

# Science Driver: Bio-Transport Computations

## Computing of Transport Processes in Biological Systems

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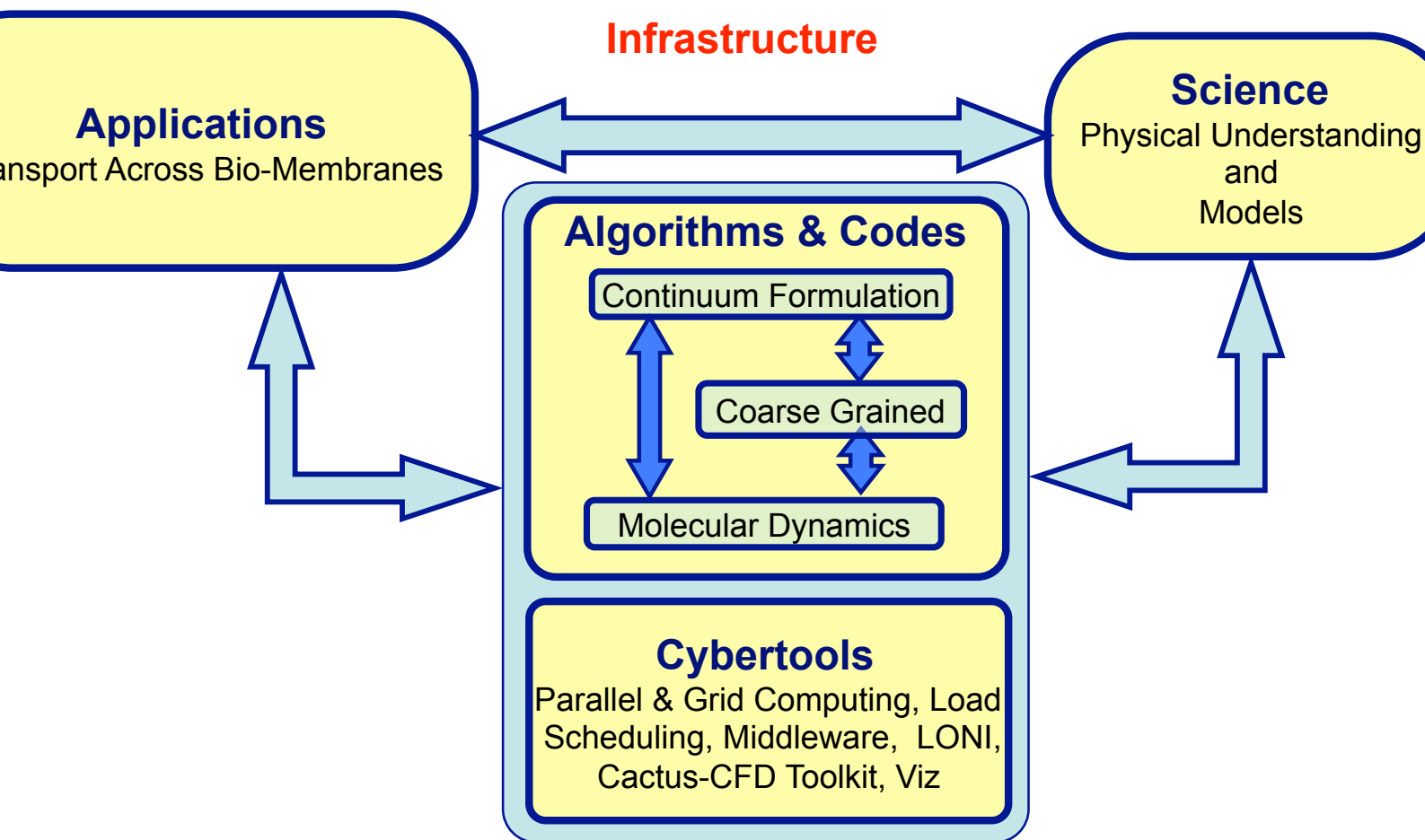
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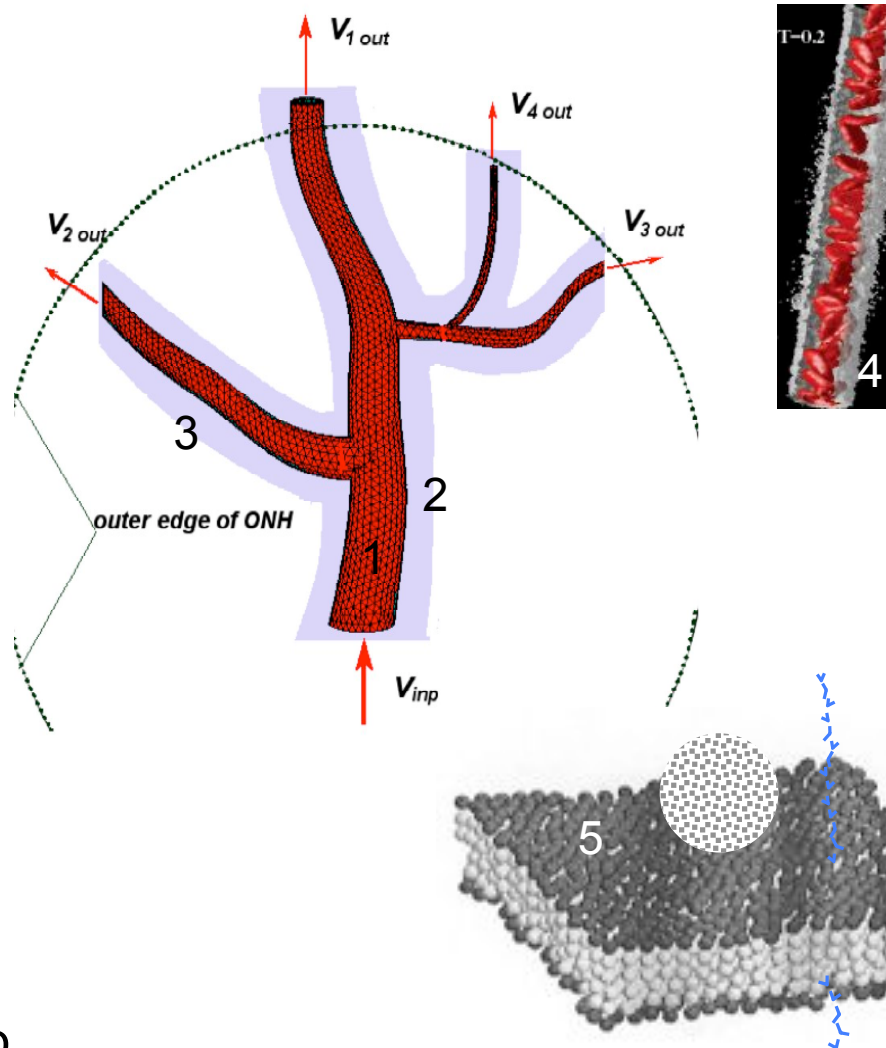
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# Prediction and understanding of oxygen transport in biological systems

1. Continuum flow in larger vessels-Navier Stokes
2. Porous media transport across vessel walls & tissues-Brinkmann
3. Structural deformation of vessels/tissues-
4. Particle flow in capillaries-Lattice-Boltzmann
5. Atomistic transport across cellular interfaces-Molecular Dynamics
6. Upscaling from atomistic to continuum



Development of **computationally efficient numerical methods** or **algorithms** needed for biological transport calculations

- ✓ Structural calculation using a meshless particle method
- ✓ Flow-Structure Interaction (FSI) methodology using Immersed Boundary Method (IBM)



Year

Contributing to **improved science-understanding of small molecule flow/transport physics** under asymmetric concentrations and applied stresses

- ✓ Asymmetric calculations of molecule/particle transport across lipid bilayers



Year

Contributing to **improved computational infrastructure-** collaborating with the cybertools group responsible for developing **the CFD toolkit**

- ✓ Development of cactus-compatible routines for transport and flow calculations
- ✓ Validation Studies



Year

Contributing to improved science-understanding of oxygen flow/transport physics under elevated pressures

## Continuum flow and transport calculations

Multiblock structured grid with continuous grid lines across block interfaces

Fractional step algorithm with staggered grid locations for the velocity (stored at cell faces)

Pressure-poisson equation for pressure

Consistent second order differencing for diffusion and pressure terms and upwind biased differencing for the convective terms

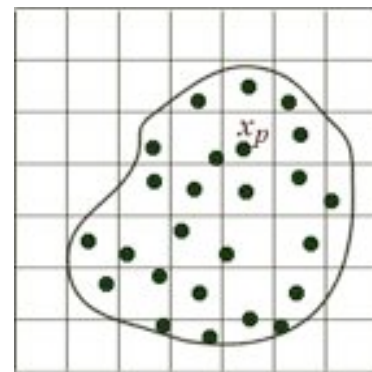
Explicit and implicit second order temporal differencing

### Flow-structure interaction

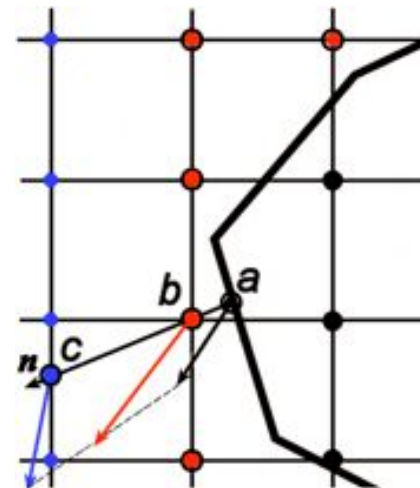
- Particle-based meshless calculations for structural deformations (called material point method-MPM)

Immersed Boundary Methodology (IBM) for resolving boundary conditions along moving interfacial surfaces

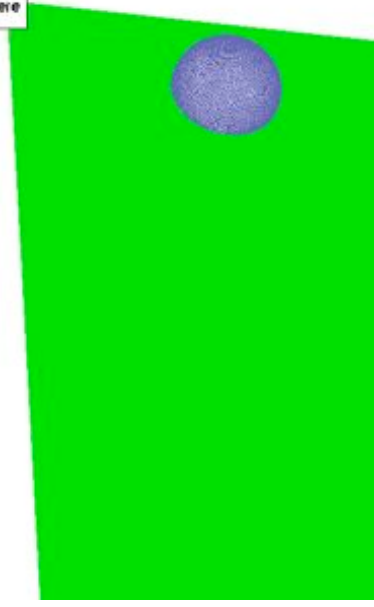
### Flow-Structure Interaction for Biosystems



Background grid for solution of momentum equations



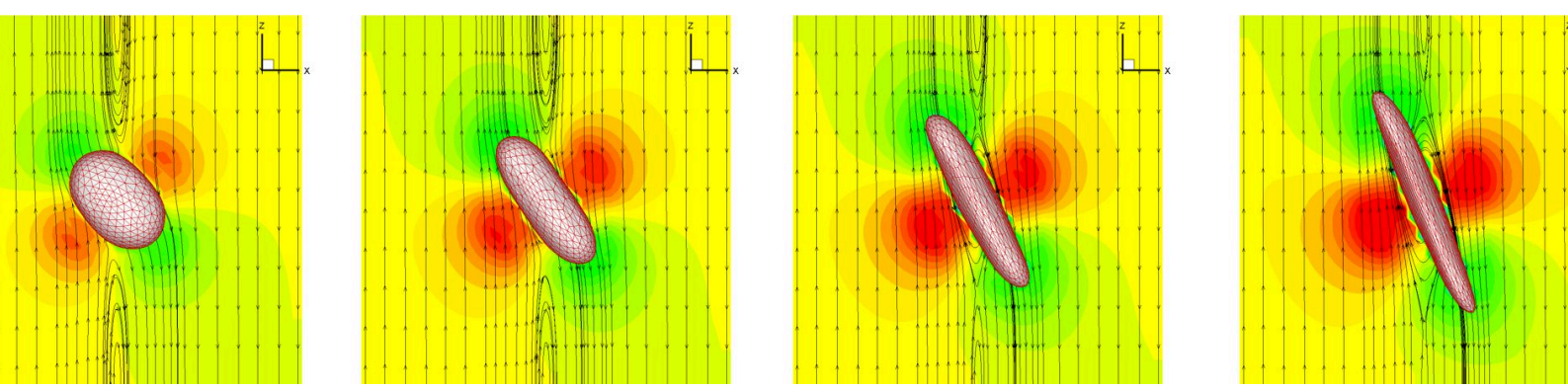
14. Dropping a sphere  
Re=50



## Material-Point Method (MPM) for structural deformations

- ★ Arbitrary distribution of points on the solid body/surface
- ★ Material points are solved (deformation & stress) on a background grid that is independent from the fluid grid
- ★ Flow-structure coupling through boundary/interface conditions
- ★ Flow around deforming surface handled through IBM

Time = 1.00



- ✿ Collaborating with the WP4 group for the development of a CFD Toolkit;
- ✿ Finite volume, multi block;
- ✿ Data array structure consistent with current structure in Cactus;
- ✿ Multi-block grid from commercial grid generators;
- ✿ Baseline code developed for laminar flow; several benchmarks being run to provide WP4 input-output files for Toolkit verification and validation;
- ✿ Long term plans are to transition to the Toolkit for the biosystems transport simulation;

- ★ Implemented suggestions for improved performance of parallel code—seen improvements
- ★ Discussions ongoing with Viz groups to get better access to better visualization codes (WP3)
- ★ Discussions ongoing on use of a Lattice Boltzmann code for particle simulations
- ★ Discussions ongoing on most effective ways of doing CFD-MD coupling

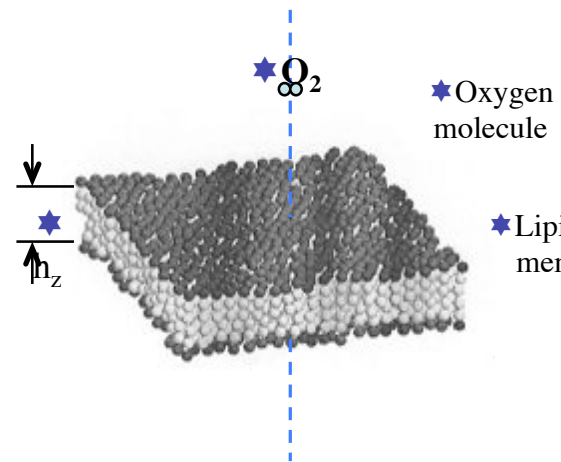
Diffusion rate and permeability coefficients across cell walls and tissues for different conditions are generally not known reliably (difficulty in in situ measurements)

Specifically designed MD simulations under different conditions can provide:

Atomistic insight and molecular mechanism underlying the transport of  $O_2$  across a lipid bilayer membrane in order to determine which details are important for the permeation process.

Derive the oxygen diffusivities,  $D_{O_2}$ , inside the homogeneous region of a lipid bilayer.

Derive permeation rates,  $P_{O_2}$ , indirectly via computation of the free energy and diffusion rate profiles of a  $O_2$  molecule across the lipid bilayer.

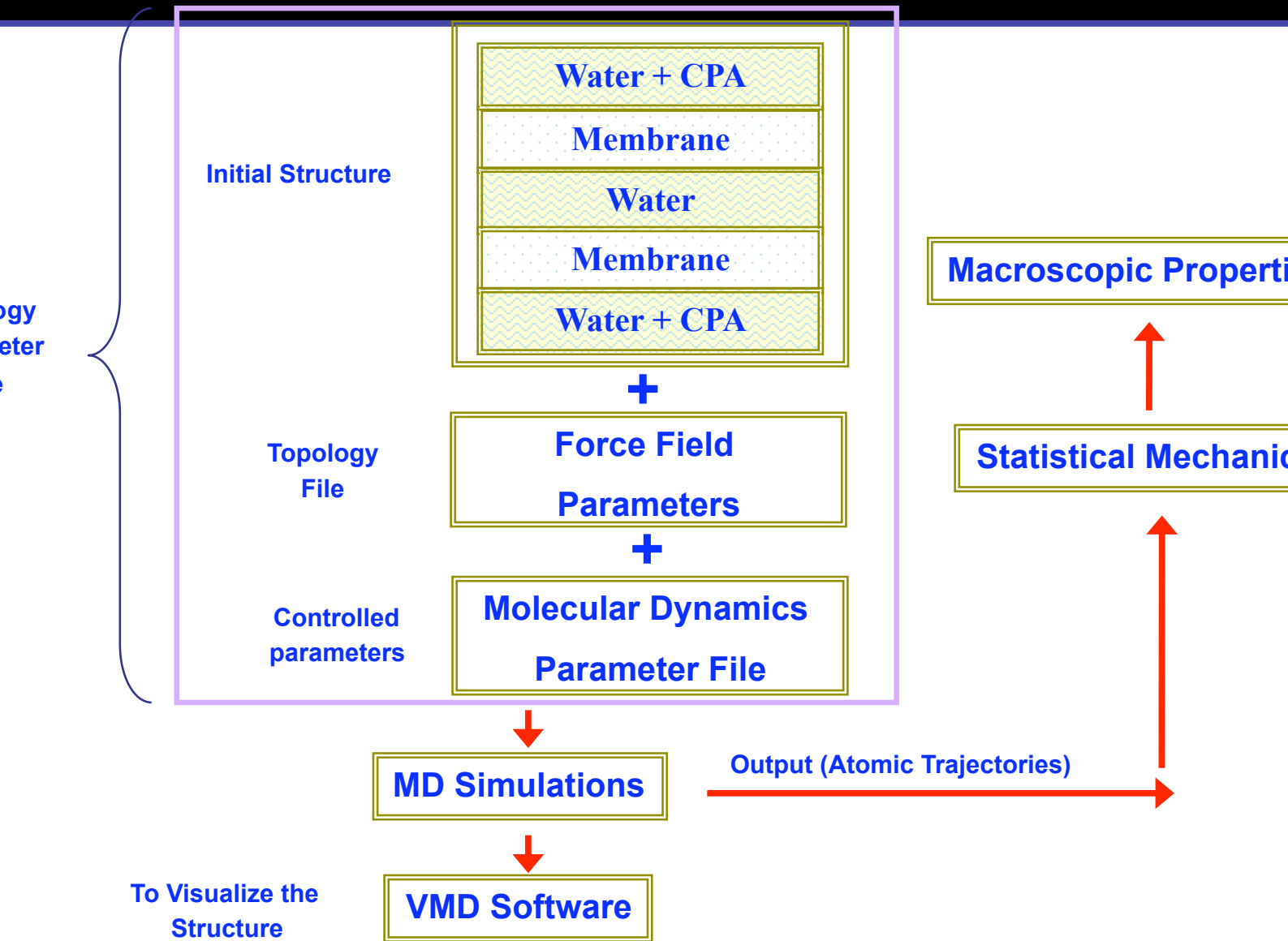


$$P = \frac{1}{\int_{z_1}^{z_2} \frac{\exp(\Delta G(z) / RT)}{D(z)} dz}$$

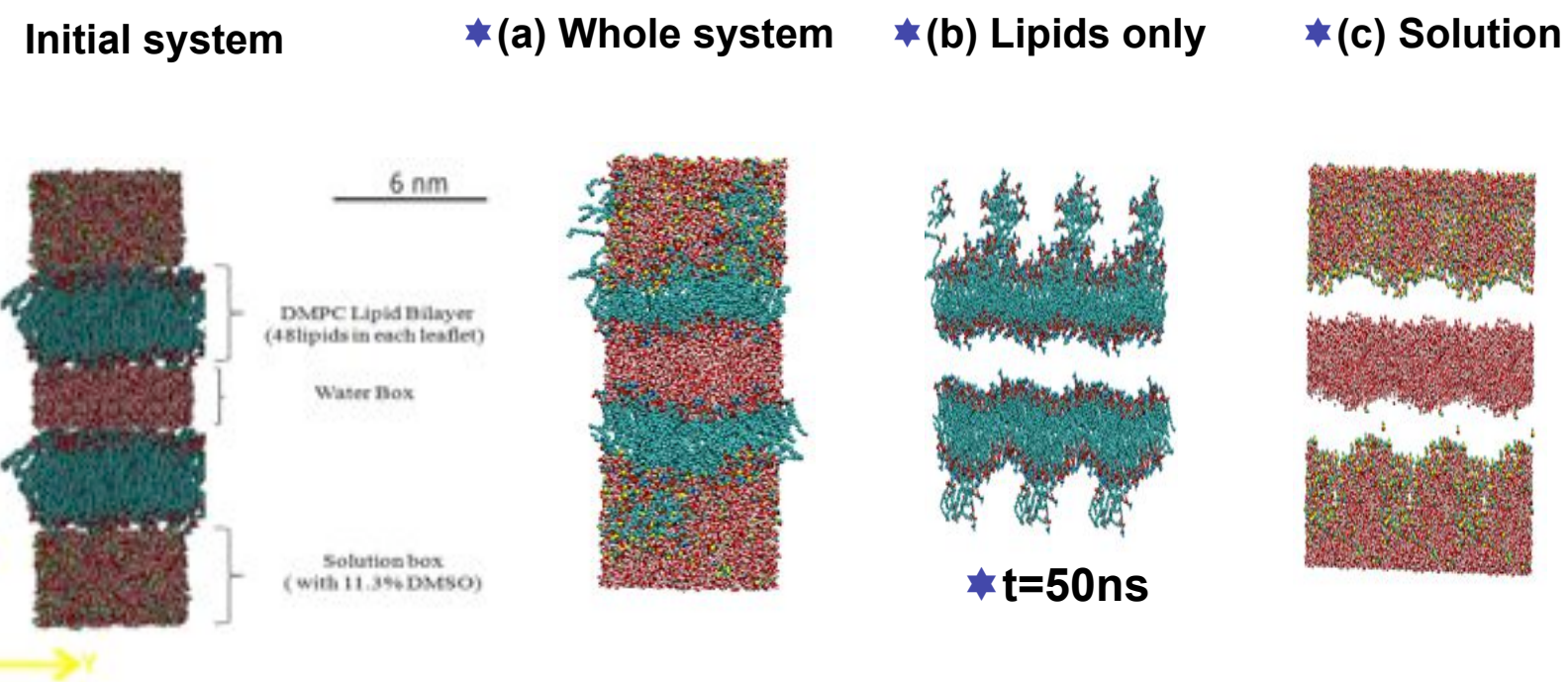
$$D = \frac{1}{3} \int_0^{\infty} \langle v(0) \cdot v(t) \rangle dt$$



# MD Simulation Using "GROMACS"

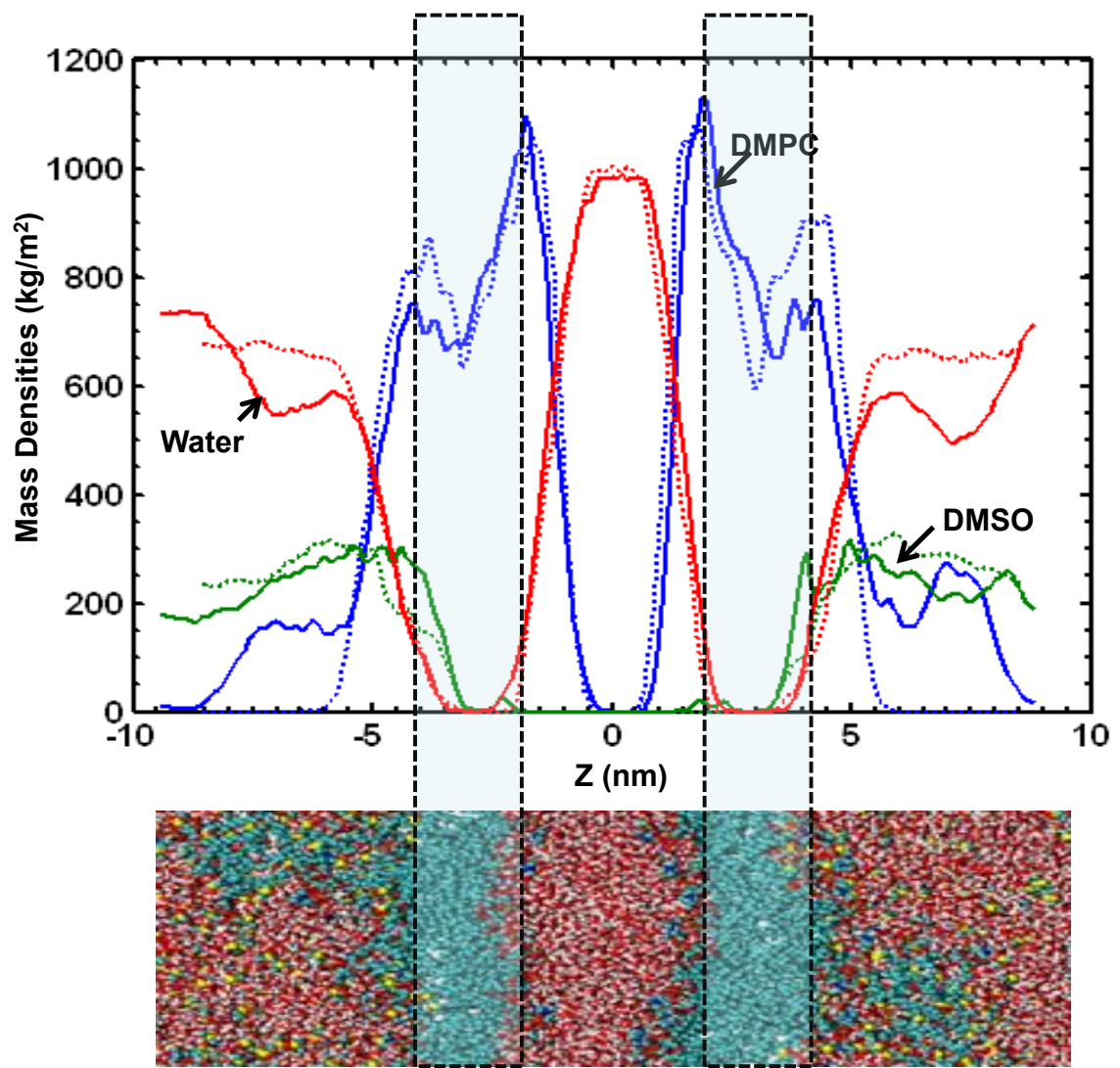


# Structural changes in Lipid Bilayers



- ◆ No penetration of water molecules
- ◆ Data analyzed for mass density profiles, radial distribution functions, tail order parameters, and water orientation profile

# Mass density profiles of : DMPC, DMSO, and wa



0ns profiles: dotted line ,50ns profiles: solid line

## CFD

- ✓ Improvements to the IBM (pressure interpolation)
- ✓ Working on the MPM for greater robustness (implicit, parallel)
- ✓ Simulation of transport in flexible tubes

## MD

- ✓ Simulation of small molecules across lipid bi-layers

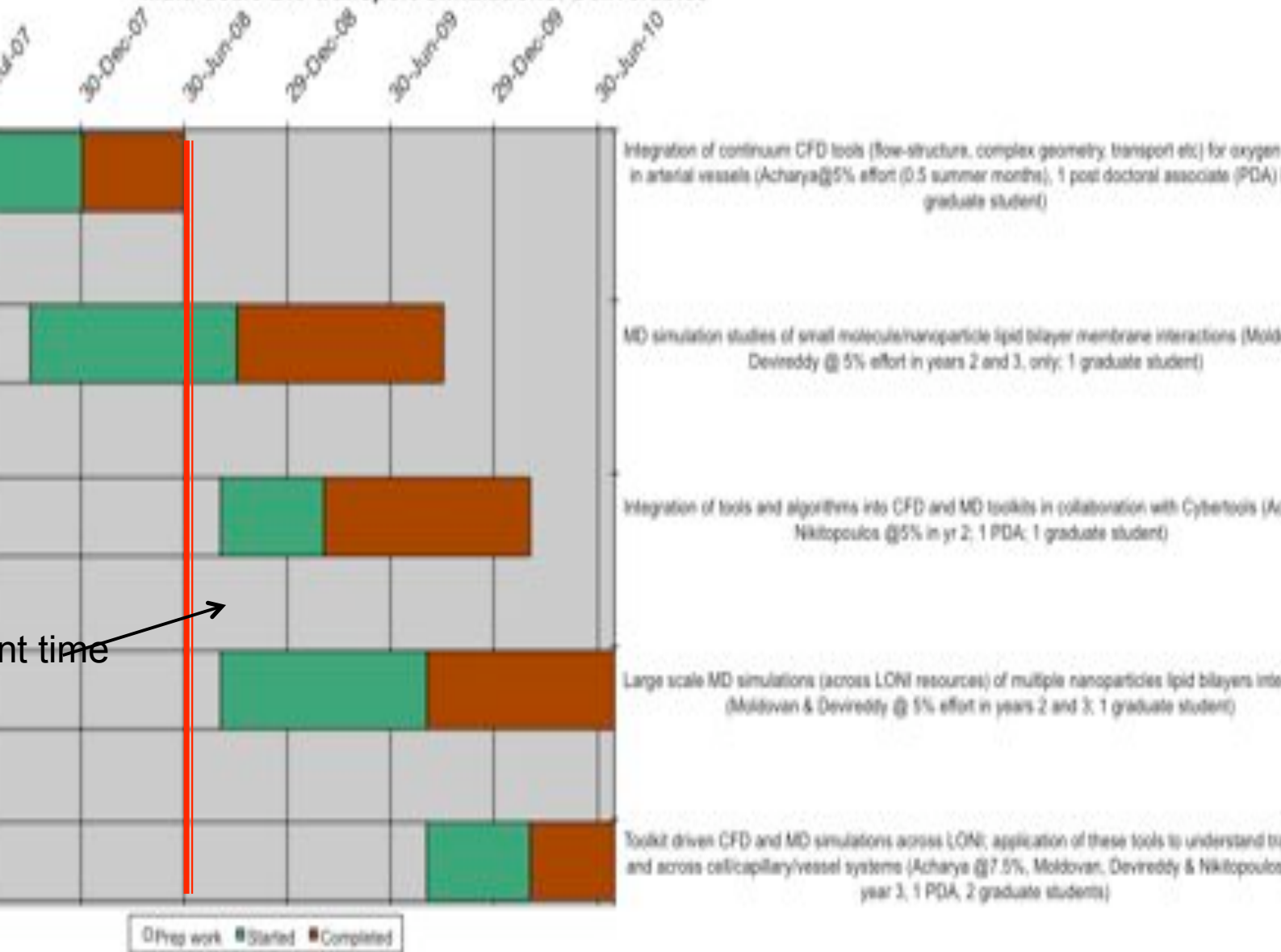
## Collaboration with WP4

- ✓ Regular meetings with the WP4 team
- ✓ Development of a simplified CFD code with data array structure consistent with Cactus for implementation as part of the CFD Toolkit

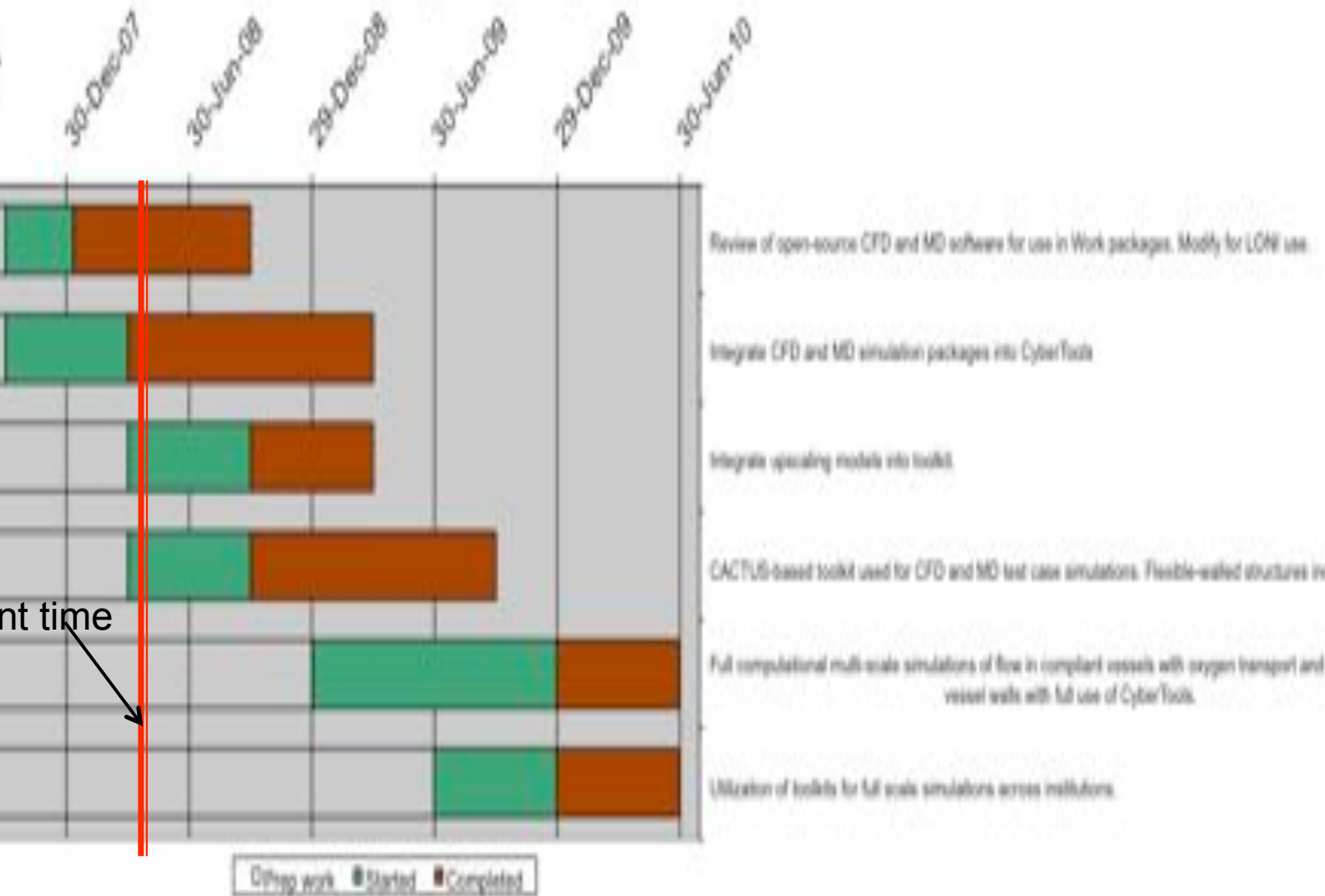
## CFD-MD Coupling

- ✓ Discussion on coupling strategy and approaches

### Multi-scale Bio-transport Simulations: Deliverables



## Bio-Transport & Cybertools Co-ordination



Development of improved CFD methodologies for biological systems (complex geometries, moving boundaries, multi-scale phenomena)

Utilization of CFD and MD methodologies for improved understanding of transport processes in biological systems

Supporting the development of Toolkit infrastructure for open source, scalable community usage

CFD-MD integration for resolving/integrating atomistic effects

Future interactions will also include the visualization groups and the portals group