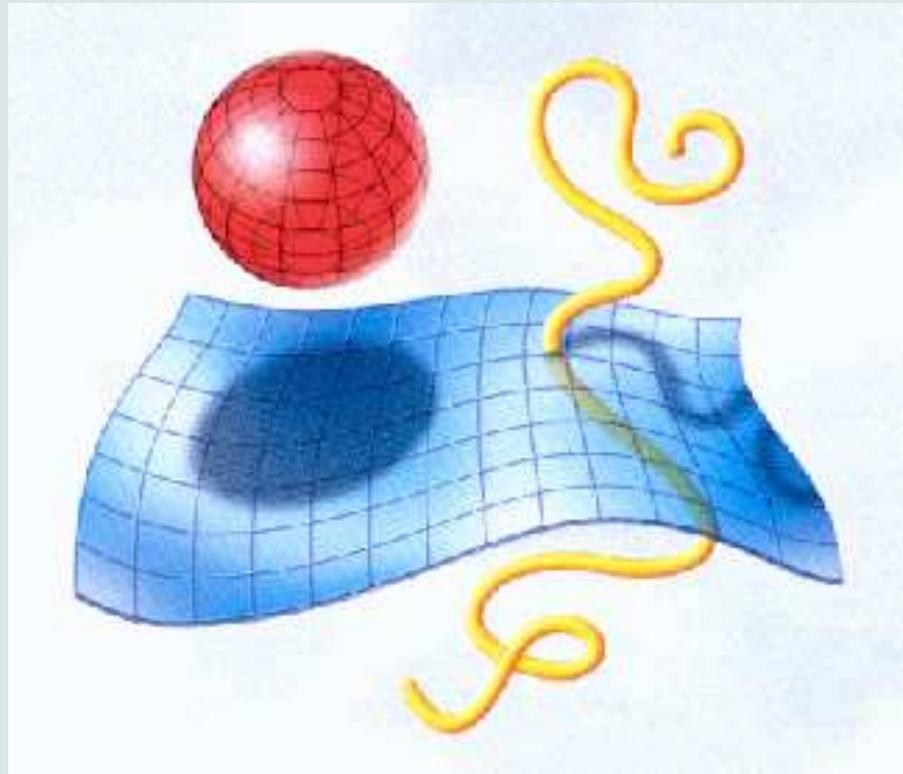




Mesoscale simulation of blood flow in microcapillaries

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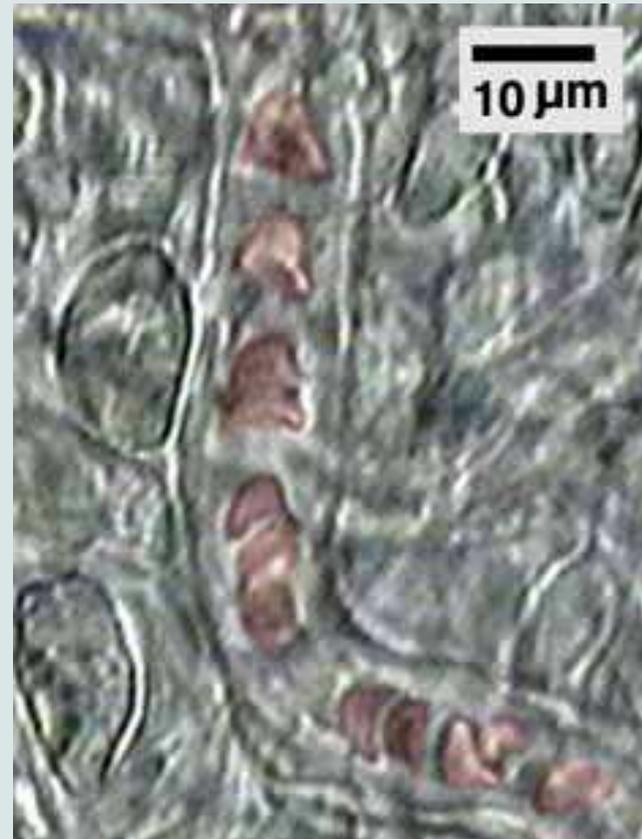
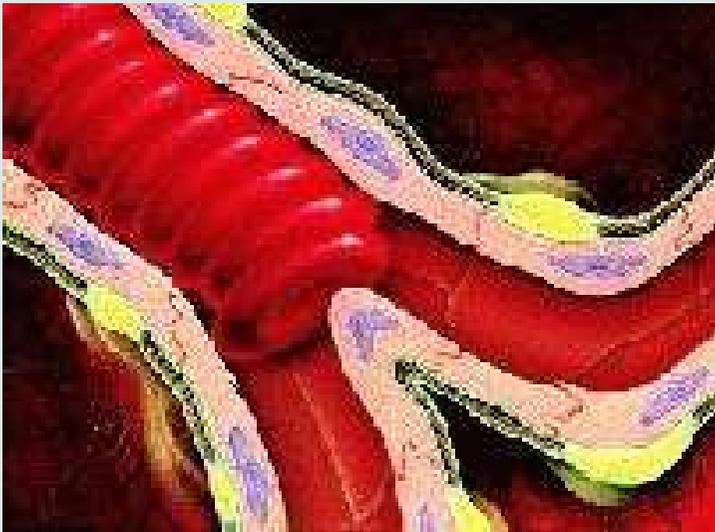
Work done during the last year in the Soft Matter Theory group in Jülich. Involved people include

- Hiroshi Noguchi (now in Tokyo)
- J. Liam McWhirter (now in Helsinki)
- Julia Fornleitner
- Dmitry Fedosov
- Matti Peltomäki
- Prof. Gerhard Gompper

Soft Matter Hydrodynamics

Cells and vesicles in flow:

- Red blood cells in microvessels:



Diseases such as diabetes reduce deformability of red blood cells!

Mesoscale Simulations

Complex fluids: length- and time-scale gap between

- atomistic scale of solvent
- mesoscopic scale of dispersed particles (colloids, polymers, membranes)

—→ **Mesoscale Simulation Techniques**

Basic idea:

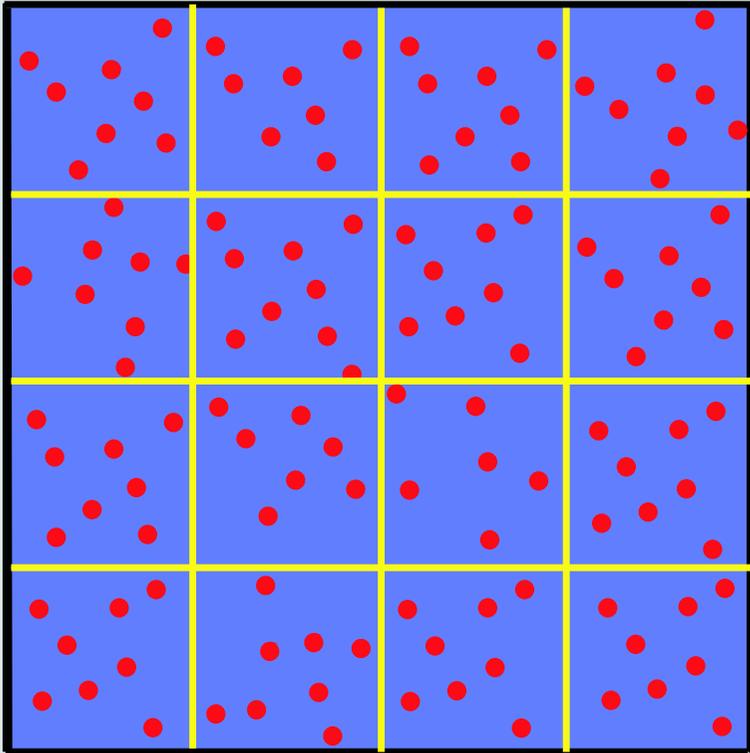
- drastically simplify dynamics on molecular scale
- respect conservation laws for mass, momentum, energy

Examples:

- Lattice Boltzmann Method (LBM)
- Dissipative Particle Dynamics (DPD)
- Multi-Particle-Collision Dynamics (MPC)

Mesoscale Simulations: MPC

Multi-Particle-Collision Dynamics (MPC)



- coarse grained fluid
- point particles
- off-lattice method
- collisions inside “cells”
- thermal fluctuations

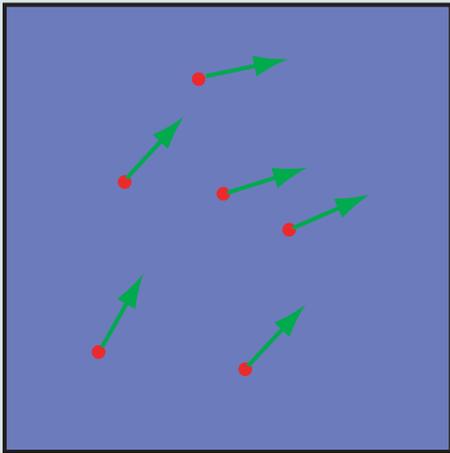
A. Malevanets and R. Kapral, J. Chem. Phys. **110** (1999)

A. Malevanets and R. Kapral, J. Chem. Phys. **112** (2000)

Mesoscale Simulations: MPC

Flow dynamics: Two step process

Streaming



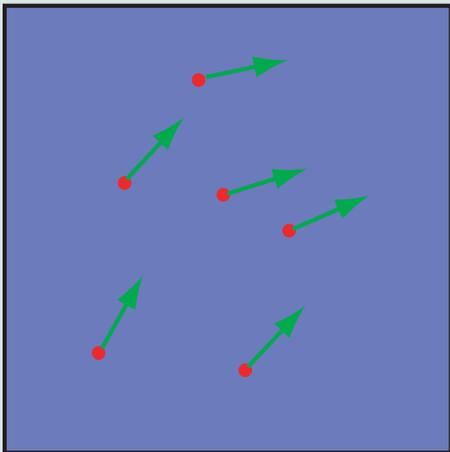
- ballistic motion

$$\mathbf{r}_i(t + h) = \mathbf{r}_i(t) + \mathbf{v}_i(t)h$$

Mesoscale Simulations: MPC

Flow dynamics: Two step process

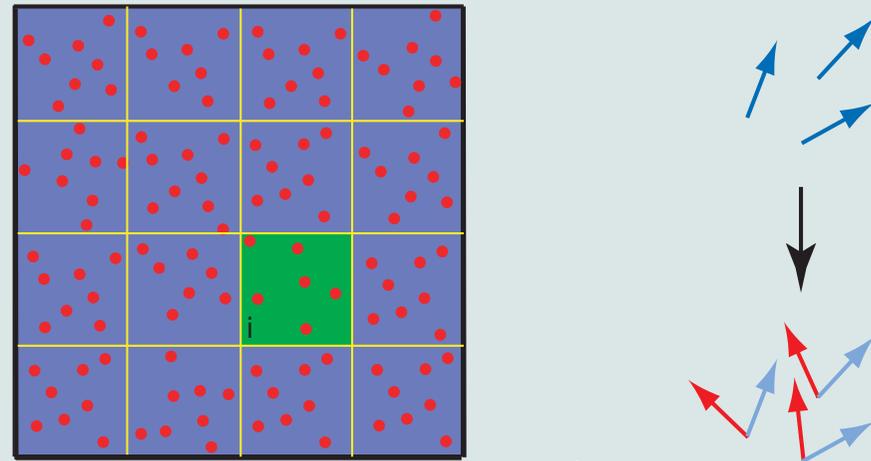
Streaming



- ballistic motion

$$\mathbf{r}_i(t + h) = \mathbf{r}_i(t) + \mathbf{v}_i(t)h$$

Collision



- mean velocity per cell

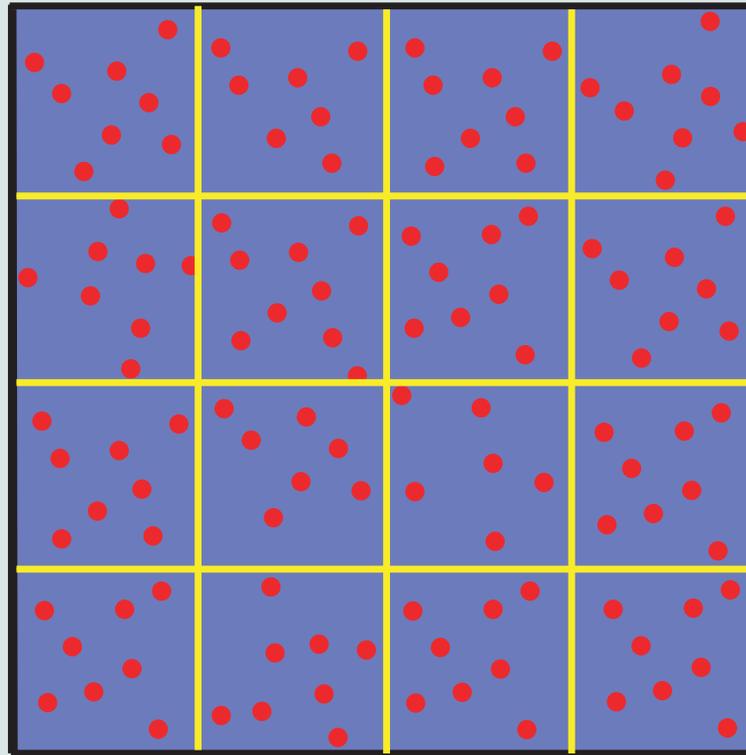
$$\bar{\mathbf{v}}_i(t) = \frac{1}{n_i} \sum_{j \in C_i} \mathbf{v}_j(t)$$

- rotation of relative velocity by angle α

$$\mathbf{v}'_i = \bar{\mathbf{v}}_i + \mathbf{D}(\alpha)(\mathbf{v}_i - \bar{\mathbf{v}}_i)$$

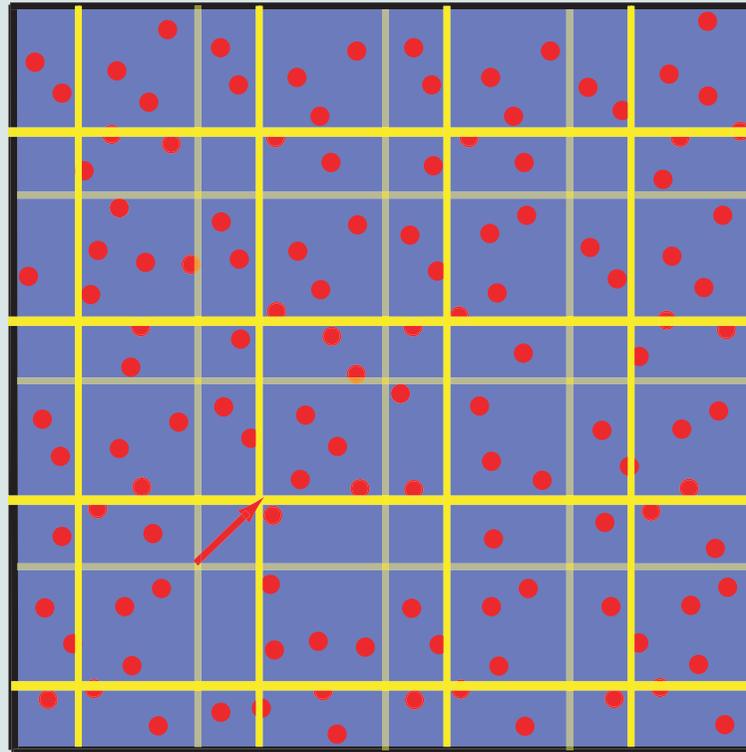
Mesoscale Simulations: MPC

- Lattice of collision cells: breakdown of **Galilean invariance**
- Restore Galilean invariance exactly: **random shifts** of cell lattice



Mesoscale Simulations: MPC

- Lattice of collision cells: breakdown of **Galilean invariance**
- Restore Galilean invariance exactly: **random shifts** of cell lattice



Mesoscale Simulations: Reynolds and Schmidt Numbers

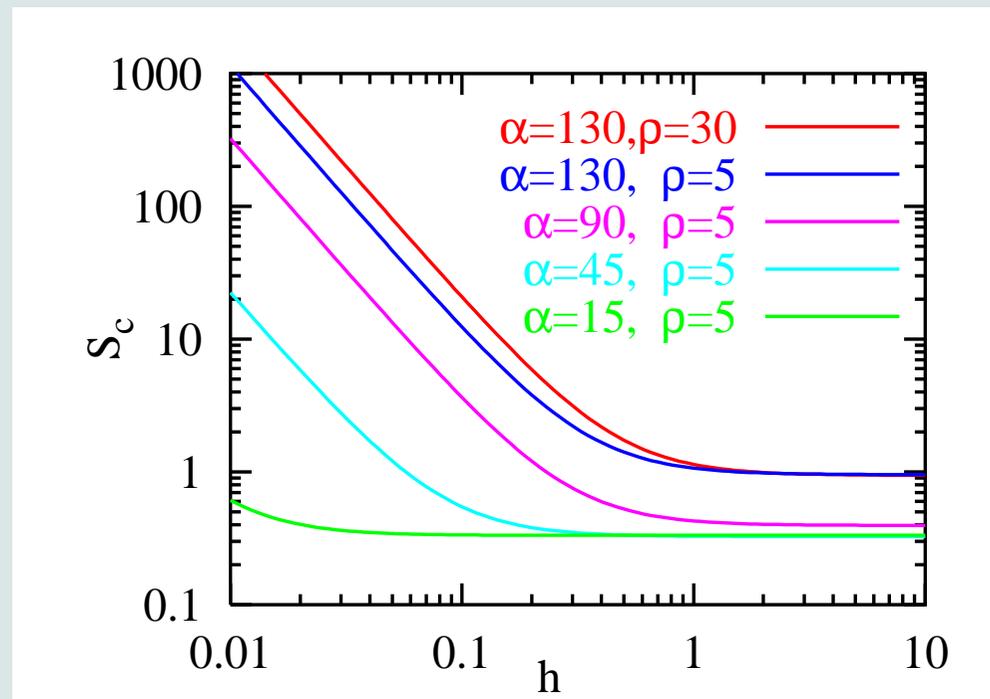
- **Reynolds number** $Re = v_{\max}L/\nu \sim$ inertia forces / friction forces

For soft matter systems with characteristic length scales of μm :

$$Re \simeq 10^{-3}$$

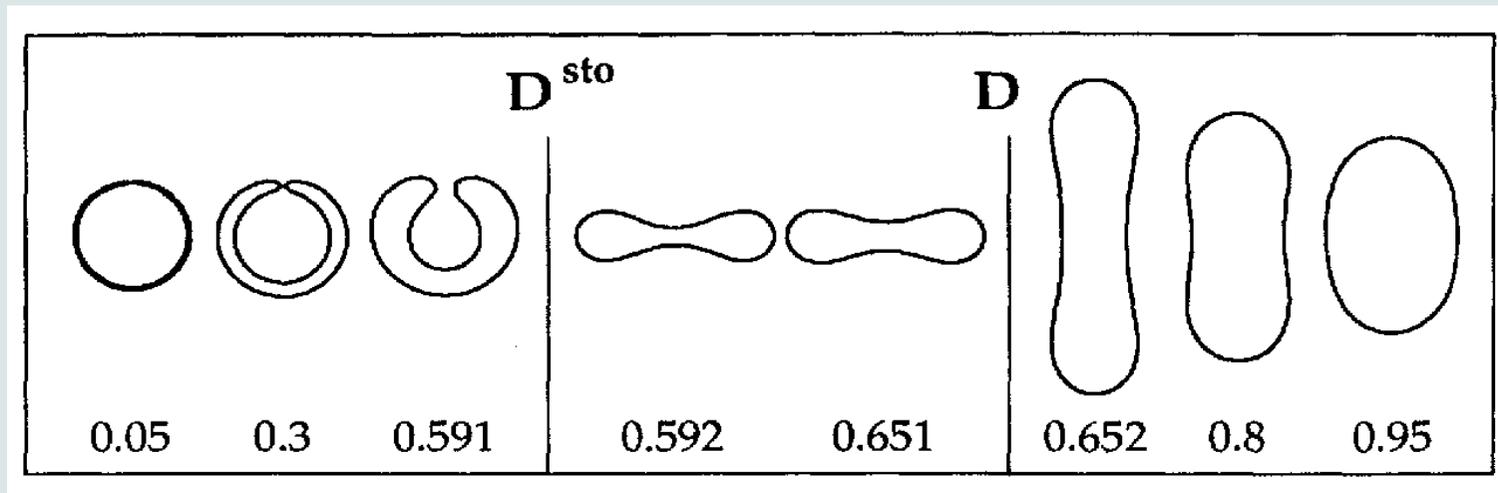
- **Schmidt number** $Sc = \nu/D \sim$ momentum transp. / mass transp.

Gases: $Sc \simeq 1$, liquids: $Sc \simeq 10^3$



Membranes and Vesicles: Equilibrium Shapes

Minimize curvature energy for fixed area $A = 4\pi R_0^2$ and reduced volume $V^* = V/V_0$, where $V_0 = 4\pi R_0^3/3$:



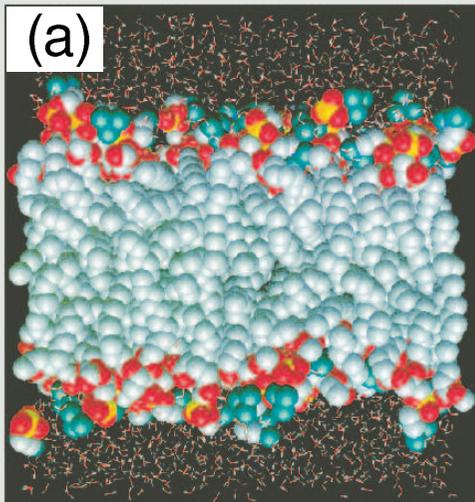
stomatocyte

discocyte

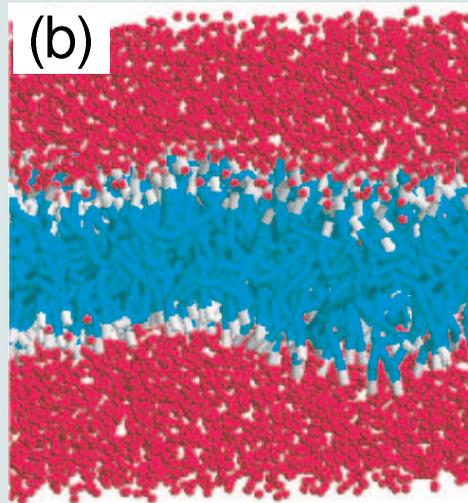
prolate

U. Seifert, K. Berndl, and R. Lipowsky, Phys. Rev. A 44 (1991)

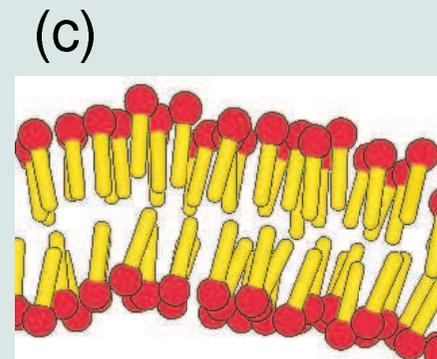
Membranes and Vesicles: Different Length Scales



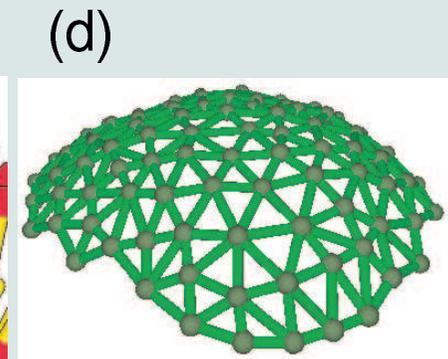
atomistic



coarse-grained

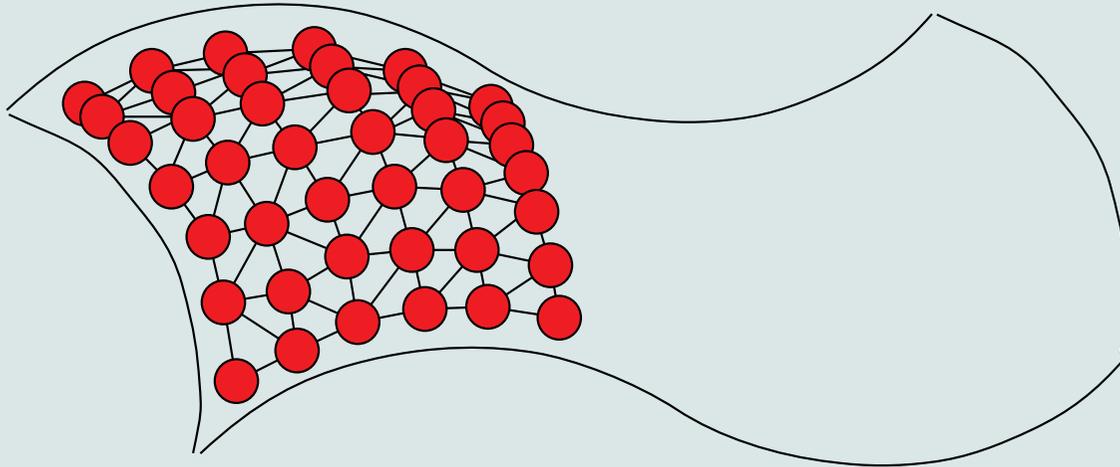


solvent-free



triangulated

Membranes and Vesicles: Triangulated Surfaces



- Closed 2D surface in 3D.
- N vertices (membrane particles)
- $3(N - 2)$ edges
- $2(N - 2)$ faces

Two- and many-body potentials:

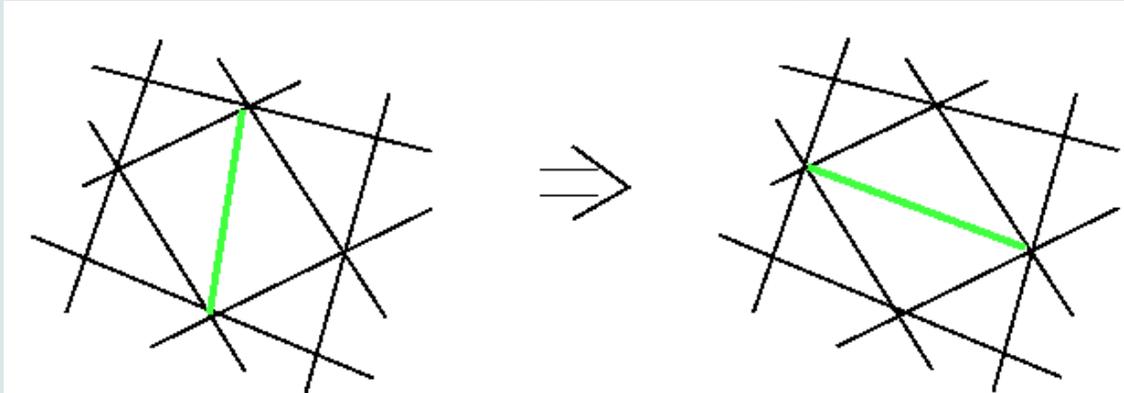
- Bending rigidity κ
- Pairwise soft repulsion
- Pairwise soft attraction (in the presence of an edge)
- Surface area constraint
- Volume constraint

G. Gompper, D.M.Kroll, J. Phys. Cond. Matt **9** (1997);

G. Gompper, D.M. Kroll, in *Statistical Mechanics of Membranes and Surfaces* (World Scientific, 2004)

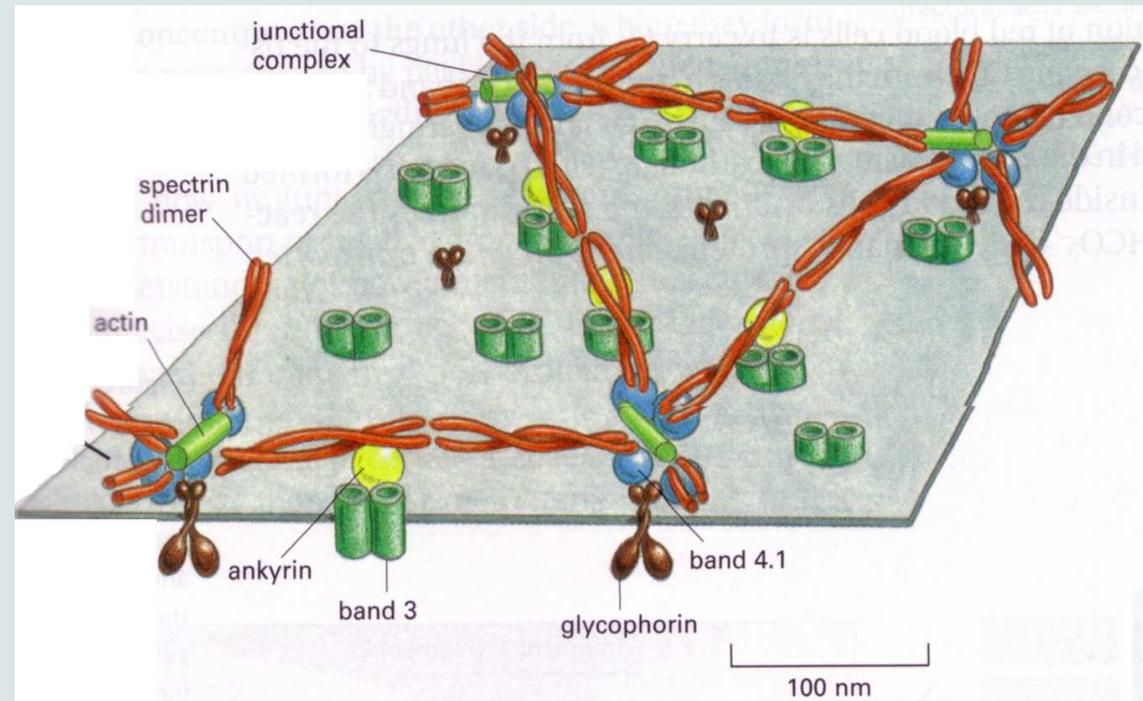
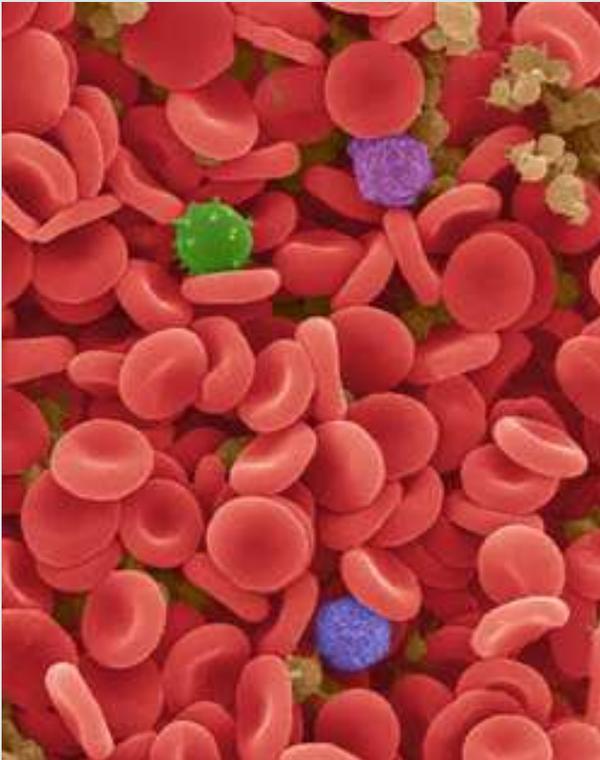
Membranes and Vesicles: Fluid Vesicles

In fluid vesicles, the triangulation is dynamic:



The membrane viscosity can be controlled through the bond-flip rate.

Membranes and Vesicles: Red Blood Cells



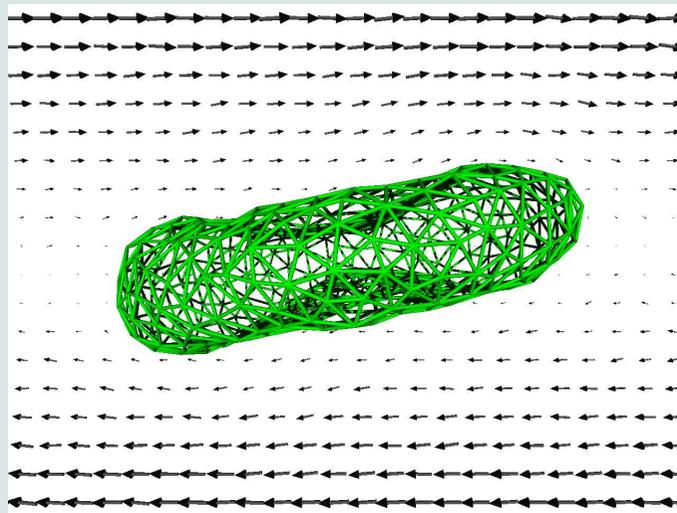
- Spectrin network (no bond flips) induces shear elasticity μ of composite membrane
- For real RBCs: elastic parameters: $\kappa/k_B T = 50$, $\mu R_0^2/k_B T = 5000$

Membranes and Vesicles: Coupling to the Fluid

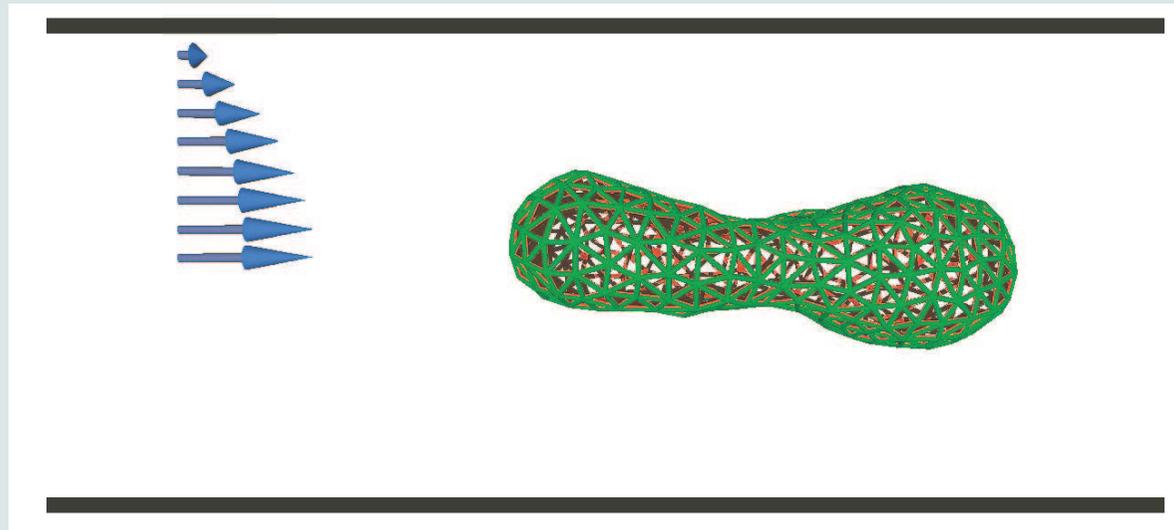
Interaction between membrane and fluid:

- Streaming step:
bounce-back scattering of solvent particles on triangles
- Collision step:
membrane vertices are included in MPC collisions

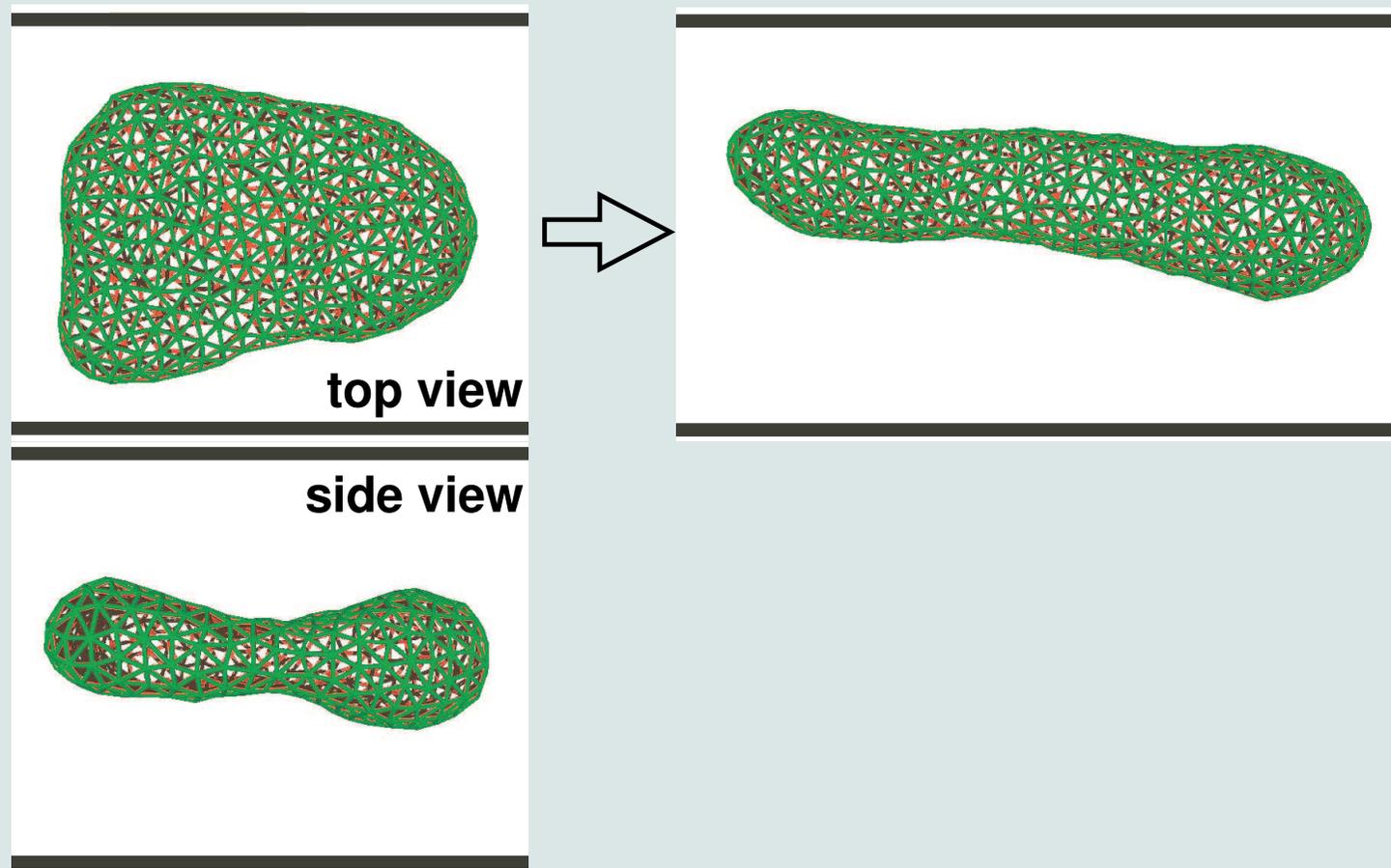
implies **impenetrable membrane** with **no-slip boundary conditions**.



Capillary Flow: Results for one vesicle



Capillary Flow: One Fluid Vesicle

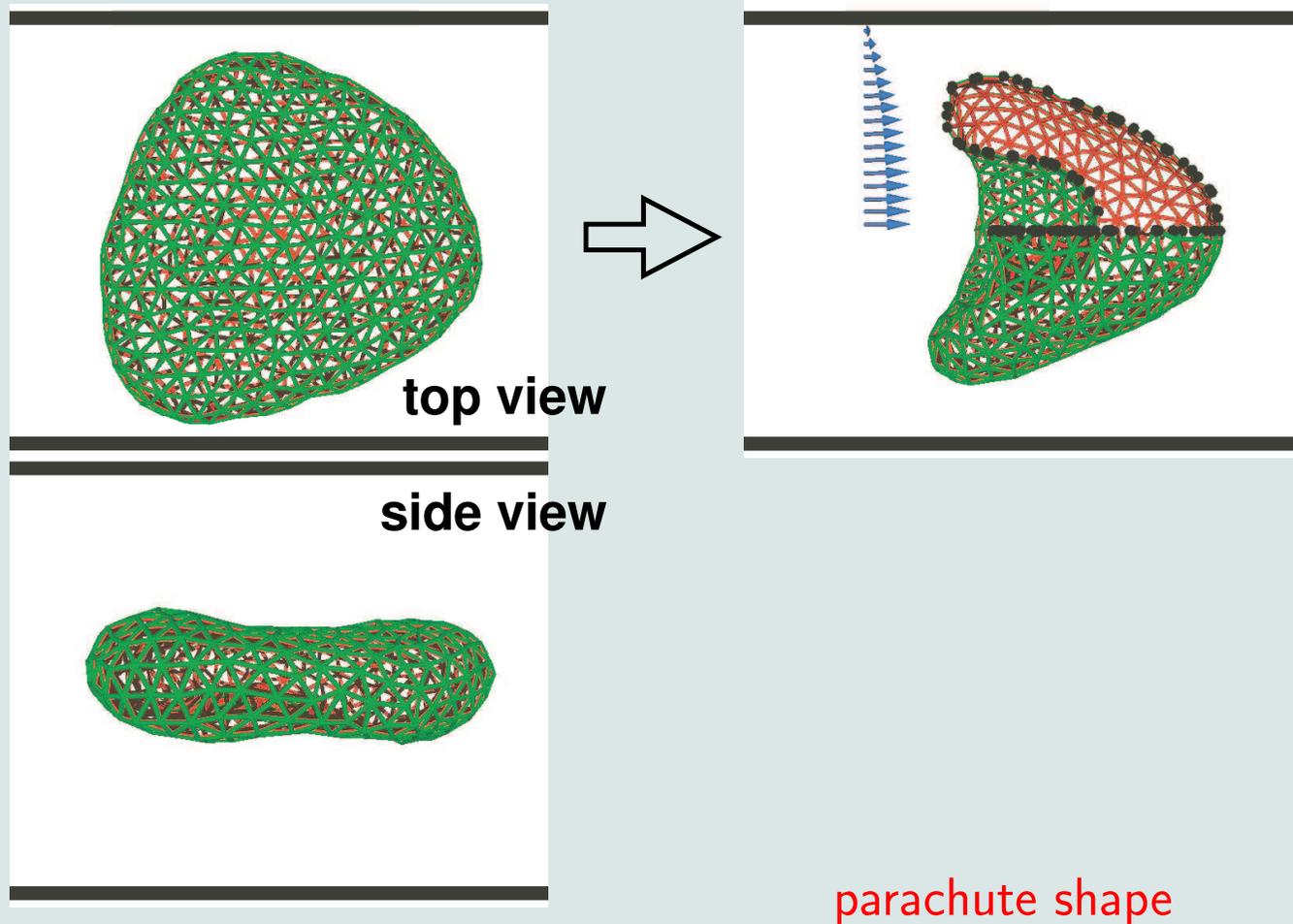


- small flow velocities: vesicle axis **perpendicular** to capillary axis \longrightarrow **no axial symmetry!**
- discocyte-to-prolate transition with increasing flow

Capillary Flow: One Elastic Vesicle

Elastic vesicle:

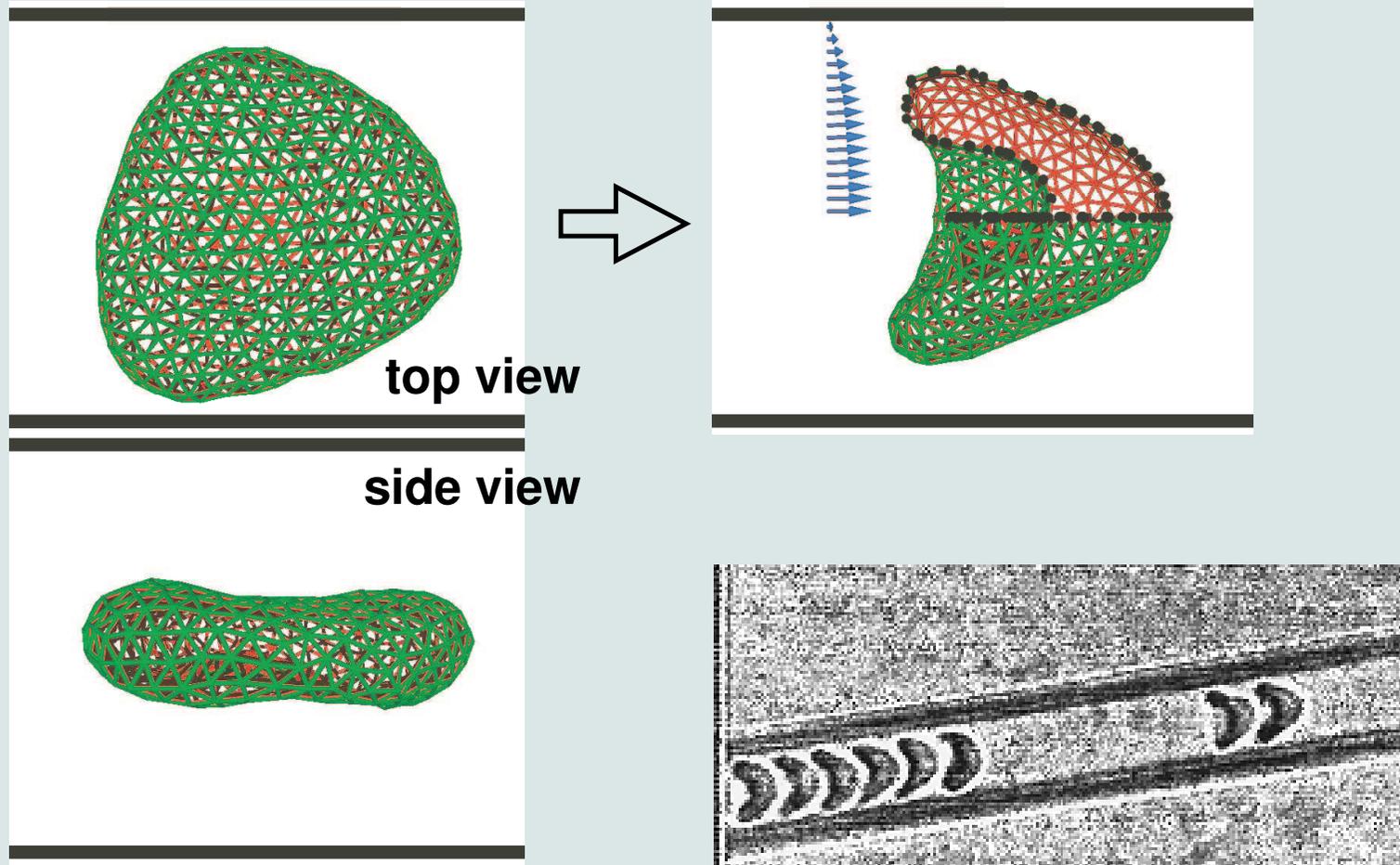
- curvature and shear elasticity ($\kappa = 20 k_B T$, $\mu = 110 k_B T / R_0^2$)
- model for red blood cells



Capillary Flow: One Elastic Vesicle

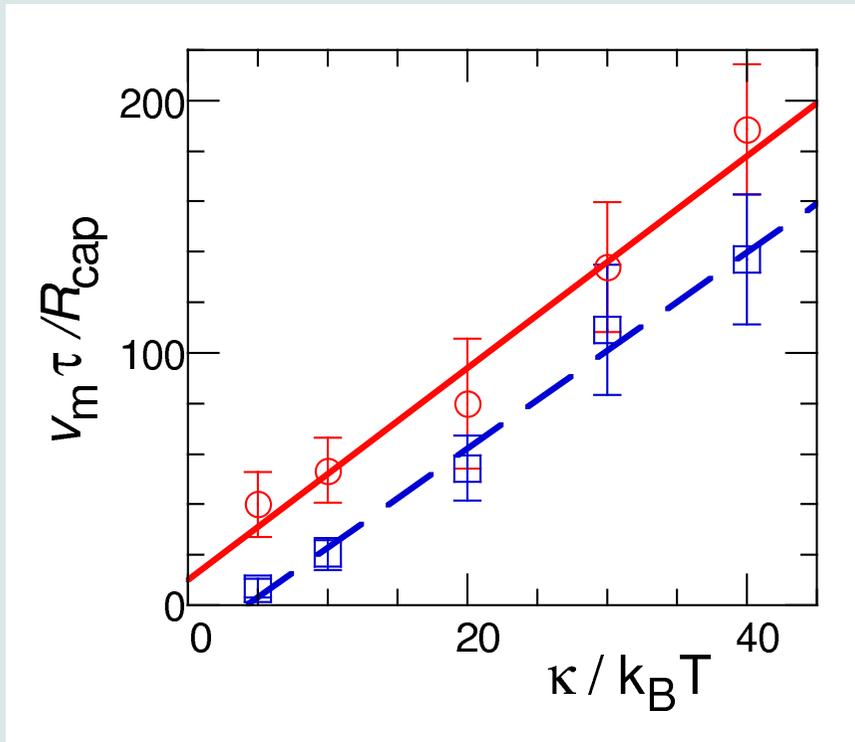
Elastic vesicle:

- curvature and shear elasticity
- model for red blood cells

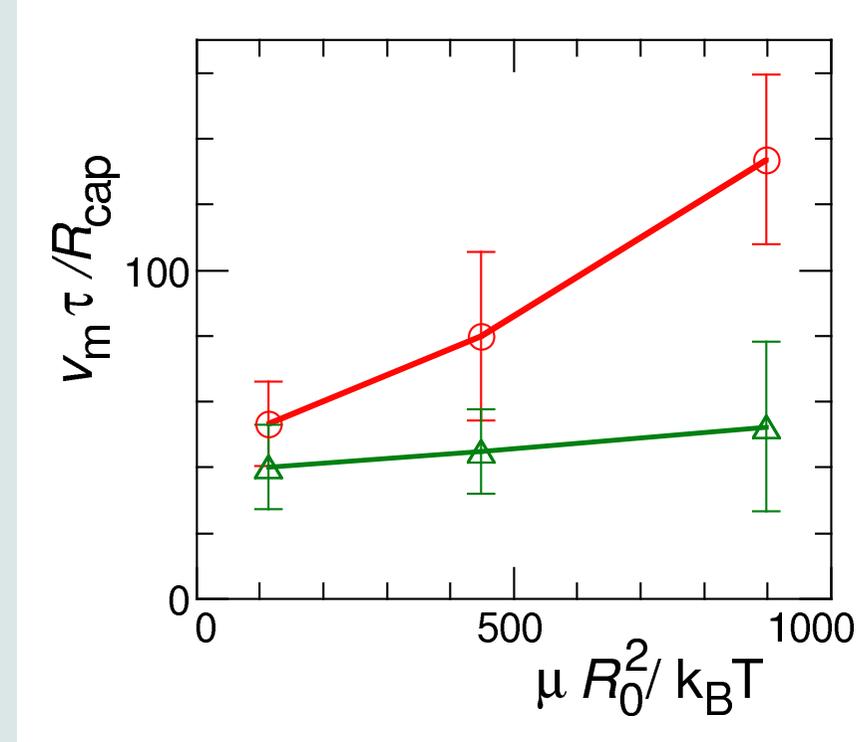


Capillary Flow: One Elastic Vesicle

The discocyte-to-parachute transition takes place as a function of flow velocity $v_m \tau / R_{cap}$



bending rigidity ($\mu = 110 k_B T / R_0^2$)

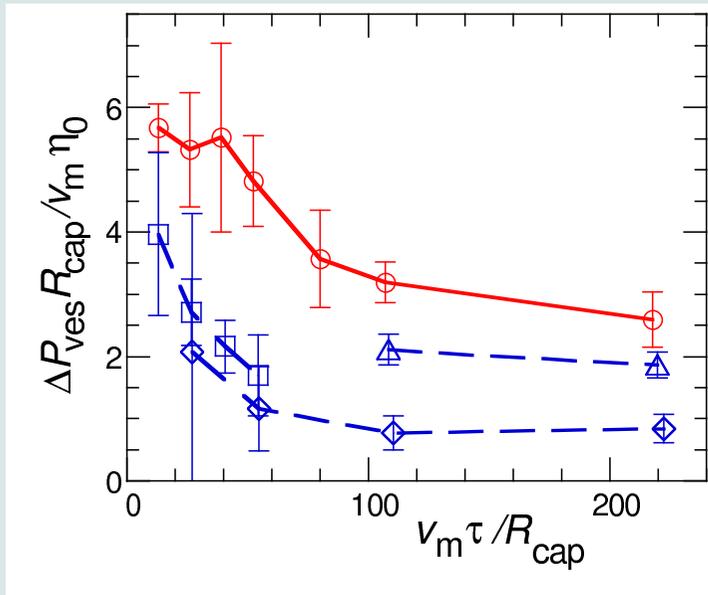


shear modulus ($\kappa = 20 k_B T$)

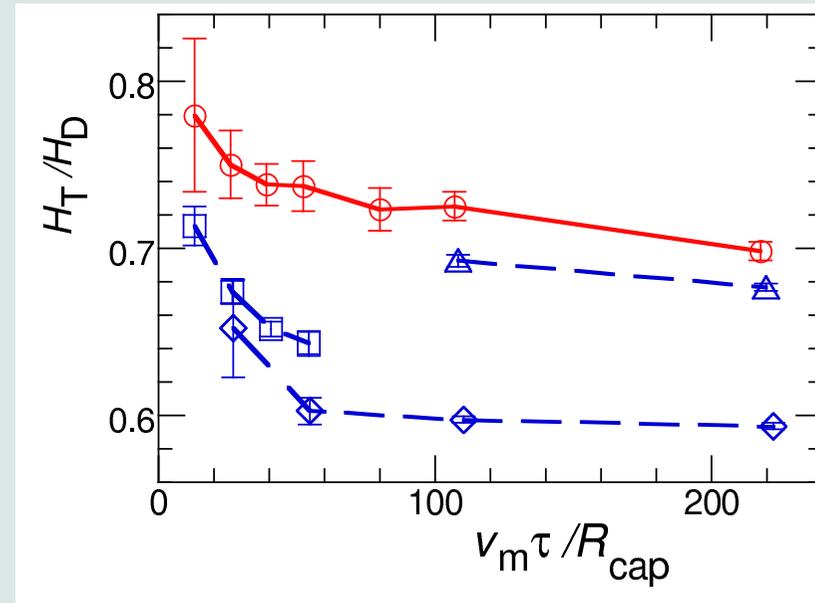
Implies for RBCs: $v_{trans} \simeq 0.2 \text{ mm/s}$ for $R_{cap} = 4.6 \mu\text{m}$

Capillary Flow: One Elastic Vesicle

Pressure drop due to vesicle



Hematocrit ratio $H_T / H_D = v_m / v_{ves}$



elastic vesicle

fluid vesicle:

□ discocyte

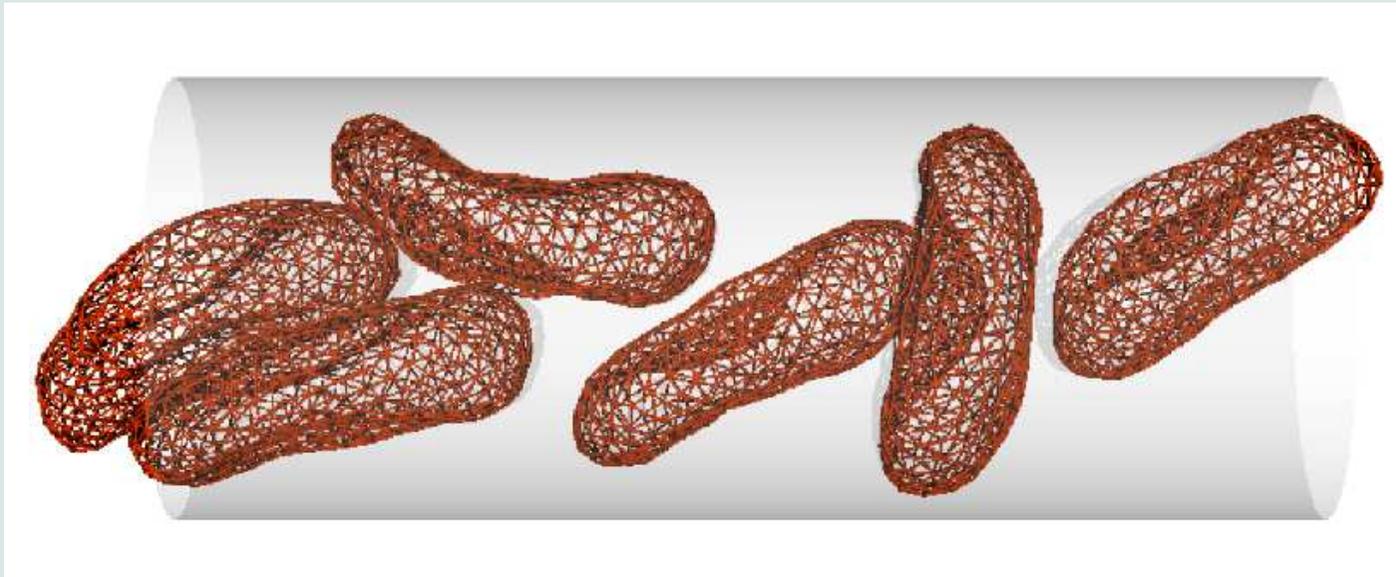
△ parachute

◇ prolate

vesicle velocity **larger** than fluid velocity (Fåhræus effect)

$$\Delta P = \frac{8\eta_0(v_o - v_m)L_z}{R_{cap}^2}$$

Capillary Flow: Results for several vesicle



Capillary Flow: Clustering & Alignment

Physiological conditions: Hematocrit (volume fraction of RBCs) $H = 0.45$

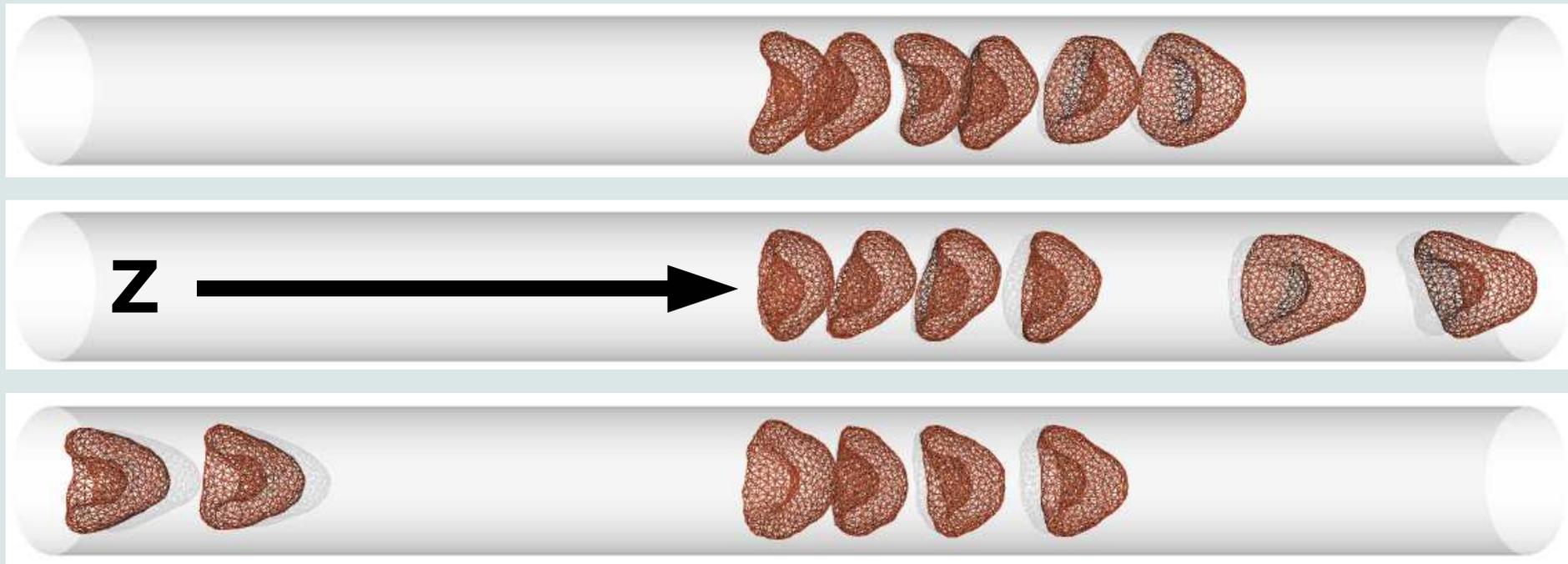
Lower in narrow capillaries $H_T = 0.1...0.2$

Therefore: Hydrodynamic interactions between RBCs very important

Note: No direct attractive interactions considered!

Capillary Flow: Clustering & Alignment

Low hematocrit H_T :

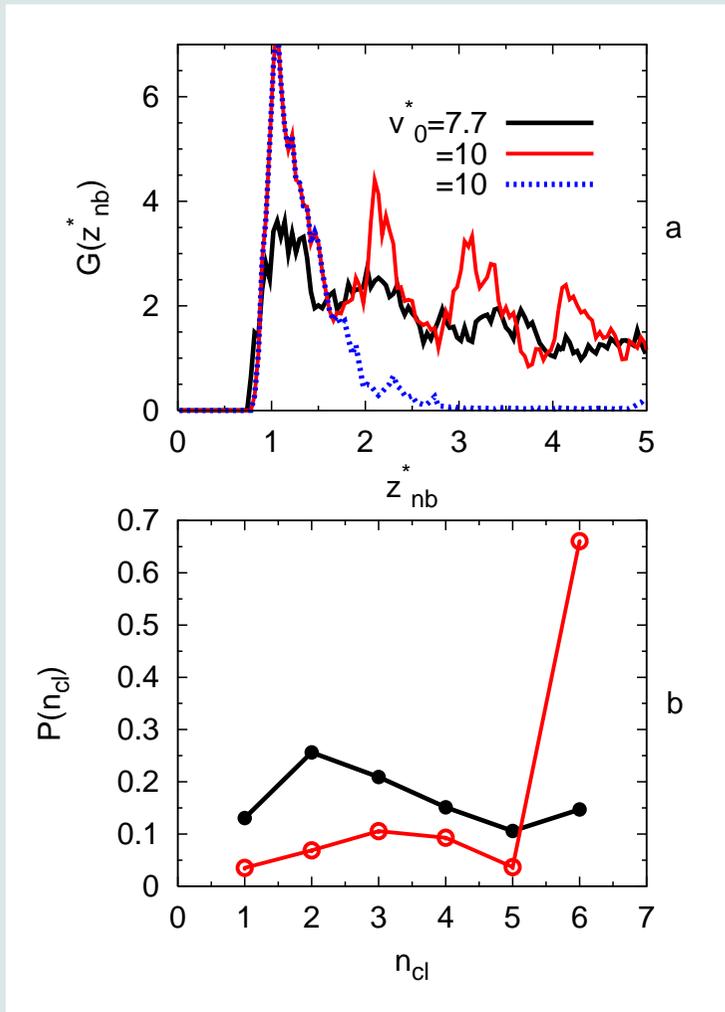


- Single vesicles more deformed \rightarrow move faster
- Effective hydrodynamic attraction stabilizes clusters

J.L. McWhirter, H. Noguchi, G. Gompper, Proc. Natl. Acad. Sci. **106** (2009)

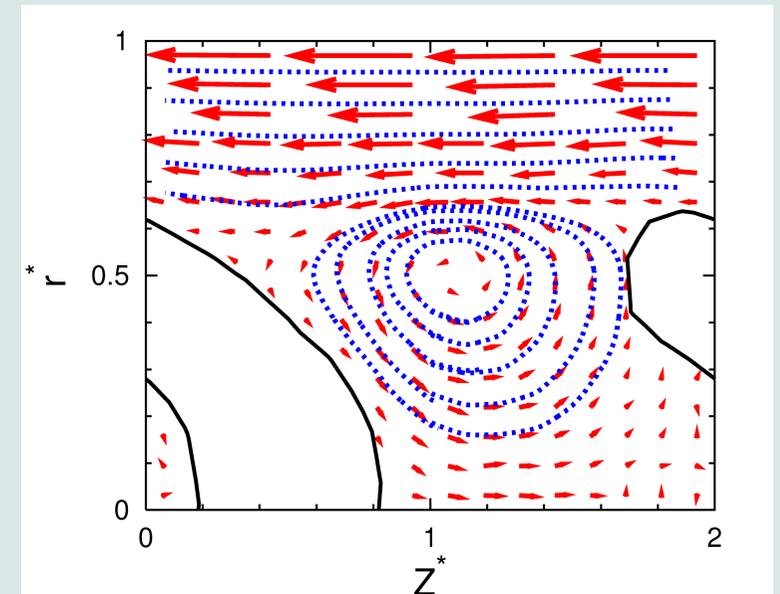
Capillary Flow: Clustering & Alignment

Low hematocrit H_T :



Positional
correlation
function

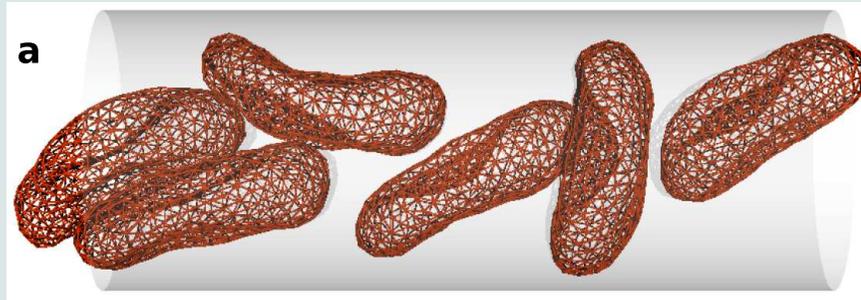
Probability for
cluster size n_{cl}



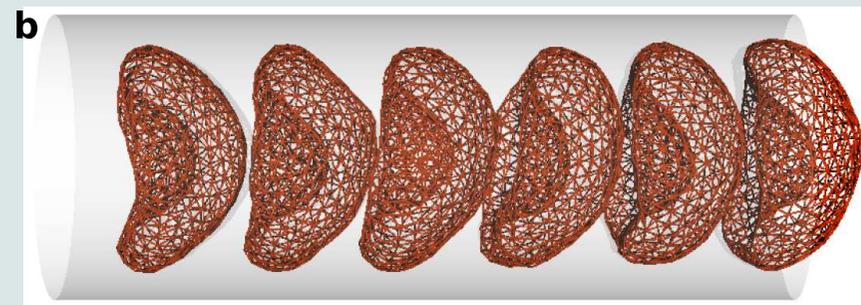
Clustering tendency **increases**
with increasing flow velocity

Capillary Flow: Clustering & Alignment

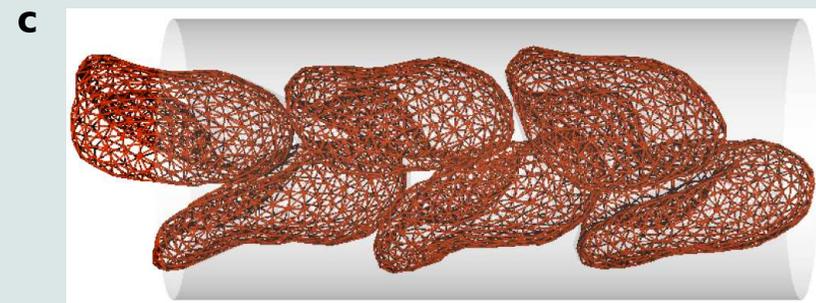
High hematocrit H_T :



disordered discocyte



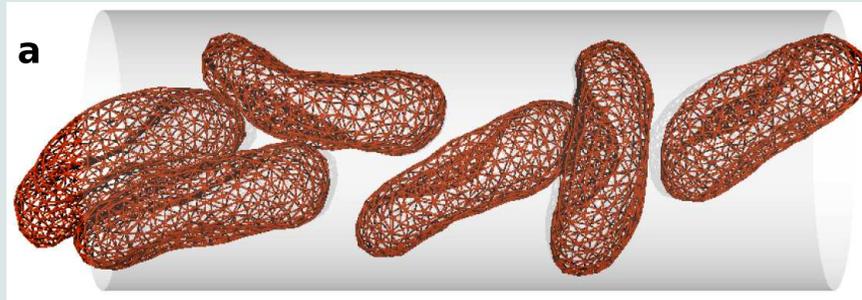
aligned parachute



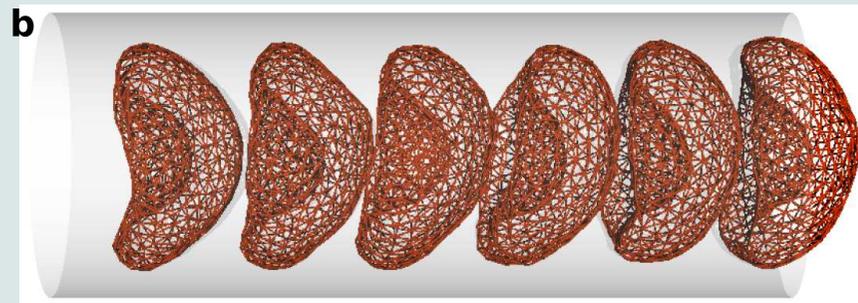
zig-zag slipper

Capillary Flow: Clustering & Alignment

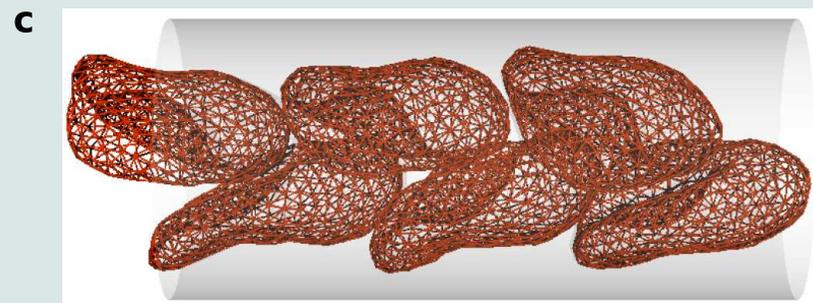
High hematocrit H_T :



disordered discocyte



aligned parachute



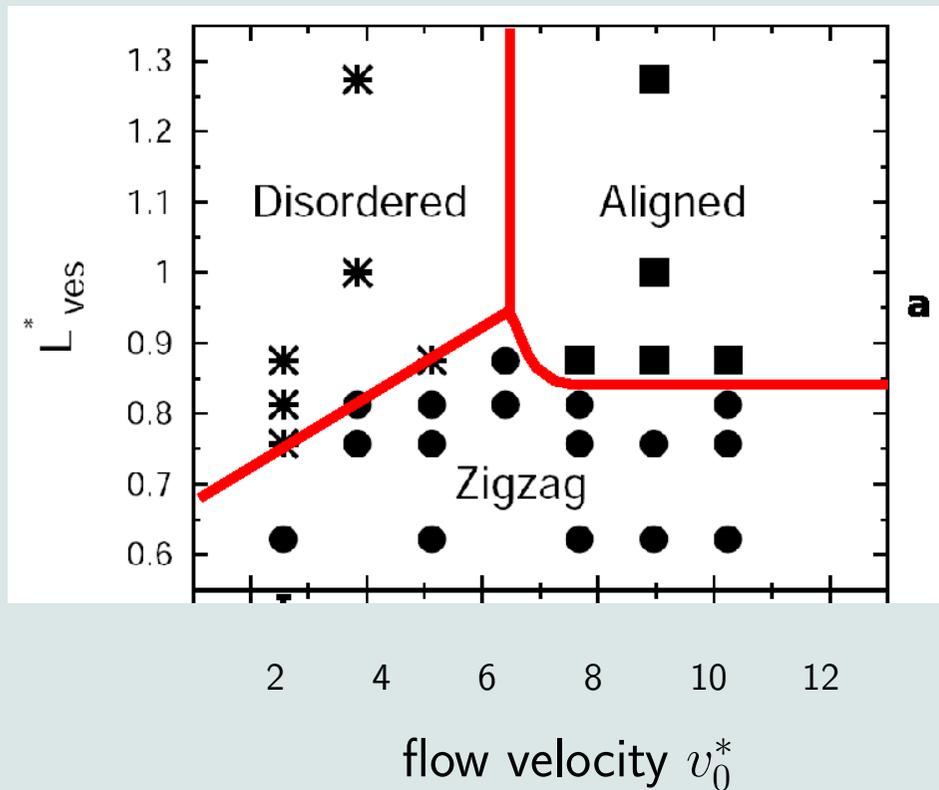
zig-zag slipper



Skalak, Science, 1969

Capillary Flow: Clustering & Alignment

Phase diagram:



$$\text{Hematocrit } H_T = 0.28/L_{ves}^*$$

Transition to zig-zag phase despite **higher** flow resistance than aligned-parachute phase!

Summary

- [Mesoscale simulation techniques](#) are powerful tool to bridge the length- and time-scale gap in complex fluids
- Multi-particle-collision dynamics well suited for hydrodynamics of [embedded particles](#): colloids, polymers, vesicles, RBCs
- Red blood cells in [capillary flow](#): shear elasticity implies parachute shapes, hydrodynamic clustering and alignment

[Review](#): G. Gompper, T. Ihle, D.M. Kroll, R.G. Winkler, Adv. Polym. Sci. **221**, 1 (2009)