DOI: 10.1111/ith.15697

REVIEW ARTICLE

Ith

Regulation of coagulation by tissue factor pathway inhibitor: Implications for hemophilia therapy

Alan E. Mast¹ Volfram Ruf^{2,3}

¹Versiti Blood Research Institute, Milwaukee, Wisconsin, USA

²Center for Thrombosis and Hemostasis, Johannes Gutenberg University Medical Center, Mainz, Germany

³Department of Immunology and Microbiology, Scripps Research, La Jolla, California, USA

Correspondence

Alan E. Mast, Versiti Blood Research Institute, PO Box 2178, Milwaukee, WI 53201-2178. USA. Email: aemast@versiti.org

Wolfram Ruf, Center for Thrombosis and Hemostasis, Johannes Gutenberg University Medical Center, Langenbeckstraße 1, 55131 Mainz, Germany. Email: ruf@uni-mainz.de or ruf@scripps. edu

Funding information Novo Nordisk

Abstract

Tissue factor pathway inhibitor (TFPI) is an alternatively spliced anticoagulant protein that primarily dampens the initiation phase of coagulation before thrombin is generated. As such, TFPI's actions are localized to cells expressing TF and to sites of injury, where it is an important regulator of bleeding in hemophilia. The major splice isoforms TFPI α and TFPIβ localize to different sites within and surrounding the vasculature. Both forms directly inhibit factor Xa (FXa) via their Kunitz 2 domain and inhibit TF-FVIIa via their Kunitz 1 domain in a tight complex primarily localized to cells. By forming complexes localized to distinct cellular microenvironments and engaging additional cell surface receptors, TFPI alters cellular trafficking and signaling pathways driven by coagulation proteases of the TF pathway. TFPI α , which circulates in complex with FV and protein S, also serves an inhibitor of FXa independent of the TF initiation complex and prevents the formation of an active prothrombinase. This regulation of thrombin generation in the context of vessel injury is effectively blocked by antibodies to Kunitz 2 domain of TFPI and exploited as a therapy to restore efficient hemostasis in hemophilia.

KEYWORDS

blood coagulation, hemophilia, hemostasis, signal transduction, tissue factor pathway inhibitor

INTRODUCTION 1 |

Tissue factor pathway inhibitor (TFPI) is an alternatively spliced anticoagulant protein present in plasma,¹ platelets,^{2,3} and extracellular matrix,⁴ as well as on the surface of endothelial cells,^{5,6} monocytes, and macrophages.⁷ It is a multifunctional Kunitz-type serine protease inhibitor that acts at several steps of the blood coagulation cascade.⁸ TFPI primarily inhibits initiation of coagulation, dampening procoagulant stimuli before thrombin is generated.⁹⁻¹¹ As such, it is an important regulator of bleeding in hemophilia,¹²⁻¹⁴ and antibodies that block TFPI activity are under development for hemophilia treatment.¹⁵⁻¹⁸ In phase 2 clinical studies, an anti-TFPI antibody prevented bleeding episodes in people with hemophilia A and B, with or without inhibitors.¹⁷ As an inhibitor of blood coagulation, TFPI also alters cellular trafficking and signaling pathways driven by coagulation proteases of the TF pathway.^{19,20} Here, we review the biochemistry and physiology underpinning the effects of TFPI in blood coagulation and cellular signaling pathways, and how these may be altered by anti-TFPI therapeutic strategies.

Manuscript handled by: Patricia Liaw

Final decision: Patricia Liaw, 07 March 2022

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2 | TFPI BIOCHEMISTRY

2.1 | Overview of TFPI isoforms and structures

mRNA encoding three isoforms of TFPI are produced in humans. The isoforms, TFPI α , TFPI β , and TFPI δ , result from three alternative splicing events that produce distinctive C-terminal ends of the TFPI α and TFPI β proteins. However, *in vivo* production of TFPI δ has not been demonstrated and its physiological relevance is uncertain.²¹ This variation in the C-termini targets the individual isoforms to different locations within the vasculature and alters their anticoagulant activities.⁸ The two major isoforms have identical N-termini consisting of an acidic stretch of amino acids followed by two Kunitz-type serine protease inhibitory domains (Figure 1). These two Kunitz domains (K1 and K2) inhibit the tissue factor-factor VIIa (TF-FVIIa) catalytic complex in a factor Xa (FXa)-dependent manner. The isoform structures diverge after the K2 domain (Figure 1). TFPI α has a third Kunitz domain (K3) that binds protein S (PS),²² followed by a stretch of

basic amino acids that bind glycosaminoglycans and other negatively charged polymers.^{23,24} The basic C-terminus of TFPI α also binds to an acidic region of the FV B-domain allowing it to inhibit early forms of the FVa-FXa catalytic complex (prothrombinase) that assemble before thrombin is generated.¹⁰ TFPI β has a stretch of amino acids that encode a glycosylphosphatidylinositol (GPI) cell-surface attachment sequence.²⁵ TFPI β rapidly inhibits TF-FVIIa procoagulant activity when both proteins are expressed on the same cell.¹⁹

2.2 | Localization of TFPI isoforms

The distinctive C-termini of TFPI α and TFPI β localize them to different portions of the vasculature. TFPI α circulates in plasma as a soluble protein at ~0.4 nM (~12.5 ng/ml).^{1,26} TFPI circulates in association with PS and FV.^{27,28} Consistently, the plasma concentration of TFPI α correlates with the plasma concentrations of PS and FV.²⁶ TFPI α is also a heparin-releasable protein, and its plasma concentration rapidly



FIGURE 1 TFPI isoforms in humans. Yellow residues represent cysteine residues and disulfide bridges. Blue residues are basic. Red residues are acidic. a, active; F, factor; GPI, glycophosphatidylinositol; K, Kunitz domain; TFPI, tissue factor pathway inhibitor. Adapted from Maroney SA and Mast AE. J Thromb Haemost. 2015;13(S1):S200–S207¹¹³

increases two- to four-fold following heparin infusion.^{19,29} It has been suggested that the heparin-releasable pool of TFPI α is bound to the endothelial surface via interactions between its basic C-terminal region and glycosaminoglycans within the endothelial glycocalyx. TFPI α also interacts with cell surface glycosaminoglycan receptors,²⁰ which can be either GPI-anchored (glypican 3)³⁰ or transmembrane domainanchored (syndecan-3 and syndecan-4)^{31,32} through its third Kunitz and carboxyl terminal domains.³³ However, TFPI α is not detectable on the surface of cultured endothelial cells or human kidney endothelium, and recent data indicate the extracellular matrix is the major source of heparin releasable TFPI α .⁴ This localizes TFPI α within the extravascular space, where it is available to regulate the activity of endogenously expressed TF on cells exposed to blood following endothelial injury.

Although TFPI α is not present on the endothelial surface, it is present within granule-like structures distinct from Weibel-Palade bodies in cultured Ea.hy926 cells.^{4,34} TFPI α is also present within platelets.³ It is released from activated platelets and accumulates at the site of vascular injury, with a portion localizing to the platelet surface.^{2,3} Interestingly, TFPI α is not present in platelet α -granules,³ and, accordingly, the platelet TFPI α concentration does not correlate with platelet PS or FV, both of which are within α -granules.²⁶ TFPI α anticoagulant function is also regulated by polyphosphates that are released from activated platelets.^{10,35}

TFPI β is localized to the surface of endothelial cells and monocytes via a GPI anchor.^{5,7,25} Expression of TFPI β on the surface of cultured endothelial cells is not altered by heparin,^{4,36} and TFPI β does not contribute to the heparin-releasable pool of TFPI observed in vivo.³⁷ TFPI β is not present in platelets³ or within the extracellular matrix.⁴

3 | TFPI INHIBITS COAGULATION THROUGH DISTINCT BIOCHEMICAL MECHANISMS

TFPI inhibits coagulation proteases at several steps of the blood coagulation cascade (Figure 2).^{8,9} The direct binding of K2 to the FXa active site provides the foundation for its anticoagulant activities,³⁸ the inhibition of TF-FVIIa performed by TFPI α and β ,¹¹ and the inhibition of early forms of prothrombinase performed by only TFPI α .¹⁰ These catalytic complexes assemble early in a procoagulant response, and their inhibition by other anticoagulant proteins, such as antithrombin³⁹⁻⁴¹ and activated protein C.^{42,43} The inability to overcome TFPI inhibitory activity through generation of additional FXa via the FVIIIa-FIXa catalytic complex reduces thrombin generation and results in the bleeding experienced by hemophilia patients.⁴⁴ This provides the mechanistic basis for inhibition of TFPI as a treatment for hemophilia bleeding.

3.1 | TFPI inhibition of TF-FVIIa

TF-FVIIa initiates the blood coagulation cascade by activating FX of the common blood coagulation pathway¹¹ and FIX of the intrinsic

blood coagulation pathway⁴⁵ (Figure 2). The inhibition of TF-FVIIa is mediated by direct binding of K1 to the FVIIa active site.³⁸ However, K1 is a poor inhibitor of TF-FVIIa and efficient inhibition requires FXa.¹¹ Thus, TFPI has been described as a two-stage inhibitor where K2 binds to FXa, and in a subsequent inhibitory step the TFPI-FXa complex inhibits TF-FVIIa. However, the rate-limiting step for inhibition of TF-FVIIa is the inhibition of FXa.⁴⁶ This indicates that instead of occurring in a two-step process, TFPI simultaneously inhibits the ternary TF-FVIIa-FXa complex in a single reaction occurring immediately after FX is activated by TF-FVIIa but before FXa dissociates from the ternary TF-FVIIa-FXa complex.⁴⁶ In this manner, TFPI blocks TF-FVIIa mediated generation of prothrombinase, FVa-FXa, in the common blood coagulation pathway. However, because TFPI does not inhibit activation of FIX by TF-FVIIa, TF-FVIIa-mediated initiation of blood coagulation can also proceed through the intrinsic pathway. Additionally, the TF-FVIIa-FXa ternary complex can directly activate FVIII in a process that is poorly inhibited by TFPI and occurs before the dissociation of the nascent product FXa.⁴⁷ This provides a mechanism for TF-FVIIa-mediated activation of both antihemophilic factors and production of the FVIIIa-FIXa intrinsic tenase complex independent of thrombin providing feedback activation of FVIII.47

Because TFPI α and TFPI β have K1 and K2, they both inhibit TF-FVIIa through this FXa-dependent mechanism. The localization of TFPI β to cell surfaces via its GPI anchor greatly enhances its TF-FVIIa inhibitory activity; soluble forms of TFPI that mimic TFPI β and contain only K1 and K2 are much weaker inhibitors.¹⁹ Similarly, the TFPI α K3 domain binds to PS,^{22,48} which contains a GIa domain that localizes TFPI α to cell surfaces and enhances its ability to inhibit FXa.⁴⁹ Additionally, about two-thirds of plasma TFPI α is C-terminally degraded and associated with lipoproteins.³⁷ This pool of plasmatruncated TFPI α has reduced anticoagulant activity in TF-FVIIainitiated plasma coagulation assays. However, it is a potent inhibitor of the propagation phase in thrombin generation assays.⁵⁰ At this point, the physiological importance of lipoprotein-associated plasma TFPI α remains uncertain.

3.2 | TFPIα inhibition of early forms of prothrombinase

Prothrombinase is the FVa-FXa catalytic complex that converts prothrombin to thrombin at the convergence of the intrinsic and extrinsic blood coagulation pathways. The discovery that TFPI α is the only TFPI isoform present in platelets³ and recognition of amino acid homology between the TFPI α C-terminus and the FV B-domain⁹ led to description of the inhibitory mechanism of early forms of prothrombinase (i.e., those that assemble before thrombin is generated¹⁰ and that involves FXa dependent activation of FV).⁵¹

In this mechanism, the TFPI α basic region binds tightly to partially activated forms of FV that lack this basic region but still contain the acidic region of the B-domain, allowing interaction with TFPI α as well as protein S.⁵² This allows prothrombinase



FIGURE 2 Presence of TFPI isoforms result in the inhibition of the TF-FVIIa-FXa ternary complex. (A) The TF-FVIIa complex activates FIX (part of the intrinsic coagulation pathway) and FX (part of the common coagulation pathway). This results in plasma membrane association of FXa and FVa, forming the prothrombinase complex, and subsequent cleavage of prothrombin to thrombin. (B) The K2 domain of TFPIα binds to FXa, which supports inhibition of the TF-FVIIa complex. The binding of the K3 domain to PS supports membrane association of TFPI α and thus further promotes inhibition of the TF-FVIIa complex. The basic C-terminus of TFPI α can also interact with FVa, resulting in inhibition of the early prothrombinase complex. (C) TFPI β localizes to the cell surface via its GPI anchor, increasing its ability to inhibit FXa. a, active; C, carboxy; F, factor; GPI, glycophosphatidylinositol; K, Kunitz; N, amino; TF, tissue factor; TFPI, tissue factor pathway inhibitor

inhibition mediated through K2 binding of the FXa active site and a lower affinity interaction between conserved uncharged amino acids in the TFPI α C-terminus and the FV heavy chain in the region of FV arginine 506.53 This later interaction is weakened in patients with the FV Leiden mutation, which decreases the threshold for initiation of coagulation and thereby increases thrombotic risk in these patients.^{54,55} Thrombin rapidly cleaves and removes the entire FV B-domain⁵⁶ after which TFPI α can no longer inhibit prothrombinase. However, the binding of TFPI α to

FV slows this thrombin cleavage,⁵⁷ further contributing to TFPI α anticoagulant activity. Three FV variants that lack the basic region of the B-domain have been identified: FV east Texas,^{58,59} FV Amsterdam,⁶⁰ and FV Atlanta.⁶¹ TFPI α binds tightly to these forms of FV, and consequently, patients have greatly increased plasma TFPIa concentrations and an associated bleeding disorder. Thus, it appears that $TFPI\alpha$ inhibition of prothrombinase is a physiologically important anticoagulant activity occurring at this early point in the coagulation cascade when nascent forms

3.3 | TFPI inhibition of thrombin generation in the cell-based model of coagulation

The cell-based model proposed by Hoffman and Monroe emphasizes that blood coagulation is regulated by the properties of the cell surfaces on which coagulation reactions occur.⁶³ In line with this model, the presence of TFPI α and TFPI β on different cell surfaces allows inhibition at different stages of the procoagulant response¹⁹ (Figure 2). TFPI β inhibits the initiation of coagulation on endothelial cells that express intravascular TF following inflammatory stimuli or reactive oxidant stress,⁶⁴ while TFPI α in the extracellular matrix is positioned to dampen the initiation of coagulation by the TF-bearing cell that is exposed to circulating blood following vascular injury.⁴ However, TFPI is a poor inhibitor of TF-mediated activation of FIX and FVIII.^{46,47} which can diffuse to the surface of activated platelets accumulating at the injury site. The FVIIIa-FIXa catalytic complex on the platelet then activates FX.⁶⁵ The FXa then interacts with FVa released from platelets to form prothrombinase.⁶³ Because some forms of FVa released from platelets lack the basic region of B-domain while retaining the acidic region, ^{13,66} platelet or plasma TFPI α can block the procoagulant response by inhibition of early prothrombinase, thereby preventing the propagation of clotting by the subsequent burst of thrombin generation.¹⁰ There is evidence that this process is enhanced by interactions between TFPI α and PS that enable localization of TFPI α to the platelet surface.⁶⁷ Thus, TFPI prevents inappropriate coagulation by acting early in a procoagulant response on both the TF-bearing cell and the platelet.

By exerting anticoagulant activity at these early stages of the procoagulant response, TFPI is the primary physiological regulator of bleeding in hemophilia,¹⁴ and blocking TFPI activity is an effective approach for hemostatic prophylaxis in these patients.¹⁶⁻¹⁸ Several therapeutics that block TFPI have been evaluated in clinical trials. An aptamer, BAX 499, which bound to K1, K3, and the TFPI α C-terminus, was withdrawn from development in 2012 because it paradoxically increased bleeding frequency in people with hemophilia.⁶⁸ This aptamer increased plasma TFPI levels up to 25-fold, possibly by causing release of endothelial stores of TFPI and/or by altering its clearance. Because the K2 domain was not blocked, the circulating aptamer-bound TFPI could potentially exert anticoagulant activity by inhibiting FXa. A monoclonal antibody, BAY1093448, that bound to K1 and K2 was terminated in clinical development in 2020 because of thrombotic events

in three people with hemophilia. There are two monoclonal antibodies, concizumab and marstacimab, that bind to K2 and are currently in phase 3 clinical trials. Marstacimab was given to 58 people with hemophilia in phase 2 studies with no episodes of thrombosis.⁶⁹ Concizumab was given to 54 people with hemophilia in phase 2 studies with no episodes of thrombosis.¹⁷ Phase 3 studies of concizumab were paused in March 2020 after three enrolled subjects with hemophilia had nonfatal thrombotic events. These patients also had preexisting risk factors and used concomitant hemostatic medication with FVIII or FVIIa.⁷⁰ Phase 3 studies resumed in August 2020 after safety issues were addressed and guidelines for the management of bleeding episodes with concomitant hemostatic agents and updates to the concizumab prophylactic dosing regimen were implemented.

Importantly, targeting TFPI does not alter the anticoagulant activity of activated protein C or antithrombin, which inhibits later steps of the coagulation cascade, and are still in place to limit thrombosis. Although inhibition of TFPI activity on both the TF-bearing cell and the platelet likely contributes to its efficacy in hemophilia prophylaxis, the cell-based model of coagulation suggests that diminished activation of FX on the platelet surface is a major contributor to bleeding in patients with hemophilia.⁶³ In support of this model, inhibition of platelet TFPI α alone is sufficient for restoring hemostasis in a mouse model of hemophilia,⁷¹ suggesting that specifically targeting platelet TFPI α rather than endothelial TFPI β may help to restore hemostasis while limiting the potential for thrombosis.

4 | ROLE OF TF IN CELLULAR SIGNALING

The TF coagulation pathway is directly connected to cell signaling by protease-activated receptors (PARs) located on the surface of endothelial cells and platelets within the vasculature, as well as on the surface of monocytes/macrophages, other immune and epithelial cells in the extravascular space. PARs are activated by TFassociated proteases FVIIa and FXa, as well as by thrombin. The TF-FVIIa catalytic complex can directly activate PAR2 in a reaction independent of FX, and prolonged endosomal signaling through this pathway is dependent on the association with integrin $\beta 1$ through a binding site in the FVIIa protease domain (Figure 3).^{72,73} Direct TF-FVIIa signaling through PAR2 is particularly relevant for extravascular processes, such as epithelial cell migration, wound healing responses, and tumor progression.^{72,74-76} In addition, TF-FVIIa directly influences growth factor signaling and angiogenesis^{77,78} and alters cell-cell interactions independently of PAR2 by cleaving ephrin receptors.⁷⁹

In a distinct signaling pathway, TF-FVIIa generated nascent product FXa (TF-FVIIa-FXa) cleaves PAR2 in the presence of the endothelial protein C receptor (EPCR).^{80,81} This signaling pathway plays a pivotal role in the innate immune system and is essential for the induction of interferon responses downstream of Toll-like receptor 4 signaling.^{81,82} In this context, TF signaling is regulated at the level



FIGURE 3 Signaling and interactions of TF and TFPI. TFPI binds coreceptors that support its cell-surface functions. Binding to GPIanchored glycosaminoglycan receptors (e.g., glypicans) via the K3 domain mediates membrane association. GPI-anchored TFPIα or TFPIβ can translocate to sphingolipid-rich raft domains and caveolae. Binding to receptors involved in internalization and degradation of FXa (e.g., lipoprotein receptor related protein 1 and VLDLR) can modulate cellular responses to various stimuli. TFPlβ is a GPI anchored via its Cterminal domain and appears to regulate cell-intrinsic functions of TF, such as immune signaling of FXa in complex with TF-FVIIa via protease activated receptor (PAR) 2 in the presence of the endothelial protein C receptor (EPCR) or signaling of the TF-FVIIa associated with integrin receptors via PAR2 after endosomal uptake. a, active; C, carboxy; F, factor; GPI, glycophosphatidylinositol; K, Kunitz; LRP1, low density lipoprotein receptor-related protein-1; TF, tissue factor; TFPI, tissue factor pathway inhibitor; VLDLR, very low-density lipoprotein receptor

of EPCR by the anticoagulant protein C pathway with FV and PS as cofactors.⁸³ This PAR2 signaling pathway requires only very low concentrations of FVIIa⁸⁰ and may therefore occur not only in intravascular but also extravascular milieus when TF-expressing cells are exposed to exudated plasma components. Moreover, monocytes and macrophages autonomously synthesize FVII and FX, and macrophage polarization in the tumor microenvironment is critically dependent on macrophage FX synthesis and activation of PAR2.⁸⁴ The synthesis of coagulation factors by immune cells is differentially regulated. Whereas FVII is constitutively expressed in tissue resident macrophages of the peritoneal cavity and lungs,^{85,86} FX is transcriptionally induced in response to inflammatory stimuli.⁸⁴ The innate immune cell signaling roles of the TF-VIIa complex can therefore involve either cell autonomous, extravascular FX synthesis or FX from plasma sources.

TF-initiated coagulation also induces procoagulant signaling by thrombin activating PARs on endothelial and immune cells. Intravascular thrombin signaling is counterbalanced by multiple mechanisms, including the endothelial cell-localized anticoagulant protein C pathway as well as plasma anticoagulants, platelet receptors and fibrin(ogen).⁸⁷ There is also expanding evidence that thrombin generation occurs in extravascular locations and can control stem cell activity through PAR1 signaling at steady state and in response to injury. $^{\it 88}$ Similarly, tissue macrophages synthesize FV and assemble a functional prothrombinase complex.⁸⁹ No clear roles have emerged for TFPI in controlling these thrombin-dependent vascular and extravascular signaling events.

ROLE OF TFPI IN MODULATING 5 | **TF-DEPENDENT CELLULAR SIGNALING**

TFPI is a relatively weak inhibitor of TF-FVIIa.^{11,90} However, its affinity for TF-FVIIa is greatly increased by binding to PS in reactions that lack FX but include FIX.⁹¹ Similarly, TFPI associated with the matrix can support cell adhesion through interaction with TF-FVIIa expressed by tumor cells. Remarkably, this process studied in cell culture is entirely independent of FXa.⁷⁶ Thus, there are no compelling data that regulation of extravascular signaling functions of the TF-FVIIa complex are influenced by antibody blockade of TFPI binding to FXa.

TFPI is a potent inhibitor of the TF-FVIIa-FXa ternary complex and thereby modulates cellular signaling at vascular interfaces.²⁰ TFPI also decreases TF-initiated thrombin generation and may thereby dampen intravascular coagulation signaling mediated by thrombin in the context of sepsis. Indeed, leukocyte proteases released in this context are known to degrade TFPI and thus contribute to disseminated intravascular coagulation in severe infections.92,93

Pathological mechanisms that release TF from the tight inhibition by TFPI are not restricted to TFPI degradation in infections. TF-dependent cell signaling is also stimulated in the context of autoimmune disease, where antiphospholipid antibodies destabilize the TF-FVIIa-FXa complex inhibited by TFPI. Remarkably, blocking TFPI with polyclonal antibodies did not trigger but rather prevented the same autoimmune signaling,⁷ emphasizing that TF control on immune cells can be achieved by alternative mechanisms in the absence of functional TFPI (e.g., by TF cellular degradation or release on extracellular vesicles).⁹⁴

The two major TFPI isoforms distinctly modulate cell signaling through their different mechanisms for cell-surface association (Figure 3). TFPI α binds GPI-anchored co-receptors on the cell surface, whereas TFPI β is directly GPI-anchored to cellular microdomains.^{3,4} The mode for cell-surface association determines the subcellular and membrane microdomain localization of TFPI isoforms, which, in turn, affects intracellular trafficking relevant for signaling and protein degradation. TFPI α also binds to transmembrane receptors mediating the cellular uptake and degradation of FXa (e.g., lipoprotein receptor related protein)⁹⁵ or regulating angiogenesis (i.e., very low density lipoprotein receptor [VLDLR]).⁹⁶ In this manner, soluble TFPI α from various sources may bind to target cells and modulate cellular responses. In contrast, cell-intrinsic functions of TF appear to be predominantly regulated by GPI-anchored TFPI β and/or endogenously synthesized TFPI α tightly bound to the cell surface through GPI-anchored receptors.^{20,25,36,97}

Monocytes express low levels of TF in the absence of inflammatory stimuli that can assemble into a TF-FVIIa-FXa complexes inhibited by membrane-anchored TFPI.⁷ Genetic evidence indicates that EPCR is required for the formation of the quaternary TF-FVIIa-FXa-TFPI complex on the monocyte cell surface.⁸¹ Antiphospholipid antibodies interact with EPCR and release the TF-FVIIa-FXa complex from TFPI inhibition, triggering TF and PAR1/2-dependent procoagulant responses and pathogenic signaling.⁸¹ Thus, TFPI has a paradoxical prothrombotic role by priming responses in the autoimmune antiphospholipid syndrome.⁷ However, genetic deletion of TFPI from monocytes has no apparent prothrombotic effect in other models of experimental thrombosis, indicating that TFPI has specific functions in disease pathologies that are unrelated to its physiological regulatory roles.

Endogenously synthesized, membrane-anchored TFPI efficiently regulates signaling of the TF-FVIIa-FXa complex in cytokine stimulated endothelial cells,²⁰ in line with data demonstrating highly efficient inactivation of cellular TF by TFPIβ.²⁵ In contrast, exogenously added TFPIα appears to be relatively ineffective in blocking cell signaling in comparison to suppressing TF-initiated coagulation.²⁰ Thus, TFPI isoforms have different potencies in regulating TF-dependent cell signaling. However, loss of TFPI in endothelial cells, similar to the observations in monocytes, does not necessarily result in increased thrombosis because deletion of TFPI in endothelial cells downregulates TF mRNA expression in the vessel wall.⁹⁸

6 | IMPLICATIONS OF TFPI LOSS ON SIGNALING PATHWAYS, AND PATHOLOGICAL AND PHYSIOLOGICAL PROCESSES

Deletion of K1 in mice causes embryonic lethality,⁹⁹ implicating TFPI as a crucial regulator of TF activity during developmental processes.

In addition, more detailed studies on mid-gestational lethality of TFPI-deficient mouse embryos demonstrate roles of TFPI in vascular development and angiogenesis. TFPI deletion causes vascular abnormalities specifically in the central nervous system and defects in the cerebrovascular development are prevented by genetic deletion of FV,¹⁰⁰ suggesting that TFPI is an important regulator of thrombin-dependent signaling events that modulate cerebrovascular development. These vascular development defects cannot be rescued by genetic overexpression of platelet TFPI α ,¹⁰¹ which is efficient in preventing death during early embryonic development. In addition, genetic reduction of TFPI expression leads to thrombotic perinatal lethality in mice carrying the prothrombotic FV_{Leiden} mutation,¹⁰² further emphasizing the crucial role of TFPI as a regulator of the common coagulation pathway during pre- and postnatal development.

The extent to which TFPI regulates vascular processes by modulating TF pathway signaling is not fully understood. TF supports postnatal vascular development in part through PAR2 signaling, and TFPI-like inhibitors specifically suppress hypoxia-induced angiogenic sprouting⁷⁸ and tumor angiogenesis.¹⁰³ Locally applied high concentrations of TFPI also suppress tumor growth, whereas FXa inhibitors were not effective in this study.¹⁰³ It is conceivable that under pathological conditions with elevated TF levels, such as in cancer and pathological angiogenesis, extravascular TFPI is not produced at sufficiently high levels to control TF signaling and that these pharmacological inhibitory strategies restore functional control of increased TF expression and pathological signaling. In addition, TFPI interactions with endothelial cells modulate proangiogenic growth factor signaling¹⁰⁴ and interaction of TFPI with VLDLR induces endothelial cell apoptosis.⁹⁶ These regulatory roles of TFPI have been mapped to carboxyl terminal domains of TFPIa, which are not directly targeted by anti-hemophilic strategies to neutralize TFPI's inhibitory functions towards FXa.

TFPI degradation is observed in pathological processes. The connecting region between K1 and K2 is protease sensitive,⁹³ and therapeutic intervention with fibrinolytics leads to TFPI degradation with a shift toward a procoagulant state on monocytes.¹⁰⁵ Proteolytic inactivation of TFPI serves important functions in host defense against intravascular pathogens,¹⁰⁶ but may ultimately lead to disseminated intravascular coagulation in severe sepsis.¹⁰⁷ Whereas TFPI degradation plays a role in the host defense to infection and inflammatory processes, loss of TFPI function leads to pathologies and disease particularly in the context of concomitant activation or upregulation of TF.

TFPI is expressed by cancer cell lines *in vitro* and bound to transmembrane syndecan- 3^{32} or GPI-anchored glypican- 3^{108} receptors. Cancer cell-associated TFPI appears to play a minor role in regulating cancer cell TF clotting activity and rather supports leukemia dissemination by promoting chemokine-dependent cell motility in patient-derived cell lines.¹⁰⁸ In contrast, downregulation of TFPI β enhances breast cancer cell line migration,¹⁰⁹ supporting the notion that TFPI isoforms regulate distinct aspects of cancer biology. Consistently, deletion of presumably membrane

anchored TFPI in endothelial cells enhances experimental metastasis in mice.⁹⁸ Thus, the role of TFPI in these processes is diverse and does not point to uniform functions in disease pathologies. TFPI also plays critical regulatory roles within the vessel wall and TFPI deletion exacerbates atherosclerosis,¹¹⁰ whereas overexpression in smooth muscle cells prevents atherosclerosis and injury-induced hyperplasia in mouse models.^{111,112} There is no evidence that the intravascular neutralization of TFPI for improved hemostasis in hemophilia influences these regulatory functions of vessel wall expressed TFPI in vascular inflammation.

7 | CONCLUSIONS

Structural differences in the C-terminal regions of TFPI α and TFPI β splice isoforms affect how they localize to cells or circulate in blood and regulate the coagulation cascade. The C-terminal region of TFPI α binds tightly to nascent forms of FVa, allowing for inhibition of prothrombinase, in a reaction that also involves binding of K2 to the FXa active site. The C-terminal region of TFPI β encodes a GPI-anchor attachment sequence localizing it to the cell surface where it is a highly effective inhibitor of TF-FVIIa complexes that assemble on the same cells. Because TF-FVIIa can be inhibited by K1 of TFPI α when bound to PS in the absence of FXa, therapeutic strategies directed to K2 may avoid, in part, the crucial function of TFPI in inhibiting TF-dependent coagulation reactions and cell signaling.

Directly or indirectly membrane anchored forms of TFPI regulate activity of TF-FVIIa and the TF-FVIIa-FXa coagulation initiation complex and their functions in cell signaling. TF-FVIIa regulates cellular functions occurring outside the vasculature, such as angiogenesis, tumor biology, and inflammation. In these extravascular processes, TFPI is particularly important for control of the TF-FVIIa-FXa coagulation initiation complex. The TFPI-TF-FVIIa-FXa complex has remarkable stability and forms rapidly to control excessive signaling of TF; however, other mechanisms may regulate TF on immune or vascular cells, and anti-TFPI antibodies do not act within the extravascular space.

Within the intravascular space, TFPI exerts anticoagulant activity early in the coagulation cascade, before thrombin is generated, and at steps that do not require FVIII or FIX. Inhibition of TFPI allows amplification of coagulation through pathways that bypass FVIII and FIX. Therefore, TFPI is an attractive target for management of hemophilia-related bleeding. Currently, several anti-TFPI antibodies that target the intravascular activity of TFPI are in clinical trials for hemophilia prophylaxis. They can be dosed subcutaneously and effectively prevent bleeding episodes in patients with hemophilia A and B, with or without inhibitors.

ACKNOWLEDGMENTS

Editorial support (in the form of figure preparation and referencing) was provided by Ashfield MedComms GmbH (Mannheim, Germany), and was supported by Novo Nordisk as an independent investigator-led initiative.

CONFLICTS OF INTEREST

A.E.M. receives research funding from Novo Nordisk, has received honoraria for serving on Novo Nordisk and Vega Therapeutics Advisory Boards, and honoraria for educational seminars from Novo Nordisk. W.R. reports intellectual property and royalty income, consultancy and research support from ARCA biopharma, research support from ICONIC therapeutics, honoraria for educational seminars from Novo Nordisk, and affiliation with Meru Vasimmune.

AUTHOR CONTRIBUTION

Alan E. Mast and Wolfram Ruf wrote the manuscript and provided the outline of the manuscript figures.

ORCID

Alan E. Mast ⁽¹⁾ https://orcid.org/0000-0003-3740-0318 Wolfram Ruf ⁽¹⁾ https://orcid.org/0000-0002-6064-2166

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How to cite this article: Mast AE, Ruf W. Regulation of coagulation by tissue factor pathway inhibitor: Implications for hemophilia therapy. *J Thromb Haemost.* 2022;20:1290–1300. doi:10.1111/jth.15697