Recombinant full-length tissue factor pathway inhibitor fails to bind to the cell surface: implications for catabolism in vitro and in vivo

Guyu Ho, Masaaki Narita, George J. Broze Jr, and Alan L. Schwartz

Tissue factor pathway inhibitor (TFPI) plays a key role in the regulation of tissue factor-initiated blood coagulation secondary to loss of the integrity of the blood vessel wall. TFPI is a naturally occurring Kunitz-type protease inhibitor that inhibits coagulation factor Xa and, in a factor Xa-dependent manner, mediates feedback inhibition of the factor VIIa/tissue factor catalytic complex. In vivo full-length TFPI is thought to be primarily bound to the vascular endothelium and the high affinity binding requires an intact carboxy terminus. Here we describe a full-length TFPI molecule, expressed in mouse C127 cells (TFPI\(^{127}\)), which exhibits virtually no cellular binding yet contains the intact carboxy terminus. This TFPI (TFPI\(^{127}\)) is neither internalized nor degraded via the TFPI endocytic receptor, LDL-receptor-related protein. Pharmacokinetic studies of TFPI\(^{127}\) in vivo demonstrate a 10-fold prolongation in the plasma half-life, compared with that of bacterial recombinant TFPI. (Blood. 2000;95:1973-1978)

© 2000 by The American Society of Hematology

Materials and methods

Materials

Iodogen was purchased from Pierce (Rockville, IL). \(^{125}\)I-Iodide was from Amersham Company (Buckinghamshire, UK). Human Factor Xa was from Enzyme Research (South Bend, IN) and Spectrozyme Xa was from American Diagnostica, Inc (Greenwich, CT). Bovine serum albumin was purchased from Calbiochem-Novabiochem. Affi-gel 10 was from Bio-rad (Hercules, CA). Tissue culture media and plasticware were obtained from Life Technologies, Inc (Rockville, MD). Recombinant TFPI from Escherichia coli was provided by Monsanto Company (St Louis, MO). CHO, SK hepatoma, and 293 cell-derived TFPI, was purified as described below.

Purification of TFPI\(^{127}\).—TFPI\(^{127}\) was affinity-purified from the conditioned media of mouse C127 cells that had been transfected with a bovine papilloma virus containing wild-type human TFPI cDNA. In brief, cells producing TFPI\(^{127}\) were cultured in serum-free media containing aprotinin (10 µg/mL; Sigma) and the media were replaced every 2 to 3 days. Harvested media were concentrated using an Amicon ultraconcentrator (YM 30; Amicon, Bedford, MA) and the concentrates were absorbed to a 2H8 monoclonal anti-TFPI-agarigel column (0.5 \(\times\) 3 cm) and eluted with 100 mmol/L glycine, pH 2.2. The pH of the eluant was adjusted to pH 7.5 using 1 mol/L Tris-HCl, pH 8.0, and applied to a 1 mL HiTrap...
heparin-agarose column (Pharmacia-LKB Biotechnology, Piscataway, NJ), which had been equilibrated in 0.1 mol/L NaCl 0.05 mol/L Tris-HCl, pH 7.5. The column was developed with a linear NaCl gradient from 0.1 to 1.0 mol, at 0.5 mL/min over 10 column volumes. Fractions were assayed for the presence of TFPI by a modified enzyme linked immunosorbent assay (ELISA) method using 2 independent monoclonal antibodies, 2H8 and 2B12.5 The desired fractions were pooled, concentrated using Centricon microconcentrators (Amicon), and stored at −80°C. The protein concentration was determined using the modified ELISA method with TFPI of known concentration as standard.

**Amidolytic assays of factor Xa activity**

In a 96-well microtiter plate, 1 nmol/L of factor Xa was incubated with various concentrations of TFPI (0.3-300 pmol) in 135 μL of 0.1 mol/L NaCl, 0.05 mol/L Tris-HCl, 0.5% bovine serum albumin (BSA), pH 7.5 (TBSA) at room temperature for 30 minutes. Fifteen microliters of 1 mmol/L Spectrozyme Xa chromagenic substrate was then introduced (final concentration 100 μmol/L) and the change in absorbance at 405 nm was determined 10 minutes thereafter.

**Tissue factor pathway inhibitor functional assay of Vila/TNF inhibition**

The end point assay of anti-VIIa/TNF activity (American Diagnostica) was used to compare TFPI C127 and bacterially expressed TFPI at equimolar concentrations.

**Factor Xa-induced coagulation of plasma**

In a fibrometer (BBL, Cockeysville, MD), 50 μL of rabbit brain cephalin prepared as described by the manufacturer (Sigma), 50 μL of CaCl2 (25 mmol/L), 50 μL of various amounts of TFPI C127 (25 ng to 300 ng) in TBSA, and 50 μL of human factor Xa (0.2 nmol/L) were incubated at 37°C. After 30 seconds, 50 μL of normal human plasma (George King Biochemical) were added, and the degree of apparent factor Xa inhibition was determined by comparing the clotting to a factor Xa standard curve.

**Cell culture**

Human hepatoma HepG2 cells, mouse fibroblast PEA 13 cells,19 and mouse brain endothelial cells (bend-3)20 were cultured in Dulbecco’s modified Eagle’s medium (DMEM), supplemented with 10% fetal calf serum, 100 U/mL of penicillin, and 100 μg/mL of streptomycin. Cells were incubated at 37°C in humidified air containing 5% CO2.

**Protein iodination**

Proteins (5-25 μg) were iodinated using the IODOGEN method.21 Specific radioactivities were typically 1 to 10 × 103 cpm/pmol of protein.

**Binding and degradation assays**

Binding of 125I-TFPI to cells was performed in suspension. Cells were displaced from Petri-dishes by incubation with 3 mmol/L EDTA in phosphate-buffered saline (PBS) at room temperature for 5 minutes. Cells were then pelleted at low speed (1000 rpm × 5 minutes) on a table top centrifuge (Sorvall RT6000B) to remove EDTA and resuspended in the assay buffer (DMEM/3% BSA). 4 × 105 cells in a volume of 300 μL were added with 2 nmol/L of 125I-TFPI in the presence or absence of 100 mol excess of cold ligands. After rocking at 4°C for 2 hours, cell suspensions were gently layered on 900 μL of fetal calf serum in microfuge tubes and spun at 14 000 rpm × 1 minute in a table-top microfuge. The cell pellet was counted in a gamma counter (Cobra II auto-gamma, Packard Instrument, Meriden, CT). Nonspecific binding was determined in the presence of excess unlabeled ligand as specified in the figure legends.

Degradation assays were performed by washing cell monolayers (in 12-well dishes) twice with the assay buffer. 0.5 mL of assay buffer containing 2 nmol/L 125I-TFPI was then added to each well. After incubation at 37°C for the indicated periods, the overlying medium was removed and proteins were precipitated by the addition of BSA to 5 mg/mL and trichloroacetic acid to 20%. Degradation of ligand was defined as the appearance of radioactive fragments in the overlying medium that were soluble in trichloroacetic acid. Fragmentation of 125I-TFPI at 37°C in cell-free wells was subtracted from the corresponding samples.

For ligand competition binding and degradation assays (Table 2), HepG2 and CHO cells were assayed as monolayers as described previously7,18; 125I-TFPI was at 20 nmol/L for the binding assays and at 0.6 nmol/L for the degradation assays.

**Plasma clearance of 125I tissue factor pathway inhibitors in mice**

Twelve- to 16-week-old BALB/c mice (weighing 20-25 g) were anesthetized with sodium pentobarbital (1 mg/20 g of body weight) during the course of experiments. Approximately 15 pmol of radiolabeled protein alone or with unlabeled 39 kd protein in 100 μL of sterile saline was injected into the tail vein over 30 seconds. In studies in which heparin was administered, heparin was injected 20 minutes after 125I-TFPI administration. At the indicated times, 40 to 50 μL of blood was collected by periorbital bleeding. The blood samples were centrifuged and 15 μL of the plasma fractions was spotted onto 3 MM filter paper (Whatman Inc, Clifton, NJ), precipitated with 10% trichloroacetic acid, rinsed with ethanol, and radioactivity determined. The initial plasma concentration (time = 0) of 125I-TFPI was extrapolated from the measured points. At the end of each experiment, animals were killed and the liver, kidney, spleen, and lungs were removed, blotted, weighed, and the radioactivity determined.

**Metabolic incorporation of 35SO4 and immunoprecipitation of TFPI C127**

One 75-cm2 flask of C127 cells was labeled for 96 hours with 37 MBq (1 mCi) of Na3[35S]SO4 (Amersham Corp) in 10 mL of sulfate-deficient DMEM (-SO4) supplemented with 2% dialyzed fetal calf serum, in the absence of G418. The culture medium was then concentrated using Centricon microconcentrators (Amicon). Monoclonal antibody 2H8-conjugated affigel-10 beads were added to the concentrates that were then rocked overnight to allow for binding. The beads were subsequently collected by centrifugation, washed 3 times with PBS, boiled in SDS-sample buffer (to release bound TFPI), and the bead-free portion was analyzed by SDS-PAGE.

**Electrophoresis and Western blotting**

SDS-PAGE was performed using 10% separating and 5% stacking gels and proteins were transferred to nitrocellulose membrane. The membrane was blocked with 5% nonfat milk for 30 minutes, followed by incubation with a 1:100 dilution of rabbit antiserum raised against a synthetic peptide matching the carboxy terminal 12 amino acid residues of the mature TFPI protein.5 The immunoreactivity was detected using ECL reagents (Amersham Corp).

**Results**

**Purification of recombinant tissue factor pathway inhibitor from mouse C127 cells**

Previous studies have shown that TFPI, when expressed in mammalian systems, is susceptible to proteolysis at its carboxy terminus.1,2,2 Thus, mouse C127 cells expressing TFPI were grown in media containing the protease inhibitor, aprotinin. TFPI was purified from the media via TFPI monoclonal antibody-affinity chromatography. This was followed by heparin-agarose affinity which separates the full-length TFPI from the truncated species by charge interactions between the heparin and the positively charged carboxy terminus of TFPI. Figure 1 shows the elution of TFPI C127 from the heparin column with a linear NaCl gradient. TFPI C127...
eluted with 2 peaks at NaCl concentrations of 0.4 and 0.6 mol/L, respectively. Because full-length TFPI elutes at 0.6 mol/L salt, only the fractions collected at or greater than 0.6 mol/L NaCl (fractions 12-15) were pooled for subsequent studies. To confirm that the purified TFPI C127 was full length, TFPI C127 was subjected to Western blot analysis using antibodies directed against the last 12 amino acid residues of TFPI. As shown in the inset of Figure 1, the TFPI was recognized by these antibodies, confirming that the carboxy terminus was intact.

The activity of TFPI C127 was assessed in a direct amidolytic assay of factor Xa activity, as well as an end point assay of anti-VIIa/TF activity. As shown in Figure 2, TFPI C127 was able to inhibit factor Xa activity in a concentration-dependent manner with complete inhibition of factor Xa at approximately 3 nmol/L, which was essentially identical to that of bacterial recombinant TFPI. Using a global anticoagulant assay, TFPI C127 inhibited factor Xa-induced plasma coagulation with IC50 approximately 10 nmol/L, compared with approximately 4 nmol/L for the bacterial recombinant TFPI. The disparity in IC50 values observed here is consistent with the previous observations that TFPI expressed in prokaryotic systems exhibits higher anticoagulant activities than that expressed in eukaryotic systems.

Table 1. Binding of 125I-TFPI C127 to various cell lines

<table>
<thead>
<tr>
<th>Cell Line</th>
<th>Specific Binding (fmol/10^6 cells)</th>
</tr>
</thead>
<tbody>
<tr>
<td>HepG2</td>
<td>Not detectable</td>
</tr>
<tr>
<td>Bend</td>
<td>Not detectable</td>
</tr>
<tr>
<td>PEA 13</td>
<td>Not detectable</td>
</tr>
</tbody>
</table>

HepG2, Bend, and PEA 13 cells were incubated in suspension with 2 nmol 125I-TFPI C127 in the presence or absence of 100 mol excess of cold TFPI C127 for 2 hours at 4°C. The specific radioactivity associated with cells was determined (open bars). Binding of bacterial recombinant 125I-TFPI to these cells was performed in the identical manner and used as control. Values represent the means of duplicate determinations.

Cellular binding and degradation of tissue factor pathway inhibitor in C127 cells

We have previously shown that bacterial recombinant TFPI binds to human hepatoma HepG2 cells with high affinity and that truncation of sequences distal to the second Kunitz domain abolishes the binding. Surprisingly, the full-length 125I-TFPI C127 exhibited no binding to hepatoma HepG2 cells, or to mouse vascular endothelial Bend cells, or mouse fibroblast PEA 13 cells (Table 1). Bacterial recombinant 125I-TFPI, however, exhibited abundant binding to these cells (Table 1). Furthermore, 125I-TFPI C127 was not taken up/degraded by HepG2 cells in contrast to anti-VIIa/TF activity. As shown in Figure 2, TFPI C127 was able to inhibit factor Xa activity in a concentration-dependent manner with complete inhibition of factor Xa at approximately 3 nmol/L, which was essentially identical to that of bacterial recombinant TFPI. Using a global anticoagulant assay, TFPI C127 inhibited factor Xa-induced plasma coagulation with IC50 approximately 10 nmol/L, compared with approximately 4 nmol/L for the bacterial recombinant TFPI. The disparity in IC50 values observed here is consistent with the previous observations that TFPI expressed in prokaryotic systems exhibits higher anticoagulant activities than that expressed in eukaryotic systems.

Figure 1. Profile of heparin-agarose chromatography of TFPI C127. Heparin-agarose chromatography was carried out using a 0.1 to 1 mol/L NaCl gradient in 0.05 mol/L Tris-HCl, pH 7.5. Elution of TFPI C127 was monitored by a modified ELISA as described in the “Materials and Methods.” Inset: Western blot analysis of TFPI C127. Purified TFPI C127 (1 µg) was run on a 10% SDS-PAGE gel and transferred to nitrocellulose membrane. The membrane was blotted with rabbit antiserum containing a peptide antibody against the last 12 amino acid residues of TFPI, and developed with ECL reagents.

Figure 2. Inhibition of factor Xa amidolytic activity by TFPI C127. Factor Xa (1 nmol) was incubated with increasing concentrations of TFPI C127 (0-3 nmol) at room temperature for 30 minutes. The remaining factor Xa activity was assessed using the chromogenic substrate Spectrozyme Xa. Bacterial recombinant TFPI (bTFPI) was assayed in the identical manner and used as control.

Figure 3. Degradation of 125I-TFPI C127 by HepG2 cells. HepG2 cells growing in petri dishes were incubated with 0.6 nmol of 125I-TFPI C127 at 37°C for the indicated times. At each time point, the overlaying medium was removed and proteins were precipitated by trichloroacetic acid (TCA). The TCA-soluble radioactivity (ie, degradation products) was determined. Degradation of bacterial recombinant 125I-TFPI by HepG2 cells was performed in the identical manner and served as a control. Numbers represent the means of duplicate determinations. Total degraded 125I-TFPI at 4 hours was less than 10% input.
protein, a molecule known to inhibit all ligand interactions with LRP,
prolongs the beta-phase elimination of bacterial recombinant TFPI. In addition, postinjection of 100 units of heparin raises the plasma 125I-TFPI level from approximately 5% of the initial plasma concentration to approximately 100%. When 50 mg/kg of 39-kd protein was coadministered with 125I-TFPI to mice, 125I-TFPI clearance was unaltered (Figure 4A). Postinjection of 100 units heparin after 125I-TFPI administration increased the plasma 125I-TFPI concentration only minimally (Figure 4B). These data suggest that neither LRP nor heparin-releasable binding to the vascular endothelium plays a significant role in TFPI clearance in vivo.

Tissue factor pathway inhibitor sulfation

Binding of TFPI to the cell surface is primarily due to charge interactions between the positively charged carboxy terminus and negatively charged cell surface heparan sulfate proteoglycans (HSPGs). One possibility to account for the inability of TFPI to bind to the cell surface is that the carboxy terminus may interact with the negatively charged clusters within TFPI itself such that the carboxy terminus is no longer available for interaction with cell surface HSPGs. It has been reported that TFPI expressed in human kidney 293 cells (TFPI293), but not that expressed in Chinese hamster ovary cells (TFPICHO) or hepatoma cells (TFPIhepatoma), contains the negatively charged sulfate. To determine the relationship between TFPI sulfation and its binding property, TFPIs expressed in 293, CHO, and hepatoma cells were examined for cellular binding. As shown in Table 3, sulfated TFPI293 was unable to bind to fibroblast PEA 13 cells, whereas nonsulfated TFPICHO and TFPIhepatoma both exhibited abundant binding.

To examine whether TFPI293 was indeed sulfated, C127 cells were metabolically labeled with 35SO4. TFPI293 was immuno precipitated from the media and analyzed by SDS-PAGE, followed by either autoradiography or Western blot. As shown in Figure 5, a 35SO4-labeled band migrated at the identical position as TFPI293 and was recognized by a TFPI monoclonal antibody, demonstrating sulfate incorporation into TFPI293.

Discussion

Recombinant human TFPI has been studied extensively in terms of its biochemical and biologic properties, as well as its pharmacokinetics. The highly basic carboxyl terminus is necessary for efficient anticoagulant function, yet confers on TFPI rapid clearance in vivo. Herein we describe a TFPI molecule with an intact carboxy terminus that, nevertheless, exhibits an inability to bind to the cell surface or be internalized and degraded via the endocytic receptor LRP, and displays a markedly prolonged plasma half-life in vivo.

Table 3. Cellular binding of 125I-TFPI expressed in 293, hepatoma, and CHO cells

<table>
<thead>
<tr>
<th>Ligand</th>
<th>Sulfation 26</th>
<th>Specific Binding (fmol/10^6 cells)</th>
</tr>
</thead>
<tbody>
<tr>
<td>TFPI293</td>
<td>+</td>
<td>not detectable</td>
</tr>
<tr>
<td>TFPIhepatoma</td>
<td>−</td>
<td>510</td>
</tr>
<tr>
<td>TFPICHO</td>
<td>−</td>
<td>430</td>
</tr>
</tbody>
</table>

PEA 13 cells were incubated in suspension with 2 nM 125I-TFPI293, or 125I-TFPICHO, or 125I-TFPIhepatoma in the presence or absence of 100 mol excess of cold ligands at 4°C for 2 hours and the bound-specific radioactivity determined. Values represent the means of duplicate determinations.
were incubated with \( {\text{S}^{35}}\text{O}_4 \) for 96 hours, after which the culture medium was collected and immunoprecipitated with anti-TFPI as described under “Materials and Methods.”

Binding of full-length TFPI to the cell surface is dependent on its highly positively charged carboxyl terminus that interacts with the negatively charged cell surface HSPGs. We have previously shown that TFPI expressed in bacteria binds with high affinity to hepatoma HepG2 cells, fibroblast PEA13 cells, and microvascular endothelial cells. Similarly, TFPI expressed in hepatoma and CHO cells displays similar binding properties (Table 3). To our surprise, TFPI expressed in mouse C127 fibroblast cells was devoid of cell surface binding activity but remained biologically active as judged by the inhibition of (1) factor Xa amidolytic activity (Figure 2), of (2) factor Xa-induced plasma coagulation, and (3) factor VIIa/TF activity. One potential explanation for this loss of binding activity is that secondary to intramolecular interactions, TFPI’s carboxyl terminus is no longer available for interaction with cell surface HSPGs. It is of note that, although TFPI\(^{C127}\) does not bind to cell surface HSPGs, it binds to heparin-agarose (Figure 1) with the same affinity as bacterial recombinant TFPI (ie, both TFPI species are eluted from heparin-agarose at identical salt concentrations (0.6 mol/L NaCl) (6, data not shown). The fact that the affinity of TFPI is 40-fold greater for heparin than for HSPGs may account for the high affinity binding of both species of TFPI to heparin and their discordant binding to cell surface HSPGs.

Girard et al\(^{29}\) have shown that serine 2 of TFPI is phosphorylated after its expression in C127 cells. However, TFPI with a serine 2 to alanine mutation, when expressed in these cells, did not regain the binding properties of wild type TFPI (data not shown). This finding suggests that this negatively charged phosphate group (ie, at serine 2) does not interfere with availability of the carboxyl terminus. TFPI contains 3 potential N-linked and 1 O-linked glycosylation sites. Smith et al\(^{26}\) have shown that metabolic labeling of 293 cells with \( {\text{H}^{35}}\text{S}\text{O}_4 \) yielded sulfated TFPI, whereas sulfated TFPI was not produced by CHO cells. They further determined that TFPI\(^{293}\) carries Asn-linked sulfated oligosaccharides: SO\(_4\)-GalNAc\(_1\)\(2\)GlcNAc\(_1\)\(2\)Man. Because TFPI\(^{293}\) was unable to bind to the cell surface (Table 3), it is possible that sulfated oligosaccharides on TFPI may interact with its own positively charged carboxyl terminus, thus abrogating TFPI’s interaction with the cell surface. Attempts to quantitatively remove linked oligosaccharides or sulfate via incubation of nonadenatured TFPI\(^{C127}\) with N-glycanase, O-glycanase or sulfatase and assess biological function have proven unsuccessful. Thus, whether TFPI\(^{C127}\) possesses the same sulfation pattern as TFPI\(^{293}\) will require further study.

TFPI\(^{C127}\) exhibited a 10-fold increase in plasma half-life compared with bacterial recombinant TFPI (Figure 4A). Because the administration of heparin did not significantly increase the plasma \( {\text{I}^{125}}\text{TFPI}\) concentration, secondary to release from the vascular endothelium, or did the administration of the 39-kd protein inhibit its clearance (Figure 4), the prolonged half-life of \( {\text{I}^{125}}\text{TFPI}\) is likely due to its inability to bind to either cell surface HSPGs or LRP. These results are consistent with our in vitro findings (Figure 3 and Table 1). However, it is not clear at present whether this observation can be accounted for by alterations in protein sulfation alone, because \( {\text{I}^{125}}\text{TFPI}\) and \( {\text{I}^{125}}\text{TFPI}\) have similar rates of plasma clearance. Nonetheless, the clearance of \( {\text{I}^{125}}\text{TFPI}\), albeit markedly prolonged, did result in elimination of 50% at approximately 20 minutes. Therefore, additional mechanisms (eg, hepatic glycoprotein receptors) may be operative.

Under physiologic conditions, TFPI is primarily synthesized and secreted by vascular endothelial cells. Because the major pool of TFPI in vivo is associated with the endothelial cell surface, it appears that endothelial TFPI maintains the carboxy terminus available for interaction with cell surface HSPGs. Assuming that sulfation does play a role in TFPI binding to cells, TFPI produced by endothelial cells in vivo may well have to be under- or unsulfated to maintain its cellular binding. However, it is not presently known whether endogenous TFPI is sulfated or its degree of sulfation. Of note, human endothelial cells express significantly lower (less than 10%) GalNAc- and sulfotransferase activities than those expressed in 293 cells. These 2 enzymes are the key activities required for the addition of sulfated oligosaccharides to TFPI.\(^{26}\)

The physiologic significance of the cell-associated TFPI pool may relate to both its rapid mobilization into the plasma as well as its role as a mediator in the cellular uptake and degradation of factor Xa. In addition, it may serve to down-regulate the coagulant activity of factor VIIa/TF complex via transfer to caveolae. Endogenous TFPI within the circulation, however, is likely of less physiological importance, as it is largely truncated at various positions within the carboxy terminal domain and hence exhibits much reduced anticoagulant activity.\(^{5}\) The nonbinding full-length TFPI we have characterized in this study may therefore offer a therapeutic advantage when high plasma TFPI concentrations are required to effectively modulate TF-induced blood coagulation.
References


Recombinant full-length tissue factor pathway inhibitor fails to bind to the cell surface: implications for catabolism in vitro and in vivo

Guyu Ho, Masaaki Narita, George J. Broze, Jr and Alan L. Schwartz