

Binding strength and activation state of single fibrinogen-integrin pairs on living cells

Rustem I. Litvinov*, Henry Shuman†, Joel S. Bennett‡, and John W. Weisel*§

*Department of Cell and Developmental Biology, †Department of Physiology, and ‡Hematology–Oncology Division of the Department of Medicine, University of Pennsylvania School of Medicine, Philadelphia, PA 19104-6058

Communicated by William F. DeGrado, University of Pennsylvania School of Medicine, Philadelphia, PA, April 3, 2002 (received for review December 4, 2001)

Integrin activation states determine the ability of these receptors to mediate cell–matrix and cell–cell interactions. The prototypic example of this phenomenon is the platelet integrin, α IIb β 3. In unstimulated platelets, α IIb β 3 is inactive, whereas exposing platelets to an agonist such as ADP or thrombin enables α IIb β 3 to bind ligands such as fibrinogen and von Willebrand factor. To study the regulation of integrin activation states at the level of single molecules, we developed a model system based on laser tweezers, enabling us to determine the rupture forces required to separate single ligand-receptor pairs by using either purified proteins or intact living cells. Here, we show that rupture forces of individual fibrinogen molecules and either purified α IIb β 3 or α IIb β 3 on the surface of living platelets were 60 to 150 pN with a peak yield strength of 80–100 pN. Platelet stimulation using either ADP or the thrombin receptor-activating peptide enhanced the accessibility but not the adhesion strength of single α IIb β 3 molecules, indicating that there are only two states of α IIb β 3 activation. Thus, we found it possible to use laser tweezers to measure the regulation of forces between individual ligand-receptor pairs on living cells. This methodology can be applied to the study of other regulated cell membrane receptors using the ligand-receptor yield strength as a direct measure of receptor activation/inactivation state.

The detailed biochemistry of receptor–ligand interactions can be determined from solution and/or surface studies, but these results do not take into account the response of receptor-mediated cell functions to externally applied forces encountered *in vivo*. This consideration is particularly relevant for integrins because of the cellular capacity to regulate the state of integrin activation. The extent to which cells regulate integrin function is highly variable. In some cellular environments, the ability of a specific integrin to support cellular adhesion may be constitutive, whereas in others, the integrin may be inactive or only partially active (1). Thus, it has been concluded that integrins exist in a variety of activation states, although the basis for this conclusion has generally been indirect, based on whole-cell adhesion assays and the interaction of integrins with specific monoclonal antibodies. Accordingly, direct studies of the activation states of individual receptors are important. In addition, such studies can reveal the relative contribution to integrin regulation of changes in receptor conformation (affinity modulation) vs. receptor clustering (avidity; ref. 2).

The platelet integrin α IIb β 3 (GPIIb/IIIa), which is inactive on resting platelets but is activated by agonists such as ADP and thrombin, is the prototypic example of adhesion receptor modulation. This tight regulation of α IIb β 3 activity is imperative to prevent the spontaneous formation of thrombi. In this paper, we demonstrate a model employing laser tweezers to determine the force between single ligand-receptor bonds either in a purified protein system or on the surface of living, agonist-reactive cells. Laser tweezers are sensitive and accurate at the lower end of the force spectrum (0–150 pN; ref. 3); thus, they are a suitable system with which to study integrin–ligand interactions, because the binding forces involved have previously been reported to be in this range (4, 5). By using this methodology, we have demonstrated that there is only one α IIb β 3 activation state that is

relevant to the biological endpoint, platelet aggregation or adhesion, and that cellular stimulation increases the probability, but not the strength, of ligand binding.

Materials and Methods

Description of the Laser Tweezers-Based Model System to Study Receptor–Ligand Interactions. Laser tweezers are an optical system that use laser light to trap and manipulate dielectric particles such as small beads or cells (6–8). External forces applied to the trapped particle can be accurately measured because the angular deflection of the laser beam is directly proportional to the lateral force applied to the particle (9–11). To measure the bond strength between purified α IIb β 3 and fibrinogen, we covalently attached α IIb β 3 molecules to polyacrylamide-coated, spherical, silica pedestals anchored to a glass surface and covalently linked fibrinogen to smaller latex beads, one of which was trapped and moved toward and away from the receptor-coated pedestal, contacting it repeatedly (Fig. 1). The position of the optical trap was oscillated in a triangular waveform at 50 Hz with a peak-to-peak amplitude of 0.8 μ m. The separation of the pedestal and the bead then was reduced in 10-nm steps with the piezostage until repeated contacts were observed. The distance between the bottom of the trapped bead and the coverslip was controlled so that bead-surface (as opposed to bead-pedestal) touching and linkages were excluded. Laser beam deflection, sensed by a photodetector, was displayed as a voltage signal, which was converted to a force value by using a Stokes' law calibration (11) with a calibration coefficient (pN/V) determined for every experiment. The measurements were arranged to test bond strength at a high repetition rate, so that reasonable statistics could be obtained. Rupture forces after contact were collected and displayed as normalized force histograms for each experimental condition. The results of many experiments under similar conditions were averaged so that each histogram represented from 10^4 to $>2 \times 10^5$ contacts. The experimental protocol and subsequent analysis were validated with streptavidin and biotin because the interaction of these molecules has been studied by other methods. The rupture force distribution for streptavidin and biotin measured with laser tweezers ranged between 10 pN and >150 pN, which is consistent with the results of similar experiments using a biomembrane force probe (12); this result is in agreement with atomic force microscopy data that the rupture forces for this ligand-receptor pair may be beyond 150 pN (13, 14).

Preparing α IIb β 3- and Fibrinogen-Coated Surfaces. Both integrin and fibrinogen were covalently attached to surfaces, so that we would not have to worry about the possibility of pulling them off the surfaces with the laser tweezers. Integrin-coated pedestals were built by using plain silica microspheres 1.4 or 2.2 μ m in diameter. A working suspension of the microspheres (5% vol/

Abbreviation: TRAP, thrombin receptor-activating peptide.

§To whom reprint requests should be addressed at: Department of Cell and Developmental Biology, University of Pennsylvania School of Medicine, Room 1054, BRB II/III, 421 Curie Boulevard, Philadelphia, PA 19104-6058. E-mail: weisel@mail.cellbio.med.upenn.edu.

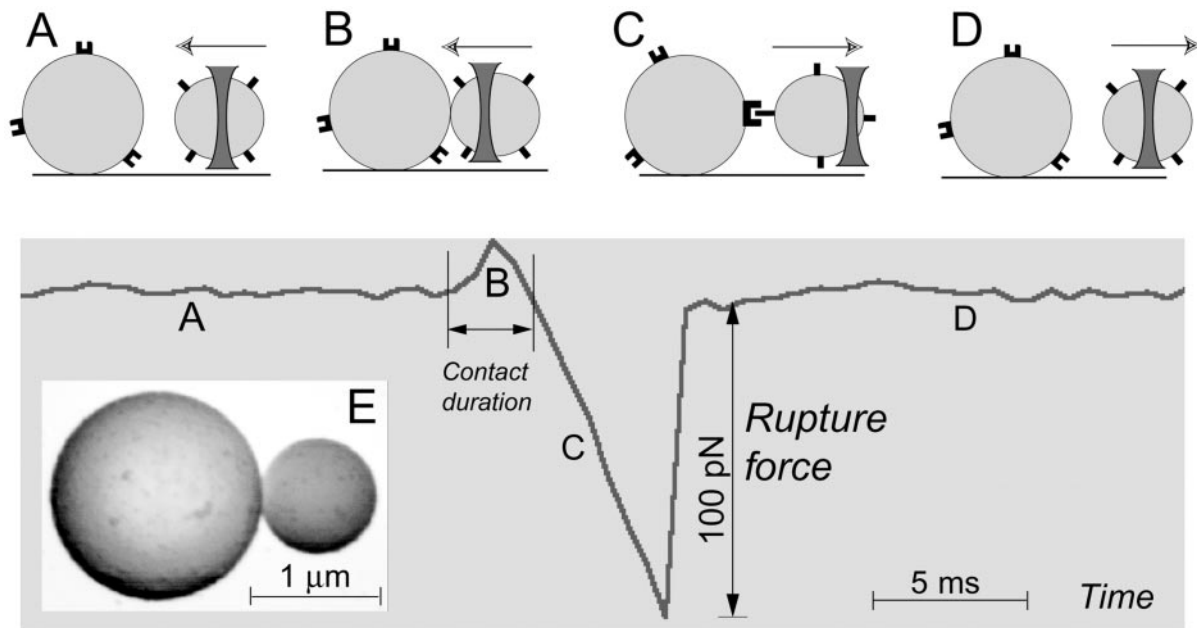


Fig. 1. Data trace for a typical interaction of fibrinogen and purified $\alpha\text{IIb}\beta_3$ as measured with laser tweezers. The force that the laser trap exerted on the latex bead can be partitioned into four parts (A–D). The latex bead was trapped near the center of the laser beam while moving toward (Upper A) or away (Upper D) from the silica pedestal. In the absence of contact between the bead and pedestal the measured force was small. (Upper B) At the moment of contact the pedestal stopped the motion of the latex bead while the laser beam continued to move in the same direction (left). The laser trap exerted a positive, compressive force on the pedestal and latex bead. The trap motion then reversed, and the compressive force declined to zero. Peak B (Lower) represents contact duration time between the surfaces. (Upper C) When the pedestal and latex bead bind, either specifically or nonspecifically, the bead position remained nearly constant as the laser trap continued to move to the right. The force on the bead increased in the negative direction almost linearly until the pedestal-bead bond was ruptured and the force rapidly returned to nearly zero. If no attachment occurred, there was no negative force. (Lower E) Scanning electron microscope image of the fibrinogen-coated latex bead (right) attached to the integrin-coated pedestal.

vol) was made in 30% (wt/vol) acrylamide/bis-acrylamide containing 2% (vol/vol) N,N,N',N' -tetramethylethylenediamine and 2% (vol/vol) amyl acetate. The microspheres were uniformly spread on a clean, dry microscope glass coverslip and air dried. Then, a freshly prepared saturated solution of ammonium persulfate ($2\ \mu\text{l}$) was smeared uniformly over the surface. The polyacrylamide-coated surface was activated by a 10% (vol/vol) solution of glutaraldehyde (4 h at room temperature), followed by washing with 20 volumes of 0.055 M borate (“binding”) buffer, pH 8.5. A solution of purified human $\alpha\text{IIb}\beta_3$ ($1\ \text{mg}\ \text{ml}^{-1}$), which had been dialyzed against the binding buffer containing 0.15 M NaCl and 1 mM CaCl_2 , was then applied for 12–16 h at 4°C . Human fibrinogen was covalently bound to $0.93\ \mu\text{m}$ carboxylate-modified latex beads with 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide as a crosslinking agent in a two-step procedure (based on TechNote no. 205, Bangs Laboratories, Carmel, IN) with minor modifications. The surface density of fibrinogen, determined by using ^{125}I -labeled protein, was near the saturation point of $(11 \pm 2) \times 10^{-9}\ \mu\text{g}/\mu\text{m}^2$; nonetheless, the fraction of reactive molecules with a conformation and orientation compatible with binding was indeterminate. Fibrinogen-coated beads were used at a final concentration of about $10^7\ \text{ml}^{-1}$.

Laser Tweezers Setup. The optical tweezers used for these experiments were assembled from a Nikon Diaphot 300 inverted microscope, $100 \times 1.3\ \text{N.A.}$ Fluor lens and a Spectra-Physics FCBar Nd:YAG laser. The tilt of the incoming laser beam at the back focal plane of the microscope objective, and consequently the trap position, was manually changed or computer controlled through a two-dimensional acousto-optical deflector (Brimrose, Baltimore, MD). The lateral forces that the trap exerted on a bead were measured with a quadrant detector conjugate to the

back focal plane of the condenser (9) and were calibrated from the low-frequency component of the Brownian motion. Manipulations were visualized by a video charge-coupled device camera. All experiments were conducted at a trap stiffness of $0.2\ \text{pN/nm}$ as computed from the bandwidth of the Brownian motion. Both the force calibration and trap stiffness were routinely confirmed by the Stokes’ force method (11). LABVIEW software (National Instruments, Austin, TX) was used to control and record laser beam deflection, to move the piezoelectric stage (Queensgate, Berkshire, U.K.), and to analyze data subsequently off-line.

Results

Observed signals using purified proteins could be experimentally partitioned into three categories: ligand-receptor binding and unbinding, nonspecific interactions, and small optical artifacts. The force distribution histogram of the interactions between intact purified $\alpha\text{IIb}\beta_3$ and fibrinogen (Fig. 2*Ab*) revealed a major peak at 60 to 150 pN with a maximum bin at 90–100 pN, the latter defined as the yield strength, i.e., the average force required to rupture an $\alpha\text{IIb}\beta_3$ -fibrinogen bond. The fraction of contacts with rupture forces $>60\ \text{pN}$, assumed to be the fraction of specific integrin–fibrinogen interactions, was as large as 3% for freshly purified intact $\alpha\text{IIb}\beta_3$. In control experiments, $\alpha\text{IIb}\beta_3$ was pretreated with 1 mM EDTA to prevent its specific association with fibrinogen (15). The force distribution histogram under this condition revealed no rupture forces $>60\ \text{pN}$ (Fig. 2*Aa*), indicating that high-affinity interactions did not occur.

Interactions between control beads and pedestals not coated with receptor-ligand pairs resulted in lower rupture forces that depended on the specific surface treatment. Optical artifacts observed with or without trapped latex beads produced signals that appeared as forces below 10 pN. The weakest interactions

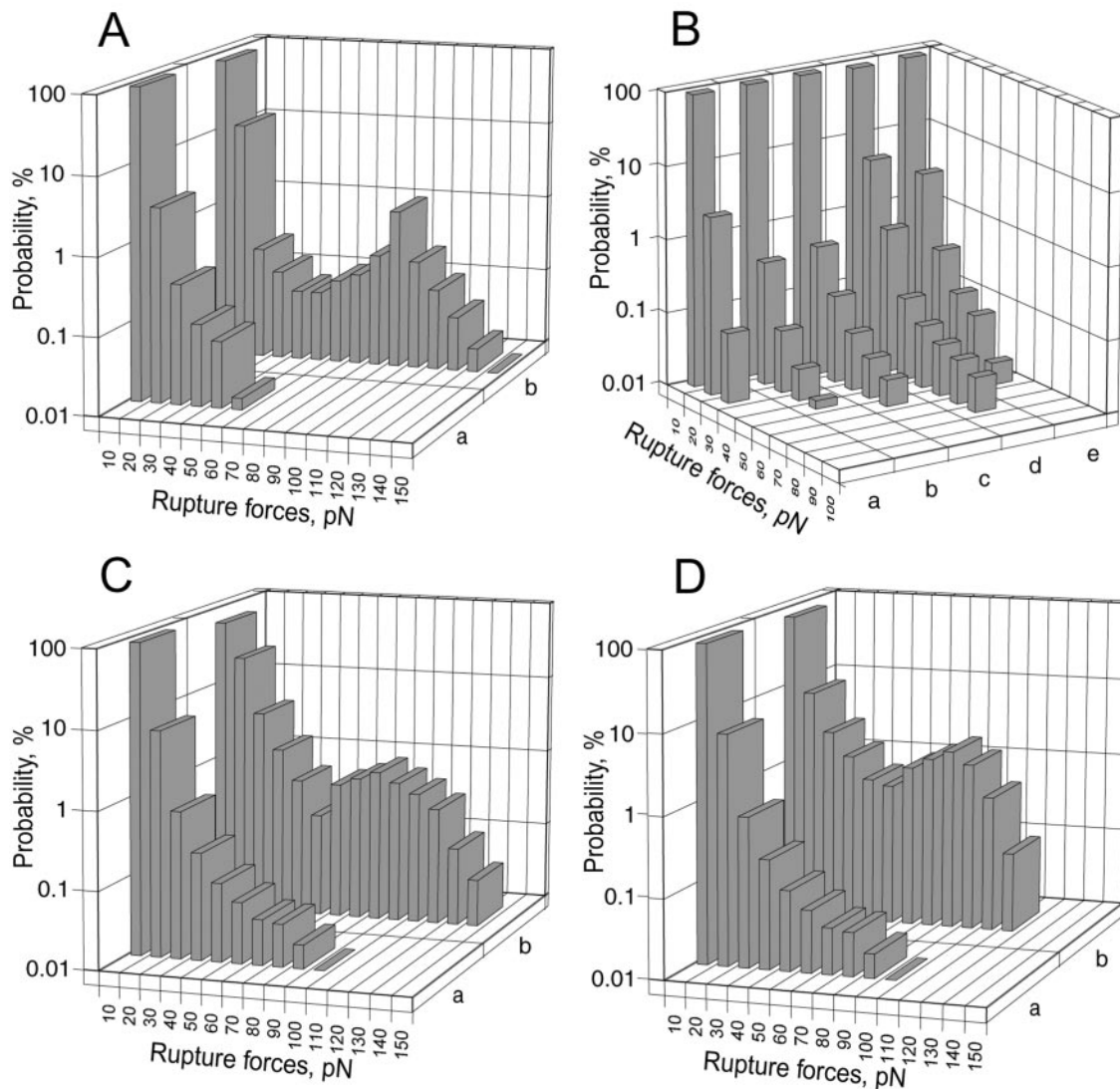


Fig. 2. Comparative force histograms of various surface-surface interactions measured by laser tweezers. The individual forces measured during each contact-detachment cycle under similar experimental conditions were collected into 10-pN wide bins. The number of events in each bin was plotted against the average force for that bin after normalizing by the total number of interaction cycles. The percentage of events in a particular force range (bin) represented the probability of rupture events at that tension. (A) Comparison of the force distributions of the interactions between fibrinogen-coated beads and EDTA-pretreated α I**IIb** β 3 molecules (a) or native immobilized α I**IIb** β 3 (b). (B) Force distribution histograms of the nonspecific, control surface-surface interactions. (a) BSA-coated beads with untreated polyacrylamide-coated pedestals. (b) BSA-coated beads with BSA-coated pedestals. (c) Fibrinogen-coated beads with BSA-coated pedestals. (d) BSA-coated beads with α I**IIb** β 3-coated pedestals. (e) Fibrinogen-coated beads with EDTA-pretreated α I**IIb** β 3 bound to pedestals. (C) Comparison of the force distributions of the interactions between fibrinogen-coated beads and α I**IIb** β 3 on the surface of resting platelets (a) or platelets stimulated with 10 μ M ADP (b). (D) Comparison of the force distributions of the interactions between fibrinogen-coated beads and α I**IIb** β 3 on the surface of resting platelets (a) or platelets stimulated with 50 μ M TRAP (b).

with rupture forces below 30 pN were observed for untreated polyacrylamide-coated pedestals in contact with BSA-coated beads (Fig. 2Ba). Pedestal surfaces were slightly more reactive after covalent coupling with BSA (Fig. 2Bb) and more reactive when coupled with fibrinogen (Fig. 2Bc) or α I**IIb** β 3 (Fig. 2Bd). For comparison, beads coated with α I**IIb** β 3 in the presence of EDTA again led to low-rupture forces (Fig. 2Be). The overall probability of weak nonspecific forces was higher than for specific interactions, such that rupture forces for specific interactions between α I**IIb** β 3 and fibrinogen were consistently >60 pN.

This conclusion was confirmed independently by probing the specificity of the forces with well characterized inhibitors of

the interaction between fibrinogen and α I**IIb** β 3 (16). The appearance of specific rupture forces >60 pN was prevented (Fig. 3A) by the presence of tirofiban, a tyrosine-based molecule that mimics the three-dimensional structure of the integrin inhibitor RGD (17), abciximab, a murine-human chimeric Fab fragment of the monoclonal antibody 7E3 that inhibits platelet aggregation (18), the H12 peptide from the carboxyl terminus of the fibrinogen γ -chain (19), or the α I**IIb** β 3-specific monoclonal antibody A2A9 (20).

Fibrinogen binding to α I**IIb** β 3 on the platelet surface is absolutely regulated by platelet agonists (1). Therefore, we next studied interactions between a fibrinogen-coated bead and α I**IIb** β 3 on the surface of living platelets. A platelet was trapped from a suspension of gel-filtered cells and attached to a 5- μ m diameter silica pedestal coated with polylysine (Fig. 4). A

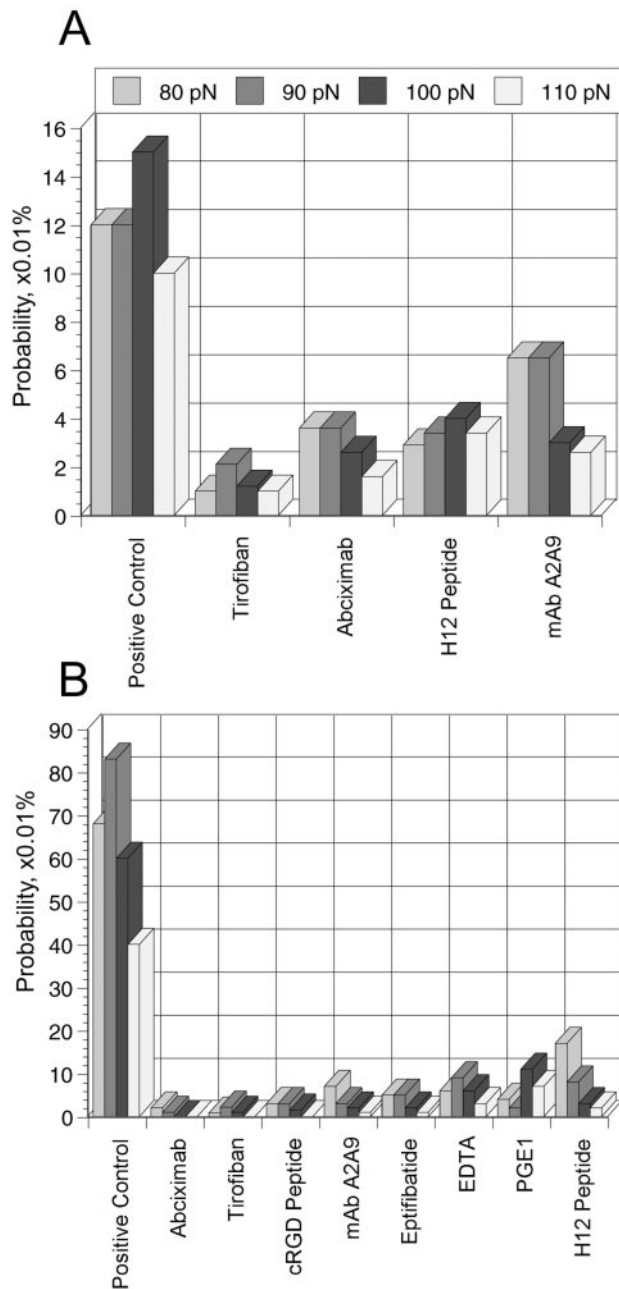


Fig. 3. Effect of specific α I**IIb** β 3 antagonists on the rupture forces between fibrinogen and α I**IIb** β 3. (A) Interactions between purified α I**IIb** β 3 and fibrinogen in the presence or absence (positive control) of inhibitors: tirofiban (40 μ M), abciximab (100 μ g/ml), H12–fibrinogen-binding inhibitor dodecapeptide (1 mM), abciximab (100 μ g/ml), H12–fibrinogen-binding inhibitor dodecapeptide (1 mM), and mAb A2A9– α I**IIb** β 3-binding specific “antiaggregant” monoclonal antibody (100 μ g/ml). (B) Force distribution of the interactions between fibrinogen and platelets stimulated by 10 μ M ADP in the presence or absence of the following inhibitors: abciximab (20 μ g/ml), tirofiban (40 μ M), cRGD–cyclic RGD peptide (80 μ M), eptifibatide (100 μ g/ml), EDTA–Na₂–ethylendiaminetetraacetic acid (1 mM), PGE1–prostaglandin E1 (2 μ M), and H12–fibrinogen-binding inhibitor dodecapeptide (1 mM). Because positive control data were collected separately for each experimental series with inhibitors, the probabilities displayed here were different from the data collected from the entire sum total of experimental data shown in Fig. 2.

fibrinogen-coated latex bead was then trapped and brought into intermittent contact with the platelet in the same way as for the experiments with purified proteins.

Force distribution histograms for the interactions between quiescent platelets and surface-bound fibrinogen were similar to

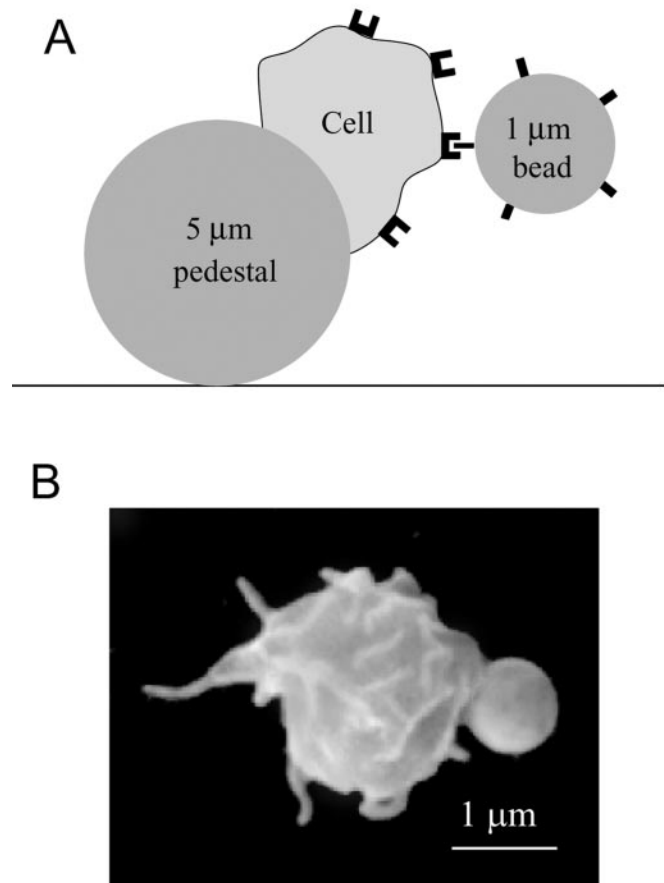


Fig. 4. Schematic representation (A) and scanning electron microscope image (B) of the interaction between an immobilized platelet and a fibrinogen-coated bead.

those for the control histograms obtained between fibrinogen and BSA or BSA and purified α I**IIb** β 3, except for rare events for which the magnitude of the recorded rupture forces reached a maximum of 100 pN (Fig. 2*C*_a).

On the other hand, platelet stimulation with ADP substantially affected their interaction with the fibrinogen-coupled beads, producing a major peak of rupture forces at 60 to 140 pN with a yield strength of 80–90 pN, without substantial change in lower nonspecific forces (Fig. 2*C*_b). Moreover, there was an ADP concentration-dependent increase in the probability of specific interactions having rupture forces >60 pN, with the greatest fraction observed after platelet stimulation with 10 μ M ADP (Fig. 5*A*). At the same time, the average yield strength was constant at all ADP concentrations, with a value of about 80 pN (Fig. 5*A*). To confirm that the ADP-stimulated interactions were specific for α I**IIb** β 3 and fibrinogen, two types of platelet inhibitors were used: competitive inhibitors of fibrinogen binding described above [abciximab, tirofiban, eptifibatide, A2A9, cyclic RGD-peptide (cRGD), and H12] and inhibitors of platelet metabolism (EDTA and PGE1). As shown in Fig. 3*B*, each of the inhibitors substantially decreased the frequency of attachments between platelets stimulated by 10 μ M ADP and fibrinogen-coated beads, with more than a 10-fold reduction in the probability of high-yield force interactions.

ADP is a “weak” platelet agonist. To determine whether stimulating platelets with a “strong” agonist results in similar fibrinogen-binding characteristics, we used the thrombin receptor-activating peptide (TRAP) as the platelet agonist. Like ADP, TRAP induced fibrinogen binding to α I**IIb** β 3 with an

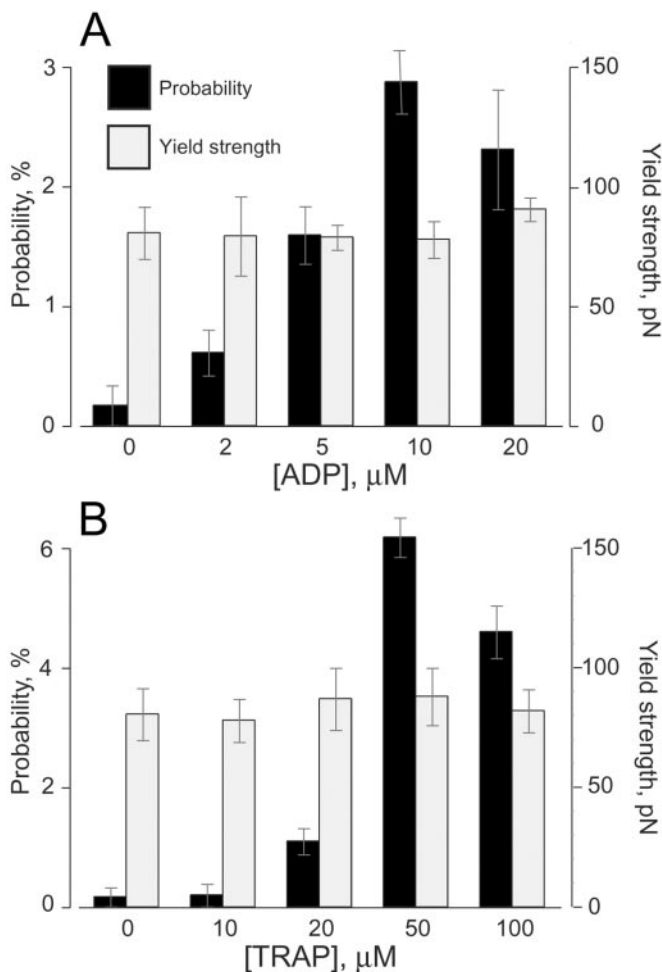


Fig. 5. Probability and average yield strength of specific (>60 pN) fibrinogen- α IIb β 3 interactions plotted against the concentration of either ADP (A) or TRAP (B).

average yield strength of 80–90 pN (Fig. 2*D**b*). Moreover, increasing the TRAP concentration increased the probability, but not the average yield strength, of fibrinogen binding (Fig. 5*B*), implying that affinity of α IIb β 3 for ligands is independent of the nature of the platelet stimulus. Nevertheless, TRAP, at a concentration of 50 μ M, induced significantly higher maximal probability of platelet interaction with fibrinogen consistent with its ability to induce a redistribution of α IIb β 3 molecules from an intracellular pool to the platelet membrane (21–23).

Discussion

By using laser tweezers, we found that α IIb β 3 resides on the platelet surface in either of two activation states, depending on the state of platelet stimulation. Nonetheless, because platelets express 80,000 or more copies of α IIb β 3 on their surface (24), it is possible that the rupture forces we measured represented the breaking of multiple rather than individual α IIb β 3-fibrinogen bonds. Two arguments support the view that our measurements represent individual bimolecular ligand-receptor binding events. First, the histograms of the distribution of rupture forces of multiple interactions should appear as a series of quantized peaks that are multiples of a single value of force and have probabilities inversely proportional to the number of bonds (25). However, apart from nonspecific interactions, we observed only a single well defined peak in the force histograms, suggesting that this peak represents individual ligand-receptor bonds. In quan-

titative terms, if we were to assume that the 100 pN peak we observed with a 3% probability represents the interactions of two ligand-receptor pairs, then the presumed bimolecular interactions at 50 pN would have a hypothetical square root probability of about 17%, whereas the observed probability was about 0.1% (Fig. 2*A**b*), indicating that the peak does indeed represent individual ligand-receptor interactions. Second, if multiple bonds were formed during a contact, it is unlikely that all would be ruptured simultaneously; rather, it is more probable that they would be ruptured sequentially so that multiple steps should have been observed during bond breakage. However, the rupture events we observed all occurred as a single step, at least with the time resolution of 0.5 ms.

Some of the events we measured may have resulted from pulling receptors out of the cell membrane, protein uprooting, or from disrupting integrin-receptor binding to the cytoskeleton (26), but such occurrences are unlikely for the following reasons. If receptor molecules are not linked to the cytoskeleton, tethers form because of reservoir buffering membrane tension rather than protein uprooting (27). Consistent with this report, we frequently observed tethers in resting or exhausted platelets after pulling the attached fibrinogen-coated beads apart from platelets. In activated platelets, although we occasionally observed tethers, the frequency was substantially lower compared with nonactivated cells. In addition, for most individual bead-platelet pairs, we observed multiple repeated binding and rupture signals, suggesting that the plasma membrane of the platelet was not drastically modified by either protein uprooting or cytoskeletal disruption.

Direct quantification of bimolecular interactions between fibrinogen and α IIb β 3 has not been reported previously, but two papers are relevant to our data. Goldsmith *et al.* (28) used external hydrodynamic force to break pairs of α IIb β 3-coated latex beads formed in the presence of free fibrinogen. They found that a substantial fraction of the fibrinogen-bridged beads separated at forces in the range of 70–150 pN, similar to the rupture forces we measured. However, a direct comparison between our work and theirs should be avoided because of substantial differences in the experimental models. Lee and Marchant used atomic force microscopy to measure the force required to rupture bonds between an immobilized RGD-containing peptide (GSSSGRGDSPA) and α IIb β 3 on the surface of adherent platelets (29). They reported that a value of 90 ± 45 pN represented the single molecular interaction between the RGD-ligand and the integrin at a loading rate of 18,000 pN/s, a value essentially identical to that used in our experiments (20,000 pN/s). The results reported in both of these papers are consistent with our data and support the view that the forces we measured are characteristic of bimolecular ligand-integrin interactions. Although not comparable in quantitative terms, our data are also in good qualitative agreement with recent work (30) in which the adhesion of ADP-activated vs. resting platelets to fibrinogen-coated latex beads was studied at different shear rates.

Our results have a number of physiological implications. First, the portion of contacts that lead to specific interactions between fibrinogen and α IIb β 3 is directly related to but not identical to the fraction of α IIb β 3 molecules with accessible exposed ligand-binding sites. On the other hand, the yield strength or, in terms of cell biology, the adhesion strength of activated receptor is an intrinsic property of the activated conformation of α IIb β 3 and is manifest as a distinctive range of forces with a particular set of loading rates (31). Therefore, the laser tweezers experiments allow a clear discrimination between accessibility and affinity of receptor-ligand binding. The probability of a specific interaction between fibrinogen and inactivated α IIb β 3 is low because the conformation of α IIb β 3 does not permit fibrinogen binding. Platelet stimulation by agonists increases the number of α IIb β 3

molecules with accessible fibrinogen-binding sites, but has no effect on the bond strength between fibrinogen and the activated form of the integrin. Thus, our results demonstrate that α IIB β 3 activation is an all-or-none phenomenon; each α IIB β 3 molecule resides on the platelet in either a completely on or a completely off conformation, which is consistent with structural data (32–34).

Second, in experiments with purified α IIB β 3, the fraction of specific interactions is related to the percentage of α IIB β 3 complexes in an active conformation and with the proper orientation on the pedestal surface. Because many experiments in the literature were performed with purified α IIB β 3, it is important to know whether the results are relevant to those obtained under physiologic conditions in live cells. The force histograms of fibrinogen binding to purified α IIB β 3 and to α IIB β 3 on the platelet surface were similar with regard to both the fraction of specific interactions and the average adhesion strength. It is important to mention that the force loading rate used in this work (20,000 pN/s) has the same order of magnitude that is imposed on platelets under physiologic shear rates (several hundred s⁻¹), which is still low enough for α IIB β 3 to interact with surface-bound fibrinogen (35).

Third, the results from these experiments lead to an important addition to the cell adhesion paradigm. The remarkable number of weak interactions produced by quiescent platelets suggests that α IIB β 3-mediated platelet adhesion may be the result of the combination of numerous low-affinity attractive forces of non-specific origin, together with forces arising from specific bond formation between ligand and receptor. The nonspecific protein-protein interactions may bring a ligand toward α IIB β 3 molecule and provide the spatial congruence required to initiate and/or reinforce specific bonding. Moreover, they may account for the ability of resting platelets to adhere spontaneously to fibrinogen-coated surfaces (1).

Our results indicate that at least for the integrin α IIB β 3, cells regulate the number of accessible binding sites, not the affinity of ligand binding. Because we have found that it is possible to measure and compare forces between individual ligand-receptor pairs on living cells, it will now be possible to determine whether this observation applies to the other members of the integrin superfamily.

We thank Yasuharu Takagi, Gaston Vilaire, and Chandrasekaran Nagaswami for their valuable methodological assistance. This work was supported by National Institutes of Health Grants HLBI 57407 and HLBI 30954.

- Bennett, J. S. (1996) *Trends Cardiovasc. Med.* **16**, 31–36.
- Hato, T., Pampori, N. & Shattil, S. J. (1998) *J. Cell Biol.* **141**, 1685–1695.
- Visscher, K. & Block, S. M. (1998) *Methods Enzymol.* **298**, 460–489.
- Lehenkari, P. P. & Horton, M. A. (1999) *Biochim. Biophys. Res. Commun.* **259**, 645–650.
- Stout, A. L. (2001) *Biophys. J.* **80**, 2976–2986.
- Ashkin, A. (1992) *Biophys. J.* **61**, 569–582.
- Ashkin, A. (1997) *Proc. Natl. Acad. Sci. USA* **94**, 4852–4860.
- Svoboda, K. & Block, S. M. (1994) *Annu. Rev. Biomol. Struct.* **23**, 247–285.
- Smith, S. B., Cui, Y. & Bustamante, C. (1996) *Science* **271**, 795–799.
- Allersma, M. W., Gittes, F., deCastro, M. J., Stewart, R. J. & Schmidt, C. F. (1998) *Biophys. J.* **74**, 1074–1085.
- Visscher, K., Gross, S. P. & Block, S. M. (1996) *IEEE J. Select. Topics Quant. Elec.* **2**, 1066–1076.
- Merkel, R., Nassoy, P., Leung, A., Ritchie, K. & Evans, E. (1999) *Nature (London)* **397**, 50–53.
- Florin, E.-L., Moy, V. T. & Gaub, H. E. (1994) *Science* **264**, 415–417.
- Moy, V. T., Florin, E.-L. & Gaub, H. E. (1994) *Science* **266**, 257–259.
- Brass, L. F., Shattil, S. J., Kunicki, T. J. & Bennett, J. S. (1985) *J. Biol. Chem.* **260**, 7875–7881.
- Bennett, J. S. (2001) *Annu. Rev. Med.* **52**, 161–184.
- Cook, J. J., Bednar, B., Lynch, J. L., Gould, R. J., Egberon, M. S., Halczenko, W., Duggan, M. E., Hartman, G. D., Lo, M.-W., Murphy, G. M., et al. (1999) *Cardiovasc. Drug Rev.* **17**, 199–224.
- Coller, B. S. (1985) *J. Clin. Invest.* **76**, 101–108.
- Kloczkiak, M., Timmons, S., Lukas, T. J. & Hawiger, J. (1984) *Biochemistry* **23**, 1767–1774.
- Bennett, J. S., Hoxie, J. A., Leitman, S. F., Vilaire, G. & Cines, D. B. (1983) *Proc. Natl. Acad. Sci. USA* **80**, 2417–2421.
- Woods, V. L., Jr., Wolf, L. E. & Keller, D. M. (1986) *J. Biol. Chem.* **261**, 15242–15251.
- Niiya, K., Hodson, E., Bader, R., Byers-Ward, V., Koziol, J. A., Plow, E. P. & Ruggeri, Z. M. (1987) *Blood* **70**, 475–483.
- Suzuki, H., Kaneko, T., Sakamoto, T., Nakagawa, M., Miyamoto, T., Yamada, M. & Tanoue, K. (1994) *J. Electron Microsc. J.* **43**, 282–289.
- Wagner, C. L., Mascelli, M. A., Neblock, D. S., Weisman, H. F., Coller, B. S. & Jordan, R. E. (1996) *Blood* **88**, 907–914.
- Pierres, A., Benoleil, A.-M. & Bongrand, P. (1998) *Cell Adhes. Commun.* **5**, 375–395.
- Shao, J. Y. & Hochmuth, R. M. (1999) *Biophys. J.* **77**, 587–596.
- Raucher, D. & Sheetz, M. P. (1999) *Biophys. J.* **77**, 1992–2002.
- Goldsmith, H. L., McIntosh, F. A., Shahin, J. & Frojmovic, M. M. (2000) *Biophys. J.* **78**, 1195–1206.
- Lee, I. & Marchant, R. E. (2000) *Microsc. Microanal.* **2**, Suppl. 6, 974–975.
- Bonnefoy, A., Liu, Q., Legrand, C. & Frojmovic, M. M. (2000) *Biophys. J.* **78**, 2834–2843.
- Evans, E. (2001) *Annu. Rev. Biophys. Biomol. Struct.* **30**, 105–128.
- Hantgan, R. R., Paumi, C., Rocco, M. & Weisel, J. W. (1999) *Biochemistry* **38**, 14461–14474.
- Hantgan, R. R., Rocco, M., Nagaswami, C. & Weisel, J. W. (2001) *Protein Sci.* **10**, 1614–1626.
- Vinogradova, O., Haas, T., Plow, E. F. & Qin, J. (2000) *Proc. Natl. Acad. Sci. USA* **97**, 1450–1455.
- Savage, B., Saldivar, E. & Ruggeri, Z. M. (1996) *Cell* **84**, 289–297.