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Tissue Factor in Cardiovascular Disease: Pathophysiology and Pharmacological Intervention

Erik W. Holy, MD and Felix C. Tanner, MD

Cardiovascular Research, Physiology Institute, University of Zurich, Zurich, Switzerland

Centre for Integrative Human Physiology, University of Zurich, Zurich, Switzerland

Cardiology, Cardiovascular Centre, University Hospital Zurich, Zurich, Switzerland

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Address for correspondence:

Felix C. Tanner, MD

Cardiology, Cardiovascular Centre, University Hospital Zurich

Rämistrasse 100, 8091 Zurich, Switzerland

Phone: +41 44 255 1111, Fax: +41 255 8701

email: felix.tanner@access.uzh.ch

Abstract

Tissue factor (TF), also known as thromboplastin or coagulation factor III, is the major trigger of the coagulation cascade and thereby crucially involved in the maintenance of vascular hemostasis. By binding factor VIIa, the resulting TF:VIIa complex activates the coagulation factors IX and X ultimately leading to fibrin and clot formation. In the vessel wall, TF expression and activity is detectable in vascular smooth muscle cells and fibroblasts and, at a much lower level, in endothelial cells, and can be induced by various stimuli including cytokines, biogenic amines, or growth factors. In addition, TF is found in the bloodstream in circulating cells such as monocytes or leukocytes, in TF containing microparticles, and as a soluble spicing isoform. Besides its well known extracellular role as a trigger of coagulation, TF also functions as a transmembrane receptor, and TF dependent intracellular signalling events regulate the expression of genes involved in cellular responses such as proliferation and migration. TF indeed appears to be involved in the pathogenesis of neointima formation and tumor growth, and increased levels of TF have been detected in patients with cardiovascular risk factors or coronary artery disease as well as in those with cancer. Therefore, pharmacological or genetic inhibition of TF may be an attractive target for the treatment of cardiovascular disease and cancer. Different strategies for inhibition of TF have been developed such as inhibition of TF expression and blockade of TF action. Clinical applications of such strategies need to be tested in appropriate trials, in particular for evaluating the advantages of targeted versus systemic delivery of the inhibitors.

1. Tissue factor: structure and function

1.1. TF protein structure

The TF (thromboplastin, CD142) gene located on chromosome 1 at p21-22 spans a 12.4 kb region and contains six exons encoding a single transmembrane polypeptide chain composed of 263 amino acids (Kao et al., 1988; Mackman et al., 1989; Morrissey et al., 1989; Scarpati et al., 1987). The TF protein is divided into three domains: an extracellular (219 amino acids), a membrane-spanning (23 amino acids) and a cytoplasmic (21 amino acids) region (Breitenstein et al., 2010). TF is considered a class I integral membrane protein since the amino-terminus of the protein is located extracellularly while the cytoplasmic tail contains the carboxy-terminus. Mobility studies on sodium dodecyl sulfate gels reveal that the molecular weight of the fully glycosylated protein is about 45'000 kDa (Broze et al., 1985). The extracellular domain provides two high affinity sites for the binding of factor VII to TF. This binding is crucial for induction of the conformational changes required to induce the catalytic activity of factor VII (Edgington et al., 1991; Rehemtulla et al., 1991). In addition, anchoring of the TF protein to the membrane via the hydrophobic transmembrane domain has been demonstrated to be essential for procoagulant activity of the TF protein. However, the nature of the anchoring domain does not seem to play a role (Paborsky et al., 1991). Indeed, deletion of the cytoplasmic tail is not paralleled by an impaired TF procoagulant activity (Paborsky et al., 1991). Therefore, recent studies focussed on possible non-hemostatic functions of this domain. Experiments performed mainly in fibroblasts and cancer cell lines revealed that phosphorylation of the cytoplasmic tail activates intracellular signaling

pathways and induces cellular responses which are described in the following chapter.

In 2003, an alternatively spliced (as) isoform of TF mRNA lacking the fifth exon and exhibiting a frameshift mutation in the carboxy-terminal region was characterized (Bogdanov et al., 2003). This TF protein lacks membrane anchorage and is detectable in the plasma as a soluble TF protein. While the asTF protein has been described in different species and cell types, its contribution to the TF procoagulant activity in plasma as well as thrombus formation in vivo remains barely understood (Bogdanov et al., 2006; Mackman, 2007).

1.2. Tissue factor: key trigger of coagulation

In intact vessels, blood circulates as a fluid due to the physiological balance between natural procoagulant and anticoagulant factors. The clotting cascade defines a complex system of zymogens and enzymatic reactions designed to limit blood loss and ultimately restore vascular integrity by formation of a clot following vessel injury. Initially, initiation of coagulation was thought to be triggered by two distinct pathways: the intrinsic, or contact activation pathway and the extrinsic, or tissue factor (TF) pathway. Even though the two pathways are independently activated, the distinction between both of them has been blurred since they share a final common pathway and, in addition, there is crosstalk between the two systems. The intrinsic pathway is activated when blood or plasma comes in contact with subendothelial connective tissue or negatively charged surfaces and involves circulating factors such as factor XII, factor XI, factor IX, prekallikrein, and high molecular weight kininogen (Gailani and Renne, 2007). However, it has been questioned that this pathway is required for maintenance of normal haemostasis, since congenital deficiency in the involved

factors, for example factor XII, does not result in an increased bleeding tendency or clinically apparent pathologies in humans.

Coagulation through the TF pathway is activated when blood interacts with a TF expressing cell surface, such as it occurs after endothelial erosion, plaque rupture, or other vascular injury (Fig. 1). The initial step consists in the reversible binding of the zymogen factor VII (FVII) or the activated FVII (FVIIa) to membrane-bound TF (Mackman, 2009; Mackman et al., 2007), resulting in the formation of both TF:FVII and TF:FVIIa complexes. Only the latter, however, possesses the enzymatic activity required for further activation of downstream proteases. Once bound to TF, the inactive FVII is converted to FVIIa via limited proteolysis by several plasma proteases, such as XIIa, Xa, IXa, or thrombin, generating a positive feedback loop (Steffel et al., 2006). The TF:FVIIa complex is indeed a potent activator of the coagulation cascade and is composed of TF acting as a cofactor with regulatory functions, and the serine protease FVIIa exerting catalytic functions (Breitenstein et al., 2010). The TF:FVIIa complex activates factor IX to factor IXa which in association with VIIIa catalyzes the conversion of factor X to Xa. In addition, factor X is directly converted to factor Xa by the TF:FVIIa complex. Factor Xa in association with factor Va and divalent calcium forms the so called prothrombinase complex which cleaves prothrombin to form thrombin (Breitenstein et al., 2010). Thrombin exerts direct effects on coagulation through conversion of fibrinogen to fibrin, activation of platelets, and activation of various proteases of the coagulation cascade such as FVII, FXI, and FV representing an important auto-feedback loop. Besides its well known role in clot formation, the functions of thrombin extend from coagulation to vascular remodelling by stimulating proliferation of vascular smooth muscle and endothelial cells as well as secretion of growth factors ultimately leading to

restoration of vascular integrity (Ferrara and Davis-Smyth, 1997; Maragoudakis et al., 2002; Tsopanoglou et al., 2002; Tsopanoglou et al., 2004).

1.3. Tissue factor: a signaling receptor

The tight link between hemostasis and progression of vascular alterations such as inflammation or atherosclerosis provides evidence for complex interactions between the coagulation cascade and vascular pathophysiology. Mainly because of its similarity to class II cytokine receptors recent studies suggested a non-hemostatic role of TF in different biological processes through induction of intracellular signaling events (Bazan, 1990). The following chapter provides an overview of TF mediated cell signaling (Fig. 1).

Evidence for TF:VIIa induced signaling is derived from experiments in which FVIIa binding to TF was shown to induce intracellular calcium transients in different types of TF expressing cells (Camerer et al., 1996; Rottingen et al., 1995). This observation was followed by studies demonstrating that binding of factor VIIa to TF induces the activation of the three major mitogen activated protein (MAP) kinase family members p38 MAP kinase, p42/p44 MAP kinase, and c-Jun N-terminal kinase (JNK) (Camerer et al., 1999; Poulsen, et al., 1998). In fibroblasts, the TF-FVIIa induced activation of Src-like kinases and PI3-kinase leads to activation of the downstream Rho-like guanosine triphosphatases Rac and Cdc42 resulting in cytoskeletal reorganization (Versteeg et al., 2000).

Hence, by activating different kinases involved in signal transduction, the TF:VIIa complex regulates the transcription of numerous genes involved in physiological and pathophysiological processes such as cell migration, cell growth, or apoptosis. Indeed, binding of factor VIIa to TF induces activation of key transcription factors

such as Egr-1 or nuclear factor kappa B and the activation of the RNA polymerase A (Camerer, et al., 1999; Pendurthi et al., 1997). Moreover, microarray analysis of the gene expression profiles revealed increased mRNA levels of growth factors (i.e. connective tissue growth factors CCN1 and CCN2, fibroblast growth factor 5) and inflammatory cytokines (i.e. interleukin-1 β , interleukin-8) following factor VIIa and TF interaction (Camerer et al., 2000; Pendurthi, et al., 2000).

Interestingly, induction of signal transduction events was demonstrated to depend on the catalytic activity of factor VIIa, because inactivated factor VII or anti-TF antibodies blunt these effects (Camerer et al., 1999; Camerer et al., 1996; Pendurthi et al, 2000; Poulsen et al., 1998). Consistent with this observation, the TF:VIIa complex initiates intracellular signalling via G-protein-coupled protease-activated receptors (PARs) (Riewald & Ruf, 2002; Siegbahn, 2000). Currently, 4 different PARs have been characterized: PAR1, PAR2, PAR3, and PAR4. There is growing evidence that TF:VIIa signaling is mediated mainly by PAR2 and to a lesser extent by PAR1, while PAR3 and PAR4 do not seem to be involved. For instance, in fibroblasts derived from PAR1 knockout mice, the FVIIa induced response only occurred when TF and PAR2 were co-expressed (Camerer et al., 2000). Moreover, data obtained in a carcinoma cell line demonstrate that antibodies targeting PAR2, but not PAR1, blunt TF:VIIa induced IL-8 expression and smooth muscle cell migration (Marutsuka et al., 2002).

Even though not essential for TF:VIIa induced activation of MAP kinases and gene expression, the cytoplasmic domain of TF is implicated in cytoskeletal organization and cell migration (Bromberg, Konigsberg, Madison, Pawashe, & Garen, 1995; Mueller & Ruf, 1998). Protein kinase C mediated phosphorylation of the cytoplasmic domain at the serine residues Ser253 and Ser258 enables the interaction of the cytoplasmic tail with the actin-binding protein 280 thereby regulating actin filament rearrangement (Ott et al., 1998; Zioncheck et al., 1992).

Studies performed in mice lacking the cytoplasmic tail of TF demonstrated the crucial role of the cytoplasmic domain for vascular smooth muscle migration and vascular remodelling after injury in vivo (Ott et al., 2005).

Since increasing evidence suggests an essential role of TF as an important signal transducer beyond its well-known functions in coagulation, developing specific inhibitors that suppress either TF:VIIa mediated signaling or its coagulant function may offer interesting therapeutic approaches for the treatment of TF associated diseases.

2. Tissue factor in cardiovascular disease

2.1. TF distribution

2.1.1. Vascular TF

Under normal hemostatic conditions, TF is only barely expressed in endothelial cells (Fig. 2). Inflammatory cytokines such as tumour necrosis factor- α (TNF- α) (Holy et al., 2005) or interleukin-1 β (Napoleone et al., 1997) enhance TF expression. This induction of TF protein is mainly mediated by the mitogen-activated protein (MAP) kinases p38, extracellular-signal-regulated kinase (ERK), and c-Jun N-terminal kinase (JNK) which activate transcription factors such as nuclear factor kappa B, activator protein-1, or early growth-response gene product-1, all of them involved in transcription of the TF gene (Bavendiek et al., 2002; Mackman, 1997; Mechtcheriakova et al., 2001). Depending on the stimulus, induction of TF expression occurs via activation of either all three MAP kinases (Holy et al., 2009; Steffel et al., 2006; Steffel et al., 2005) or only a single pathway (Mechtcheriakova et al., 2001). Protein kinase C as well as the Rho-kinase pathway have also been shown to mediate the induction of endothelial TF expression (Eto et al., 2002; Mechtcheriakova et al., 2001; Zhang et al., 2007). Whereas the MAP kinase, Rho-kinase, and protein kinase C pathways regulate TF expression in a positive manner, the phosphoinositide 3 kinase pathway exerts a negative regulatory effect on TF expression (Blum et al., 2001; Steffel et al., 2005). Indeed, downstream targets of phosphoinositide 3 kinase are able to regulate TF expression at both the transcriptional and the translational level. Akt and glycogen synthase kinase 3 β regulate TF transcription (Eto et al., 2005), whereas the mammalian target of rapamycin and p70S6 kinase inhibit TF translation (Guba et al., 2005; Steffel et al.,

2005). An additional mechanism affecting the regulation of TF protein expression is stabilization of TF messenger RNA. A recent study in endothelial cells demonstrated that a decrease in TF promoter activity may be counteracted by an increased TF mRNA stability leading to an overall enhanced TF protein expression (Holy et al., 2009).

In contrast to endothelial cells, vascular smooth muscle cells constitutively express TF providing a hemostatic barrier after vascular injury (Fig. 2; Breitenstein et al., 2008; Camici et al., 2006; Schechter et al., 1997; Wilcox et al., 1989). Similar to endothelial cells, TF expression in vascular smooth muscle cells is enhanced by several receptor ligands such as CD40 ligand, histamine, thrombin, bacterial endotoxin, oxidized LDL, or C-reactive protein (Camici et al., 2006; Kamimura et al., 2004; Llorente-Cortes et al., 2004; Taubman et al., 1993; Wu et al., 2008). Even though not investigated to the same extent as in endothelial cells, the MAP kinase and phosphoinositide 3-kinase pathways regulate TF expression also in vascular smooth muscle cells.

2.1.2. Blood-borne TF

More recently, TF has been detected at low levels in plasma and urine, and these levels increase in several inflammatory disease conditions such as atherosclerosis, disseminated intravascular coagulation, glomerulonephritis, sepsis, or diabetes.

Monocytes are thought to represent a major source of the plasmatic TF pool. Monocytes constitutively express TF and its expression is enhanced after exposure to numerous inflammatory mediators (Cai et al., 2007; Cermak et al., 1993; Guha & Mackman, 2002; Lewis et al., 1995; Luyendyk et al., 2008; Mach et al., 1997). Indeed, elevated levels of TF have been measured in plasma during septic states and may account for the thrombotic complications encountered in those patients

(Osterud & Bjorklid, 2001; ten Cate et al., 2000). Endotoxin-induced TF expression is mediated via MAP kinases (Guha & Mackman, 2002; Lewis et al., 1995) and the consecutive activation of the transcription factors EGR-1, c-Fos/c-Jun, and nuclear factor kappa B. In addition, endotoxin also increases TF mRNA stability (Brand et al., 1991).

TF has also been described in circulating granulocytes. Eosinophils, neutrophils, and basophils have been found to express TF (Kambas et al., 2008; Maugeri, Brambilla et al., 2006; Maugeri et al., 2006; Moosbauer et al., 2007; Muller et al., 2003). Both the transcription and the translocation of TF to the cell surface are enhanced upon stimulation of these cells with granulocyte/macrophage colony-stimulating factor or platelet-activating factor (Moosbauer et al., 2007). These findings may explain the pro-thrombotic state associated with hypereosinophilia (Cugno et al., 2009).

Even though TF-containing platelets have been described by several groups (Falati et al., 2003; Muller et al., 2003; Panes et al., 2007; Perez-Pujol et al., 2005; Zillmann et al., 2001) the role of platelet derived TF remains debatable (Butenas et al., 2005). In platelets, TF is encountered in different compartments including the membrane, the matrix of α -granules, and the canalicular system; it is translocated to the cell surface upon activation by different agonists. Since TF messenger RNA is not found in human megakaryocytes, the source of platelet derived TF remains a matter of debate. One hypothesis suggests that TF containing microparticles derived from activated monocytes, leucocytes (Rauch et al., 2000), or endothelial cells (Aras et al., 2004; Camera et al., 2003) fuse with the platelet membrane through a receptor mediated process (Falati et al., 2003; Rauch et al., 2000) and thereby deliver TF into platelets. However, other studies demonstrated the presence of a spliced TF mRNA in activated human platelets confirming an increased TF protein synthesis and activity in ADP stimulated platelets (Camera et al., 2003).

TF containing microparticles have been well characterized in plasma. Besides endothelial and vascular smooth muscle cells (Abid Hussein et al., 2003; Lechner et al., 2007; Moosbauer et al., 2007; Schechter et al., 2000), monocytes and platelets are the main source of TF containing microparticles (Hron et al., 2007; Siddiqui et al., 2002). Despite studies demonstrating the role of microparticle derived TF in the thrombin generation *in vitro* (Sturk-Maquelin et al., 2003), doubts remain about its contribution to thrombus formation *in vivo* (Biro et al., 2003; Day et al., 2005).

The full-length form of TF mRNA contains 6 exons and splicing of exon 5 results in a frame-shift mutation generating a soluble alternatively spliced TF protein lacking the transmembrane domain (Bogdanov et al., 2003). Apart from plasma, alternatively spliced TF has been found in several tissues and cell types including the lung, placenta, vascular smooth muscle cells, and keratinocytes. Whereas it remains unclear whether alternatively spliced TF exerts pro-coagulant activity (Szotowski et al., 2005) more recent reports have revealed a role of alternatively spliced TF in angiogenesis by promoting endothelial cell migration and differentiation (Censarek et al., 2007; Y. He et al., 2008).

2.1.3. Cardiac TF

In contrast to skeletal myocytes, cardiac myocytes express the full length form of TF under basal conditions (Hartzell et al., 1989; Mackman et al., 1993). Studies performed in mice expressing low levels of TF revealed a potential role of TF in the maintenance of cardiac hemostasis, since these mice had increased cardiac hemorrhages and hemosiderin deposition ultimately leading to cardiac fibrosis and a reduction in left ventricular function as compared to wild type animals (Pawlinski et al., 2002). Phagocytosis of hematin derived from hemorrhaging erythrocytes is thought to play a key role in this process. Interestingly, mice lacking factor IX had

normal hearts, underlining the specific role of the TF:VIIa complex in hemostatic protection of the heart.

2.2. TF in cardiovascular disease

2.2.1. TF in thrombus formation

Although TF is critically involved in initiation of the coagulation cascade, questions regarding the role and contribution of the different TF pools in arterial thrombus formation remain open.

Under normal hemostatic conditions, TF is constitutively expressed in the vascular smooth muscle layer of the media and in fibroblasts present in the vessels as well as in the adventitia surrounding the vessels (Drake et al., 1989; Fleck et al., 1990). Hence, disruption of blood vessel integrity promotes the interaction between circulating FVII and the vascular TF pool leading to initiation of the clotting cascade (Fig. 2). Moreover, the release of inflammatory triggers following vascular injury or plaque rupture induces the expression of TF in endothelial cells as well as in circulating cells thereby potentiating a full coagulation response (Steffel et al., 2006). In contrast to the initiation phase of coagulation, the mechanisms promoting the propagation of clot growth are not as evident yet.

Blood-borne TF may contribute to the propagation of the clotting reaction. This consideration is based on the fact that the maintenance of coagulation requires a constant activity of TF as a cofactor and that the formed fibrin clot isolates and prevents the initially formed TF:VIIa complexes from reacting with circulating new inactive coagulation factors (Hathcock & Nemerson, 2004). This hypothesis is supported by the observation that TF is detected at very low levels in the blood unable to trigger a coagulant response whereas the TF concentration reaches functionally significant levels on the surface of a forming thrombus (Bogdanov et al.,

2003; Chou et al., 2004; Jesty & Beltrami, 2005). In addition, TF has been found embedded in murine (Day et al., 2005; Hoffman et al., 2006) and human (Wysokinski et al., 2004) thrombi. The role of blood-borne TF in coagulation is strengthened by an in vitro study demonstrating that the sub-picomolar TF concentrations found in blood cause an increase in thrombin formation under flow, but not under no-flow conditions (Okorie et al., 2008).

On the other hand, recent experiments performed in mathematical as well as whole blood models revealed different stages of TF requirement in the clotting process and showed that the propagation of coagulation is TF independent once thrombin has started to be formed (Orfeo et al., 2005). By abrogating TF activity with specific antibodies targeting either VIIa or TF, 3 phases in the clotting process could be distinguished: a first short (<10 seconds) period of absolute TF:VIIa dependence; a second period (10-240 seconds) of partial TF:VIIa dependence that decreases with the progress of the reaction, and a period which is TF:VIIa independent beginning after 2 minutes. In addition, this study revealed that the initial activation of procoagulant proteases by the TF:VIIa complex is sufficient to maintain a pool of activated procoagulant catalysts able to maintain a sustained coagulation response in a TF independent manner.

Since venous thrombosis is not always associated with vessel damage, circulating TF may play an important role in this pathology. In a model of jugular vein thrombosis inhibition of TF reduced thrombus growth in the uninjured vein (Himber et al., 2003). However, in another study, hematopoietic cell-derived TF did not affect venous thrombosis in an inferior vena cava thrombosis model in the mouse (Day et al., 2005). Hence, the exact role of the different TF pools and their contributions to the propagation of clot formation remains controversial and further studies are definitely required.

2.2.2. TF in inflammation and atherosclerosis

Inflammation is tightly linked to the development of atherosclerosis (Hansson, 2005). Elevated levels of inflammatory cytokines known to induce TF expression such as TNF- α and interleukins are observed in early atherosclerotic lesions. In the early stages of atherogenesis, TF mRNA and antigen is expressed in macrophages (Thiruvikraman et al., 1996; Wilcox et al., 1989), and as the lesions progress, TF is increasingly expressed in other cell types such as endothelial cells and smooth muscle cells (Fig. 2). Within the necrotic core (Marmur et al., 1996), it is predominantly associated with TF containing microparticles released by foam cells, monocytes, lymphocytes, and smooth muscle cells; in addition, it is detected in the extracellular space (Mallat et al., 2000). During plaque rupture, the content of the core comes in contact with the circulating blood and TF present within the plaque as well as expressed by activated endothelial and smooth muscle cells triggers thrombus formation leading to an acute vascular event. Indeed, higher levels of TF are measured in atherosclerotic plaques of patients suffering from an acute coronary syndrome as compared to patients exhibiting stable angina (Annex et al., 1995).

Recent studies implicate that TF and TF:VIIa signaling may also play a role in the pathogenesis of atherosclerosis. The TF:VIIa complex has been shown to be a strong chemotactic stimulus for vascular smooth muscle cells (Sato et al., 1996). Inhibition of TF:VIIa by overexpression of tissue factor pathway inhibitor leads to a consecutive reduction of vascular smooth muscle cell migration and impairs vascular remodeling in the mouse in vivo (Sato et al., 1999; Singh et al., 2001). Since TF:VIIa mediated cell signalling induces a number of gene products involved in cell proliferation and migration, it is very likely that similar effects occur in TF expressing cells within atherosclerotic lesions. TF:VIIa signalling indeed enhances the expression of the

connective tissue growth factors CCN1 and CCN2 in fibroblasts and vascular smooth muscle cells. Similar to TF, CCN1 and CCN2 mRNA levels are highly expressed in atherosclerotic lesions of apoE^{-/-} mice and in human atherosclerotic plaques (Oemar & Luscher, 1997; Oemar et al., 1997). Hence, it is conceivable that by inducing the expression of CCN1 and CCN2 and possibly other mediators as well, TF:Vlla promotes plaque progression via enhanced extracellular matrix accumulation and vascular smooth muscle cell proliferation. In addition, a role of CCN1 and CCN2 in monocyte and platelet adhesion to the subendothelial matrix after endothelial injury has also been described (Jedsadayamata et al., 1999; Schober et al., 2002).

In response to inflammatory mediators, TF transcription can be induced in different cell types. On the other hand, TF expressed on these cells appears to regulate the inflammatory response. During sepsis, the TF pathway induces a lethal inflammatory exacerbation independent of thrombin generation and clot formation (Creasey et al., 1993; Taylor et al., 1991). Inhibition of the TF:Vlla complex is paralleled by a decrease in interleukin secretion and a reduction in inflammatory infiltrations in several tissue during sepsis (Miller et al., 2002). This finding is confirmed by the observation that mice suffering from sepsis and expressing low TF levels display reduced interleukin levels and an increased survival compared with control mice (Pawlinski et al., 2004). Studies performed in PAR2 deficient mice suggest that the pro-inflammatory TF:Vlla signaling response is likely mediated via PAR2 (Pawlinski et al., 2004). However it is worth noting that deficiency in PAR1 or PAR2 alone has no effect and that a reduction in inflammation and mortality is only observed in the context of a combined inhibition of thrombin generation and PAR2 receptor, indicating a tight link between coagulation and inflammation.

2.2.3. TF in acute coronary syndromes

High levels of TF have been detected in coronary atherectomy specimens from patients with unstable angina or myocardial infarction (Ardissino et al., 1997). In line with this, plasma TF levels are elevated in patients with acute coronary syndromes compared with controls (Misumi et al., 1998; Soejima et al., 1996) and correlate with an unfavourable outcome in these patients (Soejima et al., 1999). In addition, patients with unstable angina or non-ST-elevation myocardial infarction and a high TIMI score (≥ 4) exhibit increased TF plasma levels as compared to those with a low TIMI score (< 3) (Lee et al., 2005).

Percutaneous transcatheter angioplasty and coronary stent implantation has become the gold standard in the treatment of culprit lesions in acute coronary syndromes. The local inflammatory reaction following balloon dilatation and stent implantation (Brunetti et al., 2007) induces the release of TF from the dissected plaque. Thus, elevated levels of TF positively correlate with the incidence of restenosis after percutaneous interventions, presumably due to promigratory effects of TF on vascular smooth muscle cells (Tutar et al., 2003). Despite the reduction in restenosis following deployment of drug eluting stents (DES) when compared to bare metal stents, stent thrombosis rates have not been decreased with the use of DES and may even be higher (Iakovou et al., 2005). One possible explanation may be that the drugs eluting from the stents which are intended to inhibit restenosis by reduction of vascular smooth muscle cell migration and proliferation exert unwanted biological effects on these or other cell types in the vasculature, in particular endothelial cells. The drugs applied for coating the first and second generation stents, like rapamycin, everolimus, or zotarolimus, enhance TF expression in vascular cells and accelerate arterial thrombus formation via inhibition of the mTOR/p70S6K pathway (Camici et al., 2010; Steffel et al., 2005). Similarly, paclitaxel, another widespread stent coating drug was shown to increase endothelial TF expression and activity through activation

of the MAP kinase JNK (Stahli et al., 2006). Hence, these findings may raise important issues in the context of drug eluting stent design since coating a stent with a drug exerting not only antiproliferative but also antithrombotic properties may offer new options in the interventional treatment of vascular disease.

2.2.4. TF and cardiovascular risk factors

Diabetes mellitus is associated with an increase in plasmatic TF activity (El-Ghoroury et al., 2008), TF antigen levels (Limet et al., 2004), and TF containing microparticles (Diamant et al., 2002). In line with these observations, improvement of glycemic control or application of oral antidiabetics such as the peroxisome-proliferator-activated receptor- γ agonist rosiglitazone downregulate TF protein expression (Golledge et al., 2007) in these patients. Consistent with this, high glucose increases TF antigen and activity in monocytes derived from healthy subjects (Stegenga et al., 2006; Vaidyula et al., 2006). A similar effect of high glucose on TF expression was also observed in human endothelial cells (Boeri et al., 1989). In diabetes, hyperglycaemia induces the formation of advanced glycation end-products (AGE), which upregulate TF expression in endothelial cells and monocytes via nuclear factor kappa B (Bierhaus et al., 1997; Bierhaus et al., 1997; Kislinger et al., 2001). Not surprisingly, higher levels of AGE and TF were observed in the vasculature of diabetic mice, and blockade of the receptor for AGE significantly reduced the increased TF expression in the aorta of these mice (Kislinger et al., 2001).

Patients with hyperlipidemia display an increased risk of developing thrombosis (Durrington, 2003). Elevated low density lipoprotein (LDL) levels are associated with an increased TF plasma activity (Sambola et al., 2003). Oxidized LDL triggers TF protein synthesis in monocytes (Wada et al., 1994), endothelial cells (Fei et al., 1993), and vascular smooth muscle cells (Penn et al., 1999). In contrast, high-density

lipoproteins (HDL) reduce the thrombin-induced TF expression in endothelial cells through PI3K pathway activation (Viswambharan et al., 2004). Moreover, 3-hydroxy-3-methylglutaryl-CoA reductase inhibitors (statins) reduce TF expression in monocytes (Markle et al., 2003), endothelial cells (Eto et al., 2002), and vascular smooth muscle cells (Brandes et al., 2003). These findings have been confirmed by studies performed in ApoE^{-/-} mice in which simvastatin and rosuvastatin treatment reduced TF antigen expression in atherosclerotic lesions independent of their lipid lowering effects (Bea et al., 2003; Monetti et al., 2007). Fibrates, which are also in use as lipid lowering drugs in the treatment of hyperlipidemia, also exert pleiotropic effects on vascular TF expression beyond their lipid lowering properties (Marx et al., 2001).

In contrast to normotensive subjects, hypertensive patients display increased plasma levels of TF antigen (Felmeden et al., 2003), which can be lowered by an appropriate antihypertensive management (Koh et al., 2004). Several mechanisms are thought to mediate this enhanced TF expression observed in arterial hypertension. For instance, the chronic exposure to shear stress induces the activation of transcription factors and consequently the expression of TF in endothelial cells (Dielis et al., 2005). On the other hand, angiotensin II increases TF expression in monocytes (M. He et al., 2006), endothelial cells (Dielis et al., 2005), and vascular smooth muscle cells via the angiotensin II type 1 receptor. Consistent with this observation, ACE inhibitors and angiotensin II type 1 receptor antagonists reduce TF plasma activity in hypertensive patients and inhibit endotoxin-induced TF expression in monocytes (Napoleone et al., 2000).

3. Therapeutic implications

The fate of a forming thrombus is determined by the balance of numerous factors, including vessel wall, coagulation system, and aggregating platelets. In addition to the classical antithrombotic agents, drugs targeting TF and the TF:FVIIa complex have been developed, since targeting the initiation phase of thrombosis may prevent thrombus formation more efficiently. Considering the non-hemostatic properties of the TF:FVIIa complex, inhibition of TF may also interfere with the pro-migratory and pro-proliferative effects of TF.

Despite promising experimental studies suggesting an effective reduction in thrombosis with a lower bleeding risk as compared to other coagulation inhibitors such as heparin or direct thrombin inhibitors, approaches to inhibit TF not yet been applied successfully in humans, and further large scale clinical trials as well as novel concepts of delivery are required.

3.1. Inhibition of TF production

Inhibition of TF production can be achieved by genetic and pharmacological intervention at both the transcriptional and the post-transcriptional level (Fig. 3). Hairpin ribozymes degrade TF mRNA and decrease TF induction in vascular smooth muscle cells (Cavusoglu et al., 2002). Antisense oligonucleotides hybridize to the complementary TF mRNA sequence and thereby inhibit translation of the TF protein (Stephens & Rivers, 1997). Dimethyl sulfoxide, used for long-term storage of haematopoietic stem cells, inhibits arterial thrombus formation by impairing TF activity (Camici et al., 2006). Further, cardiac glycosides such as digoxin and ouabain inhibit TF translation by lowering the intracellular potassium concentration through impaired Na^+/K^+ -ATPase activity (Stahli et al., 2007). Even more interesting, the

class III anti-arrhythmic drug amiodarone inhibits the translation of the TF protein in human vascular cells and impairs arterial TF expression as well as arterial thrombus formation in mice in vivo (Breitenstein et al., 2008). These pleiotropic effects of amiodarone may at least in part account for the beneficial action of the drug on prognosis of patients with heart failure and coronary artery disease (Breitenstein et al., 2008). More recently, the phytosterole guggulsterone as well as the green tea polyphenol epigallocatechin-3-gallate, at concentrations detected in plasma after oral ingestion, were observed to inhibit TF induction through inhibition of MAP kinase activation in human vascular cells and to reduce TF expression in the carotid artery of mice in vivo (Gebhard et al., 2009; Holy, Stampfli et al., 2009). These studies demonstrate that both synthetic and naturally occurring drugs can inhibit the induction of TF in human vascular cells as well as thrombus formation in vivo. Such effects may at least in part account for the established beneficial actions of the above mentioned drugs on the prognosis of patients with cardiovascular disease. Hence, clinical trials assessing the application of these drugs in the context of TF inhibition and thrombus prevention should be performed. Ideally, the drugs would have to be applied in a targeted local manner in order to achieve a maximal effect in the vessel territory at risk for thrombosis while reducing systemic bleeding complications.

3.2. Anti-TF antibodies

Administration of anti-TF antibodies (Fig. 3) delays thrombus formation in a rabbit carotid thrombosis model and reduces the infarction area following ligation of coronary arteries (Erlich et al., 2000; Ragni et al., 1996). A clinical trial (PROXIMATE-TIMI 27) performed in 26 subjects suffering of stable coronary disease tested the impact of a chimeric monoclonal antibody targeting human TF at the factor X binding site (Sunol-cH36/ALT836) (Morrow et al., 2005). Preliminary results

showed a dose dependent anticoagulant effect with only minor bleeding. Application of Sunol-cH36/ALT836 after endarterectomy in chimpanzees reduced local thrombus formation and improved vessel patency after 30 days without increasing bleeding time or surgical blood loss, indicating that early inhibition of coagulation with TF inhibitory antibodies reduces arterial thrombus formation effectively after vascular injury (Jiao et al., 2010).

Considering the role of TF signalling in tumor angiogenesis, growth, and metastasis, inhibition of TF may also translate into anti-neoplastic effects. Indeed, administration of a monoclonal anti-TF antibody (CNTO 859) reduced breast cancer metastasis and tumor growth in xenograft models (Ngo et al., 2007).

3.3. Recombinant tissue factor pathway inhibitor (rTFPI)

TFPI is the physiological inhibitor of TF activity and the balance between TF and TFPI is critical for thrombus formation (Pedersen et al., 2005). By binding the active site of FXa, TFPI interferes with the activity of the TF:VIIa complex (Fig. 3). Administration of rTFPI inhibits thrombus formation at the site of balloon-induced arterial injury and reduces plaque thrombogenicity in humans (Badimon et al., 1999; Harker et al., 1996; St Pierre et al., 1999). In addition, local adenoviral overexpression of TFPI in arteries increases the resistance to thrombus formation (Nishida et al., 1999) and reduces intimal hyperplasia in injured vessels (Zoldhelyi et al., 2001).

Despite of these positive actions, application of rTFPI may be limited because of its apoptotic effects observed in cultured endothelial and vascular smooth muscle cells, which may contribute to instability of atherosclerotic plaques (Fu et al., 2008; Hamuro et al., 1998). Moreover, a randomized controlled study including 1754 patients with severe sepsis and high international normal ratio (INR) found no effect on all-cause

mortality (Abraham et al., 2003). Therefore, additional trials are needed with patients exhibiting cardiovascular diseases other than severe sepsis and being in a better general condition.

3.4. Nematode anticoagulant protein (NAPc2)

Recombinant nematode anticoagulant protein c2 (rNAPc2), isolated from the saliva of the haematophagous hookworm *Ancylostoma caninum* is a factor Xa dependent small protein inhibitor of the TF:VIIa complex which binds to a site on factor Xa that is distinct from the catalytic centre (exo-site) (Fig. 3; Bergum et al., 2001). In a double blind placebo-controlled trial, rNAPc2 was found to be safe and well-tolerated at doses ranging from 0.3 to 5 µg/kg body weight in preventing thrombin generation during coronary angioplasty in combination with aspirin, clopidogrel, and heparin (Moons et al., 2003). A more recent phase II clinical study (ANTHEM-TIMI-32) demonstrated that in patients with non ST-elevation acute coronary syndrome managed with standard antithrombotics and an early invasive approach, additional proximal inhibition of the coagulation cascade with rNAPc2 was well tolerated and not associated with an increase in major or minor bleedings as compared with a placebo treatment (Fluture et al., 2007; Giugliano et al., 2007). A novel finding in the ANTHEM-TIMI-32 trial was that the doses of rNAPc2 suppressing thrombin generation through 48 h were associated with a ≥50% reduction in ischemia on continuous ECG. This provides a link between the mechanism of action of this drug and the hypothesized benefit in patients with plaque rupture and vessel wall injury. However, the trial was not powered for such an assessment and larger trials are required to confirm the hypothesis that rNAPc2 reduces clinical events. The fact that rNAPc2 has a half-life of 50 to 60 hours and can be administered subcutaneously,

prolonged inhibition of thrombin generation by rNAPc2 for several weeks may have the potential to improve the recommended discharge regimen after ACS. However, immune responses after administration may limit its clinical application, since rNAPc2 is a non-human protein. Anti-rNAPc2 immunoglobulins were indeed observed in one subject out of 20 patients receiving rNAPc2. Although these antibodies were not biologically active and without clinical relevance, the fact that they occurred in a high percentage of patients will have to be considered in future trials.

4. Conclusions

Major progress has been achieved in decrypting the biological effects and clinical relevance of TF in the context of cardiovascular disease. TF is a transmembrane glycoprotein, the main trigger of coagulation and, beyond this, an intracellular signal transducer. Several signalling pathways, specific for both the cell type and the stimulus, regulate TF induction and its cellular distribution. The discovery of blood-borne TF has launched a new debate about its *in vivo* relevance, since the relative contribution of vessel-wall as compared to blood-borne TF for thrombus formation and propagation remains unclear.

Elevated levels of TF activity are present in patients with cardiovascular risk factors and in subjects suffering of coronary artery disease. TF expression is up-regulated in atherosclerotic plaques, and cellular TF as well as extracellular TF contained in microparticles come in contact with the blood during endothelial erosion or plaque rupture. Therefore, TF contributes critically to the development of acute vascular events such as myocardial infarction or stroke. On the other hand, TF may promote the progression of atherosclerosis by enhancing vascular smooth muscle cell migration and proliferation.

Since targeting the initiation phase of the clotting cascade may be more effective in preventing thrombus formation than at later stages, several strategies have been developed to inhibit TF expression or activity. Despite promising experimental results, the clinical benefit of these new approaches in comparison with current antithrombotic regimens remains to be demonstrated in large scale trials. Since the major part of TF activity is cell-associated and induced by vascular inflammation, a targeted local inhibition of TF may well prove to be the most effective approach with a minimal effect on the systemic balance of thrombosis and bleeding.

Figure Legends

Figure 1:

TF exerts extracellular effects as main trigger of coagulation as well as intracellular effects on signal transduction.

Figure 2:

Expression of TF in different cell types during the pathogenesis of atherosclerotic lesions.

Figure 3:

Different approaches for inhibition of TF.

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