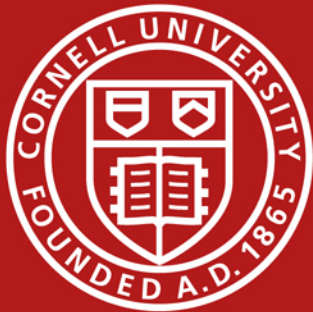


# Multiscale model of platelet tethering and adhesion: similarities with the leukocyte adhesion cascade



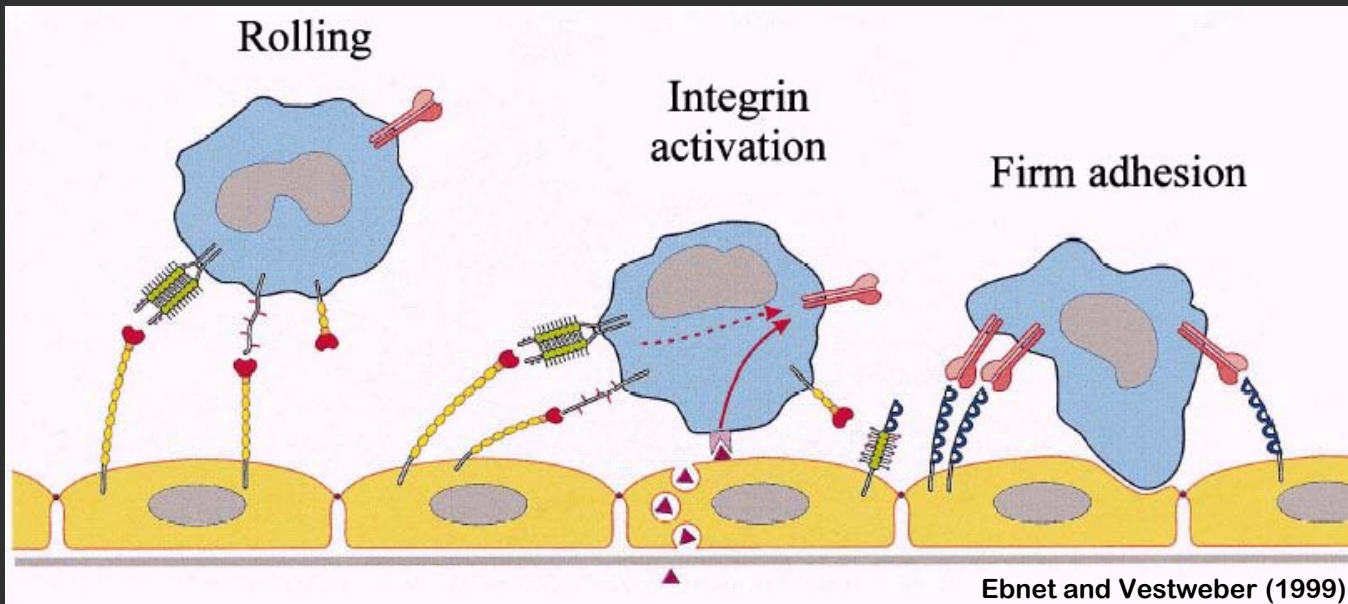
Cornell University

Weiwei Wang<sup>1</sup>, John P. Lindsey II<sup>1</sup>, Jonathan B. Freund<sup>3</sup>,  
Thomas G. Diacovo<sup>2</sup>, Michael R. King<sup>1</sup>

1. Department of Biomedical Engineering, Cornell University
2. Department of Pediatrics, Columbia University Medical Center, Irving Cancer Research Institute
3. Departments of Mechanical Science & Engineering and Aerospace Engineering, University of Illinois at Urbana-Champaign



# The adhesion cascade of circulating leukocytes

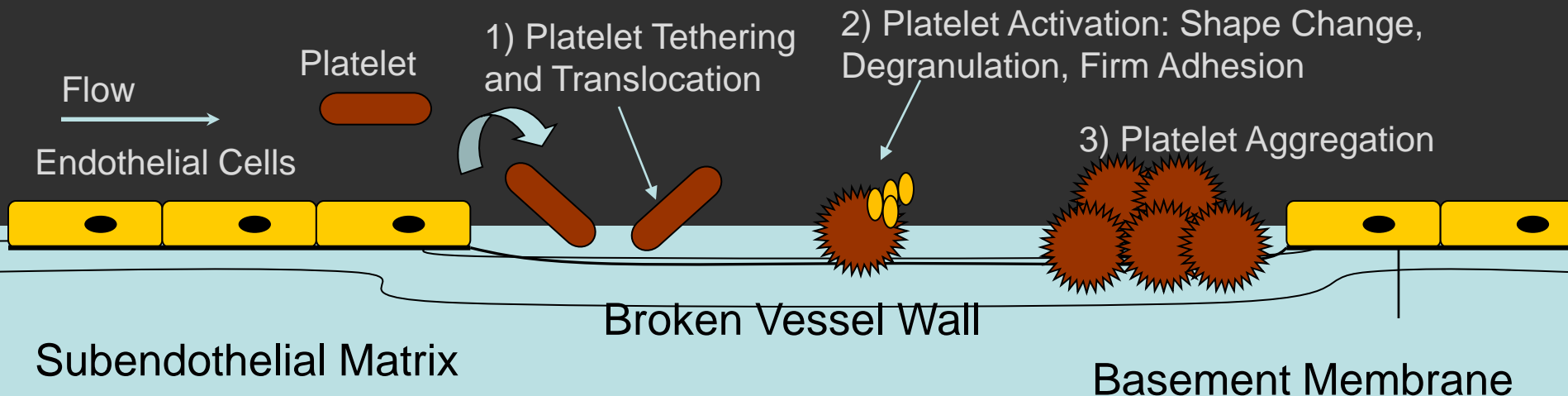
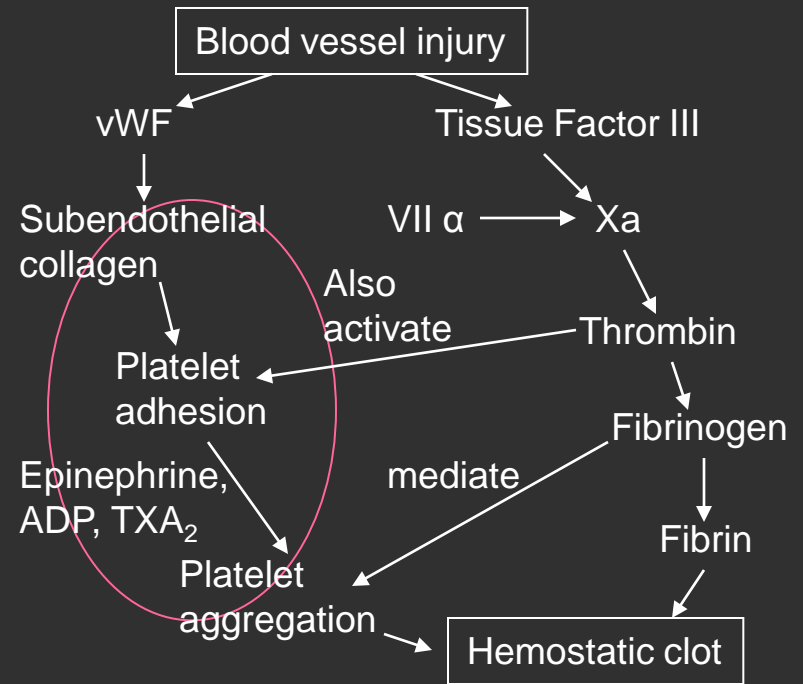


***Mild inflammation  
in live mouse venule***

King et al., *Microcirculation* (2003)  
King et al., *Annals of BME* (2004)  
King et al., *Phys. Fluids* (2005)  
Lamkin-Kennard et al., *Biorheology* (2005)

• **Hemostasis :**

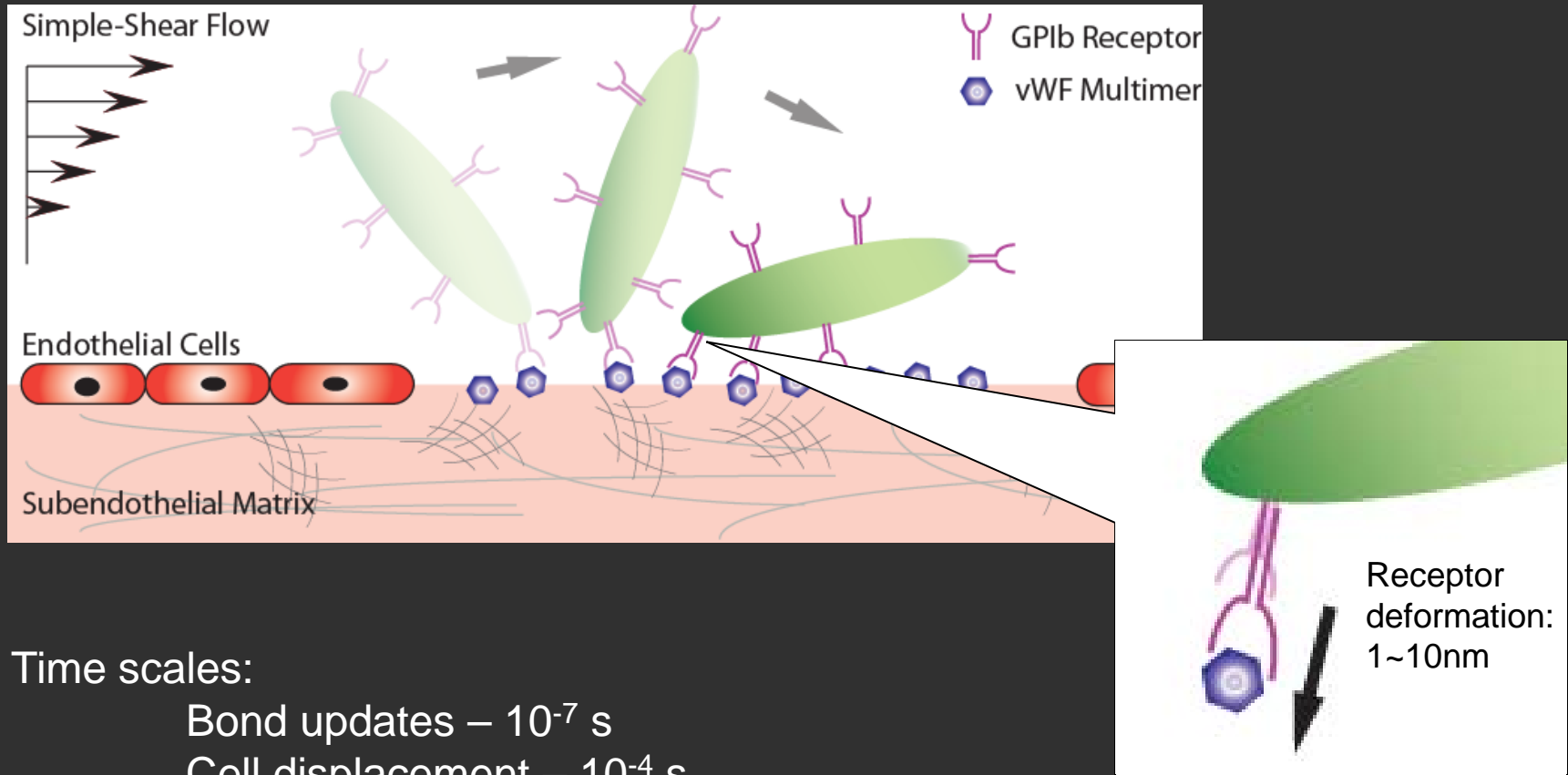
a self defensive mechanism in response to blood vessel injuries to prevent excessive blood loss. It describes the physiological process of blood clotting at injured vessels





Length Scale:

Platelet Translocation:  $\sim 1\mu\text{m}$



Time scales:

Bond updates –  $10^{-7}$  s

Cell displacement –  $10^{-4}$  s

Thrombus growth – 1 – 100 s



Here we consider the **GPIb-vWF** bonding which mediate initial tethering of platelets to the area of vessel injury.

Receptor-Ligand bond formation/breakage:

Using Monte Carlo method to determine each bond formation/breakage event.

$$P_f = 1 - \exp(-k_f \Delta t)$$

$$P_r = 1 - \exp(-k_r \Delta t)$$

$$k_f = k_{f,2-D}^0 v_s \exp\left(\sigma \left| x_b - l_b \right| \frac{\gamma - 0.5 \left| x_b - l_b \right|}{k_b T}\right)$$

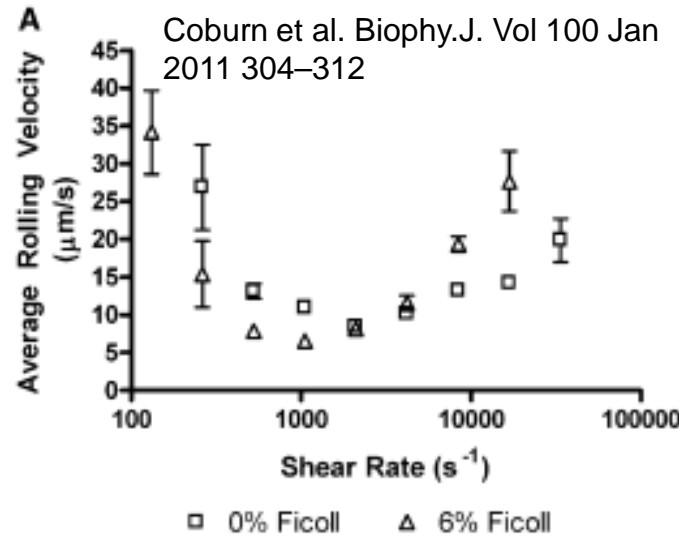
$$k_r = k_r^0 \exp\left(\frac{\gamma F_b}{k_B T}\right) \quad \text{Bell Model}$$

Parameter (unit)	Definition	Value	Reference
$l_b$ (nm)	Equilibrium GPIb $\alpha$ -VWF bond length	128	[Singh 2006, Fox 1988]
$\gamma$ (nm)	Reactive compliance	0.71	[Arya 2002]
$k_{f,2-D}^0$ (s <sup>-2</sup> /μm)	Intrinsic cross-linking formation rate constant	0.05	Determined by a series of simulations
$k_r$ (s <sup>-1</sup> )	Intrinsic dissociation constant	5.47	[Arya 2005]
$\sigma$ (pN/nm)	Spring Constant	10	[Chtcheglova 2004]



Cornell University

# Single Platelet: Rolling



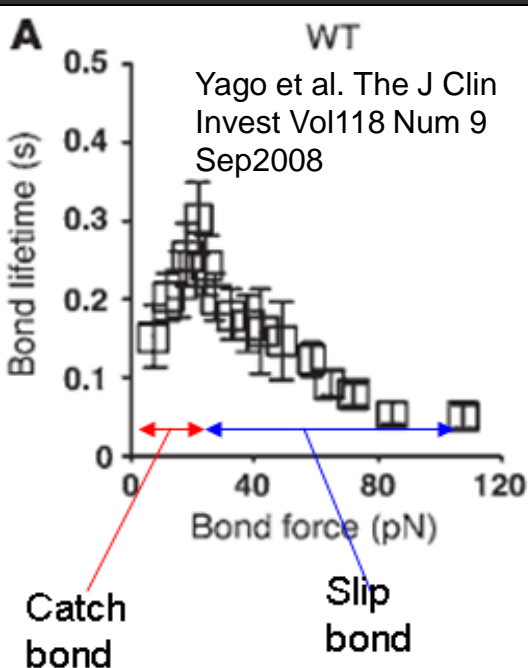
The proposed “catch bond”, “slip bond” switch for the GPIIb-vWF is caused by shear/tether induced vWF conformation change. The resulted dissociation kinetics are:

Auton et al. argued that, there is a combination of NG (native, folded state) and IG (open, intermediate state) forms of vWF in the system. With the detailed description:

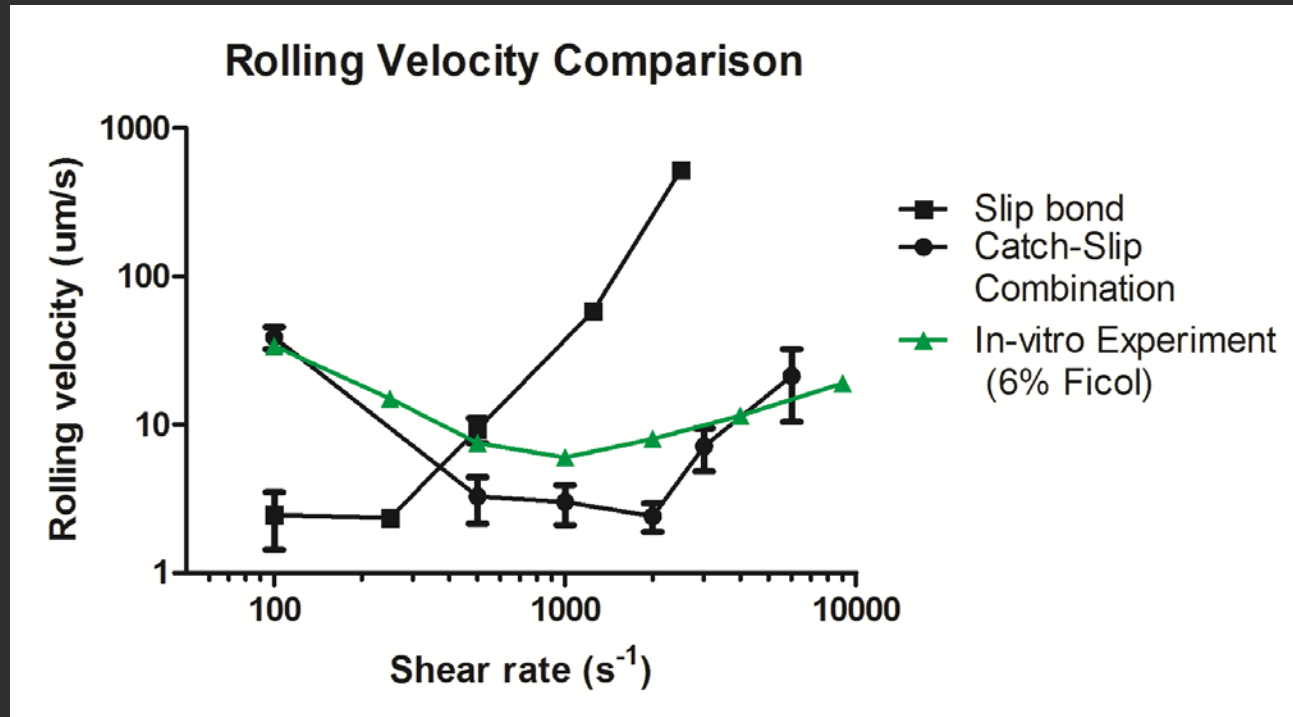
$$k_r = \frac{1}{1 + \Phi} k_N(f) + \frac{\Phi}{1 + \Phi} k_I(f), \quad \Phi = IG / NG$$

$$\Phi(f) = k_{NG \rightarrow IG}^0 / k_{IG \rightarrow NG}^0 * \exp\left(\frac{\gamma' f}{k_B T}\right)$$

$$k_N(f) = k_{N,off}^0 \exp\left(\frac{y_N f}{k_B T}\right) \quad k_I(f) = k_{I,off}^0 \exp\left(\frac{y_I f}{k_B T}\right)$$



Parameter (unit)	Definition	Value	Reference
$k_{NG \rightarrow IG}^0 / k_{IG \rightarrow NG}^0$	unstressed equilibrium constant for NG – IG states	0.4	[Auton 2010]
$\gamma'$ (nm)	force compliance for two states ratio	0.45	[Auton 2010]
$k_{N,off}^0$ ( $\text{s}^{-1}$ )	intrinsic dissociation constant for NG state	4.9	[Auton 2010]
$k_{I,off}^0$ ( $\text{s}^{-1}$ )	intrinsic dissociation constant for IG state	1.84	[Auton 2010]
$y_N$ (nm)	force compliance of dissociation for NG state	-0.23	[Auton 2010]
$y_I$ (nm)	force compliance of dissociation for IG state	0.039	[Auton 2010]



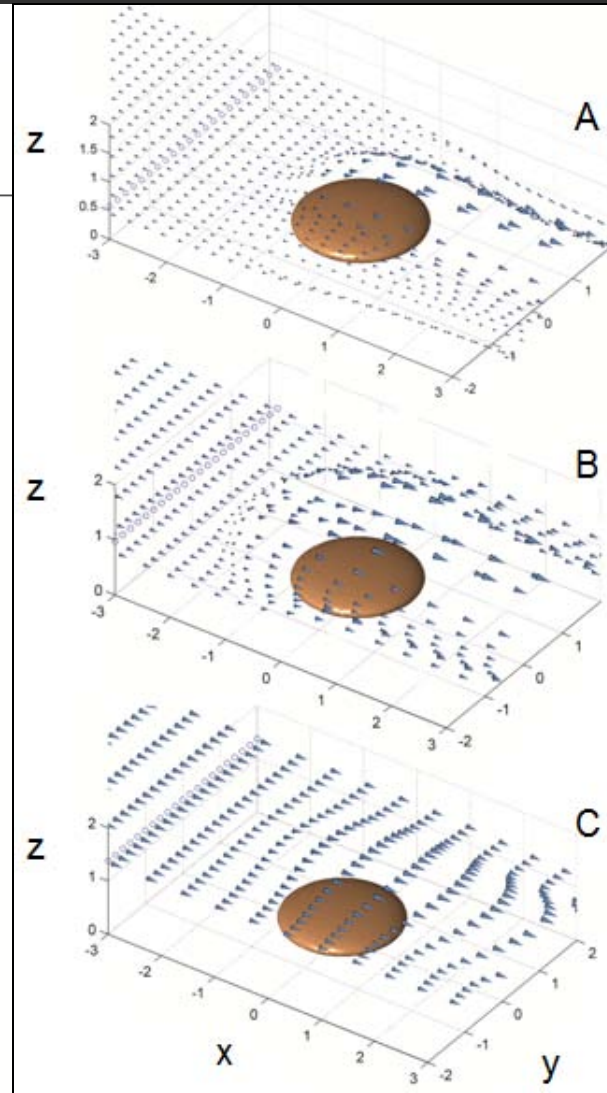
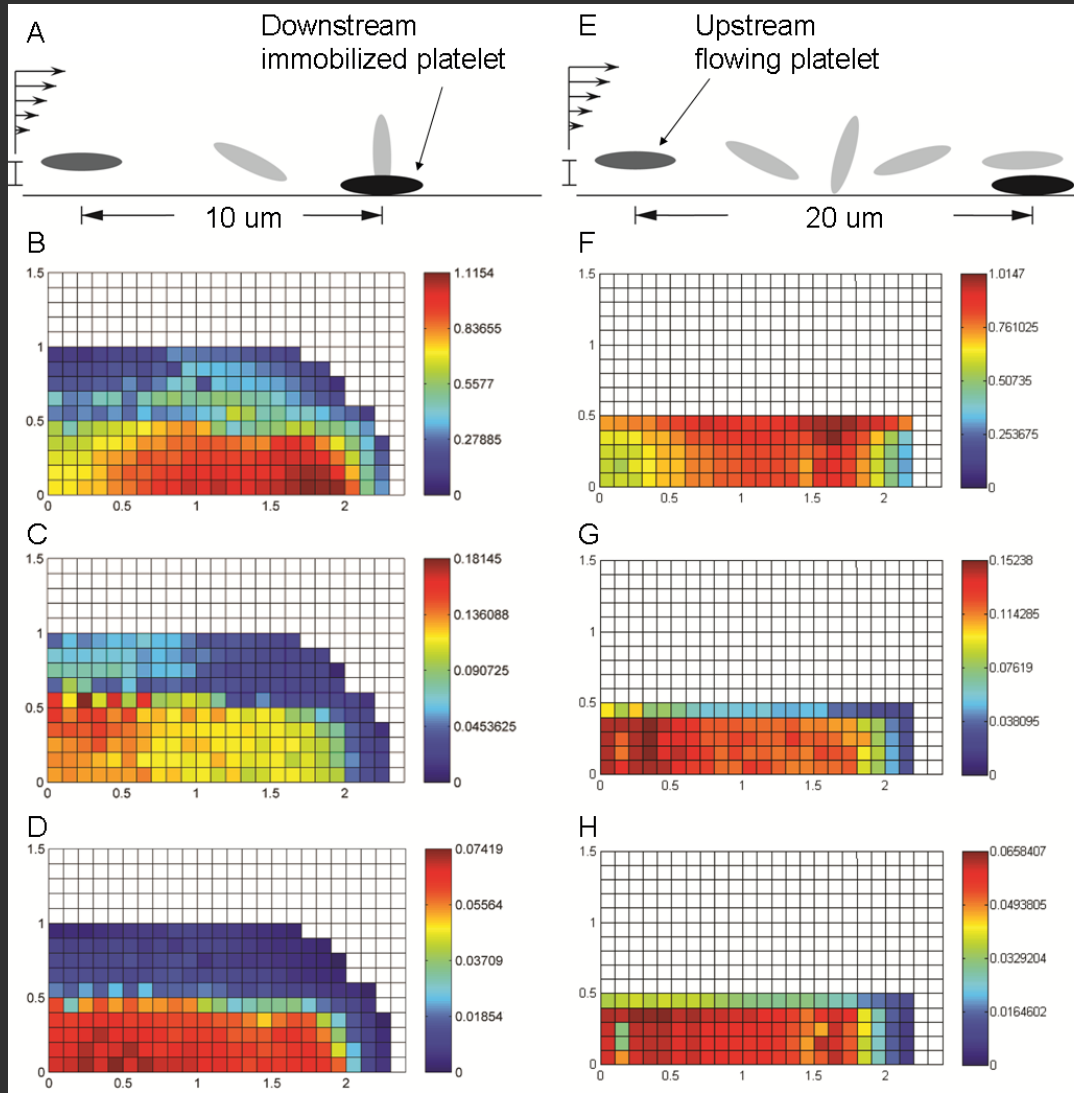
Experimental data from Coburn et al. Biophys. J. Vol 100 Jan 2011 304–312, using human platelet cells. Experiments added 6% ficol to increase viscosity.





# Binary collisions between two platelets

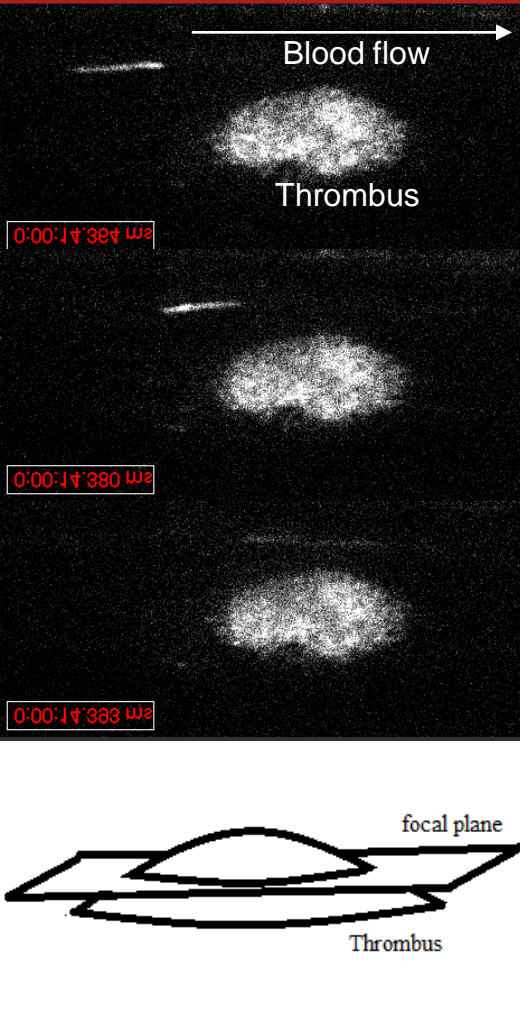
Two Platelets: One adherent to the injury site and one freely flowing



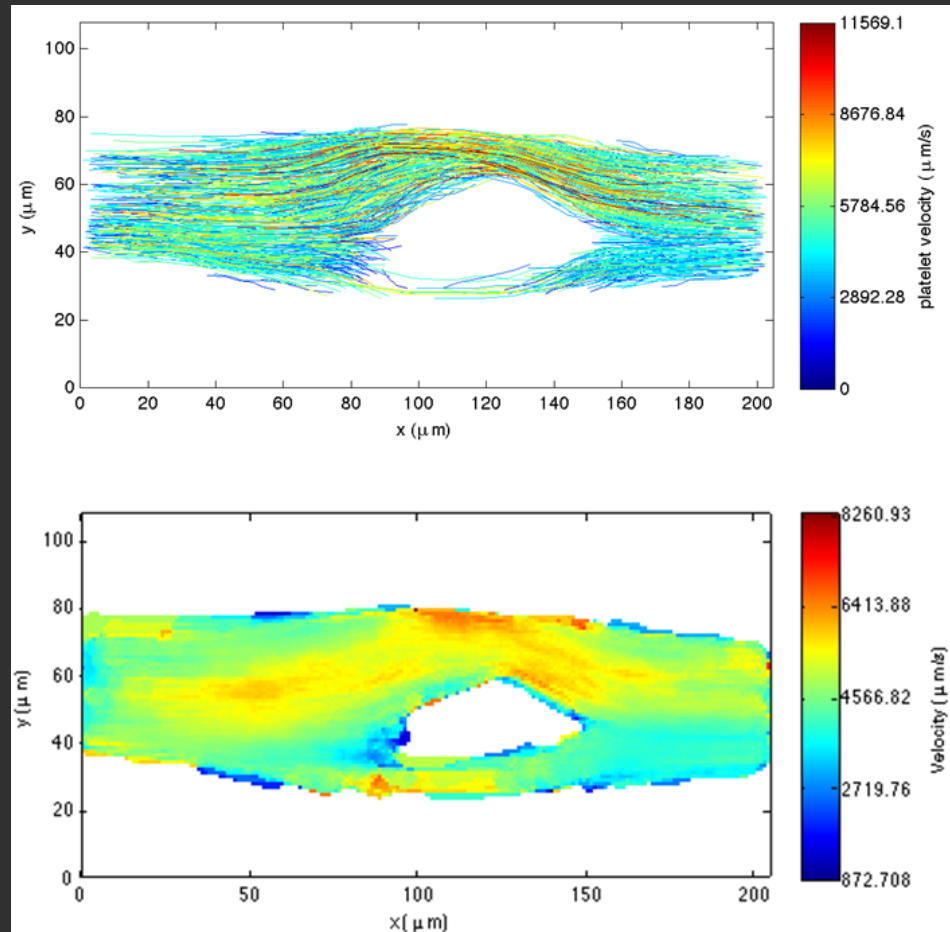
Contact time

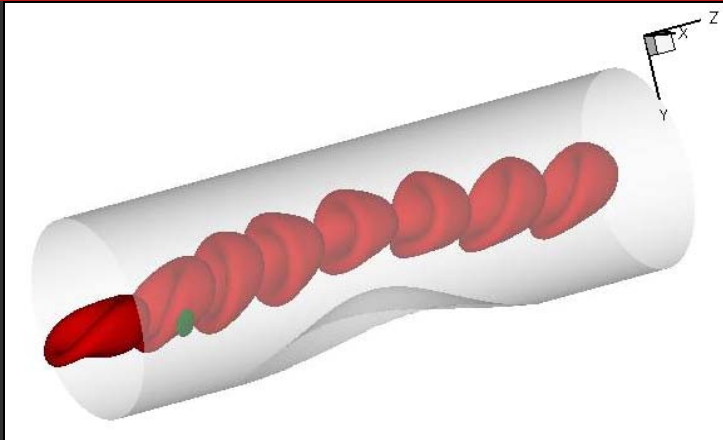
Max contact area

Contact area time integral

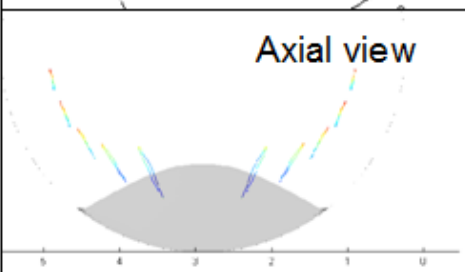
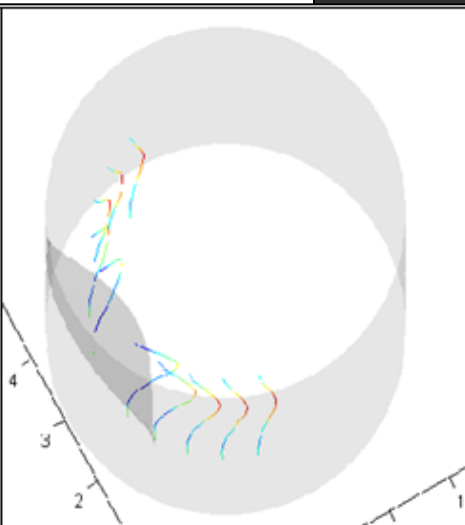
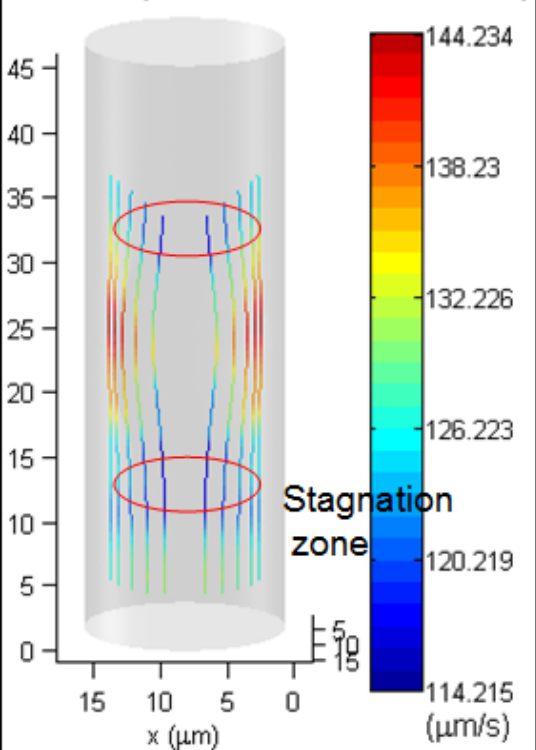


Digital analysis of platelet flow around a platelet-rich thrombus *in vivo*. (~3600 frames, 54 s real time)  
*Up*: cell trajectories color coded by velocity magnitude.  
*Down*: 2-D color map of velocity magnitude.

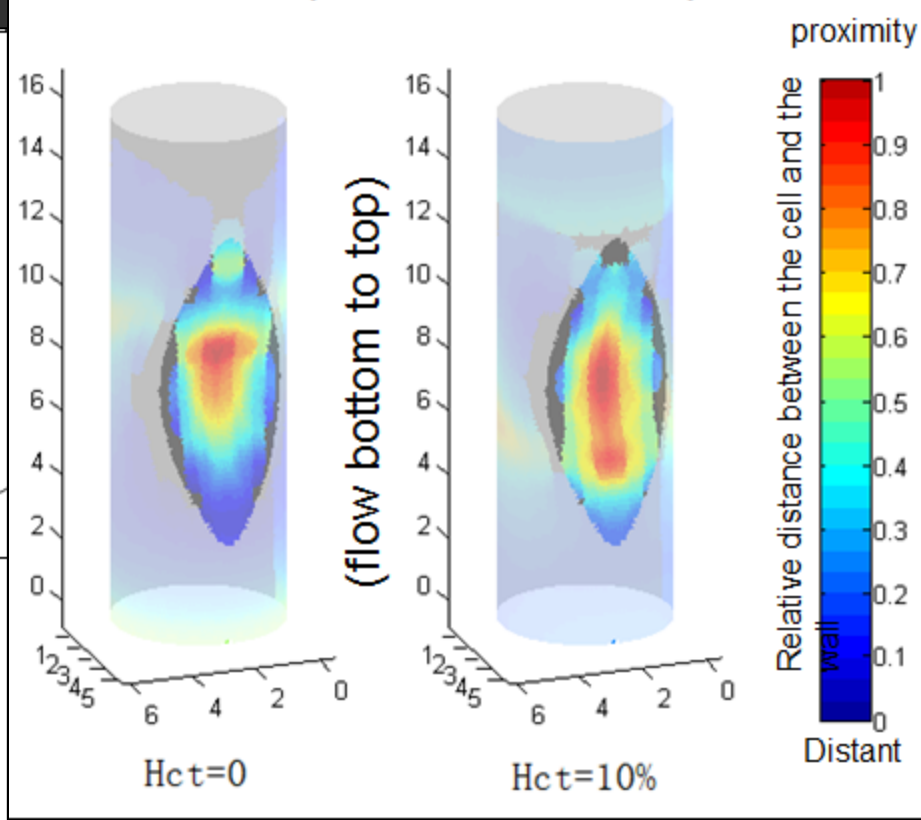




Platelet trajectories colored with velocity



Platelet Deposition Potential Map





## **Platelet Adhesive Dynamics**

### **(multiscale hydrodynamics + molecular kinetics):**

- Flow characteristics of platelet shaped cells
- Characterizing particle-particle collision phenomena
- Model can differentiate between two different kinetic models

## **Vessel scale hydrodynamic calculations**

- Platelet velocity drops upstream and downstream the thrombus
- RBCs enhance platelet deposition onto the thrombus surface.

## **Thrombus self-stabilizing dynamics in vivo:**

- Thrombus shifts towards low shear stress region
- Characterizing the flow pattern around a human-like thrombus in vivo

*Funded by NIH Grant No. HL097971*