

Digital Blood in Massively Parallel CPU/GPU Systems for the Study of Platelets deposition

Kotsalos Christos

Supervisors: Prof. B. Chopard Dr. J. Latt

Scientific and Parallel Computing Group (SPC)

High fidelity and fast simulations of <u>deformable blood cells</u> using a combined Finite Element Immersed Boundary Lattice Boltzmann method (FE-IB-LBM)



FEM for the blood cells

LBM for the blood plasma

IBM for the Fluid-Solid Interaction

CPU/ GPU images from flaticon.com







Normal Artery

Diseased Artery

Thrombus formation

Why to simulate at this level?



Prof. Bagchi Group | Drug delivery



Malaria







Normal capillary



Sickle Cell Anemia

Blood in numbers

The average adult has a blood volume of roughly 5 liters = $5 \times 10^{6} \text{ mm}^{3}$

Huge amount of deformable bodies: <u>fast</u> <u>simulations</u> <u>are one-way</u>

- 1 mm³ of blood contains:
- O(10⁶) red blood cells
- O(10⁵) platelets
- O(10³) white blood cells



Wikipedia: Blood

Plasma is the liquid component of blood occupying almost 55% of the whole blood. It is a Newtonian fluid **just like water**.

40-45% of the whole blood volume is occupied by red blood cells (RBCs). This volume fraction is called **hematocrit**.

Numerical methods:



Mass-lumped FEM (npFEM) for the deformable bodies (developed in C++/ CUDA)



Lattice Boltzmann method (LBM) for the blood plasma (Palabos in C++/ MPI)



Immersed boundary method (IBM) for the coupling (Palabos in C++/ MPI)

<u>www.palabos.org</u> : Parallel Lattice Boltzmann Solver open-source CFD solver (by SPC Lab, UniGe)

Mass-lumped FEM (nodal projective FEM)



Implicit Euler time integration: update rule

$$egin{aligned} egin{aligned} egi$$

Newton's 2nd law per vertex

Variational Implicit Euler formulation

 $C = \alpha_D \mathbf{M} + \beta_D \widetilde{\mathbf{H}}$

$$\min_{\boldsymbol{x}_{n+1}} \frac{1}{2h^2} \left\| \widetilde{\mathbf{M}}^{\frac{1}{2}} (\boldsymbol{x}_{n+1} - \boldsymbol{y}_n) \right\|_F^2 + \sum_i E_i(\boldsymbol{x}_{n+1})$$

Mass-lumped FEM (npFEM)

$$\min_{\boldsymbol{x}_{n+1}} g(\boldsymbol{x}_{n+1}) = \frac{1}{2h^2} \left\| \widetilde{\mathbf{M}}^{\frac{1}{2}} (\boldsymbol{x}_{n+1} - \boldsymbol{y}_n) \right\|_F^2 + \sum_i E_i(\boldsymbol{x}_{n+1})$$

4 different kinds to describe a blood cell:

- Area Conservation
- Global Volume Conservation
- Bending rigidity
- Material model (modified Skalak)

Discretized by – piecewise linear basis functions

Quasi-Newton:
$$m{x}_{n+1}^{k+1} = m{x}_{n+1}^k - lpha \widetilde{\mathbf{H}}^{-1}
abla g(m{x}_{n+1}^k)$$

For more details:

Bridging the computational gap between mesoscopic and continuum modeling of red blood cells for fully resolved blood flow Journal of Computational Physics 2019 https://doi.org/10.1016/j.jcp.2019.108905

Stretching experiment



Optical tweezers experiment by Dao, Li & Suresh

Shear flow experiment: Tank-treading



Poiseuille flow: Parachute-like shape





Multiple blood cells simulations



How to split the load to the available CPUs & GPUs ?

Presentation of the Hybrid Version of our computational framework

Load balancing



Straightforward to partition a static homogeneous grid CPUs deal with grid points (LBM) & Lagrangian points (IBM)



The blood cells are distributed once at the beginning to the available GPUs

They can be spatially everywhere

MPI point-to-point communication

The fluid solver sends forces & collision data to the solid solver

The solid solver communicates the state at t+1

GPU acceleration



Fit bodies in the Streaming Multiprocessors to gain from the fast shared-memory and thread synchronization





1 SMX deals with 1 CUDA block per time

Genericity & modularity of our framework



Shear flow: >500 blood cells | Hematocrit 35% | Box 50x50x50 µm³





Focus on Pathological conditions, like Diabetes (swollen RBCs)





Performance measures



Weak Scaling





Profiling of npFEM

Reference case study Ht 35%, Box 50x50x50 μm³



RBCs:	476
PLTs:	95

1 GPU node - npFEM [msec]	11
1 CPU node - npFEM [msec]	70
Speedup of npFEM	6.4
5 CPU node - npFEM [msec]	14





Future plans:

- Port the Fluid solver (and/or FSI) on GPUs using CUDA
- Check scalability of our solver for multiple deformable bodies O(10⁶ 10⁹)
- More investigation of platelet deposition through in-vitro and in-silico experiments

1 mm³ of blood contains:

O(10⁶) RBCs





Thank you very much