

Diana Magnabosco^{1,2}, Henk van Ooijen¹, Bart Bakker¹, Rene van den Ham¹, Luca Formaggia²

1. Philips Research, Molecular Diagnostic Department, Eindhoven, Netherlands

2. Politecnico di Milano, Mathematical Department, Milan, Italy

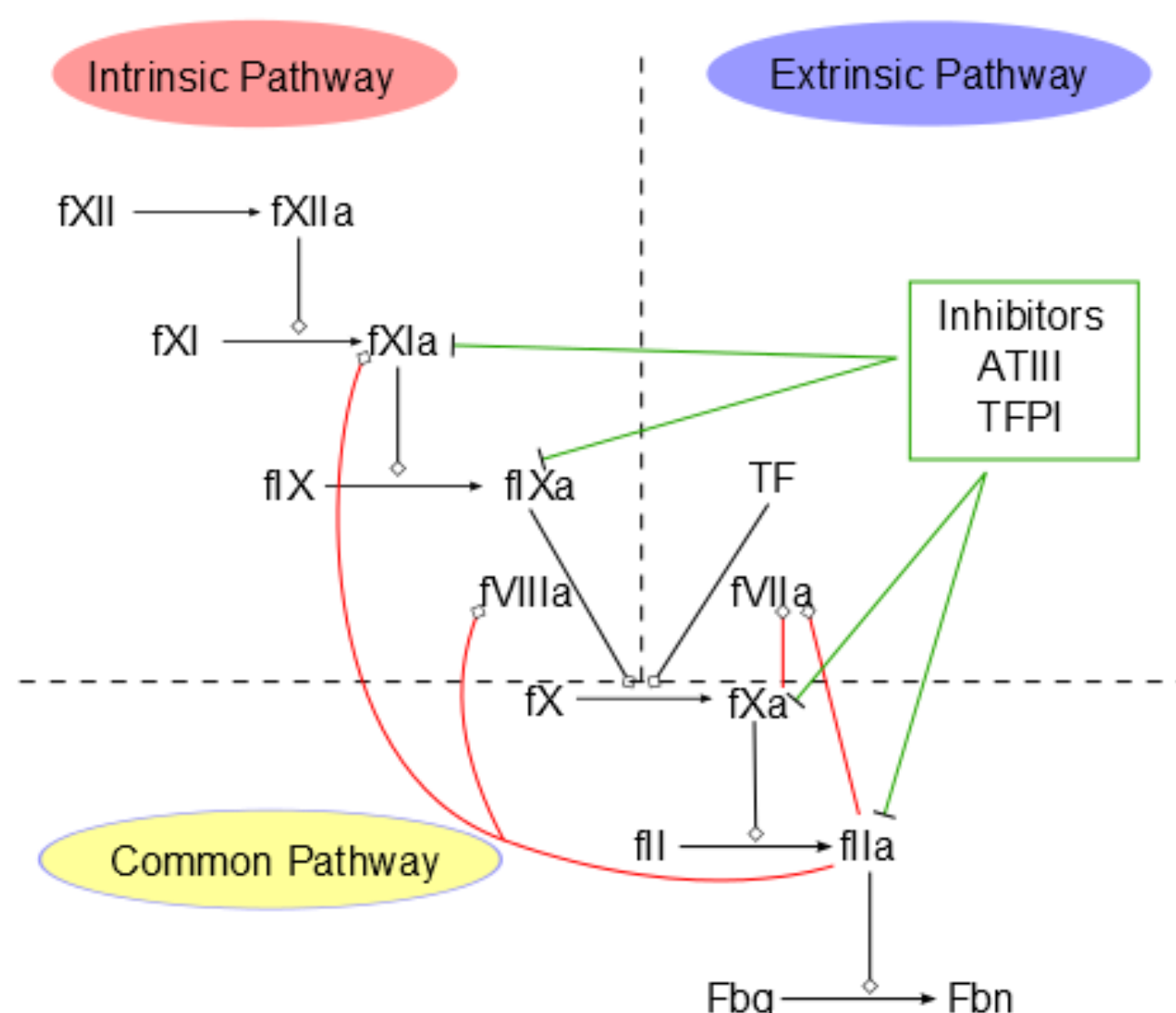
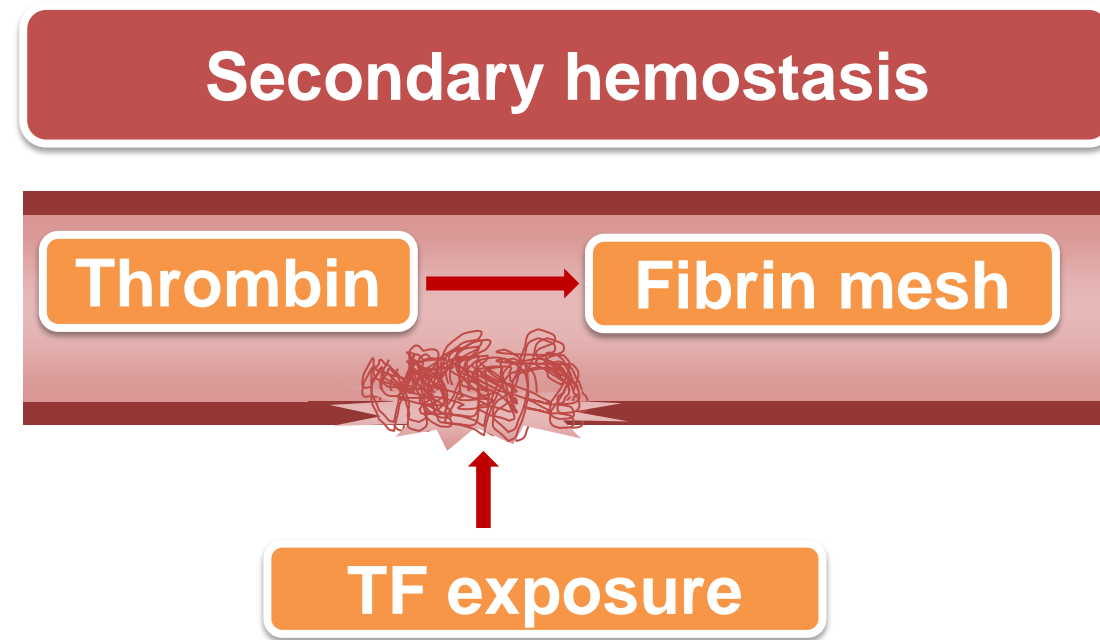
✉ diana.magnabosco@mail.polimi.it, henk.van.ooijen@philips.com

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Biological background

Enzyme reactions, blood flow and diffusion in human vasculature play interacting and fundamental roles in blood coagulation. In this complex mechanism, the balance between blood and clot is a delicate equilibrium between fluid and solid state, whose tight regulation is vital to avoid pathologies such as bleeding and thrombosis.

Exposure of tissue factor after injury triggers the coagulation network, (Figure 1) i.e. a cascade of different consecutively activating protein reactions with feedback loops (red) and inhibitors (green).



Study of the impact of:

- Tissue factor initial concentration (TF_0)
- Wound size (L_w)
- Shear rate (γ)
- Diffusion (D)

Output parameters

- Thrombogram (lag time (t_{lag}), time to peak (t_{max}), maximum concentration (c_{max}) of thrombin)
- TGD (total amount of generated thrombin on the domain)
- Time to clot (t_{clot}) and clot size (A_{clot})

Figure 1: Coagulation network with activation of proteins, positive feedback loops (red) and inhibitors (green)

Mathematical and numerical methods

Coagulation network

- Plasma species in Ω , $t > 0$
 $\frac{\partial c_p(x,t)}{\partial t} - D\nabla^2 c_p(x,t) + \mathbf{u}(x,t) \cdot \nabla c_p(x,t) = R_p(c(x,t))$
- Membrane species on Γ_{down} , $t > 0$
 $\frac{\partial c_m(x,t)}{\partial t} = R_m(c(x,t))$
- Coupling by boundary conditions on Γ_{down} , $t > 0$
 $-\mathbf{n} \cdot D\nabla c_{pm}(x,t) = R_{pm}(c(x,t))$

Linear finite elements method with streamline and crosswind numerical diffusion

46 PDEs for plasma species
11 ODEs on the wall for membrane species
1 ODE in the domain for fibrin

Transport of dilutes species interface
Boundary ODEs and DAEs interface
Domain ODEs and DAEs interface

Blood flow

- Modified Navier-Stokes equations in Ω , $t > 0$
 $\rho \frac{\partial \mathbf{u}}{\partial t} + \rho \mathbf{u} \cdot \nabla \mathbf{u} = \nabla \cdot [-p\mathbf{I} + \mu(x,t)(\nabla \mathbf{u} + \nabla^T \mathbf{u})]$
 $\nabla \cdot \mathbf{u} = 0$
- Viscosity depending on fibrin
 $\mu(x,t) = \mu([Fbn]) = \begin{cases} \mu_{blood} & \text{if } [Fbn] < [Fbn]^* \\ \mu_{clot} & \text{if } [Fbn] \geq [Fbn]^* \end{cases}$

P1+P1 with streamline and crosswind numerical diffusion

Laminar flow interface

Non-linear system: Newton method

Linear system: MUMPS

Time integration: BDF

Mesh: Finer near the wound where gradients are relevant

Tolerances: $\epsilon_{rel}=10^{-3}$ $\epsilon_{abs}=10^{-7}$

Parameters		
L_x	1000 μm	$[Fbn]^*$ 600 nM
L_y	200 μm	TF_0 90 fmol/ m^2
L_w	$L_x/32$	μ_{blood} 0.0035 kg/ms
D	$5 \cdot 10^{-11}$ m^2/s	μ_{clot} 1 kg/ms
γ	1 1/s	ρ 1000 kg/ m^3

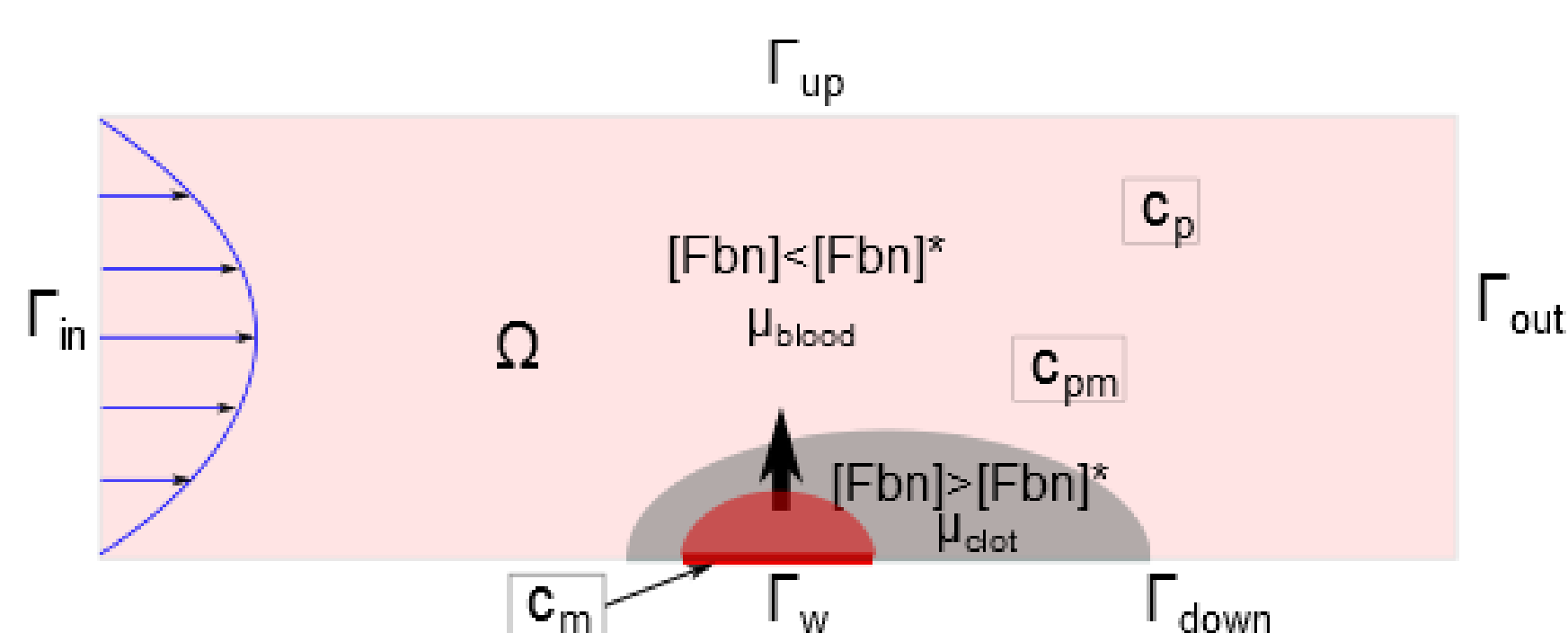
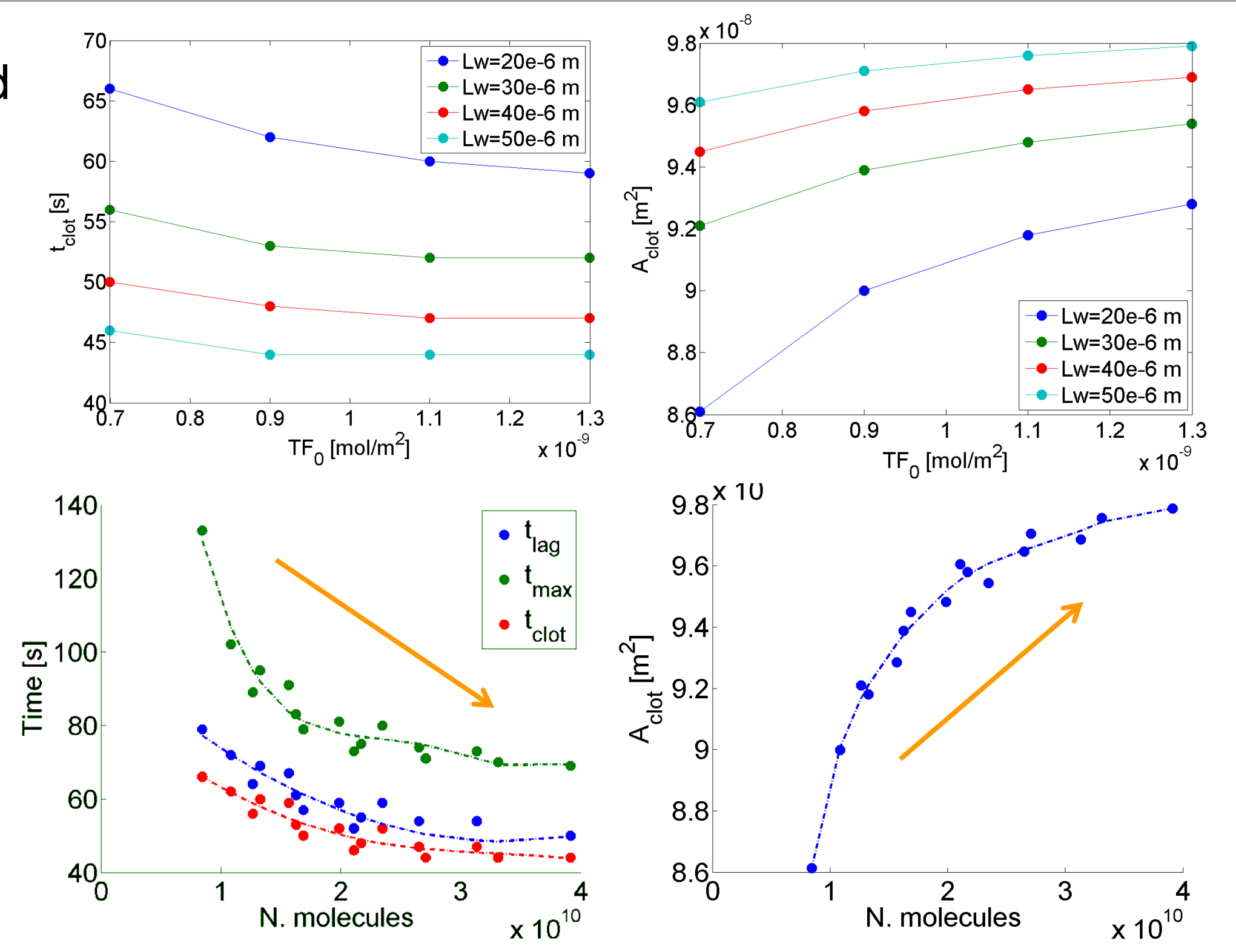


Figure 2: Geometry, boundary conditions and clot formation.

Results

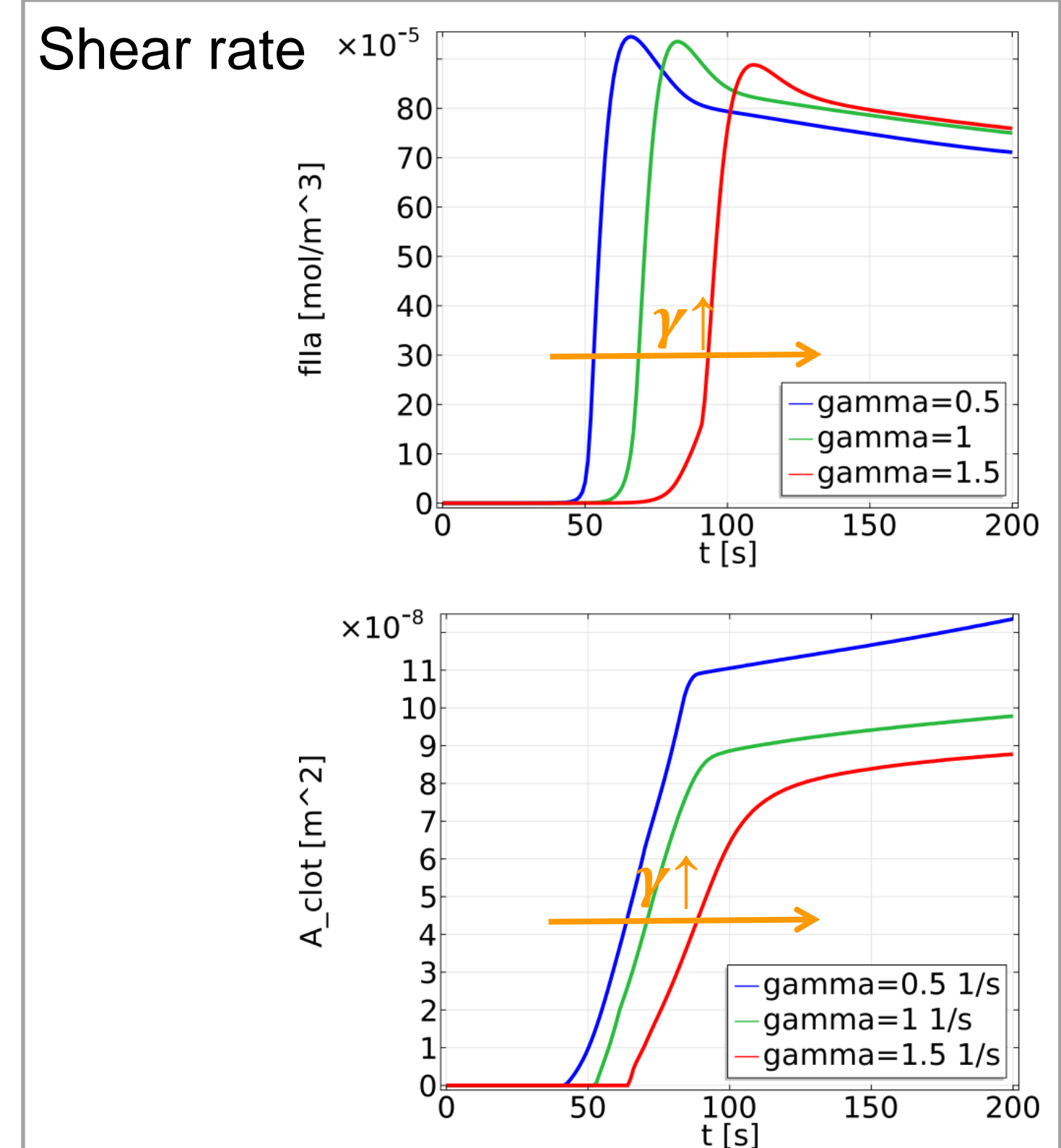
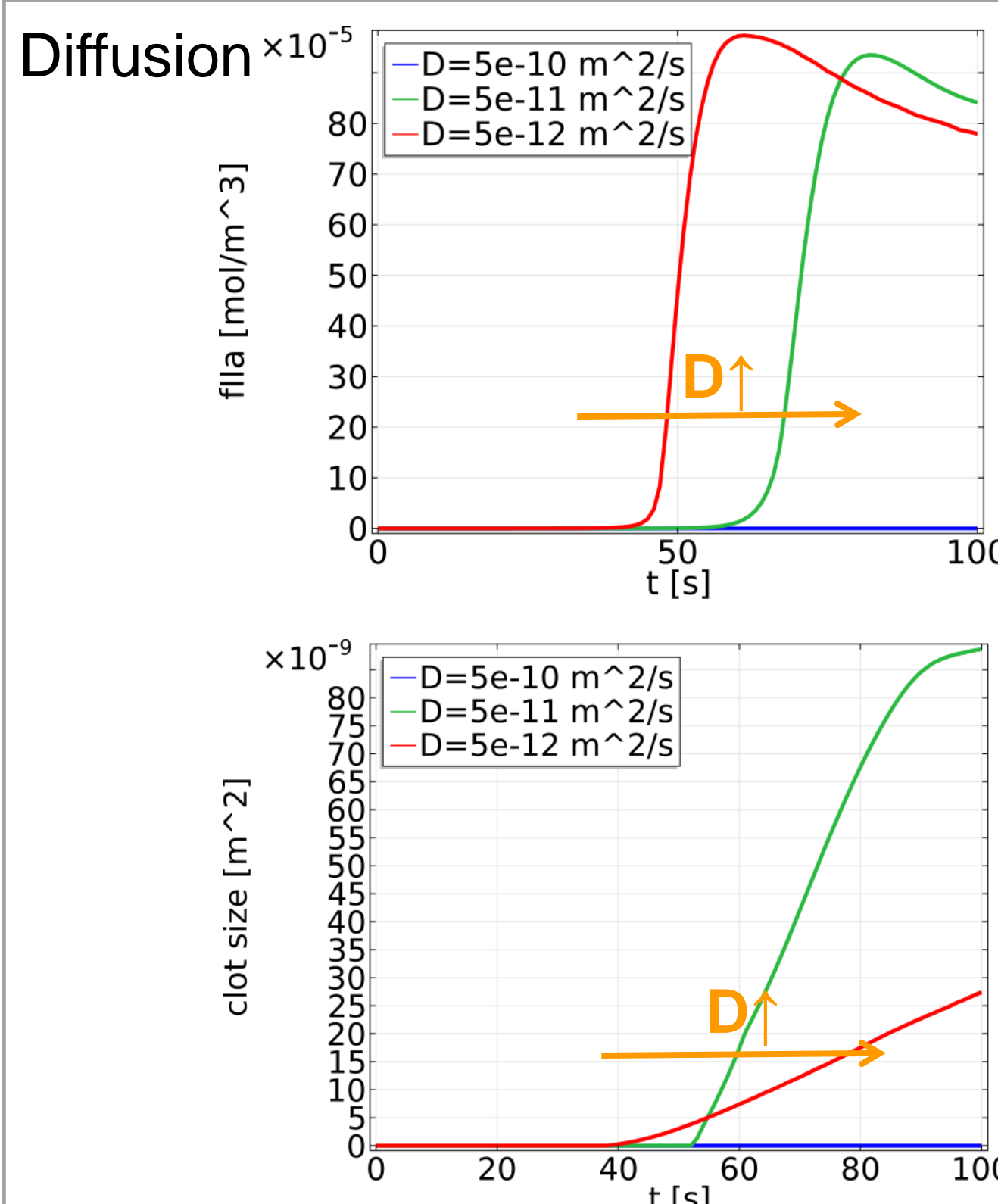
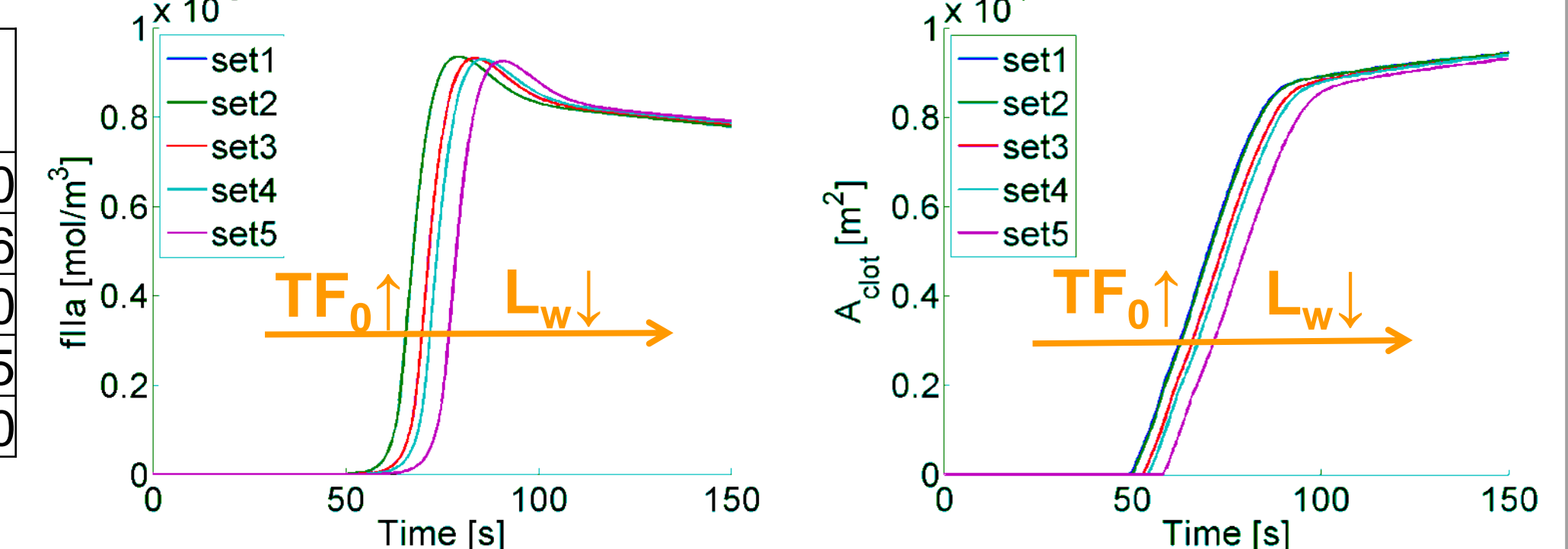
Wound size L_w and tissue factor initial concentration TF_0

TF_0 or $L_w \uparrow$	
t_{lag}	\downarrow
t_{max}	\downarrow
t_{clot}	\downarrow
c_{max}	\uparrow
A_{clot}	\uparrow
TGD	\uparrow



Constant number of molecules of tissue factor on the wound

Set	TF_0 [fmol/ cm^2]	L_w [μm]
1	67.5	40
2	70	38.6
3	90	30
4	110	24.5
5	135	20



Conclusions

- Increasing TF_0 or L_w results in the thrombin burst being stronger and earlier, hence leading to the formation of an early and bigger clot.
- Output parameters are more sensitive to variation in L_w than TF_0 , while keeping the total number of TF molecules fixed. Blood flow in combination with diffusion plays a pivotal role since the inactivated zymogens in the fluid are in contact with the same number of molecules, but stretched over a larger injury site, meaning a larger contact time/region.
- Flow and diffusion have a limiting role on the mechanism. If their values are bigger, the activation is less intense and delayed. The clot is formed later, farther from the wound and is smaller. The incoming inactive species brought by the flow are not enough to balance the removal of the active ones and as a result the balance between the pro-coagulant and anti-coagulant effects of increased flow and diffusion tips in favour of the anti-coagulant effect.

References

M. Anand, K. Rajagopal and K. R. Rajagopal, "A Model Incorporating Some of the Mechanical and Biochemical Factors Underlying Clot Formation and Dissolution in Flowing Blood," *Journal of Theoretical Medicine*, vol. 5, 183–218 (2003).

A. Sequeira, R. Santos and T. Bodnár, "Blood coagulation dynamics: mathematical modeling and stability results," *Mathematical Biosciences and Engineering*, vol. 8, 425–443 (2011).

A. M. Shibeko, E. S. Lobanova, M. A. Panteleev, and F. I. Ataullakhanov, "Blood flow controls coagulation onset via the positive feedback of factor VII activation by factor Xa.," *BMC systems biology*, vol. 4, p. 5, Jan. 2010.