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## Functional impact of oxidative post-translational modifications on fibrinogen and fibrin clots

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### Abstract

Fibrinogen is a circulating multifunctional plasma protein vital for hemostasis. Activation of the coagulation cascade converts soluble fibrinogen to insoluble polymerized fibrin, which, along with platelets, forms the hemostatic clot. However, inappropriate formation of fibrin clots may result in arterial and venous thrombotic disorders that may progress to life-threatening adverse events. Often thrombotic disorders are associated with inflammation and the production of oxidants. Fibrinogen represents a potential target for oxidants and several oxidative post-translational modifications that influence fibrinogen structure and function have been associated with disease pathogenesis. Here, we review various oxidative modifications of fibrinogen and the consequences of these modifications on protein structure, ability to form fibrin and the resulting alterations on fibrin architecture, viscoelastic and biochemical properties that may contribute to disease.

### Keywords

fibrinogen; fibrin; coagulation; thrombosis; oxidation; nitrotyrosine

## Introduction to fibrinogen and fibrin structure and biochemistry

Fibrinogen is synthesized in hepatocytes and secreted into the blood [1] where it circulates with a half-life of about 3 days [2]. Fibrinogen is a 340 kDa hexamer, composed of two pairs of three non-identical chains termed A $\alpha$ , B $\beta$ , and  $\gamma$ . Specific sites in each of these chains are more subject to oxidative modification (Figure 1) as elaborated below. Together the chains comprise a symmetrical molecule composed of one globular E region flanked on each side by globular D regions [3–6] that are connected by three-stranded  $\alpha$ -helical coiled-coils [4–6]. The E region, which is composed of all three chains, contains fibrinopeptides A (FpA) and B (FpB) [4–6]. Cleavage of these peptides by thrombin exposes knobs 'A' and 'B', resulting in the formation of fibrin monomers [7–11]. Positively charged 'A' and 'B' knobs have complementary negatively charged binding sites within the  $\gamma$  and  $\beta$  nodules in the D

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regions of adjoining monomers termed holes 'a' and 'b' [12, 13]. Knob-hole associations each result in the formation of half-staggered, double-stranded protofibrils [14, 15], which associate laterally to form fibrin fibers [14, 16], and ultimately the branching network structure of the hemostatic or thrombotic clot [17]. The formation of fibrin clots using either isolated fibrinogen or platelet poor plasma (PPP) after the addition of thrombin and  $\text{Ca}^{2+}$  can be monitored spectrophotometrically by following turbidity changes over time (Figure 2A). This assay yields some generalized structural and kinetic information about fibrin clot formation, but does not provide specifics on properties such as rate of FpA and FpB release, which requires HPLC methodology [18], or fibrin clot structure, which can be obtained by scanning electron microscopy (Figure 2B) [19] or confocal microscopy [20].

Fibrin is covalently crosslinked by the transglutaminase Factor XIII (FXIII), which provides stability and elasticity to the clot [20–23]. FXIII is activated (FXIIIa) by thrombin through cleavage of the N-terminal activation peptide [24]. FXIIIa crosslinks fibrin via the formation of  $\epsilon$ -( $\gamma$ -glutamyl) lysine bonds within and between  $\alpha$  and  $\gamma$  chains, creating  $\gamma$ -dimers,  $\alpha$ -polymers, and  $\alpha$ - $\gamma$ -heteropolymers [25, 26]. The rate and extent of FXIIIa crosslinking of fibrin, made by addition of thrombin plus FXIII and  $\text{Ca}^{2+}$  to either isolated fibrinogen or PPP, can be determined by quenching the reaction and running the products on an SDS PAGE gel with crosslinked fibrin products identified and quantified by western blot analysis and densitometry (Figure 2C).

Physiological lysis of the fibrin clot proceeds through the enzyme plasmin, which is converted from its inactive zymogen precursor, plasminogen, by the enzyme, tissue plasminogen activator (tPA) [27]. This reaction requires fibrin as a cofactor [28] and the kinetics of lysis can be followed by changes in turbidity (Figure 2A). Plasmin binds and cleaves fibrin at several high-affinity lysine residues, one of which is located in the flexible region of the carboxy termini of the A $\alpha$  chains termed  $\alpha$ C regions [29, 30] subsequently exposing additional lysine residues that plasmin further cleaves into smaller fragments [29]. Endogenous inhibitors of fibrinolysis target both tPA and plasmin, and include plasminogen activator inhibitor-1 (PAI-1),  $\alpha_2$ -antiplasmin ( $\alpha_2$ -PI), and thrombin-activatable fibrinolysis inhibitor (TAFI).

Fibrin is a highly extensible, viscoelastic polymer [22], allowing it to deform appropriately in response to shear stress. Changes in clot structure due to post-translational modifications (PTMs), elevated fibrinogen concentration, or interactions with other proteins have been shown to alter the viscoelastic properties of fibrin clots. Specifically, increased clot stiffness was significantly associated with coronary artery disease [31], as well as those at risk for thrombotic events such as smokers [32] and diabetics [33]. Traditionally, rheometry has been used to measure the elastic (stiffness) and inelastic (viscosity) properties of fibrin clots *in vitro*, although some studies have also utilized thromboelastography. Defined by the storage modulus ( $G'$ ) and loss modulus ( $G''$ ), respectively, these properties dictate the ability of the clot to store or dissipate energy due to the application of applied shear stress. The ratio of the loss modulus to the storage modulus,  $\tan \delta$ , is also used as a measure of the relative proportion of inelastic to elastic components. The relationships between viscoelastic properties, fibrin clot structure, and pathology remain unclear. Most studies have observed that increased fiber density or decreased porosity, coupled with decreased individual fiber diameter results in stiffer clots with delayed lysis, fostering a prothrombotic state [20, 31, 33]. In other cases, clot structural heterogeneities, including fiber clusters or bundles, have been observed as a consequence of PTMs and have varying effects on viscoelastic properties [32].

In addition to forming the fibrous network structure of a thrombus, fibrinogen also plays an important role in platelet aggregation. Individual platelets bind residues (400–411) located

within the carboxy termini of the  $\gamma$  chains of fibrin(ogen) via their integrin receptor,  $\alpha$ IIb $\beta$ 3 [34]. These linkages tether platelets together resulting in platelet aggregation.

Fibrinogen is a critical protein in the formation of the clot, both in the fibrin network and platelet aggregation, which are ultimately required for the generation of the hemostatic thrombus. Perturbations in these functions may influence the formation and properties of the fibrin network and promote pathological states, including thrombosis and thromboembolism. Indeed, epidemiological studies have indicated that increased levels of circulating fibrinogen are an independent predictor of coronary heart disease and in some cases, of premature death from cardiovascular disease [35–38]. Despite these sound correlations, a causative association between high levels of fibrinogen and cardiovascular disease has not been firmly established. Evidence suggests that oxidative post-translational modifications may contribute to alterations in fibrinogen function and affect pathology. Here we review the impact of oxidative modifications on the structure and function of fibrinogen and discuss their relationship with various disease states.

## Oxidative modifications of fibrinogen

Oxidative stress has been widely implicated in physiological processes such as aging, in disease pathogenesis ranging from carcinogenesis to atherogenesis, and the etiology of arterial and venous thrombosis. Proteins are major targets for oxidants and fibrinogen, which comprises a large percentage of plasma proteins (~4 %) is a likely target for oxidative post-translational modifications.

### Fibrinogen oxidation

Fibrinogen was identified as the primary oxidized protein in the plasma of smokers, an at-risk group for lung cancer and thrombosis, and levels of oxidized fibrinogen were significantly increased in smokers as compared with controls [32, 39]. In thromboelastogram studies, clots made from whole blood collected from smokers shortly after smoking 2 cigarettes demonstrated decreased R value (lag phase), increased  $\alpha$  angle (rate of fibrin polymerization), and increased G (stiffness) as compared with non-smokers and smokers before cigarette use [40]. In addition, clots formed from PPP of smokers post-cigarette use demonstrated increased fiber density and decreased fiber diameter compared with the control groups. Elevated plasma fibrinogen levels increased the risk for Alzheimer's disease [41], increased brain atrophy in Alzheimer's disease patients [42], and were associated with cognitive decline [43]. Previous studies have shown that oxidized proteins localize to sites of neurodegeneration [44]. Oxidation of the fibrinogen  $\gamma$  chain precursor protein quantified as carbonylation was increased in the plasma of subjects with Alzheimer's disease [45]. Together these data suggest that inflammation and the production of oxidants may play overlapping roles in thrombosis and neurodegenerative diseases.

The potential effects of oxidation on fibrinogen and fibrin function have been explored in several studies that exposed fibrinogen to oxidants *in vitro*. Early studies using hematoporphyrins, photosensitive dyes, and chloramines to oxidize fibrinogen reported a dose-dependent inhibition of thrombin-induced fibrin polymerization [46, 47]. Photo-oxidized fibrin monomer did not bind fibrinogen as well as nascent fibrin monomer despite similar kinetics and amount of FpA and FpB released between oxidized and nascent fibrinogen [47]. Oxidation of fibrinogen by exposure to methylene blue photosensitive dye and light impaired polymerization of des A fibrin and rendered des AB NDSK (N-terminal disulphide knot (NDSK) region of fibrinogen lacking both FpA and FpB) incapable of binding to fibrinogen, with no effect on binding to D-dimer [48].

These results suggest that knob 'A'-hole 'a' interactions are affected by oxidation. His<sup>16</sup> in the  $\beta$  chain of fibrinogen was found to be modified in these oxidation studies, leading these authors to conclude that this amino acid was part of knob 'A' (Figure 1; Table 1) [48]. However, we now recognize [49] that His<sup>16</sup> is part of knob 'B', suggesting that photo-oxidation of fibrinogen may also modify other previously unidentified sites, specifically those that affect A:a interactions. Oxidized fibrinogen also affects lysis. Pro-urokinase was activated faster in the presence of oxidized fibrin, and plasmin lysed oxidized fibrin at a faster rate implying a pro-fibrinolytic function for oxidized fibrinogen [50].

Metal catalyzed oxidation of fibrinogen results in the formation of dityrosine crosslinks [51] and a dose-dependent increase in protein carbonyl levels [51, 52], which may affect several amino acid residues such as histidine, proline, arginine, and lysine [53]. Oxidation of fibrinogen with iron-ascorbate delayed release of FpA and FpB, concomitant with delayed fibrin polymerization and a lower maximum absorbance during turbidity assays [52]. Western blot analysis showed that the A $\alpha$  and B $\beta$  chains are susceptible to carbonylation following metal-catalyzed oxidation, while the  $\gamma$  chain remained unaffected [54]. ADP-induced platelet aggregation and adhesion were diminished with oxidized compared with nascent fibrinogen [51]. However, another study found that iron-ascorbate oxidation of fibrinogen increased platelet aggregation [55]. This study also demonstrated decreased catalytic efficiency of tPA conversion of plasminogen to plasmin, which requires fibrin as a cofactor. The authors proposed that decreased fibrinogen  $\alpha$ -helical content caused by carbonylation of lysine residues may explain these findings. Moreover, aspirin-induced lysine acetylation prevented the effects of oxidation supporting a role for fibrinogen lysine residues in platelet aggregation. Both studies utilized similar oxidation systems and observed similar carbonyl levels per mg of protein. However, the shorter exposure time (0–1 hr [55] vs. 0–20 hr [51]) may have accounted for these differences. Although further experimentation is required, it is reasonable to assume that the number of amino acid residues, which amino acids, and the magnitude of oxidation per residue are significantly less during short exposures. Moreover, formation of reactive carbonyls indicated only one type of oxidative modification and it is possible that other oxidative modifications during prolonged exposures may account for the observed differences. These findings reinforce a common deficit in studies of protein oxidation that requires attention. The correlation between functional outcomes and protein oxidation should include precise identification of the modified amino acids, the chemical nature of the modification, as well as the extent of modification.

Fibrinogen oxidation with hypochlorite (HOCl) decreased the rate of fibrin polymerization and maximum absorbance by turbidity assays [56]. Treatment with hypochlorite dose dependently delayed lysis time, which may be due to the observed increase in fiber density, decrease in fiber diameter, and decrease in pore size, but not changes in plasminogen activation. Treatment of fibrinogen with hypochlorite also weakened clot stiffness, i.e. decreased  $G'$  and  $G''$ . The maximum strain and maximum modulus in strain hardening experiments were lower for oxidized compared with nascent fibrin clots. The authors identified three oxidized methionine residues, Met<sup>476</sup>, Met<sup>367</sup>, and Met<sup>78</sup> located on the A $\alpha$ , B $\beta$ , and  $\gamma$  chains, respectively (Figure 1, Table 1). Quantification of these sites identified Met<sup>476</sup> as the most prominent modification, with 150  $\mu$ M hypochlorite producing 73 % oxidation of the Met<sup>476</sup> residue. Met<sup>476</sup> is located within the  $\alpha$ C region of the A $\alpha$  chain and is thought to contribute to lateral aggregation of fibrin fibers [57]. Impairment of lateral aggregation by oxidation of Met<sup>476</sup> may result in clots with thinner fibers and a higher fiber density, ultimately reducing fibrinolysis and weaker clot structure (Table 1).

## Fibrinogen nitration

Protein 3-nitrotyrosine has been identified as both a marker and a functional post-translational modification of proteins during disease processes. Nitrated fibrinogen, (primarily tyrosine nitration within the  $\beta$  chain of the molecule) was initially identified in the plasma of subjects with acute respiratory distress syndrome [58]. Nitrated fibrinogen has been detected in the plasma of subjects with lung cancer [39] and with end stage renal disease [59]. Elevated levels of nitrated fibrinogen have been quantified in coronary artery disease subjects as compared with healthy controls [60]. Increased levels of plasma nitrated fibrinogen were also documented in subjects with venous thromboembolism (VTE) subjects compared with non-VTE subjects [61]. When compared with the lowest quartile, subjects in the highest quartile of nitrated fibrinogen had a significantly increased risk of VTE [61]. In previous studies aiming to identify biomarkers of VTE, MPO was significantly elevated in PE subjects, suggesting that nitrative and oxidative intermediates are produced during VTE [62]. Together these studies may indicate that nitrated fibrinogen is a potential diagnostic biomarker for VTE. Overall, increased circulating levels of nitrated fibrinogen have been associated with arterial [60] and venous thrombotic diseases [61], as well as those at risk for thrombosis, such as smokers [32]. Neutrophils and monocytes, which are capable of producing nitrating intermediates [63], are key mediators in the initial phases of venous thrombus formation [64], and the progression of the thrombus during atherosclerosis [65, 66]. Therefore appreciating the potential consequences of fibrinogen nitration will be of interest and could potentially provide mechanistic insights that link oxidative modifications and risk for thrombosis.

The functional effects of fibrinogen nitration have been explored both *in vitro*, [60] in smokers, [32] and in healthy subjects exposed to low levels of lipopolysaccharide [67]. When fibrinogen was nitrated *in vitro* with either 3-morpholinopyridone (SIN-1), which produces nitric oxide and superoxide that react to form peroxynitrite, or MPO, hydrogen peroxide, and nitrite, a decrease in lag time and an increase in both the rate of fibrin polymerization and final absorbance was observed during turbidity assays when compared with unmodified fibrinogen [60]. In contrast, fibrinogen treated with MPO and hydrogen peroxide alone (oxidizing conditions), impaired fibrin polymerization, consistent with the aforementioned exposure to HOCl [56]. The clot structure formed from nitrated fibrinogen was composed of thick, twisted fiber bundles and large pores, whereas fibrinogen treated with MPO and hydrogen peroxide generated a homogenous structure composed of thin fibers. Even though nitration and oxidation of fibrinogen produce disparate effects on clot structure,  $G'$  was similar between the two groups and significantly lower than control, indicating a decrease in clot stiffness. Although no differences in lysis rate or lysis product formed were seen between control and nitrated fibrinogen *in vitro*, injection of microemboli composed of fibrin treated with SIN-1 into mice followed by bolus tPA injection showed enhanced lytic susceptibility compared to emboli of nascent fibrin. No differences were observed between nitrated and control fibrinogen for FpA or FpB release, FXIII crosslinking, or platelet aggregation. Expanding these findings, Parastatidis et al. [68] generated isolated fibrinogen from subjects with coronary artery disease that were depleted of nitrated fibrinogen molecules by the use of immunoaffinity removal procedures. As a control, identical plasma samples that retained the nitrated molecules were also generated by immunodepletion with a nonspecific immunoglobulin. Thrombin-induced fibrin clot formation showed a significant increase in the rate of fibrin polymerization and maximum absorbance in turbidity assays in fibrinogen that retained the nitrated fibrinogen molecules as compared to those where nitrated fibrinogen molecules were removed.

Smoking is a risk factor for thrombosis [69], and has been associated with increased production of oxidants [32, 70]. In smokers, elevated nitrated fibrinogen levels correlated



with a dose-dependent increase in the rate of fibrin polymerization that was reversed when nitrated molecules were removed via immunoprecipitation with anti-nitrotyrosine antibodies [32]. Scanning electron micrographs showed fiber clustering that was also reversed by immunodepletion. Nitrated fibrinogen levels were positively correlated with  $G'$ ,  $G''$ ,  $\tan \delta$ , and inversely correlated with fibrinolysis rate. Tyrosine residues Tyr<sup>292</sup> and Tyr<sup>422</sup> were identified by mass spectrometry as the primary nitrated residues in a majority of the smoker samples analyzed. These residues are located within the carboxy terminus of the B $\beta$  chain of fibrinogen near hole 'B', which may be involved in lateral aggregation (Figure 1; Table 1). Use of the 'B' knob mimetic peptide accelerated lateral aggregation in all samples, but was positively correlated with nitrated fibrinogen levels. The effects of nitration *in vivo* in smokers show some similarities to the *in vitro* modified fibrinogen, like increased rate of fibrin polymerization, as well as some disparities. These differences might be related to the tyrosines susceptible to nitration in each system. Additional studies to identify nitrated tyrosines within fibrinogen *in vitro* would be necessary to explore this hypothesis.

The relationship between inflammation and fibrinogen nitration was explored by administering low levels of lipopolysaccharide to healthy human subjects, which resulted in increased production of several protein mediators of inflammation such as TGF- $\beta$ , C-reactive protein, and myeloperoxidase (MPO) [67]. Nitrated fibrinogen was also increased in this challenge model, peaking at 72-hrs post-injection, much later than the other inflammatory mediators. In addition, nitration of fibrinogen post-lipopolysaccharide injection resulted in increased rate of clot formation. This study implies that nitration results as a direct consequence of inflammatory challenge, nitrating intermediates are produced for prolonged periods of time following insult, and tyrosine nitration affects fibrinogen function, providing evidence for the role of inflammation and protein nitration in thrombotic disorders.

Although the aforementioned studies demonstrated that nitrated fibrinogen accelerated clot formation kinetics, some studies have reported opposing effects [71, 72]. Peroxynitrite treatment of fibrinogen dose-dependently decreased the rate of fibrin polymerization, increased lag time, and decreased maximum absorbance during clot polymerization assays [71, 72]. Additionally, platelet aggregation and adhesion were dose-dependently impaired by peroxynitrite treatment of fibrinogen compared to nascent fibrinogen [73]. However, plasmin degradation of fibrin clots was slower in treated fibrinogen, which is consistent with previous nitrated fibrinogen studies [32]. Peroxynitrite is an oxidant and a nitrating agent capable of oxidizing cysteine and tryptophan residues and producing dityrosine crosslinks and carbonyl-modified residues [74, 75]. Previous studies have reported that oxidized fibrinogen that was subsequently nitrated resulted in a reversal of the impaired fibrin polymerization, producing polymerization rates similar to nitration alone [60]. Thus, the effects of peroxynitrite on clotting kinetics may be due to the effects of other oxidative modifications, rather than solely nitration [47, 60]. These modifications may occur in fibrin monomer association sites [48], or within the fibrinogen integrin  $\alpha$ IIb $\beta$ 3 binding sites located within the  $\gamma$  chain [34]. Again, these studies indicate that *in vitro* modeling of the functional effects of fibrinogen oxidative modifications need to carefully consider the sites, magnitude and type of modification accounting for the presence of other off-target modifications.

### Cysteine modifications of fibrinogen

All 58 cysteine residues in fibrinogen are disulfide bonded [76]. However, several disulfide bonds, including  $\gamma$  Cys<sup>23</sup>-A $\alpha$  Cys<sup>45</sup>, A $\alpha$  Cys<sup>442</sup>-A $\alpha$  Cys<sup>472</sup>,  $\gamma$  Cys<sup>326</sup>- $\gamma$  Cys<sup>339</sup> and the symmetrical bonds A $\alpha$  Cys<sup>28</sup>,  $\gamma$  Cys<sup>8</sup>, and  $\gamma$  Cys<sup>9</sup> were capable of being reduced *in vitro* by thioredoxin and dithiothreitol [77, 78]. Reduction by thioredoxin or dithiothreitol resulted in

prolonged clotting time in the absence of effects of FpA release, indicating alterations in fibrin polymer formation [77, 78]. In addition, plasminogen was more activated in the presence of partially reduced fibrinogen compared with non-reduced [79].

The low molecular weight thiol, glutathione (GSH) and its S-nitrosated analogue S-nitrosoglutathione (GSNO) are capable of undergoing disulphide exchange reactions with disulphide-bonded cysteine residues, generating S-glutathionylated cysteine. In addition, GSNO can modify reduced cysteine residues to S-nitrosocysteine. *In vitro* treatment of purified fibrinogen with GSNO, GSH, or oxidized glutathione (GSSG) dose dependently delayed clotting time, reduced the rate of fibrin polymerization, and reduced final turbidity [80]. The authors proposed that the effects of the GSH derivatives were due to S-glutathionylation reactions within the  $\alpha$ C region of fibrinogen, although the S-glutathionylated sites were not identified by sequencing or mass spectrometry [80, 81]. Akhter et al. also reported impaired fibrin polymerization upon exposure of fibrinogen to GSNO. However, CD spectroscopy and tryptophan quenching experiments showed that GSNO did not participate in S-nitrosation or S-glutathionylation reactions, but instead, reversibly interacted with the  $\alpha$ C region of fibrinogen, causing allosteric conformational changes in the protein that affected polymerization [81]. Additional studies have investigated the role of GSNO treatment on fibrin clot structure. Fiber density decreased and fiber diameter increased following GSNO treatment of PPP [82].

However, at high concentrations of GSNO (3.75 mM), fibrin did not polymerize, and instead formed aggregate clusters. Although the authors did not perform turbidity assays, previous studies have found that the formation of clusters and increased fiber diameter both resulted in increased turbidity [32], which would be in contrast to the previously described reports [80, 81]. These conflicting results may be due to GSNO interactions or reactions with other proteins within the plasma, specifically those that play a role in fibrin clot formation, such as thrombin [19], FXIII [19], and FXII [83].

To date, neither S-glutathionylated nor S-nitrosated fibrinogen has been detected *in vivo*. The *in vitro* effects of glutathione analogs on fibrinogen function may suggest that fibrinogen becomes modified or interacts with excess glutathione produced in response to production of oxidants by neutrophils and macrophages during thrombosis or other inflammatory conditions [84]. Future studies focusing on the role of glutathione and fibrinogen in may help clarify the roles of the oxidant/antioxidant systems in thrombosis and inflammation.

## Conclusion

Oxidative modifications of fibrinogen have the potential to influence the kinetics of fibrin formation as well as the structure and biomechanical properties of fibrin, ultimately producing dysfunctional hemostatic clots. Some of these modifications have been identified *in vivo*, providing further evidence for a relationship between the production of oxidants and thrombosis. However, the precise mapping of sites of *in vivo* modifications has not been fully achieved, creating considerable reservations about the significance of these modifications in physiological or pathological settings. Moreover, the *in vitro* exposures do not always reproduce the *in vivo* modifications. As such caution should be exercised in interpreting the functional effects of these modifications on fibrinogen function. An additional question concerning the effects of oxidative modifications on the interactions of fibrinogen with other proteins, such as the platelet integrin receptor  $\alpha$ IIb $\beta$ 3 remains. Future studies should aim to identify the sites of oxidative modifications *in vivo* and, if possible, employ fibrinogen extracted from humans or animal models to determine the effects on fibrinogen function and protein interactions. These studies can then lead towards a greater understanding of the role of fibrinogen modifications in health and disease.

## Abbreviations

<b>FpA/FpB</b>	Fibrinopeptide A/B
<b>PPP</b>	Platelet poor plasma
<b>PTM</b>	Post-translational modification
<b>tPA</b>	Tissue plasminogen activator
<b>VTE</b>	Venous thromboembolism
<b>MPO</b>	Myeloperoxidase
<b>GSH</b>	Glutathione
<b>GSSG</b>	Oxidized glutathione
<b>GSNO</b>	S-nitrosoglutathione

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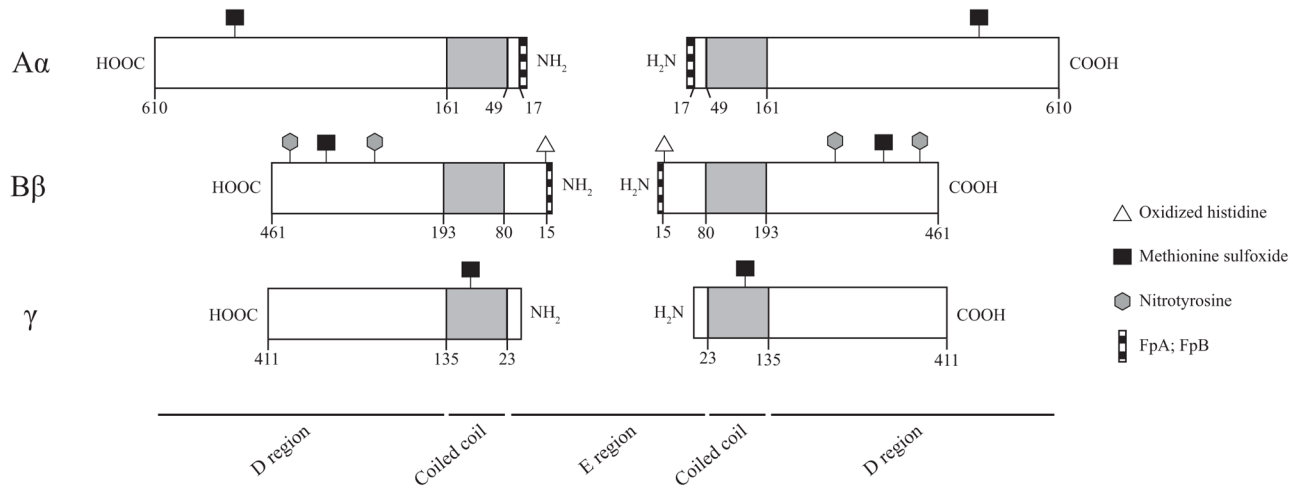


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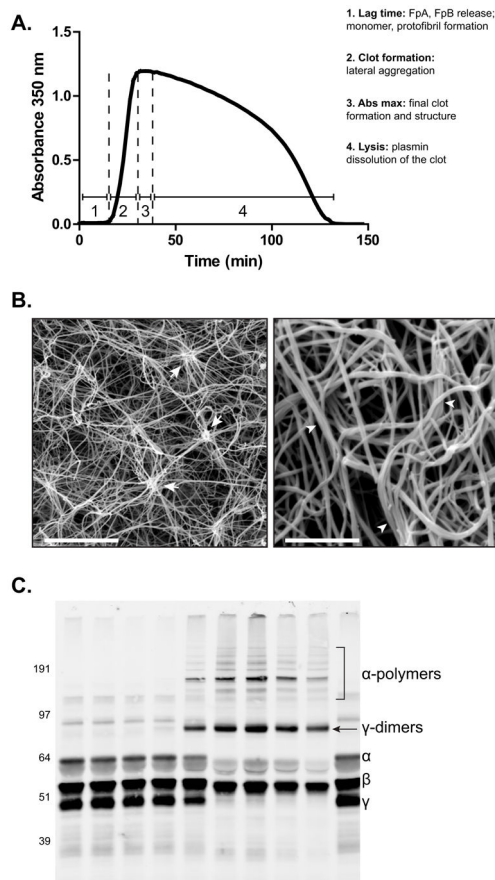
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**Figure 1.**

Sites of oxidative post-translational modifications on fibrinogen. Schematic representation of the A $\alpha$ , B $\beta$ , and  $\gamma$  chains of fibrinogen drawn to scale in length. Residue numbering follows the removal of the signal peptide. FpA and FpB (stripes) are localized towards the N-termini of the A $\alpha$  and B $\beta$  chains, respectively. Coiled-coil domains are represented in grey. His<sup>16</sup> is located within the ‘B’ knob. Met<sup>476</sup>, Met<sup>367</sup>, and Met<sup>78</sup> are localized to the  $\alpha$ C region, the carboxy termini of the B $\beta$  chains, and the coiled-coils of the  $\gamma$  chain, respectively. Both Tyr<sup>292</sup> and Tyr<sup>422</sup> are localized within the carboxy termini of the B $\beta$  chains, however, Tyr<sup>292</sup> is located near the ‘b’ holes.





**Figure 2.**

Techniques for measuring functional properties of fibrinogen. A) Turbidity assay showing both clot formation and lysis over time as a function of changes in absorbance. Thrombin,  $\text{CaCl}_2$ , and either tPA alone or tPA and plasminogen are added simultaneously to PPP or isolated fibrinogen, respectively and clot formation and lysis time are monitored by changes in absorbance at 350 nm as a function of time. 1. The lag phase represents the formation of fibrin monomers, oligomers, and protofibrils. 2. Protofibrils that have reached a sufficient length laterally aggregate to generate fibers, causing a rapid increase in absorbance. 3. Eventually the clot is fully formed, which is represented by the maximum absorbance. 4. Subsequently, with binding of both plasminogen and t-PA to fibrin, plasmin lyses the clot, indicated by a decrease in absorbance. B) (Left) Scanning electron micrograph of a clot generated from isolated fibrinogen containing high levels of nitrated fibrinogen molecules. Magnification bar represents 10  $\mu\text{m}$ . Arrows indicate clusters, defined as 8 single fibers crossing at a single point. (Right) Scanning electron micrograph of a clot generated from the PPP of a PE subject. Magnification bar represents 2  $\mu\text{m}$ . Arrowheads designate fiber bundles, defined as 3 or more fibers aligning together for greater than 1  $\mu\text{m}$ . Fiber density is commonly measured manually by counting the number of fibers in a small, designated area repeated over the entirety of the micrograph. Fiber diameter is also measured manually, often using Image J (Rasband, W.S., ImageJ, U. S. National Institutes of Health, Bethesda, Maryland, USA, <http://imagej.nih.gov/ij/>, 1997–2011) or other software that is capable of determining distance. Here, the widths of many fibers are measured within the micrograph either in designated areas or at random. C) Western blot of FXIIIa crosslinking fibrin in PPP. Addition of thrombin and  $\text{CaCl}_2$  in PPP initiates both fibrin polymerization and FXIII activation. In isolated fibrinogen, FXIII must also be added to induce crosslinking. FXIIIa

crosslinking of fibrin creates both  $\gamma$ -dimers and  $\alpha$ -polymers. The reaction is allowed to proceed for the desired time point(s) followed by quenching with concentrated sample buffer containing a reducing agent and boiling. The products of the reaction are then run on an SDS PAGE gel followed by Western blot analysis with an anti-fibrinogen antibody. Lane 1) 0 hr 2) 5 min 3) 15 min 4) 30 min 5) 1 hr 6) 2 hr 7) 4 hr 8) 8 hr 9) 24 hr 10) No thrombin or  $\text{CaCl}_2$ .

**Table 1**

Site-specific oxidative modifications of fibrinogen and effects on fibrin function and clot structure.

	<i>In vitro</i> photo-oxidized fibrinogen [47, 48]	<i>In vitro</i> HOCl-oxidized fibrinogen [56]	<i>Ex-vivo</i> plasma of smokers and non-smokers [32]
<b>Type of modification</b>	Oxidized histidine	Methionine sulphoxide	3-Nitrotyrosine
<b>Site of modification</b>	B $\beta$ His <sup>16</sup> ; unknown A $\alpha$	* A $\alpha$ Met <sup>476</sup> ; B $\beta$ Met <sup>367</sup> ; $\gamma$ Met <sup>78</sup>	** B $\beta$ Tyr <sup>292</sup> ; B $\beta$ Tyr <sup>422</sup>
<b>Monomer association</b>	Increased	NA	NA
<b>Fibrin polymerization</b>	Decreased	Decreased	Increased
<b>Final turbidity</b>	Decreased	Decreased	Increased
<b>Fibrin clot lysis</b>	NA	Slower	Slower
<b>Viscoelastic properties</b>	NA	Decreased stiffness and viscosity	Increased stiffness and viscosity
<b>Fibrin clot structure</b>	NA	Increased fiber density; decreased fiber diameter; decreased pore size	Increased fiber clusters

\* Most abundant modification;

\*\* Most frequent modification.