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Proteins, Platelets, and Blood Coagulation at Biomaterial Interfaces

Li-Chong Xu¹, James Bauer, Christopher A. Siedlecki^{1,2}, and A Contribution from the Hematology at Biomaterial Interfaces Research Group

¹Department of Surgery, Biomedical Engineering Institute, The Pennsylvania State University, College of Medicine, Hershey, PA, 17033

²Department of Bioengineering, Biomedical Engineering Institute, The Pennsylvania State University, College of Medicine, Hershey, PA, 17033

Abstract

Blood coagulation and platelet adhesion remain major impediments to the use of biomaterials in implantable medical devices. There is still significant controversy and question in the field regarding the role that surfaces play in this process. This manuscript addresses this topic area and reports on state of the art in the field. Particular emphasis is placed on the subject of surface engineering and surface measurements that allow for control and observation of surface-mediated biological responses in blood and test solutions. Appropriate use of surface texturing and chemical patterning methodologies allow for reduction of both blood coagulation and platelet adhesion, and new methods of surface interrogation at high resolution allow for measurement of the relevant biological factors.

1. Introduction

Biomaterials have been routinely used in all fields of medicine and surgery for a range of medical applications from short term dressing to long term implants, and play a critical role in the treatment of disease and the improvement of health care^{1,2}. Despite the wide use of biomaterials in clinical environments, the biocompatibility of the materials is still far from ideal and a variety of adverse reactions such as inflammation, fibrosis, infection, and thrombosis, may be triggered^{3–5}. The biocompatibility of implants and successful application of these medical devices are largely dependent on the biological events occurring at the surfaces. Probing these specific and direct interactions between biomaterials and tissue components will markedly improve the field's understanding of the mechanisms of biocompatibility and thereby to improve the development of new biomaterials⁶.

* Author to whom correspondence should be addressed: Christopher A. Siedlecki, Ph.D., Departments of Surgery and Bioengineering, Pennsylvania State University College of Medicine, Biomedical Engineering Institute, Mail Code H151, 500 University Drive, Hershey, PA 17033, Phone: (717) 531-5716, Fax: (717) 531-4464, csiedlecki@psu.edu.

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Blood is often the first body fluid that comes into contact with blood-contacting devices. Blood-material interactions trigger a complex series of events including protein adsorption, platelet adhesion and activation, coagulation, and thrombosis. Rapid adsorption of plasma proteins is the first event occurring on biomaterial surface during blood/material interactions, and leads to activated, adsorbed proteins that can catalyze, mediate, or moderate the subsequent biological responses to biomaterials⁷⁻⁹. Surface-induced thrombosis is the main problem impeding development of long-term blood contacting devices^{10,11}. Thrombus formation on device surfaces is partially due to platelet-mediated reactions¹² and partially due to coagulation of blood plasma¹³.

Biological responses are a complex process governed by many factors, but it is widely accepted that surface properties of a biomaterial dictate the biological response. Surface properties such as chemistry, topography, surface free energy, elasticity, and charge may moderate protein and cell interactions, and ultimately the host response. For example, platelet adhesion and activation on biomaterial surfaces is influenced by surface properties such as energy¹⁴, charge¹⁵, and composition¹⁶. However, platelet adhesion and activation are largely mediated by proteins such as fibrinogen^{17,18}. As plasma proteins can rapidly adsorb onto the material surface to form a “conditioning film” following blood contact, this adsorbed protein layer may minimize the direct effect of biomaterial surface properties on cell responses, and the surface-biology interactions are mediated by proteins (Figure 1).

The aim of this work is to outline the current understanding of the phenomena of thrombosis with particular emphasis placed on cardiovascular biomaterials. As surface chemistry and topography of a biomaterial are known to be the important parameters that influence the biological responses to surfaces, this manuscript focuses on the protein adsorption, platelet adhesion, and blood coagulation at biomaterial interfaces with different surface properties.

2. Protein adsorption and platelet adhesion on biomaterial surfaces

2.1 Plasma proteins and platelet adhesion

When a biomaterial comes into contact with blood, a layer of plasma proteins typically adsorbs to the surface and mediate biological responses to biomaterials. Blood contains many hundreds of proteins with a wide range of biological functions and activity, and present in vastly different concentrations. Albumin, immunoglobulins and fibrinogen are the most abundant proteins in plasma and represent more than 50% of all plasma proteins. In considering surface-induced thrombosis, albumin is generally considered to be inert towards platelet adhesion and activation¹⁹, while fibrinogen is a central protein in the process of biomaterial-induced thrombosis²⁰⁻²². Other plasma proteins such as fibronectin^{23,24}, vitronectin^{25,26}, and von Willebrand factor (vWF)^{27,28}, have been shown to be capable of mediating platelet adhesion to materials when pre-adsorbed to the surface. These adsorbed proteins can bind to platelets via cell membrane receptors such as GPIIb/IIIa (known as integrin $\alpha_{IIb}\beta_3$), GPIb, and other receptors, and such binding will induce platelet adhesion and aggregation. Adherent platelets become activated and mediate clotting events such as platelet aggregation and formation of thrombus.

The ability of proteins to promote platelet adhesion varies with substrates and flow condition. Fibrinogen has been identified as the major protein mediating platelet adhesion biomaterials at low shear stress²⁹ and plays a prominent role in development of surface-induced thrombosis due to its multi-functional role serving as a ligand for platelet adhesion, linking platelet aggregates together, and stabilizing thrombi as fibrin polymer^{20,30,31}. vWF circulates in plasma as a series of heterogeneous multimers, mediating platelet tethering, translocation and finally adhesion to areas of injured endothelium under higher shear stress conditions. Immobilization and shear induce conformational changes in vWF, and enhance the binding of vWF to glycoprotein Ib alpha on platelet cell membrane, resulting in platelet adhesion^{32,33}. Fibronectin plays little or no role in mediating platelet adhesion on material surfaces pre-adsorbed with normal plasma or fibrinogen, but it does support platelet adhesion in fibrinogen-depleted plasma³⁴. Vitronectin is a multifunctional 75-kD glycoprotein that is present in plasma and the extracellular matrix, and functions as a complement regulatory promoting growth and attachment of cells. Vitronectin can bind to platelets via membrane protein GPIIb/IIIa, and plays a role in platelet adhesion and formation of stable platelet aggregates^{35,36}.

2.2 Surface properties influence protein adsorption

A number of different factors have been shown to influence the composition and function of the adsorbed proteins on implanted biomaterials³⁷. Among these are the solution concentration of the proteins, the properties of the surface and the intrinsic “surface activity” of the protein^{38–40}. Other studies show time dependent factors involved in protein adsorption and function on biomaterials^{41–44}. A fundamental premise in the field of biomaterials is that the surface properties of a biomaterial affect the adsorption, the structure and the function of proteins^{45,46}. Surface properties that have been shown to be important (1) surface free energy and wettability, (2) surface chemistry and functional group, and (3) surface topography and roughness^{37,47,48}.

2.2.1 Surface energy and wettability—The introduction of a biomaterial surface in blood creates a new interface between cellular and fluid components of blood and the material. The thermodynamic quantity of surface free energy of both solid and liquid phases leads to the interfacial free energy at the interface, which can be used to assess the biocompatibility of biomaterials⁴⁹. In the case of blood plasma-biomaterial interface, it has been suggested that the interfacial free energy should be in the order of 1–3 dyne/cm for long-term compatibility with blood⁵⁰. Protein adsorption is an energy-driven process that involves the properties of both the substrate surface and the proteins themselves. Calculation of the interfacial energy of protein interaction with surface from the surface free energy of a biomaterial is helpful to quantitatively understand the mechanisms of protein adsorption on varied surfaces^{51–53}. It has been reported that there is a correlation between the amount of protein adsorbed on the polymeric materials such as poly(methylmethacrylate) (PMMA), polystyrene (PS), and poly(dimethylsiloxane) (PDMS) and the total surface energy of the substrate, i.e., adsorption is higher on low energy substrates⁵⁴.

Surface wettability driven by surface energy is a key factor in determining adsorption kinetics and the amount of proteins adsorbed on the surface^{55,56}. Wettability can be

characterized by the Dupré work of water adhesion to surfaces, $W = \gamma_{LV}(1 + \cos\theta)$, where the interfacial energy of water $\gamma_{LV} = 71.97 \text{ mJ/m}^2$ at 25°C ^{57,58}. Generally water contact angle θ is measured and represented by the water adhesion tension τ , ($\tau = \gamma_{LV} \cos\theta$), to estimate the surface wettability (hydrophilic or hydrophobic). Vogler et al. defined the terms “hydrophilic” and “hydrophobic” from experimental observation, with hydrophilic surfaces exhibiting advancing water contact angle $< 65^\circ$, and hydrophobic surfaces exhibiting water contact angle $> 65^\circ$ ^{59,60}. A more precise definition of the relative terms hydrophobic and hydrophilic for use in biomaterials was further derived from calculation of wetting energetics, where hydrophilic surface wetting expends $> 1.3 \text{ kJ/mole-of-surface-sites}$, and hydrophobic surface wetting expends $< 1.3 \text{ kJ/mole-of-surface-sites}$. Considering the hydrogen bond of surrounding water involved in surface wetting, hydrophilic surfaces wet with > 1 hydrogen bond per water molecule whereas hydrophobic surfaces wet with < 1 hydrogen bond per water molecule ⁶¹.

In general, hydrophobic surfaces tend to adsorb larger amount of proteins than hydrophilic⁶² as the hydrophilic surface is strongly bounded with water molecules that are difficult to be displaced by adsorbing proteins, while protein readily displaces water from the hydrophobic surface to become adsorbed⁵¹. The above experimental and theoretical observations of surface hydrophilic/hydrophobic transiting at $\theta = 65^\circ$ may also explain the sharp changes in the protein adsorption to material surfaces with water contact angles centered around 65° in diverse circumstances^{53,63–65}. *Cha et al.* measured the difference in the amount of protein adsorbed at solid-vapor and solid-liquid interfaces and found it decreased with the adhesion tension τ from maximal values measured on the methyl-terminated self-assembled monolayers surface through zero (no protein adsorption in excess of bulk solution concentration) near $\tau = 30 \text{ mN/m}$ ($\theta = 65^\circ$)⁵³. Other independent experiments show that the protein adsorption capacity decreases monotonically with increasing adsorbent hydrophilicity to a limit of detection near $\tau = 30 \text{ mN/m}$ ($\theta = 65^\circ$) for all protein/surface combinations studied ^{64,65}. All these results suggest that the structured interfacial water (with thickness $\sim 2\text{--}3 \text{ nm}$) plays an important role in protein adsorption⁶⁶.

Protein adsorption on surfaces can be understood from the aspect of interfacial energetics and measurement of adsorbed amount, but also from the direct measurement of the strength of the molecular interaction between protein and the material surface. The force measurement techniques of an atomic force microscopy (AFM) using a protein-modified probe tip have been powerful tools to analyze the actual interaction forces between protein and material surfaces⁶⁷. Since the normal AFM tip is very sharp with radii of end in the range of $10\text{--}20 \text{ nm}$, force measurement using a protein-modified model probe is generally regarded as individual proteins interacting with surface. Such characteristics of individual proteins in interaction with surface offer much insight into the environment presented by biomaterial interfaces.

Table 1 shows the adhesion forces between model self-assemble monolayer (SAM) surfaces and plasma proteins albumin, coagulation factor XII, and coagulation factor XIIa. (FXII and FXIIa are involved in contact activation of the coagulation cascade). All proteins exhibit strong adhesion to the hydrophobic methyl-terminated octadecyltrichlorosilane (OTS) surface and weak adhesion to water-wettable glass surfaces, despite FXII supposedly being uniquely

attracted to glass surfaces. The force measurements were extended to look at a range of water wettabilities in a polymer by treating low density polyethylene (LDPE) in a glow-discharge plasma⁶⁸. The contact angle and adhesion forces between albumin, FXII, and fibrinogen modified tips with the substrate were measured. All of the proteins studied exhibited a step increase in adhesion force as the contact angles of the surface increased above $\theta \sim 60^\circ$ ($\tau < 36.4$ dyn/cm). The step occurred within a narrow range of wettabilities over $\sim 60\text{--}65^\circ$. Figure 2 illustrates the trend of adhesion force of fibrinogen interaction with LDPE surface. ANOVA shows that adhesive forces between the poorly wettable surfaces are not different from each other, adhesive forces between the wettable surfaces are not different from each other, but there are substantial differences between the two groups, as illustrated in Table 2. The presence of the step increase in adhesion forces was also observed in interactions of proteins and a series of SAM-surfaces. For SAMs with $\theta < 66^\circ$ (-OH, -CONH₂, and -EG₃OH), weak adhesion was observed, while for $66^\circ < \theta < 104^\circ$ (-CH₃, -OPh, -CF₃, -CN, -OCH₃), strong adhesion was observed and increased with the molecular weight of the proteins⁶⁹. Results suggested that the effect of surface wettability on protein adhesion is actually quite straightforward and that subtle changes in wettability not affect protein adhesion unless that change yields a transition across this $\theta=60\text{--}65^\circ$ region⁶⁸.

2.2.2 Surface chemical structure and functional group—Surface chemistry (elemental constituents and functional group) gives rise to surface properties such as wettability and charge, and influences the extent of protein adsorption, denaturation, and functional activity. It plays an important role in biocompatibility of implants⁸. To reduce unwanted-protein adsorption on surfaces, intensive research efforts has gone towards engineering surfaces with functional groups to repel proteins. Ostuni *et al.* described that typical characteristics of protein-resistant surface should be hydrophilic, neutral charge, containing hydrogen-bond acceptors, but without hydrogen-bond donors⁷⁰. Although there appears to be exceptions, it is believed that the identification of protein resistant chemical structure and/or functional group of surface can help to control protein adsorption and biological events following adsorption, and thus enhance the biocompatibility of biomaterials⁸.

Surface chemical structure and functional groups affect the extent of protein adsorption, denaturation, and epitope exposure^{8,71–74}. Compared to the complexity of polymer surface chemistry, SAMs have precisely controlled density and conformation of single or multiple functional groups, and can be easily prepared with different terminal groups. Therefore, SAMs applied on model surfaces are used to study surface chemistry effects on protein adsorption and biological responses^{75,76}. Hydrophobic functionalized surfaces tend towards more protein adsorption than the hydrophilic surfaces⁷⁷. Aminated surfaces are able to form hydrogen bonds with proteins while negative charged groups may react less with proteins compared to other surface^{71,74}. Kidoaki and Matsuda⁷⁸ measured the adhesion forces of blood plasma proteins and above SAMs modified surfaces by AFM and revealed that the hydrophobic methyl functional group exhibited the highest forces for all measured proteins (fibrinogen, albumin, and IgG), while carboxyl functional group was lowest, with fibrinogen exhibiting higher adhesion than albumin and IgG. SAMs also allow production of surfaces containing different functional groups with varying for studies of protein adsorption and

biological responses^{79,80}. For example, binary mixing of alkanethiol SAMs terminated with -COOH and one of the end groups -CH₃, -OH or -NH₂ on gold surfaces influences the surface charge, and alters the secondary structure of covalently bound fibronectin⁸¹. Mixing longer chain methyl- and shorter chain hydroxyl-terminated alkanethiols at different ratios produced a series of surfaces containing different percentages of components. Fibrinogen adsorption, platelet adhesion and activation were found to decrease with increasing -OH groups (increasing hydrophilicity), and there is good correlation between fibrinogen adsorption and platelet adhesion¹⁴.

In contrast to SAMs, surface polymerization can be used to create polymer brushes, where an array of macromolecular chains (polymer) are bonded to a substrate surface at one end, while in an extended conformation to minimize segment-segment overlaps⁸². The polymer chains that extend to the grafting surface can exhibit properties distinctly different from chains in solution so that the substrate surface grafted such polymers can possess novel properties, and have been applied to many areas including medical applications^{83,84}. One of most effective polymer brushes is poly(ethyleneoxide) (PEO) or poly(ethyleneglycol) (PEG), which are considered to possess excellent blood compatibility. PEG is widely used in the production of biomaterials because of a resistance to protein adsorption, weak immunogenicity, and good compatibility with living cells⁸⁵. *In vitro* and *in vivo* experiments show that PEG coatings on PVC drain tubes significantly suppressed plasma protein adsorption and platelet adhesion, leading to reduced risk of thrombus formation, tissue damage, and cytotoxic effects, suggesting PEG grafted surfaces have great potential for clinical applications⁸⁶. Protein resistance of PEG is related to grafting density, chain length, molecular weight, protein size, polymer-protein interactions and protein-surface interactions⁸⁷⁻⁹¹. Generally, it is believed that PEG protein-resistance arises from steric repulsion, chain mobility, and excluded volume effects⁹²⁻⁹⁵. Jin et al.⁹⁶ recently described the protein adsorption mechanism on PEG-grafted polyethylene surface, stating that dense PEG brush can release more trapped water molecules to resist protein adsorption. However, other responses to PEG-modified materials still remain unclear. It should be noted that unanticipated responses such as hypersensitivity reactions caused by PEG-modified surfaces have been reported. There is also evidence that PEG surface may activate the complement system involved in the foreign body reaction to these biomaterials^{97,98} so it is not clear that resisting protein adsorption does not mean that protein interactions cannot still occur.

2.2.3 Surface topography and roughness—Surface topography and roughness are important factors in determining the biological responses to a foreign material. Numerous studies have demonstrated that micro- and nanoscale structured surfaces influence cell behaviors⁹⁹⁻¹⁰¹ such as orientation¹⁰², morphology¹⁰³, adhesion¹⁰⁴, proliferation¹⁰⁵, function regulation¹⁰⁶, and gene expression¹⁰⁷. Topographical surfaces also influence adhesion of bacteria^{108,109} and platelets^{110,111}, and may affect protein adsorption¹¹². The engineering of surfaces with well-defined micro- and nanoscale topographies has become relevant as a strategy to control biological responses^{113,114}.

There is evidence that surface topography might influence the amount of protein adsorbed on surface and alter the adsorption ratio spatial distribution, conformation, and surface binding affinity of different proteins¹¹⁴. As the dimensions of proteins are the nanometer

range, nanoscale topographies are thought to influence proteins, whereas micrometer sized topographic surfaces would appear smooth at the protein scale, minimizing the effect of these topographies on protein behavior. However, there is some evidence that even micro-sized topographies can influence protein interactions with surfaces, and greater amounts of proteins were found on surfaces bearing micrometer scale roughness^{112,115}. In fact, the adsorption of proteins on topographical surfaces appears to be a result of complex processes influenced by surface wettability, surface chemistry, and surface topography, including dimension and shape of features and roughness. However, when assessing the field as a whole, there appears to be no general trend in the amount of protein adsorbed onto topographically-defined surfaces, even when analyzing just a single type of protein.

Surface texturing affects the surface wettability and surface contact area without changing the bulk properties of the material or the surface chemistry. Surface texturing of hydrophobic materials can increase the surface hydrophobicity as evidenced by water contact angles up to $>150^\circ$, termed superhydrophobic, due to enhancing water repellency by trapped air between structures, whereas hydrophilic materials can be made more wettable with texturing (so-called superhydrophilic) due to the increased contact area of the liquid with surface¹¹⁴. Superhydrophobic or superhydrophilic surfaces prepared by addition of surface topography can decrease or increase protein adsorption on surfaces depending on size, shape, and composition of surface feature, and type of proteins. Interactions of proteins or cells with surfaces exhibiting extreme wettability have been reviewed elsewhere¹¹⁶.

The size of topographical feature on hydrophobic surfaces affects protein adsorption. It has been reported that superhydrophobic material surfaces bearing nanoscale topography suppress protein adsorption, while surfaces having micrometer-sized roughness increase protein adsorption^{112,117}. Koh et al.¹¹⁷ suggested that topographical parameters for reducing protein fibrinogen adsorption and platelet responses would be high aspect ratio ($>3:1$) with reduced interspacing ($<200\text{nm}$) or high density. This possible mechanism of reduction in protein adsorption is that nanoscale topography limits the contact of the protein solution to the top of the structures, reducing the surface area for protein adsorption due to the Cassie-Baxter effect, with liquid sitting on top of the rough surface and air in the hollows¹¹². Indeed, the trapped air between the structures plays an important role in reducing protein adsorption on superhydrophobic structured surfaces. Leibner *et al.*¹¹⁸ showed that air trapped within the interstices of fibular structure of ePTFE used in vascular grafts prevented intimate contact with protein solutions. Moreover, air trapped in nanoscale topography blocks the interactions of proteins with surface, and lower protein adsorption. For TiO_2 nanotube surfaces bearing either superhydrophobic or superhydrophilic microtemplated domains, less protein was found on the superhydrophobic domains in the presence of trapped air, and more was bound to superhydrophilic domains with trapped air. However, more proteins were found on superhydrophobic domains than hydrophilic domains after the air was removed by ultra-sonication¹¹⁹. These superhydrophobic nanoscale surfaces reduce protein adsorption, but also promote desorption of proteins under flow conditions¹¹². In fact, almost all proteins could be removed from some nanoscale surfaces, which might be very useful in microfluidic devices¹¹⁶.

Increasing amounts of protein are found on textured surfaces with micro- or nano-sized features. Liu¹²⁰ fabricated a three-dimensional microfluidic chip for DNA and protein analysis using the lotus leaf as a template, and found the device to have 5 times higher sensitivity than flat surface after the superhydrophobic surface was changed to hydrophilic by a simple oxygen plasma treatment. This increase in protein uptake is due to the increase in surface area by topography. Luong-Van et al.¹¹⁴ reported that protein uptake linearly increased with surface area when textured surface structures were wetted by protein solution. However, it is not always sufficient to explain observed increase in protein adsorption strictly by surface area. Chen et al.¹²¹ fabricated dot-like patterns in PDMS with protrusion diameter bottom of 12 μm , and intervals of 7.5 μm between, and found that adsorbed fibrinogen increased 46% but surface area increased only about 8%, suggesting that the surface area factor was not solely responsible for increases in protein adsorption. Increases in protein adsorption are also found with nanoscale roughness. Rechendorff et al.¹²² found fibrinogen adsorption on tantalum films increased with increasing root-mean-square roughness (2.0 to 32.9 nm) beyond the accompanying increase in surface area. Vlachopoulou et al.¹¹⁵ observed that BSA adsorption was enhanced up to 10 times on high-aspect-ratio plasma-induced nanotextured PDMS surface. Scopelliti et al.¹²³ quantified the protein adsorption on titanium with surface roughness ranging from 14 to 32 nm, and found fibrinogen and BSA adsorption 500% higher than the theoretical monolayer coverage on rough surfaces. The suggested possible mechanism for these findings is that surface nanoscale pores promote protein nucleation and aggregation so that proteins are trapped in pores, contributing to protein multilayer formation.

2.3 Fibrinogen adsorption and functional activity and platelet adhesion

Fibrinogen is a central protein in the process of biomaterial-induced thrombosis, promoting thrombus formation through binding of the platelet integrin receptor $\alpha_{\text{IIb}}\beta_3$ (GPIIb/IIIa), leading to platelet immobilization, activation and aggregation¹²⁴, and also being the precursor to fibrin, the predominant structural component in blood clotting/coagulation. Fibrinogen is a symmetric molecule, composed of two sets of three intertwined polypeptide chains termed the A α , B β and γ chains, with a total molecular weight of 340 kDa. The six chains fold together in a trinodular structure with a center globular knot (termed the E domain) containing portions of all six polypeptide chains, and flanked on each side by a larger globular structure (the D domain) containing one of the sets of three polypeptide chains. Each fibrinogen molecule possesses three pairs of potential platelet-binding peptide sequences, two RGD sequences in each of the A α chains (RGDF and RGDS) and a dodecapeptide sequence (HHLGGAKQAGDV) in each of the γ chains; the γ chain dodecapeptide sequence being the primary ligand for platelet adhesion to adsorbed fibrinogen^{22,125}. It is now becoming accepted that the conformation and availability of platelet-binding sites in fibrinogen are more important in mediating platelet adhesion than is the amount of adsorbed protein^{126,127}. The discussion of conformational structure and functional activity of adsorbed proteins in this manuscript is mainly based on AFM measurement. This is not to discount the other techniques and methods have been used to study the conformation of proteins adsorbed on biomaterial surface, such as time-of-flight secondary ion mass spectrometry with principal component analysis¹²⁸, and improved circular dichroism methods¹²⁹.

2.3.1 Time-dependent conformational change and energy profiles involved in fibrinogen adsorption on surfaces—Fibrinogen undergoes conformational changes following adsorption to biomaterial substrates^{41,130,131}. Conformational changes are both time- and surface-dependent. It was observed that structural deformation (spreading) of fibrinogen increased with material surface hydrophobicity; quantitative analysis of the different domains showed that the overall molecular length and widths of the individual D and E domains increased while the heights decreased when the material surface is more hydrophobic¹³². The time dependence of these changes in fibrinogen upon adsorption was directly observed in our lab¹³³.

Figure 3 illustrates the time-dependent conformational changes in fibrinogen on a hydrophobic highly ordered pyrolytic graphite (HOPG) material (Fig. 3a) and a hydrophilic muscovite mica substrate (Fig. 3b). Fibrinogen was continuously imaged by AFM for ~ 2 hours. Spreading curves were generated by plotting heights of D and E domains of individual fibrinogen molecules. On HOPG, a clear decrease in height was observed with time, indicating protein spreading (Fig. 3c). On mica, the height of the D and E domains increased with time (up to 60 min) eventually reaching a plateau height of 2.1 nm (Fig. 3d). Together with other studies, these data demonstrate that proteins undergo conformational changes following contact with material surfaces and these changes are dependent on the properties of the underlying substrate.

Time-dependent unfolding of protein on surface causes variations in surface-protein interaction. AFM force mode makes it possible to measure the interaction strength of protein-surface interactions by bringing a protein into contact with a material surface and then separating. Assessing the effects of contact times on protein adhesion forces reveals information about the unfolding energy involved in protein-surface interactions. Low-density polyethylene (LDPE) was modified by glow discharge plasma to yield surfaces spanning a range of water wettability. Adhesion forces between protein and substrate increased with increasing protein contact time up to 50sec, and with smaller adhesion forces measured on wettable surfaces. Changes in adhesion forces with contact time were modeled by a simple exponential equation and energy barriers for protein unfolding were calculated with the Arrhenius equation¹³⁴. The analysis yielded activation energies of 18.0–22.6 *kT* for human fibrinogen on hydrophobic surfaces, and 18.6–23.2 *kT* for hydrophilic surfaces⁶⁸. Based on the observations of changes in adhesion force at early time points and the changes in height of D and E domains in fibrinogen at ~2 hours, a two-step spreading model was proposed, with the first step being very rapid (on the order of seconds to minutes) and involving rearrangement of protein surface amino acids, and the second step taking much longer and involving rearrangement of the internal amino acids to the hydrophobic surface⁶⁸.

The energy profiles in interactions of protein/surface can be further explored through examining the interaction forces between protein and surface as a function of the loading rate using AFM measurement, where loading rate (r_f) is defined as the product of probe retraction velocity and spring constant of the cantilever. In measuring the strength of bonds between biological receptor molecules and their ligands, the bond strengths measured under a variety of loading rates are governed by the barriers in the energy landscape along the force-driven, bond dissociation pathway^{135–137}. Using the Bell model¹³⁸, dissociation rates

and energy barriers in biological molecule reactions can be quantified. These measurements can be extended to probe protein-surface interactions and reveal the energy profiles in interactions of fibrinogen with surfaces¹³⁹.

Dynamic forces were measured between colloid materials modified with SAMs of different wettability and fibrinogen. Figure 4 illustrates the adhesive forces between fibrinogen and these model surfaces as a function of loading rate. The linear relationship between adhesion forces and the $\ln(r_f)$ for wettable surfaces (bare glass and 3-aminopropyltrichlorosilane (APS)-coated) suggests a single energy barrier for interactions of fibrinogen with these hydrophilic surfaces (Figs. 4a and b), while two line segments with differing slopes between hydrophobic surfaces and fibrinogen (Figs. 4c and d) suggests that there are multiple energy barriers and multiple transition states during interactions of fibrinogen with hydrophobic surfaces¹³⁹. Dynamic force spectroscopy analysis was performed to fit the plots of adhesion force against $\ln(r_f)$ by following equation:

$$F = \frac{kT}{\chi_\beta} \ln(r_f) - \frac{kT}{\chi_\beta} \ln\left(\frac{k_{\text{off}} kT}{\chi_\beta}\right)$$

where k is the Boltzmann constant, T is the absolute temperature, k_{off} is the thermal off-rate at zero force, and χ_β is the effective distance (\AA) between the bound and transition states along the direction of applied force that measures the width of the energy well that traps the interacting molecules in the bond state. The results show that the off-rate (k_{off}) for protein interactions with hydrophilic surfaces lie in the range of 1.35–2.92 s^{-1} , about 3–10 fold greater than those for protein interactions with hydrophobic surfaces at the lower loading rates (Table 3). The bond lifetimes on the hydrophobic surfaces are longer than they are on hydrophilic surfaces, consistent with increased adhesion forces observed between proteins and hydrophobic surfaces. This analysis of dynamic forces reveals energy profiles in interactions of protein and surfaces, providing insights into the role of surface wettability on protein interaction with biomaterials.

2.3.2 Time-dependent fibrinogen functional activity and platelet adhesion—

Conformational change following protein adsorption results in transient exposure of functional epitopes and time-dependent platelet adhesion. Studies of the availability of functional epitopes in adsorbed proteins help to understand platelet responses to surfaces. Although many techniques have been applied to analyze the protein structure, immunological methods using antibodies directly recognize the functional/activity changes in proteins^{41,140}. AFM with antibody-functionalized probes offers a unique method to recognize and identify adsorbed proteins^{141–143}. It allows for examination of multiple proteins at the molecular scale^{144,145}, and also probes the conformation, orientation, and activity of adsorbed proteins with appropriate choice of antibodies¹⁴⁶. As the γ -chain dodecapeptide sequence in fibrinogen is the primary ligand for platelet adhesion to adsorbed fibrinogen, the availability of the dodecapeptide can be viewed as an indicator of the functional activity of fibrinogen as related to the conformation and/or orientation of the adsorbed protein^{18,41,147}.

The force measurement mode of AFM provides an approach to study fibrinogen functional activity as recognized by monoclonal antibodies (mAb) at differing residence times. An AFM probe coupled with mAb that recognizes the last 20 amino acids of the γ chain of fibrinogen (γ 392–411) was used to measure the interaction forces between mAb and adsorbed fibrinogen. Figure 5(a) illustrates the probability of this mAb encountering its antigen in the platelet-binding region as a function of fibrinogen residence time after adsorption to a hydrophilic muscovite mica surface. Data demonstrate that the activity of fibrinogen is time-dependent, with the maximum likelihood of recognition occurring at around 45 min and decreasing at longer residence times¹⁴⁸. Macro-scale assessments of platelet adhesion using the LDH assay showed that platelet adhesion varied with residence time of fibrinogen, and also reached a peak at ~45 min (Fig. 5b), correlating well with the molecular scale AFM results.

Protein adsorption to biomaterials occurs from protein mixtures in most biomedical applications. The presence of other proteins results in adsorption competition¹⁴⁹, and also alters the conformational structure and dynamics of adsorbed proteins. Our studies show that inclusion of albumin into a fibrinogen solution moves the peak biological activity of fibrinogen to earlier residence time points¹⁵⁰. With a protein concentration ratio of 50:50 (Fibrinogen:BSA) in binary protein solutions, the peak biological activity of adsorbed fibrinogen shifts to ~15 min and activity decreases with increasing residence time. In the case of 10% fibrinogen in a binary protein solution, no peaks in fibrinogen activity were observed, and the activity remained essentially steady at a lower value. Measurement of platelet adhesion carried out under the same conditions show correlation between changes in activity of fibrinogen and platelet adhesion.

It should be noted that the studies of protein conformation were often carried on the model substrates such as mica and gold while the conformation of fibrinogen response to biomedical related polymeric surface is less reported. There is evidence that conformation of protein is related to chemistry and surface properties of polymeric biomaterials. Berglin et al.¹⁵¹ reported that the conformation of fibrinogen adsorbed to acrylic polymeric substrate is related to polymer chain flexibility and fibrinogen maintained a more native conformation on the flexible polymer. More work is needed to focus on protein conformational response and platelet adhesion to individual biomedical polymeric surfaces.

2.4 Protein adsorption and platelet adhesion on polyurethane biomaterial surface

2.4.1 Microphase separation structure and protein adsorption—The surface chemical structure of polymeric biomaterials is often complicated compared to model materials. Block copolymers are composed of different segments having different properties, and are widely used in biomedical applications. One of the most important structures formed in block copolymers is found in polyurethane materials, and is termed microphase separation. This microphase separation creates a surface microenvironment where chemical composition and physicochemical properties are distributed spatially, and therefore have the potential to induce different levels of protein adsorption/activity across the material surface. Polyurethane (PU) and poly(urethane urea) (PUU) block copolymers are used in a variety of blood-contacting medical devices due to their broad range of mechanical properties, fatigue

resistance and relatively-good hemocompatibility^{152–154}, and these properties are believed to arise from the presence of separated microphases in the copolymers. Microphase separation is a result of thermodynamic immiscibility of the polar hard segments (made from diisocyanate and either a diol or a diamine) and the relatively non-polar soft segments (often a polyether but increasingly PU materials are being made from polycarbonates or mixed PTMO/PDMS precursors), where hard domains act as thermoplastic crosslinks and reinforcing agents dispersed in a soft segment rich background matrix¹⁵⁵.

Numerous studies have shown that the Microphase separation structure influences biological responses to surfaces, including protein adsorption^{156–159}, platelet adhesion/activation^{160,161}, and cell attachment^{162,163}. We measured adhesion forces of a protein modified AFM probe with a PUU surface, and correlated mechanical properties with local adhesion forces. Results showed that low adhesion forces were associated with hard domain regions¹⁶⁴. A nanogold-labeled protein conjugate was further used to visualize individual protein adsorption to the separate microstructures on PUU surfaces (Figure 6). Analysis of the image shows an ~ 50% increase in the number of proteins per unit area on the soft segment material compared to the hard segment regions, suggesting preferential adsorption to the less polar, more hydrophobic, soft segment region¹⁶⁴.

2.4.2 Effect of microphase chemistry of polyurethane on protein adsorption and platelet adhesion—A variety of polyurethanes have been developed and synthesized for meeting varied requirements in medical device applications¹⁵⁵. Principle candidate PU biomaterials generally consist of similar hard segments, but with different hard segment content and varying soft segment chemistry. The soft segments used in PU include polyether, aliphatic polycarbonate (PC), and polydimethylsiloxan (PDMS). These different soft segments produce different phase separation structures and affect material properties^{165,166}, resulting in different biological responses including fibrinogen functional activity and platelet adhesion¹⁶⁷.

As a first effort at relating polymer microphase chemistry and structure, protein activity and subsequent platelet adhesion, we assessed the properties of a series of polyurethane materials derived from polycarbonate (PC-PU40), polytetramethylene oxide (PTMO-PU40), and PDMS (PDMS-PU series) chemistries. The number following the polymer chemistry description is the hard segment content of the material. Figure 7 summarizes the fibrinogen adsorption as recognized by polyclonal antibody, functional activity as recognized by mAb, and platelet adhesion on these materials. The amount of adsorbed fibrinogen recognized generally increased with hard segment content except for one material, the PDMS-PU32.5 (Fig. 7a). This trend was roughly consistent with platelet adhesion results (Fig. 7c). However, larger amounts of fibrinogen were measured on the PC-PU40 and PTMO-PU40 surfaces despite the fact that fewer platelets adhered on these materials. This inconsistency between platelet adhesion and fibrinogen adsorption suggests that the amounts of fibrinogen adsorbed may not be the key factor determining platelet adhesion. Rather, the availability of γ chain dodecapeptide in fibrinogen is more important. Results showed that fewer specific binding events recognized by the mAb against fibrinogen γ 392–411 were measured with proteins adsorbed on PC-PU40 and PTMO-PU40 materials (Fig. 7b). Generally, the activity of fibrinogen measured by the mAb was consistent with platelet adhesion on the three types

of polymers tested (Fig. 8). However, it should be noted that there are two other RGD sites in the α chain that may also be involved in mediating platelet adhesion, acting perhaps separately or synergistically with the C-terminus of the γ chain¹²⁶.

2.4.3 Protein adsorption and platelet adhesion with dynamic restructuring polyurethane biomaterial surfaces

—Microphase separation structure in segmented polyurethanes is the result of thermodynamic immiscibility of the hard and soft segments. The structure is influenced by block length and chemical composition of segments, but is also affected by the environmental conditions. Studies show that polyurethane materials undergo reorganization and reorientation in aqueous environments for minimizing surface free energy, resulting in enhancement of hard domains at the surface^{168–170}. Figure 9 illustrates sequential *in-situ* AFM phase images of a PUU film following hydration in phosphate buffered saline (PBS) for 21 hrs and shows that hard domains in PUU undergo reorientation. The analysis of phase angle distribution showed that the percentage of areas having high phase angle shift increased with hydration time, suggesting enrichment of hard domains at the surface¹⁶⁴. Hydration-induced surface enrichment of hard domains was also confirmed by measurements of local mechanical properties using a nano-indentation technique¹⁷¹. Water contact angle measurements showed that the surface becomes more wettable with hydration time due to the migration of hydrophilic domains to surface, consistent with the observation of others¹⁷². Dynamic restructuring of hard domains increases surface wettability and reduces the interaction forces between proteins and surfaces. For example, the water contact angle at the initial stage of PUU was near 90° and adhesion force with BSA was 1.5±0.3 nN, while the contact angle decreased to 70° and adhesion force with BSA dropped to 0.5 ± 0.3 nN at 53 hrs hydration¹⁶⁴.

Dynamic restructuring of PUU hard domains of PUU also decreases the functional activity of adsorbed fibrinogen and subsequent platelet adhesion. On a series of PU biomaterials with different soft segments, higher fibrinogen activity was observed on surfaces at hydration time of 1 hr compared to samples with longer hydration times (Fig. 10a). Immuno-AFM force measurements identified the platelet binding epitope in adsorbed fibrinogen 51%, 56% and 53% of the measures for the PDMS/PHMO-PU40, PHEC-PU40 and PTMO-PU40 surfaces, respectively, when samples were hydrated in PBS for 1 hr. Increasing the hydration time of the polymers decreased this measure of activity for fibrinogen on these materials. When hydrated for 10 days prior to a 10 minute incubation with fibrinogen, only 18% (PDMS/PHMO-PU40), 14% (PHEC-PU40) and 12% (PTMO-PU40) of the force measures showed an interaction for the mAb against γ ^{392–411}. Corresponding platelet adhesion is shown in Figure 10b, and shows that regardless of the soft segment chemistry, PU materials show the highest platelet adhesion at 1 hr hydration time. This result is consistent with the higher activities of fibrinogen found on the surfaces, again showing that the availability of the γ -chain dodecapeptide in fibrinogen correlates with platelet adhesion.

Taken together, phase restructuring during hydration changes the local microenvironments and the physicochemical/mechanical properties at the interface of polyurethane. As a result, the interactions of blood with biomaterial surfaces, including platelet adhesion/activation, thrombus formation, and cell attachment/growth, is expected to change during this process.

Fibrinogen plays the dual roles of serving as a ligand for platelet adhesion to the surface and also for linking platelets together into aggregates²⁰. The decrease in platelet adhesion on PU surfaces after hydration, consistent with the measured decrease in fibrinogen bioactivity on the surface, strongly suggests that phase rearrangements in PU could affect thrombus formation. The varying degrees of phase separation/restructuring in different PU polymers would therefore be expected to affect protein adsorption, platelet adhesion/activation, cell attachment and proliferation, as well as bacterial adhesion.

2.5 Protein adsorption and platelet adhesion on textured biomaterial surface

2.5.1 Surface topography controlling biological responses to surface—Physical approaches to biomaterial design have been a promising route for controlling biological responses¹¹³. Design and control material surface topography has had a profound impact on cell and tissue behaviors including platelet adhesion and bacterial adhesion, as well as cell growth. The motivation to use physical properties to control biological response comes from nature, for example, the skin of sharks, the mussel, lotus leaf, and the inner surface of blood vessels^{173–177} have all provided templates for the fabrication of materials. These natural surfaces are micro-topographically structured and possess intrinsic non-adherence function. These materials have inspired novel biointerface materials¹⁷⁸. Materials with topographical features mimicking shark skin show resistance to marine biofouling^{108,179–181}. This topography applied into biomedical-related polymeric materials was found to disrupt biofilm formation of *Staphylococcal aureus*¹⁸² and inhibit platelet adhesion¹¹⁷, demonstrating applicability for medical devices. We developed textured surfaces having ordered array of pillars and demonstrated that these materials reduced the adhesion of *Staphylococcal* strains and inhibit biofilm formation^{183,184}. Previous studies showed these materials inhibit platelet adhesion and activation^{110,111}.

Surface features influencing biological responses include feature shape and size, distance between features, and organization of features¹⁸⁵. Submicron textured PUU surfaces (dimension/separation of 400/400 nm and 500/500 nm) consisting of ordered arrays of pillars with dimension and separation of pillars less than 1 μm produce a dramatic reduction in accessible contact area for bacteria or platelet interaction, and minimize opportunities for bacteria and platelet to be trapped in inter-pillar spaces^{111,183}. This design is consistent with the finding of other researchers that show that surface structures are most effective when in the range of 50 to 90% of the diameter or length of settling organisms¹⁸⁰ and the height of nanocylinders on structured surface should be >160 nm¹⁸⁶.

PUU textured films can be physically prepared by soft lithography, two-stage replication molding techniques where a master pattern is fabricated in silicon, a silicone mold is cast and PUU replicas created from the silicone negative¹¹¹. A variety of techniques have been applied to characterize the surfaces. Figure 11 illustrates representative SEM and AFM images of a textured PUU surface having a 500/500 nm pattern. Analysis of these surfaces revealed the top surface of this 500/500 nm pattern is only about 24.5% of the nominal surface area. Reductions in surface contact area reduce opportunity for platelet or bacteria to adhere to the surface. Another characteristic of textured surfaces can be an increase in surface hydrophobicity. Water contact angles of submicron patterned PUU surface reach

nearly 145°, near to the “superhydrophobic”, while the water contact angle for a smooth surface PUU surface was just 93°¹⁸³. The increase in hydrophobicity of textured surfaces is believed to be due to air captured in the spaces between pillars¹⁸⁷. Evidence suggests that both reduction in surface contact area and increase in hydrophobicity influence the biological events on surfaces.

2.5.2 Protein adsorption on textured biomaterial surfaces—Protein adsorption on topographically-modified surfaces depends on both the surface features and the proteins, as described in Section 2.2.3. The amount of total protein or of a selective protein is quantified to indicate the behavior of protein adsorption, often by using traditional colorimetric protein assays¹⁸⁸, the enzyme-linked immunosorbent assay (ELISA)¹¹⁷, radiolabeling¹⁸⁹, electrophoresis¹¹⁸, quartz crystal microbalance (QCM)¹²², or any of the myriad other techniques available. The main weakness to these methods is that it only provides the overall quantification of protein adsorption and often misses one of the most important characteristics of topographic surfaces – the increase in surface area. It has been suggested that protein adsorption on topographic surfaces may be non-uniform and vary locally.

Directly imaging or detecting single molecular proteins on surface is difficult, especially on polymeric biomaterial surfaces with textured topographies. With the development of nanotechnology in biology and medicine, nanoparticles conjugated with proteins have numerous applications in sensing, imaging, delivery, catalysis, therapy and control of protein structure and activity¹⁹⁰. Among nanoparticles, gold nanoparticles are receiving significant attention because their unique physical, chemical, and biological properties are quite different from the bulk of their counterparts¹⁹¹ and can be easily detected by microscopy techniques such as electron microscopy and AFM. We used nanogold particles conjugated to an anti-fibrinogen antibody and identified fibrinogen from dual-component protein solutions following adsorption and imaging by AFM¹⁴⁴. We have also characterized the adsorption of proteins coupled with 6 nm nanogold beads on textured surface by field emission scanning electron microscopy (FESEM) and transmission electron microscope (TEM) (Fig. 12). The bright spots in SEM images illustrate the nanogold beads, indicating the presence of adsorbed protein. FESEM image demonstrates that the beads distribute around the top and bottom surfaces, and more beads (i.e., proteins) were detected on the bottom surface than on the top surface of pillars. The side view of a single pillar (TEM, Fig 12b) shows that the beads (i.e., proteins) are distributed on pillar top, edge, sidewall, and bottom areas. These results suggested that protein adsorption on textured surface with submicron pillars occurred not only on the top surface of pillars, but also on the sidewall and bottom areas. As a result, the quantification of protein adsorption by other methods should be normalized to the entire surface area including sidewall of pillars and other topographic features. Figure 13 illustrates the trends of adsorption of human albumin on textured PUU films with smooth, 400/400 nm, and 500/500 nm patterns, measured by ELISA. Results demonstrated that less protein was adsorbed on textured surfaces, suggesting that submicron textured surface may produce the heterogeneous local microenvironments at pillar surface area and alters the protein adsorption behavior, resulting in different densities of protein adsorbed.

2.5.3 Platelet adhesion on textured biomaterial surface—Platelet response to topographical surface is influenced by the surface physical features and chemical composition of substrate. Koh et al.¹⁹² prepared poly(lactic-co-glycolic-acid) films with multi-walled carbon nanotubes having two different nanotube orientations, either random or vertically aligned, and found the film with vertically aligned nanotubes exhibited very low levels of fibrinogen adsorption and platelet adhesion. Reduction of platelet adhesion was regarded to be attributed to both chemical and topographical effects. They further studied the effect of topographical features on platelet adhesion and concluded that the optimal range of topographical dimensions of the pillars with height of 300–800 nm, width of 100–200 nm, and interspacing distances <200 nm¹¹⁷. Other researchers using polymer demixing methods created structured polymer films with typical feature sizes ranging from the nanometer to the micrometer, and found that increasing the feature size of the surface encourages von Willebrand factor adsorption and thus platelet adhesion and consequent thrombus formation¹⁹³. Ye et al.¹⁹⁴ used dual scale structures with nano and micro dimensions as model substrates to investigate the impact of surface topography on platelet response, and found fewer platelets adhesion and no activation on such surfaces. We measured the platelet adhesion on smooth and patterned PUU film surfaces at low shear stresses across a 0–10 dyn/cm² range. Significant reduction in platelet adhesion was found on 400/400 nm and 700/700 nm PUUs for shear stress <5 dyn/cm² compared to smooth. Additionally, non-adherent platelets did not demonstrate increased activation after transient exposure to textured surface. These results suggest that using surface topography can reduce platelet adhesion and activation¹¹¹.

Protein adsorption affects platelet adhesion and activation on textured biomaterial surface. A series of PUU film surfaces were pre-adsorbed with proteins (human albumin, fibrinogen, and platelet poor plasma (PPP)) at 37°C for 1 hr. Samples were then incubated in platelet-containing solution (2.50×10⁸/ml) at 37°C for 1 hr in a microwell plate¹⁹⁵ under static condition. Samples were labeled for fluorescence microscopy to examine platelet adhesion and activation. Activation of platelets adherent on surfaces was inferred from platelet circularity, with higher circularity indicative of less active platelets. Results showed that the both 400/400 nm and 500/500 nm textured surfaces significantly reduced platelet adhesion and activation compared to the smooth surfaces (Fig. 14). Fibrinogen increases platelet adhesion on smooth surfaces while albumin and PPP suppress adhesion, as expected. It is interesting to see that platelet adhesion on fibrinogen-adsorbed textured surfaces is lower than that on smooth surfaces, and even lower than smooth surfaces pre-adsorbed with albumin. This suggests that fibrinogen adsorbed on textured surface may have less functional activity, resulting in lower platelet adhesion. This is also evidenced by the circularity of platelets, with those adhered on fibrinogen-textured surfaces having higher circularity than platelets on smooth or textured surfaces (without protein adsorption), indicating that textured surfaces reduce platelet activation.

Surface topography can potentially affect protein orientation and conformation structure upon adsorption. There is evidence to show that the curvature of features affects the unfolding of proteins (e.g., fibrinogen) at feature edges and make receptor ligands more (or less) accessible, resulting in differing cell adhesion¹⁹⁶. As the dimension of top surface of

submicron-sized pillars is larger than dimensions of proteins which lie in the nanometer range, the top surface is not as flat as smooth, but with a slight curve, due to the fabrication. Moreover, the edges of pillar features are generally on the length scale of a protein. Therefore, both top surface and edges may affect protein adsorption and its conformation, causing different platelet responses. Characterization of protein conformational structure and functional activity at local area on top surface of features will largely contribute to understand the cell behaviors on textured surface and will be more interesting to biomaterial scientists.

3. Blood coagulation at biomaterial interface

3.1 Blood coagulation cascade and contact activation of FXII

The process of biomaterial associated thrombosis consists of both platelet-mediated reactions (platelet adhesion, activation, and aggregation) and coagulation of blood plasma. Plasma protein interactions with surfaces trigger the coagulation cascade of blood, resulting in thrombus production and formation of a fibrin clot. Coagulation involves a series of self-amplifying, zymogen-enzyme conversions which are traditionally grouped as the intrinsic and extrinsic pathways^{197,198}. Both pathways are initiated separately but merge into a common pathway leading to thrombin (FIIa), which hydrolyses fibrinogen into fibrin, which assembles and causes plasma to become clotted^{198–201}. The extrinsic pathway is responsible for hemostatic control and response to vascular injury while the intrinsic coagulation cascade has little physiological significance under normal conditions. However, it is widely believed that the intrinsic pathway is triggered by blood contact with artificial materials and it is an important cause for poor hemocompatibility of biomaterials^{202–204}.

The initiation of the intrinsic pathway is generally referred as contact activation, and primarily mainly involves four proteins: coagulation factor XII (FXII, Hageman factor), prekallikrein (PK, Fletcher factor), high-molecular weight kininogen (HMWK, Fitzgerald factor), and coagulation factor XI (FXI, Plasma Thromboplastin Antecedent). The traditional biochemistry of contact activation shows that FXII is converted to the active enzyme form FXIIa. FXIIa can be produced by at least three different biochemical reactions (Fig. 15): (1) contact autoactivation process - FXII interacts with a procoagulant surface and converts into active enzyme form FXIIa through autoactivation due to conformational structural change upon binding of FXII to the surface ($\text{FXII} \rightarrow \text{FXIIa}$)²⁰⁵; (2) reciprocal activation - the generated FXIIa can in turn cleave PK bound to the surface as a complex with HMWK to produce Kal which enzymatically acts on FXII upon binding on surface²⁰⁶; (3) autohydrolysis or self-amplification - FXIIa hydrolyzes FXII to produce FXIIa ($\text{FXIIa} + \text{FXII} \rightarrow 2\text{FXIIa}$)²⁰⁷. Ultimately, FXIIa activates FXI bound at the surface as complex with HMWK to generate FXIa leading to propagation of subsequent coagulation cascade reactions^{198,208,209}.

Each of these three biochemical reactions can be observed in defined solutions, however, all three reactions do not occur in plasma to the same extent. For example, autohydrolysis is thought to be a contributing reaction in buffer solutions of FXIIa, but does appear to be a significant reaction in plasma, because the preferred FXIIa substrates PK and FXI are not available in neat buffer solutions of FXII²⁰⁷. Mathematical modeling of the intrinsic cascade

predicts that the primary mechanism for activation of coagulation involves autoactivation of FXII by the procoagulant surface or kallikrein-mediated reciprocal activation of FXII, but FXIIa-induced self-amplification of FXII is insignificant²¹⁰. In plasma, FXIIa is principally produced by autoactivation which is then substantially increased by reciprocal-activation pathways. Analysis of the amounts of FXIIa produced showed that kal-mediated reciprocal-activation pathway principally contributes to FXIIa generation at 75%²¹¹.

FXII contact activation is surface-dependent. A variety of parameters including surface energy or wettability, surface chemistry, and surface area have been well investigated on contact activation^{212–214}. Since common observations clearly show that plasma coagulation is more efficient in activation by contact with anionic^{215,216} or hydrophilic surfaces^{13,217}, it was concluded that contact autoactivation of FXII was more specific for hydrophilic surfaces than hydrophobic surfaces based on traditional biochemistry theory. However, experimental evidence demonstrated that hydrophobic and hydrophilic surfaces have nearly equal autoactivation properties in neat – buffer solution of FXII, i.e. contact activation of FXII is not specific to anionic hydrophilic surfaces in neat buffer^{214,218}. In fact, contact activation of FXII in neat-buffer solution exhibits a parabolic profile when scaled as a function of surface energy. Nearly equal activation is observed at both extremes of activator water wettability, and falls through a broad minimum with water contact angle in the range of $20 < \tau < 40$ dyn/cm ($55^\circ < \theta < 75^\circ$), suggesting that materials with surface energy in this range exhibit minimal activation of blood-plasma coagulation (Fig. 16)²¹³. A similar result was obtained by preparing mixtures of various thiol-based self-assembled monolayers at different concentrations and the mixtures of thiols appear to prolong coagulation time beyond that seen with the background container alone (unpublished results).

Contribution of surface area to contact activation of FXII is inconclusive. The plausible explanation consistent with current understanding of coagulation-cascade biochemistry is that procoagulant stimulus arising from the activation complex of the intrinsic pathway is dependent on activator surface area. In plasma, FXII autoactivation can be studied *in vitro* using activator surface-area titration of plasma coagulation and the titration plot of clotting time against activator surface areas is generally characterized to be asymptotic and leading to a lower clotting time plateau which depends on activator surface energy^{213,219}. The total amount of FXIIa yield is also characterized by the FXIIa titration curve and results show that total contact activation of FXII scales with increasing surface area of procoagulant and the normalized amount of enzyme produced per-unit-area procoagulant is constant²¹¹. All these results suggested that contact activation of FXII increased with surface area in plasma depending on surface energy. However, recent results showed that activation of FXII dissolved in buffer, protein cocktail, heat-denatured serum, and FXI deficient plasma did not exhibit activator surface-area dependence, suggesting activator surface-area dependence observed in contact activation of plasma coagulation does not solely arise at the FXII activation step of the intrinsic pathway²²⁰. Other data show that contact activation of FXII either in plasma or in buffer is surface-area dependence for hydrophilic activator but independence for hydrophobic activator. This is most likely a result of synergistic effects of protein interaction (adsorption/desorption) and contact activation on surfaces.

Contact activation of blood FXII is moderated by the protein composition of the fluid phase and protein adsorption competition on surfaces depending on surface energy. Zhuo et al.²²¹ found that FXIIa yield arising from FXII contact activation on hydrophilic surfaces in a protein cocktail was much greater than that obtained in buffer solution containing only FXII, but it was contrary on the hydrophobic surface. Results suggest that contact activation in the presence of proteins unrelated to the plasma coagulation cascade leads to an apparent specificity for hydrophilic surfaces due to a relative diminution of activation at hydrophobic surfaces and an enhancement at hydrophilic surfaces. It was also found that the rate of FXIIa accumulation decreased with time leading to a steady-state FXIIa yield for both hydrophilic and hydrophobic surfaces in protein and buffer solutions, suggesting that activation competed with a yet unknown autoinhibition reaction to inhibit the conversion of FXII to FXIIa. Thus plasma proteins play a dual role in moderating contact activation of the plasma coagulation cascade. The principle role is inhibiting FXII contact with activating surfaces, but these proteins also displace FXIIa from an activating surface into solution where FXIIa is involved in the subsequent steps of the plasma coagulation cascade²²². The importance of protein adsorption competition in contact activation was also observed in moderating interactions between FXII and PK, which are components of the reciprocal-activations²¹⁸. All these observations implicate protein adsorption competition is an important mediator of contact activation and must be included in any comprehensive mechanisms of surface-induced blood coagulation.

3.2 Blood coagulation in response to surface with chemical heterogeneity

The effects of surface chemistry or surface energy on blood coagulation is often investigated using surfaces with well-defined homogeneous chemistry such as silane or thiol SAMs with various terminating functional groups^{212,213,223}. However, nanometer-scale patterning or domain formation with different chemistry is of interest as the block copolymers such as polyurethane have been widely used in biomedical applications. The microphase separation structures with nanoscale chemical domains contribute to the hemocompatibility of block biomaterial. Polyurethanes have shown to reduce contact activation with lower amidolytic activities by FXII and kallikrein compared to glass in plasma²²⁴. Our studies indicated that the microphase separation of polyurethanes produces a local surface microenvironment and influences protein and cellular responses to the surfaces^{164,171}, suggesting the nanoscale chemistry heterogeneity may influence the blood coagulation. Miller et al.²²⁵ investigated silane-based mixed films on glass substrates having nanoscale chemical heterogeneity formed by the stepwise adsorption of an amine-terminated silane, 3-aminopropyltriethoxysilane (APS), and a methyl-terminated silane, n-butyltrichlorosilane (BTS), and found that the mixed films consists of 400 nm islands of an amine-terminated silane dispersed in a butyltrichlorosilane background on glass substrates and such nanometer-scale chemical heterogeneity were inefficient at activating the intrinsic coagulation of human blood plasma when compared to their one-component control surfaces. Results suggested that the nanoscale engineering of surface may provide a route to biomaterials with improved hemocompatibility.

With the development of nanotechnology, it is possible to create surfaces having well-defined surface chemistry combined with nanometer scale patterning. Figure 17 illustrates

an example of linear patterned surface chemistry consisting of carboxyl-terminated and methyl-terminated thiols on gold-coated glass coverslips prepared by microcontact printing. The wider bands in the friction trace and retrace microscopy images correspond to the octadecanethiol stamped areas, while the thinner bands correspond to the 11-mercaptopundecanoic acid backfilled areas. The cumulative area of 11-mercaptopundecanoic acid backfilled areas was estimated to be 40% by comparing band widths. The measured water contact angle of patterned surface is $73\pm 4^\circ$, while the contact angles for octadecanethiol and mercaptopundecanoic acid controlled surfaces are $96\pm 1^\circ$ and $24\pm 1^\circ$, as expected. The plasma coagulation characteristics of nanoscale chemical patterned surface were assayed as coagulation time of platelet poor plasma incubated with gold coverslips. Results show that nanometer-scale patterning of surfaces leads to statistically-significant increases in coagulation times when compared to carboxyl surface area (Fig 17a) or simply to the measured fraction of carboxyl in mixed film, relative to surfaces with equivalent, macro-scale areas of carboxyl and methyl-terminated thiol chemisorption. The increased coagulation times by patterning suggest that the patterning of the surface may affect activation of the intrinsic coagulation cascade and that it is not simply a matter of the cumulative carboxyl-terminated thiol backfilled area.

4. Summary and prospective

Biocompatibility is the central theme for the design and application of polymeric biomaterials. Blood coagulation and thrombosis resulting from blood-material interactions remain a challenge in the use of blood-contacting materials. Protein adsorption is a critical early event during the interaction of blood with implanted biomaterials and it is the adsorbed proteins, rather than the surface itself, that mediates the subsequent biological responses, including platelet adhesion and thrombus formation on surfaces. However, surface properties including surface chemistry, surface energy, and surface topography determines the protein composition, structure, function of the adsorbed protein layer. Thus the biological events occurring on biomaterial surface are the results of interactions of material surface, proteins, and cells at interface. The future of research in this field will rely on a better analysis and understanding of the influence of surface properties from microscale to nanoscale on the quality, quantity, and conformation of proteins adsorbed on surfaces. The development of new techniques or the adaption of existing techniques as well as the development of new biomaterials with specific features will largely help to obtain these information.

Protein undergoes the conformational structural change upon adsorption, e.g., fibrinogen, and influences the subsequent biological responses. It would be more important to reveal the conformation state or functional activity of protein adsorbed than the amount of protein in mediating the platelet adhesion and activation¹²⁷. The current understanding of plasma protein adsorption on biomaterial surfaces has primarily focused on an overall measurement of protein adsorption, primarily due to limitations in the analysis techniques, especially on topographical surfaces, while the precise effects of surface properties on protein structure at molecular scale is largely unknown. It is of fundamental interest to the biomaterials community to understand how molecular protein structure and activity are correlated to specific surface properties. This may call for new measurement techniques and analysis methods to measure the individual protein adsorption and its functional activity on surfaces,

and to understand the role of surface property in protein adsorption and subsequent biological responses.

The formation of a thrombus on the surface of a blood-contacting material in vivo involves both the activation of the intrinsic coagulation cascade and the adhesion/activation of platelets. Platelet adhesion and contact activation are generally studied separately. However, platelet activation can trigger FXII-mediated contact activation on the surface, leading to generation of FXIIa-antithrombin and FXIa-antithrombin complex, and contributing to clot formation²²⁶. It is also reported that contact activation and platelet adhesion have a strong synergistic effect on coagulation on blood-contacting materials even though these events in isolation are not sufficient to induce substantial thrombus formation²²⁷. These results may motivate the study of contact activation and blood coagulation on biomaterial surfaces with incorporation of platelet, proteins, and surfaces. The systematic investigation of biological responses to surfaces will be the great beneficial for biomaterial designs with improved biocompatibility.

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Highlights

We highlight the current knowledge in the field of blood-materials interactions

We highlight the role of surface chemistry, energy and topography in mediating blood-material interactions

We highlight novel methods for measuring biological responses

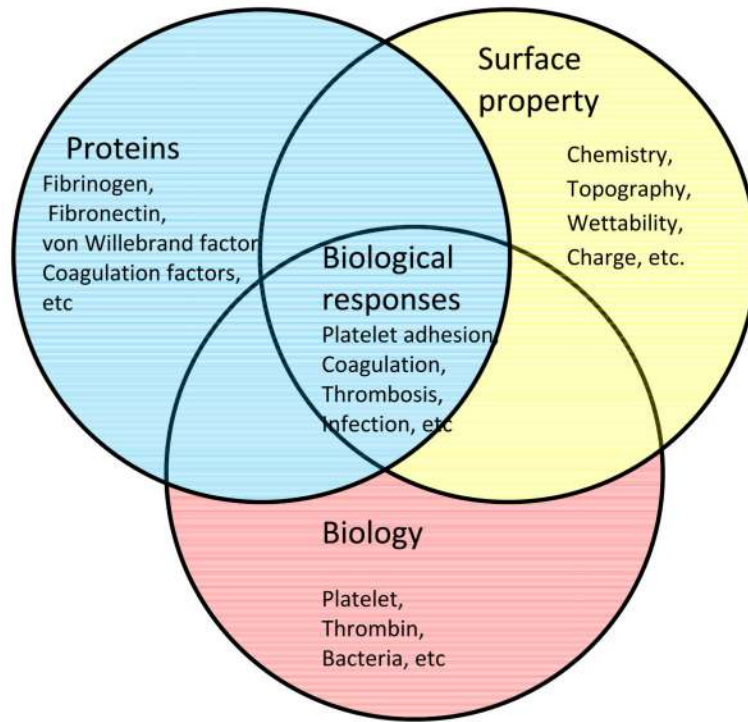


Figure 1. The interaction between biomaterials and biological entities at the interface is influenced by surface property and proteins.

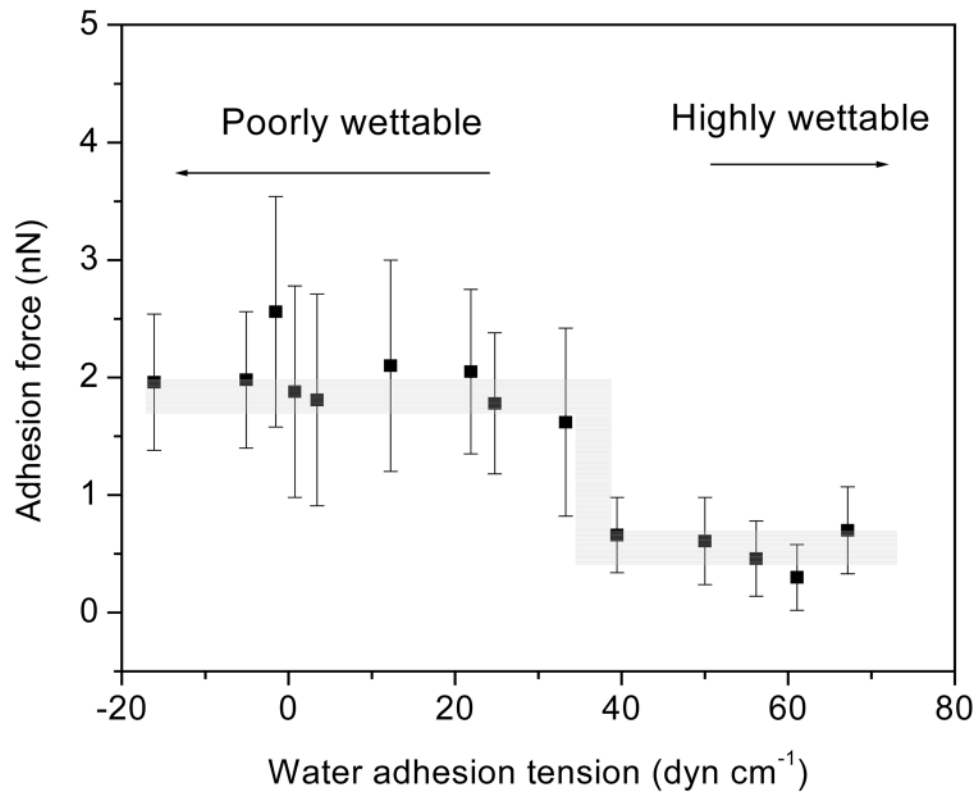


Figure 2. Average adhesion forces of human fibrinogen coated AFM tips to LDPE surfaces with different water adhesion tensions. (error bar: standard deviation) (Reprint from Xu and Siedlecki⁶⁸ with permission from Elsevier)

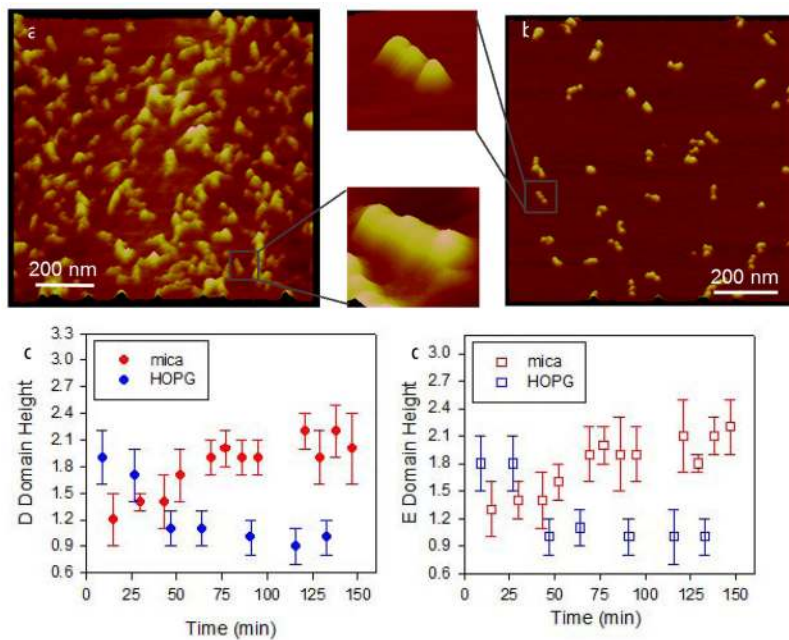


Figure 3. Time dependent spreading of fibrinogen on model substrates, (a) HOPG and (b) muscovite mica. Individual molecules could clearly be observed as seen in the higher magnification images. Spreading of the molecules was followed by measuring heights of the (c) D domains and (d) E domains for ~ 2 hrs following adsorption. (error bar: standard deviation) (Reproduced from Agnihotri and Siedlecki¹³³ with permission from ACS)

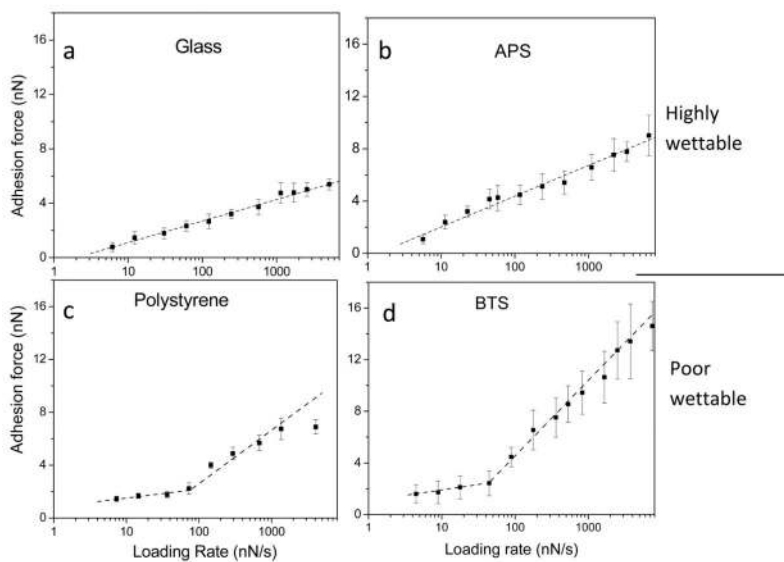


Figure 4. Adhesion forces of fibrinogen and colloid surfaces under different loading rates. High wettable surfaces: (a) glass and (b) 3-aminopropyltrichlorosilane (APS), and poorly wettable surfaces: (c) polystyrene and (d) n-butyltrichlorosilane (BTS). Multiple energy barriers are present between fibrinogen and hydrophobic surfaces while a single energy barrier is present between fibrinogen and hydrophilic surfaces. (error bar: standard deviation) (Reprint from Xu and Siedlecki¹³⁹ with permission from ACS)

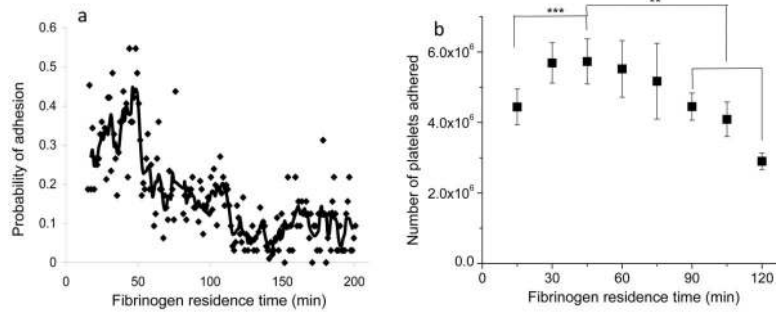


Figure 5.

(a) Probability of interaction between a mAb against γ chain 392–411 and its antigen on fibrinogen measured by AFM. The solid line, the average of 5 points, is included for visualization purposes. (b) Platelet adhesion data from multiple experiments ($n \geq 6$ for each time point) showing changes in platelet adhesion as a function of fibrinogen residence time on mica substrates. (**: $p < 0.01$, ***: $p < 0.001$, error bar: standard deviation) (Reprint from Soman, Rice and Siedlecki¹⁴⁸ with permission from ACS)

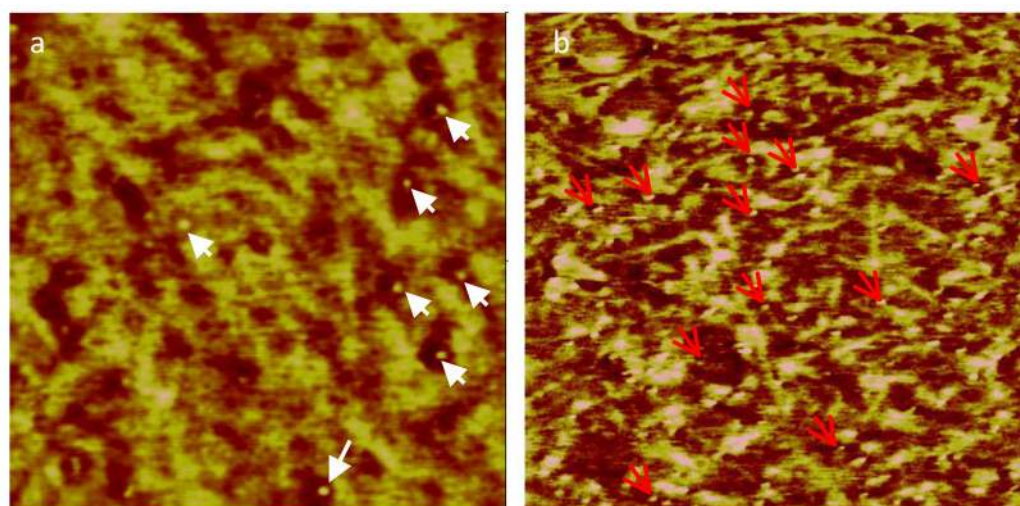


Figure 6. AFM images of nanogold conjugated BSA adsorbed on PUU surface imaged under (a) ambient condition and (b) in PBS buffer. Gold beads indicated by arrows are seen in phase image where lighter color represents hard domains. Quantification reveals more than 2 fold increase in number of labeled proteins per unit area on the soft segment regions compared to the hard segment. (Scan size: $500 \times 500 \text{ nm}^2$, reprint from Xu and Siedlecki¹⁶⁴ with permission from Wiley)

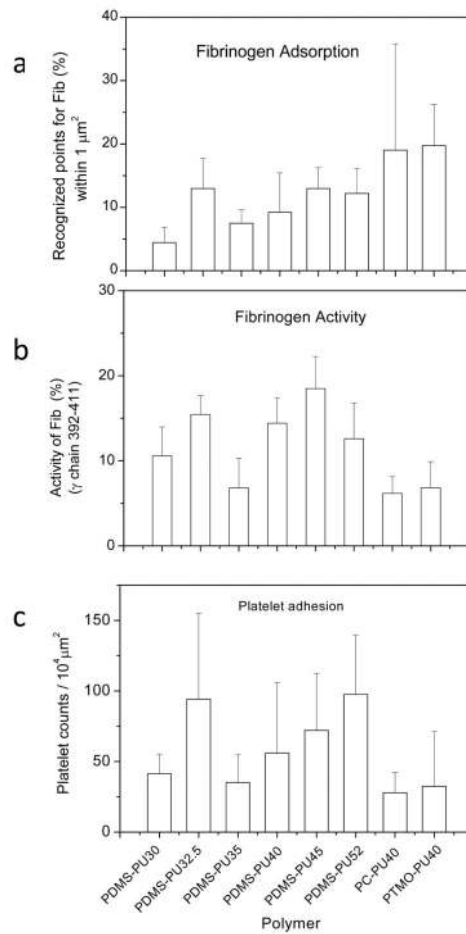


Figure 7.

(a) Fibrinogen adsorption on PUs as recognized by a polyclonal antibody; (b) Functional activity of adsorbed fibrinogen detected by a monoclonal antibody against the fibrinogen γ chain 392–411; (c) platelets adhesion on PU surfaces. Note: The number following the type of PU is the percentage hard segment content. (mean \pm standard deviation) (Reprint with permission from ¹⁶⁷)

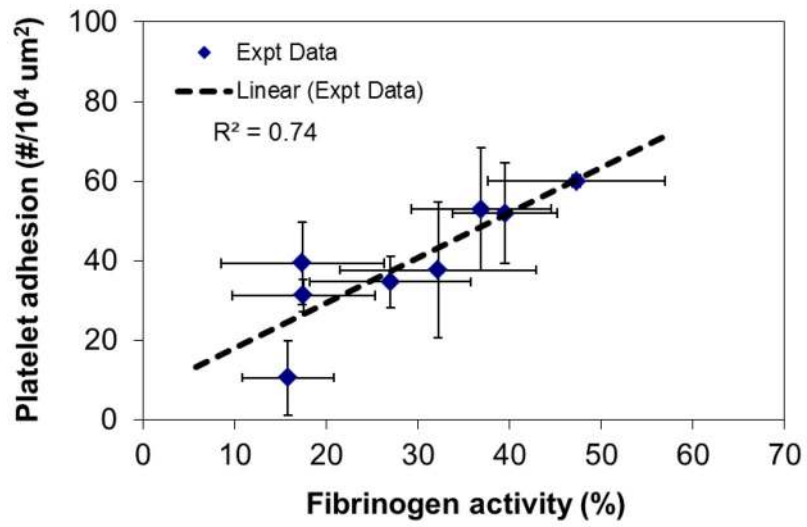


Figure 8. Relationship of platelet adhesion and fibrinogen bioactivity on polyurethane surfaces. (mean \pm standard deviation)

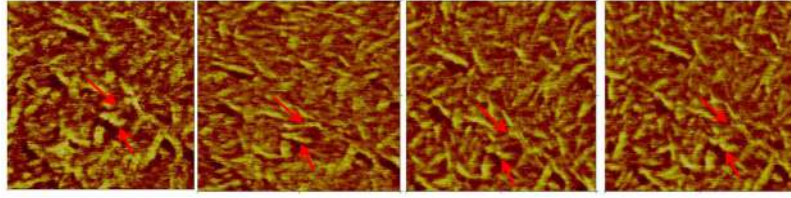


Figure 9. Sequential phase images of PUU under PBS buffer hydrated for 21 hrs, showing hard domain reorientation and reorganization due to hydration. The images were captured at approximately 9 min intervals and high phase angle shifts (lighter colors) represent the “harder” regions. The lines point to landmarks that can be used for reference in monitoring the dynamic changes.

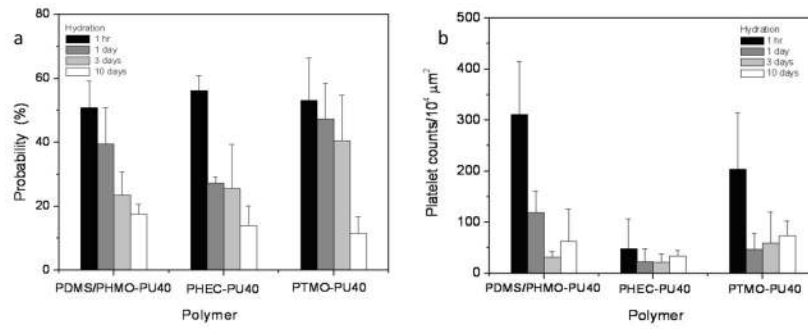


Figure 10.

(a) Probability of recognition by mAb shows the functional activity of fibrinogen on polyurethane surfaces following hydration for 1 hr, 1 day, 3 days and 10 days, (b) platelet adhesion on polyurethane surfaces following hydration with same duration. (reprint from reference¹⁷¹). (error bar: standard deviation) (reprint from Xu, Runt, and Siedlecki¹⁷¹ with permission from Elsevier)

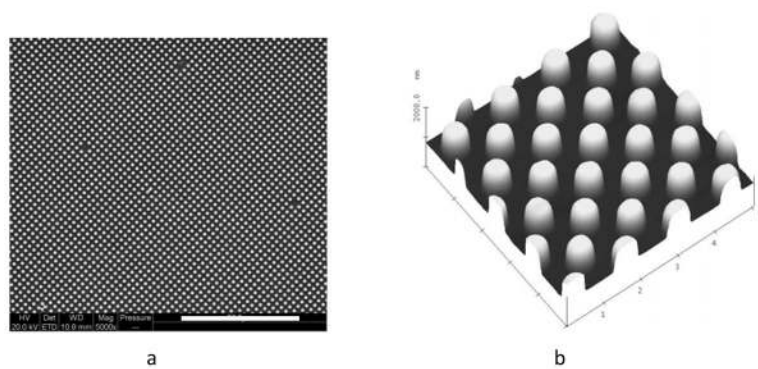


Figure 11. (a) SEM and (b) AFM images showing topography of textured PUU film surface with 500/500 nm pattern. (Reprint from Xu and Siedlecki¹⁸³ with permission from Elsevier)

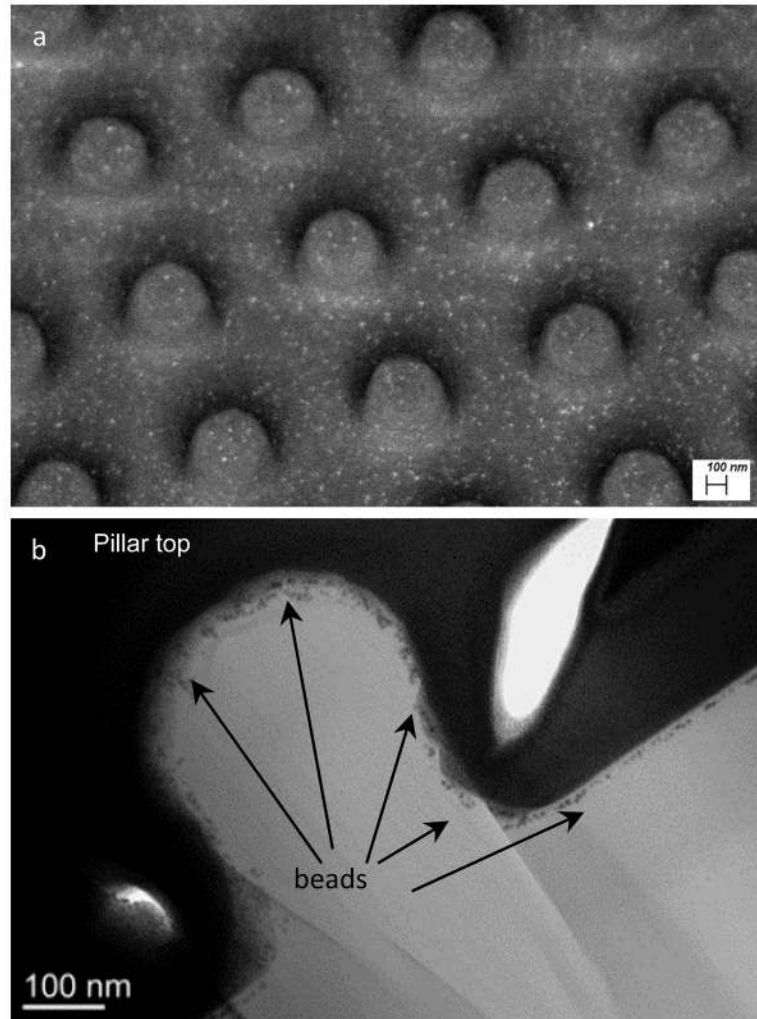


Figure 12. (a) FESEM with 30 degree angle and (b) TEM images of 500/500 nm patterned PUU surfaces adsorbed with nano-gold conjugated BSA, showing the distribution of protein adsorbed on textured pillar surface.

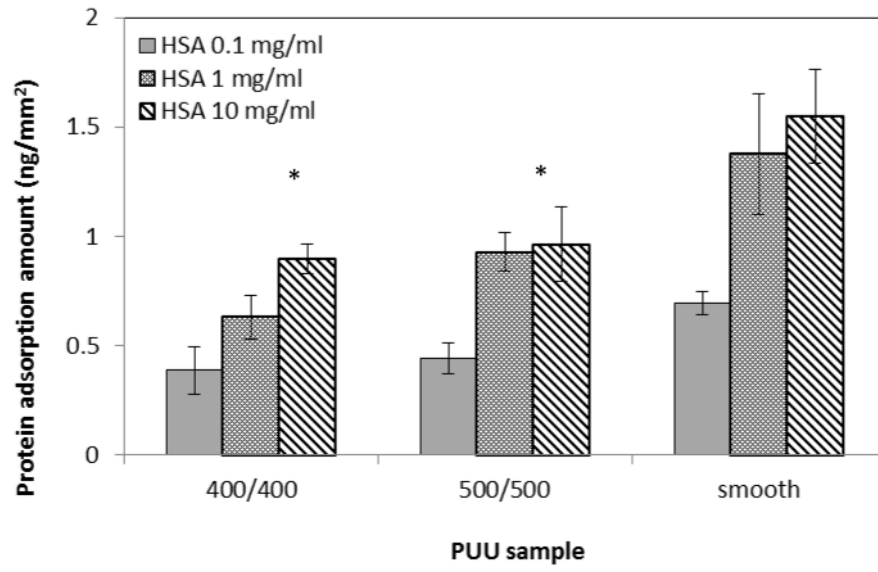


Figure 13. Human albumin adsorption on textured PUU surfaces at different concentrations. (*: $p < 0.05$, error bar: standard deviation)

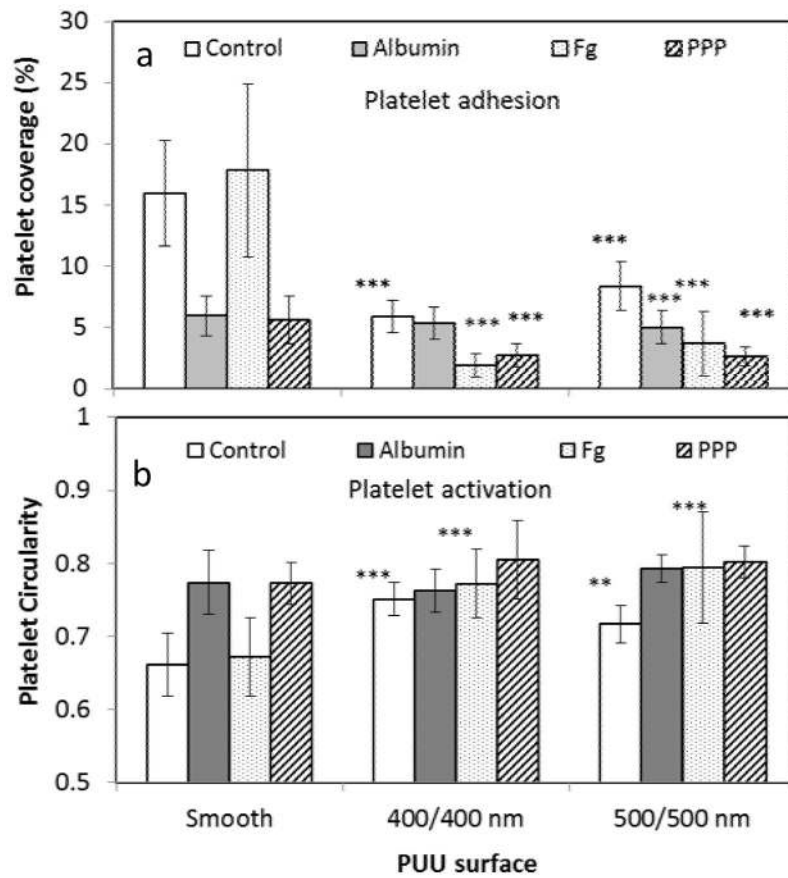


Figure 14. (a) Platelet adhesion and (b) activation on PUU surfaces with pre-adsorbed with plasma proteins. The statistical analysis is performed between textured and smooth PUU samples. (**: $p < 0.01$; ***: $p < 0.001$, error bar: standard deviation)

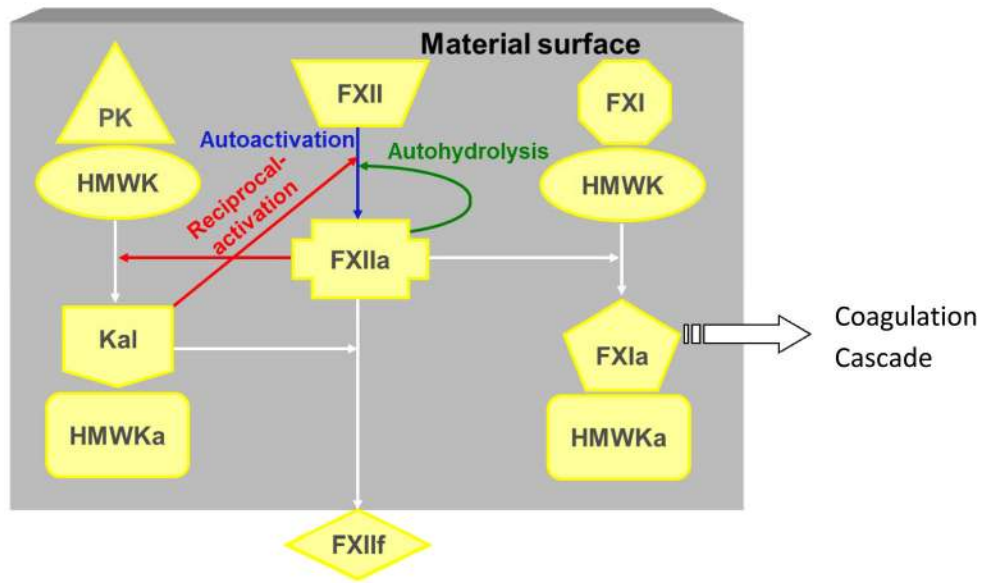


Figure 15. Surface-mediated interactions in contact activation of plasma coagulation. PK: prekallikrein; HMWK: high molecular weight kinnogen; Kal: kallikrein; FXII: factor XII; FXI: Factor XI. Suffixes “a” and “f” represent activated and fragmented form. (Reprint from Vogler and Siedlecki ¹⁹⁸ with permission from Elsevier)

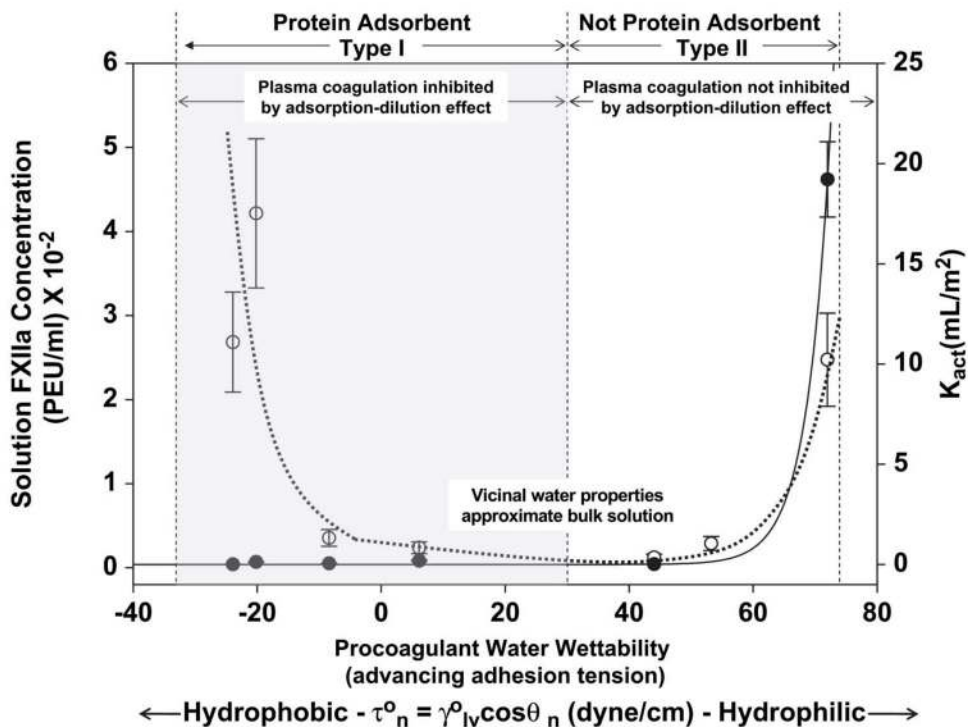


Figure 16. Comparison of FXII activation in neat-buffer solution (left-hand ordinate, dashed line) to catalytic potential in plasma K_{act} (right-hand ordinate, solid line) for activator (procoagulant) particles exhibiting different surface energy (abscissa) expressed as water adhesion tension $\tau_a^o = \gamma_{lv}^o \cos \theta$ in dyne/cm (where γ_{lv}^o is water interfacial tension in dyne/cm and θ is the advancing contact angle) (reprint from ²¹³ with permission from Elsevier).

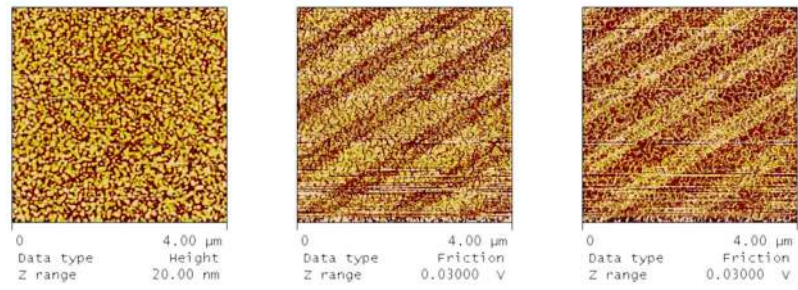


Figure 17.

From left to right – height, friction mode retrace, and friction mode trace AFM images of patterned thiol surfaces. Bare gold surface was stamped with a linear pattern PDMS stamp inked with octadecanethiol and backfilled with 11-mercaptopundecanoic acid. The light colored bands in the friction retrace image (middle) correspond to the stamped octadecanethiol.

Table 1

Contact angle and adhesive force values for protein modified AFM probes

Surface	OTS	APTES	Glass
Advancing Water Contact Angle (°)	110	59	<10
Standard Si ₃ N ₄ AFM Probe	6.5 ± 4.8	1.3 ± 1.1	0.8 ± 0.2
Adhesion force (nN) (mean ± std. dev.)			
FXII-Modified AFM Probe	4.4 ± 2.1	2.0 ± 0.9	0.5 ± 0.6
FXIIa-Modified AFM Probe	3.4 ± 1.1	0.5 ± 0.4	0.5 ± 0.4
Albumin-Modified AFM Probe	5.6 ± 4.2	1.1 ± 0.7	0.5 ± 0.4

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Table 2

ANOVA analysis of adhesion forces for fibrinogen and LDPE surfaces with different water adhesion tensions (τ , dyn.cm⁻¹) and water contact angle (θ , degree). $\tau = \gamma_{LV} \cos \theta$ (Reprint from Xu and Siedlecki⁶⁸ with permission from Elsevier).

		← Highly wettable					τ	Poorly wettable →					
τ	θ	67.2	61.1	56.2	50.0	39.4	33.3	24.8	21.9	12.3	3.43	0.76	-16.1
		22.7	33.0	39.5	46.6	57.2	62.8	70.1	72.5	80.3	87.3	89.4	102.8
67.2	22.7		***	NS	NS	NS	***	***	***	***	***	***	***
61.1	33.0			NS	***	***	***	***	***	***	***	***	***
56.2	39.5				NS	***	***	**	***	***	**	***	***
50.0	46.6					NS	***	***	***	***	***	***	***
39.4	57.2						***	***	***	***	**	***	***
33.3	62.8							NS	***	***	NS	NS	NS
24.8	70.1								NS	NS	NS	NS	NS
21.9	72.5									NS	NS	NS	NS
12.3	80.3										***	NS	NS
3.43	87.3											NS	NS
0.76	89.4												NS
-16.1	102.8												

NS : Not significant
 *** : significant p <0.001
 ** : significant p <0.01
 * : significant p <0.05

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Parameters for dynamic model of fibrinogen interaction with surface. (Reprint from Xu and Siedlecki¹³⁹ with permission from ACS)

Table 3

Wettability	Surface	Water contact angle (°)	Loading rate (nN/s)	$\chi\beta$	k_{off} (s ⁻¹)	τ (s)
Highly wettable	Glass	11.3±1.7	6–5100	0.06	2.92	0.34
	Polystyrene (plasma treated)	22.3±1.6	6–3100	0.02	1.99	0.50
	APS	57.2±1.7	5–6500	0.04	1.35	0.74
Poorly wettable	Polystyrene	82.2±1.5	6–100 100–4100	0.13 0.04	0.23 4.98	4.35 0.20
	BTS	90.4±1.2	4–100 100–7400	0.11 0.02	0.19 5.50	5.12 0.18
	OTS	103.0±1.9	5–100 100–3200	0.07 0.01	0.42 13.8	2.38 0.07