

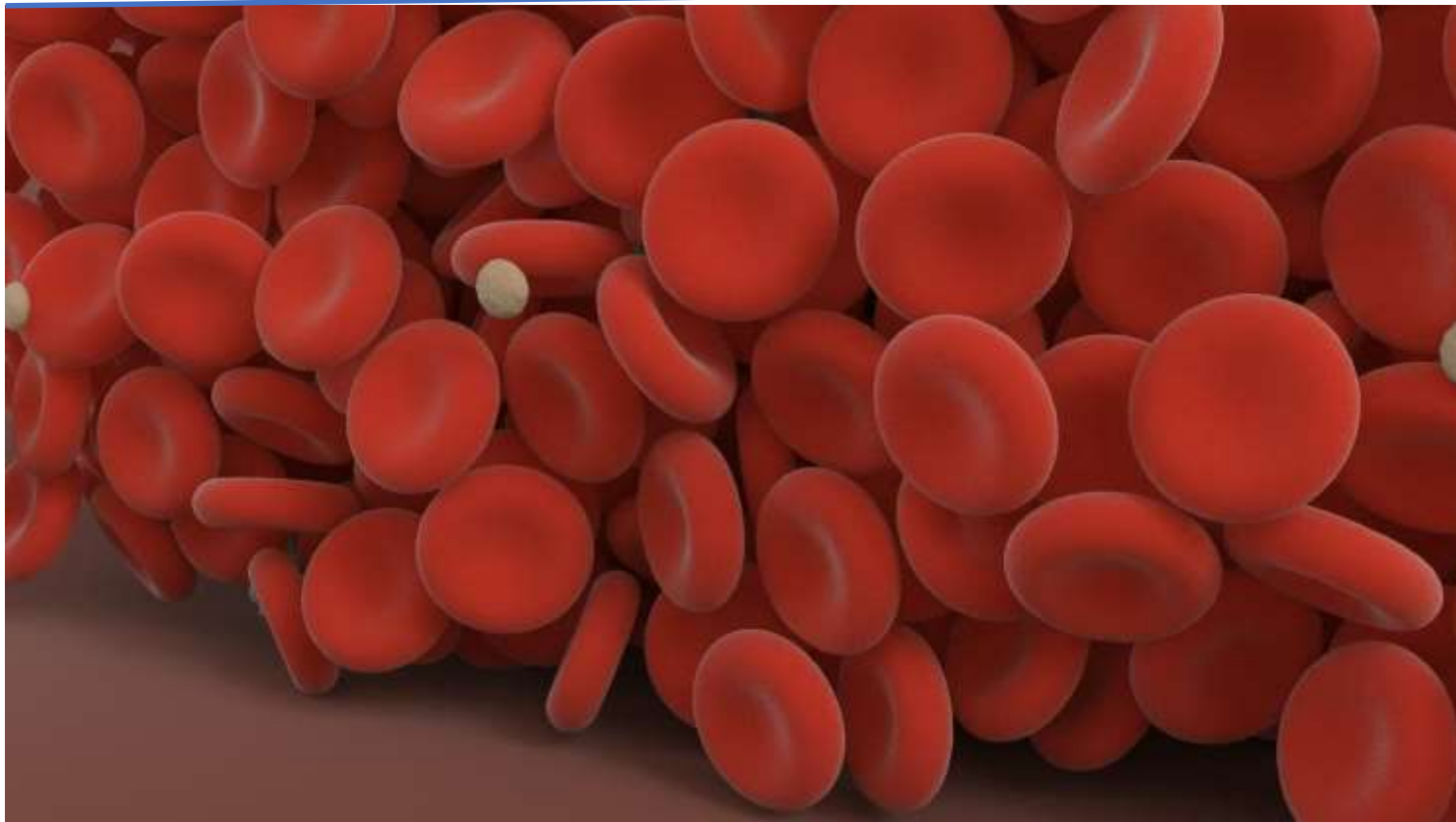
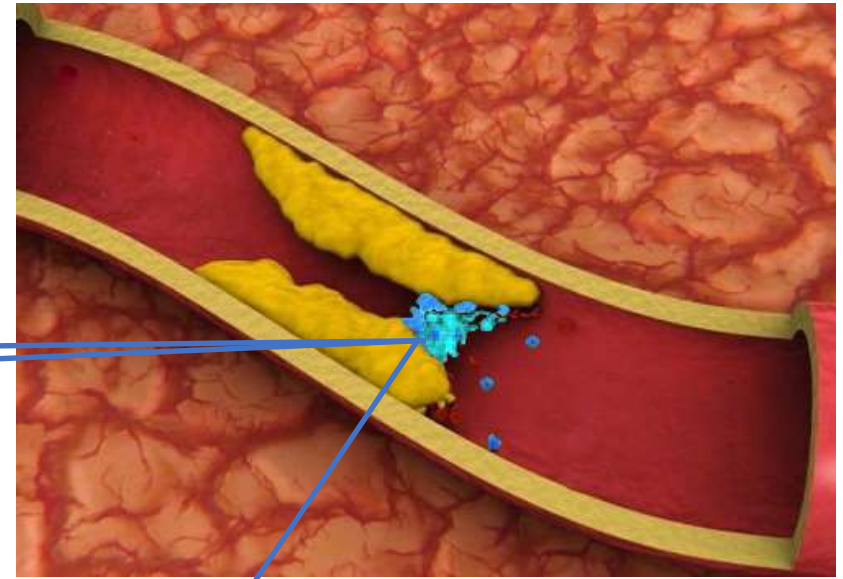
Multi-Scale Investigation of High Shear Rate Thrombus Formation

G. Zavodszky^{1,2}, B.J.M. van Rooij¹, A.G. Hoekstra¹

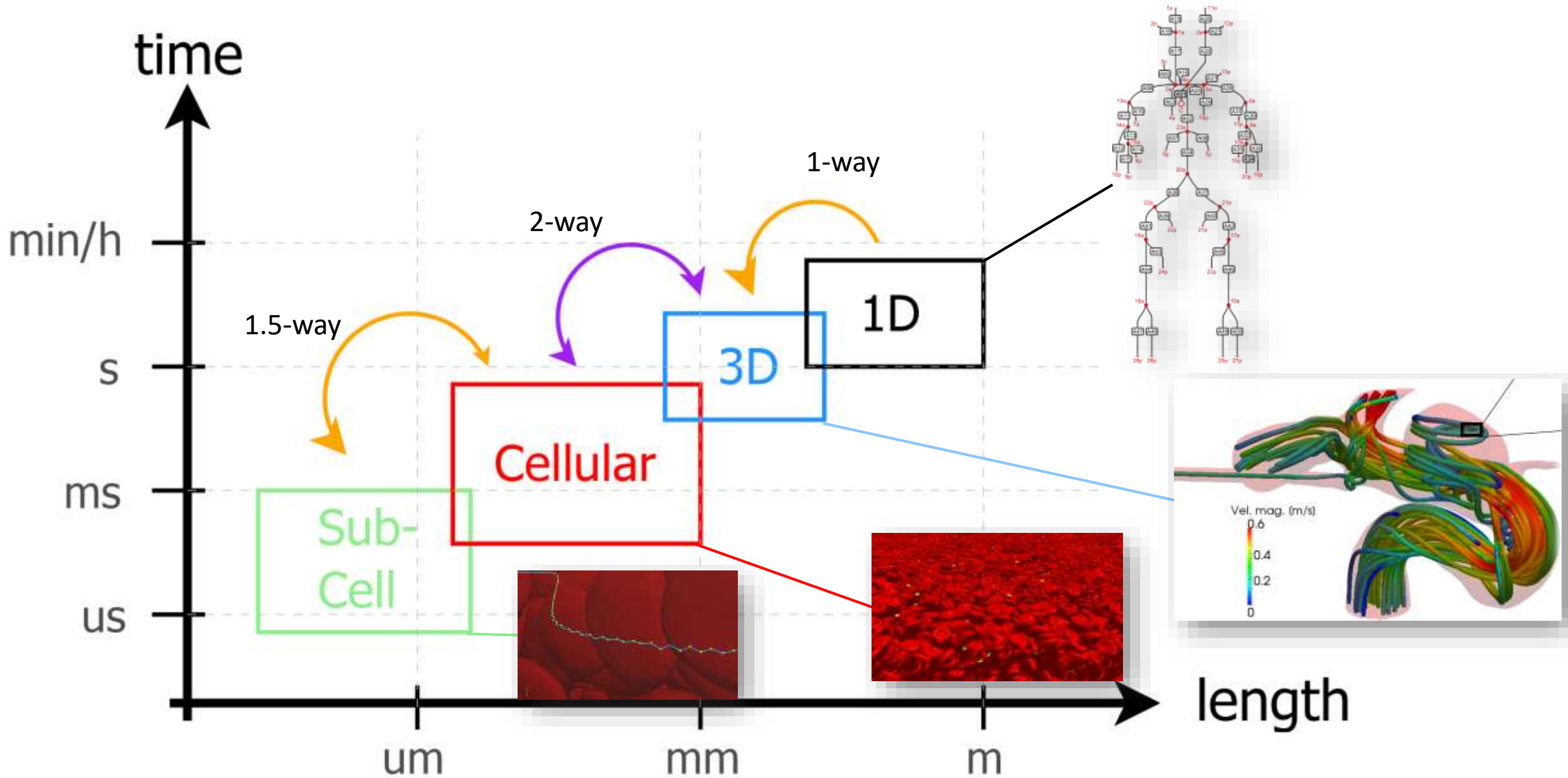
¹*Computational Science Lab, University of Amsterdam, The Netherlands*

²*Budapest University of Technology and Economics, Hungary*

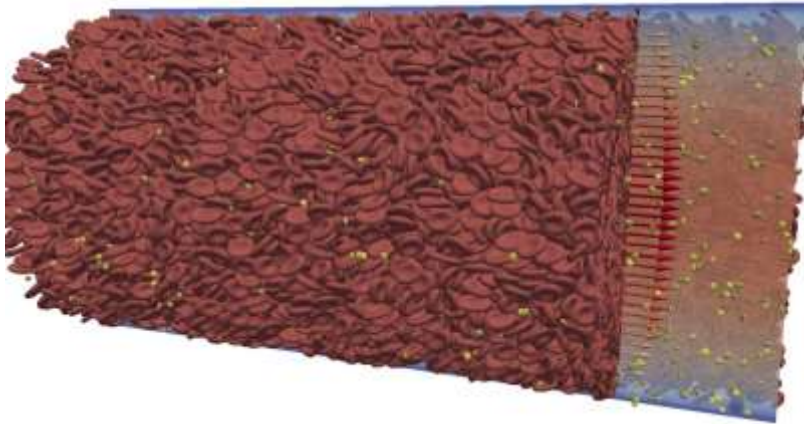
High shear-rate thrombus formation



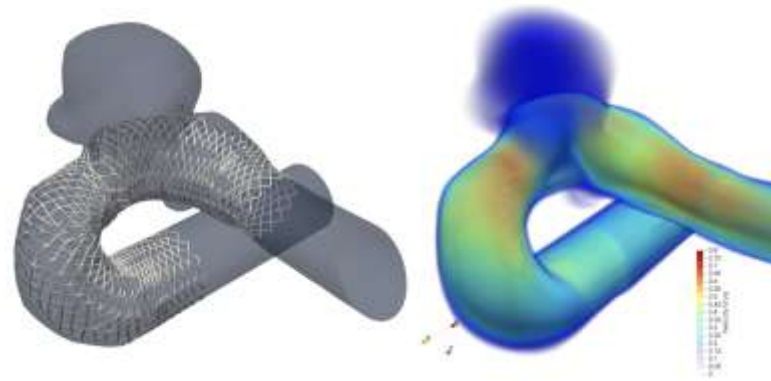
Scale-separation map



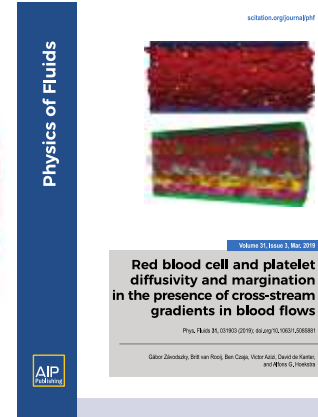
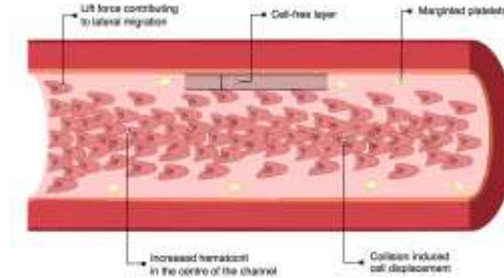
HemoCell *cellular*



HemoFlow *macroscopic*



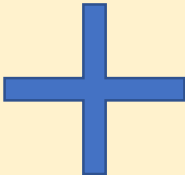
rheoModel *coarse-grained*



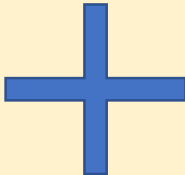
Závodszy, et al. (2017) *Frontiers in Physiology*, 8, 563
 Závodszy, et al. (2017) *Procedia Comput Sci.* 108, 159.
 Alowayyed et al. (2018) *Journal of Comp. Sci.*, 24, 1-7.
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 Mountrakis et al. (2016) *EPL (Europhysics Letters)* 114.1: 14002.
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Závodszy, et al.. (2013). *Int. J. of Heat and Fluid Flow*, 44, 276
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 Csippa, et al. (2018). *Computers in biology and medicine* 103: 244-251.
 Závodszy, et al. (2019). *Int. J. for Num. M. in Bio. Eng.*, under review

Závodszy, et al. (2019) *Physics of Fluids*, 31 (3), 031903



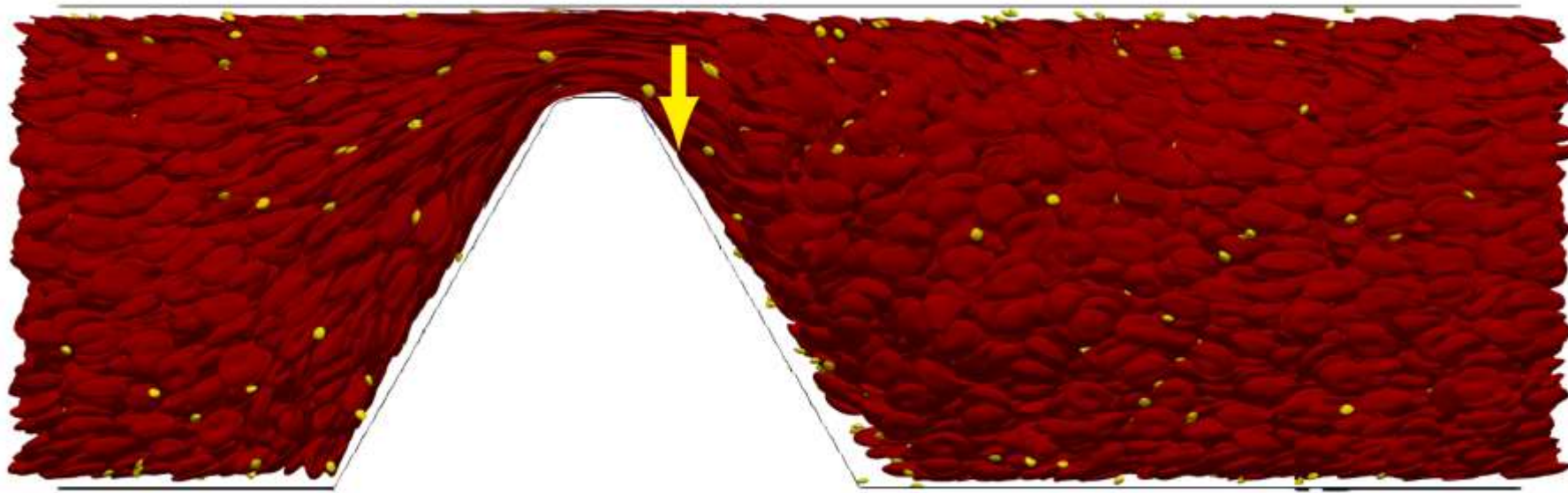
vWF model
Under development



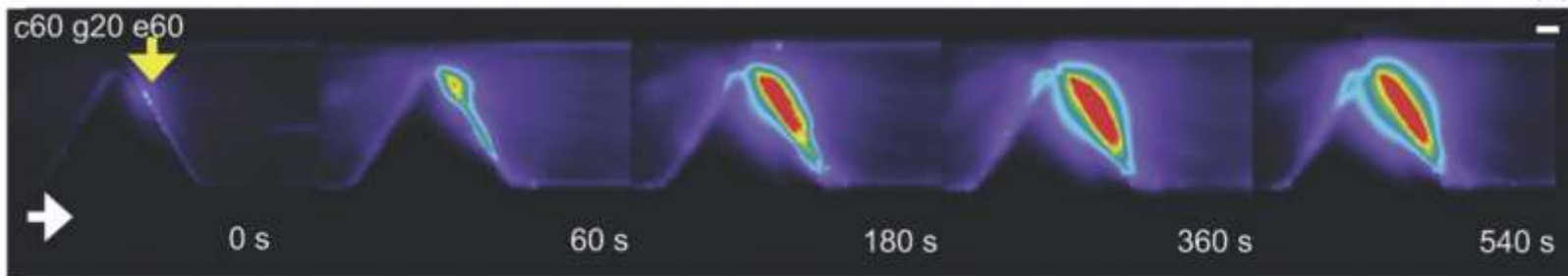
CEPAC
Under development

Cellular modeling – shear conditions

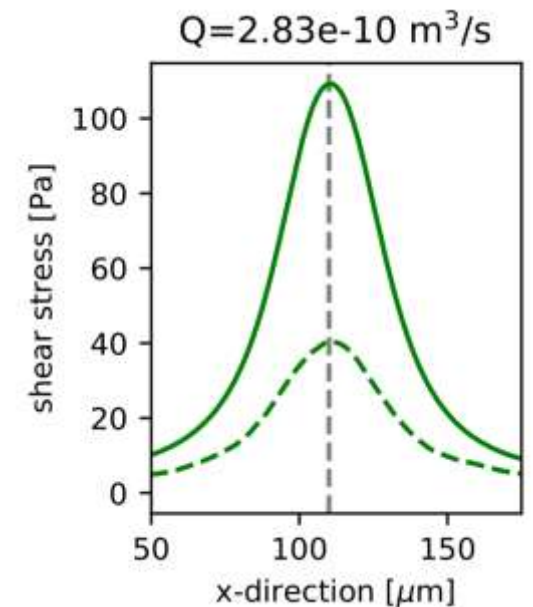
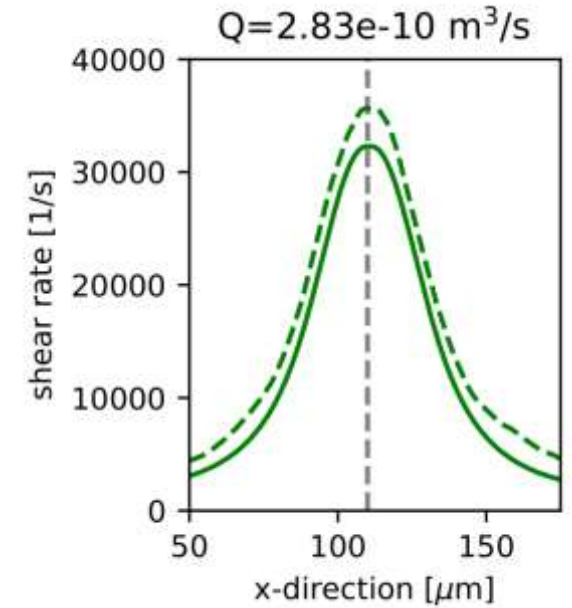
vs. continuum flow model



van Rooij, et al. (2019) J. Royal Society Interface, accepted



Tovar-Lopez and Nesbitt et al., Lab Chip, 2010, 10, 291-302



Cellular modeling - platelet margination

$Re \sim 1.0$

$V_{max} = 0.5 \text{ cm/s}$

$D = 50 \text{ } \mu\text{m}$

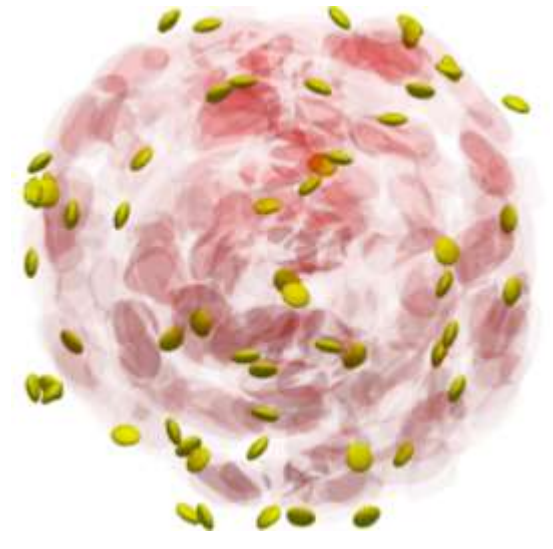
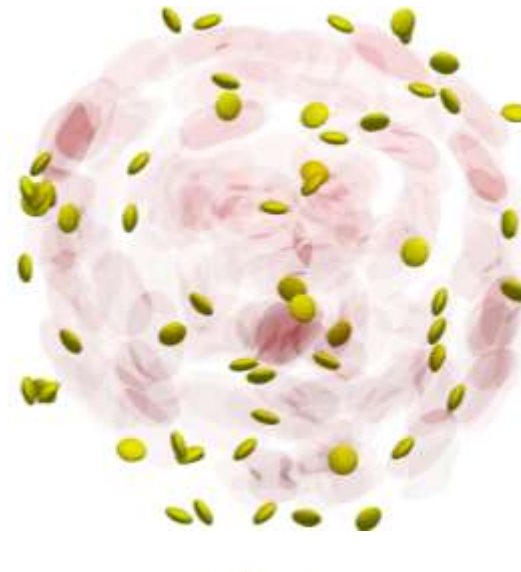
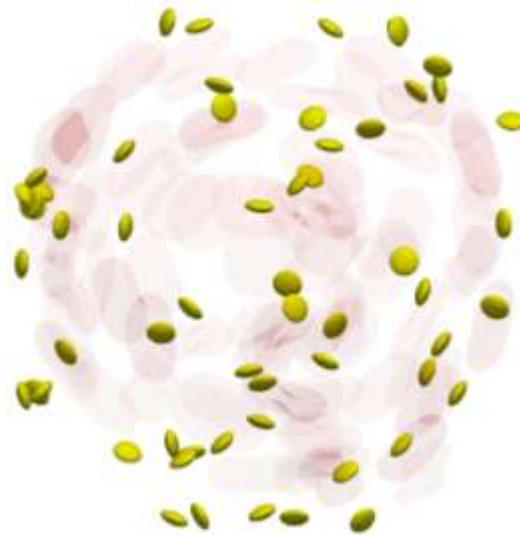
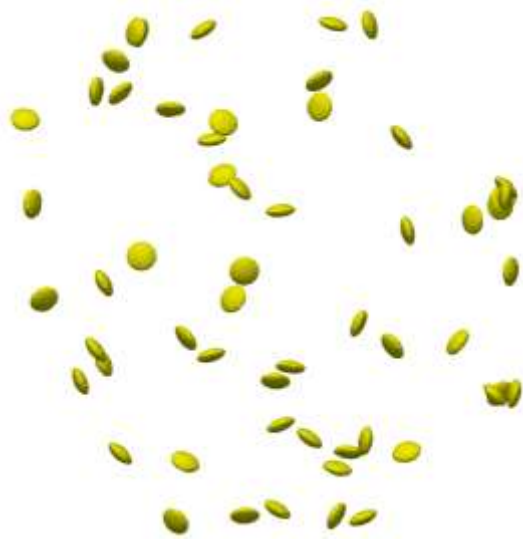
H = 0 %

H = 5 %

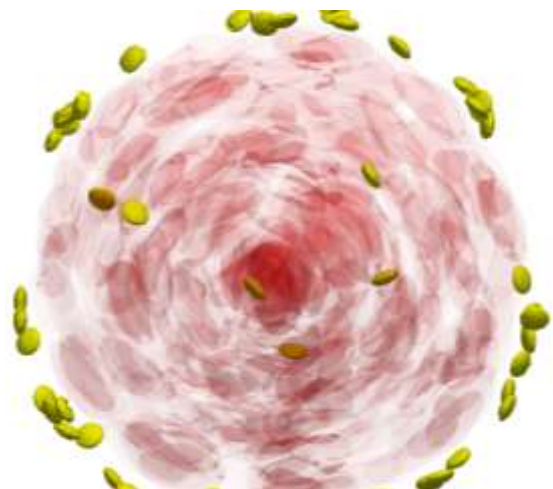
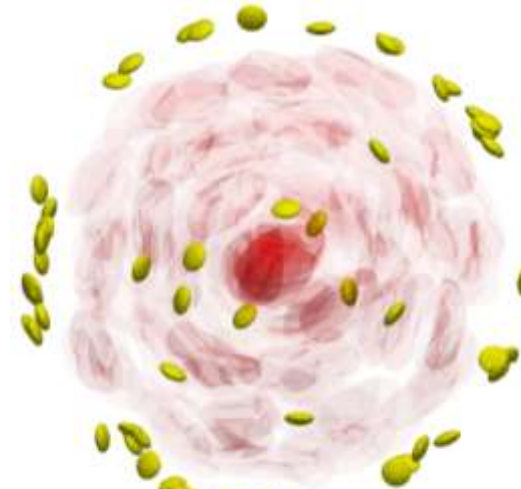
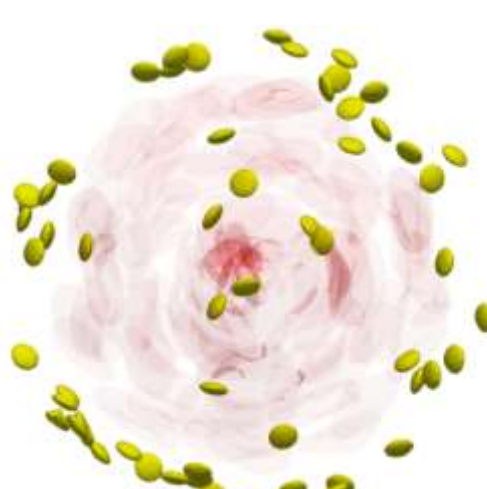
H = 10 %

H = 20 %

t = 0 s



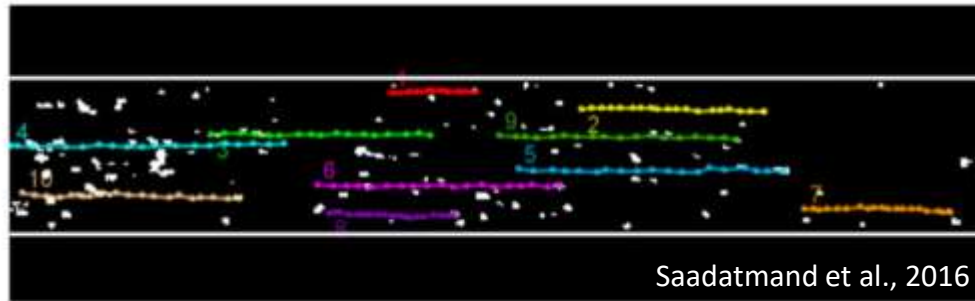
t = 1.3 s



Margination from cell pair collisions

Experiment:

- Following both RBC and small tracer particle trajectories
- Straight channel flow



Experiment:

$$D_{rr_RBC} = 1.2e-8 \text{ cm}^2/\text{s}$$

Simulation (using the same parameters) with [HemoCell](#):



HemoCell:

$$D_{rr_RBC} = 1.15e-8 \text{ cm}^2/\text{s}$$

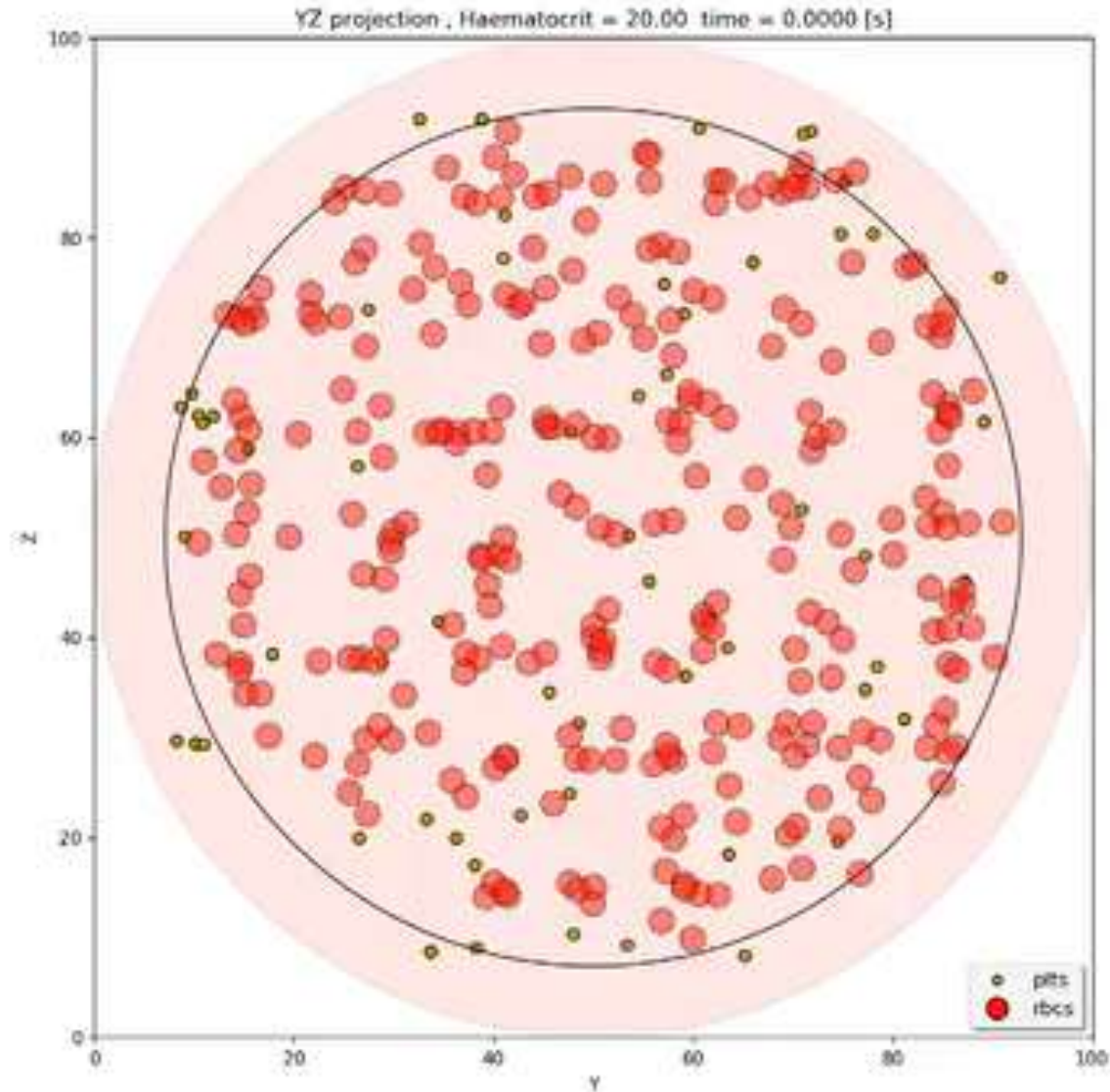
Cell radial dispersion:

$$D_{rr} = \frac{1}{N} \sum_{i=1}^N \frac{\langle (R_{i,r}(t) - R_{i,r}(0))^2 \rangle}{2t}$$

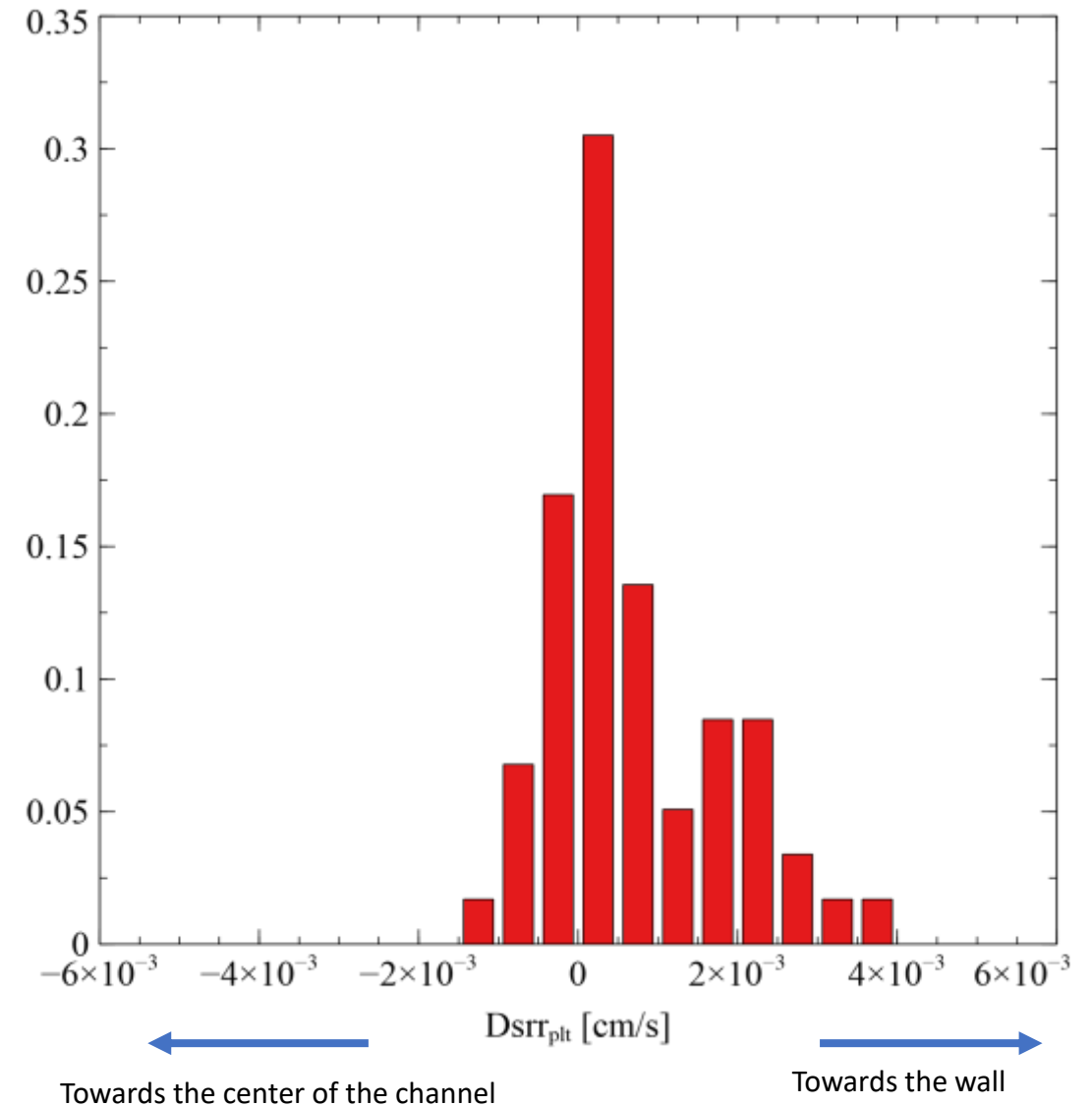
Cell radial drift velocity:

$$D_{srr} = \frac{1}{N} \sum_{i=1}^N \frac{\langle R_{i,r}(t) - R_{i,r}(0) \rangle}{t}$$

Preferential diffusivity

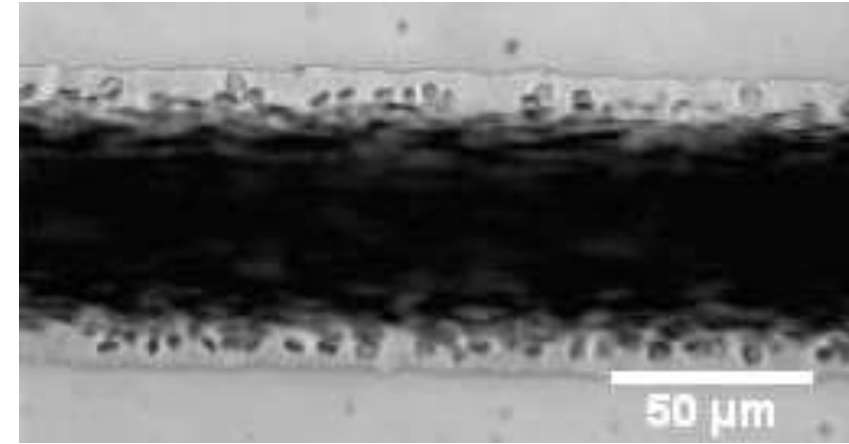
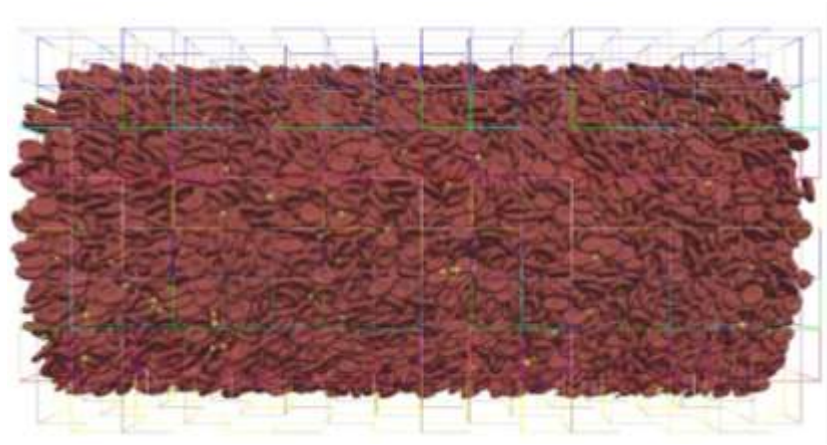


Video: David de Kanter



Coarse-grained model from statistics

22 x

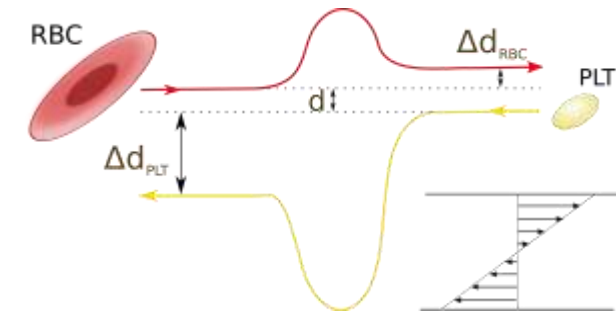


$D = 50..70 \text{ } \mu\text{m}$
 $H = 7\% \dots 36\%$
 $Re = 0,04 \dots 2,0$



Extract various parameters
in a statistical sense in 242
points.

Závodszy, et al. (2019) Physics of Fluids



$$\frac{\partial \Phi}{\partial t} = -\frac{\partial}{\partial r} \left[V(r)\Phi - D(r) \frac{\partial \Phi}{\partial r} \right]$$

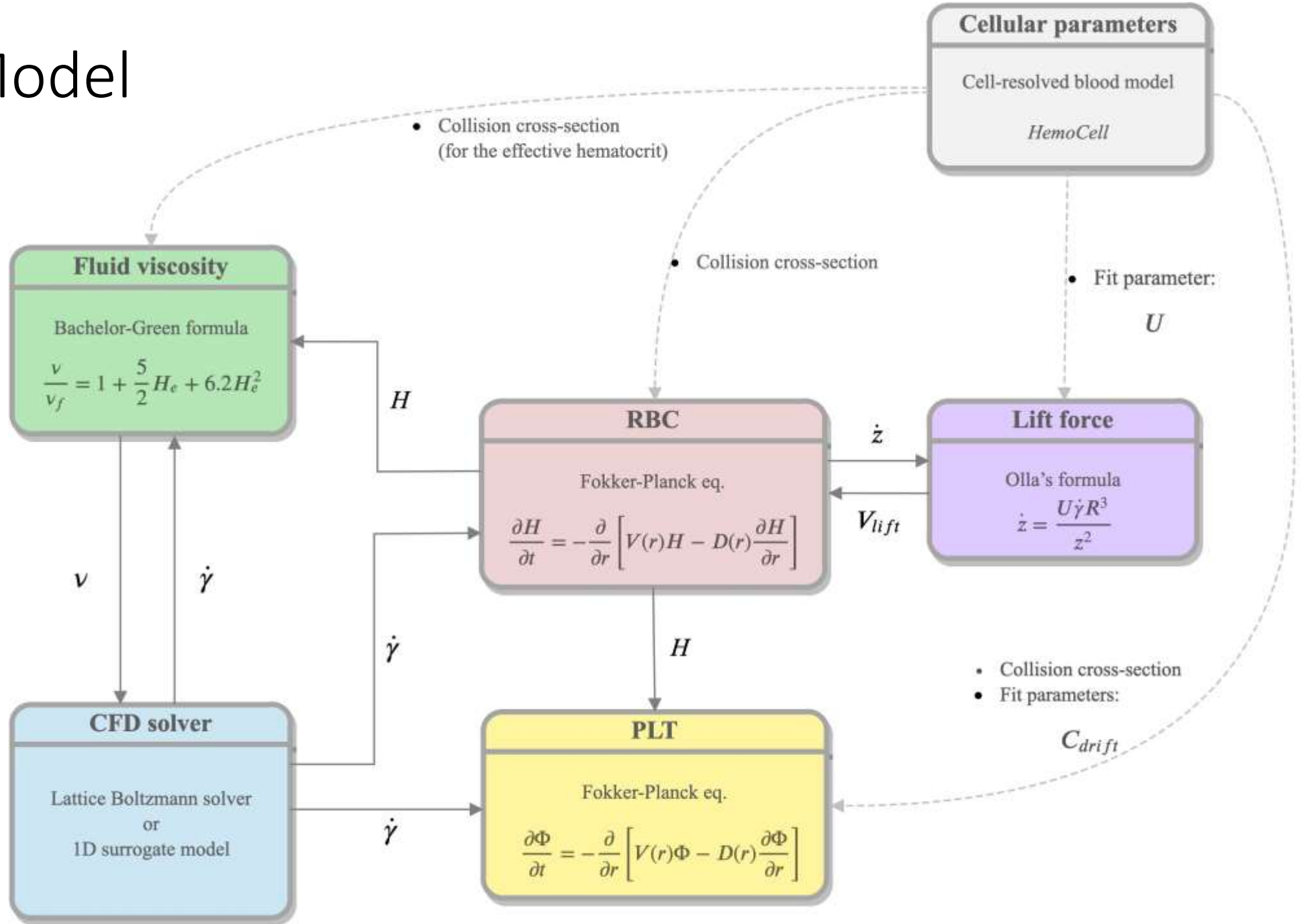
$$V_{lift} = -\frac{U\dot{\gamma}R^3}{z^2}$$

$$V_{asym} = -C * a(\dot{\gamma})^2 * \dot{\gamma} * \frac{\partial H}{\partial r}$$

$$D_{self} = C * a(\dot{\gamma})^2 * \dot{\gamma} * H$$

$$D_{saturation} = C * H^2$$

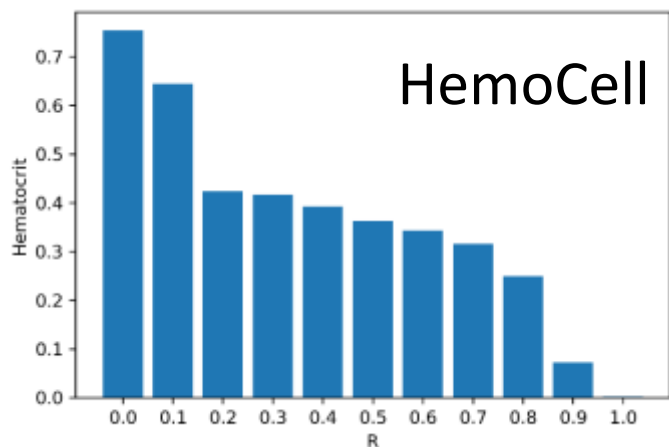
rheoModel



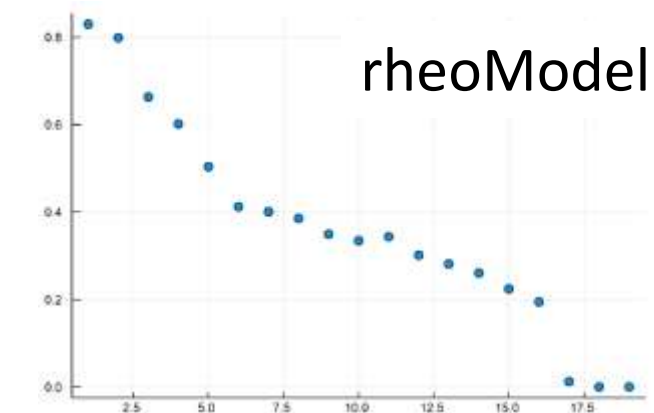
rheoModel vs. experiments

Radial distribution of cells in a vessel

($D=50\mu\text{m}$, $H=36\%$, $Re=1.5$, $t=1\text{ s}$)

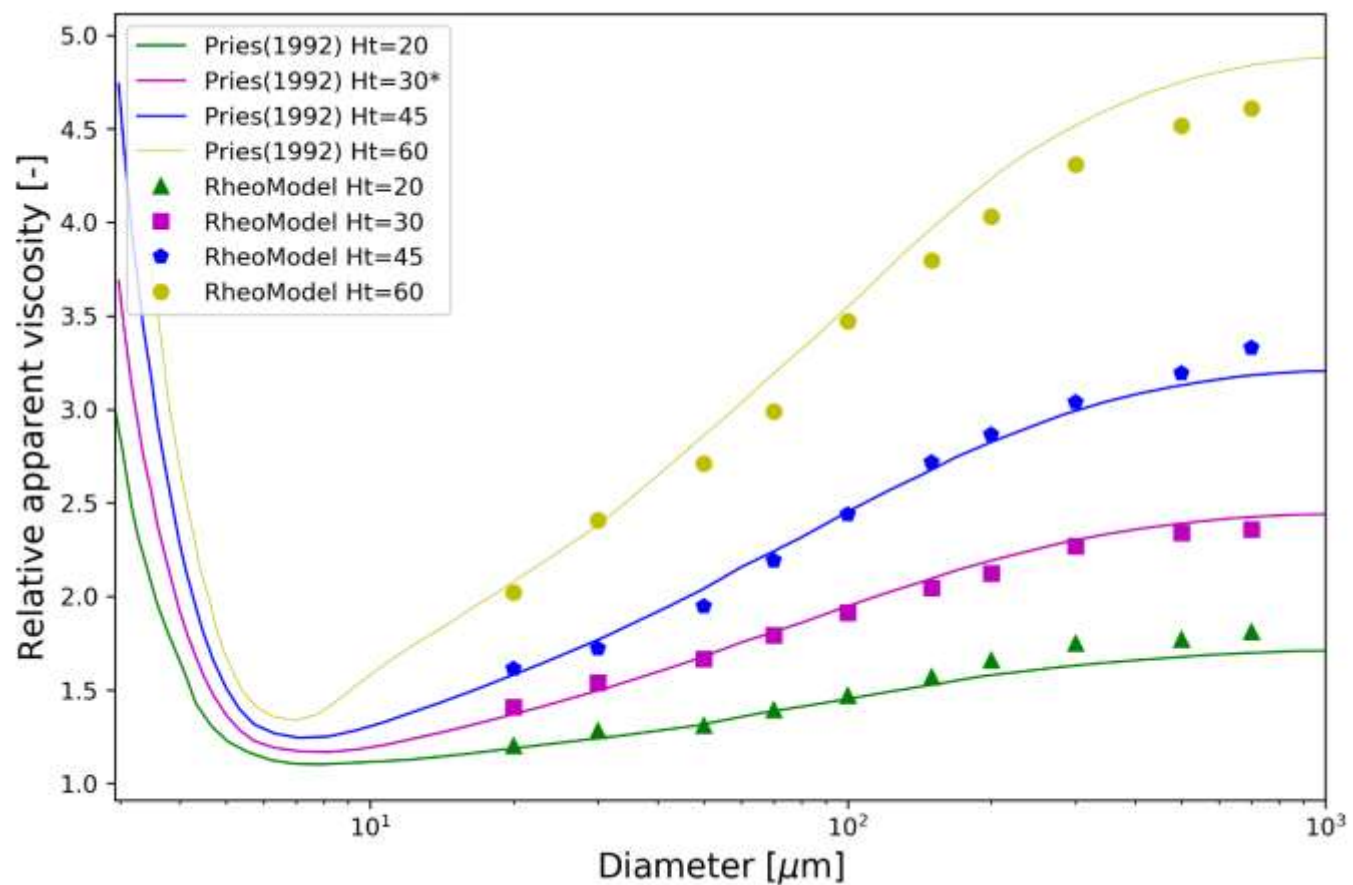


Computation: 9 days on 128 cores (Cartesius)

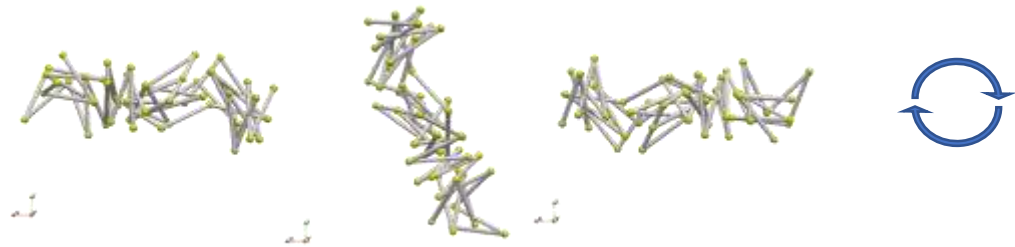


Computation: ~5 s on my notebook

Viscosity of blood in a vessel section



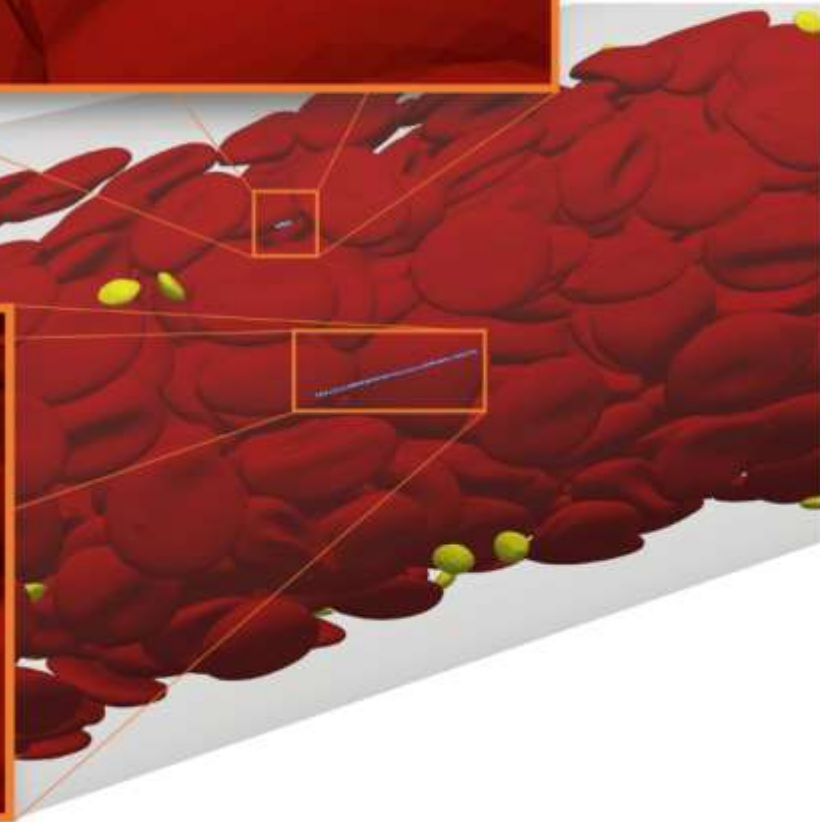
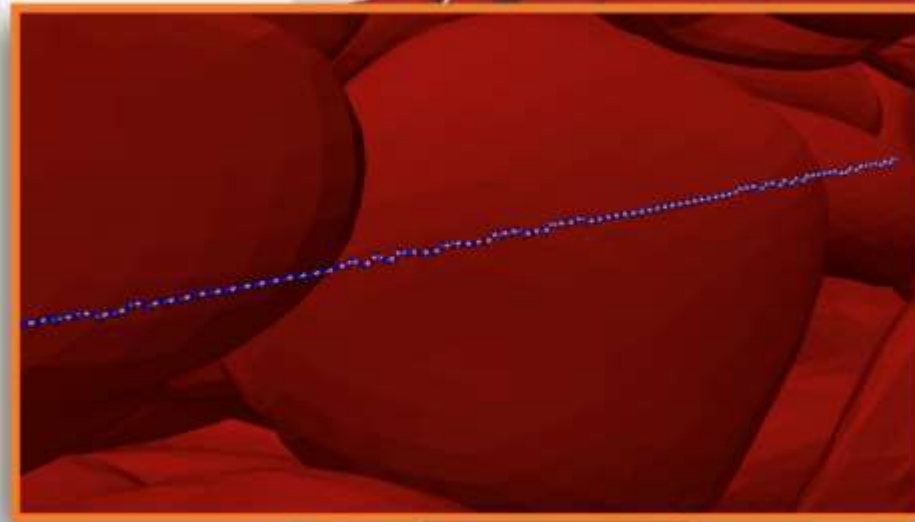
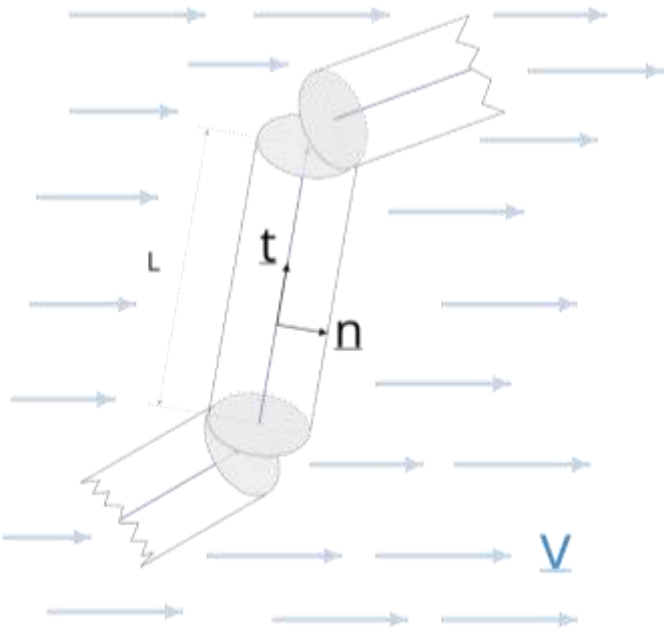
Low shear ($< 2000 \text{ s}^{-1}$)
Tumbling



Medium shear ($2000 - 6000 \text{ s}^{-1}$)
Cyclic uncoiling-recoiling

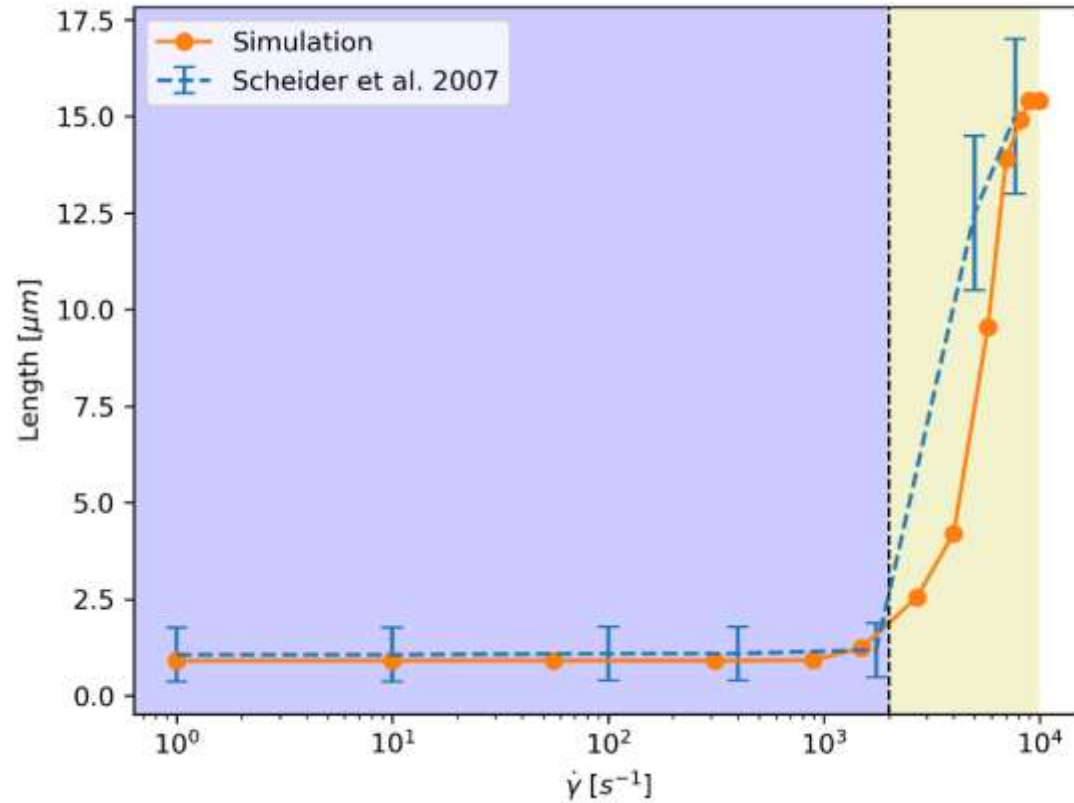


High shear ($> 6000 \text{ s}^{-1}$)
Stable uncoiled state



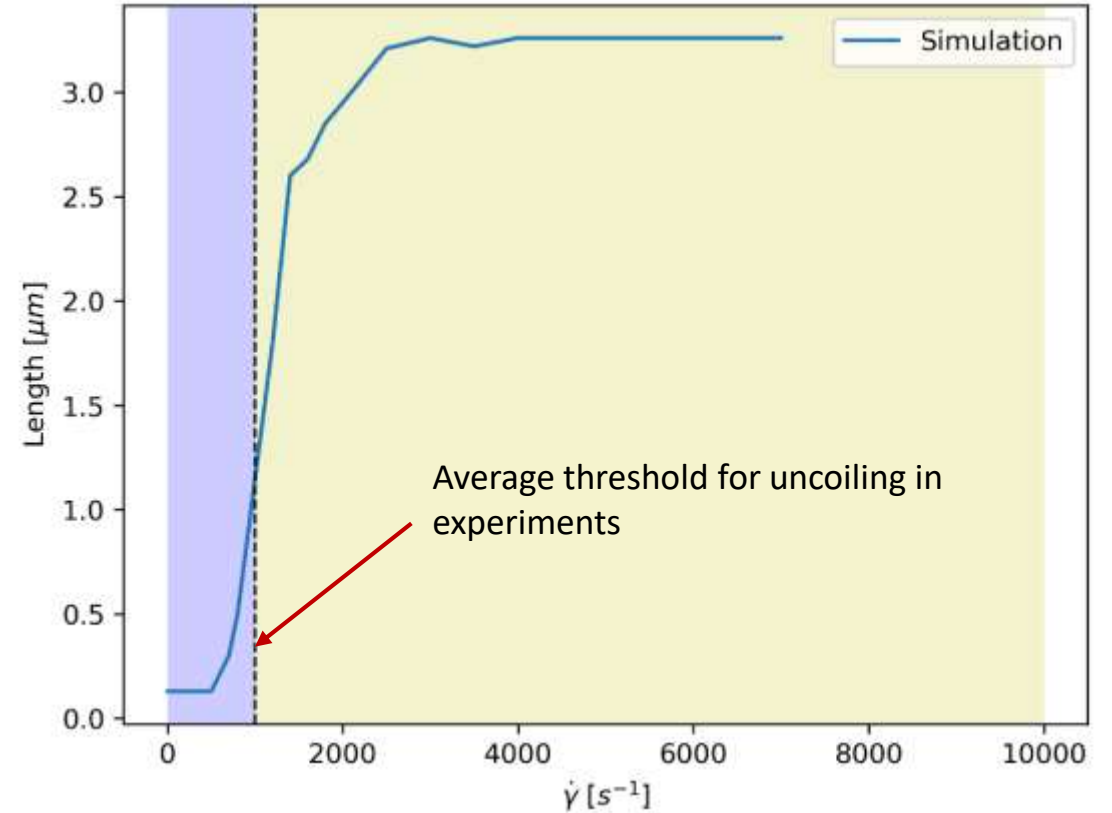
vWF mechanical model

Uncoiling in pure shear flow



Comparison to: Schneider et al. (2007),
PNAS

Uncoiling in shear flow while head fixed to wall

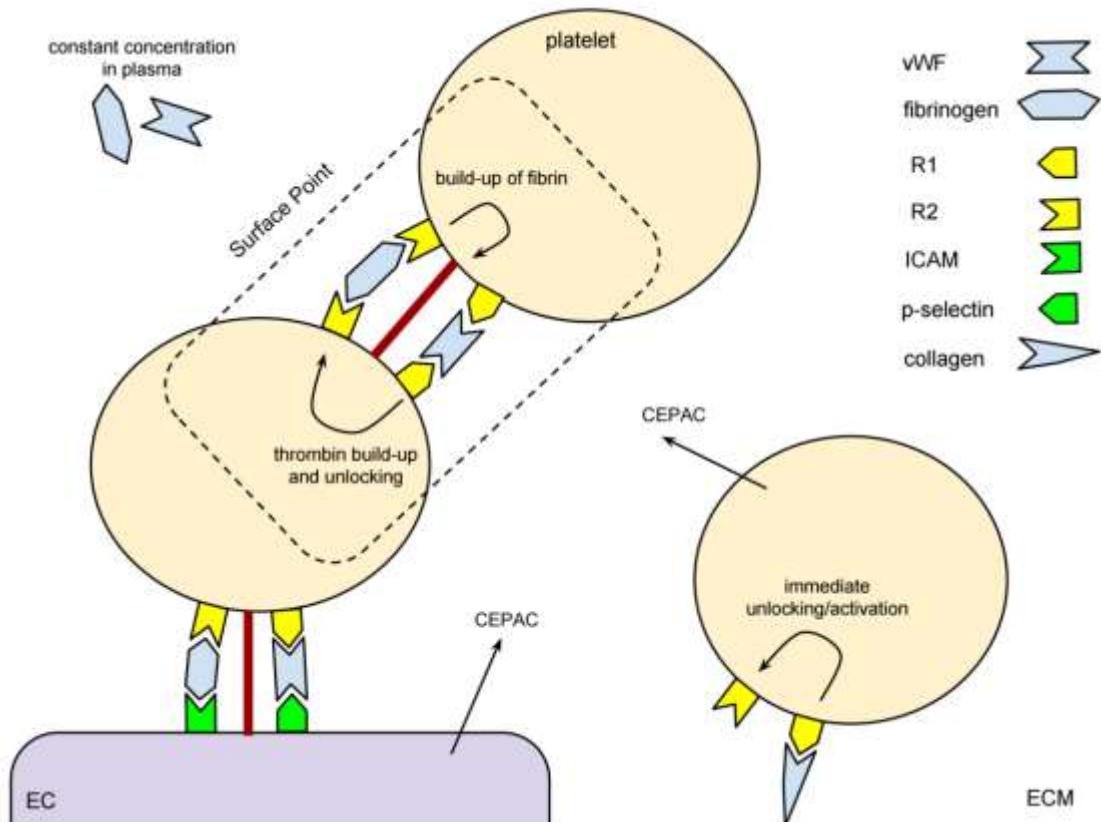


Comparison to: Fu et al. (2017),
Nature Communications

Note: the spherical chain representation cannot match both with the same parameters.

Conclusion: allowing the elements of the chain to change alignment compared to the flow might be important.

CEPAC - Combined Effect for Pro- and Anti-coagulants



R1 receptor → weak bonds

- Platelet receptor to vWF: Gp Ib/Ia [~25000 on a platelet]
- Integrin receptor to vWF: α IIb β 3 [~50000 on a platelet]
- Platelet receptor to collagen: Gp VI (+Gp Ia/IIa) [~4000 on platelet]

↓
Platelet adhesion and rolling over surface

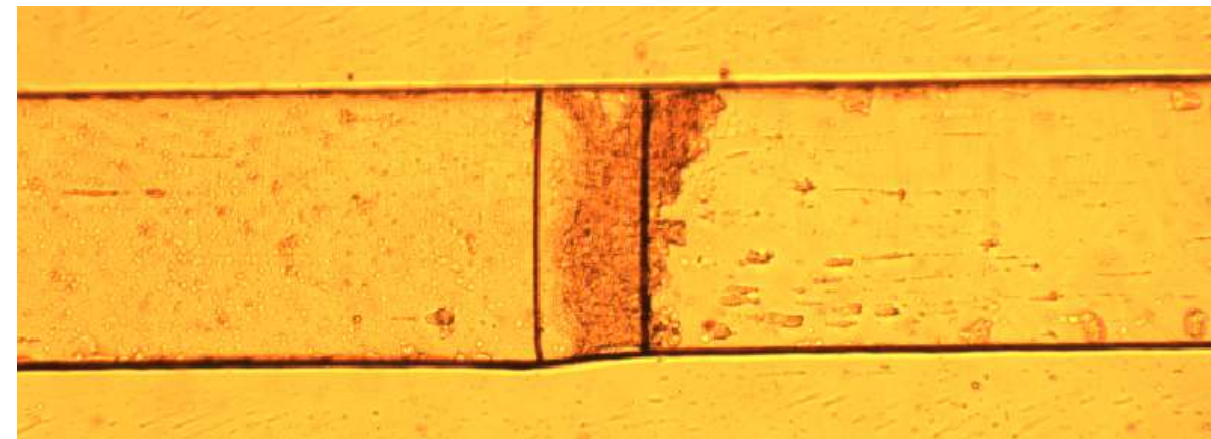
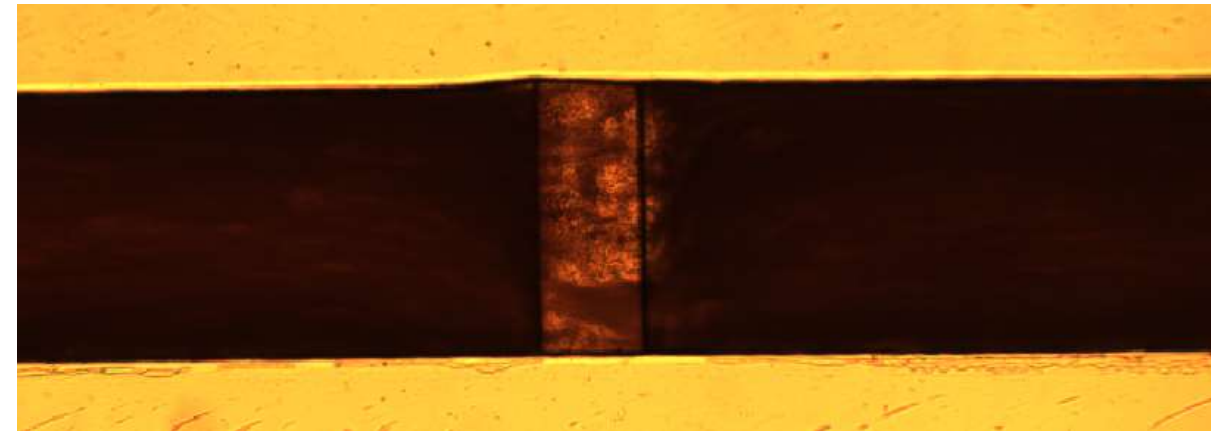
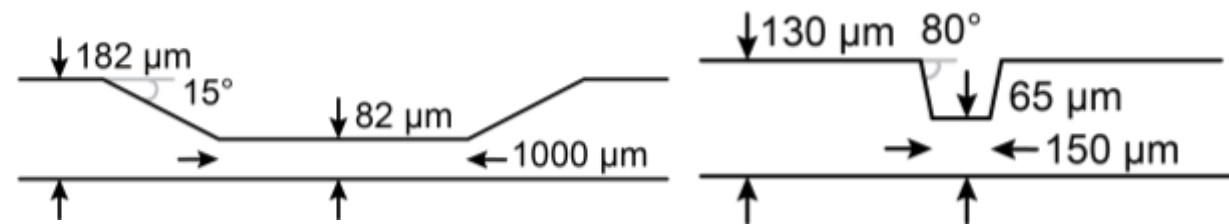
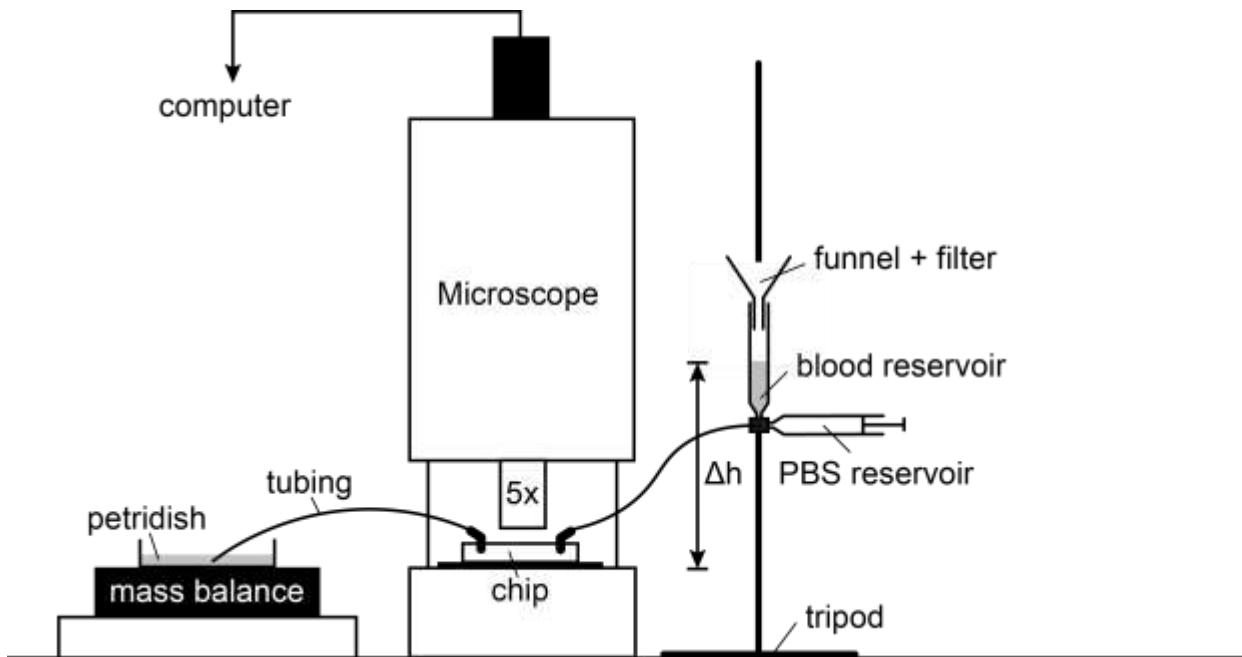
More important in arteries (>500 s⁻¹) and arterioles (1000-1800 s⁻¹) than in veins (<200 s⁻¹)

R2 receptor → strong bonds

- Integrin receptor to fibrinogen: α IIb β 3 [~50000 on a platelet]
- Integrin receptor to collagen via vWF: α 2 β 1 [~1000-4000 on a platelet]

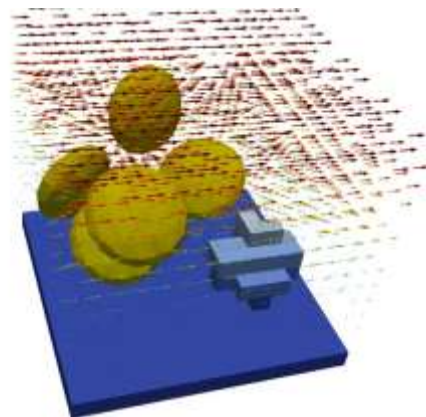
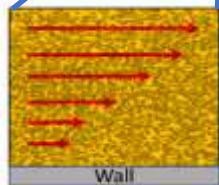
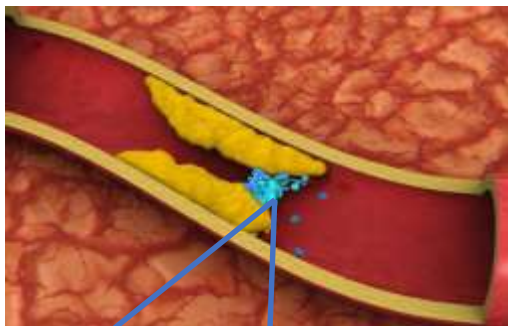
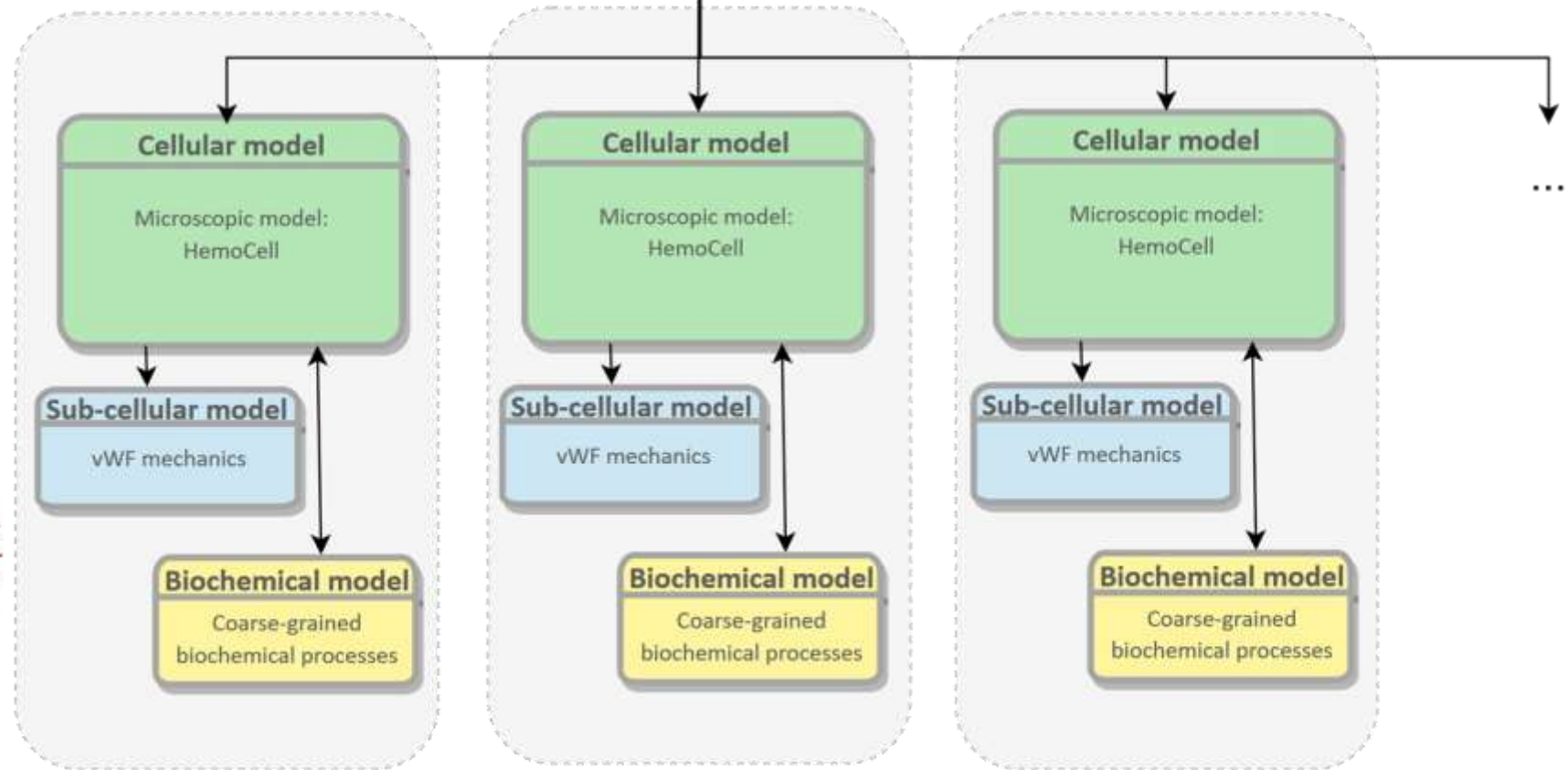
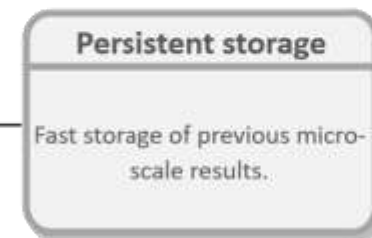
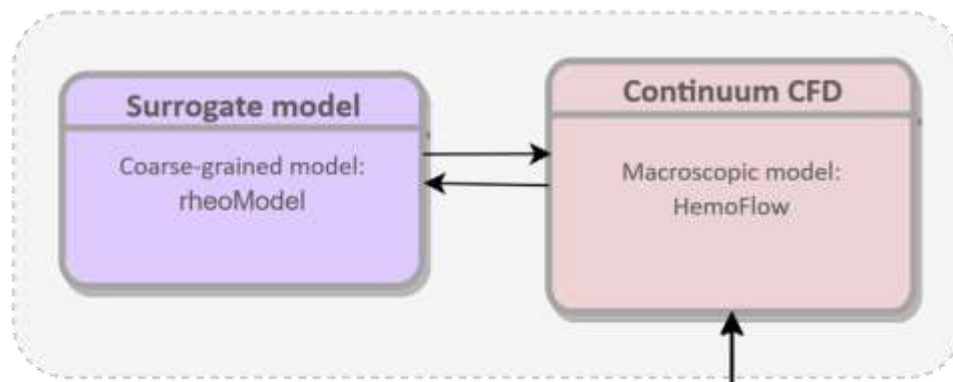
↓
Platelet aggregation

First target experiments



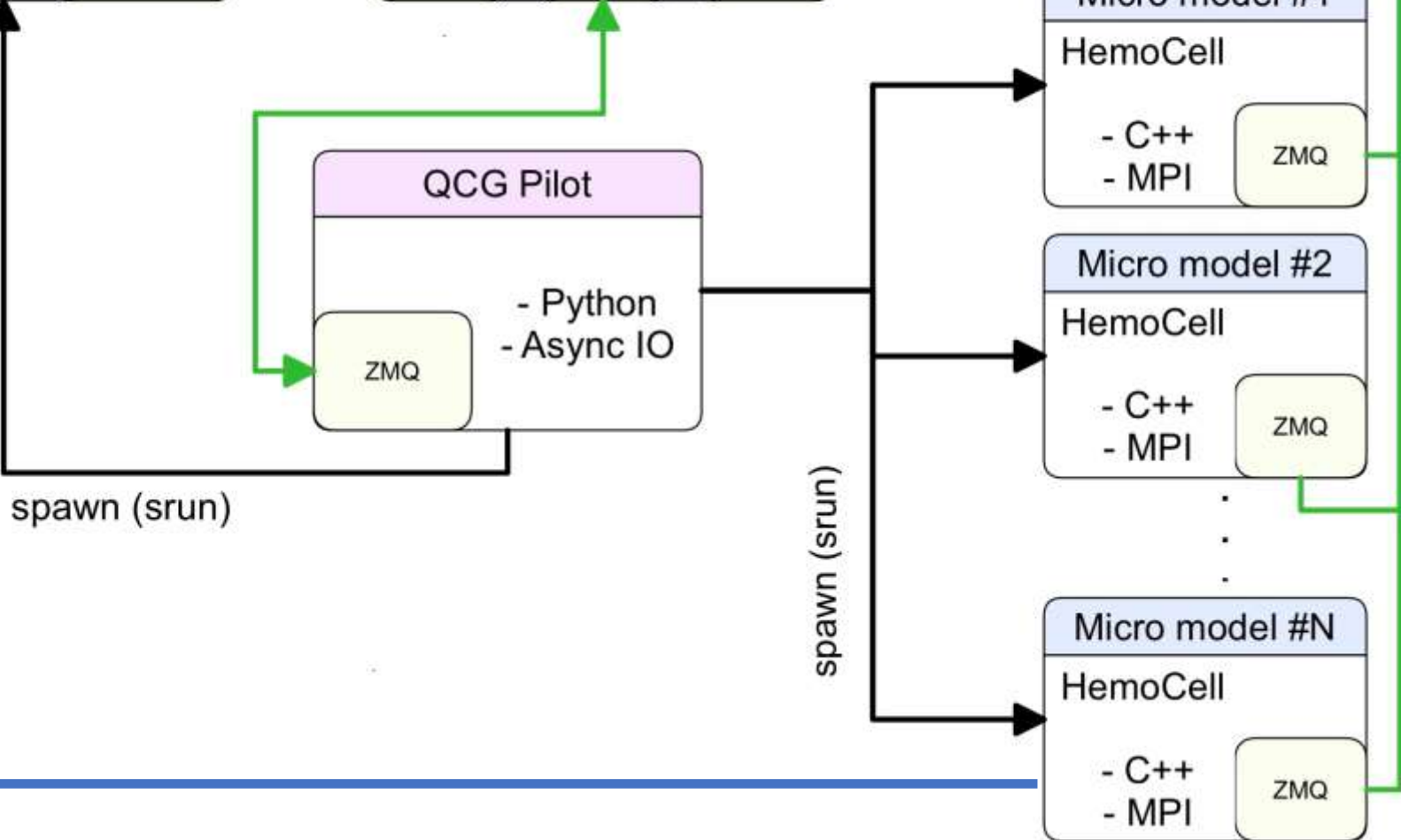
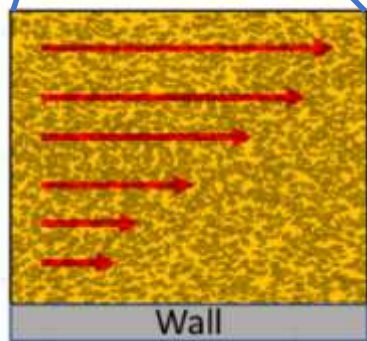
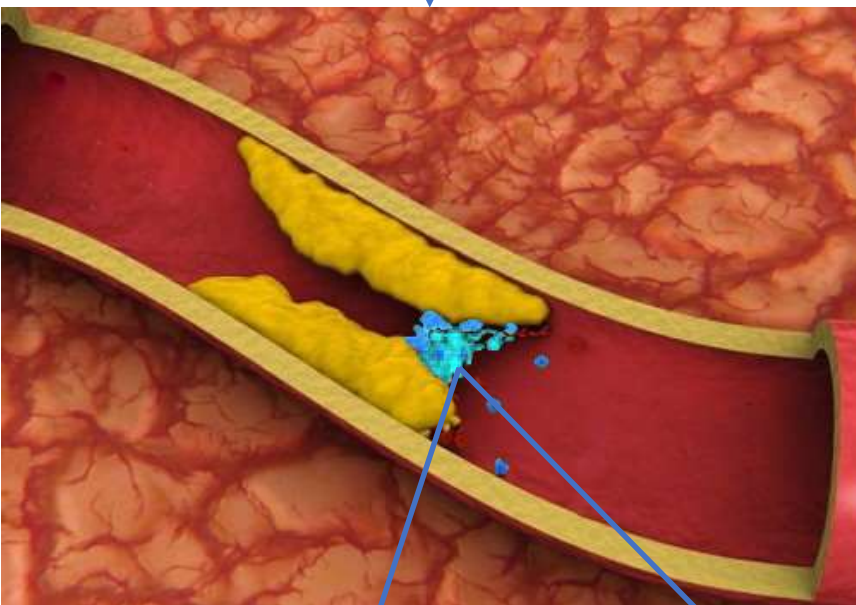
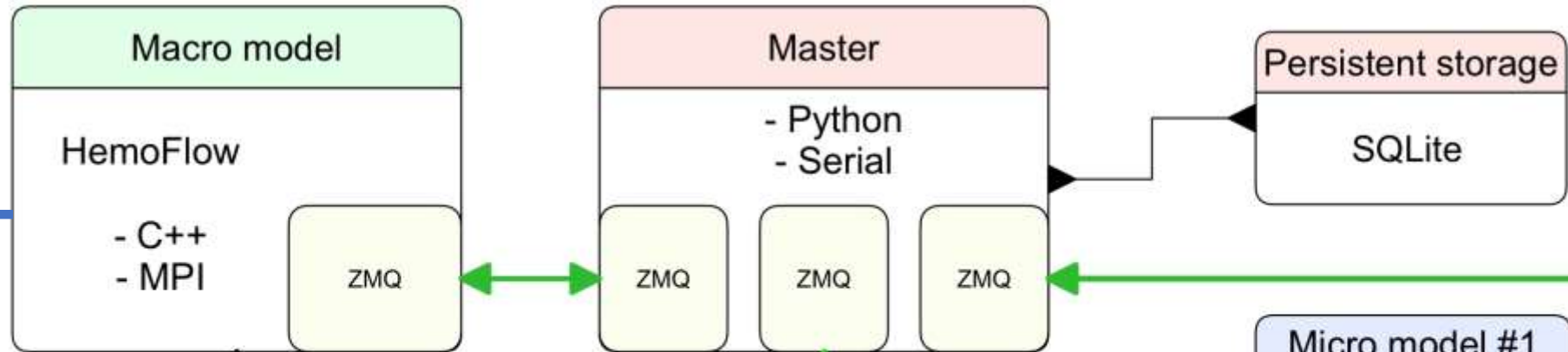
HemoScale

Model level



HemoScale

Implementation level



Thank you for your attention!



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