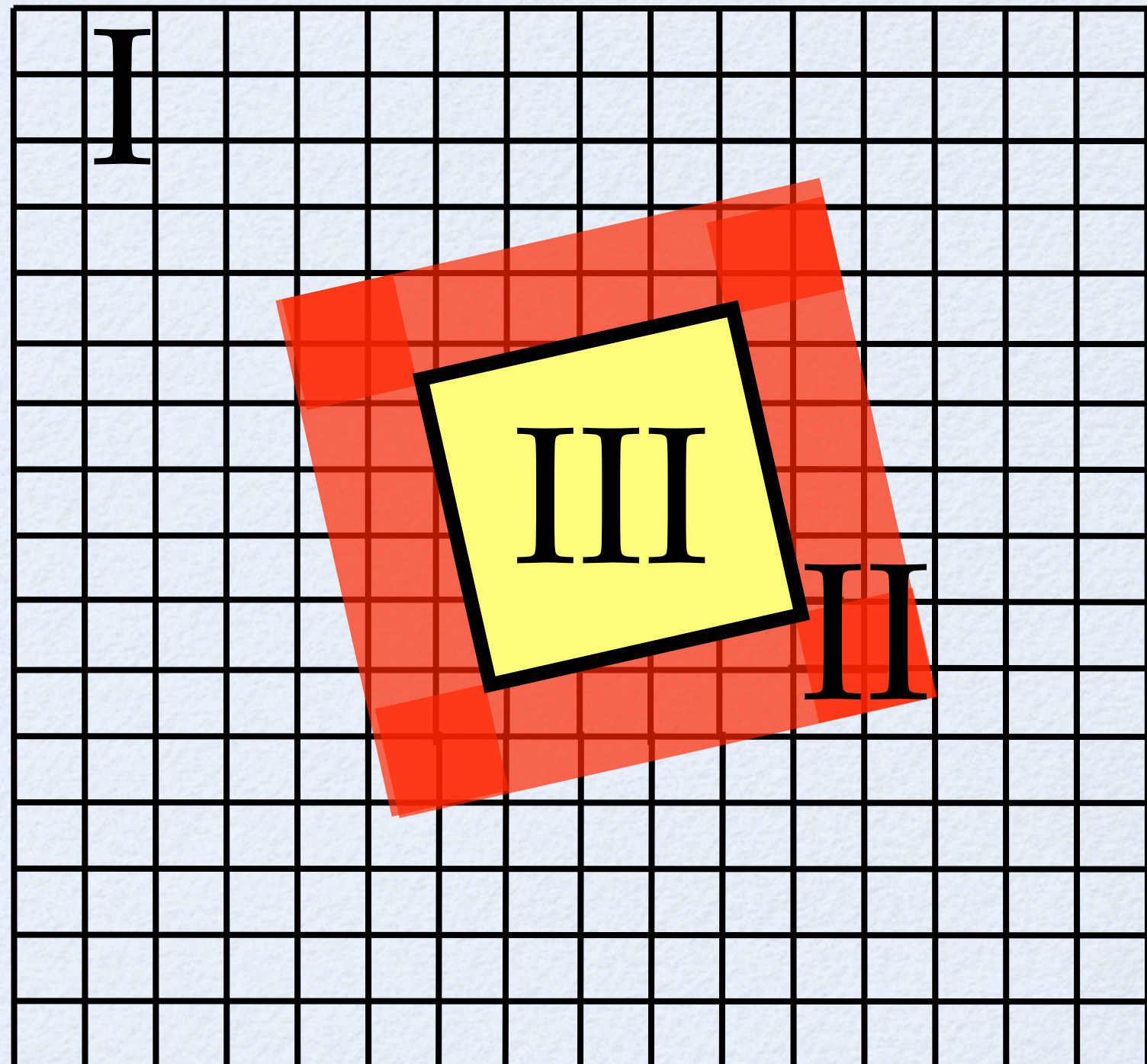
Bayati B., Owhadi H., Koumoutsakos P, J. Chem. Phys., 2010



# Boundary Conditions = Coupling

COUPLING Different Physics in Space / Time

|                       | I     | II                 | III    |
|-----------------------|-------|--------------------|--------|
| No Slip               | FLUID | Sharp Force        | -      |
| No Slip + PENALTY/IBM | Fluid | Smooth Force       | -      |
| Flow-Structure        | Fluid | Smooth/Sharp Force | Solid  |
| Multiscale I          | Fluid | Fluid+MD           | MD     |
| Multiscale II         | Fluid | Fluid+DPD          | DPD+MD |

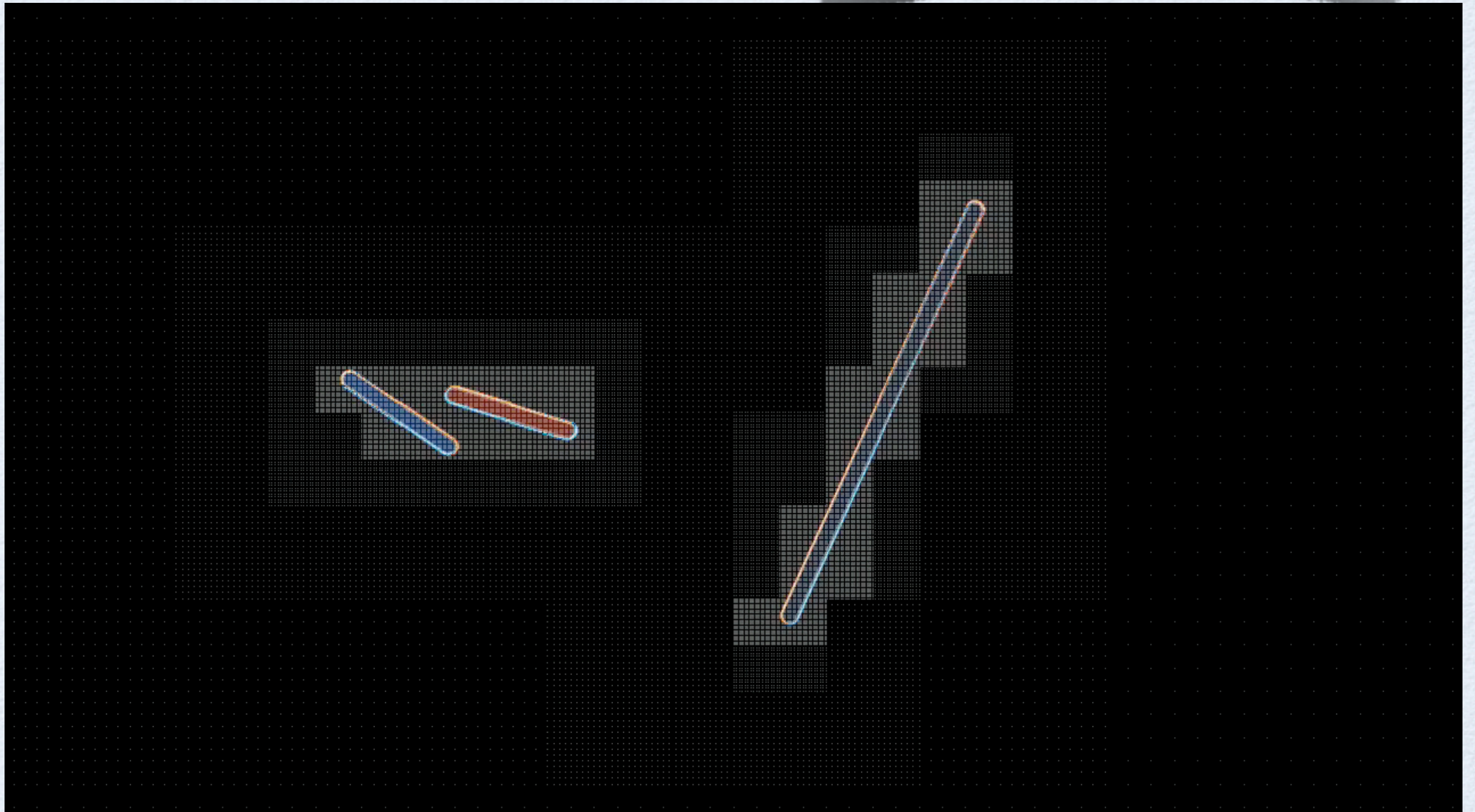




# COMPRESSIBLE FLOWS

Moving Boundaries

Brinkman Penalization for  
Compressible Flow

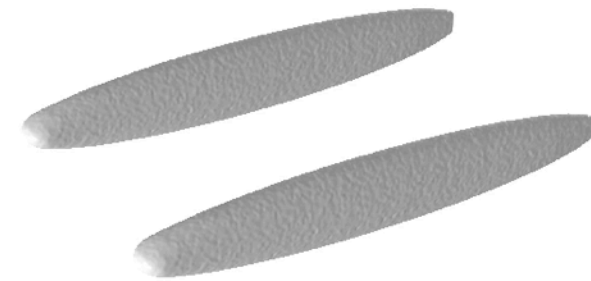




# FISH SCHOOLING



1 FISH

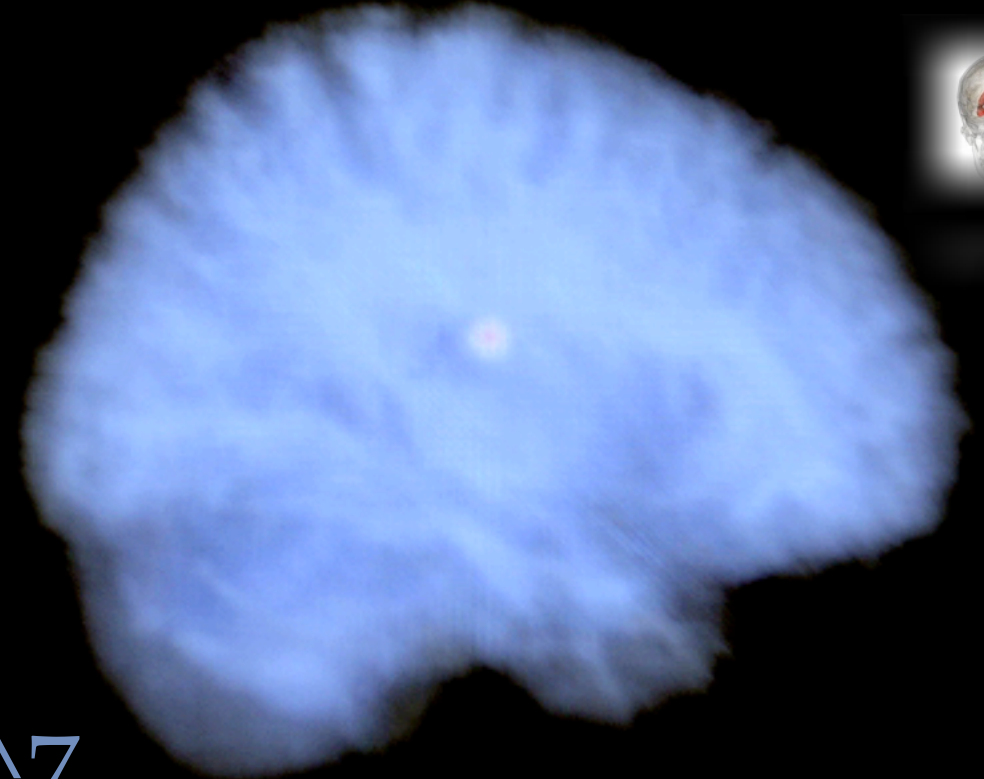
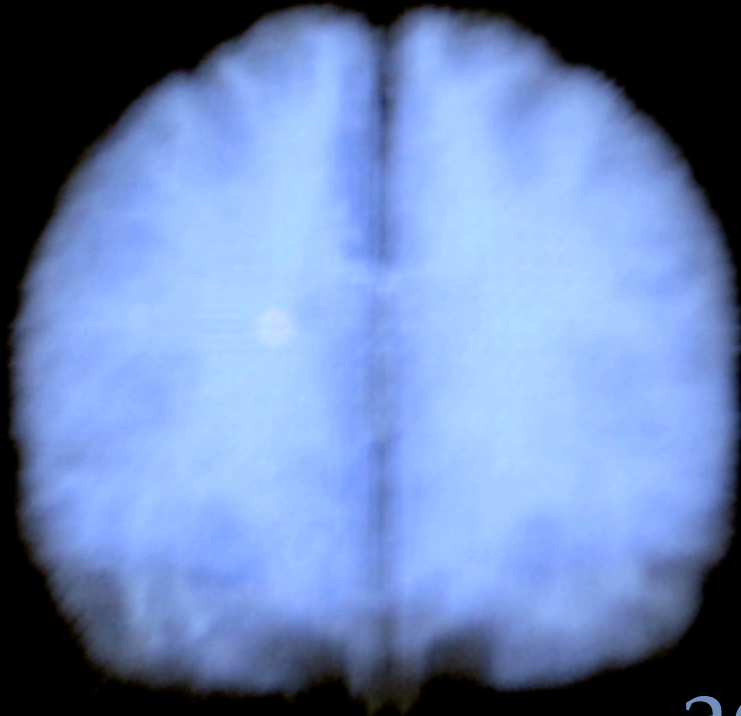


2 FISH (OBVIOUSLY)

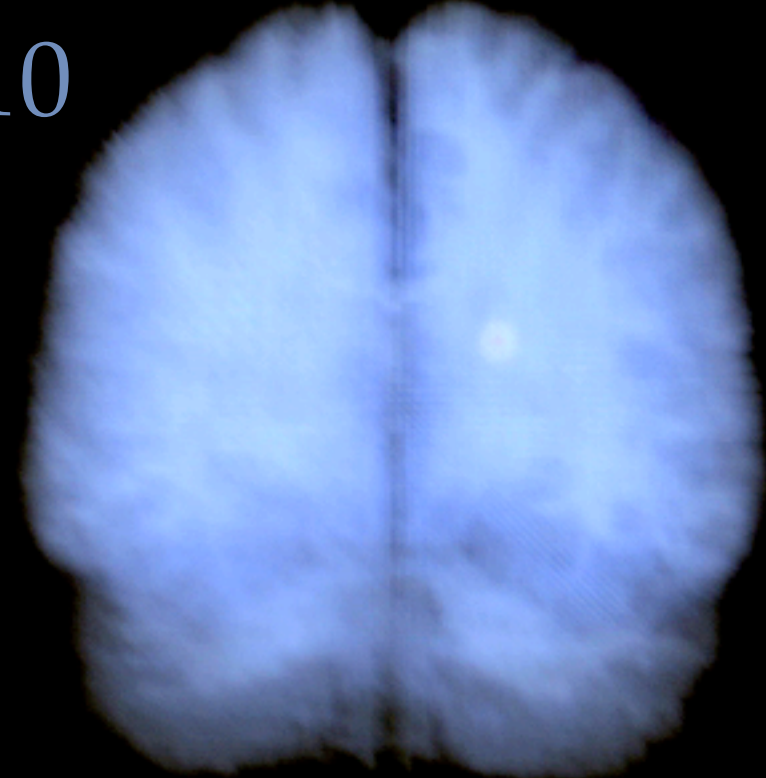
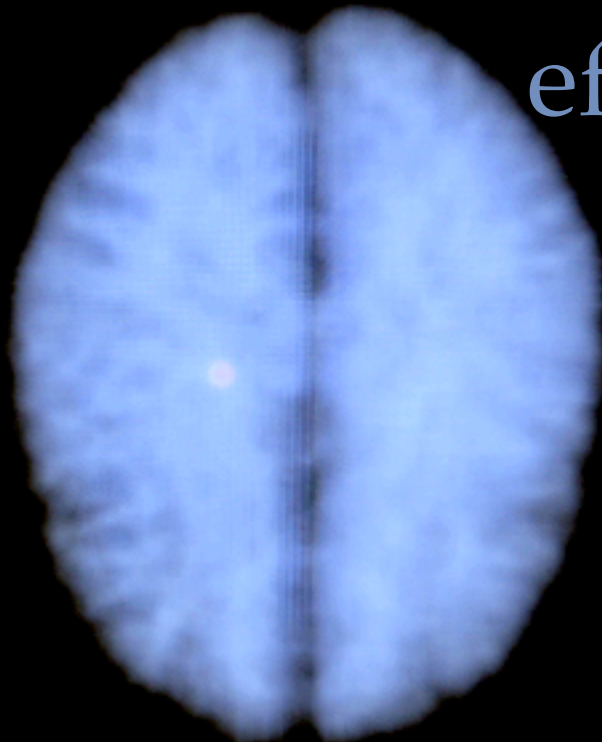


# Simulations of Gliomas using MRAG

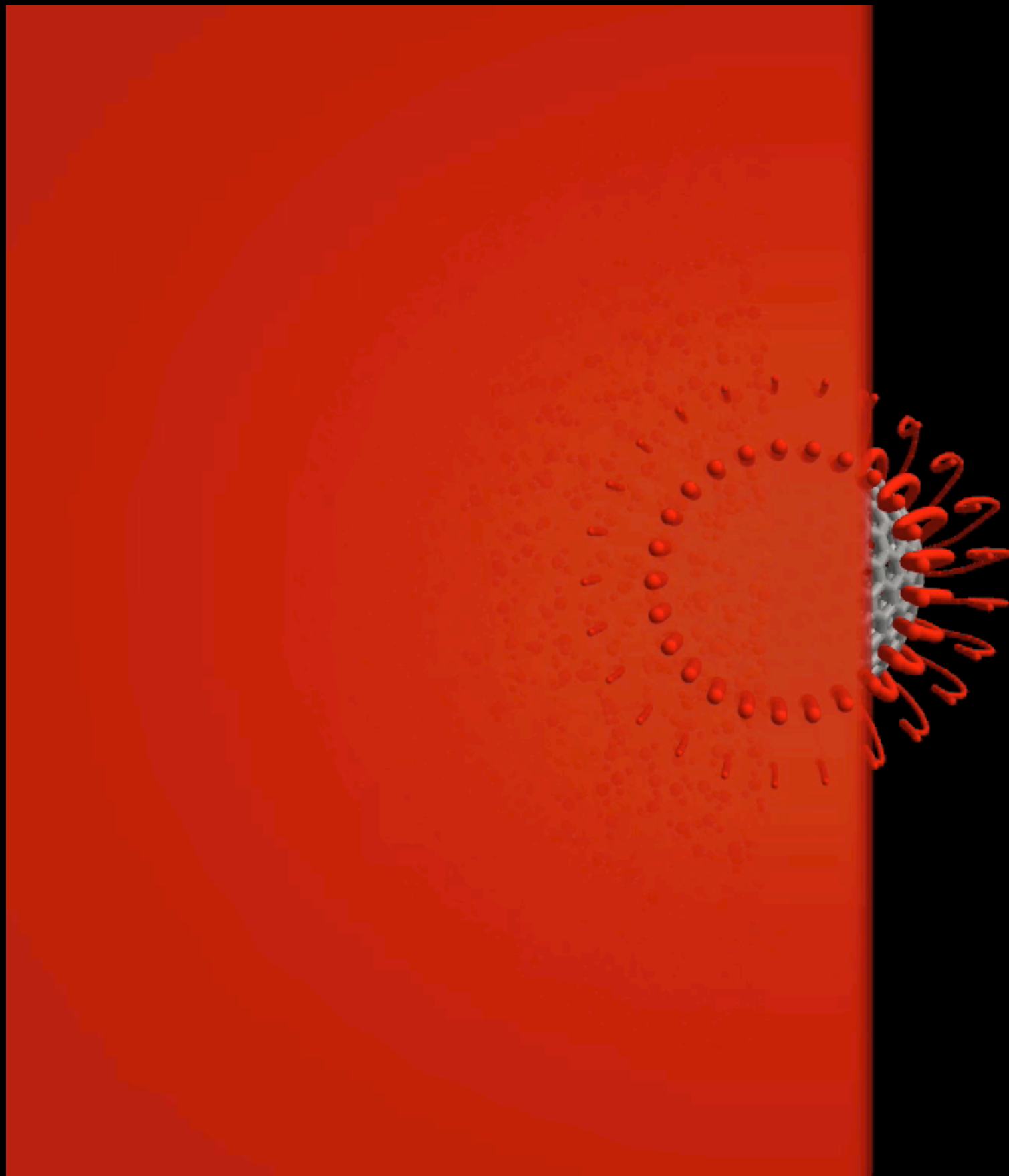
Time: 0.00 years



actual  $M = 10^7$   
effective  $M = 10^{10}$









# COMPUTATIONAL Challenges

"Although **X** seem to be a small step from **Y** computation, in fact, they represent a huge step. They discard the most essential and appealing properties of **Y** computation:

**understandability, predictability, and determinism.**

**X** as a model of computation, are wildly nondeterministic, and the job of the programmer becomes one of pruning that nondeterminism."

-- *'The Problem with X, Edward A. Lee, UC Berkeley, 2006*

**X = threads**

**Y = sequential**

**X = Multi-scale**

**Y = single-scale**



Bergdorf



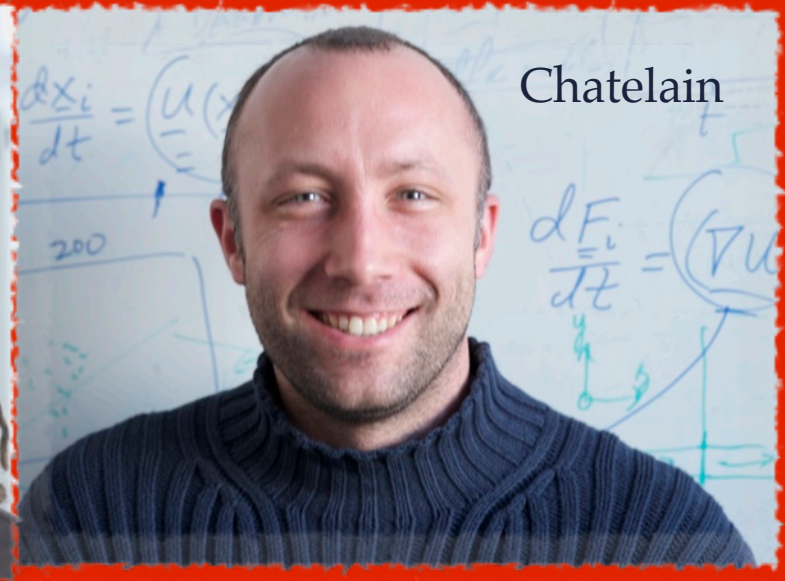
Bayati



Rossinelli



Chatelain



Hedjazialhosseini

