Supplemental Data

Active Site-labeled Prothrombin Inhibits Prothrombinase In Vitro and Thrombosis In Vivo

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FIGURE 18. Inhibition of thrombin generation in platelet-poor human plasma as a function of FPR-ProT at three different TF concentrations. Thrombin generation curves were measured in a total volume of 125 μ l containing 80 μ l platelet-poor human plasma by CAT as described under "Experimental Procedures". Thrombin generation was initiated by 40 pM TF (*A*), 7 pM TF (*B*) or 1.75 pM TF (*C*) in the presence of 30 μ M PSPCPE vesicles, 40 μ g/ml corn trypsin inhibitor, 16 mM CaCl₂, 0.34 mM fluorogenic thrombin substrate, and varying concentrations of FPR-ProT (μ M): 0 (*gray*), 0.2 (*blue*), 0.4 (*green*), 0.6 (*orange*), 0.8 (*red*) and 1.0 (*violet*). All concentrations are final concentrations in the reaction mixtures.



FIGURE 2S. Low TF-initiated thrombin generation in platelet-poor human plasma at varying concentrations of F1, F1.2, FPR-Pre 1, and FPR-thrombin. Thrombin generation curves were measured in a total volume of 125 μ l containing 80 μ l platelet-poor human plasma by CAT as described under "Experimental Procedures". Final concentrations were: 1.75 pM TF, 30 μ M PSPCPE vesicles, 40 μ g/ml corn trypsin inhibitor, 16 mM added CaCl₂, 0.34 mM substrate and 0 (*gray*), 0.2 (*blue*), 0.4 (*green*), 0.6 (*orange*), 0.8 (*red*) or 1.0 (*violet*) μ M F1 (*A*), F1.2 (*B*), FPR-Pre 1 (*C*) or FPR-thrombin (*D*).



FIGURE 3S. Effect of FPR-ProT, F1, F1.2, and FPR-Pre 1 on factor Xa-initiated thrombin generation in platelet-poor human plasma. *A*. Thrombin generation curves were measured in a total volume of 125 μ l containing 80 μ l platelet-poor human plasma by CAT as described under "Experimental Procedures". Thrombin generation was initiated by addition of 0.7 nM human factor Xa in the presence of 30 μ M PSPCPE vesicles, 40 μ g/ml corn trypsin inhibitor , 16 mM CaCl₂, 0.34 mM substrate and 0 (*gray*), 0.2 (*blue*), 0.4 (*green*), 0.6 (*orange*), 0.8 (*red*), 1 (*violet*) or 1.2 (*vellow*) μ M FPR-ProT. *B*. Thrombin generation in platelet-poor human plasma determined as in A, but with 1.2 μ M of the following ProT derivatives: none (*red*); F1 (*orange*); F1.2 (*violet*); FPR-Pre 1 (*blue*). All concentrations are final concentrations in the reaction mixtures.



FIGURE 4S. Competitive equilibrium binding of F1.2 to [6F]FPR-thrombin and native Pre 2 or FPR-Pre 2. The fractional change in fluorescence ($\Delta F/F_o$) of 103 nM [6F]FPR-thrombin as a function of total F1.2 concentration ([F1.2]_o). *A*. in the absence (•) and presence of 3.5 (•) or 10 (\blacktriangle) μ M native Pre 2. *B*. as in *A* in the absence (•) or presence (•) of 10 μ M FPR-Pre 2. The *lines* represent the simultaneous fits by the cubic competitive binding equation, with K_D for F1.2 binding to [6F]FPR-thrombin in *A* of 5.5 \pm 0.5 μ M and in *B* of 5.8 \pm 0.5 μ M, and K_D for F1.2 binding to native Pre 2 in *A* of 0.04 \pm 0.03 μ M, and for F1.2 binding to FPR-Pre 2 in *B* of 3.6 \pm 0.6 μ M. The stoichiometric factor was fixed at 1. Titrations were performed as described in "Experimental Procedures."



FIGURE 5S. Time-courses of cleavage of native Pre 2 or FPR-Pre 2 in the presence of F1.2 by prothrombinase. Concentration of Pre 2 (*A*) or FPR-Pre 2 (*B*) disappearance (•) and appearance of the reaction product, the thrombin B-chain (\circ). Reactions were performed at 37 °C with 2 µM Pre 2 or FPR-Pre 2 in the presence of 2 µM F1.2, 0.7 nM FXa, 50 nM FVa, 50 µM lipid vesicles (PSPCPE), and 60 µM DAPA. The *lines* represent the fits by single exponentials for both substrate depletion and product formation, with the parameters given in "Results". Reactions were performed and analyzed by SDS-gel electrophoresis and quantitative densitometry as described in "Experimental Procedures".



SCHEME 1S. Pathways of native Pre 2 and FPR-Pre 2 activation in the absence and presence of F1.2. Values in *black* are those of Kamath and Krishnaswamy (Ref. 59), and values in *red* are those supplied by the authors. Reprinted in modified form with permission. © 2008 The American Society for Biochemistry and Molecular Biology. All rights reserved.



FIGURE 6S. Time-course simulations of the mechanism in Scheme 1S for reactions of Pre 2 and FPR-Pre 2 with 0.7 nM prothrombinase. A. Simulated time-courses for equimolar mixtures (1 μ M) of native Pre 2 and F1.2, B. FPR-Pre 2 and F1.2, or C. 1 μ M each of Pre 2 and FPR-Pre 2 with 2 μ M F1.2. An expanded view of the initial 100 s of C is shown in D. Formation and disappearance of enzyme-bound complexes are represented as *solid lines* for species derived from native Pre 2 (Pre 2, *blue*; Pre 2·F1.2, *red*; thrombin, *black*; thrombin·F1.2, *green*), and as *dotted lines* for species derived from FPR-Pre 2 (FPR-Pre 2, *blue*; FPR-Pre 2·F1.2, *red*; FPR-thrombin, *black*; FPR-thrombin·F1.2, *green*). Simulations were done with KinTek Explorer (Refs. 64, 65), with bimolecular on-rate constants fixed at 10⁹ M⁻¹ s⁻¹ and off-rate constants to give the rapid-equilibrium dissociation constants shown in Scheme 1S.



FIGURE 7S. Accumulation of platelets and [AF660]-F1.2 following laser-induced injury. Platelets (*green squares*) and [AF660]-F1.2 (*blue triangles*) were monitored for 5 min following laser injury. Notice platelet accumulation follows the reported triphasic pattern and is unaffected by the presence of [AF660]-F1.2.



FIGURE 8S. Time to reach maximum platelet deposition following laser-induced injury in the presence of [AF660]FPR-ProT or F1.2. [AF660]-F1.2 or [AF660]FPR-ProT was titrated into animals prior to laser injury. Median time to achieve maximal platelet signal at each concentration of F1.2 (*blue circles*) and [AF660]FRP-ProT (*red triangles*). Notice the time to peak platelet accumulation increases with the dose of [AF660]FPR-ProT, but is minimally affected by increasing dosage of the Gla domain-containing F1.2.