Platelet binding to polymerizing fibrin is avidity driven and requires activated α IIb β 3 but not fibrin cross-linking

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Key Points

- Platelet interaction with polymerizing fibrin is avidity driven and requires activated αllbβ3 but not fibrin cross-linking.
- The mechanism by which αllbβ3 interacts with polymerizing fibrin differs subtly from the interaction of αllbβ3 with fibrinogen.

The molecular basis of platelet-fibrin interactions remains poorly understood despite the predominance of fibrin in thrombi. We have studied the interaction of platelets with polymerizing fibrin by adding thrombin to washed platelets in the presence of the peptide RGDW, which inhibits the initial platelet aggregation mediated by fibrinogen binding to aIIb_β3 but leaves intact a delayed increase in light transmission (delayed wave; DW) as platelets interact with the polymerizing fibrin. The DW was absent in platelets from a patient with Glanzmann thrombasthenia, indicating a requirement for α IIb β 3. The DW required α IIbb3 activation and it was inhibited by the α IIb β 3 antagonists eptifibatide and the monoclonal antibody (mAb) 7E3, but only at much higher concentrations than needed to inhibit platelet aggregation initiated by a thrombin receptor activating peptide (T6). Surface plasmon resonance and scanning electron microscopy studies both supported fibrin having greater avidity for αIIbβ3 than fibrinogen rather than greater affinity, consistent with fibrin's multivalency. mAb 10E5, a potent inhibitor of T6-induced platelet aggregation, did not inhibit the DW, suggesting that fibrin differs from fibrinogen in its mechanism of binding. Inhibition of factor XIII–mediated fibrin cross-linking by >95% reduced the DW by only 32%. Clot retraction showed a pattern of inhibition similar to that of the DW. We conclude that activated aIIb_{β3} is the primary mediator of platelet-fibrin interactions leading to clot retraction, and that the interaction is avidity driven, does not require fibrin cross-linking, and is mediated by a mechanism that differs subtly from that of the interaction of α IIb β 3 with fibrinogen.

Introduction

Whereas vascular injury leads to sufficient thrombin (Thr) generation within 20 seconds to produce fibrin deposition,^{1,2} and whereas Thr also rapidly activates factor XIII, resulting in fibrin cross-linking via reciprocal transamidation of the C-terminal γ -chain peptides from adjacent fibrinogen molecules, platelets are likely to primarily encounter polymerizing fibrin and cross-linked fibrin in developing and mature thrombi.³⁻⁶ Moreover, the interactions of platelets with these fibrin species are also likely to contribute to clot retraction, which stabilizes clots and diminishes their susceptibility to fibrinolysis.⁷⁻¹⁰ Because patients with Glanzmann thrombasthenia, who lack or have an abnormal α Ilb β 3 receptor, have diminished or absent clot retraction, it is likely that α Ilb β 3 is required for fibrin interaction and clot retraction.¹¹⁻¹³

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For original data, please contact lbuitrago@rockefeller.edu.

The full-text version of this article contains a data supplement.

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Investigators have, however, reported that platelet interactions with fibrin can be supported by GPVI (reviewed by Slater et al,¹⁴ and Perrella et al,¹⁵ and Xu et al¹⁶) and GPIb, either directly or through interactions with von Willebrand factor,¹⁷⁻²³ and thus, these receptors may also participate in the interaction of platelets with polymerizing fibrin and clot retraction.

The interaction between the allbß3 Arg-Gly-Asp (RGD)-binding pocket, formed jointly by the α IIb β -propeller and β 3 β -I domains, and the fibrinogen γ -chain C-terminal sequence 404 to 411, has been defined at the atomic level, 24,25 but the sites on α Ilb β 3 involved in fibrin binding may be different from those that mediate fibrinogen binding because: (1) y-chain Lys406, which interacts with α IIb Asp224,²⁵ is cross-linked to γ -Gln398/Gln399 in crosslinked fibrin,^{26,27} eliminating its ability to make an ionic interaction with Asp224; (2) other sites on fibrinogen and α Ilb β 3, including sites exposed when fibrinogen is converted into fibrin,28,29 have been suggested to support interactions between fibrin and αllbβ3^{30,31}; (3) monomeric fibrin made from γ'/γ' -fibrinogen, which lacks the γ -12 sequence, can interact with purified α IIb β 3³¹; (4) treating allbβ3 with EDTA under conditions that lead to irreversible loss of the ability to bind fibrinogen does not inhibit clot retraction^{32,33}; (5) in contrast to the ability of unactivated α Ilb β 3 to support adhesion to immobilized, intact fibrinogen,34 our recent data indicate that platelet allbß3 activation is required to support adhesion to D-dimer, a plasmin degradation product of cross-linked fibrin that contains the cross-link site⁶; (6) platelet interactions with fibrin also differ from those with fibrinogen with regard to ligand valency. Thus, polymerized fibrin's marked multivalency may contribute to greater avidity, even with the same receptor-ligand affinity.35,36

To study the molecular basis of the interaction of platelets with polymerizing, cross-linked fibrin as well as the impact of fibrin's multivalency, we developed an assay to separate the α IIb β 3-mediated interactions with fibrinogen from those with polymerizing fibrin. We also used surface plasmon resonance (SPR) to compare the affinity of monomolecular α IIb β 3 for fibrinogen vs fibrin.

Materials and methods

Reagents

All reagents were obtained from Sigma-Aldrich unless stated otherwise. We previously described our monoclonal antibodies (mAbs) 10E5 (anti- α Ilb cap domain),³⁷ 7E3 (anti- β 3 near the MIDAS),³⁸⁻⁴⁰ 6D1 (anti-GPIb),⁴¹ and 6F1 (anti- α 2 β 1).⁴² Details on mAb PT25-2, which activates α Ilb β 3,^{43,44} the Thr receptor activating peptide SFLLRN (T6), collagen, collagen-related peptide (CRP), human Thr, ristocetin, soluble human fibrin, fibrinogen, fibrin(ogen) fragments D98 and p-dimer,⁶ peptides GPRP, RGDW, and KQAGDV, acid-citrate-dextrose, and buffers, are in the supplemental data.

Preparation of human platelets

Healthy volunteers. This study was approved by The Rockefeller University Institutional Review Board, and it is Helsinki-compliant. Blood was obtained from healthy volunteers who provided informed consent and who had not taken antiplatelet medications during the 2 weeks before blood collection. The blood was anticoagulated with one-sixth volume of acid-citrate-dextrose. Details on preparation of platelet-rich plasma and washed platelets are provided in the supplemental data. For platelet aggregation studies and studies of the interaction of platelets with polymerizing fibrin, washed platelets were stirred in an aggregometer cuvette and activated with 25 μ M T6 or 0.2 U/mL Thr at the indicated time of agonist injection (TI). The time to the onset of the increase in light transmission is reported as TA. α Ilb β 3 antagonists (RGDW, γ 12-like peptide [KOAGDV], eptifibatide, abciximab, mAb 10E5, and mAb 7E3) were added as indicated 20 minutes prior to initiating aggregation.

Glanzmann thrombasthenia patient. See the supplemental data for case history. Light transmission aggregometry demonstrated no aggregation in response to ADP, epinephrine, collagen, or arachidonic acid, but a normal response to ristocetin. Flow cytometry demonstrated virtually no binding of mAb 7E3. Reverse transcription polymerase chain reaction of platelet RNA identified a homozygous β 3 C575R mutation.

Clot retraction and SEM

Washed platelets (300 μ L; 3 \times 10⁸/mL) were added to a cuvette containing a mixture of 2 mM CaCl₂ and 0.2 U/mL Thr, and clot retraction was monitored by taking photographs of the clots at timed intervals and then analyzing the images for the area of the clot with NIH ImageJ software. For scanning electron microscopy (SEM), clot retraction was performed on a coverslip as described in the supplemental data. The platelet/fibrin suspensions were fixed with 2.5%v glutaraldehyde, 4%v paraformaldehyde, 0.2 M sucrose, and 0.1 M sodium cacodylate, pH 7.4 for 30 minutes at room temperature. They were then deposited on a high-precision glass 1.5 H coverslip, and the samples were dehydrated, dried, sputter-coated with \approx 20 nm gold/palladium particles, and imaged with a field-emission scanning electron microscope at 4 kV as described in the supplemental data.

SPR

The interaction of full-length, purified $\alpha IIb\beta 3$ with either human fibrinogen or soluble fibrin was assessed by analyzing changes in response units using an SPR instrument (Biacore 8K) as described in the supplemental data. Specific binding between the proteins was calculated by subtracting the reference sensorgram from the sample. Data were analyzed with the Biacore Insight Evaluation 3.0.12.15655 software.

Statistical analysis

Results are expressed as means \pm standard deviation (SD) with the number of observations indicated as n. Data were analyzed using GraphPad Prism 6.07. Comparison between samples was determined using the Student *t* test. Differences were considered significant at $P \leq .05$.

Results

The peptide RGDW eliminates the Thr-induced, fibrinogen-mediated, initial wave of platelet aggregation and reveals a delayed wave of increased light transmission (DW) that is mediated by polymerizing fibrin

Because adding Thr to washed platelets results in platelet activation, rapid binding of fibrinogen to $\alpha IIb\beta 3$, platelet

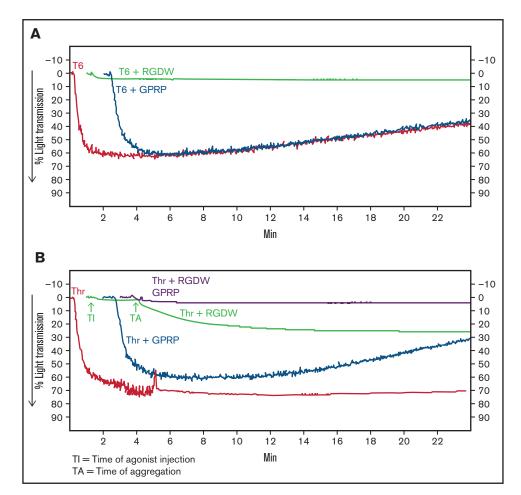
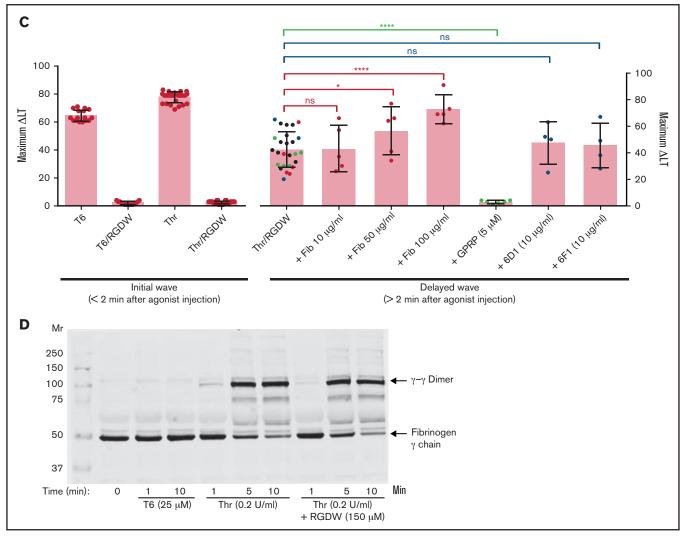


Figure 1. Human platelets treated with the peptide RGDW show a fibrin-mediated delayed wave (DW) of increased light transmission when activated with Thr; fibrinogen enhances the DW, and platelet activation is required for interaction with polymerized fibrin. Washed platelets (2 × 10⁸/mL) were treated with 150 µM RGDW and/or the fibrin polymerization inhibitor Gly-Pro-Arg-Pro (GPRP) for 20 minutes at room temperature and then activated with either (A) 25 µM of a Thr PAR-1 receptor activating peptide (SFLLRN; T6) or (B) 0.2 U/mL Thr in an aggregometer cuvette. Changes in light transmission were measured at 37°C with stirring. (C) Quantitation and statistical analysis of the corresponding maximal change in light transmission (ΔLT) of the T6- or Thr-induced initial wave (<2 minutes after agonist injection) in the presence or absence of RGDW (left panel), and Thr-induced DW (>2 minutes after agonist injection), presented as mean ± SD. Where indicated, samples were treated with human fibrinogen (10, 50, and 100 µg/mL; red dots), GPRP (5 µM; green dots), or mAbs 6D1 or 6F1 (both at 10 µg/mL; blue dots). Data are presented as mean ± SD. Statistical analyses were performed using Student t test between samples tested in the same experiments with and without the indicated intervention (colorcoded). *P < .05; ****P < .0001. ns, nonsignificant. (D) In parallel studies, the reaction was stopped at the indicated times by adding one-third volume of 3× sodium dodecyl sulfate (SDS) sample buffer. Samples were subjected to SDS-polyacrylamide gel electrophoresis under reducing conditions and immunoblotted with an mAb specific for the fibrinogen γ -chain. (E) Addition of fibrinogen at 10 μ g/mL (blue tracing), 50 μ g/mL (purple tracing), 100 μ g/mL (black tracing) enhanced the DW. (F) Effect of added fibrinogen on the time to DW onset. Fibrinogen at the indicated concentrations was added to washed platelet samples treated with 150 µM RGDW in an aggregometer cuvette at 37°C with stirring. After 60 seconds, 0.2 U/mL Thr was added (time of injection), and the time of DW onset was recorded. Data were presented as mean ± SD. Significant differences (P < .05) between groups were estimated using Student t test. *P < .05, **P < .01. (G) Washed platelets were treated with 15 µg/mL PT25-2 for 20 minutes at room temperature. Then, soluble fibrin (100 µg/mL) was added, and the change in light transmission was monitored. All tracings and the immunoblot are representative of at least 3 independent experiments.

aggregation (manifested as an initial wave of increased light transmission), initiation of fibrin polymerization, and activation of factor XIII, we sought to inhibit the rapid interactions between platelets and fibrinogen so that we could observe possible later-occurring interactions between platelets and polymerizing fibrin. In dose-response studies, we found that the peptide RGDW profoundly inhibited platelet aggregation of washed platelets initiated by T6 at 5 to 10 μ M and essentially eliminated it at concentrations of 50 μ M or higher (Figure 1A,C; supplemental

Figure 1A). When Thr (0.2 U/mL) was used to activate the platelets, the rapid platelet aggregation response could be eliminated by RGDW at concentrations >5 to 10 μ M, and a DW of increased light transmission appeared after several minutes (Figure 1B,C; supplemental Figure 1B). Higher doses of RGDW decreased the slope and the maximal increase in light transmission of the DW, but even at 150 to 200 μ M, RGDW failed to inhibit the DW completely (supplemental Figure 1B). Inhibiting fibrin polymerization, as shown by a reduction in γ - γ dimers





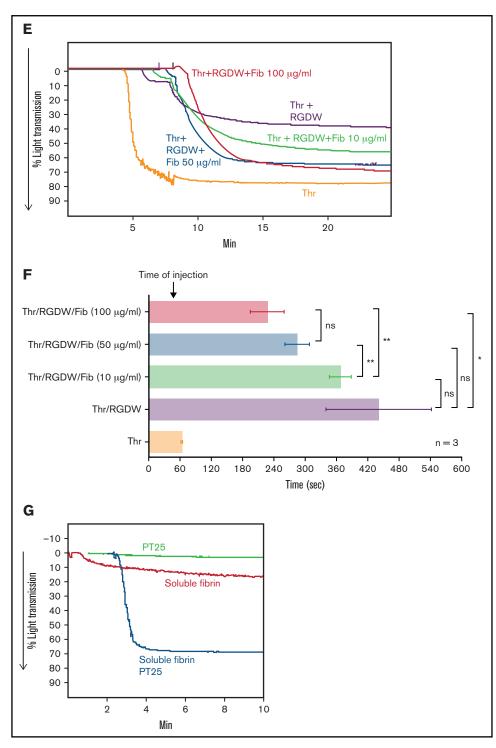
(supplemental Figure 2A), with the peptide GPRP (5 μ M) had no effect on T6-induced platelet aggregation (Figure 1A) and only a minor effect on the initial wave of Thr-induced aggregation in the absence of RGDW, but it eliminated the DW when Thr was combined with RGDW at 150 μ M (Figure 1B-C), indicating that the DW requires fibrin polymerization. Immunoblot analysis of samples removed at different times after adding Thr showed that fibrin cross-linking, as judged by the appearance of γ - γ dimers, began by 1 minute, was >50% complete by 5 minutes, and nearly was complete by 10 minutes (Figure 1D).

The above experiments relied on fibrinogen released from platelets, and so fibrinogen levels were much lower than in plasma ($\approx\!15$ ug/mL 45 compared with 2-4 mg/mL). Adding 10 µg/mL, 50 µg/mL, or 100 µg/mL of fibrinogen to platelets enhanced the DW in a dose-dependent manner (Figure 1C,E). Figure 1F shows that the time to the DW varied inversely as a function of the concentration of fibrinogen, with the greater variability in the absence of added fibrinogen probably reflecting interindividual differences in platelet fibrinogen levels and release of fibrinogen.

Integrin α IIb β 3 activation is required for platelet-fibrin interactions

To assess if integrin activation is required for its interaction with polymerizing fibrin, we used soluble fibrin derived from plasma clots by dissociation using nondenaturing conditions. Under acidic conditions, the fibrin is soluble, but it reassembles into fibrin strands when the solution is neutralized. Immunoblot analysis of the soluble fibrin identified all 3 fibrinogen chains, with >50% of the γ -chains migrating as γ - γ dimers, indicating that it simulates an intermediate time point during fibrin formation and cross-linking (supplemental Figure 2B). Adding soluble fibrin to platelets led to a small (\approx 15%) increase in light transmission over 10 minutes (Figure 1G). Activating α Ilb β 3 with 15 µg/mL PT25-2 dramatically enhanced the increase in light transmission, increasing from \approx 15% to 70%.

T6 and Thr primarily induce platelet-platelet interactions, whereas Thr/RGDW primarily induces platelet-fibrin interactions. SEM of washed platelets at 30 seconds and 2 minutes after treatment with 25 μM T6 showed primarily platelet-





platelet interactions forming variably sized aggregates, whereas platelets treated with 0.2 U/mL Thr for the same period of time showed a mixture of platelet aggregation and platelet interaction with polymerizing fibrin (Figure 2A). In the presence of RGDW, however, there were few platelet-platelet interactions and a predominance of platelet-fibrin interactions. In the Thr/RGDW group, at high

magnification (Figure 2A right 2 panels; Figure 2B), one could see that the fibrin strands had extended lengths of interactions with the surface of single platelets, with some strands interacting >1 µm or more. In addition, Thr/RGDW-treated platelets exhibited a rounder shape and fewer filopodia and lamellipodia than their Thr-treated counterparts. At 4 minutes after Thr treatment in the absence of

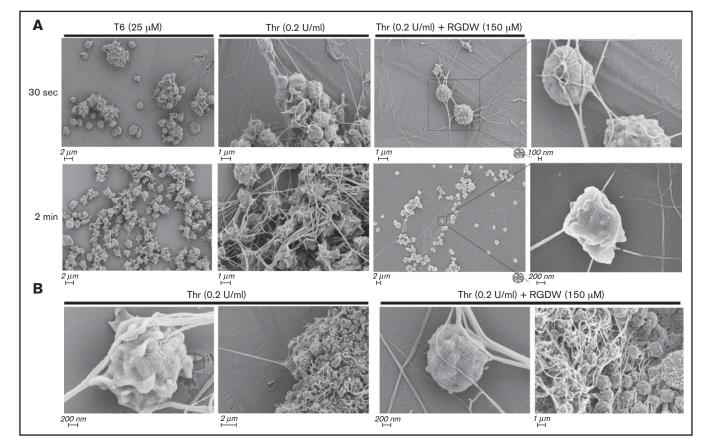


Figure 2. SEM demonstrates that RGDW inhibits platelet-platelet interactions, but not platelet interactions with polymerizing fibrin in the DW. Representative SEM images of washed platelets $(2 \times 10^8/mL)$ stimulated in the aggregometer with the indicated agonist, with or without pretreatment with 150 μ M RGDW. (A) Samples fixed 30 seconds (top row) or 2 minutes (lower row) after initiation of aggregation or onset of the DW. (B) Samples fixed 4 minutes after activation with Thr alone (left 2 panels) or Thr with RGDW (right 2 panels).

RGDW (Figure 2B), there was extensive, tight, platelet-platelet aggregate formation and extensive platelet-fibrin interactions. In contrast, at 4 minutes after the onset of the DW with Thr/RGDW treatment, there was a predominance of single platelet-fibrin interactions.

Platelet glycoprotein receptors GPVI, $\alpha 2\beta 1$, and GPIb play relatively small, if any, roles in mediating the DW

To assess the role of GPVI on the DW, we incubated washed platelets with 2 different inhibitors of platelet aggregation initiated by the interaction of CRP with GPVI, PP2 (100 mM), and Syk inhibitor IV (1 mM). Both inhibitors essentially eliminated CRP-induced aggregation of washed platelets but had minimal effect on the DW (Figure 3A). Consistent with a lack of GPVI ligand engagement, platelet samples obtained during the DW did not demonstrate evidence of signaling downstream of GPVI ligand engagement as judged by Syk or PLC γ 2 phosphorylation (supplemental Figure 3). Antibody 6F1 (10 µg/mL), which inhibits $\alpha 2\beta$ 1-mediated adhesion to collagen,⁴² had little if any impact on the DW (Figures 3B and 1C). Antibody 6D1 (10 µg/mL), which is a potent inhibitor of von Willebrand factor binding to GPIb,⁴¹

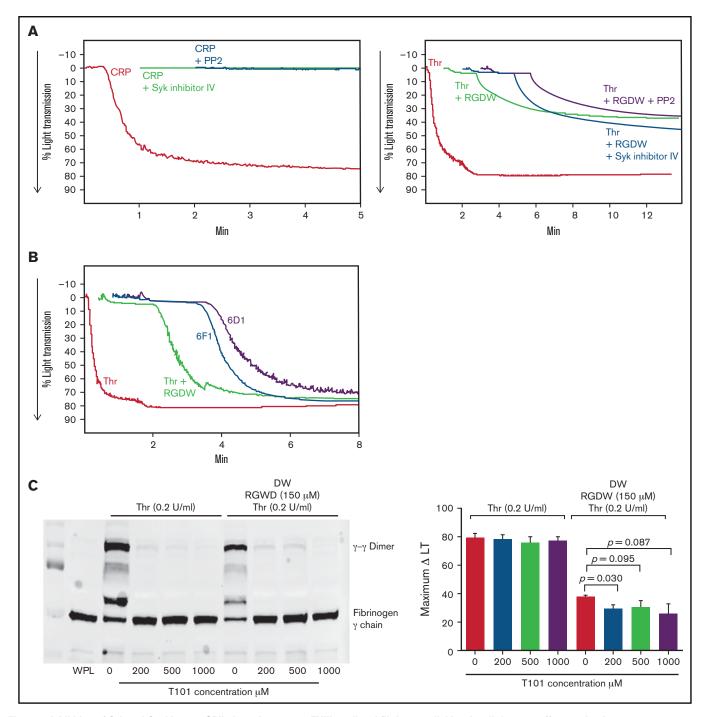
had a modest effect on the DW, reducing the maximal amplitude of the DW by 5% (Figures 3B and 1C).

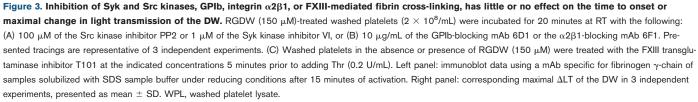
Inhibiting factor XIII-mediated fibrin crosslinking has little impact on the DW

We used the factor XIII inhibitor T101⁴⁶ to test whether crosslinking is required for the platelet-fibrin interactions leading to the DW. Increasing doses of T101 reduced fibrin cross-linking by >95% as judged by the formation of γ - γ dimers (Figure 3C; supplemental Figure 4A). At 200, 500, and 1000 μ M, it decreased the DW by 24% (P = .03), 19% (P = .095), and 32% (P = .087; n = 3) (Figure 3C; supplemental Figure 4B).

Integrin $\alpha \text{IIb}\beta 3$ is the dominant receptor mediating the DW

To assess the role of α Ilb β 3 in the DW, we studied a patient with Glanzmann thrombasthenia whose platelets have virtually no surface expression of the receptor. As shown in Figure 4A, unlike the control platelets, which aggregated in response to both T6 and Thr and had a prominent DW when stimulated with Thr and RGDW, the patient's platelets did not respond to any of these stimuli and did not demonstrate a DW. Even in the presence of preformed soluble fibrin, platelet activation by either T6 (Figure 4B) or PT25-2 (Figure





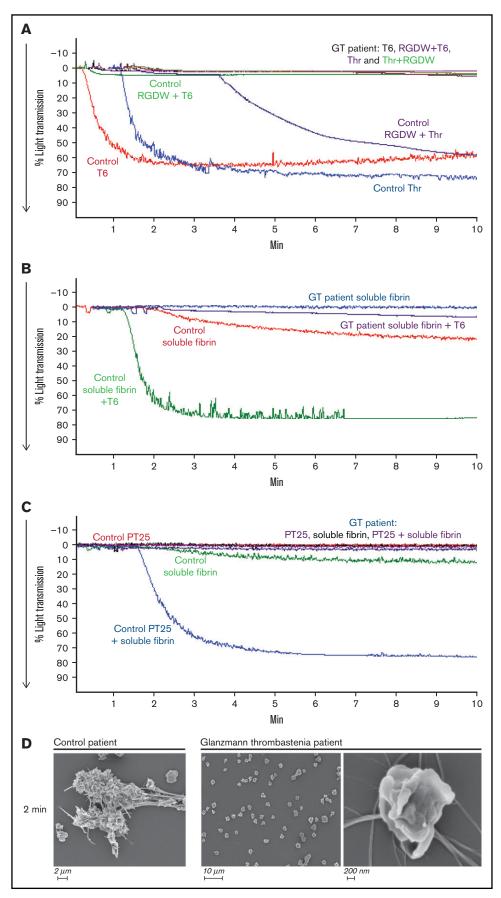


Figure 4.

4C) failed to produce an increase in light transmission. Concordantly, SEM images demonstrated that platelets from the Glanzmann thrombasthenia patient did not interact with fibrin (Figure 4D).

Inhibition of the DW correlates with inhibition of clot retraction

To assess whether the interaction between platelets and polymerizing fibrin in our assay is similar to the interaction between platelets and fibrin that results in clot retraction, we compared the results of our DW assay studies with those found with clot retraction. We found a correspondence between the impact of RGDW in delaying, but not preventing clot retraction (Figure 5A). SEM images demonstrated that in the absence of RGDW, as with Thr-induced platelet aggregation, there are masses of aggregated platelets interacting with thick fibrin strands connecting the platelet aggregates. In the presence of RGDW, there were smaller platelet aggregates, more single platelet-fibrin interactions, and a denser web of fibrin strands (Figure 5B).

To assess whether clot retraction, like the DW, requires integrin activation, we added soluble fibrin to washed platelets in the absence and presence of T6 or PT25-2. In the absence of either activator, there was no clot retraction. In contrast, adding either T6 or the α Ilb β 3-selective and specific activator PT25-2 resulted in clot retraction (Figure 5C).

Small-molecule and mAb α IIb β 3 antagonists variably inhibit the DW, whereas D-dimer and D98 both inhibit the DW

The γ -chain 406-411 KQAGDV peptide, which mediates the interaction of fibrinogen with α IIb β 3 by attaching to the α IIb β 3 RGD-binding pocket,²⁵ dramatically inhibited T6-induced platelet aggregation at 100 to 200 μ M, and partially inhibited T6-induced binding of the mAb PAC-1 (which has higher affinity for activated α IIb β 3 than does fibrinogen, and which binds to the RGD-binding pocket),⁴⁷ but had no effect on the DW at concentrations up to 500 μ M (Figure 6A-C). The RGDW peptide was similarly highly potent in inhibiting T6-induced platelet aggregation, less potent at inhibiting PAC-1 binding, and least potent in inhibiting the DW. Eptifibatide, a high-affinity α IIb β 3 antagonist that also binds in the RGDbinding pocket,⁴⁸ nearly eliminated T6-induced platelet aggregation at just 5 μ M, fully eliminated PAC-1 binding at 100 μ M, but required 500 μ M for near-complete inhibition of the DW (Figure 6A-C).

We then tested 2 different mAbs with nanomolar affinities for $\alpha llb\beta 3$ that are potent inhibitors of fibrinogen binding to activated platelets and platelet aggregation. The mAb 7E3 binds primarily near the RGD-binding pocket in $\beta 3$,³⁷⁻⁴⁰ whereas mAb 10E5 binds primarily to the αllb cap domain.^{37,48} Both antibodies nearly completely inhibited platelet aggregation initiated by T6 at 10 µg/mL (Figure 6D; supplemental Figure 1C,E), but they showed dramatically different

effects on the DW, with 7E3 producing $\approx 75\%$ inhibition at 50 to 100 µg/mL, and 10E5 failing to inhibit the DW at concentrations up to 100 µg/mL (Figure 6E; supplemental Figure 1D,F). SEM analysis confirmed that 7E3 at 100 µg/mL dramatically inhibited the interaction of platelet with fibrin, whereas 10E5 at the same concentration did not inhibit the interaction (Figure 6F). Of particular note, although mAb 7E3 inhibited the DW, abciximab, the chimeric Fab fragment of 7E3, which shares the same antibody binding site, and dramatically inhibits T6-induced platelet aggregation at 5 µg/mL (Figure 6D), produced only minimal inhibition of the DW at concentrations up to 100 µg/mL (Figure 6E).

To localize the binding site for activated platelets on polymerizing fibrin, we added purified D-dimer at 200 μ g/mL and found that it dramatically inhibited the DW, as did D98 at the same concentration (Figure 6G).

SPR studies with purified, monomeric $\alpha IIb\beta 3$ show similar affinities for immobilized fibrinogen and fibrin, in both the absence and presence of priming by mAb PT25-2

In order to assess the relative contributions of affinity and avidity differences in the interaction of allbβ3 with fibrinogen and fibrin, we measured the affinity of purified, monomeric allbB3 to immobilized fibrinogen and immobilized soluble fibrin in the absence and presence of the activating mAb PT25-2 (Figure 7A-B). We found that the binding of $\alpha IIb\beta 3$ to both fibrinogen and fibrin best fit a heterogeneous ligand binding model (χ^2 values between 3 and 27), which is based on the interaction between 1 analyte, in our data α Ilb β 3, with 2 different ligands (or potentially a single ligand with 2 different conformations), in our case either fibrinogen or fibrin.⁴⁹ The data fit a 1:1 model least well (χ^2 values between 65 and 619), whereas the fit to a bivalent analyte model was similar to, but not guite as good as, the heterogeneous model (χ^2 values between 4 and 601) (supplemental Figure 5A-B). The k_{a1} for fibrinogen (1.21 \times 10⁴) was similar to the k_{a1} for fibrin (1.34 \times 10^4), as were their k_{d1} values (2.79×10^{-3} and 1.96×10^{-3} , respectively), resulting in very similar K_{d1} values (2.31 \times 10⁻⁷ and 1.47 \times 10⁻⁷, respectively). Both the ka2 and kd2 values for both ligands were substantially higher than the k_{a1} and k_{d1} values, but the values for fibrinogen and fibrin were again similar, resulting in similar K_{D2} values (5.02 imes 10^{-7} and 3.79×10^{-7}). Priming allb β 3 with PT25-2 increased the ka1 values for both fibrinogen and fibrin approximately threefold without materially changing the k_{d1} values, resulting in K_{d1} values of 8.50×10^{-8} and 4.78×10^{-8} , respectively. Priming lowered both the k_{a2} and the k_{d2} values to similar extents, resulting in similar K_{D2} values (2.31 \times 10⁻⁷ and 1.37 \times 10⁻⁷, respectively). In the presence of either mAb 7E3 or mAb 10E5, the binding of allbß3 to fibrinogen (response units) was dramatically decreased (supplemental Figure 6A). In contrast, as with their differential impact on the DW, mAb 7E3 was better able to inhibit the binding of α IIb β 3 to

Figure 4. Platelets from a patient with Glanzmann thrombasthenia do not aggregate with T6 or Thr, do not demonstrate a DW in response to Thr + RGDW, and do not interact with soluble fibrin when activated with T6 or mAb PT25-2. Fibrinogen (50 μ g/mL) was added to washed platelets (2 × 10⁸/mL) from a healthy donor (Control) or a patient with Glanzmann thrombasthenia (GT), and then the platelets were treated with the following: (A) T6 (25 μ M), Thr (0.2 U/mL), T6 (25 μ M) + RGDW (150 μ M), or Thr (0.2 U/mL) + RGDW (150 μ M). (B) Soluble fibrin (100 μ g/mL) was added to washed control or patient platelets and then the platelets were either untreated or treated with 25 μ M T6. (C) Washed control or patient platelets (2 × 10⁸/mL) were treated with 15 μ g/mL PT25-2 for 20 minutes at room temperature, and then soluble fibrin (100 μ g/mL) was added and the change in light transmission was monitored. (D) SEM of samples from a healthy donor and the GT patient 2 minutes after the onset of the DW in the healthy donor.

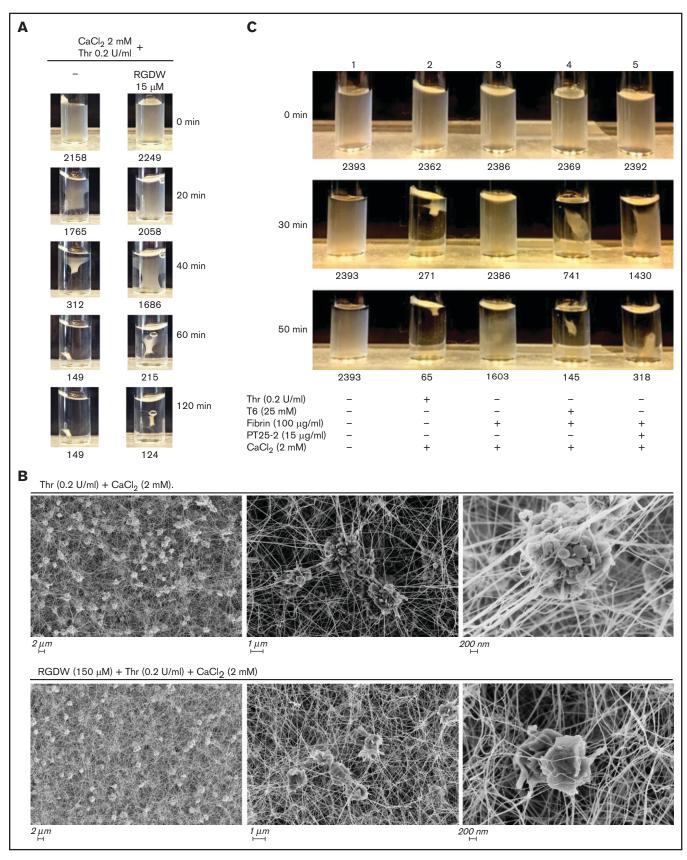


Figure 5.

fibrin (supplemental Figure 6B). A repeat experiment gave very similar results.

Discussion

Our data demonstrate that the interaction of Thr-activated platelets with fibrinogen, and the ensuing platelet aggregation, can be differentiated from the interaction of Thr-activated platelets with polymerizing fibrin by including the RGDW peptide; the latter eliminates the initial platelet aggregation response, but leaves intact a DW of platelet interactions with fibrin, which in turn can be eliminated by a peptide (GPRP) that prevents fibrin polymerization.⁵⁰ We found that the DW has an absolute requirement for allbß3 because it was absent when platelets from a patient with Glanzmann thrombasthenia were substituted for control platelets. The DW could be inhibited by the potent αllbβ3 antagonists eptifibatide and mAb 7E3, but much higher concentrations were required than those needed to inhibit fibrinogen-mediated platelet aggregation induced by T6. To differentiate whether the requirement for higher concentrations of the inhibitors was due to αllbβ3 having higher affinity or avidity, or both, for fibrin vs fibrinogen, we performed SPR studies in which monomeric α IIb β 3 in solution bound to either immobilized fibrinogen or soluble fibrin. We found that the binding to each best fit a heterogeneous ligand binding model and that the k_{a1} , k_{d1} , k_{a2} , and k_{d2} values were similar for both fibrinogen and fibrin, suggesting that the affinity of the receptor was similar for both proteins. By inference, this suggests that the stronger interaction with fibrin reflects increased avidity due to the simultaneous engagement of multiple allbß3 receptors. Our SEM studies of the interaction of platelets with polymerizing fibrin are consistent with this interpretation because platelets make intimate contact with extended regions of the polymerizing fibrin strands, in some cases several microns in length. Whereas α Ilb β 3 receptors are only \approx 200 Å apart, on average, on the platelet surface, the interacting polymerizing fibrin strands likely contact many allbß3 receptors. This would translate into dramatically enhanced avidity as the cumulative off-rate is the product of the individual off-rates of the interactions.35 Although our SPR data fit the heterogeneous model, in which there are 2 different ligands and a single analyte, they also fit the bivalent analyte model, in which the analyte makes 2 interactions, nearly as well. These results are consistent with the complex pattern of fibrinogen and fibrin interaction with allbß3 in which there are multiple steps and interacting sites on the receptor and ligand.

The difference in the ability of mAb 7E3 and its chimeric Fab fragment, abciximab, to inhibit the DW is striking because they share the same antibody binding site.⁵¹ We previously showed that mAb 7E3 binds bivalently to α IIb β 3 on platelets,⁵² and thus, its off-rate will be the product of the off-rates for each binding site. In contrast, as a Fab fragment, abciximab can only make monovalent contact and thus will have lower avidity even though the affinity of the individual interactions should be the same. Thus, the bivalent binding of mAb 7E3 is sufficient to overcome the greater avidity of the interaction of α IIb β 3 with polymerizing fibrin than with fibrinogen. Of note, Collet et al also found that that abciximab did not prevent the interaction of platelets with fibrin during in vitro Thr-induced clotting, but by inhibiting platelet-platelet interactions, it facilitated lysis by tissue plasminogen activator.^{7,53,54} Similar results were found by Braaten et al using an RGD peptide.⁵⁵

Surprisingly, mAb 10E5, a potent inhibitor of fibrinogen binding to $\alpha IIb\beta3$ (K_D \approx 16 nM), and of platelet aggregation initiated by multiple agonists, 37 did not inhibit the DW even at high concentrations. This raises the possibility that fibrin binding is less dependent on interacting with the αIIb cap domain, which is 10E5's epitope, 48 than the $\beta3$ specificity-determining loop and $\beta1\text{-}\alpha1$ loop in the I domain, which is where 7E3 binds. 39,40 These new data are in accord with our previous finding that 10E5 is also less potent at inhibiting platelet adhesion to D-dimer and clot retraction.

We previously reviewed the potential for the fibrinogen γ 404-411 sequence to mediate the interaction of α IIb β 3 with p-dimer,⁶ and the same analysis applies to cross-linked fibrin because in both cases the Lys406 will not be available for an ionic interaction with α IIb Asp224. Thus, it is possible that the γ 407-411 peptide can bind to the β 3 subunit via coordination of the MIDAS Mg²⁺ by the Asp410 carboxyl group and via an interaction of Val411 (through an intermediary water molecule) with the ADMIDAS (Adjacent to MIDAS) Ca²⁺,²⁵ if the peptide retains sufficient accessibility and flexibility. The ability of fibrinogen fragment D98, which lacks the γ 407-411 sequence, to inhibit both clot retraction⁶ and the DW does not exclude a potential role for γ 407-411, because D98's inhibition may reflect it containing a necessary ancillary binding site for fibrin binding, or allosteric inhibition if it interacts with the polymerizing fibrin.

Litvinov et al used optical trap-based force spectroscopy to study the interaction of α Ilb β 3 (activated with MnCl₂) with fibrinogen or fibrin monomer.³¹ They found that the interaction with fibrin had higher binding probability and higher rupture force but noted that a substantial percentage of the α Ilb β 3-fibrin interactions was probably multimolecular. Thus, they could not formally differentiate between affinity and avidity effects. They also found that an RGD peptide, eptifibatide, and abciximab were less effective in inhibiting the interaction with fibrin compared with fibrin, which is in accord with our finding with the DW.

We found little evidence of a major contribution of either GPVI or GPIb to the DW because neither inhibitor of GPVI signaling nor GPIb binding to von Willebrand factor had a strong impact. Moreover, we could not identify evidence of GPVI signaling as a result of the DW. Our data differ from those of others who used different assays,¹⁴⁻¹⁶ and we did not perform studies of the role of GPIb

Figure 5. As with the DW, RGDW only partially inhibits clot retraction; α IIb β 3 activation is required for retraction of preformed fibrin. (A) Washed platelets (3 × 10⁸/mL) were treated with the RGDW (150 μ M) for 20 minutes at room temperature and then transferred to an aggregometer cuvette containing 0.2 U/mL Thr and 2 mM CaCl₂ to initiate clot retraction. The area of the clot is shown underneath each photograph and expressed in square pixels. (B) SEM images of clot retraction after 15 minutes. (C) Washed platelets were untreated (sample 1) or stimulated to undergo clot retraction with the following: Thr + CaCl₂ (sample 2), or soluble fibrin + CaCl₂ alone (sample 3), or in combination with 15 μ g/mL of the mAb PT25-2 (sample 4), or 25 μ M T6 (sample 5). Photographs of the time course of clot retraction are shown. The area of the clot is shown underneath each photograph and expressed in square pixels. Data shown in panels A and C are representative of at least 3 independent experiments.

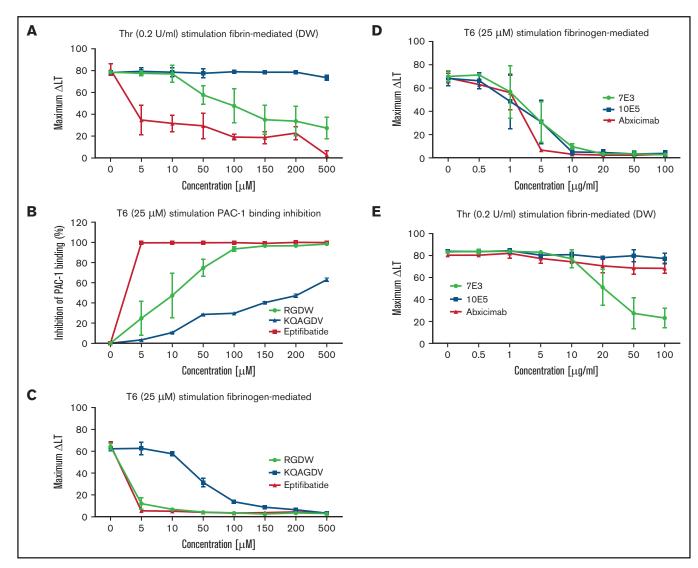


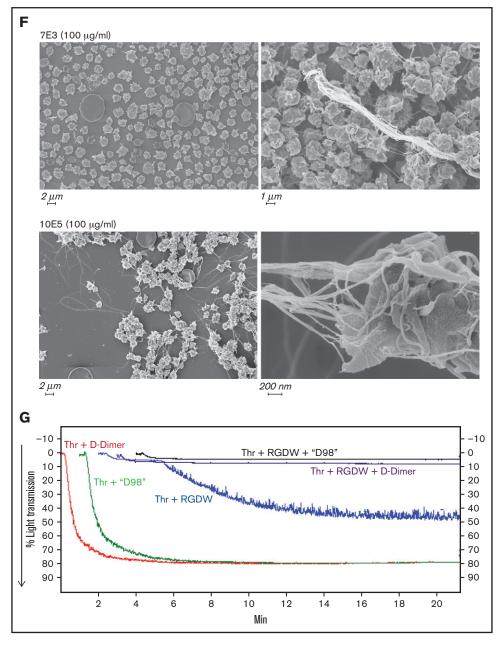
Figure 6. Integrin α **IIb** β **3 antagonists differentially affect the DW and platelet-polymerizing fibrin interactions, and p-dimer and D98 inhibit the DW.** Washed platelets (2 × 10⁸/mL) were treated for 20 minutes at room temperature with increasing doses of the peptides RGDW or KOAGDV, eptifibatide, the mAbs 7E3 or 10E5, or the chimeric mAb 7E3 Fab fragment abciximab. After incubation, platelets were treated with 25 μ M T6 (A,D) or 0.2 U/mL Thr (C,E) and stirred in an aggregometer at 37°C. The change in light transmission 15 minutes after adding the agonist was recorded as the maximal Δ LT. (B) In parallel studies, untreated and α IIb β 3 antagonist-treated platelets (2 × 10⁶/mL) were activated with T6 (25 μ M) under static conditions without stirring for 5 minutes and incubated with Alexa⁴⁸⁸-labeled PAC-1 for 20 minutes, after which platelet-associated fluorescence was analyzed by flow cytometry. The data are reported as the mean \pm SD inhibition of the geometric mean fluorescence in the absence of each antagonist. (F) SEM of samples of washed platelets 2 minutes after the onset of the DW in the presence of either mAb 7E3 or 10E5. (G) Adding purified p-dimer or D98 at 200 μ g/mL to the washed platelets inhibited the DW. One of 3 similar experiments with p-dimer and D98 (mean DW \pm SD without and with p-dimer equal 42 \pm 6.5 and 5.3 \pm 3.2 [*P* < .005]; mean DW \pm SD without and with D98 equal 42 \pm 6.5 and 1.7 \pm 0.5 [*P* < .005]).

under controlled high shear force conditions, so additional studies are required to assess the relative contributions of these other receptors.

We found that platelets showed relatively little interaction with soluble fibrin unless activated with T6 or PT25-2. These data are in accord with our previous data demonstrating that platelet activation is required for adhesion to D-dimer,⁶ and data from multiple groups that platelet activation is required for clot retraction.⁵⁶⁻⁵⁸ We previously found that D-dimer can inhibit clot retraction as can a polyclonal antibody to D-dimer.⁶ In the present study, we also found that

D-dimer can inhibit the DW, another similarity between these phenomena.

There are conflicting data on whether FXIII activity is required for platelet-mediated clot retraction, which requires platelet interactions with polymerizing fibrin and force generation.⁵⁹⁻⁶⁶ Results with mouse blood also vary, with Kasahara et al reporting that clot retraction requires cross-linked fibrin,^{67,68} whereas Kattula et al found equivalent retraction in wild-type and F13a^{-/-} mice.⁴⁶ In the presence of red cells, the effect of FXIII on entrapping red cells via α -polymer formation complicates the analysis, but this does not





bear on the effect of FXIII on the interaction of α IIb β 3 with fibrin.^{46,69,70} Although we found evidence of fibrin cross-linking before the DW, inhibiting FXIII-mediated cross-linking by >95% reduced the DW by only 32%.

Our data have potentially important clinical significance. In particular, the higher avidity of fibrin for α Ilbβ3 than fibrinogen means that the current α Ilbβ3 antagonists may not be effective in inhibiting platelet fibrin interactions, even when they dramatically inhibit platelet-fibrinogen interactions and platelet aggregation. For example, eptifibatide reaches plasma levels of at most 2 to 3 μ M,⁷¹ which is enough to profoundly inhibit T6-induced platelet aggregation and PAC-1 binding, but only inhibit the DW by ~50%. Abciximab is even less potent in inhibiting the DW at the peak concentration it reaches in vivo after

bolus administration.⁷² It is thus possible that new α Ilb β 3 antagonists with very high affinity may be able inhibit platelet-fibrin interactions, and potentially be more effective as antithrombotic agents.

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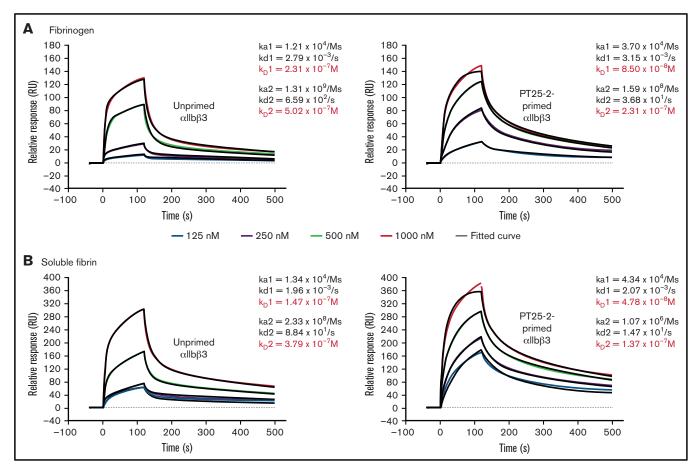


Figure 7. SPR analysis demonstrates that α**IIbβ3 has similar affinity for fibrinogen and fibrin.** Purified human α**IIb**β3 binding to (A) immobilized fibrinogen and (B) soluble fibrin. Where indicated, purified integrin was treated with 15 µg/mL PT25-2 at room temperature for 20 minutes. Kinetic values were obtained by fitting data to a heterogenous ligand interaction model.

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Authorship

Contribution: L.B. designed, performed, and analyzed functional experiments and wrote the manuscript; S.L. performed and analyzed SEM experiments; O.B. contributed the clinical assessment of Glanzmann thrombasthenia patient and wrote the manuscript; J.P. analyzed SPR data and wrote the manuscript; B.C. designed,

oversaw, and analyzed the data and had primary responsibility for writing the manuscript.

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